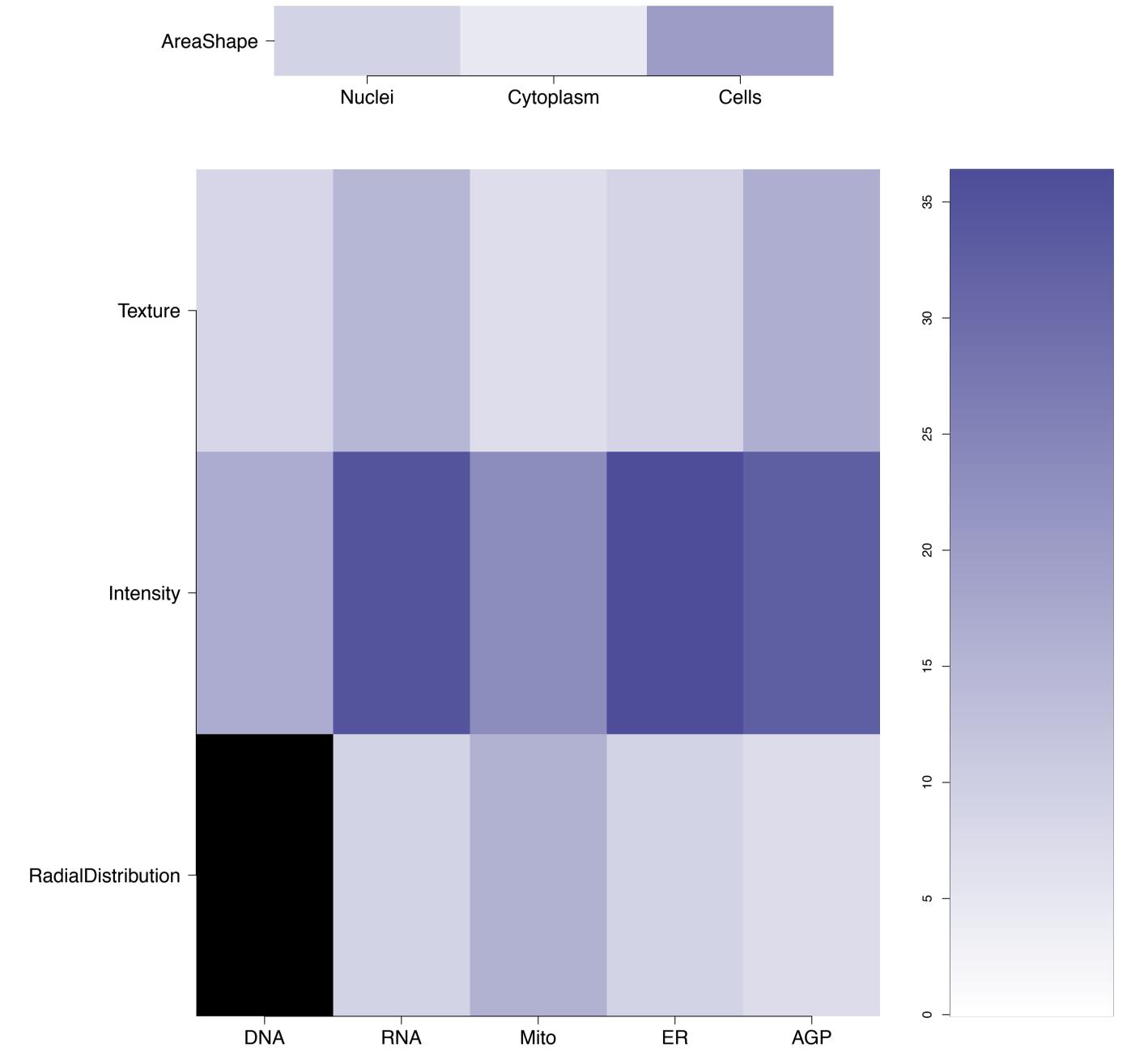


What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a feature z-score across genes in the cluster) Black means no feature is available in the category



ER

RNA

box means the matching

compound is

positively/negatively

correlated with the cluster

Chemical

structure

between

compound

the gene

compound signature

(95th DMSO

replicate correlation

is 0.52)

against the

gene using

L1000

profiling

experiment? (Yellow and red lines correspond to top/bottom

1st and 5th percentile DMSO correlation to all the genes)

Which individual morphological features are distinguishing in the gene relative to the untreated samples? Blue/Red means the feature has a positive/negative z-score. Size is proportional to the z-score value.

untreated samples. Black means a mismatch; i.e. active (= high

z-score in magnitude) in the compound, and either inactive (=

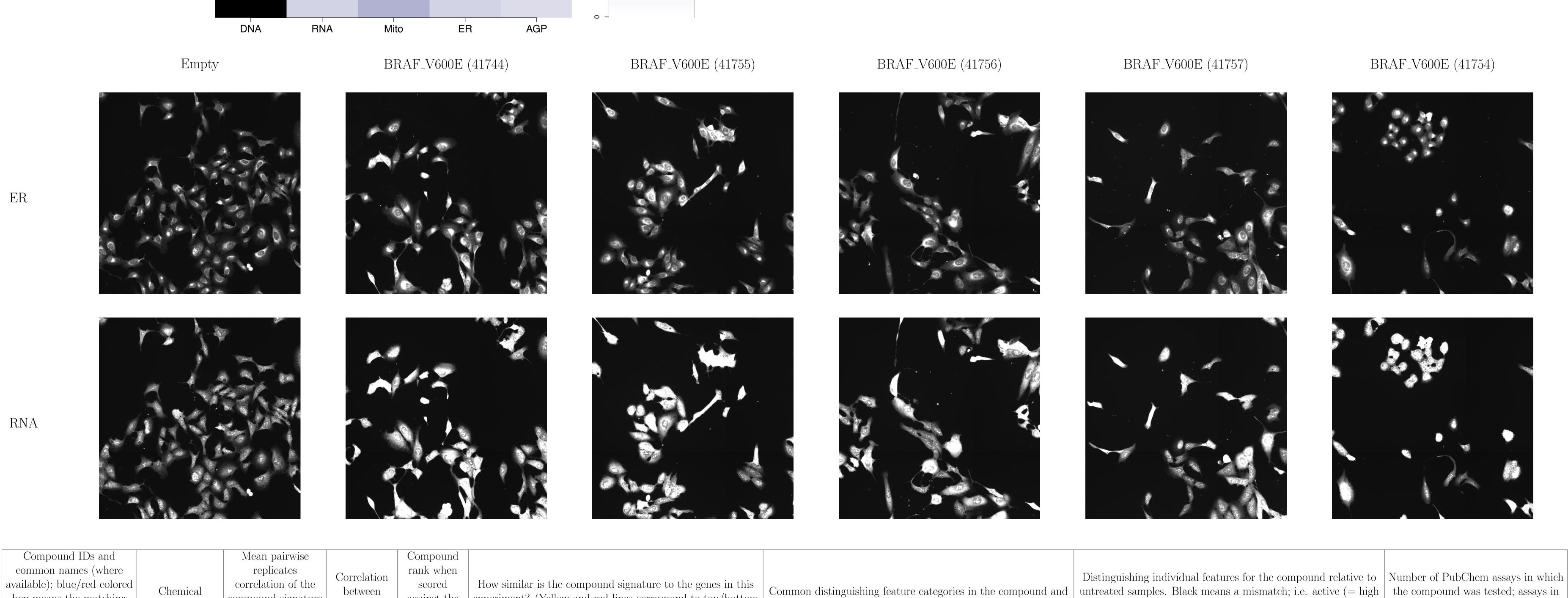
small z-score in magnitude) or oppositely active in the gene

the compound was tested; assays in

which the compound was active are

itemized





the gene relative to the untreated samples

