Breast cancer (n = 84)
Cervical cancer (n = 28)
Colorectal cancer I (n = 26)
Colorectal cancer II (n = 101)
Lung cancer II (n = 43)
Prostate cancer I (n = 131)
Gastric cancer (n = 25)
Lung cancer I (n = 20)
Neuroblastoma (n = 47)

Percentage (horizontal axis) of:
- Splicing factor genes with expression levels significantly correlated with the sample-wise TIN-estimates
- Genes in each of 1,000 random gene sets (amount of gene sets on vertical axes) with expression levels significantly correlated with the sample-wise TIN-estimates
- Splicing factor genes with expression levels significantly correlated with each of 1,000 permutations of the TIN-estimates (amount of permuted datasets on vertical axes)