

# mrr\_project\_bio

2025-11-10

## Classification of Cancer outcome using Genetic and Clinical data

### Introduction

This project investigates the effect of genetic and clinical variables on the survival outcome of breast cancer patients. The data contains records of more than 1000 breast cancer patients from several research institutions. Clinical data contains patient-related and tumor-related information. Additionally mRNA gene expression data is available for each patient. The gene expression data has been processed to include only the top 5000 most variable genes on the transformed scale ( $\log_2(\text{counts} + 1)$ ).

We consider that the main outcome variable of interest is **vital\_status**, which is defined in clinical data.

### Load data

```
load("mrr_bio.Rdata")
```

```
# genetic data: n x p matrix  
dim(GeneX) # 1231 x 5000
```

```
## [1] 1231 5000
```

```
str(GeneX)
```

```
## num [1:1231, 1:5000] 0 0 10.92 3.17 5 ...  
## - attr(*, "dimnames")=List of 2  
## ..$ : chr [1:1231] "EW-A2FS-01A-11R-A17B" "OL-A6VR-01A-32R-A33J" "E9-A226-01A-21R-A157" "A8-A08H-0  
## ..$ : chr [1:5000] "CLEC3A" "SCGB2A2" "CPB1" "GSTM1" ...
```

```
colnames(GeneX)[1:10] # first 10 names of genes
```

```
## [1] "CLEC3A" "SCGB2A2" "CPB1" "GSTM1" "TFF1" "SCGB1D2"  
## [7] "KCNJ3" "MUCL1" "LINC00993" "ANKRD30A"
```

```
# clinical data  
dim(clinical_data) # 1231 x 24
```

```
## Loading required namespace: S4Vectors
```

```
## NULL
```

```
head(clinical_data)
```

```
## DataFrame with 6 rows and 24 columns
##           initial_weight tissue_type laterality
##           <numeric> <character> <character>
## EW-A2FS-01A-11R-A17B          110      Tumor      Left
## OL-A6VR-01A-32R-A33J          380      Tumor      Right
## E9-A226-01A-21R-A157          510      Tumor      Left
## A8-A08H-01A-21R-A00Z          250      Tumor      Left
## D8-A27H-01A-11R-A16F          170      Tumor      Right
## D8-A3Z6-01A-11R-A239           60      Tumor      Left
##           tissue_or_organ_of_origin age_at_diagnosis
##           <character> <integer>
## EW-A2FS-01A-11R-A17B      Breast, NOS          15302
## OL-A6VR-01A-32R-A33J      Breast, NOS          17702
## E9-A226-01A-21R-A157      Breast, NOS          16511
## A8-A08H-01A-21R-A00Z      Breast, NOS          24137
## D8-A27H-01A-11R-A16F      Breast, NOS          26588
## D8-A3Z6-01A-11R-A239      Breast, NOS          20629
##           primary_diagnosis prior_treatment
##           <character> <character>
## EW-A2FS-01A-11R-A17B Infiltrating duct ca..      No
## OL-A6VR-01A-32R-A33J Infiltrating duct ca..      No
## E9-A226-01A-21R-A157 Infiltrating duct ca..      No
## A8-A08H-01A-21R-A00Z Infiltrating duct ca..      No
## D8-A27H-01A-11R-A16F Infiltrating duct ca..      No
## D8-A3Z6-01A-11R-A239 Lobular carcinoma, NOS      No
##           diagnosis_is_primary_disease ajcc_pathologic_t morphology
##           <logical> <character> <character>
## EW-A2FS-01A-11R-A17B          TRUE          T2      8500/3
## OL-A6VR-01A-32R-A33J          TRUE          T1b      8500/3
## E9-A226-01A-21R-A157          TRUE          T2      8500/3
## A8-A08H-01A-21R-A00Z          TRUE          T2      8500/3
## D8-A27H-01A-11R-A16F          TRUE          T2      8500/3
## D8-A3Z6-01A-11R-A239          TRUE          T3      8520/3
##           classification_of_tumor sites_of_involvement
##           <character> <list>
## EW-A2FS-01A-11R-A17B      primary Breast, Left Upper O..
## OL-A6VR-01A-32R-A33J      primary      Breast, NOS
## E9-A226-01A-21R-A157      primary Breast, Left Lower I..
## A8-A08H-01A-21R-A00Z      primary Breast, Left Upper I..
## D8-A27H-01A-11R-A16F      primary Breast, Right Upper ..
## D8-A3Z6-01A-11R-A239      primary      Breast, NOS
##           days_to_last_follow_up follow_ups_disease_response
##           <integer> <character>
## EW-A2FS-01A-11R-A17B          1604      TF-Tumor Free
## OL-A6VR-01A-32R-A33J          1220      TF-Tumor Free
## E9-A226-01A-21R-A157          1048      WT-With Tumor
## A8-A08H-01A-21R-A00Z           0      TF-Tumor Free
## D8-A27H-01A-11R-A16F          397      TF-Tumor Free
## D8-A3Z6-01A-11R-A239          563      TF-Tumor Free
##           race      gender      ethnicity
##           <character> <character> <character>
```

```
## EW-A2FS-01A-11R-A17B black or african ame.. female not hispanic or latino
## OL-A6VR-01A-32R-A33J black or african ame.. female not hispanic or latino
## E9-A226-01A-21R-A157 white female not hispanic or latino
## A8-A08H-01A-21R-A00Z not reported female not reported
## D8-A27H-01A-11R-A16F white female not hispanic or latino
## D8-A3Z6-01A-11R-A239 white female not hispanic or latino
## vital_status age_at_index days_to_birth age_is_obfuscated
## <character> <integer> <integer> <logical>
## EW-A2FS-01A-11R-A17B Alive 41 -15302 FALSE
## OL-A6VR-01A-32R-A33J Alive 48 -17702 FALSE
## E9-A226-01A-21R-A157 Dead 45 -16511 FALSE
## A8-A08H-01A-21R-A00Z Alive 66 -24137 FALSE
## D8-A27H-01A-11R-A16F Alive 72 -26588 FALSE
## D8-A3Z6-01A-11R-A239 Alive 56 -20629 FALSE
## bcr_patient_barcode primary_site
## <character> <list>
## EW-A2FS-01A-11R-A17B TCGA-EW-A2FS-01A Breast
## OL-A6VR-01A-32R-A33J TCGA-OL-A6VR-01A Breast
## E9-A226-01A-21R-A157 TCGA-E9-A226-01A Breast
## A8-A08H-01A-21R-A00Z TCGA-A8-A08H-01A Breast
## D8-A27H-01A-11R-A16F TCGA-D8-A27H-01A Breast
## D8-A3Z6-01A-11R-A239 TCGA-D8-A3Z6-01A Breast
## disease_type
## <list>
## EW-A2FS-01A-11R-A17B Adenomas and Adenoca.,Adnexal and Skin App.,Fibroepithelial Neop.,...
## OL-A6VR-01A-32R-A33J Adenomas and Adenoca.,Adnexal and Skin App.,Fibroepithelial Neop.,...
## E9-A226-01A-21R-A157 Adenomas and Adenoca.,Adnexal and Skin App.,Fibroepithelial Neop.,...
## A8-A08H-01A-21R-A00Z Adenomas and Adenoca.,Adnexal and Skin App.,Fibroepithelial Neop.,...
## D8-A27H-01A-11R-A16F Adenomas and Adenoca.,Adnexal and Skin App.,Fibroepithelial Neop.,...
## D8-A3Z6-01A-11R-A239 Adenomas and Adenoca.,Adnexal and Skin App.,Fibroepithelial Neop.,...
```

```
colnames(clinical_data) # variables
```

```
## [1] "initial_weight" "tissue_type"
## [3] "laterality" "tissue_or_organ_of_origin"
## [5] "age_at_diagnosis" "primary_diagnosis"
## [7] "prior_treatment" "diagnosis_is_primary_disease"
## [9] "ajcc_pathologic_t" "morphology"
## [11] "classification_of_tumor" "sites_of_involvement"
## [13] "days_to_last_follow_up" "follow_ups_disease_response"
## [15] "race" "gender"
## [17] "ethnicity" "vital_status"
## [19] "age_at_index" "days_to_birth"
## [21] "age_is_obfuscated" "bcr_patient_barcode"
## [23] "primary_site" "disease_type"
```

```
# outcome variable
y = clinical_data$vital_status
table(y)
```

```
## y
## Alive Dead
## 1029 201
```

## Exploratory data analysis: example

```
# summary statistics of first 10 genes  
apply(GeneX[,1:10],2, summary)
```

```
##          CLEC3A  SCGB2A2      CPB1      GSTM1      TFF1  SCGB1D2      KCNJ3  
## Min.      0.000000 0.00000 0.000000 0.000000 0.000000 0.000000 0.000000  
## 1st Qu.    1.000000 7.48779 4.523562 0.000000 6.679463 5.614710 2.807355  
## Median    4.523562 11.70434 7.686501 5.491853 10.796040 9.273796 5.930737  
## Mean      5.925758 11.12788 8.313829 5.357693 9.712977 9.026789 6.687515  
## 3rd Qu.   10.175539 15.06285 11.151939 10.344845 13.207240 12.274950 10.321917  
## Max.      19.455835 22.36921 22.745581 15.874141 18.927850 21.479643 16.672715  
##          MUCL1  LINC00993  ANKRD30A  
## Min.      0.000000 0.000000 0.000000  
## 1st Qu.    5.894767 5.643856 5.554589  
## Median    8.945444 10.512740 10.338736  
## Mean      9.151482 8.864887 8.650306  
## 3rd Qu.   12.404866 12.230770 11.990273  
## Max.      20.547814 15.914922 16.758184
```

```
# clinical variables: gender  
unique(clinical_data$gender) # unique values
```

```
## [1] "female" "male"  NA
```

```
table(clinical_data$gender) # frequency
```

```
##  
## female  male  
##   1217    13
```