## Significance Analysis of Prognostic Signatures

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## 1 Analysis

From MetaGxBreast, we selected patients from studies which tracked overall survival. We identified 4213 patients from 9 datasets.

Number of samples
117
170
1971
238
88
118
1073
198
240
4213

```
## Error in lapply(CompleteLists, function(x) intersect(x, genes.in.pooled.intersect)): error
in evaluating the argument 'X' in selecting a method for function 'lapply': Error: object
'CompleteLists' not found
## Error in eval(expr, envir, enclos): object 'GO.gene.sets.intersect' not found
## Error in lapply(GO.gene.sets.intersect, function(x) gsub(""", "geneid.", : error in evaluating
the argument 'X' in selecting a method for function 'lapply': Error: object 'GO.gene.sets.intersect'
not found
## Error in sapply(GO.gene.sets.intersect, length): error in evaluating the argument 'X' in
selecting a method for function 'sapply': Error: object 'GO.gene.sets.intersect' not found
## Error in do.call(rbind, lapply(GO.gene.sets.intersect, function(x) {: error in evaluating
the argument 'args' in selecting a method for function 'do.call': Error in lapply(GO.gene.sets.intersect
function(x) {:
## error in evaluating the argument 'X' in selecting a method for function 'lapply': Error:
object 'GO.gene.sets.intersect' not found
## Error in nrow(candidateGeneSets): object 'GO.gene.sets.matrix' not found
```