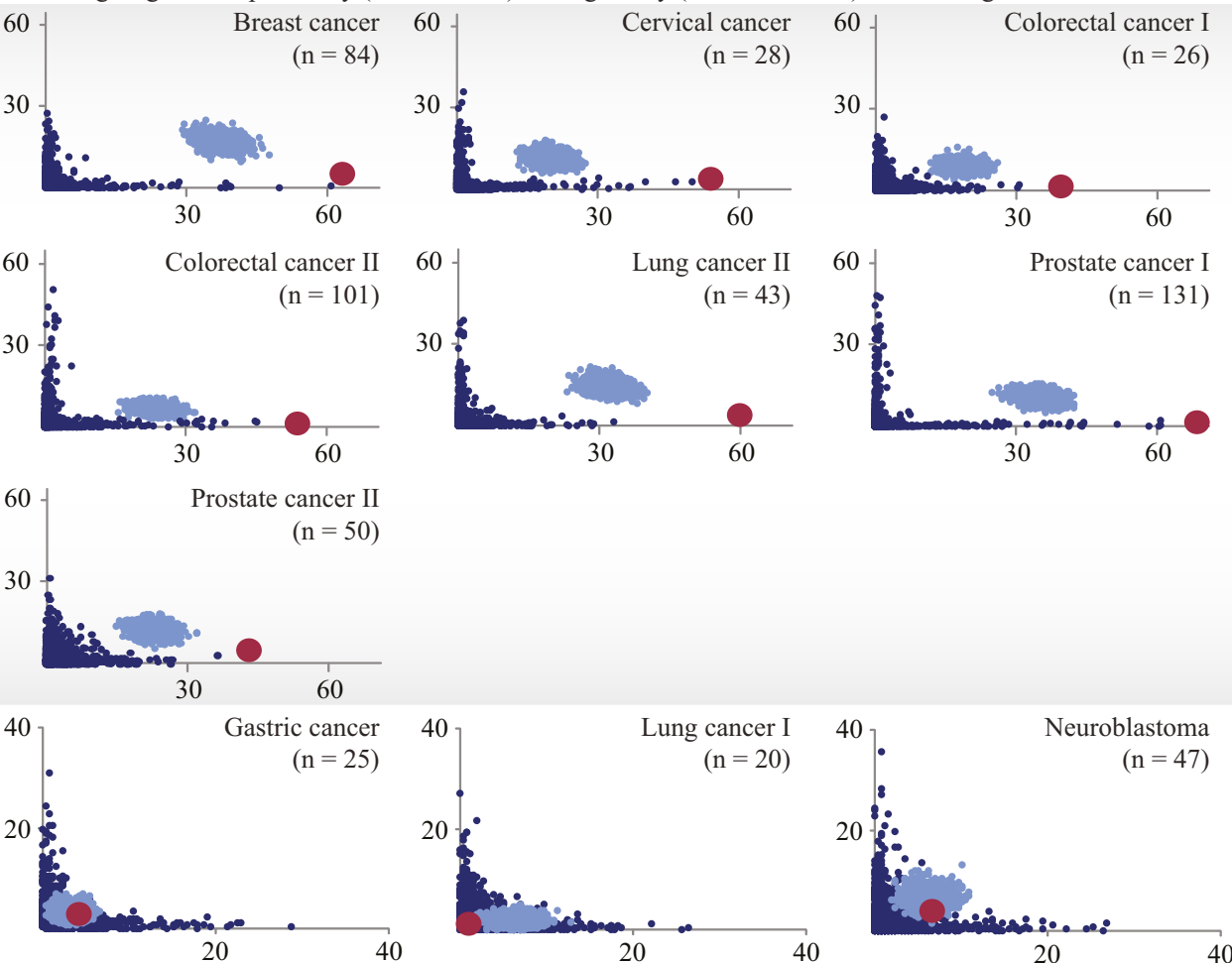


Percentage significant positively (vertical axes) vs. negatively (horizontal axes) correlated genes:



Percentage significantly correlated genes ($P < 0.05$; Pearson correlation) comparing:

- Expression levels of splicing factor genes and sample-wise TIN-estimates
- Expression levels of genes in each of 1,000 random gene sets and sample-wise TIN-estimates
- Expression levels of splicing factor genes and each of 1,000 permutations of the TIN-estimates