Package 'MetaGxOvarian'

July 9, 2024

Type Package Title Transcriptomic Ovarian Cancer Datasets Version 1.25.0 Date `r Sys.date()` Description A collection of Ovarian Cancer Transcriptomic Datasets that are part of the MetaGxData package compendium. License Artistic-2.0 Depends Biobase, AnnotationHub, ExperimentHub, SummarizedExperiment, R (>= 3.6.0)Imports stats, lattice, impute Suggests testthat, xtable, rmarkdown, knitr, BiocStyle, markdown **Encoding** UTF-8 VignetteBuilder knitr NeedsCompilation no biocViews ExpressionData, ExperimentHub, CancerData, Homo_sapiens_Data, ArrayExpress, GEO, NCI, MicroarrayData, ExperimentData LazyData yes RoxygenNote 7.1.1 git_url https://git.bioconductor.org/packages/MetaGxOvarian git_branch devel git_last_commit 2043cd1 git_last_commit_date 2024-04-30 Repository Bioconductor 3.20 Date/Publication 2024-07-09 Author Michael Zon [aut], Vandana Sandhu [aut], Christopher Eeles [ctb], Benjamin Haibe-Kains [aut, cre]

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attention day

days_to_death

Description

This is a note to inform package users that the days_to_death variable is also valid for living pateints. In this case, the value in days_to_death is the number of days since the last follow-up appointment.

Format

A field included in various data files in the this package.

duplicates

a list containing the names of patients that are believed to be dulicates across datasets

Description

The object is a list where each element is a patient ID that is believed to be a duplicate of a patient in another dataset. Patients are designated as duplicated if they have Spearman correlations greater than or equal to 0.98 with other patient expression profiles

Format

A list with 130 elements, each of which is a patient ID.

E.MTAB.386	Angiogenic mRNA and microRNA gene expression signature predicts
	a novel subtype of serous ovarian cancer.

Description

Ovarian cancer is the fifth leading cause of cancer death for women in the U.S. and the seventh most fatal worldwide. Although ovarian cancer is notable for its initial sensitivity to platinum-based therapies, the vast majority of patients eventually develop recurrent cancer and succumb to increasingly platinum-resistant disease. Modern, targeted cancer drugs intervene in cell signaling, and identifying key disease mechanisms and pathways would greatly advance our treatment abilities. In order to shed light on the molecular diversity of ovarian cancer, we performed comprehensive transcriptional profiling on 129 advanced stage, high grade serous ovarian cancers. We implemented a, re-sampling based version of the ISIS class discovery algorithm (rISIS: robust ISIS) and applied it to the entire set of ovarian cancer transcriptional profiles. rISIS identified a previously undescribed patient stratification, further supported by micro-RNA expression profiles, and gene set enrichment analysis found strong biological support for the stratification by extracellular matrix, cell adhesion, and angiogenesis genes. The corresponding "angiogenesis signature" was validated in ten published independent ovarian cancer gene expression datasets and is significantly associated with overall survival. The subtypes we have defined are of potential translational interest as they may be relevant for identifying patients who may benefit from the addition of anti-angiogenic therapies that are now being tested in clinical trials.

Format

```
experimentData(eset):
Experiment data
Experimenter name: Bentink S, Haibe-Kains B, Risch T, Fan J-B, Hirsch MS, Holton
Laboratory: Bentink, Matulonis 2012
Contact information:
Title: Angiogenic mRNA and microRNA gene expression signature predicts a novel su
```

```
URL:
  PMIDs: 22348002
  Abstract: A 212 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      Illumina humanRef-8 v2.0 expression beadchip
   platform shorttitle:
      Illumina humanRef-8 v2.0
  platform_summary:
      illuminaHumanv2
   platform_manufacturer:
     Illumina
  platform_distribution:
      commercial
  platform_accession:
     GPL6104
   version:
      2015-09-22 19:06:44
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: ILMN_1343291 ILMN_1651228 ... ILMN_1815951 (12449
    total)
 varLabels: probeset gene EntrezGene.ID best_probe
```

Details

varMetadata: labelDescription

 1
 1
 1
 1
 1
 1
 1
 1

 DFCI.106
 DFCI.107
 DFCI.108
 DFCI.109
 DFCI.111
 DFCI.1110
 DFCI.1111
 DFCI.1112

 1
 1
 1
 1
 1
 1
 1
 1
 1

 DFCI.113
 DFCI.114
 DFCI.115
 DFCI.116
 DFCI.117
 DFCI.118
 DFCI.119
 DFCI.12

1	1	1	1	1	1	1	1
					DFCI.125	_	-
1	1	1	1	1	1	1	1
DFCI.128	DFCI.129		DFCI.130		DFCI.132	DFCI.14	DFCI.15
1	1	1	1	1	1	1	1
DFCI.16	DFCI.17	DFCI.18	DFCI.19	DFCI.2	DFCI.20	DFCI.21	DFCI.22
1 DFCI.23	1 DFCI.24	1 DFCI.25	1 DFCI.26	1 DFCI.27	1 DFCI.28	1 DFCI.29	1 DFCI.3
Drc1.23	DFC1.24 1	Drc1.23	Drc1.20	Drc1.27 1	Drc1.20	Drc1.29	Drc1.3 1
DFCI.30	DFCI.31	DFCI.32	DFCI.33	DFCI.34		DFCI.36	DFCI.37
1	1	1	1	1	1	1	1
DFCI.38	DFCI.39	DFCI.4	DFCI.40	DFCI.41	DFCI.42	DFCI.44	DFCI.45
1	1	1	1	1	1	1	1
DFCI.46	DFCI.47	DFCI.48	DFCI.49	DFCI.50	DFCI.51	DFCI.52	DFCI.53
1	1	1	1	1	1	1	1
DFCI.54	DFCI.55 1	DFCI.56 1	DFCI.57	DFCI.58 1	DFCI.59 1	DFCI.6 1	DFCI.60
1 DFCI.61	DFCI.62	DFCI.63	1 DFCI.64	DFCI.65	_	_	1 DFCI.68
Drc1.01 1	DFC1.02	Drc1.03	Drc1.04 1	Drc1.05	Drc1.00 1	Drc1.07	1 DFC1.00
DFCI.69	DFCI.7	DFCI.70	(Other)	Ŧ	±	±	±
1	1	1	30				
sample_ty tumor 129 histologi ser	-	:					
129							
primarysi ov	te:						
129							
summarygr high 129	ade:						
summaryst	age:						
early la							
1 1	.28						
tumorstag							
2 3							
1 109	ТЭ						
substage:							
a b		's					

5 12 93 19 age_at_initial_pathologic_diagnosis: Min. 1st Qu. Median Mean 3rd Qu. Max. 21.00 50.00 66.00 60.71 72.00 95.00 days_to_death: Min. 1st Qu. Median Mean 3rd Qu. Max. 3.9 516.9 917.1 1007.0 1401.0 2724.0 vital_status: deceased living 73 56 debulking: optimal suboptimal NA's 98 28 3 uncurated_author_metadata:

Source.Name: DFCI-100///Ch

Source.Name: DFCI-

Source.Name: DFCI-1

Source.Name: DFCI-103///

Source.Name: DFCI-104///0

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Source.Name: DFCI-107///0

Source.Name: DFCI-108///

Source.Name: DFCI-109///Ch

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Source.Name: DFCI-110/

Source.Name: DFCI-111///Ch

Source.Name: DFCI-112///

Source.Name: DFCI-113///

Source.Name: DFCI-11 Source.Name: DFCI-115///0 Source.Name: DFCI-116///Ch Source.Name: DFCI-117/ Source.Name: DFCI-118///Characteristics.Age.: Age <has_measurement <Measurement <ha Source.Name: DFCI-119/// Source.Name: DFCI-11/// Source.Name: DFCI-120///Characteristics.Age.: Age <has_measurement <Measurement Source.Name: DFCI-121/ Source.Name: DFCI-12 Source.Name: DFCI-123///0 Source.Name: DFCI-124/ Source.Name: DFCI-125/ Source.Name: DFCI-1 Source.Name: DFCI-127///Characteristics.Age.: Age <has_measurement <Measurement Source.Name: DFCI-128/ Source.Name: DFCI-129///Characteristics.Age.: Age <has_measurement <Measurement Source.Name: DFCI-12// Source.Name: DFCI-130///Characteristics.Age.: Age <has_measurement <Measurement <ha Source.Name: DFCI-131///Characteristics.Age.: Age <has_measurement <Measurement Source.Name: DFCI-132///Characteristics.Age.: Age <has_measurement <Measurement <h Source.Name: DFCI-13/ Source.Name: DFCI-14, Source.Name: DFCI-

E.MTAB.386

Source.Name: DFCI Source.Name: DFCI-17/, Source.Name: DFCI-18// Source.Name: DFCI-19// Source.Name: DH Source.Name: DFCI-20/ Source.Name: DFCI-Source.Name: DFCI-22///Characteristics.Age.: Age <has_measurement <Measurement Source.Name: DFCI-23/// Source.Name: DFCI-24///Ch Source.Name: DFCI-25/// Source.Name: DFCI-20 Source.Name: DFCI-27/ Source.Name: DFCI-2 Source.Name: DFCI-29/ Source.Name: DFCI-2 Source.Name: DFCI-30 Source.Name: DFCI-31/, Source.Name: DFCI-32 Source.Name: DFCI-33, Source.Name: DFCI-34, Source.Name: DFCI-35/ Source.Name: DFCI-

E.MTAB.386

- Source.Name: DFCI-38/// Source.Name: DFCI-39/// Source.Name: DFCI-Source.Name: DFCI-40/ Source.Name: DFCI-41, Source.Name: DFCI-42/ Source.Name: DFCI-44, Source.Name: DFCI-Source.Name: DFCI-46/ Source.Name: DFCI-47, Source.Name: DFCI-Source.Name: DFCI-49 Source.Name: DFCI-Source.Name: DFCI-50, Source.Name: DFCI-51/// Source.Name: DFCI-52/ Source.Name: DFCI-53//, Source.Name: DFCI-54/// Source.Name: DFCI-55, Source.Name: DFCI-56///
 - Source.Name: DFCI-57//
 - Source.Name: DFCI-58,
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 - Source.Name: DFCI-60

Source.Name: DFCI-6

Source.Name: DFCI-62///Characteristics.Age.: Age <has_measurement <Measurement

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- Source.Name: DFCI-64
- Source.Name: DFCI-65///
 - Source.Name: DFCI-6
 - Source.Name: DFCI-
- Source.Name: DFCI-68/
- Source.Name: DFCI-69//

Source.Name: DFC

- Source.Name: DFCI-70,
 - Source.Name: DFCI-71

Value

An expression set

GSE12418

Expression analysis of stage III serous ovarian adenocarcinoma distinguishes a sub-group of survivors.

Description

It is difficult to predict the clinical outcome for patients with ovarian cancer. However, the use of biomarkers as additional prognostic factors may improve the outcome for these patients. In order to find novel candidate biomarkers, differences in gene expressions were analysed in 54 stage III serous ovarian adenocarcinomas with oligonucleotide microarrays containing 27,000 unique probes. The microarray data was verified with quantitative real-time polymerase chain reaction for the genes TACC1, MUC5B and PRAME. Using hierarchical cluster analysis we detected a subgroup that included 60% of the survivors. The gene expressions in tumours from patients in this sub-group of survivors were compared with the remaining tumours, and 204 genes were found to

be differently expressed. We conclude that the sub-group of survivors might represent patients with favourable tumour biology and sensitivity to treatment. A selection of the 204 genes might be used as a predictive model to distinguish patients within and outside of this group. Alternative chemotherapy strategies could then be offered as first-line treatment, which may lead to improvements in the clinical outcome for these patients.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Partheen K, Levan K, Osterberg L, Horvath G.Expression analyst
  Laboratory: Partheen, Horvath 2006
  Contact information:
  Title: Expression analysis of stage III serous ovarian adenocarcinoma distinguish
  URL:
  PMIDs: 16996261
  Abstract: A 177 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      SWEGENE H_v2.1.1_27k
  platform_shorttitle:
      SWEGENE H_v2.1.1_27k
   platform_summary:
      PartheenMetaData
   platform_manufacturer:
      other
   platform_distribution:
      non-commercial
   platform_accession:
      GPL5886
   version:
      2015-09-22 19:07:14
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 28 29 ... 29999 (11304 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

assayData: 11304 features, 54 samples Platform type: ------Available sample meta-data:

```
alt_sample_name:
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                       1 1
             1
    1
       1
                                    1 1
                                              1 1
                                                           1
 134LC 1426LB 1487DB 1528DC 1538DC 1567DB 1568DC 1574LC0
                                                     164DC
                                                           1658DC
    1
           1
              1
                        1
                              1
                                     1
                                           1
                                                  1
                                                        1
                                                               1
1760LB 1805DB
             193DC
                   198DC
                           202DC
                                211DC
                                         26DC
                                               272DC
                                                     405LB
                                                           436DC
    1
               1
                     1
                             1
                                          1
                                                 1
                                                      1
                                                               1
        1
                                     1
                                  47DC 480DC0
 452DC
       454LC
              45LA0
                    462DB
                          46LB0
                                              489DC
                                                     505DB
                                                           541DC
    1
        1
               1
                     1
                              1
                                    1
                                       1
                                               1
                                                      1
                                                               1
 559DC
        563LA
             626DC
                     662DC
                           719DC 742LC0
                                       755LC
                                               759DC
                                                      76DC
                                                            789DC
              1
                     1
                           1 1
                                          1
                                               1
                                                      1
                                                               1
    1
        1
  83LC 918DB0 988LC0
                     99LC0
      1
                     1
    1
             1
sample_type:
tumor
  54
histological_type:
ser
54
primarysite:
ov
54
summarystage:
late
 54
tumorstage:
3
54
substage:
b c
19 35
age_at_initial_pathologic_diagnosis:
  Min. 1st Qu. Median Mean 3rd Qu.
                                  Max.
 35.00 51.25 59.50 59.56 69.75
                                  84.00
pltx:
У
54
```

os_binary:

long short 20 34 debulking: optimal suboptimal 13 41

uncurated author metadata: title: 1035LA0///geo_accession: GSM311973///status: Public on Aug 12 2008///submiss title: 1047LB///geo_accession: GSM311974///status: Public on Aug 12 2008///subr title: 1059LB0///geo_accession: GSM311975///status: Public on Aug 12 2008///submiss title: 1177DB///geo_accession: GSM311976///status: Public on Aug 12 2008 title: 1178LB0///geo_accession: GSM311977///status: Public on Aug 12 2008///submiss title: 1180DB///geo_accession: GSM311978///status: Public on Aug 12 2008 title: 1186DB0///geo_accession: GSM311979///status: Public on Aug 12 2008// title: 123DC///geo_accession: GSM311945///status: Public on Aug 12 200 title: 1242LC0///geo_accession: GSM311980///status: Public on Aug 12 2008///submis title: 1274LC///geo_accession: GSM311981///status: Public on Aug 12 2008///sub title: 134LC///geo_accession: GSM311946///status: Public on Aug 12 2008///sub title: 1426LB///geo_accession: GSM311982///status: Public on Aug 12 2008///subr title: 1487DB///geo_accession: GSM311983///status: Public on Aug 12 2008 title: 1528DC///geo_accession: GSM311984///status: Public on Aug 12 200 title: 1538DC///geo_accession: GSM311985///status: Public on Aug 12 200 title: 1567DB///geo_accession: GSM311986///status: Public on Aug 12 2008 title: 1568DC///geo_accession: GSM311987///status: Public on Aug 12 200 title: 1574LC0///geo_accession: GSM311988///status: Public on Aug 12 2008///submis title: 164DC///geo_accession: GSM311947///status: Public on Aug 12 200 title: 1658DC///geo_accession: GSM311989///status: Public on Aug 12 200

title: 1760LB///geo_accession: GSM311990///status: Public on Aug 12 2008///subr title: 1805DB///geo_accession: GSM311991///status: Public on Aug 12 2008 title: 193DC///geo_accession: GSM311948///status: Public on Aug 12 200 title: 198DC///geo_accession: GSM311949///status: Public on Aug 12 200 title: 202DC///geo_accession: GSM311950///status: Public on Aug 12 200 title: 211DC///geo_accession: GSM311951///status: Public on Aug 12 200 title: 26DC///geo_accession: GSM311938///status: Public on Aug 12 200 title: 272DC///geo_accession: GSM311952///status: Public on Aug 12 200 title: 405LB///geo_accession: GSM311953///status: Public on Aug 12 2008///subr title: 436DC///geo_accession: GSM311954///status: Public on Aug 12 200 title: 452DC///geo_accession: GSM311955///status: Public on Aug 12 200 title: 454LC///geo_accession: GSM311956///status: Public on Aug 12 2008///sub title: 45LA0///geo_accession: GSM311939///status: Public on Aug 12 2008///submiss title: 462DB///geo_accession: GSM311957///status: Public on Aug 12 2008 title: 46LB0///geo_accession: GSM311940///status: Public on Aug 12 2008///submiss title: 47DC///geo_accession: GSM311941///status: Public on Aug 12 200 title: 480DC0///geo_accession: GSM311958///status: Public on Aug 12 2008/ title: 489DC///geo_accession: GSM311959///status: Public on Aug 12 200 title: 505DB///geo_accession: GSM311960///status: Public on Aug 12 2008 title: 541DC///geo_accession: GSM311961///status: Public on Aug 12 200 title: 559DC///geo_accession: GSM311962///status: Public on Aug 12 200 title: 563LA///geo_accession: GSM311963///status: Public on Aug 12 2008///subr title: 626DC///geo_accession: GSM311964///status: Public on Aug 12 200 title: 662DC///geo_accession: GSM311965///status: Public on Aug 12 200

title: 719DC///geo_accession: GSM311966///status: Public on Aug 12 200 title: 742LCO///geo_accession: GSM311967///status: Public on Aug 12 2008///submise title: 755LC///geo_accession: GSM311968///status: Public on Aug 12 2008///sub title: 759DC///geo_accession: GSM311969///status: Public on Aug 12 200 title: 76DC///geo_accession: GSM311942///status: Public on Aug 12 200 title: 789DC///geo_accession: GSM311942///status: Public on Aug 12 200 title: 83LC///geo_accession: GSM311943///status: Public on Aug 12 2008///sub title: 918DB0///geo_accession: GSM311971///status: Public on Aug 12 2008///sub title: 988LC0///geo_accession: GSM311972///status: Public on Aug 12 2008///submise title: 99LC0///geo_accession: GSM311944///status: Public on Aug 12 2008///submise

Value

An expression set

GSE12470

Gene expression profiling of advanced-stage serous ovarian cancers distinguishes novel subclasses and implicates ZEB2 in tumor progression and prognosis.

Description

To elucidate the mechanisms of rapid progression of serous ovarian cancer, gene expression profiles from 43 ovarian cancer tissues comprising eight early stage and 35 advanced stage tissues were carried out using oligonucleotide microarrays of 18,716 genes. By non-negative matrix factorization analysis using 178 genes, which were extracted as stage-specific genes, 35 advanced stage cases were classified into two subclasses with superior (n = 17) and poor (n = 18) outcome evaluated by progression-free survival (log rank test, P = 0.03). Of the 178 stage-specific genes, 112 genes were identified as showing different expression between the two subclasses. Of the 48 genes selected for biological function by gene ontology analysis or Ingenuity Pathway Analysis, five genes (ZEB2, CDH1, LTBP2, COL16A1, and ACTA2) were extracted as candidates for prognostic factors associated with progression-free survival. The relationship between high ZEB2 or low CDH1 expression and shorter progression-free survival was validated by real-time RT-PCR experiments of 37 independent advanced stage cancer samples. ZEB2 expression was negatively correlated with CDH1 expression in advanced stage samples, whereas ZEB2 knockdown in ovarian adenocarcinoma SKOV3 cells resulted in an increase in CDH1 expression. Multivariate analysis showed that high ZEB2 expression was independently associated with poor prognosis. Furthermore, the prognostic effect of E-cadherin encoded by CDH1 was verified using immunohistochemical analysis of an independent advanced stage cancer samples set (n = 74). These findings suggest that the expression of epithelial-mesenchymal transition-related genes such as ZEB2 and CDH1 may play important roles in the invasion process of advanced stage serous ovarian cancer.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Yoshihara K, Tajima A, Komata D, Yamamoto T, Kodama S, Fujiwan
  Laboratory: Yoshihara, Tanaka 2009
  Contact information:
  Title: Gene expression profiling of advanced-stage serous ovarian cancers disting
  URL:
  PMIDs: 19486012
  Abstract: A 253 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      Agilent-012097 Human 1A Microarray (V2) G4110B (Feature Number version)
  platform_shorttitle:
      Agilent G4110B
   platform_summary:
      hgug4110b
   platform_manufacturer:
      Agilent
   platform_distribution:
      commercial
   platform_accession:
      GPL887
   version:
      2015-09-22 19:08:17
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 3 5 ... 22571 (15999 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

assayData: 15999 features, 53 samples Platform type: ------Available sample meta-data:

alt_sample_name:	
	Advanced serous ovarian cancer 11
1	1
Advanced serous ovarian cancer 15	Advanced serous ovarian cancer 17
1	1
Advanced serous ovarian cancer 18	Advanced serous ovarian cancer 2
1	1
Advanced serous ovarian cancer 20	Advanced serous ovarian cancer 23
1	1
Advanced serous ovarian cancer 24	Advanced serous ovarian cancer 25
1	1
Advanced corous overian cancer 27	Advanced serous ovarian cancer 36
Advanced Serous Ovarran Cancer 27	Advanced Serous Ovarian cancer 50
±	L 20
_	Advanced serous ovarian cancer 38
1	1
Advanced serous ovarian cancer 39	Advanced serous ovarian cancer 42
1	1
Advanced serous ovarian cancer 43	Advanced serous ovarian cancer 45
1	1
Advanced serous ovarian cancer 46	Advanced serous ovarian cancer 49
1	1
Advanced serous ovarian cancer 50	Advanced serous ovarian cancer 51
1	1
Advanced serous ovarian cancer 52	Advanced serous ovarian cancer 53
1	1
Advanced serous ovarian cancer 54	Advanced serous ovarian cancer 55
1	1
Advanced serous ovarian cancer 56	Advanced serous ovarian cancer 57
1	1
-	Advanced serous ovarian cancer 6
Advanced Serous Ovarran cancer 50	Advanced Serous Ovarran cancer 0
	-
	Advanced serous ovarian cancer 61
1	1
Advanced serous ovarian cancer 62	Advanced serous ovarian cancer 64
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Advanced serous ovarian cancer 7	Early serous ovarian cancer 28
1	1
Early serous ovarian cancer 32	Early serous ovarian cancer 33
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Early serous ovarian cancer 35	Early serous ovarian cancer 5
1	1
Early serous ovarian cancer 65	Early serous ovarian cancer 8
1	1
- Early serous ovarian cancer 9	Peritoneum normal 12
1	1
- Peritoneum normal 15	Peritoneum normal 16
1	1
1	L

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Peritoneum normal 21
1
Peritoneum normal 3
1
Peritoneum normal 4
1
```

```
Peritoneum normal 18
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Peritoneum normal 23
1
Peritoneum normal 30
1
Peritoneum normal 7
```

```
sample_type:
healthy tumor
    10    43
```

histological_type: ser NA's 43 10

```
primarysite:
ov
53
```

summarystage: early late NA's 8 35 10

tumorstage: 1 NA's 8 45

```
uncurated_author_metadata:
```

11	Icurated	author_n	letauata	4:				
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	title:	Advanced	serous	ovarian	cancer	18///geo_accession:	GSM312143///status:	Puk
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title: Advanced serous ovarian cancer 27///geo_accession: GSM312158///status: Pub

title: Advanced serous ovarian cancer 2///geo_accession: GSM312138///st title: Advanced serous ovarian cancer 36///geo_accession: GSM312147///status: Pub title: Advanced serous ovarian cancer 37///geo_accession: GSM312148///status: Pub title: Advanced serous ovarian cancer 38///geo_accession: GSM312149///status: Pub title: Advanced serous ovarian cancer 39///geo_accession: GSM312159///status: Pub title: Advanced serous ovarian cancer 42///geo_accession: GSM312160///status: Pub title: Advanced serous ovarian cancer 43///geo_accession: GSM312150///status: Pub title: Advanced serous ovarian cancer 45///geo_accession: GSM312161///status: Pub title: Advanced serous ovarian cancer 46///geo_accession: GSM312162///status: Pub title: Advanced serous ovarian cancer 49///geo_accession: GSM312151///status: Pub title: Advanced serous ovarian cancer 50///geo_accession: GSM312163///status: Pub title: Advanced serous ovarian cancer 51///geo_accession: GSM312165///status: Pub title: Advanced serous ovarian cancer 52///geo_accession: GSM312167///status: Pub title: Advanced serous ovarian cancer 53///geo_accession: GSM312168///status: Pub title: Advanced serous ovarian cancer 54///geo_accession: GSM312152///status: Pub title: Advanced serous ovarian cancer 55///geo_accession: GSM312170///status: Publi title: Advanced serous ovarian cancer 56///geo_accession: GSM312171///status: Pub title: Advanced serous ovarian cancer 57///geo_accession: GSM312153///status: Pub title: Advanced serous ovarian cancer 58///geo_accession: GSM312172///status: Pub title: Advanced serous ovarian cancer 60///geo_accession: GSM312173///status: Pub title: Advanced serous ovarian cancer 61///geo_accession: GSM312154///status: Pub title: Advanced serous ovarian cancer 62///geo_accession: GSM312174///status: Pub title: Advanced serous ovarian cancer 64///geo_accession: GSM312175///status: Pub

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Value

An expression set

GSE13876

Survival-related profile, pathways, and transcription factors in ovarian cancer.

Description

Ovarian cancer has a poor prognosis due to advanced stage at presentation and either intrinsic or acquired resistance to classic cytotoxic drugs such as platinum and taxoids. Recent large clinical trials with different combinations and sequences of classic cytotoxic drugs indicate that further significant improvement in prognosis by this type of drugs is not to be expected. Currently a large number of drugs, targeting dysregulated molecular pathways in cancer cells have been developed and are introduced in the clinic. A major challenge is to identify those patients who will benefit from drugs targeting these specific dysregulated pathways. The aims of our study were (1) to develop a gene expression profile associated with overall survival in advanced stage serous ovarian cancer, (2) to assess the association of pathways and transcription factors with overall survival, and (3) to validate our identified profile and pathways/transcription factors in an independent set of ovarian cancers. According to a randomized design, profiling of 157 advanced stage serous ovarian cancers was performed in duplicate using approximately 35,000 70-mer oligonucleotide microarrays. A continuous predictor of overall survival was built taking into account well-known issues in microarray analysis, such as multiple testing and overfitting. A functional class scoring analysis was utilized to assess pathways/transcription factors for their association with overall survival. The prognostic value of genes that constitute our overall survival profile was validated on a fully independent, publicly available dataset of 118 well-defined primary serous ovarian cancers. Furthermore, functional class scoring analysis was also performed on this independent dataset to assess the similarities with results from our own dataset. An 86-gene overall survival profile discriminated between patients with unfavorable and favorable prognosis (median survival, 19 versus 41 mo, respectively; permutation p-value of log-rank statistic = 0.015) and maintained its independent prognostic value in multivariate analysis. Genes that composed the overall survival profile were also able to discriminate between the two risk groups in the independent dataset. In our dataset 17/167 pathways and 13/111 transcription factors were associated with overall survival, of which 16 and 12, respectively, were confirmed in the independent dataset. Our study provides new clues to genes, pathways, and transcription factors that contribute to the clinical outcome of serous ovarian cancer and might be exploited in designing new treatment strategies.

Format

```
experimentData(eset):
Experiment data
Experimenter name: Crijns AP, Fehrmann RS, de Jong S, Gerbens F, Meersma GJ, Klip
Laboratory: Crijns, van der Zee 2009
Contact information:
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Title: Survival-related profile, pathways, and transcription factors in ovarian of
  URL:
  PMIDs: 19192944
  Abstract: A 371 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
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      Operon v3 two-color
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  platform_manufacturer:
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  platform_distribution:
      non-commercial
  platform_accession:
     GPL7759
   version:
      2015-09-22 19:11:43
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  varMetadata: labelDescription
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Details

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uncurated_author_metadata:

title: Ovarian tumor sample 105 / Ovarian tumor sample 106///geo_accession:

title: Ovarian tumor sample 10 / Ovarian tumor sample 11///geo_accession:

title: Ovarian tumor sample 111 / Ovarian tumor sample 112///geo_accession: title: Ovarian tumor sample 115 / Ovarian tumor sample 117///geo_accession:

title: Ovarian tumor sample 126 / Ovarian tumor sample 127///geo_accession:

title: Ovarian tumor sample 13 / Ovarian tumor sample 14///geo_accession:

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Value

An expression set

GSE14764

A prognostic gene expression index in ovarian cancer - validation across different independent data sets.

Description

Ovarian carcinoma has the highest mortality rate among gynaecological malignancies. In this project, we investigated the hypothesis that molecular markers are able to predict outcome of ovarian cancer independently of classical clinical predictors, and that these molecular markers can be validated using independent data sets. We applied a semi-supervised method for prediction of patient survival. Microarrays from a cohort of 80 ovarian carcinomas (TOC cohort) were used for the development of a predictive model, which was then evaluated in an entirely independent cohort of 118 carcinomas (Duke cohort). A 300-gene ovarian prognostic index (OPI) was generated and validated in a leave-one-out approach in the TOC cohort (Kaplan-Meier analysis, p = 0.0087). In a second validation step, the prognostic power of the OPI was confirmed in an independent data set (Duke cohort, p = 0.0063). In multivariate analysis, the OPI was independent of the post-operative residual tumour, the main clinico-pathological prognostic parameter with an adjusted hazard ratio of 6.4 (TOC cohort, CI 1.8-23.5, p = 0.0049) and 1.9 (Duke cohort, CI 1.2-3.0, p = 0.0068). We constructed a combined score of molecular data (OPI) and clinical parameters (residual tumour), which was able to define patient groups with highly significant differences in survival. The integrated analysis of gene expression data as well as residual tumour can be used for optimized assessment of the prognosis of platinum-taxol-treated ovarian cancer. As traditional treatment options are limited, this analysis may be able to optimize clinical management and to identify those patients who would be candidates for new therapeutic strategies.

Format

```
experimentData(eset):
Experiment data
Experimenter name: Denkert C, Budczies J, Darb-Esfahani S, Gy??rffy B et al. A pr
Laboratory: Denkert, Lage 2009
Contact information:
Title: A prognostic gene expression index in ovarian cancer - validation across of
URL:
PMIDs: 19294737
Abstract: A 254 word abstract is available. Use 'abstract' method.
Information is available on: preprocessing
notes:
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    Affymetrix HG-U133A
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  platform_distribution:
      commercial
  platform_accession:
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   version:
      2015-09-22 19:13:08
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    (20967 total)
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  varMetadata: labelDescription
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Details

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Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
     n events median 0.95LCL 0.95UCL
 80.00 21.00
              4.52 4.19 NA
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Available sample meta-data:
_____
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                                    Max.
  1.00 20.75 40.50 40.50 60.25
                                    80.00
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tumor
  80
histological_type:
      clearcell
                         endo
                                        mix
                                                     other
            2
                          6
                                         1
                                                         2
           ser undifferentiated
           68
                           1
primarysite:
ov
```

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80
summarygrade:
high low
54 26
summarystage:
early late
9 71
tumorstage:
1 2 3 4
8 1 69 2
substage:
 a b c NA's
  4 6 32 38
grade:
1 2 3
3 23 54
recurrence_status:
norecurrence recurrence NA's
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                           4
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  210 660 1050 1011 1328
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2005-03-02 2006-07-26 2006-07-27 2006-07-28 2006-08-11 2006-08-18 2006-08-19
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2006-08-21
      5
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NA's
78
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Value

An expression set

GSE17260

Gene expression profile for predicting survival in advanced-stage serous ovarian cancer across two independent datasets.

Description

Advanced-stage ovarian cancer patients are generally treated with platinum/taxane-based chemotherapy after primary debulking surgery. However, there is a wide range of outcomes for individual patients. Therefore, the clinicopathological factors alone are insufficient for predicting prognosis. Our aim is to identify a progression-free survival (PFS)-related molecular profile for predicting survival of patients with advanced-stage serous ovarian cancer. Advanced-stage serous ovarian cancer tissues from 110 Japanese patients who underwent primary surgery and platinum/taxane-based chemotherapy were profiled using oligonucleotide microarrays. We selected 88 PFS-related genes by a univariate Cox model (p<0.01) and generated the prognostic index based on 88 PFS-related genes after adjustment of regression coefficients of the respective genes by ridge regression Cox model using 10-fold cross-validation. The prognostic index was independently associated with PFS time compared to other clinical factors in multivariate analysis [hazard ratio (HR), 3.72; 95% confidence interval (CI), 2.66-5.43; p<0.0001]. In an external dataset, multivariate analysis revealed that this prognostic index was significantly correlated with PFS time (HR, 1.54; 95% CI, 1.20-1.98; p = 0.0008). Furthermore, the correlation between the prognostic index and overall survival time was confirmed in the two independent external datasets (log rank test, p = 0.0010 and 0.0008). The prognostic ability of our index based on the 88-gene expression profile in ridge regression Cox hazard model was shown to be independent of other clinical factors in predicting cancer prognosis across two distinct datasets. Further study will be necessary to improve predictive accuracy of the

prognostic index toward clinical application for evaluation of the risk of recurrence in patients with advanced-stage serous ovarian cancer.

Format

```
experimentData(eset):
Experiment data
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  Laboratory: Yoshihara, Tanaka 2010
  Contact information:
  Title: Gene expression profile for predicting survival in advanced-stage serous of
  URL:
  PMIDs: 20300634
  Abstract: A 257 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
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   platform_distribution:
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   platform_accession:
      GPL6848
   version:
      2015-09-22 19:16:49
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  varMetadata: labelDescription
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Details

Available sample meta-data:

alt sample name:

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1	1	1
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Serous ovarian cancer 43	Serous ovarian cancer 44	Serous ovarian cancer 45
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(Other)	1	1
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11		
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tumor		
110		

histological_type: ser 110 primarysite: ov 110 summarygrade: high low 43 67 summarystage: late 110

tumorstage: 3 4

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93 17
substage:
         c NA's
  a b
     18
         69 17
  6
grade:
1 2 3
26 41 43
pltx:
 У
110
tax:
 У
110
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norecurrence recurrence
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                76
days_to_death:
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deceased living
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debulking:
   optimal suboptimal
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title: Serous ovarian cancer 109///geo_accession: GSM432226///status: Public on Ma title: Serous ovarian cancer 10///geo_accession: GSM43 title: Serous ovarian cancer 110///geo_accession: GSM432228///status: Public on Man title: Serous ovarian cancer 111///geo_accession: GSM432229///status: Public on Man title: Serous ovarian cancer 112///geo_accession: GSM43 title: Serous ovarian cancer 113///geo_accession: GSM432231 title: Serous ovarian cancer 114///geo_accession: GSM43223 title: Serous ovarian cancer 115///geo_accession: GSM432233 title: Serous ovarian cancer 116///geo_accession: GSM432234 title: Serous ovarian cancer 117///geo_accession: GSM43 title: Serous ovarian cancer 118///geo_accession: GSM43223 title: Serous ovarian cancer 119///geo_accession: GSM43 title: Serous ovarian cancer 11///geo_accession: GSM43 title: Serous ovarian cancer 120///geo_accession: GSM432 title: Serous ovarian cancer 122///geo_accession: GSM43224 title: Serous ovarian cancer 123///geo_accession: GSM432242 title: Serous ovarian cancer 127///geo_accession: GSM432243 title: Serous ovarian cancer 129///geo_accession: GSM43 title: Serous ovarian cancer 12///geo_accession: GSM4 title: Serous ovarian cancer 130///geo_accession: GSM432245 title: Serous ovarian cancer 131///geo_accession: GSM43 title: Serous ovarian cancer 132///geo_accession: GSM43 title: Serous ovarian cancer 134///geo_accession: GSM43 title: Serous ovarian cancer 136///geo_accession: GSM43

title: Serous ovarian cancer 137///geo_accession: GSM43 title: Serous ovarian cancer 139///geo_accession: GSM43 title: Serous ovarian cancer 140///geo_accession: GSM4322 title: Serous ovarian cancer 143///geo_accession: GSM43225 title: Serous ovarian cancer 144///geo_accession: GSM4322 title: Serous ovarian cancer 145///geo_accession: GSM432255 title: Serous ovarian cancer 146///geo_accession: GSM432256 title: Serous ovarian cancer 148///geo_accession: GSM43 title: Serous ovarian cancer 149///geo_accession: GSM43 title: Serous ovarian cancer 150///geo_accession: GSM432260 title: Serous ovarian cancer 151///geo_accession: GSM43 title: Serous ovarian cancer 154///geo_accession: GSN title: Serous ovarian cancer 156///geo_accession: GSM43 title: Serous ovarian cancer 157///geo_accession: GSM43 title: Serous ovarian cancer 15///geo_accession: GSM43 title: Serous ovarian cancer 160///geo_accession: GSM432260 title: Serous ovarian cancer 16///geo_accession: GSM43 title: Serous ovarian cancer 171///geo_accession: GSM43226 title: Serous ovarian cancer 172///geo_accession: GSM43226 title: Serous ovarian cancer 173///geo_accession: GSM4322⁻ title: Serous ovarian cancer 174///geo_accession: GSM title: Serous ovarian cancer 176///geo_accession: GSM432 title: Serous ovarian cancer 178///geo_accession: GSN title: Serous ovarian cancer 17///geo_accession: GSM43

title: Serous ovarian cancer 182///geo_accession: GSM4322 title: Serous ovarian cancer 183///geo_accession: GSM432276 title: Serous ovarian cancer 184///geo_accession: GSM43 title: Serous ovarian cancer 185///geo_accession: GSM43 title: Serous ovarian cancer 186///geo_accession: GSM43 title: Serous ovarian cancer 18///geo_accession: GSM43 title: Serous ovarian cancer 20///geo_accession: GSM43 title: Serous ovarian cancer 22///geo_accession: GSM432282 title: Serous ovarian cancer 23///geo_accession: GSM4322 title: Serous ovarian cancer 25///geo_accession: GSM43 title: Serous ovarian cancer 27///geo_accession: GSM43228 title: Serous ovarian cancer 2///geo_accession: GSM432280 title: Serous ovarian cancer 31///geo_accession: GSM title: Serous ovarian cancer 36///geo_accession: GSM title: Serous ovarian cancer 37///geo_accession: GSM43 title: Serous ovarian cancer 38///geo_accession: GSM43 title: Serous ovarian cancer 41///geo_accession: GSM43 title: Serous ovarian cancer 42///geo_accession: GSM432292 title: Serous ovarian cancer 43///geo_accession: GSM43229 title: Serous ovarian cancer 44///geo_accession: GSM43229 title: Serous ovarian cancer 45///geo_accession: GSM43 title: Serous ovarian cancer 49///geo_accession: GSM432296 title: Serous ovarian cancer 4///geo_accession: GSM43229 title: Serous ovarian cancer 50///geo_accession: GSM4

title: Serous ovarian cancer 51///geo_accession: GSM432298 title: Serous ovarian cancer 52///geo_accession: GSM432 title: Serous ovarian cancer 53///geo_accession: GSM432 title: Serous ovarian cancer 54///geo_accession: GSM43 title: Serous ovarian cancer 55///geo_accession: GSM43 title: Serous ovarian cancer 56///geo_accession: GSM432303 title: Serous ovarian cancer 57///geo_accession: GSM43 title: Serous ovarian cancer 58///geo_accession: GSM43 title: Serous ovarian cancer 60///geo_accession: GSM4323 title: Serous ovarian cancer 61///geo_accession: GSM432308 title: Serous ovarian cancer 62///geo_accession: GSM432309 title: Serous ovarian cancer 64///geo_accession: GSM43 title: Serous ovarian cancer 66///geo_accession: GSM43 title: Serous ovarian cancer 67///geo_accession: GSM43 title: Serous ovarian cancer 68///geo_accession: GSM43 title: Serous ovarian cancer 69///geo_accession: GSM43 title: Serous ovarian cancer 6///geo_accession: GSM43 title: Serous ovarian cancer 72///geo_accession: GSM43 title: Serous ovarian cancer 77///geo_accession: GSM4323 title: Serous ovarian cancer 79///geo_accession: GSM43 title: Serous ovarian cancer 7///geo_accession: GSM43 title: Serous ovarian cancer 80///geo_accession: GSM432319

Value

An expression set

GSE18520

A gene signature predictive for outcome in advanced ovarian cancer identifies a survival factor: microfibril-associated glycoprotein 2.

Description

Advanced stage papillary serous tumors of the ovary are responsible for the majority of ovarian cancer deaths, yet the molecular determinants modulating patient survival are poorly characterized. Here, we identify and validate a prognostic gene expression signature correlating with survival in a series of microdissected serous ovarian tumors. Independent evaluation confirmed the association of a prognostic gene microfibril-associated glycoprotein 2 (MAGP2) with poor prognosis, whereas in vitro mechanistic analyses demonstrated its ability to prolong tumor cell survival and stimulate endothelial cell motility and survival via the alpha(V)beta(3) integrin receptor. Increased MAGP2 expression correlated with microvessel density suggesting a proangiogenic role in vivo. Thus, MAGP2 may serve as a survival-associated target.

Format

```
experimentData(eset):
Experiment data
 Experimenter name: Mok SC, Bonome T, Vathipadiekal V, Bell A, Johnson ME, Wong KH
  Laboratory: Mok, Birrer 2009
  Contact information:
  Title: A gene signature predictive for outcome in advanced ovarian cancer identia
  URL:
  PMIDs: 19962670
 Abstract: A 110 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [HG-U133_Plus_2] Affymetrix Human Genome U133 Plus 2.0 Array
  platform_shorttitle:
      Affymetrix HG-U133Plus2
  platform_summary:
      hgu133plus2
  platform_manufacturer:
      Affymetrix|Operon
  platform_distribution:
      commercial | non-commercial
  platform_accession:
      GPL570|GPL9216
   version:
```

```
2015-09-22 19:21:25
```

```
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (42447 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 42447 features, 63 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit (formula = Surv(time, cens) ~ -1)
  10 observations deleted due to missingness
     n events median 0.95LCL 0.95UCL
  53.00 41.00 2.05 1.48 3.70
_____
Available sample meta-data:
_____
alt_sample_name:
  Min. 1st Qu. Median Mean 3rd Qu. Max.
  312.0 395.0 694.0 893.3 1040.0 2237.0
sample_type:
healthy tumor
    10
          53
histological_type:
 ser NA's
 53 10
primarysite:
ov
63
summarygrade:
high NA's
 53 10
summarystage:
late NA's
 53 10
```

```
tumorstage:
  3 NA's
  53 10
grade:
  3 NA's
  53 10
days_to_death:
  Min. 1st Qu.MedianMean 3rd Qu.Max.NA's15045063012121440450010
vital_status:
deceased living NA's
41 12 10
debulking:
optimal
    63
percent_normal_cells:
0
63
percent_stromal_cells:
0
63
percent_tumor_cells:
100
63
batch:
2004-03-12 2004-04-08 2004-04-09 2004-07-20 2004-08-12 2004-08-13 2004-09-30
       20 6 9 11 10 1 6
uncurated_author_metadata:
                                             title: Normal Ovary, 2008///geo_acc
                                             title: Normal Ovary, 2061///geo_acc
                                             title: Normal Ovary, 2064///geo_acc
                                             title: Normal Ovary, 2085///geo_acc
                                             title: Normal Ovary, 2225///geo_acc
                                             title: Normal Ovary, 2226///geo_acc
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title: Normal Ovary, 2228///geo_acc title: Normal Ovary, 2230///geo_acc title: Normal Ovary, 2234///geo_acc

title: Normal Ovary, 2237///geo_acc

title: Ovarian Tumor, 1109///geo_accession: GSM461390///status: Public on Oct 17 20 title: Ovarian Tumor, 1214///geo_accession: GSM461391///status: Public on Oct 1 title: Ovarian Tumor, 1231///geo_accession: GSM461367///status: Public on Oct 1 title: Ovarian Tumor, 1562///geo_accession: GSM461368///status: Public on Oct 17 20 title: Ovarian Tumor, 1660///geo_accession: GSM461369///status: Public on Oct 17 20 title: Ovarian Tumor, 1993///geo_accession: GSM461400///status: Public on Oct 17 20 title: Ovarian Tumor, 312///geo_accession: GSM461379///status: Public on Oct title: Ovarian Tumor, 317///geo_accession: GSM461348///status: Public on Oct 17 20 title: Ovarian Tumor, 321///geo_accession: GSM461380///status: Public on Oct title: Ovarian Tumor, 324///geo_accession: GSM461373///status: Public on Oct title: Ovarian Tumor, 332///geo_accession: GSM461349///status: Public on Oct title: Ovarian Tumor, 345///geo_accession: GSM461392///status: Public on Oct title: Ovarian Tumor, 349///geo_accession: GSM461350///status: Public on Oct 17 20 title: Ovarian Tumor, 351///geo_accession: GSM461351///status: Public on Oct 17 20 title: Ovarian Tumor, 358///geo_accession: GSM461393///status: Public on Oct title: Ovarian Tumor, 367///geo_accession: GSM461381///status: Public on Oct title: Ovarian Tumor, 377///geo_accession: GSM461374///status: Public on Oct title: Ovarian Tumor, 380///geo_accession: GSM461375///status: Public on Oct title: Ovarian Tumor, 386///geo_accession: GSM461352///status: Public on Oct title: Ovarian Tumor, 388///geo_accession: GSM461353///status: Public on Oct 17 20

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title: Ovarian Tumor, 778///geo_accession: GSM461399///status: Public on Oct title: Ovarian Tumor, 780///geo_accession: GSM461364///status: Public on Oct 17 2 title: Ovarian Tumor, 786///geo_accession: GSM461387///status: Public on Oct 17 2 title: Ovarian Tumor, 794///geo_accession: GSM461388///status: Public on Oct title: Ovarian Tumor, 799///geo_accession: GSM461365///status: Public on Oct title: Ovarian Tumor, 800///geo_accession: GSM461365///status: Public on Oct title: Ovarian Tumor, 872///geo_accession: GSM461366///status: Public on Oct title: Ovarian Tumor, 979///geo_accession: GSM461366///status: Public on Oct title: Ovarian Tumor, 970///geo_accession: GSM461371///status: Public on Oct 17 2

duplicates:

GSE18520.GSE18520_GSM462649 1 GSE18520.GSE18520_GSM462649///GSE18520.GSE18520_GSM462650 1 GSE18520.GSE18520_GSM462650 1 NA's 60

Value

An expression set

GSE19829 Gene expression profile of BRCAness that correlates with responsiveness to chemotherapy and with outcome in patients with epithelial ovarian cancer.

Description

To define a gene expression profile of BRCAness that correlates with chemotherapy response and outcome in epithelial ovarian cancer (EOC). A publicly available microarray data set including 61 patients with EOC with either sporadic disease or BRCA(1/2) germline mutations was used for development of the BRCAness profile. Correlation with platinum responsiveness was assessed in platinum-sensitive and platinum-resistant tumor biopsy specimens from six patients with BRCA

germline mutations. Association with poly-ADP ribose polymerase (PARP) inhibitor responsiveness and with radiation-induced RAD51 foci formation (a surrogate of homologous recombination) was assessed in Capan-1 cell line clones. The BRCAness profile was validated in 70 patients enriched for sporadic disease to assess its association with outcome. The BRCAness profile accurately predicted platinum responsiveness in eight out of 10 patient-derived tumor specimens, and between PARP-inhibitor sensitivity and resistance in four out of four Capan-1 clones. [corrected] When applied to the 70 patients with sporadic disease, patients with the BRCA-like (BL) profile had improved disease-free survival (34 months v 15 months; log-rank P = .013) and overall survival (72 months v 41 months; log-rank P = .006) compared with patients with a non-BRCA-like (NBL) profile, respectively. The BRCAness profile maintained independent prognostic value in multivariate analysis, which controlled for other known clinical prognostic factors. The BRCAness profile correlates with responsiveness to platinum and PARP inhibitors and identifies a subset of sporadic patients with improved outcome. Additional evaluation of this profile as a predictive tool in patients with sporadic EOC is warranted.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Konstantinopoulos PA, Spentzos D, Karlan BY, Taniguchi T et al
 Laboratory: Konstantinopoulos, Cannistra 2010 hgu95
  Contact information:
  Title: Gene expression profile of BRCAness that correlates with responsiveness to
  URT.
  PMIDs: 20547991
 Abstract: A 241 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [HG_U95Av2] Affymetrix Human Genome U95 Version 2 Array
  platform shorttitle:
      Affymetrix HG_U95Av2
  platform_summary:
      hgu95av2
  platform_manufacturer:
      Affymetrix
  platform distribution:
      commercial
  platform_accession:
      GPL570 | GPL8300
   version:
      2015-09-22 19:26:29
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-MurIL4_at (54253 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

1

1

1

1

1

1

1

1

1

1

1

1

1

1

1

1

Details

assayData: 54253 features, 70 samples Platform type: Overall survival time-to-event summary (in years): Call: survfit(formula = Surv(time, cens) ~ -1) n events median 0.95LCL 0.95UCL 40.00 3.78 2.96 5.92 70.00 Available sample meta-data: _____ alt_sample_name: Ovarian cancer_sample 1 Ovarian cancer_sample 10 Ovarian cancer_sample 11 1 1 Ovarian cancer_sample 12 Ovarian cancer_sample 13 Ovarian cancer_sample 14 Ovarian cancer_sample 15 Ovarian cancer_sample 16 Ovarian cancer_sample 17 1 Ovarian cancer_sample 18 Ovarian cancer_sample 19 Ovarian cancer_sample 2 1 1 Ovarian cancer_sample 20 Ovarian cancer_sample 21 Ovarian cancer_sample 22 1 Ovarian cancer_sample 23 Ovarian cancer_sample 24 Ovarian cancer_sample 25 1 1 Ovarian cancer_sample 26 Ovarian cancer_sample 27 Ovarian cancer_sample 28 1 Ovarian cancer_sample 29 Ovarian cancer_sample 3 Ovarian cancer_sample 30 1 1 Ovarian cancer_sample 31 Ovarian cancer_sample 32 Ovarian cancer_sample 33 Ovarian cancer_sample 34 Ovarian cancer_sample 35 Ovarian cancer_sample 36 1 1 Ovarian cancer_sample 37 Ovarian cancer_sample 38 Ovarian cancer_sample 39 1 Ovarian cancer_sample 4 Ovarian cancer_sample 40 Ovarian cancer_sample 41 1 1 Ovarian cancer_sample 42 Ovarian cancer_sample 43 Ovarian cancer_sample 44 1 Ovarian cancer_sample 45 Ovarian cancer_sample 46 Ovarian cancer_sample 47 1 1

Ovarian cancer_sample 48 Ovarian cancer_sample 49 Ovarian cancer_sample 5

Ovarian cancer_sample 50 Ovarian cancer_sample 51 Ovarian cancer_sample 52

1

1

1

1

```
Ovarian cancer_sample 53 Ovarian cancer_sample 54 Ovarian cancer_sample 55
                                                                         1
                       1
                                                1
Ovarian cancer_sample 56 Ovarian cancer_sample 57 Ovarian cancer_sample 58
                                                                         1
Ovarian cancer_sample 59 Ovarian cancer_sample 6 Ovarian cancer_sample 60
                                                                         1
                       1
                                                1
Ovarian cancer_sample 61 Ovarian cancer_sample 62 Ovarian cancer_sample 63
                                                1
                                                                         1
Ovarian cancer_sample 64 Ovarian cancer_sample 65 Ovarian cancer_sample 66
                                                1
                                                                         1
                       1
Ovarian cancer_sample 67 Ovarian cancer_sample 68 Ovarian cancer_sample 69
                                                                         1
                                                1
 Ovarian cancer_sample 7 Ovarian cancer_sample 70 Ovarian cancer_sample 8
                                                                         1
 Ovarian cancer_sample 9
batch:
2001-09-14 2001-12-14 2002-08-20 2003-09-09 2003-09-18 2009-08-14
         7
                    4
                           14
                                   13
                                                    4
                                                               2.8
days_to_death:
  Min. 1st Qu. Median Mean 3rd Qu.
                                          Max.
   30.0 667.5 1125.0 1170.0 1522.0 3450.0
primarysite:
ov
70
sample_type:
tumor
   70
uncurated_author_metadata:
            title: Ovarian cancer_sample 10///geo_accession: GSM495148///status: Pu
            title: Ovarian cancer_sample 11///geo_accession: GSM495149///status: Pu
                 title: Ovarian cancer_sample 12///geo_accession: GSM495150///statu
            title: Ovarian cancer_sample 13///geo_accession: GSM495151///status: Pu
            title: Ovarian cancer_sample 14///geo_accession: GSM495152///status: Pu
            title: Ovarian cancer_sample 15///geo_accession: GSM495153///status: Pu
            title: Ovarian cancer_sample 16///geo_accession: GSM495154///status: Pu
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title: Ovarian cancer_sample 39///geo_accession: GSM495177///status: Public on Jul title: Ovarian cancer_sample 3///geo_accession: GSM495141///status title: Ovarian cancer_sample 40///geo_accession: GSM495178///status: Public of title: Ovarian cancer_sample 41///geo_accession: GSM495179///status: Public on Jul title: Ovarian cancer_sample 42///geo_accession: GSM495180///status: Public of title: Ovarian cancer_sample 43///geo_accession: GSM495181///status: Public on Jul title: Ovarian cancer_sample 44///geo_accession: GSM495182///status: Public or title: Ovarian cancer_sample 45///geo_accession: GSM495183///status: Public or title: Ovarian cancer_sample 46///geo_accession: GSM495184///status: Public on Jul title: Ovarian cancer_sample 47///geo_accession: GSM495185///status: Public or title: Ovarian cancer_sample 48///geo_accession: GSM495186///status: Public on Jul title: Ovarian cancer_sample 49///geo_accession: GSM495187///status: Public or title: Ovarian cancer_sample 4///geo_accession: GSM495142///status title: Ovarian cancer_sample 50///geo_accession: GSM495188///status: Public on Jul title: Ovarian cancer_sample 51///geo_accession: GSM495189///status: Public or title: Ovarian cancer_sample 52///geo_accession: GSM495190///status: Public or title: Ovarian cancer_sample 53///geo_accession: GSM495191///status: Public or title: Ovarian cancer_sample 54///geo_accession: GSM495192///status: Public or title: Ovarian cancer_sample 55///qeo_accession: GSM495193///status: Public on Jul title: Ovarian cancer_sample 56///geo_accession: GSM495194///status: Public on C title: Ovarian cancer_sample 57///geo_accession: GSM495195///status: Public on C title: Ovarian cancer_sample 58///geo_accession: GSM495196///status: Public on Ju title: Ovarian cancer_sample 59///geo_accession: GSM495197///status: Public title: Ovarian cancer_sample 5///geo_accession: GSM495143///status: Pu

title: Ovarian cancer_sample 60///geo_accession: GSM495198///status: Public on Ju title: Ovarian cancer_sample 61///geo_accession: GSM495200///status: Public on Ju title: Ovarian cancer_sample 63///geo_accession: GSM495201///status: Public on Ju title: Ovarian cancer_sample 63///geo_accession: GSM495202///status: Public on Ju title: Ovarian cancer_sample 64///geo_accession: GSM495203///status: Public on Ju title: Ovarian cancer_sample 65///geo_accession: GSM495203///status: Public on title: Ovarian cancer_sample 66///geo_accession: GSM495204///status: Public on title: Ovarian cancer_sample 66///geo_accession: GSM495205///status: Public on Ju title: Ovarian cancer_sample 66///geo_accession: GSM495206///status: Public on Ju title: Ovarian cancer_sample 68///geo_accession: GSM495206///status: Public on Ju title: Ovarian cancer_sample 68///geo_accession: GSM495206///status: Public on Ju title: Ovarian cancer_sample 69///geo_accession: GSM495206///status: Public on Ju title: Ovarian cancer_sample 69///geo_accession: GSM495206///status: Public on Ju title: Ovarian cancer_sample 69///geo_accession: GSM495208///status: Public on Ju title: Ovarian cancer_sample 70///geo_accession: GSM495144///status: Public title: Ovarian cancer_sample 70///geo_accession: GSM495145///status: Public title: Ovarian cancer_sample 8///geo_accession: GSM495146///status: Public

vital_status: deceased living 40 30

Value

An expression set

GSE20565 A genomic and transcriptomic approach for a differential diagnosis between primary and secondary ovarian carcinomas in patients with a previous history of breast cancer.

Description

The distinction between primary and secondary ovarian tumors may be challenging for pathologists. The purpose of the present work was to develop genomic and transcriptomic tools to further refine the pathological diagnosis of ovarian tumors after a previous history of breast cancer.Sixteen paired breast-ovary tumors from patients with a former diagnosis of breast cancer were collected. The genomic profiles of paired tumors were analyzed using the Affymetrix GeneChip Mapping 50 K Xba Array or Genome-Wide Human SNP Array 6.0 (for one pair), and the data were normalized with ITALICS (ITerative and Alternative normaLIzation and Copy number calling for affymetrix Snp arrays) algorithm or Partek Genomic Suite, respectively. The transcriptome of paired samples was analyzed using Affymetrix GeneChip Human Genome U133 Plus 2.0 Arrays, and the data were normalized with gc-Robust Multi-array Average (gcRMA) algorithm. A hierarchical clustering of these samples was performed, combined with a dataset of well-identified primary and secondary ovarian tumors. In 12 of the 16 paired tumors analyzed, the comparison of genomic profiles confirmed the pathological diagnosis of primary ovarian tumor (n = 5) or metastasis of breast cancer (n = 7). Among four cases with uncertain pathological diagnosis, genomic profiles were clearly distinct between the ovarian and breast tumors in two pairs, thus indicating primary ovarian carcinomas, and showed common patterns in the two others, indicating metastases from breast cancer. In all pairs, the result of the transcriptomic analysis was concordant with that of the genomic analysis. In patients with ovarian carcinoma and a previous history of breast cancer, SNP array analysis can be used to distinguish primary and secondary ovarian tumors. Transcriptomic analysis may be used when primary breast tissue specimen is not available.

Format

```
experimentData(eset):
Experiment data
 Experimenter name: Meyniel JP, Cottu PH, Decraene C, Stern MH, Couturier J, Lebig
  Laboratory: Meyniel, Sastre-Garau 2010
  Contact information:
  Title: A genomic and transcriptomic approach for a differential diagnosis between
  URT:
  PMIDs: 20492709
 Abstract: A 277 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform title:
      [HG-U133_Plus_2] Affymetrix Human Genome U133 Plus 2.0 Array
  platform_shorttitle:
      Affymetrix HG-U133Plus2
  platform_summary:
      hgu133plus2
  platform_manufacturer:
      Affymetrix
  platform_distribution:
      commercial
  platform accession:
      GPL570 | GPL2005 | GPL6801
```

```
version:
    2015-09-22 19:33:01
featureData(eset):
An object of class 'AnnotatedDataFrame'
featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (42447 total)
varLabels: probeset gene EntrezGene.ID best_probe
varMetadata: labelDescription
```

Details

assayData: 42447 features, 140 samples Platform type: Available sample meta-data:

alt_sample_name:

ait_sar	npie_name:				
Breast	metastasis	in	the	ovary_OC01_ARN0016	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0017	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0020	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0029	-
Breast	metastasis	in	the	ovary_OC01_ARN0035	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0046	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0051	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0053	-
Breast	metastasis	in	the	ovary_OC01_ARN0055	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0060	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0069	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0073	-
Breast	metastasis	in	the	ovary_OC01_ARN0077	[HG-U133_Plus_2]
Breast	metastasis	in	the	ovary_OC01_ARN0079	
Breast	metastasis	in	the	ovary_OC01_ARN0081	1 [HG-U133_Plus_2]

Breast metastasis in the ovary_OC01_ARN0083 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0092 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0097 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0098 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0099 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0102 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0104 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0112 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0120 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0121 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0123 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0126 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0141 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0142 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0143 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0145 [HG-U133_Plus_2] 1 Breast metastasis in the ovary_OC01_ARN0146 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0153 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0162 [HG-U133_Plus_2] Breast metastasis in the ovary_OC01_ARN0201 [HG-U133_Plus_2] Ovarian carcinoma OC01 ARN0001 [HG-U133 Plus 2] Ovarian carcinoma_OC01_ARN0002 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0004 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0005 [HG-U133_Plus_2]

1 Ovarian carcinoma_OC01_ARN0007 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0008 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0009 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0010 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0011 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0012 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0013 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0015 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0022 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0023 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0025 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0028 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0030 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0032 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0034 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0036 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0037 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0038 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0039 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0041 [HG-U133_Plus_2] Ovarian carcinoma OC01 ARN0042 [HG-U133 Plus 2] Ovarian carcinoma_OC01_ARN0045 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0049 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0057 [HG-U133_Plus_2]

Ovarian carcinoma_OC01_ARN0058 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0061 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0062 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0063 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0064 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0066 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0067 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0070 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0072 [HG-U133_Plus 2] 1 Ovarian carcinoma_OC01_ARN0075 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0076 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0080 [HG-U133_Plus_2] 1 Ovarian carcinoma OC01 ARN0084 [HG-U133 Plus 2] 1 Ovarian carcinoma_OC01_ARN0085 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0089 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0091 [HG-U133_Plus_2] 1 Ovarian carcinoma_OC01_ARN0093 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0095 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0096 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0100 [HG-U133_Plus_2] Ovarian carcinoma OC01 ARN0101 [HG-U133 Plus 2] Ovarian carcinoma_OC01_ARN0103 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0105 [HG-U133_Plus_2] Ovarian carcinoma_OC01_ARN0106 [HG-U133_Plus_2]

```
1
Ovarian carcinoma_OC01_ARN0107 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0108 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0109 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0111 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0113 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0114 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0115 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0116 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0118 [HG-U133_Plus_2]
                                              1
Ovarian carcinoma_OC01_ARN0119 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0124 [HG-U133_Plus_2]
                                               1
Ovarian carcinoma_OC01_ARN0125 [HG-U133_Plus_2]
                                              1
                                         (Other)
                                              41
```

```
sample_type:
tumor
 140
histological_type:
clearcell endo mucinous
                           other
                                             NA's
                                    ser
                 7
                                      71
      6
            6
                              6
                                              44
primarysite:
other ov
  44
       96
summarygrade:
high low NA's
 63 33 44
summarystage:
early late NA's
  27
     67 46
```

tumorstage: 3 2 4 NA's 1 9 52 18 15 46 substage: c NA's а b 14 10 55 61 grade: 1 2 3 NA's 6 27 63 44 batch: 2006-06-01 2006-06-27 2006-06-28 2006-06-29 2006-06-30 2006-07-20 2008-03-06 21 18 37 20 36 7 1 uncurated_author_metadata: title: Breast metastasis in the ovary_OC01_ARN0016 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0017 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0020 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0029 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0035 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0046 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0051 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0053 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0055 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0060 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0069 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0073 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0077 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0079 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0081 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0083 [HG-U133_Plus_2]///geo_accession

title: Breast metastasis in the ovary_OC01_ARN0092 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0097 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0098 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0099 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0102 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0104 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0112 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0120 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0121 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0123 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0126 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0141 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0142 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0143 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0145 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0146 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0153 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0162 [HG-U133_Plus_2]///geo_accession title: Breast metastasis in the ovary_OC01_ARN0201 [HG-U133_Plus_2]///geo_accession title: Ovarian carcinoma_OC01_A

title: Ovarian carcinoma_OC01_ARN0002

title: Ovarian carcinoma_OC01_ARM

title: Ovarian carcinoma_OC01_A

title: Ovarian carcinoma_OC01_ARM

title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_AB title: Ovarian carcinoma_OC01_A title: Ovarian carcinoma_OC01_A title: Ovarian carcinoma_OC01_ARN00 title: Ovarian carcinoma_OC01_A title: Ovarian carcinoma_OC01_ARN0030 title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_A title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARN0041 title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARN0045 title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM

title: Ovarian carcinoma_OC01_ARN005

title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_A title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_AH title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARN0076 [H title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARN008 title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARN0091 [H title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARN00 title: Ovarian carcinoma_OC01_AF title: Ovarian carcinoma_OC01_ARN010 title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM title: Ovarian carcinoma_OC01_ARM

title: Ovarian carcinoma_OCO1_ARN title: Ovarian carcinoma_OCO1_ARN0109 title: Ovarian carcinoma_OCO1_ARN0113 title: Ovarian carcinoma_OCO1_ARN0114 [HO title: Ovarian carcinoma_OCO1_ARN0115 title: Ovarian carcinoma_OCO1_ARN015 title: Ovarian carcinoma_OCO1_ARN015

```
duplicates:
GSE20565.GSE20565_GSM516722 GSE20565.GSE20565_GSM516741
1 1
NA's
138
```

Value

An expression set

GSE2109

IGC EXpression Project for Oncology

Description

EXpression Project for Oncology, International Genomics Consortium, www.intgen.org

Format

```
experimentData(eset):
Experiment data
  Experimenter name: EXpression Project for Oncology, International Genomics Consor
 Laboratory: exp0, IGC 2005
  Contact information:
  Title: IGC EXpression Project for Oncology
  URL:
  PMIDs: PMID unknown
  Abstract: A 8 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [HG-U133_Plus_2] Affymetrix Human Genome U133 Plus 2.0 Array
   platform_shorttitle:
      Affymetrix HG-U133Plus2
   platform_summary:
      hgu133plus2
   platform_manufacturer:
      Affymetrix
  platform distribution:
     commercial
  platform accession:
     GPL570
   version:
      2015-09-22 19:40:35
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (42447 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

1	1
Omentum - 8240	Ovary - 101094
1	1
1 Ovary - 101109 1	0vary - 101120 1
Ovary - 101150	Ovary - 1018
1	1
Ovary - 1040 1	
Ovary - 112866	Ovary - 112867
1	1
Ovary - 118662	Ovary - 118671
1	1
Ovary - 1241	Ovary - 1270
1	1
Ovary - 129660	Ovary - 129669
1	1
Ovary - 1311	Ovary - 1313
1	1
Ovary - 1323	Ovary - 133643
1	1
Ovary - 133651	Ovary - 1351
1	1
Ovary - 151614	Ovary - 151622
1	1
Ovary - 161465	Ovary - 161524
1	1
Ovary - 161525	Ovary - 161534
1	1
Ovary - 1636	Ovary - 1639
1	1
Ovary - 1643	Ovary - 170809
1	1
Ovary - 174931	Ovary - 174936
1	1
Ovary - 180953	Ovary - 184837
1	1
Ovary - 187243	Ovary - 187246
1	1
Ovary - 187251	Ovary - 187253 1
Ovary - 191413	Ovary - 191424
1	1
Ovary - 195198	Ovary - 199399
1	1
Ovary - 199400	Ovary - 202030
1	1
Ovary - 202041	Ovary - 20284

1	1
Ovary - 20285	Ovary - 20296
1	1
Ovary - 20307 1	
Ovary - 20323 1	
Ovary - 20326	Ovary - 20329
1	1
Ovary - 207532 1	
Ovary - 209709	Ovary - 209714
1	1
Ovary - 209718	Ovary - 211371
1	1
Ovary - 211372	Ovary - 211395
1	1
Ovary - 211409	Ovary - 21758
1	1
Ovary - 219571	Ovary - 219581
1	1
Ovary - 219590	Ovary - 219604
1	1
Ovary - 21981	Ovary - 22218
1	1
Ovary - 226414 1	
Ovary - 228537	Ovary - 228549
1	1
Ovary - 231863	Ovary - 234328
1	1
Ovary - 234329 1	
Ovary - 235692 1	
Ovary - 23862 1	- Ovary - 23884 1
Ovary - 23904	Ovary - 23930
1	1
Ovary - 23934 1	- Ovary - 23936 1
Ovary - 23938	Ovary - 241181
1	1
Ovary - 241187	Ovary - 241196
1	1
Ovary - 241198	Ovary - 241199
1	1
Ovary - 242929	(Other)

1 105 sample_type: tumor 204 histological_type: endo clearcell mucinous other 11 9 28 59 ser undifferentiated NA's 85 2 10 primarysite: other ov NA's 23 178 3 summarygrade: high low NA's 91 31 82 summarystage: early late NA's 37 87 80 tumorstage: 1 2 3 4 NA's 20 14 58 18 94 substage: a b c NA's 17 22 79 86 grade: 1 2 3 4 NA's 11 20 83 8 82 age_at_initial_pathologic_diagnosis: Min. 1st Qu. Median Mean 3rd Qu. Max. 25.00 45.00 55.00 58.82 65.00 85.00 batch: 2004-12-03 2004-12-04 2004-12-07 2005-02-11 2005-03-03 2005-03-10 2005-03-11 3 3 1 1 1 1 1 2005-03-15 2005-03-16 2005-03-17 2005-03-19 2005-03-22 2005-04-13 2005-04-26 3 1 2 4 2 1 5 2005-04-29 2005-05-10 2005-06-01 2005-06-03 2005-06-08 2005-06-17 2005-08-05 2 2 5 3 3 6 3 2005-08-09 2005-08-11 2005-09-07 2005-09-09 2005-09-13 2005-11-02 2005-11-04

6 1 3 3 6 3 1 2005-11-15 2005-11-18 2005-12-02 2006-01-24 2006-01-26 2006-02-07 2006-02-28 3 1 4 2 1 1 1 2006-03-06 2006-03-14 2006-04-18 2006-04-20 2006-05-16 2006-06-08 2006-07-26 2 2 1 2 3 1 2 2006-07-28 2006-09-12 2006-09-14 2006-10-10 2006-10-24 2006-10-31 2006-11-09 1 2 1 1 9 5 10 2006-11-21 2006-11-30 2006-12-07 2007-01-12 2007-02-09 2007-03-07 2007-03-09 1 6 3 1 1 8 1 2007-03-15 2007-05-01 2007-05-03 2007-05-15 2007-05-18 2007-05-30 2007-06-12 4 2 3 4 2 2 1 2007-07-27 2007-09-05 2007-09-07 2007-09-11 2007-09-12 2008-02-15 2008-02-21 2 3 1 4 4 1 3 2008-02-27 2008-03-04 2008-05-13 2008-05-16 2008-05-23 2 1 4 4 5

uncurated_author_metadata:

title: Omentum -

title: Ovary - 170809///geo_accession: GSM137917///status: Public on Sep 28 2006///

```
duplicates:
GSE2109.GSE2109_GSM76554 GSE2109.GSE2109_GSM76567 NA's
1 1 202
```

Value

An expression set

GSE26193

miR-141 and miR-200a act on ovarian tumorigenesis by controlling oxidative stress response.

Description

Although there is evidence that redox regulation has an essential role in malignancies, its impact on tumor prognosis remains unclear. Here we show crosstalk between oxidative stress and the miR-200 family of microRNAs that affects tumorigenesis and chemosensitivity. miR-141 and miR-200a target p38?? and modulate the oxidative stress response. Enhanced expression of these microR-NAs mimics p38?? deficiency and increases tumor growth in mouse models, but it also improves

the response to chemotherapeutic agents. High-grade human ovarian adenocarcinomas that accumulate miR-200a have low concentrations of p38?? and an associated oxidative stress signature. The miR200a-dependent stress signature correlates with improved survival of patients in response to treatment. Therefore, the role of miR-200a in stress could be a predictive marker for clinical outcome in ovarian cancer. In addition, although oxidative stress promotes tumor growth, it also sensitizes tumors to treatment, which could account for the limited success of antioxidants in clinical trials.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Mateescu B, Batista L, Mariani O, Meyniel J, Cottu PH, Sastre-
  Laboratory: Mateescu, Mechta-Grigoriou 2011
  Contact information:
  Title: miR-141 and miR-200a act on ovarian tumorigenesis by controlling oxidative
  URL:
  PMIDs: 22101765
  Abstract: A 149 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [HG-U133_Plus_2] Affymetrix Human Genome U133 Plus 2.0 Array
   platform_shorttitle:
      Affymetrix HG-U133Plus2
   platform_summary:
      hgu133plus2
   platform_manufacturer:
      Affymetrix
   platform_distribution:
      commercial
   platform_accession:
      GPL570
   platform technology:
      in situ oligonucleotide
   version:
      2015-09-22 19:44:56
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (42447 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 42447 features, 107 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
```

n events median 0.95LCL 0.95UCL 107.00 76.00 3.05 2.50 4.56

```
-----
```

Available sample meta-data:

alt_sample_name:

alt_sample_name:		
Ovarian carcinoma 1	Ovarian carcinoma 10 Ovarian	carcinoma 100
1	1	1
Ovarian carcinoma 101	Ovarian carcinoma 102 Ovarian	carcinoma 103
1	1	1
Ovarian carcinoma 104	Ovarian carcinoma 105 Ovarian	carcinoma 106
1	1	1
Ovarian carcinoma 107	Ovarian carcinoma 11 Ovaria	n carcinoma 12
1	1	1
Ovarian carcinoma 13	Ovarian carcinoma 14 Ovaria	n carcinoma 15
1	1	1
Ovarian carcinoma 16	Ovarian carcinoma 17 Ovaria	n carcinoma 18
1	1	1
Ovarian carcinoma 19	Ovarian carcinoma 2 Ovaria	n carcinoma 20
1	1	1
Ovarian carcinoma 21	Ovarian carcinoma 22 Ovaria	n carcinoma 23
1	1	1
Ovarian carcinoma 24	Ovarian carcinoma 25 Ovaria	n carcinoma 26
1	1	1
Ovarian carcinoma 27	Ovarian carcinoma 28 Ovaria	n carcinoma 29
1	1	1
Ovarian carcinoma 3	Ovarian carcinoma 30 Ovaria	n carcinoma 31
1	1	1
Ovarian carcinoma 32	Ovarian carcinoma 33 Ovaria	n carcinoma 34
1	1	1
Ovarian carcinoma 35	Ovarian carcinoma 36 Ovaria	n carcinoma 37
1	1	1
Ovarian carcinoma 38	Ovarian carcinoma 39 Ovaria	an carcinoma 4
1	1	1
Ovarian carcinoma 40	Ovarian carcinoma 41 Ovaria	n carcinoma 42
1	1	1
Ovarian carcinoma 43	Ovarian carcinoma 44 Ovaria	n carcinoma 45
1	1	1
Ovarian carcinoma 46	Ovarian carcinoma 47 Ovaria	n carcinoma 48
1	1	1

Ovarian carcinoma 49	Ovarian carcinoma 5	Ovarian carcinoma 50
1	1	1
- Ovarian carcinoma 51 1	Ovarian carcinoma 52	Ovarian carcinoma 53 1
- Ovarian carcinoma 54 1	Ovarian carcinoma 55 1	Ovarian carcinoma 56 1
- Ovarian carcinoma 57 1	Ovarian carcinoma 58 1	Ovarian carcinoma 59 1
- Ovarian carcinoma 6 1	Ovarian carcinoma 60 1	Ovarian carcinoma 61 1
Ovarian carcinoma 62	Ovarian carcinoma 63	Ovarian carcinoma 64
1	1	1
- Ovarian carcinoma 65 1	Ovarian carcinoma 66 1	Ovarian carcinoma 67 1
Ovarian carcinoma 68	Ovarian carcinoma 69	Ovarian carcinoma 7
1	1	1
Ovarian carcinoma 70	Ovarian carcinoma 71	Ovarian carcinoma 72
1	1	1
Ovarian carcinoma 73	Ovarian carcinoma 74	Ovarian carcinoma 75
1	1	1
Ovarian carcinoma 76	Ovarian carcinoma 77	Ovarian carcinoma 78
1	1	1
Ovarian carcinoma 79	Ovarian carcinoma 8	Ovarian carcinoma 80
1	1	1
Ovarian carcinoma 81	Ovarian carcinoma 82	Ovarian carcinoma 83
1	1	1
Ovarian carcinoma 84	Ovarian carcinoma 85	Ovarian carcinoma 86
1	1	1
Ovarian carcinoma 87	Ovarian carcinoma 88	Ovarian carcinoma 89
1	1	1
Ovarian carcinoma 9	Ovarian carcinoma 90	Ovarian carcinoma 91
1	1	1
(Other) 8		
sample_type: tumor 107		
histological_type: clearcell endo 6 8	mucinous other 8 6	ser 79
summarygrade: high low 67 40		

summarystage:

```
early late
  31 76
tumorstage:
1 2 3 4
20 11 59 17
substage:
 a b cNA's
 16 12 62 17
grade:
1 2 3
7 33 67
days_to_tumor_recurrence:
  Min. 1st Qu. Median Mean 3rd Qu.
                                     Max.
   3.0 340.5 584.0 1108.0 1525.0 7386.0
recurrence_status:
norecurrence recurrence
        27
                   80
days_to_death:
  Min. 1st Qu. Median Mean 3rd Qu. Max.
     3 668 1096 1520 2220
                                     7386
vital_status:
deceased living
    76 31
batch:
2006-06-01 2006-06-27 2006-06-28 2006-06-29 2006-06-30 2006-07-20 2008-03-06
      15 14
                      23 16 21
                                                  3
                                                             1
2009-03-18 2009-03-19
       4
            10
uncurated_author_metadata:
     title: Ovarian carcinoma 100///geo_accession: GSM643032///status: Public on M
          title: Ovarian carcinoma 101///geo_accession: GSM643033///status: Publi
     title: Ovarian carcinoma 102///geo_accession: GSM643034///status: Public on M
         title: Ovarian carcinoma 103///geo_accession: GSM643035///status: Public
         title: Ovarian carcinoma 104///geo_accession: GSM643036///status: Public
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title: Ovarian carcinoma 105///geo_accession: GSM643037///status: Public title: Ovarian carcinoma 106///geo_accession: GSM643038///status: Public or title: Ovarian carcinoma 107///geo_accession: GSM643039///status: Public on Nov 01 title: Ovarian carcinoma 10///geo_accession: GSM642942///status: Public on title: Ovarian carcinoma 11///geo_accession: GSM642943///status: Public title: Ovarian carcinoma 12///geo_accession: GSM642944///status: Public title: Ovarian carcinoma 13///geo_accession: GSM642945///status: Publi title: Ovarian carcinoma 14///geo_accession: GSM642946///status: Public title: Ovarian carcinoma 15///geo_accession: GSM642947///status: Public title: Ovarian carcinoma 16///geo_accession: GSM642948///status: Public title: Ovarian carcinoma 17///geo_accession: GSM642949///status: Public of title: Ovarian carcinoma 18///geo_accession: GSM642950///status: Public on No title: Ovarian carcinoma 19///geo_accession: GSM642951///status: Public title: Ovarian carcinoma 1///geo_accession: GSM642933///status: Pub title: Ovarian carcinoma 20///geo_accession: GSM642952///status: Public on Nov (title: Ovarian carcinoma 21///geo_accession: GSM642953///status: Public title: Ovarian carcinoma 22///geo_accession: GSM642954///status: Public title: Ovarian carcinoma 23///geo_accession: GSM642955///status: Public title: Ovarian carcinoma 24///geo_accession: GSM642956///status: Public of title: Ovarian carcinoma 25///geo_accession: GSM642957///status: Public of title: Ovarian carcinoma 26///geo_accession: GSM642958///status: Public of title: Ovarian carcinoma 27///geo_accession: GSM642959///status: Public title: Ovarian carcinoma 28///geo_accession: GSM642960///status: Public of

title: Ovarian carcinoma 29///geo_accession: GSM642961///status: Public of

title: Ovarian carcinoma 2///geo_accession: GSM642934///status: Public on title: Ovarian carcinoma 30///geo_accession: GSM642962///status: Public on No title: Ovarian carcinoma 31///geo_accession: GSM642963///status: Public title: Ovarian carcinoma 32///geo_accession: GSM642964///status: Public of title: Ovarian carcinoma 33///geo_accession: GSM642965///status: Public on Nov title: Ovarian carcinoma 34///geo_accession: GSM642966///status: Public title: Ovarian carcinoma 35///geo_accession: GSM642967///status: Publ title: Ovarian carcinoma 36///geo_accession: GSM642968///status: Public title: Ovarian carcinoma 37///geo_accession: GSM642969///status: Public title: Ovarian carcinoma 38///geo_accession: GSM642970///status: Public title: Ovarian carcinoma 39///geo_accession: GSM642971///status: Public or title: Ovarian carcinoma 3///geo_accession: GSM642935///status: Public title: Ovarian carcinoma 40///geo_accession: GSM642972///status: Public title: Ovarian carcinoma 41///geo_accession: GSM642973///status: Public title: Ovarian carcinoma 42///geo_accession: GSM642974///status: Public of title: Ovarian carcinoma 43///geo_accession: GSM642975///status: Public or title: Ovarian carcinoma 44///geo_accession: GSM642976///status: Public title: Ovarian carcinoma 45///geo_accession: GSM642977///status: Public title: Ovarian carcinoma 46///geo_accession: GSM642978///status: Public title: Ovarian carcinoma 47///geo_accession: GSM642979///status: Public title: Ovarian carcinoma 48///geo_accession: GSM642980///status: Publ title: Ovarian carcinoma 49///geo_accession: GSM642981///status: Public title: Ovarian carcinoma 4///geo_accession: GSM642936///status: Pub title: Ovarian carcinoma 50///geo_accession: GSM642982///status: Public on Nov

title: Ovarian carcinoma 51///geo_accession: GSM642983///status: Public of title: Ovarian carcinoma 52///geo_accession: GSM642984///status: Publ title: Ovarian carcinoma 53///geo_accession: GSM642985///status: Public title: Ovarian carcinoma 54///geo_accession: GSM642986///status: Public on M title: Ovarian carcinoma 55///geo_accession: GSM642987///status: Public title: Ovarian carcinoma 56///geo_accession: GSM642988///status: Public title: Ovarian carcinoma 57///geo_accession: GSM642989///status: Public title: Ovarian carcinoma 58///geo_accession: GSM642990///status: Public on Nov title: Ovarian carcinoma 59///geo_accession: GSM642991///status: Public of title: Ovarian carcinoma 5///geo_accession: GSM642937///status: Public title: Ovarian carcinoma 60///geo_accession: GSM642992///status: Public title: Ovarian carcinoma 61///geo_accession: GSM642993///status: Public on M title: Ovarian carcinoma 62///geo_accession: GSM642994///status: Publi title: Ovarian carcinoma 63///geo_accession: GSM642995///status: Public of title: Ovarian carcinoma 64///geo_accession: GSM642996///status: Public or title: Ovarian carcinoma 65///geo_accession: GSM642997///status: Public on title: Ovarian carcinoma 66///geo_accession: GSM642998///status: Public of title: Ovarian carcinoma 67///geo_accession: GSM642999///status: Public title: Ovarian carcinoma 68///geo_accession: GSM643000///status: Public title: Ovarian carcinoma 69///geo_accession: GSM643001///status: Public on No title: Ovarian carcinoma 6///geo_accession: GSM642938///status: Public title: Ovarian carcinoma 70///geo_accession: GSM643002///status: Public title: Ovarian carcinoma 71///geo_accession: GSM643003///status: Public on Nov title: Ovarian carcinoma 72///geo_accession: GSM643004///status: Public on Nov 01 2

title: Ovarian carcinoma 73///geo_accession: GSM643005///status: Public on No title: Ovarian carcinoma 74///geo_accession: GSM643006///status: Public title: Ovarian carcinoma 75///geo_accession: GSM643007///status: Public title: Ovarian carcinoma 76///geo_accession: GSM643008///status: Public of title: Ovarian carcinoma 77///geo_accession: GSM643009///status: Public of title: Ovarian carcinoma 78///geo_accession: GSM643010///status: Public on title: Ovarian carcinoma 79///geo_accession: GSM643011///status: Public on No title: Ovarian carcinoma 7///geo_accession: GSM642939///status: Public title: Ovarian carcinoma 80///geo_accession: GSM643012///status: Publ title: Ovarian carcinoma 81///geo_accession: GSM643013///status: Public on No title: Ovarian carcinoma 82///geo_accession: GSM643014///status: Public title: Ovarian carcinoma 83///geo_accession: GSM643015///status: Public of title: Ovarian carcinoma 84///geo_accession: GSM643016///status: Public of title: Ovarian carcinoma 85///geo_accession: GSM643017///status: Public of title: Ovarian carcinoma 86///geo_accession: GSM643018///status: Public or title: Ovarian carcinoma 87///geo_accession: GSM643019///status: Public title: Ovarian carcinoma 88///geo_accession: GSM643020///status: Public of title: Ovarian carcinoma 89///geo_accession: GSM643021///status: Public on title: Ovarian carcinoma 8///geo_accession: GSM642940///status: Pub title: Ovarian carcinoma 90///geo_accession: GSM643022///status: Pub title: Ovarian carcinoma 91///geo_accession: GSM643023///status: Public on Nov title: Ovarian carcinoma 92///geo_accession: GSM643024///status: Public of

Value

An expression set

GSE26712

A gene signature predicting for survival in suboptimally debulked patients with ovarian cancer.

Description

Despite the existence of morphologically indistinguishable disease, patients with advanced ovarian tumors display a broad range of survival end points. We hypothesize that gene expression profiling can identify a prognostic signature accounting for these distinct clinical outcomes. To resolve survival-associated loci, gene expression profiling was completed for an extensive set of 185 (90 optimal/95 suboptimal) primary ovarian tumors using the Affymetrix human U133A microarray. Cox regression analysis identified probe sets associated with survival in optimally and suboptimally debulked tumor sets at a P value of <0.01. Leave-one-out cross-validation was applied to each tumor cohort and confirmed by a permutation test. External validation was conducted by applying the gene signature to a publicly available array database of expression profiles of advanced stage suboptimally debulked tumors. The prognostic signature successfully classified the tumors according to survival for suboptimally (P = 0.0179) but not optimally debulked (P = 0.144) patients. The suboptimal gene signature was validated using the independent set of tumors (odds ratio, 8.75; P = 0.0146). To elucidate signaling events amenable to the appendix intervention in suboptimally debulked patients, pathway analysis was completed for the top 57 survival-associated probe sets. For suboptimally debulked patients, confirmation of the predictive gene signature supports the existence of a clinically relevant predictor, as well as the possibility of novel therapeutic opportunities. Ultimately, the prognostic classifier defined for suboptimally debulked tumors may aid in the classification and enhancement of patient outcome for this high-risk population.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Bonome T, Levine DA, Shih J, Randonovich M, Pise-Masison CA, H
  Laboratory: Bonome, Birrer 2008
  Contact information:
  Title: A gene signature predicting for survival in suboptimally debulked patients
  URL:
  PMIDs: 18593951
 Abstract: A 238 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [HG-U133A] Affymetrix Human Genome U133A Array
  platform_shorttitle:
      Affymetrix HG-U133A
  platform_summary:
```

```
hgu133a
platform_manufacturer:
    Affymetrix
platform_distribution:
    commercial
platform_accession:
    GPL96
version:
    2015-09-22 19:46:24
featureData(eset):
An object of class 'AnnotatedDataFrame'
featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (20967 total)
varLabels: probeset gene EntrezGene.ID best_probe
varMetadata: labelDescription
```

Details

```
assayData: 20967 features, 195 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
10 observations deleted due to missingness
    n events median 0.95LCL 0.95UCL
185.00 129.00 3.83 3.24 4.83
```

```
_____
```

```
Available sample meta-data:
```

alt_sample_name:							
Normal HOSE2008	Normal HOSE2061	Normal HOSE2064					
1	1	1					
Normal HOSE2085	Normal HOSE2225	Normal HOSE2226					
1	1	1					
Normal HOSE2228	Normal HOSE2230	Normal HOSE2234					
1	1	1					
Normal HOSE2237	Ovarian Cancer SO10	Ovarian Cancer SO100					
1	1	1					
Ovarian Cancer SO103	Ovarian Cancer SO106	Ovarian Cancer SO108					
1	1	1					
Ovarian Cancer SO11	Ovarian Cancer SO113	Ovarian Cancer SO115					
1	1	1					
Ovarian Cancer SO116	Ovarian Cancer SO117	Ovarian Cancer SO118					
1	1	1					
Ovarian Cancer SO12	Ovarian Cancer SO121	Ovarian Cancer SO122					

1	1	1
Ovarian Cancer SO124	Ovarian Cancer SO129	Ovarian Cancer SO13 1
_	Ovarian Cancer SO134	Ovarian Cancer SO135
Ovarian Cancer SO137	Ovarian Cancer SO141	
1 Ovarian Cancer SO148	1 Ovarian Cancer SO154	1 Ovarian Cancer SO16
1 Ovarian Cancer SO166		1 Ovarian Cancer SO173
1 Ovarian Cancer SO174		1 Ovarian Cancer SO181
1 Ovarian Cancer SO184	1 Ovarian Cancer SO185	1 Ovarian Cancer SO187
1 Ovarian Cancer SO189	1 Ovarian Cancer SO190	1 Ovarian Cancer SO193
1 Ovarian Cancer SO194	1 Ovarian Cancer SO196	1 Ovarian Cancer SO197
1 Ovarian Cancer SO2	1 Ovarian Cancer SO200	1 Ovarian Cancer SO201
1 UVALIAN CANCEL 502	1	1
Ovarian Cancer SO203	Ovarian Cancer SO205	Ovarian Cancer SO21 1
Ovarian Cancer SO211	Ovarian Cancer SO214	Ovarian Cancer SO216
Ovarian Cancer SO217	Ovarian Cancer SO218	-
Ovarian Cancer SO225	Ovarian Cancer SO227	Ovarian Cancer SO228
1 Ovarian Cancer SO229	_	1 Ovarian Cancer SO230
1 Ovarian Cancer SO231	1 Ovarian Cancer SO235	1 Ovarian Cancer SO236
1 Ovarian Cancer SO237	1 Ovarian Cancer SO241	1 Ovarian Cancer SO242
1 Ovarian Cancer SO243	1 Ovarian Cancer SO244	1 Ovarian Cancer SO246
1 Ovarian Cancer SO247	1 Ovarian Cancer SO249	1 Ovarian Cancer SO25
1	1 Ovarian Cancer SO256	1
1	1	1
Ovarian Cancer SO258 1	Ovarian Cancer SO261	Ovarian Cancer SO262 1
Ovarian Cancer SO263	Ovarian Cancer SO265	Ovarian Cancer SO267 1
Ovarian Cancer SO268	Ovarian Cancer SO272	-
Ovarian Cancer SO278	Ovarian Cancer SO279	Ovarian Cancer SO282

```
1
                                  1
                                                    1
Ovarian Cancer SO283 Ovarian Cancer SO285 Ovarian Cancer SO290
            1
                          1
                                                 1
           (Other)
              96
sample_type:
healthy tumor
10 185
histological_type:
ser NA's
185 10
primarysite:
ov
195
summarygrade:
high NA's
185 10
summarystage:
late NA's
185 10
tumorstage:
  3 4 NA's
146 36 13
substage:
 b c NA's
  9 137 49
age_at_initial_pathologic_diagnosis:
  Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
 26.00 52.00 63.00 61.54 70.00 84.00
                                           13
recurrence_status:
norecurrence recurrence
       42
             153
days_to_death:
  Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
  21.9 660.6 1164.0 1429.0 1880.0 4982.0 10
vital_status:
deceased living NA's
```

129 56 10 debulking: optimal suboptimal NA's 90 95 10 percent_normal_cells: 20-195 percent_stromal_cells: 20-195 percent_tumor_cells: 80+ 195 batch: 2003-11-04 2003-11-05 2003-11-06 2003-11-07 2003-11-20 2003-11-21 2003-12-16 14 16 9 6 10 15 17 2003-12-23 2003-12-24 2004-04-20 2004-04-21 2004-04-27 2004-09-28 2005-07-27 11 20 17 9 15 12 14 2006-11-09 10 uncurated_author_metadata: title: Norma title: Ovarian Cancer SO100///geo_accession: GSM657530///status: Public on Jan 20

title: Ovarian Cancer SO103///geo_accession: GSM657531///status: Public on Jan 20 title: Ovarian Cancer SO106///geo_accession: GSM657532///status: Public on title: Ovarian Cancer SO108///geo_accession: GSM657533///status: Public on Jar title: Ovarian Cancer SO10///geo_accession: GSM657529///status: Public on Ja title: Ovarian Cancer SO113///geo_accession: GSM657535///status: Public on Jan 20 title: Ovarian Cancer SO115///geo_accession: GSM657536///status: Public on title: Ovarian Cancer SO116///geo_accession: GSM657537///status: Public on Jar title: Ovarian Cancer SO117///geo_accession: GSM657538///status: Public on Jar title: Ovarian Cancer SO118///geo_accession: GSM657539///status: Public on Jan 20 title: Ovarian Cancer SO11///geo_accession: GSM657534///status: Public or title: Ovarian Cancer SO121///geo_accession: GSM657541///status: Public on Jan 20 title: Ovarian Cancer SO122///geo_accession: GSM657542///status: Public or title: Ovarian Cancer SO124///geo_accession: GSM657543///status: Public on Jan 20 title: Ovarian Cancer SO129///geo_accession: GSM657544///status: Public on Jan 20 title: Ovarian Cancer SO12///geo_accession: GSM657540///status: Public or title: Ovarian Cancer SO131///geo_accession: GSM657546///status: Public on Jan 20 title: Ovarian Cancer SO134///geo_accession: GSM657547///status: Public on title: Ovarian Cancer SO135///geo_accession: GSM657548///status: Public on title: Ovarian Cancer S0137///geo_accession: GSM657549///status: Public on Jar title: Ovarian Cancer SO13///geo_accession: GSM657545///status: Public on Jan 20 20 title: Ovarian Cancer SO141///geo_accession: GSM657550///status: Public on Jar title: Ovarian Cancer S0143///geo_accession: GSM657551///status: Public on Jar title: Ovarian Cancer S0148///geo_accession: GSM657552///status: Public on Jar title: Ovarian Cancer S0154///geo_accession: GSM657553///status: Public on Jan 2

title: Ovarian Cancer SO166///geo_accession: GSM657555///status: Public on Jan 20 title: Ovarian Cancer SO16///geo_accession: GSM657554///status: Public on Ja title: Ovarian Cancer S0173///geo_accession: GSM657557///status: Public on Jar title: Ovarian Cancer S0174///geo_accession: GSM657558///status: Public on Jar title: Ovarian Cancer SO17///geo_accession: GSM657556///status: Public on Ja title: Ovarian Cancer SO181///geo_accession: GSM657560///status: Public on Jan 20 title: Ovarian Cancer SO184///geo_accession: GSM657561///status: Public on Jan 20 title: Ovarian Cancer S0185///geo_accession: GSM657562///status: Public on Jan 20 title: Ovarian Cancer SO187///geo_accession: GSM657563///status: Public on Jar title: Ovarian Cancer SO189///geo_accession: GSM657564///status: Public on Jan 20 title: Ovarian Cancer SO18///geo_accession: GSM657559///status: Public of title: Ovarian Cancer S0190///geo_accession: GSM657565///status: Public on title: Ovarian Cancer S0193///geo_accession: GSM657566///status: Public on Jar title: Ovarian Cancer S0194///geo_accession: GSM657567///status: Public on Jar title: Ovarian Cancer SO196///geo_accession: GSM657568///status: Public on title: Ovarian Cancer S0197///geo_accession: GSM657569///status: Public on Jar title: Ovarian Cancer SO200///geo_accession: GSM657571///status: Public on Jar title: Ovarian Cancer SO201///geo_accession: GSM657572///status: Public on Jar title: Ovarian Cancer SO203///geo_accession: GSM657573///status: Public on title: Ovarian Cancer SO205///geo_accession: GSM657574///status: Public on Jan 20 title: Ovarian Cancer SO211///geo_accession: GSM657576///status: Public on Jar title: Ovarian Cancer SO214///geo_accession: GSM657577///status: Public on Jar title: Ovarian Cancer SO216///geo_accession: GSM657578///status: Public on Jar title: Ovarian Cancer SO217///geo_accession: GSM657579///status: Public on Jar

title: Ovarian Cancer SO218///geo_accession: GSM657580///status: Public or title: Ovarian Cancer SO21///geo_accession: GSM657575///status: Public or title: Ovarian Cancer SO224///geo_accession: GSM657581///status: Public on Jar title: Ovarian Cancer SO225///geo_accession: GSM657582///status: Public or title: Ovarian Cancer SO227///geo_accession: GSM657583///status: Public on Jar title: Ovarian Cancer SO228///geo_accession: GSM657584///status: Public on title: Ovarian Cancer SO229///geo_accession: GSM657585///status: Public on Jar title: Ovarian Cancer SO230///geo_accession: GSM657587///status: Public on title: Ovarian Cancer SO231///geo_accession: GSM657588///status: Public on title: Ovarian Cancer SO235///geo_accession: GSM657589///status: Public on Jar title: Ovarian Cancer S0236///geo_accession: GSM657590///status: Public on Jan 20 title: Ovarian Cancer S0237///geo_accession: GSM657591///status: Public on Jar title: Ovarian Cancer SO23///geo_accession: GSM657586///status: Public on Jan 20 title: Ovarian Cancer SO241///geo_accession: GSM657592///status: Public on Jar title: Ovarian Cancer SO242///geo_accession: GSM657593///status: Public on Jar title: Ovarian Cancer SO243///geo_accession: GSM657594///status: Public on Jan 20 title: Ovarian Cancer SO244///geo_accession: GSM657595///status: Public on Jar title: Ovarian Cancer SO246///geo_accession: GSM657596///status: Public on title: Ovarian Cancer SO247///geo_accession: GSM657597///status: Public on Jar title: Ovarian Cancer SO249///geo_accession: GSM657598///status: Public on Jar title: Ovarian Cancer S0250///geo_accession: GSM657600///status: Public on Jar title: Ovarian Cancer SO256///geo_accession: GSM657601///status: Public on Jar title: Ovarian Cancer S0257///geo_accession: GSM657602///status: Public on Ja title: Ovarian Cancer SO258///geo_accession: GSM657603///status: Public on Jar

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```
duplicates:

GSE26712.GSE26712_GSM657526

1

GSE26712.GSE26712_GSM657526///GSE26712.GSE26712_GSM657527

1

GSE26712.GSE26712_GSM657527

1

NA's

192
```

Value

An expression set

GSE30009

Multidrug resistance-linked gene signature predicts overall survival of patients with primary ovarian serous carcinoma.

Description

This study assesses the ability of multidrug resistance (MDR)-associated gene expression patterns to predict survival in patients with newly diagnosed carcinoma of the ovary. The scope of this research differs substantially from that of previous reports, as a very large set of genes was evaluated whose expression has been shown to affect response to chemotherapy. We applied a customized TaqMan low density array, a highly sensitive and specific assay, to study the expression profiles of 380 MDR-linked genes in 80 tumor specimens collected at initial surgery to debulk primary serous carcinoma. The RNA expression profiles of these drug resistance genes were correlated with clinical outcomes.Leave-one-out cross-validation was used to estimate the ability of MDR gene expression to predict survival. Although gene expression alone does not predict overall survival (OS; P = 0.06), four covariates (age, stage, CA125 level, and surgical debulking) do (P = 0.03). When gene expression was added to the covariates, we found an 11-gene signature that provides a major improvement in OS prediction (log-rank statistic P < 0.003). The predictive power of this 11-gene signature was confirmed by dividing high- and low-risk patient groups, as defined by their clinical covariates, into four specific risk groups on the basis of expression levels. This study reveals an 11-gene signature that allows a more precise prognosis for patients with serous cancer of the ovary treated with carboplatin- and paclitaxel-based therapy. These 11 new targets offer opportunities for new therapies to improve clinical outcome in ovarian cancer.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Gillet JP, Calcagno AM, Varma S, Davidson B et al. Multidrug
  Laboratory: Gillet, Gottesman 2012
  Contact information:
  Title: Multidrug resistance-linked gene signature predicts overall survival of pa
  URL:
  PMIDs: 22492981
 Abstract: A 244 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      TaqMan qRT-PCR Homo sapiens Low-Density Array 380
  platform_shorttitle:
      TaqMan qRT-PCR
   platform_summary:
```

```
NA

platform_manufacturer:

TaqMan

platform_distribution:

custom

platform_accession:

GPL13728

version:

2015-09-22 19:46:26

featureData(eset):

An object of class 'AnnotatedDataFrame'

featureNames: 5 6 ... 380 (363 total)

varLabels: probeset gene EntrezGene.ID best_probe

varMetadata: labelDescription
```

Details

```
103.00 57.00 3.42 2.92 5.34
```

```
Available sample meta-data:
```

```
alt_sample_name:
```

Norwegian patient 1	Norwegian patient 10	Norwegian patient 11
1	1	1
Norwegian patient 12	Norwegian patient 13	Norwegian patient 14
1	1	1
Norwegian patient 15	Norwegian patient 16	Norwegian patient 17
1	1	1
Norwegian patient 18	Norwegian patient 19	Norwegian patient 2
1	1	1
Norwegian patient 20	Norwegian patient 21	Norwegian patient 22
1	1	1
Norwegian patient 23	Norwegian patient 3	Norwegian patient 4
1	1	1
Norwegian patient 5	Norwegian patient 6	Norwegian patient 7
1	1	1
Norwegian patient 8	Norwegian patient 9	US Patient 1
1	1	1
US Patient 10	US Patient 11	US Patient 12

1	1	1
US Patient 13	US Patient 14	US Patient 15
1	1	1
US Patient 16	US Patient 17	US Patient 18
1	1	1
US Patient 19	US Patient 2	US Patient 20
1	1	1
US Patient 21	US Patient 22	US Patient 23
1	1	1
US Patient 24	US Patient 25	US Patient 26
1	1	1
US Patient 27 1	US Patient 28	US Patient 29 1
US Patient 3	US Patient 30	US Patient 31
1	1	1
US Patient 32 1	US Patient 33	US Patient 34 1
US Patient 35	US Patient 36	US Patient 37
1	1	1
US Patient 38	US Patient 39	US Patient 4
1	1	1
US Patient 40	US Patient 41	US Patient 42
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US Patient 43	US Patient 44	US Patient 45
1	1	1
US Patient 46	US Patient 47	US Patient 48
1	1	1
US Patient 49	US Patient 5	US Patient 50
1	1	1
US Patient 51	US Patient 52	US Patient 53
1	1	1
US Patient 54	US Patient 55	US Patient 56
1	1	1
US Patient 57	US Patient 58	US Patient 59
1	1	1
US Patient 6	US Patient 60	US Patient 61
1	1	1
US Patient 62	US Patient 63	US Patient 64
1	1	1
US Patient 65	US Patient 66	US Patient 67
1	1	1
US Patient 68	US Patient 69	US Patient 7
1	1	1
US Patient 70 1	US Patient 71 1	US Patient 72
US Patient 73	US Patient 74 1	US Patient 75 1
US Patient 76	US Patient 77	US Patient 78

```
1
                                 1
           (Other)
               4
sample_type:
tumor
 103
histological_type:
clearcell ser
  1
            102
summarygrade:
high low NA's
 92 9 2
summarystage:
late
103
tumorstage:
3 4
82 21
substage:
  b c NA's
  2 60 41
grade:
  1 2 3 NA's
  4 5 92 2
age_at_initial_pathologic_diagnosis:
  Min. 1st Qu. Median Mean 3rd Qu. Max.
 30.00 56.00 61.00
                      62.45 71.50
                                   87.00
days_to_death:
  Min. 1st Qu. Median Mean 3rd Qu.
                                   Max.
    24 598 1053 1156 1568
                                    4748
vital_status:
deceased living
    57
        46
debulking:
  optimal suboptimal
      81 22
```

95

uncurated_author_metadata:

title: US Pati

title: US

title: US Patient 5

title: US Patient 51///geo_accession: GSM742615///status: Public on Apr 19 2012///s

title: US Patient 54///geo_accession: GSM7426

tit

title: US Patient 57///geo_accession: GSM742621///status: Public of

title: US Patient 59///geo_accession: GSM742623///status: Public of

title: US Patient 63///geo_accessio

title: US Patient

title: US Patient 66///geo_accession: GSM742630///status

ti

title: US Patient 70///geo_accession: GSM742634///status: Public on Apr 19 20

t

title: US Patier

title: US Patient 75///geo_accession: GSM7426

title:

title: US Patient 77///geo_ad

title: US Patient 78//,

title: US Patient 79///g

Value

An expression set

GSE30161

Multi-gene expression predictors of single drug responses to adjuvant chemotherapy in ovarian carcinoma: predicting platinum resistance.

Description

Despite advances in radical surgery and chemotherapy delivery, ovarian cancer is the most lethal gynecologic malignancy. Standard therapy includes treatment with platinum-based combination chemotherapies yet there is no biomarker model to predict their responses to these agents. We here have developed and independently tested our multi-gene molecular predictors for forecasting patients' responses to individual drugs on a cohort of 55 ovarian cancer patients. To independently validate these molecular predictors, we performed microarray profiling on FFPE tumor samples of 55 ovarian cancer patients (UVA-55) treated with platinum-based adjuvant chemotherapy. Genomewide chemosensitivity biomarkers were initially discovered from the in vitro drug activities and genomic expression data for carboplatin and paclitaxel, respectively. Multivariate predictors were trained with the cell line data and then evaluated with a historical patient cohort. For the UVA-55 cohort, the carboplatin, taxol, and combination predictors significantly stratified responder patients and non-responder patients (p = 0.019, 0.04, 0.014) with sensitivity = 91%, 96%, 93 and NPV = 57%, 67%, 67% in pathologic clinical response. The combination predictor also demonstrated a significant survival difference between predicted responders and non-responders with a median survival of 55.4 months vs. 32.1 months. Thus, COXEN single- and combination-drug predictors successfully stratified platinum resistance and taxane response in an independent cohort of ovarian cancer patients based on their FFPE tumor samples.

Format

```
experimentData(eset):
Experiment data
Experimenter name: Ferriss JS, Kim Y, Duska L, Birrer M, Levine DA, Moskaluk C,Th
Laboratory: Ferriss, Lee 2012
Contact information:
Title: Multi-gene expression predictors of single drug responses to adjuvant chem
```

```
URL:
  PMIDs: 22348014
  Abstract: A 215 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform title:
      [HG-U133_Plus_2] Affymetrix Human Genome U133 Plus 2.0 Array
   platform shorttitle:
      Affymetrix HG-U133Plus2
  platform_summary:
      hgu133plus2
   platform_manufacturer:
     Affymetrix
  platform_distribution:
      commercial
  platform_accession:
     GPL570
   version:
      2015-09-22 19:50:24
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (42447 total)
 varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 42447 features, 58 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
     n events median 0.95LCL 0.95UCL
 58.00 36.00 4.19 2.70 6.17
_____
Available sample meta-data:
_____
alt_sample_name:
OV_FFPE_1 OV_FFPE_10 OV_FFPE_11 OV_FFPE_12 OV_FFPE_13 OV_FFPE_14 OV_FFPE_15
      1 1 1 1 1 1
                                                       1
OV_FFPE_16 OV_FFPE_17 OV_FFPE_18 OV_FFPE_19 OV_FFPE_2 OV_FFPE_20 OV_FFPE_21
               1
                        1
                                  1
                                                   1
       1
                                          1
                                                            1
OV_FFPE_22 OV_FFPE_23 OV_FFPE_24 OV_FFPE_25 OV_FFPE_26 OV_FFPE_27 OV_FFPE_28
```

```
1
              1
                      1
                              1
                                      1
                                              1
                                                       1
OV_FFPE_29 OV_FFPE_3 OV_FFPE_30 OV_FFPE_31 OV_FFPE_32 OV_FFPE_33 OV_FFPE_34
     1 1 1 1 1 1 1
OV_FFPE_35 OV_FFPE_36 OV_FFPE_37 OV_FFPE_38 OV_FFPE_39 OV_FFPE_4 OV_FFPE_40
      1 1 1 1 1 1 1
OV_FFPE_41 OV_FFPE_42 OV_FFPE_43 OV_FFPE_44 OV_FFPE_45 OV_FFPE_46 OV_FFPE_47
                                      1 1 1
     1 1 1
                              1
OV_FFPE_48 OV_FFPE_49 OV_FFPE_5 OV_FFPE_50 OV_FFPE_51 OV_FFPE_52 OV_FFPE_53
     1 1 1 1 1 1 1
OV_FFPE_54 OV_FFPE_55 OV_FFPE_56 OV_FFPE_57 OV_FFPE_58 OV_FFPE_6 OV_FFPE_7
     1 1
                  1 1 1 1 1
OV_FFPE_8 OV_FFPE_9
      1
        1
sample_type:
tumor
  58
histological_type:
     clearcell
                     endo mucinous
                                             other
         5
                               1
                      1
                                              1
         ser undifferentiated
                                NA's
          47
                                   2
                      1
summarygrade:
high low NA's
33 21 4
summarystage:
late
 58
tumorstage:
3 4
53 5
substage:
a b c
9 11 38
grade:
  1 2 3 NA's
  2 19 33 4
age_at_initial_pathologic_diagnosis:
 Min. 1st Qu. Median Mean 3rd Qu. Max.
38.00 53.50 62.00 62.57 72.00 85.00
```

pltx: У 58 tax: n y 4 54 neo: n 58 days_to_tumor_recurrence: Min. 1st Qu. Median Mean 3rd Qu. Max. 12.0 255.2 386.0 742.1 768.2 4208.0 recurrence_status: norecurrence recurrence NA's 6 48 4 days_to_death: Min. 1st Qu. Median Mean 3rd Qu. Max. 49.0 585.2 1010.0 1375.0 2131.0 4208.0 vital_status: deceased living 36 22 debulking: optimal suboptimal NA's 2 26 30 batch: 2009-10-07 2009-10-08 2009-10-09 2009-10-20 28 18 8 4 uncurated_author_metadata: title: OV_FFPE_10///geo_accession: GSM746870///status: Public on Aug 21 title: OV_FFPE_11///geo_accession: GSM746871///status: Public on Aug 21 2012///su title: OV_FFPE_12///geo_accession: GSM746872///status: Public on Aug 21 2012/// title: OV_FFPE_13///geo_accession: GSM746873///status: Public on Aug 21 2012// title: OV_FFPE_14///geo_accession: GSM746874///status: Public on Aug 21 2012///s title: OV_FFPE_15///geo_accession: GSM746875///status: Pu

title: OV_FFPE_16///geo_accession: GSM746876///status: Public on Aug 21 2012/ title: OV_FFPE_17///geo_accession: GSM746877///status: Public on Aug 21 2012 title: OV_FFPE_18///geo_accession: GSM746

title: OV_FFPE_19///geo_

title: OV_FFPE_22///geo_accession: GSM7

title: OV_FFPE_23///geo_accession: GSM746883///status: Public on Aug 21 2 title: OV_FFPE_24///geo_accession: GSM746884///status: Public on Aug 21 201 title: OV_FFPE_25///geo_accession: GSM746885///status: Public on Aug 21 2012// title: OV_FFPE_26///geo_accession: GSM746886///status: Public on Aug 21 2012// title: OV_FFPE_27///geo_accession: GSM746887///status: Public on Aug 21 2012//s title: OV_FFPE_27///geo_accession: GSM746887///status: Public on Aug 21 2012//s

title: OV_FFPE_29///geo_accession: GSM746889///status: Public on Aug 21 2012//
title: OV_FFPE_2///geo_accession: GSM746862///status: Public on Aug 21 2012//
title: OV_FFPE_30///geo_accession: GSM746890///status: Public on Aug 21 2012/
title: OV_FFPE_31///geo_accession: GSM746891///status: Public on Aug 21 2012///subtitle: OV_FFPE_32///geo_accession: GSM746892///status: Public on Aug 21 2012///subtitle: OV_FFPE_33///geo_accession: GSM746893///status: Public on Aug 21 2012///subtitle: OV_FFPE_34///geo_accession: GSM746894///status: Public on Aug 21 2012///subtitle: OV_FFPE_35///geo_accession: GSM746895///status: Public on Aug 21 2012///subtitle: OV_FFPE_36///geo_accession: GSM746896///status: Public on Aug 21 2012///subtitle: OV_FFPE_37///geo_accession: GSM746897///status: Public OV_FFPE_37///geo_accession: GSM746897///status: Public OV_FFPE_37///geo_accession: GSM746896///status: GSM746897///status: Public OV_FFPE_30///geo_accession: GSM746896///status: Public OV_FFPE_30///geo_accession: GSM746896//status: Public OV_FFPE_30///geo_accession: G

title: OV_FFPE_38///geo_accession: GSM746898///status: Public on Aug 21 2012///sub title: OV_FFPE_39///geo_accession: GSM746899///status: Public on Aug 21 2012// title: OV_FFPE_3///geo_accession: GSM746863///status: Public on Aug 21 2012// title: OV_FFPE_40///geo_accession: GSM746900///status: Public on Aug 21 2012// title: OV_FFPE_40///geo_accession: GSM746900///status: Public on Aug 21 2012//

title: OV_FFPE_42///geo_accession: GSM746902///status: Public on Aug 21 2012///su title: OV_FFPE_43///geo_accession: GSM746903///statu

title: OV_FFPE_44///geo_accession: GSM746904///status: Public on Aug 21 2012///suk
title: OV_FFPE_45///geo_accession: GSM746905///status: Public on Aug 21 2012///suk
title: OV_FFPE_46///geo_accession: GSM746906///status: Public on Aug 21 2012//
title: OV_FFPE_47///geo_accession: GSM746907///status: Public on Aug 21 2012//
title: OV_FFPE_48///geo_accession: GSM746908///status: Public

title: OV_FFPE_49///geo_accession: GSM746909///status: Public on Aug 21 2012/ title: OV_FFPE_4///geo_accession: GSM746864///status: Public on Aug 21 2012/ title: OV_FFPE_50///geo_accession: GSM746910///status: Public on Aug 21 2012/ title: OV_FFPE_51///geo_accession: GSM746911///status: Public on Aug 21 201 title: OV_FFPE_52///geo_accession: GSM746912///status: Public on Aug 21 2012/ title: OV_FFPE_53///geo_accession: GSM746913///status: Public on Aug 21 2012// title: OV_FFPE_54///geo_accession: GSM746914///status: Public on Aug 21 2012// title: OV_FFPE_55///geo_accession: GSM746915///status: Public on Aug 21 2012// title: OV_FFPE_56///geo_accession: GSM746915///status: Public on Aug 21 2012// title: OV_FFPE_56///geo_accession: GSM746916///status: Public on Aug 21 2012//status: Public OV_FFPE_57///geo_accession: Public OV_FFPE_57///geo_accession: Publ

title: OV_FFPE_58///geo_accession: GSM746918///status: Public on Aug 21 2012//
title: OV_FFPE_5///geo_accession: GSM746865///status: Public on Aug 21 2012

title: OV_FFPE_6///geo_accession: GSM746866///status: Public on Aug 21 2012
title: OV_FFPE_7///geo_accession: GSM746867///status: Public on Aug 21 2012,
title: OV_FFPE_8///geo_accession: GSM746868///status: Public on Aug 21 2012/
title: OV_FFPE_9///geo_accession: GSM746869///status: Public on Aug 21 2012/

Value

An expression set

GSE32062

High-risk ovarian cancer based on 126-gene expression signature is uniquely characterized by downregulation of antigen presentation pathway.

Description

High-grade serous ovarian cancers are heterogeneous not only in terms of clinical outcome but also at the molecular level. Our aim was to establish a novel risk classification system based on a gene expression signature for predicting overall survival, leading to suggesting novel therapeutic strategies for high-risk patients. In this large-scale cross-platform study of six microarray data sets consisting of 1,054 ovarian cancer patients, we developed a gene expression signature for predicting overall survival by applying elastic net and 10-fold cross-validation to a Japanese data set A (n =260) and evaluated the signature in five other data sets. Subsequently, we investigated differences in the biological characteristics between high- and low-risk ovarian cancer groups. An elastic net analysis identified a 126-gene expression signature for predicting overall survival in patients with ovarian cancer using the Japanese data set A (multivariate analysis, P = 4 ?? 10(-20)). We validated its predictive ability with five other data sets using multivariate analysis (Tothill's data set, P =1 ?? 10(-5); Bonome's data set, P = 0.0033; Dressman's data set, P = 0.0016; TCGA data set, P = 0.0027; Japanese data set B, P = 0.021). Through gene ontology and pathway analyses, we identified a significant reduction in expression of immune-response-related genes, especially on the antigen presentation pathway, in high-risk ovarian cancer patients. This risk classification based on the 126-gene expression signature is an accurate predictor of clinical outcome in patients with advanced stage high-grade serous ovarian cancer and has the potential to develop new therapeutic strategies for high-grade serous ovarian cancer patients.

Format

```
experimentData(eset):
Experiment data
Experimenter name: Yoshihara K, Tsunoda T, Shigemizu D, Fujiwara H et al. High-ri
Laboratory: Yoshihara, Tanaka 2012
```

```
Contact information:
  Title: High-risk ovarian cancer based on 126-gene expression signature is unique
  URL:
  PMIDs: 22241791
 Abstract: A 255 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      Agilent-014850 Whole Human Genome Microarray 4x44K G4112F (Probe Name vers
ion)
  platform_shorttitle:
      Agilent G4112F
  platform_summary:
     hgug4112a
   platform_manufacturer:
      Agilent
  platform_distribution:
      commercial
  platform accession:
     GPL6480
   version:
      2015-09-22 19:55:29
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: A_23_P100001 A_23_P100011 ... A_32_P99902 (30936 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 30936 features, 260 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
    n events median 0.95LCL 0.95UCL
260.00 121.00 4.93 4.11 6.58
_____
Available sample meta-data:
_____
alt_sample_name:
             116d 117d 119d
   10d 115d
                               11d 120d 122d 123d 125Rd
        1
                                                  1
    1
              1
                    1
                          1
                                1
                                      1
                                            1
                                                       1
  129d 12d 130d 132d 134d 139d 140d 143d 144d 145d
```

1	1	1	1	1	1	1	1	1	1	
1 146d	1 148d	1 150d	1 155d	1 156d	1 15d	1 160d	1 16d	1 171d	1 173d	
1	1	1304	1	1300	150	1000	100	1	1,50	
174d	178d	17d	183d	184d	185d	186d	18d	20d	22d	
1	1	1	1	1010	1000	1	1	1	1	
23d	249d	257d	25d	260d	262d	264d	266d	267d	268d	
1	1	1	1	1	1	1	1	1	1	
269d	27d	299d	2d	300d	301d	302d	303d	304d	305d2	
1	1	1	1	1	1	1	1	1	1	
306d	307d	310d	318d	319d	320d2	323d	327d	330d	331d	
1	1	1	1	1	1	1	1	1	1	
333d2	335d	337d	340d	342d	346d	347d	348d2	350d	352d	
1	1	1	1	1	1	1	1	1	1	
353d	355d	356d	357d	358d	360d	362d	363d	365d	366d	
1	1	1	1	1	1	1	1	1	1	
367d	368d2 1	36d	38d 1	41d2R	42d 1	43d 1	44d		(Other) 161	
1	T	1	T	1	Ţ	T	1	1	TOT	
<pre>sample_type: tumor 260 histological_type: ser 260 summarygrade: high low 129 131 summarystage: late</pre>										
260 tumorstag	1e •									
3 4 204 56	,									
	: D C NA D 180									
grade: 23 131 129										

pltx: Y

260 tax: y 260 days_to_death: Min. 1st Qu. Median Mean 3rd Qu. Max. 30 810 1245 1344 1710 3840 vital_status: deceased living 121 139 debulking: optimal suboptimal 103 157 uncurated_author_metadata: title: serous ovarian cancer 10d///geo_accession: GSM794865///status: Public of title: serous ovarian cancer 115d///geo_accession: GSM794867///status: Public on Ma title: serous ovarian cancer 116d///geo_accession: GSM794868///status: Public on Ma title: serous ovarian cancer 117d///geo_accession: GSM794869///status: Public or title: serous ovarian cancer 119d///geo_accession: GSM794870///status: Public or title: serous ovarian cancer 11d///geo_accession: GSM794866///status: Public of title: serous ovarian cancer 120d///geo_accession: GSM794872///status: Public or title: serous ovarian cancer 122d///geo_accession: GSM794873///status: Public on M title: serous ovarian cancer 123d///geo_accession: GSM794874///status: Public on Ma title: serous ovarian cancer 125Rd///geo_accession: GSM794875///status: Public on M title: serous ovarian cancer 129d///geo_accession: GSM794876///status: Public or title: serous ovarian cancer 12d///geo_accession: GSM794871///status: Public title: serous ovarian cancer 130d///geo_accession: GSM794877///status: Public on Ma title: serous ovarian cancer 132d///geo_accession: GSM794878///status: Public or

title: serous ovarian cancer 134d///geo_accession: GSM794879///status: Public or

title: serous ovarian cancer 139d///geo_accession: GSM794880///status: Public or title: serous ovarian cancer 140d///geo_accession: GSM794881///status: Public on title: serous ovarian cancer 143d///geo_accession: GSM794882///status: Public on M title: serous ovarian cancer 144d///geo_accession: GSM794883///status: Public on title: serous ovarian cancer 145d///geo_accession: GSM794884///status: Public on Ma title: serous ovarian cancer 146d///geo_accession: GSM794885///status: Public on Ma title: serous ovarian cancer 148d///geo_accession: GSM794886///status: Public or title: serous ovarian cancer 150d///geo_accession: GSM794888///status: Public on Ma title: serous ovarian cancer 155d///geo_accession: GSM794889///status: Public or title: serous ovarian cancer 156d///geo_accession: GSM794890///status: Public or title: serous ovarian cancer 15d///geo_accession: GSM794887///status: Public of title: serous ovarian cancer 160d///geo_accession: GSM794892///status: Public on Ma title: serous ovarian cancer 16d///geo_accession: GSM794891///status: Public of title: serous ovarian cancer 171d///geo_accession: GSM794894///status: Public on M title: serous ovarian cancer 173d///geo_accession: GSM794895///status: Public on M title: serous ovarian cancer 174d///geo_accession: GSM794896///status: Public title: serous ovarian cancer 178d///geo_accession: GSM794897///status: Public title: serous ovarian cancer 17d///geo_accession: GSM794893///status: Public of title: serous ovarian cancer 183d///geo_accession: GSM794899///status: Public on Ma title: serous ovarian cancer 184d///geo_accession: GSM794900///status: Public or title: serous ovarian cancer 185d///geo_accession: GSM794901///status: Public or title: serous ovarian cancer 186d///geo_accession: GSM794902///status: Public or title: serous ovarian cancer 18d///geo_accession: GSM794898///status: Public of title: serous ovarian cancer 20d///geo_accession: GSM794904///status: Public on

title: serous ovarian cancer 22d///geo_accession: GSM794905///status: Public on M title: serous ovarian cancer 23d///geo_accession: GSM794906///status: Public or title: serous ovarian cancer 249d///geo_accession: GSM794907///status: Public or title: serous ovarian cancer 257d///geo_accession: GSM794909///status: Public on Ma title: serous ovarian cancer 25d///geo_accession: GSM794908///status: Public of title: serous ovarian cancer 260d///geo_accession: GSM794910///status: Public on M title: serous ovarian cancer 262d///geo_accession: GSM794911///status: Public on Ma title: serous ovarian cancer 264d///geo_accession: GSM794912///status: Public on Ma title: serous ovarian cancer 266d///geo_accession: GSM794913///status: Public on Ma title: serous ovarian cancer 267d///geo_accession: GSM794914///status: Public or title: serous ovarian cancer 268d///geo_accession: GSM794915///status: Public or title: serous ovarian cancer 269d///geo_accession: GSM794916///status: Public on Ma title: serous ovarian cancer 27d///geo_accession: GSM794917///status: Public on title: serous ovarian cancer 299d///geo_accession: GSM794918///status: Public o title: serous ovarian cancer 2d///geo_accession: GSM794903///status: Public on title: serous ovarian cancer 300d///geo_accession: GSM794919///status: Public on title: serous ovarian cancer 301d///geo_accession: GSM794920///status: Public on Ma title: serous ovarian cancer 302d///geo_accession: GSM794921///status: Public or title: serous ovarian cancer 303d///geo_accession: GSM794922///status: Public or title: serous ovarian cancer 304d///geo_accession: GSM794923///status: Public on Ma title: serous ovarian cancer 305d2///geo_accession: GSM794924///status: Public on title: serous ovarian cancer 306d///geo_accession: GSM794925///status: Public on Ma title: serous ovarian cancer 307d///geo_accession: GSM794926///status: Public on Ma title: serous ovarian cancer 310d///geo_accession: GSM794927///status: Public

title: serous ovarian cancer 318d///geo_accession: GSM794928///status: Public on M title: serous ovarian cancer 319d///geo_accession: GSM794929///status: Public on Ma title: serous ovarian cancer 320d2///geo_accession: GSM794930///status: Public or title: serous ovarian cancer 323d///geo_accession: GSM794931///status: Public or title: serous ovarian cancer 327d///geo_accession: GSM794932///status: Public on title: serous ovarian cancer 330d///geo_accession: GSM794933///status: Public on title: serous ovarian cancer 331d///geo_accession: GSM794934///status: Public on Ma title: serous ovarian cancer 333d2///geo_accession: GSM794935///status: Public on title: serous ovarian cancer 335d///geo_accession: GSM794936///status: Public or title: serous ovarian cancer 337d///geo_accession: GSM794937///status: Public on title: serous ovarian cancer 340d///geo_accession: GSM794938///status: Public on title: serous ovarian cancer 342d///geo_accession: GSM794939///status: Public or title: serous ovarian cancer 346d///geo_accession: GSM794940///status: Public or title: serous ovarian cancer 347d///geo_accession: GSM794941///status: Public on Ma title: serous ovarian cancer 348d2///geo_accession: GSM794942///status: Public on title: serous ovarian cancer 350d///geo_accession: GSM794943///status: Public on M title: serous ovarian cancer 352d///geo_accession: GSM794944///status: Public on Ma title: serous ovarian cancer 353d///geo_accession: GSM794945///status: Public title: serous ovarian cancer 355d///geo_accession: GSM794946///status: Public or title: serous ovarian cancer 356d///geo_accession: GSM794947///status: Public title: serous ovarian cancer 357d///geo_accession: GSM794948///status: Public on Ma title: serous ovarian cancer 358d///geo_accession: GSM794949///status: Public or title: serous ovarian cancer 360d///geo_accession: GSM794951///status: Public on Ma title: serous ovarian cancer 362d///geo_accession: GSM794952///status: Public on Ma

title: serous ovarian cancer 363d///geo_accession: GSM794953///status: Public or title: serous ovarian cancer 365d//geo_accession: GSM794954///status: Public on title: serous ovarian cancer 366d///geo_accession: GSM794955///status: Public on title: serous ovarian cancer 367d//geo_accession: GSM794956///status: Public or title: serous ovarian cancer 368d2///geo_accession: GSM794957///status: Public on title: serous ovarian cancer 368d2///geo_accession: GSM794957///status: Public on title: serous ovarian cancer 36d///geo_accession: GSM794950///status: Public on title: serous ovarian cancer 36d///geo_accession: GSM794958///status: Public of title: serous ovarian cancer 41d2R//geo_accession: GSM794960///status: Public on N title: serous ovarian cancer 41d2R//geo_accession: GSM794960///status: Public on N title: serous ovarian cancer 43d///geo_accession: GSM794961///status: Public on N title: serous ovarian cancer 43d///geo_accession: GSM794961///status: Public on N title: serous ovarian cancer 43d///geo_accession: GSM794962///status: Public on N title: serous ovarian cancer 43d///geo_accession: GSM794963///status: Public on title: serous ovarian cancer 44d///geo_accession: GSM794963///status: Public on

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duplicates:
GSE32062.GSE32062.GPL6480_GSM794933 GSE32062.GSE32062.GPL6480_GSM794935
1 1
NA's
258
```

Value

An expression set

GSE32063	High-risk ovarian cancer based on 126-gene expression signature
	is uniquely characterized by downregulation of antigen presentation
	pathway.

Description

High-grade serous ovarian cancers are heterogeneous not only in terms of clinical outcome but also at the molecular level. Our aim was to establish a novel risk classification system based on

a gene expression signature for predicting overall survival, leading to suggesting novel therapeutic strategies for high-risk patients. In this large-scale cross-platform study of six microarray data sets consisting of 1,054 ovarian cancer patients, we developed a gene expression signature for predicting overall survival by applying elastic net and 10-fold cross-validation to a Japanese data set A (n = 260) and evaluated the signature in five other data sets. Subsequently, we investigated differences in the biological characteristics between high- and low-risk ovarian cancer groups. An elastic net analysis identified a 126-gene expression signature for predicting overall survival in patients with ovarian cancer using the Japanese data set A (multivariate analysis, P = 4?? 10(-20)). We validated its predictive ability with five other data sets using multivariate analysis (Tothill's data set, P =1 ?? 10(-5); Bonome's data set, P = 0.0033; Dressman's data set, P = 0.0016; TCGA data set, P = 0.0027; Japanese data set B, P = 0.021). Through gene ontology and pathway analyses, we identified a significant reduction in expression of immune-response-related genes, especially on the antigen presentation pathway, in high-risk ovarian cancer patients. This risk classification based on the 126-gene expression signature is an accurate predictor of clinical outcome in patients with advanced stage high-grade serous ovarian cancer and has the potential to develop new therapeutic strategies for high-grade serous ovarian cancer patients.

Format

```
experimentData(eset):
Experiment data
 Experimenter name: Yoshihara K, Tsunoda T, Shigemizu D, Fujiwara H et al. High-ri
 Laboratory: Yoshihara, Tanaka 2012
  Contact information:
  Title: High-risk ovarian cancer based on 126-gene expression signature is unique
  URL:
  PMIDs: 22241791
  Abstract: A 255 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
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      Agilent-014850 Whole Human Genome Microarray 4x44K G4112F (Probe Name vers
ion)
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      Agilent G4112F
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      hquq4112a
  platform_manufacturer:
      Agilent
  platform_distribution:
      commercial
  platform_accession:
      GPL6480
   version:
      2015-09-22 19:58:23
featureData(eset):
An object of class 'AnnotatedDataFrame'
```

```
featureNames: A_23_P100001 A_23_P100011 ... A_32_P99902 (30936 total)
varLabels: probeset gene EntrezGene.ID best_probe
varMetadata: labelDescription
```

Details

```
assayData: 30936 features, 40 samples
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Overall survival time-to-event summary (in years):
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                                                            1
222 224 226 229 230 231 274 277 278 280 281 282 283 284 285 286
 1 1 1 1 1
                    1 1 1
                                1 1 1 1 1
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287 288 289 291 292 294 297R 298R
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                 1
                    1
                        1
                             1
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tumor
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histological_type:
ser
40
summarygrade:
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17 23
summarystage:
late
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tumorstage:
3 4
31 9
substage:
 b c NA's
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3
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23 17
pltx:
 У
40
tax:
 У
40
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debulking:
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Value

An expression set

GSE44104

COL11A1 promotes tumor progression and predicts poor clinical outcome in ovarian cancer.

Description

Biomarkers that predict disease progression might assist the development of better therapeutic strategies for aggressive cancers, such as ovarian cancer. Here, we investigated the role of collagen type XI alpha 1 (COL11A1) in cell invasiveness and tumor formation and the prognostic impact of COL11A1 expression in ovarian cancer. Microarray analysis suggested that COL11A1 is a disease progression-associated gene that is linked to ovarian cancer recurrence and poor survival. Small interference RNA-mediated specific reduction in COL11A1 protein levels suppressed the invasive ability and oncogenic potential of ovarian cancer cells and decreased tumor formation and lung colonization in mouse xenografts. A combination of experimental approaches, including realtime RT-PCR, casein zymography and chromatin immunoprecipitation (ChIP) assays, showed that COL11A1 knockdown attenuated MMP3 expression and suppressed binding of Ets-1 to its putative MMP3 promoter-binding site, suggesting that the Ets-1-MMP3 axis is upregulated by COL11A1. Transforming growth factor (TGF)-beta (TGF-??1) treatment triggers the activation of smad2 signaling cascades, leading to activation of COL11A1 and MMP3. Pharmacological inhibition of MMP3 abrogated the TGF-??1-triggered, COL11A1-dependent cell invasiveness. Furthermore, the NF-YA-binding site on the COL11A1 promoter was identified as the major determinant of TGF-??1-dependent COL11A1 activation. Analysis of 88 ovarian cancer patients indicated that high COL11A1 mRNA levels are associated with advanced disease stage. The 5-year recurrence-free and overall survival rates were significantly lower (P=0.006 and P=0.018, respectively) among patients with high expression levels of tissue COL11A1 mRNA compared with those with low expression. We conclude that COL11A1 may promote tumor aggressiveness via the TGF-??1-MMP3 axis and that COL11A1 expression can predict clinical outcome in ovarian cancer patients.

Format

```
experimentData(eset):
Experiment data
 Experimenter name: Wu Y, Chang T, Huang Y, Huang H, Chou C
 Laboratory: Wu, Chou 2013
  Contact information:
  Title: COL11A1 promotes tumor progression and predicts poor clinical outcome in o
 URL:
 PMIDs: 23934190
 Abstract: A 260 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
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      Affymetrix HG-U133Plus2
   platform_summary:
      hgu133plus2
   platform_manufacturer:
      Affymetrix
  platform distribution:
      commercial
  platform accession:
     GPL570
  platform_technology:
     in situ oligonucleotide
   version:
      2015-09-22 20:02:05
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (42447 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

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duplicates: Length Class Mode 60 character character

Value

An expression set

GSE49997

Validating the impact of a molecular subtype in ovarian cancer on outcomes: a study of the OVCAD Consortium.

Description

Most patients with epithelial ovarian cancer (EOC) are diagnosed at advanced stage and have a poor prognosis. However, a small proportion of these patients will survive, whereas others will die very quickly. Clinicopathological factors do not allow precise identification of these subgroups. Thus, we have validated a molecular subclassification as new prognostic factor in EOC. One hundred and ninety-four patients with Stage II-IV EOC were characterized by whole-genome expression profiling of tumor tissues and were classified using a published 112 gene set, derived from an International Federation of Gynecology and Obstetrics (FIGO) stage-directed supervised classification approach. The 194 tumor samples were classified into two subclasses comprising 95 (Subclass 1) and 99 (Subclass 2) tumors. All nine FIGO II tumors were grouped in Subclass 1 (P = 0.001). Subclass 2 (54% of advanced-stage tumors) was significantly correlated with peritoneal carcinomatosis

and non-optimal debulking. Patients with Subclass 2 tumors had a worse overall survival for both serous and non-serous histological subtypes, as revealed by univariate analysis (hazard ratios [HR] of 3.17 and 17.11, respectively; P ??? 0.001) and in models corrected for relevant clinicopathologic parameters (HR 2.87 and 12.42, respectively; P ??? 0.023). Significance analysis of microarrays revealed 2082 genes that were differentially expressed in advanced-grade serous tumors of both subclasses and the focal adhesion pathway as the most deregulated pathway. In the present validation study, we have shown that, in advanced-stage serous ovarian cancer, two approximately equally large molecular subtypes exist, independent of classical clinocopathological parameters and presenting with highly different whole-genome expression profiles and a markedly different overall survival. Similar results were obtained in a small cohort of patients with non-serous tumors.?? 2012 Japanese Cancer Association.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Pils D1, Hager G, Tong D, Aust S, Heinze G, Kohl M, Schuster H
 Laboratory: Pils, Zeilinger 2012
  Contact information:
  Title: Validating the impact of a molecular subtype in ovarian cancer on outcomes
  URT:
 PMIDs: 22497737
 Abstract: A 276 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      ABI Human Genome Survey Microarray Version 2
  platform_shorttitle:
      ABI Human Genome
  platform_summary:
  platform_manufacturer:
      Applied Biosystems
  platform_distribution:
      commercial
  platform_accession:
      GPL2986
  platform_technology:
      in situ oligonucleotide
   version:
      2015-09-22 20:04:13
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 100027 100036 ... 10715781 (18439 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 18439 features, 204 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
  10 observations deleted due to missingness
   n events median 0.95LCL 0.95UCL
194.00 57.00 NA 3.67 NA
_____
Available sample meta-data:
_____
alt sample name:
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tumor
 204
histological_type:
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other ser NA's

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Value

An expression set

GSE51088

POSTN/TGFBI-associated stromal signature predicts poor prognosis in serous epithelial ovarian cancer.

Description

To identify molecular prognosticators and therapeutic targets for high-grade serous epithelial ovarian cancers (EOCs) using genetic analyses driven by biologic features of EOC pathogenesis. Ovarian tissue samples (n = 172; 122 serous EOCs, 30 other EOCs, 20 normal/benign) collected prospectively from sequential patients undergoing gynecologic surgery were analyzed using RNA expression microarrays. Samples were classified based on expression of genes with potential relevance in ovarian cancer. Gene sets were defined using Rosetta Similarity Search Tool (ROAST) and analysis of variance (ANOVA). Gene copy number variations were identified by array comparative genomic hybridization. No distinct subgroups of EOC could be identified by unsupervised clustering, however, analyses based on genes correlated with periostin (POSTN) and estrogen receptoralpha (ESR1) yielded distinct subgroups. When 95 high-grade serous EOCs were grouped by genes based on ANOVA comparing ESR1/WT1 and POSTN/TGFBI samples, overall survival (OS) was significantly shorter for 43 patients with tumors expressing genes associated with POSTN/TGFBI compared to 52 patients with tumors expressing genes associated with ESR1/WT1 (median 30 versus 49 months, respectively; P = 0.022). Several targets with the apeutic potential were identified within each subgroup. BRCA germline mutations were more frequent in the ESR1/WT1 subgroup. Proliferation-associated genes and TP53 status (mutated or wild-type) did not correlate with survival. Findings were validated using independent ovarian cancer datasets. Two distinct molecular subgroups of high-grade serous EOCs based on POSTN/TGFBI and ESR1/WT1 expressions were identified with significantly different OS. Specific differentially expressed genes between these subgroups provide potential prognostic and therapeutic targets. Copyright ?? 2013 Elsevier Inc. All rights reserved.

Format

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experimentData(eset):
Experiment data
Experimenter name: Karlan BY, Dering J, Walsh C, Orsulic S, Lester J, Anderson LA
Laboratory: Karlan, Slamon 2014
Contact information:
Title: POSTN/TGFBI-associated stromal signature predicts poor prognosis in serous
URL:
PMIDs: 24368280
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Details

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Value

An expression set

GSE6008

Lysophosphatidic acid-induced transcriptional profile represents serous epithelial ovarian carcinoma and worsened prognosis.

Description

Lysophosphatidic acid (LPA) governs a number of physiologic and pathophysiological processes. Malignant ascites fluid is rich in LPA, and LPA receptors are aberrantly expressed by ovarian cancer cells, implicating LPA in the initiation and progression of ovarian cancer. However, there is an absence of systematic data critically analyzing the transcriptional changes induced by LPA in ovarian cancer. In this study, gene expression profiling was used to examine LPA-mediated transcription by exogenously adding LPA to human epithelial ovarian cancer cells for 24 h to mimic long-term stimulation in the tumor microenvironment. The resultant transcriptional profile comprised a 39-gene signature that closely correlated to serous epithelial ovarian carcinoma. Hierarchical clustering of ovarian cancer patient specimens demonstrated that the signature is associated with worsened prognosis. Patients with LPA-signature-positive ovarian tumors have reduced disease-specific and progression-free survival times. They have a higher frequency of stage IIIc serous carcinoma and a greater proportion is deceased. Among the 39-gene signature, a group of seven genes associated with cell adhesion recapitulated the results. Out of those seven, claudin-1, an adhesion molecule and phenotypic epithelial marker, is the only independent biomarker of serous epithelial ovarian

carcinoma. Knockdown of claudin-1 expression in ovarian cancer cells reduces LPA-mediated cellular adhesion, enhances suspended cells and reduces LPA-mediated migration. The data suggest that transcriptional events mediated by LPA in the tumor microenvironment influence tumor progression through modulation of cell adhesion molecules like claudin-1 and, for the first time, report an LPA-mediated expression signature in ovarian cancer that predicts a worse prognosis.

Format

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  Laboratory: Murph, Mills 2009
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  Title: Lysophosphatidic acid-induced transcriptional profile represents serous ep
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  PMIDs: 19440550
  Abstract: A 247 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
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      [HG-U133A] Affymetrix Human Genome U133A Array
   platform_shorttitle:
      Affymetrix HG-U133A
   platform_summary:
      hqu133a
   platform_manufacturer:
      Affymetrix
   platform_distribution:
      commercial
   platform_accession:
      GPL96
   version:
      2015-09-22 20:07:11
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (20967 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
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Details

assayData: 20967 features, 103 samples Platform type: ------Available sample meta-data:

1

1

1

1

1

1

1

1

1

1

1

1

1

142

alt_sample_name: Ovarian_Tumor_ClearCell_CHTN-OC-004 Ovarian_Tumor_ClearCell_CHTN-OC-012 Ovarian_Tumor_ClearCell_CHTN-OC-028 Ovarian_Tumor_ClearCell_KU-OC-003 Ovarian Tumor ClearCell KU-OC-004 Ovarian Tumor ClearCell KU-OC-005 Ovarian Tumor ClearCell KU-OC-006 Ovarian Tumor ClearCell KU-OC-007 Ovarian_Tumor_Endometrioid_CHTN-OE-005 Ovarian_Tumor_Endometrioid_CHTN-OE-011 Ovarian_Tumor_Endometrioid_CHTN-OE-014 Ovarian_Tumor_Endometrioid_CHTN-OE-017 Ovarian_Tumor_Endometrioid_CHTN-OE-018 Ovarian_Tumor_Endometrioid_CHTN-OE-019 1 Ovarian_Tumor_Endometrioid_CHTN-OE-023 Ovarian_Tumor_Endometrioid_CHTN-OE-029 Ovarian_Tumor_Endometrioid_CHTN-OE-033 Ovarian_Tumor_Endometrioid_CHTN-OE-035 Ovarian_Tumor_Endometrioid_CHTN-OE-036 Ovarian_Tumor_Endometrioid_CHTN-OE-038 1 Ovarian_Tumor_Endometrioid_CHTN-OE-039 Ovarian_Tumor_Endometrioid_CHTN-OE-040 1 Ovarian Tumor Endometrioid CHTN-OE-042 Ovarian Tumor Endometrioid CHTN-OE-046 Ovarian_Tumor_Endometrioid_CHTN-OE-047 Ovarian_Tumor_Endometrioid_CHTN-OE-048 Ovarian_Tumor_Endometrioid_CHTN-OE-053 Ovarian_Tumor_Endometrioid_CHTN-OE-054 Ovarian_Tumor_Endometrioid_CHTN-OE-056 Ovarian_Tumor_Endometrioid_CHTN-OE-059 Ovarian_Tumor_Endometrioid_CHTN-OE-060 Ovarian_Tumor_Endometrioid_CHTN-OE-061 Ovarian_Tumor_Endometrioid_CHTN-OE-065 Ovarian_Tumor_Endometrioid_CHTN-OE-069 Ovarian_Tumor_Endometrioid_CHTN-OE-080 Ovarian_Tumor_Endometrioid_CHTN-OE-082 Ovarian Tumor Endometrioid CHTN-OE-087 Ovarian Tumor Endometrioid CHTN-OE-092 1 Ovarian Tumor Endometrioid JH-OE-2T Ovarian Tumor Endometrioid KU-OE-003 Ovarian_Tumor_Endometrioid_KU-OE-004 Ovarian_Tumor_Endometrioid_KU-OE-007 Ovarian_Tumor_Endometrioid_UM-OE-1T Ovarian_Tumor_Mucinous_CHTN-OM-007

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1 Ovarian_Tumor_Serous_UM-OS-07 1 Ovarian_Tumor_Serous_UM-OS-10 1 (Other) 4

4

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42 53 8

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substage: a b c d NA's 2 54 1 27 19

grade: 1 2 3 NA's 19 17 38 29

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ov 103

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100
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Value

An expression set

GSE6822

Classification of ovarian tumor samples

Description

Ouellet V, Provencher DM, Maugard CM, Le Page C, Ren F, Lussier C, Novak J, Ge B, Hudson TJ, Tonin PN, Mes-Masson A-M: Discrimination between serous low malignant potential and invasive epithelial ovarian tumors using molecular profiling. Oncogene 2005, 24:4672-4687.

Format

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experimentData(eset):
Experiment data
Experimenter name: Ouellet V, Provencher DM, Maugard CM, Le Page C, Ren F, Lussie
Laboratory: Ouellet, Mes-Masson 2005
Contact information:
Title: Classification of ovarian tumor samples
URL:
PMIDs: PMID unknown
Abstract: A 40 word abstract is available. Use 'abstract' method.
Information is available on: preprocessing
```

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notes:
  platform_title:
      [Hu6800] Affymetrix Human Full Length HuGeneFL Array
  platform_shorttitle:
      Affymetrix Hu6800
  platform_summary:
     hu6800
  platform_manufacturer:
      Affymetrix
  platform_distribution:
      commercial
  platform_accession:
      GPL80
   version:
      2015-09-22 20:07:22
featureData(eset):
An object of class 'AnnotatedDataFrame'
```

```
featureNames: A28102_at AB000114_at ... Z97074_at (6407 total)
varLabels: probeset gene EntrezGene.ID best_probe
varMetadata: labelDescription
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Details

assayData: 6407 features, 66 samples Platform type: Available sample meta-data:

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alt_sample_name:
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Ovarian	tumor	AM178	Ovarian	tumor	AM179	Ovarian	tumor	AM182	Ovarian	tumor	AM195
		1			1			1			1
Ovarian	tumor	AM196	Ovarian	tumor	AM198	Ovarian	tumor	AM200	Ovarian	tumor	AM201
		1			1			1			1
Ovarian	tumor	AM202	Ovarian	tumor	AM203	Ovarian	tumor	AM204	Ovarian	tumor	AM207
		1			1			1			1
Ovarian	tumor	AM208	Ovarian	tumor	AM209	Ovarian	tumor	AM225	Ovarian	tumor	AM226
		1			1			1			1
Ovarian	tumor	AM228	Ovarian	tumor	AM233	Ovarian	tumor	AM250	Ovarian	tumor	AM252
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Ovarian	tumor	AM253	Ovarian	tumor	AM255	Ovarian	tumor	AM256	Ovarian	tumor	AM259
		1			1			1			1
Ovarian	tumor	AM261	Ovarian	tumor	AM263	Ovarian	tumor	AM268	Ovarian	tumor	AM269

1	1	1 1
Ovarian tumor AM287 Ovarian 1	tumor AM288 Ovarian 1	tumor AM289 Ovarian tumor AM290 1 1
-		tumor AM294 Ovarian tumor AM311
-	—	tumor AM317 Ovarian tumor AM333
Ovarian tumor AM335 Ovarian	tumor AM339 Ovarian	tumor AM341 Ovarian tumor AM344
		1 1 tumor AM348 Ovarian tumor AM349
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1 Ovarian tumor AM431 Ovarian	1 tumor AM438	1 1
1	1	
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histological_type:		
clearcell		mix mucinous
11 ser undifferent	7 iated	3 1
41	3	
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summarygrade:		
high low NA's 40 15 11		
grade: 1 2 3 NA's		
1 14 40 11		
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2000-12-21 2001-05-03 2001-0 1 1	3 3	1-09-25 2001-09-26 2001-09-27 1 5 8
		2-07-24 2002-10-20 2002-10-30
4 1 2002-11-01 2002-11-13	9 7	4 10 5
2 2		

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title: Ovarian tumor AM335///geo_accession: title: Ovarian tumor AM339///geo_accession: title: Ovarian tumor AM341///geo_accession: (title: Ovarian tumor AM344///geo_accession: GSM15 title: Ovarian tumor AM345///geo_accession: title: Ovarian tumor AM347///geo_accession: GSM157286///status: Public title: Ovarian tumor AM348///geo_accession: title: Ovarian tumor AM349///geo_accession: title: Ovarian tumor AM354///geo_accession: GSM1 title: Ovarian tumor AM364///geo_accession: GSM157290// title: Ovarian tumor AM367///geo_accession: title: Ovarian tumor AM368///geo_accession: title: Ovarian tumor AM381///geo_accession: title: Ovarian tumor AM382///geo_accession: title: Ovarian tumor AM398///geo_accession: GSM157295///status: Public on Dec 31 title: Ovarian tumor AM429///geo_accession: GSM157296///status: Public on Dec 3 title: Ovarian tumor AM431///geo_accession: title: Ovarian tumor AM438///geo_accession:

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Value

An expression set

Analysis of gene expression in early-stage ovarian cancer.

Description

Gene expression profile was analyzed in 68 stage I and 15 borderline ovarian cancers to determine if different clinical features of stage I ovarian cancer such as histotype, grade, and survival are related to differential gene expression. Tumors were obtained directly at surgery and immediately frozen in liquid nitrogen until analysis. Glass arrays containing 16,000 genes were used in a dualcolor assay labeling protocol. Unsupervised analysis identified eight major patient partitions, one of which was statistically associated to overall survival, grading, and histotype and another with grading and histotype. Supervised analysis allowed detection of gene profiles clearly associated to histotype or to degree of differentiation. No difference was found between borderline and grade 1 tumors. As to recurrence, a subset of genes able to differentiate relapsers from nonrelapsers was identified. Among these, cyclin E and minichromosome maintenance protein 5 were found particularly relevant, as their expression was inversely correlated to progression-free survival (P = 0.00033 and 0.017, respectively). Specific molecular signatures define different histotypes and prognosis of stage I ovarian cancer. Mucinous and clear cells histotypes can be distinguished from the others regardless of tumor grade. Cyclin E and minichromosome maintenance protein 5, whose expression was found previously to be related to a bad prognosis of advanced ovarian cancer, appear to be potential prognostic markers in stage I ovarian cancer too, independent of other pathologic and clinical variables.

Format

```
experimentData(eset):
Experiment data
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 Laboratory: Marchini, D'Incalci 2008
  Contact information:
  Title: Analysis of gene expression in early-stage ovarian cancer.
  URL:
  PMIDs: 19047114
  Abstract: A 225 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
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  platform_shorttitle:
      Agilent G4100A cDNA
  platform_summary:
      hgug4100a
  platform_manufacturer:
      Agilent
  platform distribution:
      custom-commerical
```

```
platform_accession:
    GPL5689
platform_technology:
    spotted DNA/cDNA
version:
    2015-09-22 20:07:40
```

```
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1 2 ... 8864 (7809 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 7809 features, 83 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
     n events median 0.95LCL 0.95UCL
    83
       15 NA 12 NA
 _____
Available sample meta-data:
_____
alt_sample_name:
p0102bis sample_Ovarian tumor p0103bis sample_Ovarian tumor
                                                       1
                          1
p0112bis sample_Ovarian tumor p0114bis sample_Ovarian tumor
                          1
                                                       1
p0125bis sample_Ovarian tumor p0128bis sample_Ovarian tumor
                          1
                                                       1
p0143bis sample_Ovarian tumor p0146bis sample_Ovarian tumor
                                                       1
                          1
p0188bis sample Ovarian tumor p0208bis sample Ovarian tumor
                                                      1
                          1
p0210bis sample_Ovarian tumor p0217bis sample_Ovarian tumor
                          1
                                                       1
p057bis sample_Ovarian tumor p070bis sample_Ovarian tumor
                          1
                                                       1
p080bis sample_Ovarian tumor p091bis sample_Ovarian tumor
                          1
                                                      1
p139bis sample_Ovarian tumor p13bis sample_Ovarian tumor
                                                       1
                          1
p141bis sample_Ovarian tumor p166bis sample_Ovarian tumor
                          1
                                                       1
```

p171bis sample_Ovarian tumor p17bis sample_Ovarian tumor 1 p183bis sample Ovarian tumor p209bis sample_Ovarian tumor 1 p212bis sample_Ovarian tumor p213bis sample_Ovarian tumor 1 1 p243bis sample Ovarian tumor p246bis sample Ovarian tumor 1 1 p261bis sample Ovarian tumor p284bis sample Ovarian tumor 1 1 p293bis sample_Ovarian tumor p310bis sample_Ovarian tumor 1 p31bis sample_Ovarian tumor p320bis sample_Ovarian tumor 1 1 p331bis sample_Ovarian tumor p336bis sample_Ovarian tumor p375bis sample_Ovarian tumor p350bis sample_Ovarian tumor p382bis sample_Ovarian tumor p383bis sample_Ovarian tumor 1 1 p386bis sample_Ovarian tumor p388bis sample_Ovarian tumor 1 p398bis sample_Ovarian tumor p39bis sample_Ovarian tumor p401bis sample Ovarian tumor p414bis sample Ovarian tumor 1 1 p429bis sample_Ovarian tumor p421bis sample_Ovarian tumor 1 1 p433bis sample_Ovarian tumor p448bis sample_Ovarian tumor 1 1 p455bis sample_Ovarian tumor p459bis sample_Ovarian tumor 1 1 p462bis sample_Ovarian tumor p482bis sample_Ovarian tumor 1 1 p487bis sample_Ovarian tumor p497bis sample_Ovarian tumor 1 1 p502bis sample_Ovarian tumor p540bis sample_Ovarian tumor 1 1 p541bis sample Ovarian tumor p549bis sample_Ovarian tumor 1 1 p550bis sample_Ovarian tumor p567bis sample_Ovarian tumor 1 1 p56bis sample Ovarian tumor p573bis sample Ovarian tumor 1 1 p586bis sample_Ovarian tumor p597bis sample_Ovarian tumor p616bis sample_Ovarian tumor p63bis sample_Ovarian tumor 1 1

p646bis sample_Ovarian tumor p66bis sample_Ovarian tumor 1 1 p690bis sample_Ovarian tumor p68bis sample_Ovarian tumor 1 1 p692bis sample_Ovarian tumor p725bis sample_Ovarian tumor 1 1 p760bis sample_Ovarian tumor p73bis sample_Ovarian tumor 1 1 p770bis sample_Ovarian tumor p772bis sample_Ovarian tumor 1 1 p775bis sample_Ovarian tumor p793bis sample_Ovarian tumor 1 1 p79bis sample_Ovarian tumor p84bis sample_Ovarian tumor 1 1 p90bis sample_Ovarian tumor 1 sample_type: borderline tumor 15 68 histological_type: clearcell mucinous other endo 16 17 17 1 ser undifferentiated 31 1 primarysite: ov 83 summarygrade: high low NA's 35 33 15 summarystage: early 83 tumorstage: 1 83 substage: a b c 25 5 53 grade:

1 2 3 NA's 35 15 13 20 age_at_initial_pathologic_diagnosis: Min. 1st Qu. Median Mean 3rd Qu. Max. 21.00 43.00 50.00 51.25 61.00 87.00 recurrence_status: norecurrence recurrence 62 21 days_to_death: Min. 1st Qu. Median Mean 3rd Qu. Max. 0 1192 2248 2273 3048 5824 vital_status: deceased living 15 68 uncurated_author_metadata: title: p0102bis sample_Ovarian tumor///geo_accession: GSM214010///status: title: p0103bis sample_Ovarian tumor///geo_accession: GSM214078///status: Public title: p0112bis sample_Ovarian tumor///geo_accession: GSM214040///sta title: p0114bis sample_Ovarian tumor///geo_accession: GSM214 title: p0125bis sample_Ovarian tumor///geo_accession: GSM214009///status: title: p0128bis sample_Ovarian tumor///geo_accession: GSM214030///s title: p0143bis sample_Ovarian tumor///geo_accession: GSM214012///status: title: p0146bis sample_Ovarian tumor///geo_accession: GSM214033///status: title: p0188bis sample_Ovarian tumor///geo_accession: GSM214041// title: p0208bis sample_Ovarian tumor///geo_accession: GSM214011///status title: p0210bis sample_Ovarian tumor///geo_accession: GSM214031///status: title: p0217bis sample_Ovarian tumor///geo_accession: GSM214008///sta title: p057bis sample_Ovarian tumor///geo_accession: GSM214064///status: Publ title: p070bis sample_Ovarian tumor///geo_accession: GSM214032///status

title: p080bis sample_Ovarian tumor///geo_accession: GSM214017///status: F title: p091bis sample_Ovarian tumor///geo_accession: GSM214024///status: title: p139bis sample_Ovarian tumor///geo_accession: GSM214047///status title: p13bis sample_Ovarian tumor///geo_accession: GSM214043///status: title: p141bis sample_Ovarian tumor///geo_accession: GSM214081///status: Public title: p166bis sample_Ovarian tumor///geo_accession: GSM214013///status: Public title: p171bis sample_Ovarian tumor///geo_accession: GSM214014///status title: p17bis sample_Ovarian tumor///geo_accession: GSM214080///status: Public or title: p183bis sample_Ovarian tumor///geo_accession: GSM214015///status: title: p209bis sample_Ovarian tumor///geo_accession: GSM214090///status: Public or title: p212bis sample_Ovarian tumor///geo_accession: GSM214065///status: Publ title: p213bis sample_Ovarian tumor///geo_accession: GSM214018///status: F title: p243bis sample_Ovarian tumor///geo_accession: GSM214042///status: H title: p246bis sample_Ovarian tumor///geo_accession: GSM214055///status: Publi title: p261bis sample_Ovarian tumor///geo_accession: GSM214034///status: title: p284bis sample_Ovarian tumor///geo_accession: GSM21 title: p293bis sample_Ovarian tumor///geo_accession: GSM214035///status: Publi title: p310bis sample_Ovarian tumor///geo_accession: GSM214083///status: Public on title: p31bis sample_Ovarian tumor///geo_accession: GSM214019///status: F title: p320bis sample_Ovarian tumor///geo_accession: GSM214020///status: title: p331bis sample_Ovarian tumor///geo_accession: GSM214021///status: Publ title: p336bis sample_Ovarian tumor///geo_accession: GSM214056///status: title: p350bis sample_Ovarian tumor///geo_accession: GSM214036///status: Publi title: p375bis sample_Ovarian tumor///geo_accession: GSM214048///status: Publi

title: p382bis sample_Ovarian tumor///geo_accession: GSM214037///status: title: p383bis sample_Ovarian tumor///geo_accession: GSM214029///status: title: p386bis sample_Ovarian tumor///geo_accession: GSM214038///status: title: p388bis sample_Ovarian tumor///geo_accession: GSM214059///status: title: p398bis sample_Ovarian tumor///geo_accession: GSM214066///status: title: p39bis sample_Ovarian tumor///geo_accession: GSM214076///status: Public on title: p401bis sample_Ovarian tumor///geo_accession: GSM214022///status: H title: p414bis sample_Ovarian tumor///geo_accession: GSM214051///status: Publi title: p421bis sample_Ovarian tumor///geo_accession: GSM214023///status title: p429bis sample_Ovarian tumor///geo_accession: GSM214067///sta title: p433bis sample_Ovarian tumor///geo_accession: GSM214079///status: Public on title: p448bis sample_Ovarian tumor///geo_accession: GSM214068///status: H title: p455bis sample_Ovarian tumor///geo_accession: GSM214069///status title: p459bis sample_Ovarian tumor///geo_accession: GSM214025///status: Publi title: p462bis sample_Ovarian tumor///geo_accession: GSM214084///status: Public on title: p482bis sample_Ovarian tumor///geo_accession: GSM214050///status: F title: p487bis sample_Ovarian tumor///geo_accession: GSM214026///status: Publ title: p497bis sample_Ovarian tumor///geo_accession: GSM214052///statu title: p502bis sample_Ovarian tumor///geo_accession: GSM214070///status: Puk title: p540bis sample_Ovarian tumor///geo_accession: GSM214085///status: Public title: p541bis sample_Ovarian tumor///geo_accession: GSM214082///status: Public or title: p549bis sample_Ovarian tumor///geo_accession: GSM214086///status: Public on title: p550bis sample_Ovarian tumor///geo_accession: GSM214053///status: title: p567bis sample_Ovarian tumor///geo_accession: GSM214054///status: Publi

title: p56bis sample_Ovarian tumor///geo_accession: GSM214044///status: title: p573bis sample_Ovarian tumor///geo_accession: GSM214060///status: Publ title: p586bis sample_Ovarian tumor///geo_accession: GSM214061///status: Publ title: p597bis sample_Ovarian tumor///geo_accession: GSM214088///status: Public title: p616bis sample_Ovarian tumor///geo_accession: GSM214071///status: Publ title: p63bis sample_Ovarian tumor///geo_accession: GSM214027///status: Public title: p646bis sample_Ovarian tumor///geo_accession: GSM214087///status: Public of title: p66bis sample_Ovarian tumor///geo_accession: GSM214045///status: Publi title: p68bis sample_Ovarian tumor///geo_accession: GSM214046///status: Publ title: p690bis sample_Ovarian tumor///geo_accession: GSM214072///status: title: p692bis sample_Ovarian tumor///geo_accession: GSM214073///status: Pub title: p725bis sample_Ovarian tumor///geo_accession: GSM214057///status: Pub title: p73bis sample_Ovarian tumor///geo_accession: GSM214028///status: Publi title: p760bis sample_Ovarian tumor///geo_accession: GSM214062///statu title: p770bis sample_Ovarian tumor///geo_accession: GSM214089///status: F title: p772bis sample_Ovarian tumor///geo_accession: GSM214058///status: Pu title: p775bis sample_Ovarian tumor///geo_accession: GSM214074///status: Pub

title: p793bis sample_Ovarian tumor///geo_accession: GSM214075///status: Pu

title: p79bis sample_Ovarian tumor///geo_accession: GSM214063///status: Publi

title: p84bis sample_Ovarian tumor///geo_accession: GSM214039///status: Publ

title: p90bis sample_Ovarian tumor///geo_accession: GSM214077///status: Public on

Value

An expression set

Novel molecular subtypes of serous and endometrioid ovarian cancer linked to clinical outcome.

Description

The study aim to identify novel molecular subtypes of ovarian cancer by gene expression profiling with linkage to clinical and pathologic features. Microarray gene expression profiling was done on 285 serous and endometrioid tumors of the ovary, peritoneum, and fallopian tube. K-means clustering was applied to identify robust molecular subtypes. Statistical analysis identified differentially expressed genes, pathways, and gene ontologies. Laser capture microdissection, pathology review, and immunohistochemistry validated the array-based findings. Patient survival within kmeans groups was evaluated using Cox proportional hazards models. Class prediction validated k-means groups in an independent dataset. A semisupervised survival analysis of the array data was used to compare against unsupervised clustering results.Optimal clustering of array data identified six molecular subtypes. Two subtypes represented predominantly serous low malignant potential and low-grade endometrioid subtypes, respectively. The remaining four subtypes represented higher grade and advanced stage cancers of serous and endometrioid morphology. A novel subtype of high-grade serous cancers reflected a mesenchymal cell type, characterized by overexpression of N-cadherin and P-cadherin and low expression of differentiation markers, including CA125 and MUC1. A poor prognosis subtype was defined by a reactive stroma gene expression signature, correlating with extensive desmoplasia in such samples. A similar poor prognosis signature could be found using a semisupervised analysis. Each subtype displayed distinct levels and patterns of immune cell infiltration. Class prediction identified similar subtypes in an independent ovarian dataset with similar prognostic trends. Gene expression profiling identified molecular subtypes of ovarian cancer of biological and clinical importance.

Format

```
experimentData(eset):
Experiment data
 Experimenter name: Tothill RW, Tinker AV, George J, Brown R, Fox SB, Lade S, Johr
 Laboratory: Tothill, Bowtell 2008
 Contact information:
 Title: Novel molecular subtypes of serous and endometrioid ovarian cancer linked
 URL:
 PMIDs: 18698038
 Abstract: A 243 word abstract is available. Use 'abstract' method.
 Information is available on: preprocessing
  notes:
  platform_title:
      [HG-U133_Plus_2] Affymetrix Human Genome U133 Plus 2.0 Array
  platform_shorttitle:
      Affymetrix HG-U133Plus2
  platform_summary:
     hgu133plus2
```

```
platform_manufacturer:
    Affymetrix
    platform_distribution:
        commercial
    platform_accession:
        GPL570
    version:
        2015-09-22 20:16:32
featureData(eset):
An object of class 'AnnotatedDataFrame'
    featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
        (42447 total)
    varLabels: probeset gene EntrezGene.ID best_probe
    varMetadata: labelDescription
```

Details

```
assayData: 42447 features, 285 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
```

7 observations deleted due to missingness n events median 0.95LCL 0.95UCL 278.00 113.00 3.95 3.53 5.01

```
_____
```

Available sample meta-data:

alt_sample_name:									
X129	X146	X152	X20019	X20025	X20027	X20031	X20032	X20041	X20046
1	1	1	1	1	1	1	1	1	1
X20074	X22002	X22012	X22013	X22020	X22023	X22027	X22029	X22031	X22037
1	1	1	1	1	1	1	1	1	1
X22046	X22047	X22048	X22057	X22058	X2219	X2227	X23026	X23030	X23036
1	1	1	1	1	1	1	1	1	1
X23043	X23052	X23053	X23055	X23066	X23070	X23074	X23077	X23084	X23098
1	1	1	1	1	1	1	1	1	1
X23102	X23106	X23116	X23128	X23139	X23143	X23162	X23165	X23167	X23170
1	1	1	1	1	1	1	1	1	1
X23172	X23177	X23178	X23182	X23187	X23197	X23202	X23204	X23210	X23212
1	1	1	1	1	1	1	1	1	1
X23213	X23221	X26047	X261	X27006	X27098	X32013	X32022	X32032	X32034
1	1	1	1	1	1	1	1	1	1
X32048	X32049	X32054	X32055	X32089	X32098	X32103	X32117	X34019	X34049
1	1	1	1	1	1	1	1	1	1

```
X34066 X34078 X34080 X34085 X34086 X34090 X34102 X34103 X34111 X34113
 1 1 1 1 1 1 1 1 1 1
X34117 X34125 X34165 X34168 X34172 X34186 X34202 X34207 X34801 (Other)
      1 1 1 1 1 1 1
                                               1
   1
                                                       186
sample_type:
tumor
 285
histological_type:
endo other ser
  20 1 264
primarysite:
  ft other ov
  8 34 243
arrayedsite:
  ft other ov
  2 83
          200
summarygrade:
high low NA's
163 116 6
summarystage:
early late NA's
  42 240 3
tumorstage:
 1 2 3 4 NA's
 24 18 218 22 3
substage:
 a b c NA's
 26 19 212 28
grade:
 1 2 3 NA's
 19 97 163 6
age_at_initial_pathologic_diagnosis:
  Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
 22.00 53.00 59.00 59.62 68.00 80.00 3
pltx:
 n y NA's
 39 243 3
```

```
tax:
 n y NA's
 87 195 3
neo:
n y NA's
264 18 3
days_to_tumor_recurrence:
  Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
  0.0 300.0 450.0 618.9 810.0 4980.0 10
recurrence_status:
norecurrence recurrence NA's
    94
          188
                       3
days_to_death:
  Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
  0.0 547.5 855.0 955.1 1252.0 6420.0 7
vital status:
deceased living NA's
113 169 3
debulking:
  optimal suboptimal NA's 160 88 37
batch:
2004-12-03 2004-12-23 2005-01-12 2005-01-17 2005-01-24 2005-01-31 2005-02-21
     3 4 7 7 8 10 10
2005-03-17 2005-05-05 2005-05-09 2005-05-25 2005-05-27 2005-05-30 2005-06-02
     2 1 1 2 3 3 6
2005-06-06 2005-06-08 2005-06-16 2005-06-17 2005-06-24 2005-07-06 2005-07-15
                                            2
     4
             5
                     3
                            5 6
                                                    9
2005-07-20 2005-07-29 2005-08-03 2005-08-05 2005-08-18 2005-08-24 2005-08-26
                            3
     7
             5
                    6
                                    4
                                            8
                                                    4
2005-09-09 2005-09-14 2005-09-16 2005-09-21 2005-10-05 2005-10-26 2005-10-28
     4 6 6 4 5 2
                                                    4
2005-11-04 2005-11-09 2005-11-11 2005-11-23 2005-12-15 2005-12-21 2006-01-20
     6 3 7 4 7 8 3
2006-01-31 2006-02-08 2006-02-28 2006-04-05 2006-04-06 2006-04-12 2006-04-13
     7 3 3 7 3 7
                                                   4
2006-04-28 2006-05-03 2006-06-06 2006-06-07 2006-06-22 2006-07-07 2006-07-19
     6 9 6 3 9 4
                                                    7
```

uncurated_author_metadata:

title: X129///geo_accession: GSM250001///status: Public on Mar 01 20 title: X146///geo_accession: GSM250000///status: Public on Mar 01 2008/ title: X152///geo_accession: GSM249999///status: Public on Mar 01 2008///submi title: X20019///geo_accession: GSM249998///status: Public on Mar 01 2008 title: X20025///geo_accession: GSM249997///status: Public on Mar 01 2008///subr title: X20027///geo_accession: GSM249996///status: Public on Mar 01 2008 title: X20031///geo_accession: GSM249995///status: Public on Mar 01 title: X20032///geo_accession: GSM249994///status: Public on Mar 01 20 title: X20041///geo_accession: GSM249993///status: Public on Mar 01 2008 title: X20046///geo_accession: GSM249992///status: Public on Mar 01 2008// title: X20074///geo_accession: GSM249991///status: Public on Mar 01 2008///sub title: X22002///geo_accession: GSM249728///status: Public on Mar 01 2 title: X22012///geo_accession: GSM249990///status: Public on Mar 01 2008/ title: X22013///geo_accession: GSM249989///status: Public on Mar 01 2008///subr title: X22020///geo_accession: GSM249988///status: Public on Mar 01 2008 title: X22023///geo_accession: GSM249987///status: Public on Mar 01 2008/ title: X22027///geo_accession: GSM249725///status: Public on Mar 01 title: X22029///geo_accession: GSM249986///status: Public on Mar 01 200 title: X22031///geo_accession: GSM249985///status: Public on Mar 01 2 title: X22037///geo_accession: GSM249984///status: Public on Mar 01 2008/ title: X22046///geo_accession: GSM249983///status: Public on Mar 01 2008/ title: X22047///geo_accession: GSM249982///status: Public on Mar 01 2008///su title: X22048///geo_accession: GSM249981///status: Public on Mar 01 2008/ title: X22057///geo_accession: GSM249980///status: Public on Mar 01 2008//

title: X22058///geo_accession: GSM249979///status: Public on Mar 01 2008/ title: X2219///geo_accession: GSM249978///status: Public on Mar 01 title: X2227///geo_accession: GSM249977///status: Public on Ma title: X23026///geo_accession: GSM249976///status: Public on Mar 01 2008/ title: X23030///geo_accession: GSM249975///status: Public on Mar 01 2008/ title: X23036///geo_accession: GSM249727///status: Public on Mar 01 2 title: X23043///geo_accession: GSM249974///status: Public on Mar 01 title: X23052///geo_accession: GSM249721///status: Public on Mar title: X23053///geo_accession: GSM249973///status: Public on Mar 01 2008/ title: X23055///geo_accession: GSM249972///status: Public on Mar 01 2008/ title: X23066///geo_accession: GSM249716///status: Public on Mar 01 2 title: X23070///geo_accession: GSM249971///status: Public on Mar 01 200 title: X23074///geo_accession: GSM249970///status: Public on Mar 01 2008// title: X23077///geo_accession: GSM249969///status: Public on Mar 01 2008/ title: X23084///geo_accession: GSM249968///status: Public on Mar 01 2008/ title: X23098///geo_accession: GSM249967///status: Public on Mar 01 2008// title: X23102///geo_accession: GSM249966///status: Public on Mar 01 2 title: X23106///geo_accession: GSM249965///status: Public on Mar 01 2008///submissi title: X23116///geo_accession: GSM249964///status: Public on Mar 01 2008///su title: X23128///geo_accession: GSM249963///status: Public on Mar 01 2008/ title: X23139///geo_accession: GSM249962///status: Public on Mar 01 2008/ title: X23143///geo_accession: GSM249961///status: Public on Mar 01 2008/ title: X23162///geo_accession: GSM249960///status: Public on Mar 01 2008// title: X23165///geo_accession: GSM249959///status: Public on Mar 01 2008/

title: X23167///geo_accession: GSM249958///status: Public on Mar 01 2008/ title: X23170///geo_accession: GSM249957///status: Public on Mar 01 2008/ title: X23172///geo_accession: GSM249956///status: Public on Mar 01 2008 title: X23177///geo_accession: GSM249720///status: Public on Mar title: X23178///geo_accession: GSM249955///status: Public on Mar 01 2 title: X23182///geo_accession: GSM249954///status: Public on Mar 01 2008/ title: X23187///geo_accession: GSM249953///status: Public on Mar 01 2008/ title: X23197///geo_accession: GSM249951///status: Public on Mar 01 2008 title: X23202///geo_accession: GSM249950///status: Public on Mar 01 2008// title: X23204///geo_accession: GSM249949///status: Public on Mar 01 2008// title: X23210///geo_accession: GSM249948///status: Public on Mar 01 20 title: X23212///geo_accession: GSM249947///status: Public on Mar 01 2008///su title: X23213///geo_accession: GSM249946///status: Public on Mar 01 2008 title: X23221///geo_accession: GSM249945///status: Public on Mar 01 2008///su title: X26047///geo_accession: GSM249944///status: Public on Mar 01 2008///sub title: X261///geo_accession: GSM249943///status: Public on Mar 01 title: X27006///geo_accession: GSM249942///status: Public on Mar 01 2008/ title: X27098///geo_accession: GSM249941///status: Public on Mar 01 2008// title: X32013///geo_accession: GSM249940///status: Public on Mar 01 200 title: X32022///geo_accession: GSM249939///status: Public on Mar 01 2008/ title: X32032///geo_accession: GSM249938///status: Public on Mar 01 2008///su title: X32034///geo_accession: GSM249937///status: Public on Mar 01 2008///subr title: X32048///geo_accession: GSM249936///status: Public on Mar 01 2008// title: X32049///geo_accession: GSM249935///status: Public on Mar 01 2008///su

title: X32054///geo_accession: GSM249934///status: Public on Mar 01 2008/ title: X32055///geo_accession: GSM249933///status: Public on Mar 01 2 title: X32089///geo_accession: GSM249932///status: Public on Mar 01 2008///submiss title: X32098///geo_accession: GSM249931///status: Public on Mar 01 2008///s title: X32103///geo_accession: GSM249930///status: Public on Mar 01 2008/ title: X32117///geo_accession: GSM249715///status: Public on Mar title: X34019///geo_accession: GSM249929///status: Public on Mar 01 2008/ title: X34049///geo_accession: GSM249928///status: Public on Mar 01 200 title: X34066///geo_accession: GSM249927///status: Public on Mar 01 2008/ title: X34078///geo_accession: GSM249926///status: Public on Mar 01 2008/ title: X34080///geo_accession: GSM249925///status: Public on Mar 01 2008/ title: X34085///geo_accession: GSM249924///status: Public on Mar 01 2008/// title: X34086///geo_accession: GSM249923///status: Public on Mar 01 2008 title: X34090///geo_accession: GSM249922///status: Public on Mar (title: X34102///geo_accession: GSM249921///status: Public on Mar 01 2008///sub title: X34103///geo_accession: GSM249920///status: Public on Mar 01 2008/ title: X34111///geo_accession: GSM249919///status: Public on Mar 01 2008/ title: X34113///geo_accession: GSM249918///status: Public on Mar 01 2008// title: X34117///geo_accession: GSM249917///status: Public on Mar 01 2008/ title: X34125///geo_accession: GSM249916///status: Public on Mar 01 2008/ title: X34165///geo_accession: GSM249915///status: Public on Mar 01 2008/ title: X34168///geo_accession: GSM249914///status: Public on Mar 01 2008///su title: X34172///geo_accession: GSM249913///status: Public on Mar 01 2008/ title: X34186///geo_accession: GSM249912///status: Public on Mar 01 2008/ title: X34202///geo_accession: GSM249911///status: Public on Mar 01 2008///sub title: X34207///geo_accession: GSM249910///status: Public on Mar 01 2008// title: X34801///geo_accession: GSM249909///status: Public on Mar 01 2008//

Value

An expression set

loadOvarianDatasets

Function to load ovarian cancer SummarizedExperiment objects from the Experiment Hub

Description

This function returns ovarian cancer datasets from the hub and a vector of patients from the datasets that are duplicates based on a spearman correlation > 0.98

Usage

```
loadOvarianDatasets(
  rescale = FALSE,
  minNumberGenes = 0,
  minSampleSize = 0,
  keepCommonOnly = FALSE,
  imputeMissing = FALSE,
  removeDuplicates = FALSE
)
```

Arguments

rescale	apply centering and scaling to the expression sets (default FALSE)
minNumberGene	es
	an integer specifying to remove expression sets with less genes than this number (default 0) $% \left(\frac{1}{2}\right) =0$
minNumberEven	nts
	an integer specifying how man survival events must be in the dataset to keep the dataset (default 0)

minSampleSiz	e
	an integer specifying the minimum number of patients required in a summa-rizedExperiment (default 0)
keepCommonOn	ly
	remove entrezIDs not common to all datasets (default FALSE)
imputeMissin	g
	remove patients from datasets with missing expression values
removeDuplic	ates
	remove patients with a Spearman correlation greater than or equal to 0.98 with
	other patient expression profiles (default TRUE)

Value

a list with 2 elements. The First element named summarizedExperiments contains the datasets. The second element named duplicates contains a vector with patient IDs for the duplicate patients (those with Spearman correlation greater than or equal to 0.98 with other patient expression profiles).

Examples

```
experimentsAndDups = loadOvarianDatasets()
```

loadOvarianEsets	Function to load ovarian cancer expression sets from the Experiment
	Hub

Description

This function returns ovarian cancer datasets from the hub and a vector of patients from the datasets that are most likely duplicates

Usage

```
loadOvarianEsets(
   removeDuplicates = TRUE,
   quantileCutoff = 0,
   rescale = FALSE,
   minNumberGenes = 0,
   minSampleSize = 0,
   removeRetracted = TRUE,
   removeSubsets = TRUE,
   keepCommonOnly = FALSE,
   imputeMissing = FALSE
)
```

PMID15897565

Arguments

removeDuplic	ates					
	remove patients with a Spearman correlation greater than or equal to 0.98 with other patient expression profiles (default TRUE)					
quantileCuto	ff					
	A nueric between 0 and 1 specifying to remove genes with standard deviation below the required quantile (default 0)					
rescale	apply centering and scaling to the expression sets (default FALSE)					
minNumberGen	es					
	an integer specifying to remove expression sets with less genes than this number (default 0)					
minNumberEvents						
	an integer specifying how man survival events must be in the dataset to keep the dataset (default 0)					
minSampleSiz	e					
	an integer specifying the minimum number of patients required in an eset (de-fault 0)					
removeRetrac	ted					
	remove datasets from retracted papers (default TRUE, currently just PMID17290060 dataset)					
removeSubset	S					
	remove datasets that are a subset of other datasets (defeault TRUE, currently just PMID19318476)					
keepCommonOn	ly					
	remove probes not common to all datasets (default FALSE)					
imputeMissin	g					
	remove patients from datasets with missing expression values					

Value

a list with 2 elements. The First element named esets contains the datasets. The second element named duplicates contains a vector with patient IDs for the duplicate patients (those with Spearman correlation greater than or equal to 0.98 with other patient expression profiles).

Examples

```
esetsAndDups = loadOvarianEsets()
```

PMID15897565

Patterns of gene expression that characterize long-term survival in advanced stage serous ovarian cancers.

Description

A better understanding of the underlying biology of invasive serous ovarian cancer is critical for the development of early detection strategies and new therapeutics. The objective of this study was to define gene expression patterns associated with favorable survival.RNA from 65 serous ovarian cancers was analyzed using Affymetrix U133A microarrays. This included 54 stage III/IV cases (30 short-term survivors who lived <3 years and 24 long-term survivors who lived >7 years) and 11 stage I/II cases. Genes were screened on the basis of their level of and variability in expression, leaving 7,821 for use in developing a predictive model for survival. A composite predictive model was developed that combines Bayesian classification tree and multivariate discriminant models. Leave-one-out cross-validation was used to select and evaluate models.Patterns of genes were identified that distinguish short-term and long-term ovarian cancer survivors. The expression model developed for advanced stage disease classified all 11 early-stage ovarian cancers as long-term survivors. The MAL gene, which has been shown to confer resistance to cancer therapy, was most highly overexpressed in short-term survivors (3-fold compared with long-term survivors, and 29fold compared with early-stage cases). These results suggest that gene expression patterns underlie differences in outcome, and an examination of the genes that provide this discrimination reveals that many are implicated in processes that define the malignant phenotype.Differences in survival of advanced ovarian cancers are reflected by distinct patterns of gene expression. This biological distinction is further emphasized by the finding that early-stage cancers share expression patterns with the advanced stage long-term survivors, suggesting a shared favorable biology.

Format

```
experimentData(eset):
Experiment data
 Experimenter name: Berchuck A, Iversen ES, Lancaster JM, Pittman J, Luo J, Lee P,
  Laboratory: Berchuck, Marks 2005
  Contact information:
  Title: Patterns of gene expression that characterize long-term survival in advance
  URT:
  PMIDs: 15897565
 Abstract: A 258 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [HG-U133A] Affymetrix Human Genome U133A Array
  platform_shorttitle:
      Affymetrix HG-U133A
  platform_summary:
      hgu133a
  platform_manufacturer:
      Affymetrix
  platform_distribution:
      commercial
  platform_accession:
      GPL96
   warnings:
```

PMID15897565

```
These samples are a subset of PMID17290060.

version:

2015-09-22 20:17:53

featureData(eset):

An object of class 'AnnotatedDataFrame'

featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at

(20967 total)

varLabels: probeset gene EntrezGene.ID best_probe

varMetadata: labelDescription
```

Details

```
assayData: 20967 features, 63 samples
Platform type:
_____
Available sample meta-data:
_____
alt_sample_name:
  Min. 1st Qu. Median Mean 3rd Qu. Max.
  1761 1828 1907 2001 2032
                                   2536
sample_type:
tumor
  63
histological_type:
ser
63
primarysite:
ov
63
summarygrade:
high low NA's
 25 37 1
summarystage:
early late
  11 52
tumorstage:
1 2 3 4
7 4 48 4
grade:
```

2 3 4 NA's 1 2 35 24 1 1 age_at_initial_pathologic_diagnosis: Min. 1st Qu. Median Mean 3rd Qu. Max. 33.00 52.50 59.00 59.21 67.00 79.00 os_binary: long short NA's 24 28 11 debulking: optimal suboptimal NA's 24 28 11 batch: 2002-09-20 2002-10-23 2002-11-12 2002-12-16 2002-12-21 2003-01-03 2003-05-30 9 10 1 3 11 13 15 2003-07-02 1 uncurated author metadata: Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1761///Cancer.Type: Early st Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1762///Cancer.Type: Early st Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1763///Cancer.Type: Early sta Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1764///Cancer.Type: Early st Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1765///Cancer.Type: Early st Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1772///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1773///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1774///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1775///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1776///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1777///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1778///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1779///Cancer.Type: Lor

Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1780///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1781///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1828///Cancer.Type: Sh Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1829///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1830///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1831///Cancer.Type: Show Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1832///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1833///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1834///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1835///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1836///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1900///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1901///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1902///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1903///Cancer.Type: Early sta Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1904///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1905///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1906///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1907///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1908///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1909///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 1989///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2003///Cancer.Type: Show Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2004///Cancer.Type: Show

PMID15897565

Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2005///Cancer.Type: Show Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2019///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2020///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2021///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2026///Cancer.Type: Show Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2027///Cancer.Type: Show Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2028///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2029///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2030///Cancer.Type: Sh Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2031///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2032///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2033///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2390///Cancer.Type: Early st Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2391///Cancer.Type: Early st Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2392///Cancer.Type: Early sta Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2393///Cancer.Type: Early sta Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2394///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2395///Cancer.Type: Lor Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2396///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2397///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2398///Cancer.Type: Sh Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2399///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2400///Cancer.Type: Show Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2401///Cancer.Type: Shore Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2402///Cancer.Type: Sh

Genome.ID..File.name....0074_GenomeID_h133a_2802.cel: 2536///Cancer.Type: Early st

Value

An expression set

PMID17290060An integrated genomic-based approach to individualized treatment of
patients with advanced-stage ovarian cancer.

Description

The purpose of this study was to develop an integrated genomic-based approach to personalized treatment of patients with advanced-stage ovarian cancer. We have used gene expression profiles to identify patients likely to be resistant to primary platinum-based chemotherapy and also to identify alternate targeted therapeutic options for patients with de novo platinum-resistant disease.A gene expression model that predicts response to platinum-based therapy was developed using a training set of 83 advanced-stage serous ovarian cancers and tested on a 36-sample external validation set. In parallel, expression signatures that define the status of oncogenic signaling pathways were evaluated in 119 primary ovarian cancers and 12 ovarian cancer cell lines. In an effort to increase chemotherapy sensitivity, pathways shown to be activated in platinum-resistant cancers were subject to targeted therapy in ovarian cancer cell lines.Gene expression profiles identified patients with ovarian cancer likely to be resistant to primary platinum-based chemotherapy with greater than 80% accuracy. In patients with platinum-resistant disease, we identified expression signatures consistent with activation of Src and Rb/E2F pathways, components of which were successfully targeted to increase response in ovarian cancer cell lines.We have defined a strategy for treatment of patients with advanced-stage ovarian cancer that uses therapeutic stratification based on predictions of response to chemotherapy, coupled with prediction of oncogenic pathway deregulation, as a method to direct the use of targeted agents.

Format

```
experimentData(eset):
Experiment data
Experimenter name: Dressman HK, Berchuck A, Chan G, Zhai J, Bild A, Sayer R, Crace
Laboratory: Dressman, Lancaster 2007
Contact information:
Title: An integrated genomic-based approach to individualized treatment of patier
URL:
PMIDs: 17290060
Abstract: A 223 word abstract is available. Use 'abstract' method.
Information is available on: preprocessing
```

```
notes:
  platform_title:
      [HG-U133A] Affymetrix Human Genome U133A Array
  platform_shorttitle:
      Affymetrix HG-U133A
  platform_summary:
     hgu133a
   platform_manufacturer:
      Affymetrix
   platform_distribution:
      commercial
  platform_accession:
     GPL96
   warnings:
     This paper has been retracted.
   version:
      2015-09-22 20:19:16
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (20967 total)
```

```
varLabels: probeset gene EntrezGene.ID best_probe
varMetadata: labelDescription
```

Details

```
assayData: 20967 features, 117 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
```

n events median 0.95LCL 0.95UCL 117.00 67.00 5.26 2.79 7.48

Available sample meta-data:

alt_sample_name:

1024	1447	1451	1504	1526	1552	1578	1590	1615	1623
1	1	1	1	1	1	1	1	1	1
1665	1674	1675	1774	1784	1834	1846	1877	1913	1929
1	1	1	1	1	1	1	1	1	1
2046	2063	2064	2075	2198	2204	2324	2419	2422	2424
1	1	1	1	1	1	1	1	1	1
2465	2476	2479	2505	2542	2573	2673	2739	2802	2849
1	1	1	1	1	1	1	1	1	1

		2981								
1 872	1 922	1 D1805	1 D1837	1 D1859	1 D2098	1 D2208	1 D2332	1 D2342	1 D2358	
1 D2421	1 D2432	1 D2433	1 D2480	1 D2557	1 D2559	1 D2560	1 D2572	1 D2575		
1	1	1	1	1	1	1	1	1	1	
D2581 1	D2603 1	D2611 1	D2629 1	D2640 1	D2648 1	D2668 1	D2689 1	D2691 1	D2700 1	
D2726	D2727	D2733	D2738	D2749	D2776	D2792	M1054	M1055	M120	
1 M1241 1	1 M1390 1	1 M1503 1	M1572	1 M17 1	1 M1891 1		1 M2097 1	1 M2184 1	1 (Other) 18	
sample_t; tumor 117		-			_		_	_		
histolog ser 117	ical_typ	e:								
primarys ov 117	ite:									
summaryg high lo 57 5	w NA's									
summarys early 1 1	ate NA'									
	ge: 3 4 N. 8 17	A's 1								
	2 3 3 56	4 NA's 1 3								
days_to_ Min. 30	death: 1st Qu. 510	Median 1020	Mean 3 1496	Brd Qu. 2220	Max. 5550					
vital_st deceased 67	livin	-								

```
primary_therapy_outcome_success:
  completeresponse progressivedisease
                85
                                   32
debulking:
   optimal suboptimal
        63
                   54
batch:
2002-09-20 2002-10-23 2002-11-12 2002-12-16 2002-12-21 2003-01-03 2003-05-30
        10
                   8
                               9
                                          1
                                                      3
                                                                11
                                                                           10
2004-03-09 2004-03-16 2004-04-20 2004-05-18 2004-05-21 2004-05-27 2004-06-22
                               5
                                        15
                                                      7
                                                                 7
        16
                    6
                                                                            1
2004-06-23
         8
uncurated_author_metadata:
                        OVC.TumorID: 1024///Survival: 13///X0...alive...1...dead: 1
                       OVC.TumorID: 1447///Survival: 75///X0...alive...1...dead: 1/
                       OVC.TumorID: 1451///Survival: 132///X0...alive...1...dead: 1
                        OVC.TumorID: 1504///Survival: 108///X0...alive...1...dead:
                       OVC.TumorID: 1526///Survival: 74///X0...alive...1...dead: 1/
                       OVC.TumorID: 1552///Survival: 33///X0...alive...1...dead: 1/
                       OVC.TumorID: 1578///Survival: 33///X0...alive...1...dead: 1/
                        OVC.TumorID: 1590///Survival: 148///X0...alive...1...dead:
                       OVC.TumorID: 1615///Survival: 13///X0...alive...1...dead: 1/
                        OVC.TumorID: 1623///Survival: 147///X0...alive...1...dead:
                       OVC.TumorID: 1665///Survival: 15///X0...alive...1...dead: 1/
                        OVC.TumorID: 1674///Survival: 18///X0...alive...1...dead: 1
                      OVC.TumorID: 1675///Survival: 34///X0...alive...1...dead: 1/,
                      OVC.TumorID: 1774///Survival: 22///X0...alive...1...dead: 1/,
                        OVC.TumorID: 1784///Survival: 78///X0...alive...1...dead: 1
```

OVC.TumorID: 1834///Survival: 118///X0...alive...1...dead: 1 OVC.TumorID: 1846///Survival: 142///X0...alive...1...dead: OVC.TumorID: 1877///Survival: 119///X0...alive...1...dead: OVC.TumorID: 1913///Survival: 32///X0...alive...1...dead: 1/ OVC.TumorID: 1929///Survival: 134///X0...alive...1...dead: OVC.TumorID: 2046///Survival: 127///X0...alive...1...dead: OVC.TumorID: 2063///Survival: 16///X0...alive...1...dead: 1/, OVC.TumorID: 2064///Survival: 27///X0...alive...1...dead: 1///Ass OVC.TumorID: 2075///Survival: 87///X0...alive...1...dead: OVC.TumorID: 2198///Survival: 91///X0...alive...1...dead: OVC.TumorID: 2204///Survival: 118///X0...alive...1...dead: OVC.TumorID: 2324///Survival: 98///X0...alive...1...dead: OVC.TumorID: 2419///Survival: 107///X0...alive...1...dead: (OVC.TumorID: 2422///Survival: 20///X0...alive...1...dead: OVC.TumorID: 2424///Survival: 16///X0...alive...1...dead: 1/ OVC.TumorID: 2465///Survival: 17///X0...alive...1...dead: 1/, OVC.TumorID: 2476///Survival: 86///X0...alive...1...dead: 1/, OVC.TumorID: 2479///Survival: 95///X0...alive...1...dead: 0// OVC.TumorID: 2505///Survival: 95///X0...alive...1...dead: (OVC.TumorID: 2542///Survival: 36///X0...alive...1...dead: OVC.TumorID: 2573///Survival: 7///X0...alive...1...dead: 1// OVC.TumorID: 2673///Survival: 74///X0...alive...1...dead: 0// OVC.TumorID: 2739///Survival: 67///X0...alive...1...dead: (OVC.TumorID: 2802///Survival: 24///X0...alive...1...dead: 1/

PMID17290060

OVC.TumorID: 2849///Survival: 23///X0...alive...1...dead: 1/ OVC.TumorID: 2895///Survival: 9///X0...alive...1...dead: 1/ OVC.TumorID: 2967///Survival: 22///X0...alive...1...dead: 1 OVC.TumorID: 2981///Survival: 6///X0...alive...1...dead: 1/ OVC.TumorID: 2999///Survival: 16///X0...alive...1...dead: 1/ OVC.TumorID: 3018///Survival: 16///X0...alive...1...dead: 1/ OVC.TumorID: 3090///Survival: 16///X0...alive...1...dead: 1/, OVC.TumorID: 3102///Survival: 10///X0...alive...1...dead: 1// OVC.TumorID: 3107///Survival: 31///X0...alive...1...dead: 1/ OVC.TumorID: 3142///Survival: 18///X0...alive...1...dead: 1 OVC.TumorID: 860///Survival: 17///X0...alive...1...dead: 1/ OVC.TumorID: 872///Survival: 185///X0...alive...1...dead: 0/ OVC.TumorID: 922///Survival: 183///X0...alive...1...dead: OVC.TumorID: D1805///Survival: 9///X0...alive...1...dead: 1/ OVC.TumorID: D1837///Survival: 83///X0...alive...1...dead: 0// OVC.TumorID: D1859///Survival: 110///X0...alive...1...dead: 1 OVC.TumorID: D2098///Survival: 42///X0...alive...1...dead: 1 OVC.TumorID: D2208///Survival: 2///X0...alive...1...dead: 0/// OVC.TumorID: D2332///Survival: 27///X0...alive...1...dead: 1 OVC.TumorID: D2342///Survival: 20///X0...alive...1...dead: 1/ OVC.TumorID: D2358///Survival: 9///X0...alive...1...dead: 1 OVC.TumorID: D2421///Survival: 12///X0...alive...1...dead: 1 OVC.TumorID: D2432///Survival: 34///X0...alive...1...dead: OVC.TumorID: D2433///Survival: 49///X0...alive...1...dead: 0// OVC.TumorID: D2480///Survival: 34///X0...alive...1...dead: 1//
OVC.TumorID: D2557///Survival: 62///X0...alive...1...dead: 0//
OVC.TumorID: D2559///Survival: 5///X0...alive...1...dead: 1//
OVC.TumorID: D2560///Survival: 91///X0...alive...1...dead: 0//
OVC.TumorID: D2572///Survival: 37///X0...alive...1...dead: 0//
OVC.TumorID: D2575///Survival: 37///X0...alive...1...dead: 1//
OVC.TumorID: D2576///Survival: 17///X0...alive...1...dead: 1//
OVC.TumorID: D2581///Survival: 63///X0...alive...1...dead: 0//
OVC.TumorID: D2603///Survival: 42///X0...alive...1...dead: 1//
OVC.TumorID: D2611///Survival: 2///X0...alive...1...dead: 1//
OVC.TumorID: D2640///Survival: 36///X0...alive...1...dead: 1///
OVC.TumorID: D2640///Survival: 1///X0...alive...1...dead: 1///
OVC.TumorID: D2648///Survival: 35///X0...alive...1...dead: 1///

OVC.TumorID: D2668///Survival: 40///X0...alive...1...dead OVC.TumorID: D2689///Survival: 45///X0...alive...1...dead: 0// OVC.TumorID: D2691///Survival: 63///X0...alive...1...dead: 0// OVC.TumorID: D2700///Survival: 74///X0...alive...1...dead: 0// OVC.TumorID: D2726///Survival: 71///X0...alive...1...dead: 0/

OVC.TumorID: D2727///Survival: 53///X0...alive...1...dead: 0// OVC.TumorID: D2733///Survival: 55///X0...alive...1...dead: 0// OVC.TumorID: D2738///Survival: 68///X0...alive...1...dead: 0// OVC.TumorID: D2749///Survival: 24///X0...alive...1...dead: 1// OVC.TumorID: D2776///Survival: 10///X0...alive...1...dead: 1//

PMID19318476

OVC.TumorID: M1054///Survival: 101///X0alive1dead: 0///Assig
OVC.TumorID: M1055///Survival: 13///X0alive1dead: 0///Assigned
OVC.TumorID: M120///Survival: 35///X0alive1dead: 1///Assigr
OVC.TumorID: M1241///Survival: 95///X0alive1dead: 0///Assigned.S
OVC.TumorID: M1390///Survival: 46///X0alive1dead: 0/
OVC.TumorID: M1503///Survival: 53///X0alive1dead: 1///Assigr
OVC.TumorID: M1572///Survival: 22///X0alive1dead: 1///Assigne
OVC.TumorID: M17///Survival: 17///X0alive1dead: 0///Assigned.Sta
OVC.TumorID: M1891///Survival: 12///X0alive1dead: 0///Assigned.Stage: 4///
OVC.TumorID: M2070///Survival: 65///X0alive1dead: 0///Assigned.S
OVC.TumorID: M2097///Survival: 58///X0alive1dead: 0///Assi
OVC.TumorID: M2184///Survival: 34///X0alive1dead: 0///Assigne

Value

An expression set

PMID19318476	Microarray analysis of early stage serous ovarian cancers shows pro-
	files predictive of favorable outcome.

Description

Although few women with advanced serous ovarian cancer are cured, detection of the disease at an early stage is associated with a much higher likelihood of survival. We previously used gene expression array analysis to distinguish subsets of advanced cancers based on disease outcome. In the present study, we report on gene expression of early-stage cancers and validate our prognostic model for advanced-stage cancers. Frozen specimens from 39 stage I/II, 42 stage III/IV, and 20 low malignant potential cancers were obtained from four different sites. A linear discriminant model was used to predict survival based upon array data. We validated the late-stage survival model and show that three of the most differentially expressed genes continue to be predictive of outcome. Most early-stage cancers (38 of 39 invasive, 15 of 20 low malignant potential) were classified as

PMID19318476

long-term survivors (median probabilities 0.97 and 0.86). MAL, the most differentially expressed gene, was further validated at the protein level and found to be an independent predictor of poor survival in an unselected group of advanced serous cancers (P = 0.0004). These data suggest that serous ovarian cancers detected at an early stage generally have a favorable underlying biology similar to advanced-stage cases that are long-term survivors. Conversely, most late-stage ovarian cancers seem to have a more virulent biology. This insight suggests that if screening approaches are to succeed it will be necessary to develop approaches that are able to detect these virulent cancers at an early stage.

Format

```
experimentData(eset):
Experiment data
 Experimenter name: Berchuck A, Iversen ES, Luo J, Clarke JP, Horne H, Levine DA,
  Laboratory: Berchuck, Lancaster 2009
  Contact information:
 Title: Microarray analysis of early stage serous ovarian cancers shows profiles p
  URL:
 PMIDs: 19318476
 Abstract: A 241 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [HG-U133A] Affymetrix Human Genome U133A Array
  platform_shorttitle:
      Affymetrix HG-U133A
  platform_summary:
      hgu133a
  platform_manufacturer:
      Affymetrix
  platform_distribution:
      commercial
  platform_accession:
      GPL96
   warnings:
      These samples are a subset of PMID17290060.
   version:
      2015-09-22 20:20:30
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-HUMISGF3A/M97935_MB_at
    (20967 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 20967 features, 42 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
    n events median 0.95LCL 0.95UCL
 42.00 22.00 2.79 2.30 NA
_____
Available sample meta-data:
_____
alt sample name:
D1462 D1805 D2171 D2208 D2247 D2332 D2432 D2480 D2559 D2560 D2575 D2576 D2611
  D2629 D2640 D2648 D2736 D2749 D2776 D2792 M1025 M1054 M1055 M120 M1241 M1572
  1 1 1 1 1 1 1 1 1 1 1 1
 M17 M1777 M1891 M2184 M2515 M2807 M3035 M337 M3484 M359 M4161 M444 M503
  1 1 1 1 1 1 1 1 1 1
                                                1
                                                    1
M5668 M5775 M806
  1 1 1
sample_type:
tumor
 42
histological_type:
ser
42
summarygrade:
high low NA's
24 17 1
summarystage:
early late NA's
 2 39 1
tumorstage:
  1 2 3 4 NA's
    1 29 10 1
  1
substage:
  a b c NA's
  1 1 29 11
grade:
```

PMID19318476

1 2 3 NA's 2 15 24 1 age_at_initial_pathologic_diagnosis:
 Min. 1st Qu.
 Median
 Mean 3rd Qu.
 Max.
 NA's

 33.00
 55.00
 62.00
 61.46
 70.00
 81.00
 1
 recurrence_status: norecurrence recurrence 6 36 days_to_death: Min. 1st Qu. Median Mean 3rd Qu. Max. 30.0 367.5 825.0 1105.0 1050.0 3420.0 vital_status: deceased living 22 20 debulking: optimal suboptimal NA's 20 21 1 batch: 2004-03-09 2004-03-16 2004-04-20 2004-05-18 2004-05-21 2004-05-27 2004-06-22 14 3 4 8 6 5 1 2004-06-23 1

uncurated_author_metadata:

PMID19318476

Tumor: D2560///NEW.Response: CR///SHORT.LONG: NA///AgeDx: 60///DateDx: 5/14/1996//

Value

An expression set

TCGA.RNASeqV2

Integrated genomic analyses of ovarian carcinoma.

Description

A catalogue of molecular aberrations that cause ovarian cancer is critical for developing and deploying therapies that will improve patients' lives. The Cancer Genome Atlas project has analysed messenger RNA expression, microRNA expression, promoter methylation and DNA copy number in 489 high-grade serous ovarian adenocarcinomas and the DNA sequences of exons from coding genes in 316 of these tumours. Here we report that high-grade serous ovarian cancer is characterized by TP53 mutations in almost all tumours (96%); low prevalence but statistically recurrent somatic mutations in nine further genes including NF1, BRCA1, BRCA2, RB1 and CDK12; 113 significant focal DNA copy number aberrations; and promoter methylation events involving 168 genes. Analyses delineated four ovarian cancer transcriptional subtypes, three microRNA subtypes, four promoter methylation subtypes and a transcriptional signature associated with survival duration, and shed new light on the impact that tumours with BRCA1/2 (BRCA1 or BRCA2) and CCNE1 aberrations have on survival. Pathway analyses suggested that homologous recombination is defective in about half of the tumours analysed, and that NOTCH and FOXM1 signalling are involved in serous ovarian cancer pathophysiology.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Integrated genomic analyses of ovarian carcinoma. Nature 2011,
 Laboratory: Cancer Genome Atlas Research Network 2011
  Contact information:
  Title: Integrated genomic analyses of ovarian carcinoma.
  URL:
  PMIDs: 21720365
  Abstract: A 179 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [RNASeqV2] Illumina HiSeq RNA sequencing
   platform_shorttitle:
      Illumina HiSeq RNA sequencing
   platform_summary:
      NA
   platform_manufacturer:
      Illumina
   platform distribution:
      sequencing
  platform_accession:
      NA
  platform_technology:
     RNA sequencing
   version:
      2015-09-22 20:27:26
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: ?|100133144 ?|100134869 ... ZZZ3|26009 (20471 total)
 varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 20471 features, 261 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
5 observations deleted due to missingness
n events median 0.95LCL 0.95UCL
256.00 143.00 3.62 3.19 4.03
```

Available sample meta-data:

alt sample name: TCGA-04-1348-01A-01R-1565-13 TCGA-04-1357-01A-01R-1565-13 TCGA-04-1362-01A-01R-1565-13 TCGA-04-1364-01A-01R-1565-13 TCGA-04-1365-01A-01R-1565-13 TCGA-04-1514-01A-01R-1566-13 TCGA-04-1519-01A-01R-1565-13 TCGA-09-0364-01A-02R-1564-13 TCGA-09-0366-01A-01R-1564-13 TCGA-09-0367-01A-01R-1564-13 TCGA-09-0369-01A-01R-1564-13 TCGA-09-1662-01A-01R-1566-13 TCGA-09-1666-01A-01R-1566-13 TCGA-09-1667-01C-01R-1566-13 TCGA-09-1668-01B-01R-1566-13 TCGA-09-1669-01A-01R-1566-13 TCGA-09-1670-01A-01R-1566-13 TCGA-09-1673-01A-01R-1566-13 TCGA-09-1674-01A-01R-1566-13 TCGA-09-2044-01B-01R-1568-13 TCGA-09-2045-01A-01R-1568-13 TCGA-09-2048-01A-01R-1568-13 TCGA-09-2051-01A-01R-1568-13 TCGA-09-2054-01A-01R-1568-13 TCGA-09-2056-01B-01R-1568-13 TCGA-10-0928-01A-02R-1564-13 TCGA-10-0936-01A-01R-1564-13 TCGA-13-0730-01A-01R-1564-13 TCGA-13-0799-01A-01R-1564-13 TCGA-13-0800-01A-01R-1564-13 TCGA-13-0801-01A-01R-1564-13 TCGA-13-0890-01A-01R-1564-13 TCGA-13-0893-01B-01R-1565-13 TCGA-13-0897-01A-01R-1564-13 TCGA-13-0899-01A-01R-1564-13 TCGA-13-0913-01A-01R-1564-13 TCGA-13-0916-01A-01R-1564-13 TCGA-13-0920-01A-01R-1564-13 TCGA-13-0924-01A-01R-1564-13 TCGA-13-1403-01A-01R-1565-13 TCGA-13-1405-01A-01R-1565-13 TCGA-13-1410-01A-01R-1565-13 TCGA-13-1481-01A-01R-1565-13 TCGA-13-1497-01A-01R-1565-13

1	1
TCGA-13-1498-01A-01R-1565-13	TCGA-13-1505-01A-01R-1565-13
TCGA-13-1506-01A-01R-1565-13	TCGA-13-1507-01A-01R-1565-13
TCGA-13-1511-01A-01R-1565-13	TCGA-13-1512-01A-01R-1565-13
TCGA-13-2060-01A-01R-1568-13	TCGA-20-1682-01A-01R-1564-13
TCGA-20-1683-01A-01R-1566-13	TCGA-20-1684-01A-01R-1566-13
TCGA-20-1685-01A-01R-1566-13	TCGA-20-1687-01A-01R-1566-13
TCGA-23-1023-01A-02R-1564-13	TCGA-23-1026-01B-01R-1569-13
TCGA-23-1027-01A-02R-1564-13	TCGA-23-1029-01B-01R-1567-13
TCGA-23-1109-01A-01R-1564-13	TCGA-23-1111-01A-01R-1567-13
TCGA-23-1114-01B-01R-1566-13	TCGA-23-1120-01A-02R-1565-13
TCGA-23-1122-01A-01R-1565-13	TCGA-23-1123-01A-01R-1565-13
TCGA-23-1809-01A-01R-1566-13	TCGA-23-2077-01A-01R-1568-13
TCGA-23-2081-01A-01R-1568-13	TCGA-23-2084-01A-02R-1568-13
TCGA-24-0975-01A-02R-1565-13	TCGA-24-1103-01A-01R-1565-13
1 TCGA-24-1413-01A-01R-1565-13	1 TCGA-24-1416-01A-01R-1565-13
TCGA-24-1417-01A-01R-1565-13	1 TCGA-24-1418-01A-01R-1565-13
TCGA-24-1419-01A-01R-1565-13	1 TCGA-24-1423-01A-01R-1565-13
TCGA-24-1424-01A-01R-1565-13	1 TCGA-24-1427-01A-01R-1565-13
	1 TCGA-24-1430-01A-01R-1566-13
	TCGA-24-1467-01A-01R-1566-13
	TCGA-24-1474-01A-01R-1566-13
	1 TCGA-24-1548-01A-01R-1566-13
	1 TCGA-24-1550-01A-01R-1566-13
1 TCGA-24-1551-01A-01R-1566-13	1 TCGA-24-1552-01A-01R-1566-13

TCGA-24-1553-01A-01R-1566-13 TCGA-24-1555-01A-01R-1566-13						
TCGA-24-1556-01A-01R-1566-13 TCGA-24-1557-01A-01R-1566-13						
TCGA-24-1558-	-01A-01R-1566-	-13 TCGA-24-15	560-01A-01R-15	1 566-13 1		
TCGA-24-1562-	-01A-01R-1566-	-13	((Dther)		
		1		162		
unique_patien	nt_ID:					
TCGA-04-1348	TCGA-04-1357 1	TCGA-04-1362	TCGA-04-1364 1	TCGA-04-1365 1	TCGA-04-1514 1	
TCGA-04-1519	TCGA-09-0364 1	TCGA-09-0366	TCGA-09-0367 1	TCGA-09-0369	TCGA-09-1662 1	
TCGA-09-1666	TCGA-09-1667	-	TCGA-09-1669	-	TCGA-09-1673	
TCGA-09-1674	TCGA-09-2044	TCGA-09-2045	TCGA-09-2048	TCGA-09-2051	TCGA-09-2054	
TCGA-09-2056	1 TCGA-10-0928	1 TCGA-10-0936		TCGA-13-0799	TCGA-13-0800	
1 TCGA-13-0801	1 TCGA-13-0890	1 TCGA-13-0893		1 TCGA-13-0899	TCGA-13-0913	
1 TCGA-13-0916	1 TCGA-13-0920	1 TCGA-13-0924		1 TCGA-13-1405	1 TCGA-13-1410	
1 TCGA-13-1481	1 TCGA-13-1497	1 TCGA-13-1498	1 TCGA-13-1505	1 TCGA-13-1506	1 TCGA-13-1507	
1 TCGA-13-1511	1 TCGA-13-1512	1 TCGA-13-2060	1 TCGA-20-1682	1 TCGA-20-1683	1 TCGA-20-1684	
1 TCGA-20-1685	1 TCGA-20-1687	1 TCGA-23-1023	1 TCGA-23-1026	1 TCGA-23-1027	1 TCGA-23-1029	
1 TCGA-23-1109	1 TCGA-23-1111	1 TCGA-23-1114	1 TCGA-23-1120	1 TCGA-23-1122	1 TCGA-23-1123	
1 TCGA-23-1809	1 TCGA-23-2077	1 TCGA-23-2081	1 TCGA-23-2084	1 TCGA-24-0975	1 TCGA-24-1103	
1 TCGA-24-1413	1 TCGA-24-1416	1 TCGA-24-1417	1 TCGA-24-1418	1 TCGA-24-1419	1 TCGA-24-1423	
1 TCGA-24-1424	1 TCGA-24-1427	1 TCGA-24-1428	1 TCGA-24-1430	1 TCGA-24-1436	1 TCGA-24-1467	
1 TCGA-24-1469	1 TCGA-24-1474	1 TCGA-24-1544	1 TCGA-24-1548	1 TCGA-24-1549	1 TCGA-24-1550	
1 TCGA-24-1551	1 TCGA-24-1552	1 TCGA-24-1553	1 TCGA-24-1555	1 TCGA-24-1556	1 TCGA-24-1557	
1	1 TCGA-24-1560	1	1	1	1	
10GA-24-1558 1	10GA-24-1500 1	10GA-24-1562 1				

1

sample_type:
tumor

```
261
histological_type:
ser
261
primarysite:
other ov
  1 260
summarygrade:
high low NA's
226 29 6
summarystage:
early late NA's
  18 242 1
tumorstage:
 2 3 4 NA's
18 209 33 1
substage:
  b c NA's
 16 211 34
grade:
  1 2 3 4 NA's
  1 28 225
             1 6
age_at_initial_pathologic_diagnosis:
  Min. 1st Qu. Median Mean 3rd Qu. Max.
  34.00 51.00 58.00 58.84 66.00 87.00
pltx:
 n y NA's
 17 215 29
tax:
 n y NA's
 17 215 29
neo:
 n NA's
232 29
days_to_tumor_recurrence:
  Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
```

9.0 225.0 426.5 585.3 755.0 5480.0 19 recurrence_status: norecurrence recurrence 123 138 days_to_death: 9.0341.8878.01018.01446.05480.05 Min. 1st Qu. Median Mean 3rd Qu. Max. vital_status: deceased living NA's 143 114 4 site_of_tumor_first_recurrence: locoregional metastasis NA's 82 56 123 primary_therapy_outcome_success: completeresponse partialresponse progressivedisease stabledisease 15 147 30 15 NA's 54 debulking: optimal suboptimal NA's 60 30 171 percent_normal_cells: Min. 1st Qu. Median Mean 3rd Qu. Max. NA's 0.000 0.000 0.000 2.066 0.000 55.000 5 percent_stromal_cells: Min. 1st Qu. Median Mean 3rd Qu. Max. NA's 0.00 5.00 10.00 11.43 15.00 70.00 4 percent_tumor_cells: Min. 1st Qu. Median Mean 3rd Qu. Max. NA's 0.00 77.00 85.00 82.07 90.00 100.00 4

uncurated_author_metadata:

age_at_initial_pathologic_diagnosis: 38///anatomic_organ_subdivision: Bilateral///k

age_at_initial_

age_at_ir

age_at_initial_pathologic_diagr

age_at_initial_pathologic_diagnosis: 4

age_at_initial_pathologic_diagnosi

age_at_ir

age_at_initial_pathologic_diagnosis: 42///anatomic_organ_subdivi

age_at_initial_pathologic_diagnosis: 4

age_at_init

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age_at_initial_path
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age_at_initial_pathol

ć

```
age_at_initial_patholog
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ac

age_at_initial_p

age_at_initial_pathologic_diagnosis: 45///anatomic_or

age_at

age_at_initial_pathologic_diagnosis: 45///anato

age_at_initial_patholog

age_at_initial_pathold

age_at_initial_pathologic_diagnosis

age_at_initial_pathologic_diagnosis: 45///anatomic_organ_subdivision:

age_at_initial_pathologic_dia

age_at_initial_pathologic_diagnosis: 46///anatomic_organ_subdivision:

age_at_initial_pathologic_diagnosis: 47

age_at_initial_pathologic_diagnosis

age_at_initial_pathologic_diagnosis: 47///anatomic

age_at_initial_

age_at_initial_pathologic_diagnosis: 47///anatomic_org

age_at_initial_pathologic_diagnosis: 48///ana

age_at

age_

age_at_initial_pathologic_dia

```
age_at_initi
```

```
age_at_initial_pathologic_diagnosis: 49///anatomic_
```

age_at_initial_pathologic_diagnosis: 50///anatomic_organ_

age_at_initial_pathologic_diagno

age_at_initial_pathol

age_at_initial_pathologic_diagnosis: 50///anatomic_organ_subdivision: Left///bcr_p

age_at_initial_pathologic_diagnosis: 50///anator

age_at_initial_pathologi

age_at_initial_pathologic_diagnosis: 51///anatomic_organ_subdivision: Bilateral//

age_at_initial

age_at_initial_pathologic_diagno

age_at_ir

age_at

age_at_initial_pathologic_diagnosis: 51///anatomi

age_at_initial_pathologic_di

age_at_initial_p

age

age_at_initial_pathologic_c

age_at_initial_pathologic_diagr

age_at_

age_at_initial_pathologic_diagnosis:

age_at_initial_pathol

age_at_initial_pathologic_diagr

age_at_initial_pathologic_diagnosis: 53///anatomic_organ_suk

age_at_initial_pathologic_diagnosis:

age_at_initial_pathologic_diagnosis: 53///anatomic

age_at_initial_pathol

age_at_initial_pathologic_diagnosis: 54///anatomic_organ_subdivisior

age_at_i

age_at_initia

age_at_init

age_at_initial_pathologic_diagnosis: 54///anatomic_organ_subdivisi

age

Value

An expression set

TCGAOVARIAN

Integrated genomic analyses of ovarian carcinoma.

Description

A catalogue of molecular aberrations that cause ovarian cancer is critical for developing and deploying therapies that will improve patients' lives. The Cancer Genome Atlas project has analysed messenger RNA expression, microRNA expression, promoter methylation and DNA copy number in 489 high-grade serous ovarian adenocarcinomas and the DNA sequences of exons from coding genes in 316 of these tumours. Here we report that high-grade serous ovarian cancer is characterized by TP53 mutations in almost all tumours (96%); low prevalence but statistically recurrent somatic mutations in nine further genes including NF1, BRCA1, BRCA2, RB1 and CDK12; 113 significant focal DNA copy number aberrations; and promoter methylation events involving 168 genes. Analyses delineated four ovarian cancer transcriptional subtypes, three microRNA subtypes, four promoter methylation subtypes and a transcriptional signature associated with survival duration, and shed new light on the impact that tumours with BRCA1/2 (BRCA1 or BRCA2) and CCNE1 aberrations have on survival. Pathway analyses suggested that homologous recombination is defective in about half of the tumours analysed, and that NOTCH and FOXM1 signalling are involved in serous ovarian cancer pathophysiology.

Format

```
experimentData(eset):
Experiment data
  Experimenter name: Integrated genomic analyses of ovarian carcinoma. Nature 2011,
 Laboratory: Cancer Genome Atlas Research Network 2011
  Contact information:
  Title: Integrated genomic analyses of ovarian carcinoma.
  URL:
  PMIDs: 21720365
  Abstract: A 179 word abstract is available. Use 'abstract' method.
  Information is available on: preprocessing
  notes:
  platform_title:
      [HT_HG-U133A] Affymetrix HT Human Genome U133A Array
  platform_shorttitle:
      Affymetrix HT_HG-U133A
  platform_summary:
      hthgu133a
```

```
platform_manufacturer:
      Affymetrix
   platform distribution:
      commercial
   platform accession:
      GPL3921
   warnings:
      The following samples are likely from specimens also used in GSE26712: TCG
A.13.0725, TCGA.13.0885, TCGA.13.0887, TCGA.13.0890, TCGA.13.0886, TCGA.13
.0714, TCGA.13.0727, TCGA.13.1817, TCGA.13.1499, TCGA.13.0883
   version:
      2015-09-22 20:25:15
featureData(eset):
An object of class 'AnnotatedDataFrame'
  featureNames: 1007_s_at 1053_at ... AFFX-M27830_M_at (21260 total)
  varLabels: probeset gene EntrezGene.ID best_probe
  varMetadata: labelDescription
```

Details

```
assayData: 21260 features, 578 samples
Platform type:
Overall survival time-to-event summary (in years):
Call: survfit(formula = Surv(time, cens) ~ -1)
  21 observations deleted due to missingness
     n events median 0.95LCL 0.95UCL
 557.00 290.00 3.73 3.45 4.06
 _____
Available sample meta-data:
_____
alt_sample_name:
TCGA-01-0628-11A-01R-0362-01 TCGA-01-0630-11A-01R-0362-01
                         1
                                                     1
TCGA-01-0631-11A-01R-0362-01 TCGA-01-0633-11A-01R-0362-01
                         1
                                                     1
TCGA-01-0636-11A-01R-0362-01 TCGA-01-0637-11A-01R-0362-01
                         1
                                                    1
TCGA-01-0639-11A-01R-0362-01 TCGA-01-0642-11A-02R-0362-01
                         1
                                                     1
TCGA-04-1331-01A-01R-0434-01 TCGA-04-1332-01A-01R-0434-01
                                                     1
                         1
TCGA-04-1335-01A-01R-0434-01 TCGA-04-1336-01A-01R-0434-01
```

1

TCGA-04-1337-01A-01R-0434-01 TCGA-04-1338-01A-01R-0434-01

1	1
TCGA-04-1341-01A-01R-0434-01	TCGA-04-1342-01A-01R-0434-01
TCGA-04-1343-01A-01R-0434-01	TCGA-04-1346-01A-01R-0434-01
TCGA-04-1347-01A-01R-0434-01	TCGA-04-1348-01A-01R-0453-01
TCGA-04-1349-01A-01R-0453-01	TCGA-04-1350-01A-01R-0453-01
TCGA-04-1351-01A-01R-0453-01	TCGA-04-1353-01A-01R-1048-01
TCGA-04-1356-01A-01R-0453-01	TCGA-04-1357-01A-01R-0453-01
TCGA-04-1360-01A-01R-0453-01	TCGA-04-1361-01A-01R-0453-01
TCGA-04-1362-01A-01R-0453-01	TCGA-04-1364-01A-01R-0453-01
TCGA-04-1365-01A-01R-0453-01	TCGA-04-1367-01A-01R-0453-01
TCGA-04-1369-01A-02R-1048-01	TCGA-04-1371-01A-01R-0453-01
TCGA-04-1514-01A-01R-0502-01	TCGA-04-1516-01A-01R-1048-01
TCGA-04-1517-01A-01R-0538-01	TCGA-04-1519-01A-01R-0538-01
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TCGA-04-1536-01A-01R-0538-01	TCGA-04-1542-01A-01R-0502-01
TCGA-04-1638-01A-01R-0582-01	TCGA-04-1644-01B-01R-1048-01
TCGA-04-1646-01A-01R-0582-01	TCGA-04-1648-01A-01R-0582-01
TCGA-04-1649-01A-01R-0582-01	TCGA-04-1651-01A-01R-0582-01
TCGA-04-1652-01A-01R-0582-01	TCGA-04-1654-01A-02R-0653-01
	TCGA-09-0364-01A-02R-0362-01
TCGA-09-0365-01A-02R-0362-01	TCGA-09-0366-01A-01R-0362-01
TCGA-09-0367-01A-01R-0362-01	
TCGA-09-1659-01B-01R-0538-01 1	
	TCGA-09-1664-01A-01R-0582-01
	TCGA-09-1666-01A-01R-0538-01

1	1	
TCGA-09-1667-01C-01R-0538-01	TCGA-09-1668-01B-01R-0538-01 1	
TCGA-09-1669-01A-01R-0538-01	TCGA-09-1670-01A-01R-0564-01 1	
TCGA-09-1672-01A-01R-0564-01	TCGA-09-1673-01A-01R-0564-01 1	
_	TCGA-09-1675-01B-01R-0564-01 1	
-	TCGA-09-2044-01B-01R-0709-01	
_	TCGA-09-2048-01A-01R-0709-01 1	
_	TCGA-09-2050-01A-01R-0709-01 1	
TCGA-09-2051-01A-01R-0709-01	-	
TCGA-09-2054-01A-01R-0668-01	TCGA-09-2055-01B-01R-0709-01 1	
TCGA-09-2056-01B-01R-0668-01	TCGA-10-0925-01B-01R-0653-01 1	
-	TCGA-10-0927-01A-02R-0404-01	
-	TCGA-10-0930-01A-02R-0404-01	
-	TCGA-10-0933-01A-01R-0404-01 1	
_	TCGA-10-0935-01A-02R-0404-01 1	
_	TCGA-10-0937-01A-02R-0404-01 1	
-	TCGA-13-0714-01A-01R-0362-01 1	
-	TCGA-13-0720-01A-01R-0362-01	
TCGA-13-0723-01A-02R-0362-01	TCGA-13-0724-01A-01R-0362-01	
(Other) 479	NA's 1	
1,3	±	

unique_patient_ID:							
TCGA-01-0628	TCGA-01-0630	TCGA-01-0631	TCGA-01-0633	TCGA-01-0636	TCGA-01-0637		
1	1	1	1	1	1		
TCGA-01-0639	TCGA-01-0642	TCGA-04-1331	TCGA-04-1332	TCGA-04-1335	TCGA-04-1336		
1	1	1	1	1	1		
TCGA-04-1337	TCGA-04-1338	TCGA-04-1341	TCGA-04-1342	TCGA-04-1343	TCGA-04-1346		
1	1	1	1	1	1		
TCGA-04-1347	TCGA-04-1348	TCGA-04-1349	TCGA-04-1350	TCGA-04-1351	TCGA-04-1353		

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TCGA-04-1356	TCGA-04-1357	TCGA-04-1360	TCGA-04-1361	TCGA-04-1362	
1 TCGA-04-1365	1 TCGA-04-1367	1 TCGA-04-1369	1 TCGA-04-1371	1 TCGA-04-1514	1 TCGA-04-1516 1
TCGA-04-1517	TCGA-04-1519	TCGA-04-1525	TCGA-04-1530	TCGA-04-1536	TCGA-04-1542 1
TCGA-04-1638	TCGA-04-1644 1	TCGA-04-1646	TCGA-04-1648	_	TCGA-04-1651 1
TCGA-04-1652 1	TCGA-04-1654 1	TCGA-04-1655 1	TCGA-09-0364 1	TCGA-09-0365 1	TCGA-09-0366 1
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1	1	TCGA-09-1667 1	1	1	1
1	1	TCGA-09-1674 1	1	1	TCGA-09-2044 1
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1CGA-09-2034 1 TCGA-10-0928	1	TCGA-09-2056 1 TCGA-10-0931	1	1	1
1	1	1 TCGA-10-0938	1	1	1
1	1 TCGA-13-0724	1	(Other)	1	1
1	1	1	479		
sample_type: adjacentnorm	al ti	umor			
	8	570			
histological ser NA's 568 10	_type:				
primarysite: other ov 4 564	NA's 10				
summarygrade high low NA 480 75 :					
summarystage early late 43 520	: NA's 15				

```
tumorstage:
1 2 3 4 NA's
 16 27 436 84 15
substage:
  b c NA's
  31 448 99
grade:
  1 2 3 4 NA's
   6 69 479 1 23
age_at_initial_pathologic_diagnosis:
  Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
  26.00 51.00 59.00 59.70 68.25 89.00 10
pltx:
 n y NA's
 19 492 67
tax:
 n y NA's
 43 468 67
neo:
 n NA's
 511 67
days_to_tumor_recurrence:
  Min. 1st Qu.MedianMean 3rd Qu.Max.NA's8.0238.2443.5623.7812.05480.056
recurrence_status:
norecurrence recurrence
       279
              299
days_to_death:
  Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
8 349 881 1010 1446 5480 21
vital status:
deceased living NA's
290 270 18
site_of_tumor_first_recurrence:
               locoregional locoregional_plus_metastatic
                       153
                                                   3
                metastasis
                                                 NA's
```

	143				279		
NA	- –			ogressiv	edisease 41	stabledisease 30	
debulking: optimal subop 367	timal 140	NA's 71					
percent_normal_c Min. 1st Qu. 0.000 0.000	Median		~		NA's 19		
percent_stromal_ Min. 1st Qu. 0.00 5.00	Median				NA's 25		
percent_tumor_ce Min. 1st Qu. 0.00 75.00	Median				NA's 22		
batch: Min. 1st Qu. 9.00 13.00	Median 17.00		~		NA's 1		

uncurated_author_metadata:

age_at_initial_pathologic_diagnosis:

age_at

age_at_initial_pathologic_

age_at_initial_pathologic_diagnosis: 37///ar

age_at_initial_pathologic_diagnosis: 38///anatomic_organ_subdivision: Bilateral///k

- age_at_initial_pathologic_diagnosis: 38///anatomic_organ_subdivision: Bil
 - age_at_initial_
 - age_at_ir
 - age_at_initial_pathologic_diagnosis: 39///ana
 - age_at_initial_pathologic_dia
 - age_at_initial_pathologic_diagr
 - age_at_initial_pathologic_diagnosis: 4
 - age_at_initial_pathologic_diagnosis: 40///anatomic_organ_su
 - age_at_initial_pathologic_diagnosi
 - age_at_ir
 - age_at_initial_patho
 - age_at_initial_pathologic_diag
 - age_at_initial_pathologic_diagnosis: 4
 - age_at_initial_pathologic_diagnosis: 42///anatomic_organ_subdivi
 - age_at_initial_pat
 - age_at_initial_pathologic_diagnosis: 42///anatomic_org
 - age_at_initial_pathol
 - age_at_initial_pathologic_diagnosis: 4

age_at_ini

age

age_at_initial_pathologic_diagnosis: 4

age_at_initial

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age_at_initial_pathologic_diagno

age_at_initial_pathologic_diagnosis: 44///anatomic_c

age_at_initial_pathologic_diagr

age_at_initial_path

age_at_initial_patho

age_at_initial_pathol

ć

age_at_initial_patholog

ac

age_at_initial_p

age_at_initial_pathologic_diagnosis: 45///anatomic_or

211

age_at

age_at_initial_pathologic_diagnosis: 45///anato

age_at_initial_patholog

age_at_initial_pathold

age_at_initial_pathologic_diagnosis

age_at_initial_pathologic_diagnosis: 45///anatomic_organ_subdivision:

age_at_initial_pathologic_dia

age_at_initial_pathologic_diagnosis: 46///anatomic_organ_subdivision

age_at_initial_pathologic_diagnosis: 46///anato

age_at_initial_pathologic_diagnosis: 47

age_at_initial_pathologic

age_at_initial_pathologic_diagnosis

age_at_initial_pathologic_diagnosis

age_at_initial_pathologic_diagnosis: 47///anatomic

age_at_initial_

age_at_initial_pathologic_diagnosis: 47///anatomic_org

age_at_initial_pathologic_diagnosis: 48///ana

age_at_initial_pathologic_diagnosis

age_at_initial_pathologic_di

age_at_initial_pathologic_diagnosis: 48///ana

duplicates: Length Class Mode 578 character character

Value

An expression set