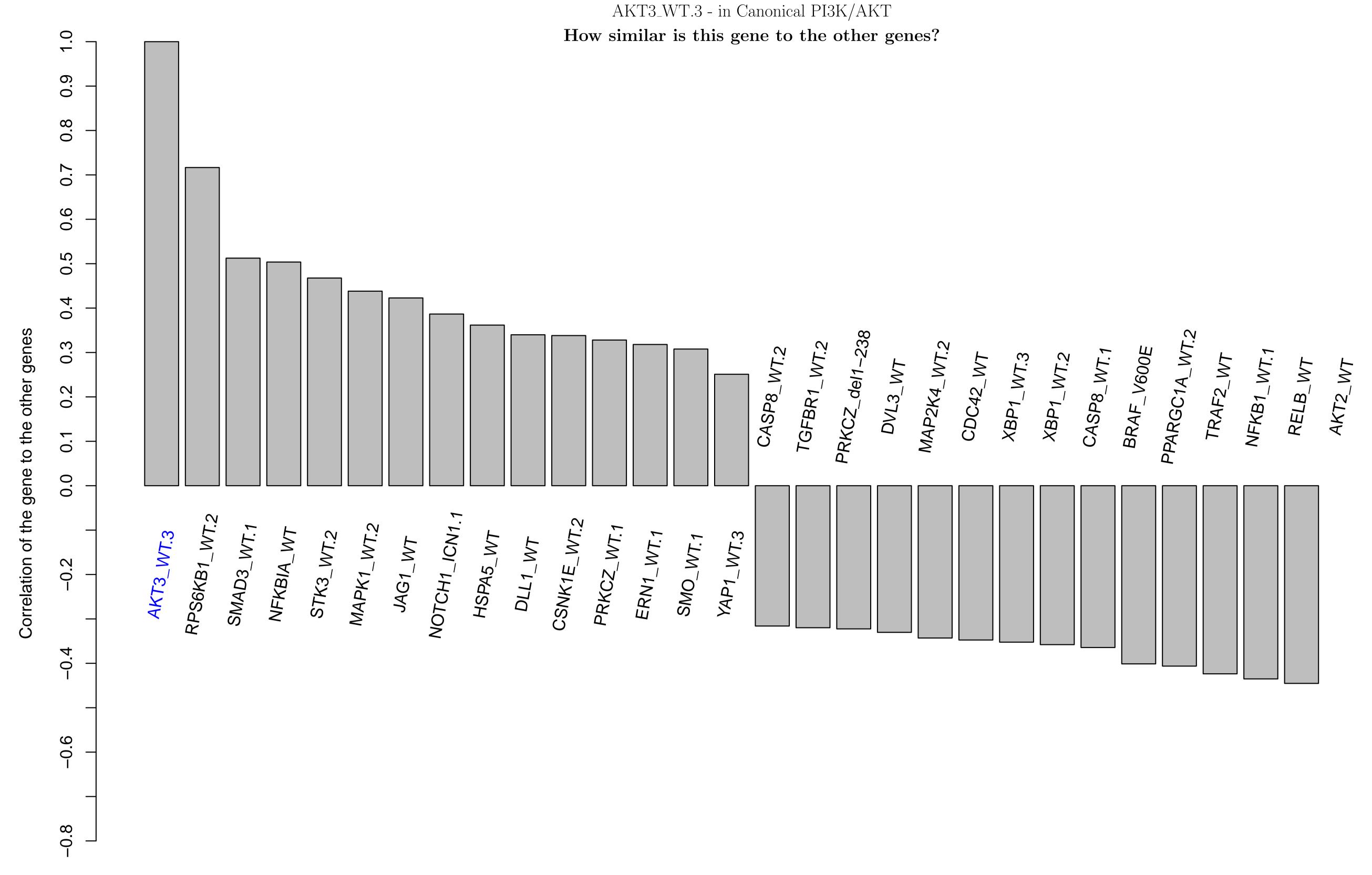
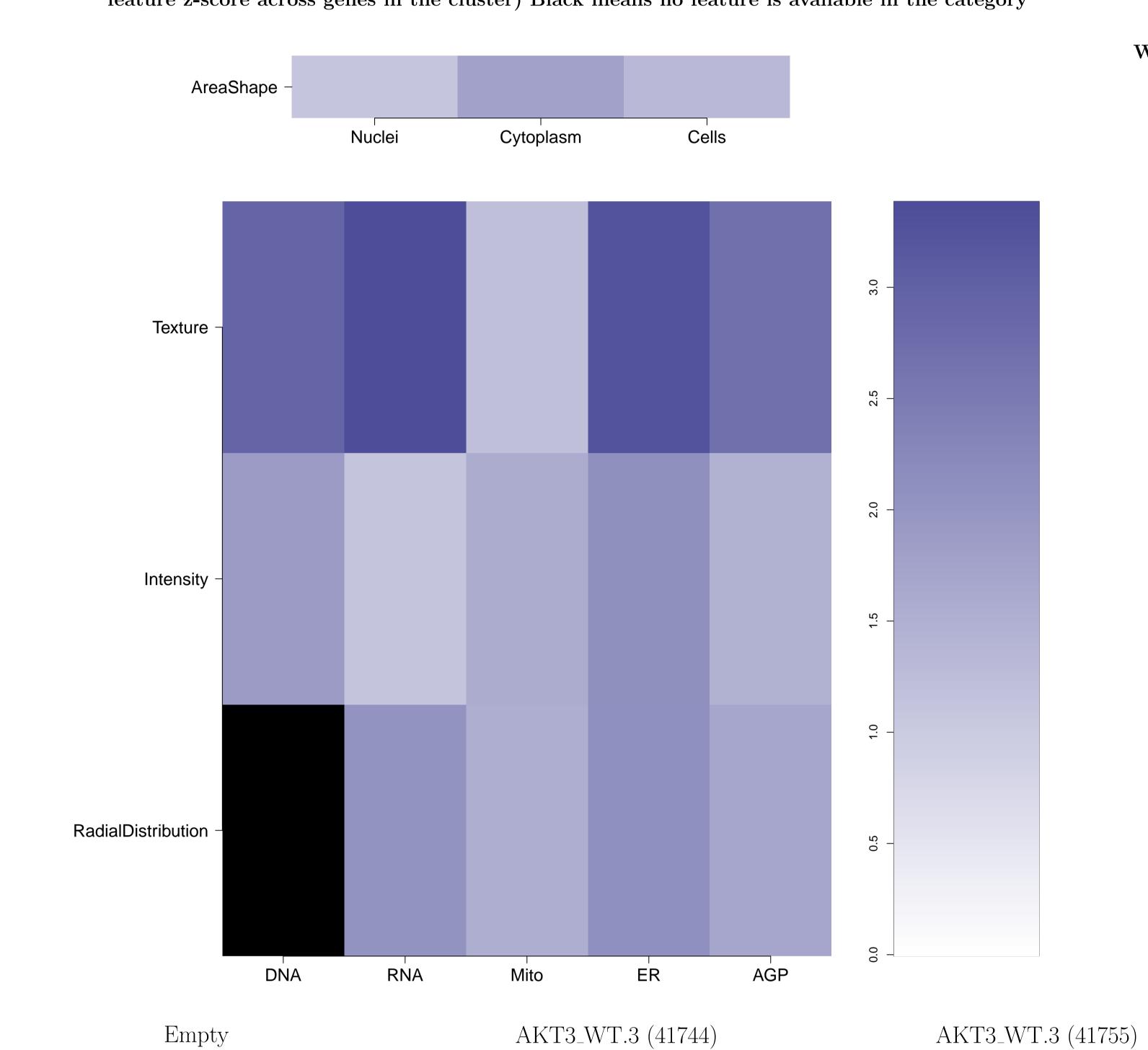
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein



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

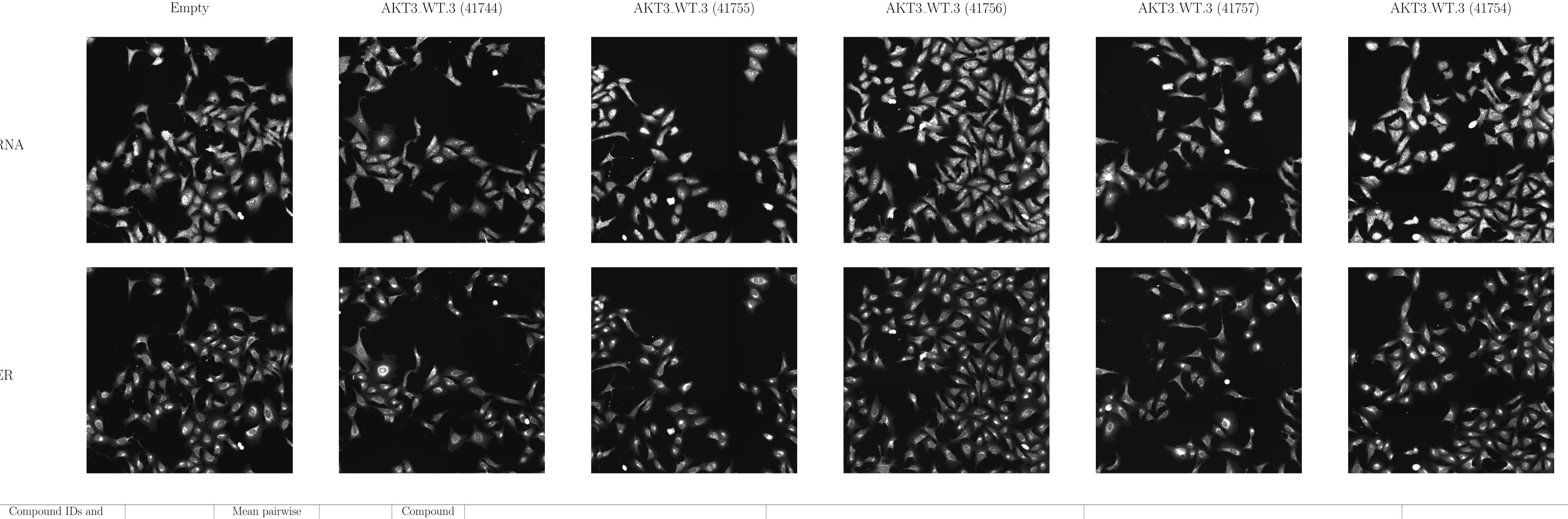


RNA

ER

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.





| common names (where available); blue/red colored box means the matching compound is positively/negatively correlated with the cluster | Chemical structure | replicates correlation of the compound signature (95th DMSO replicate correlation is 0.51) | Correlation between compound the gene |  | How similar is the compound signature to the genes in this experiment? (Yellow and red lines correspond to top/bottom 1st and 5th percentile DMSO correlation to all the genes) | Common distinguishing teature categories in the compound and | Distinguishing individual features for the compound relative to 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 |
|---|--------------------|--|---------------------------------------|--|---|--|---|------------------------------------|
|---|--------------------|--|---------------------------------------|--|---|--|---|------------------------------------|

