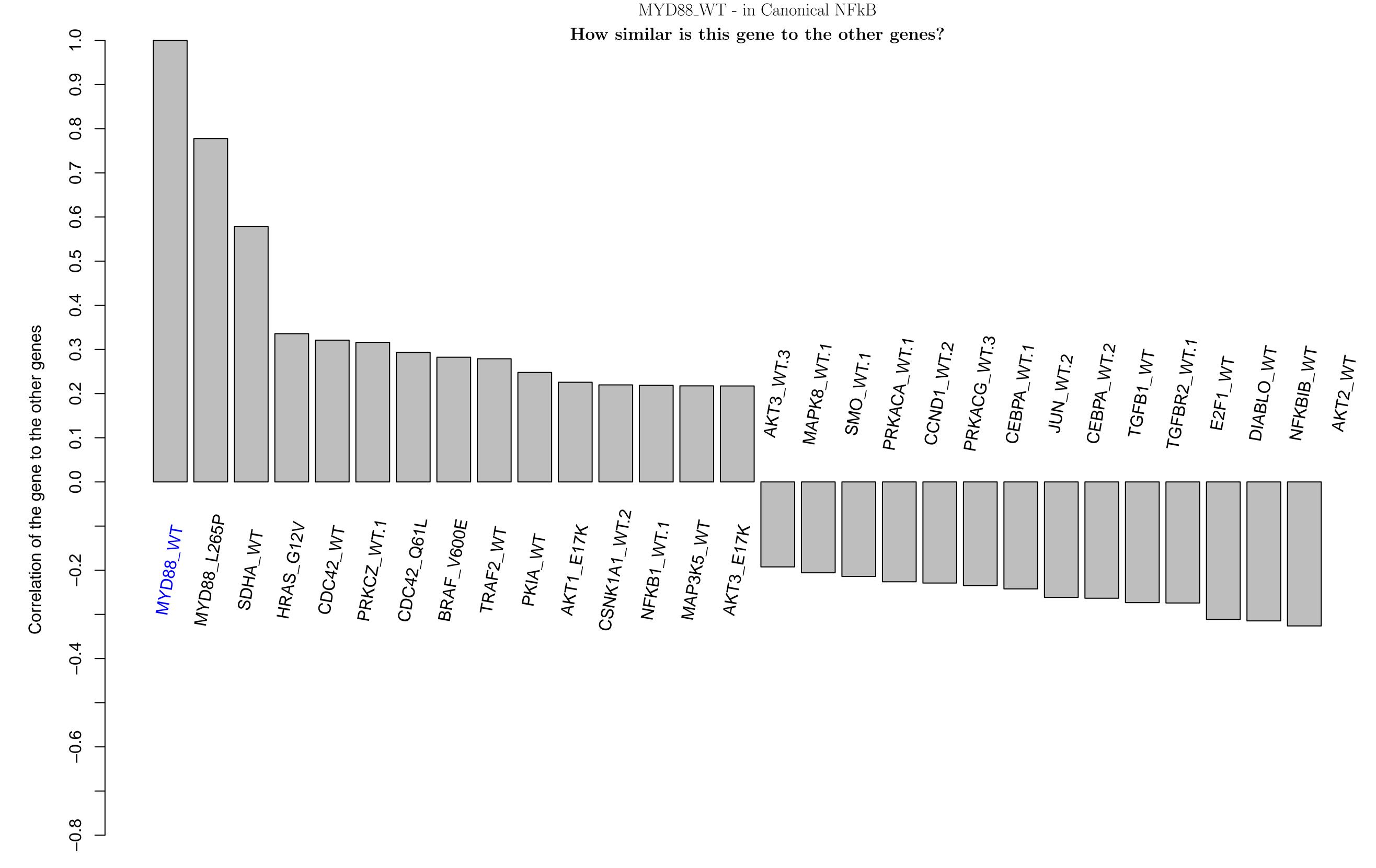
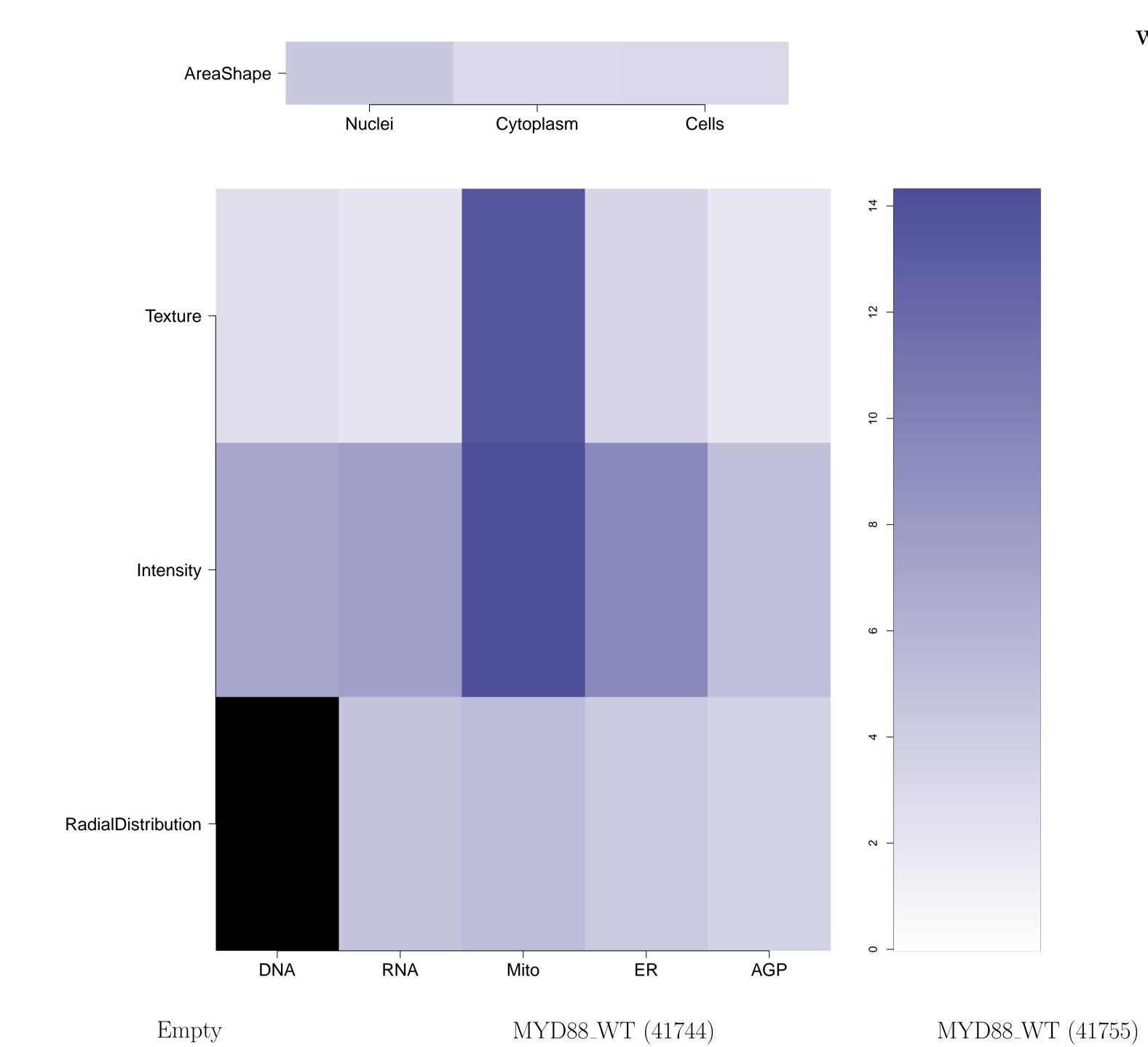
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein



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

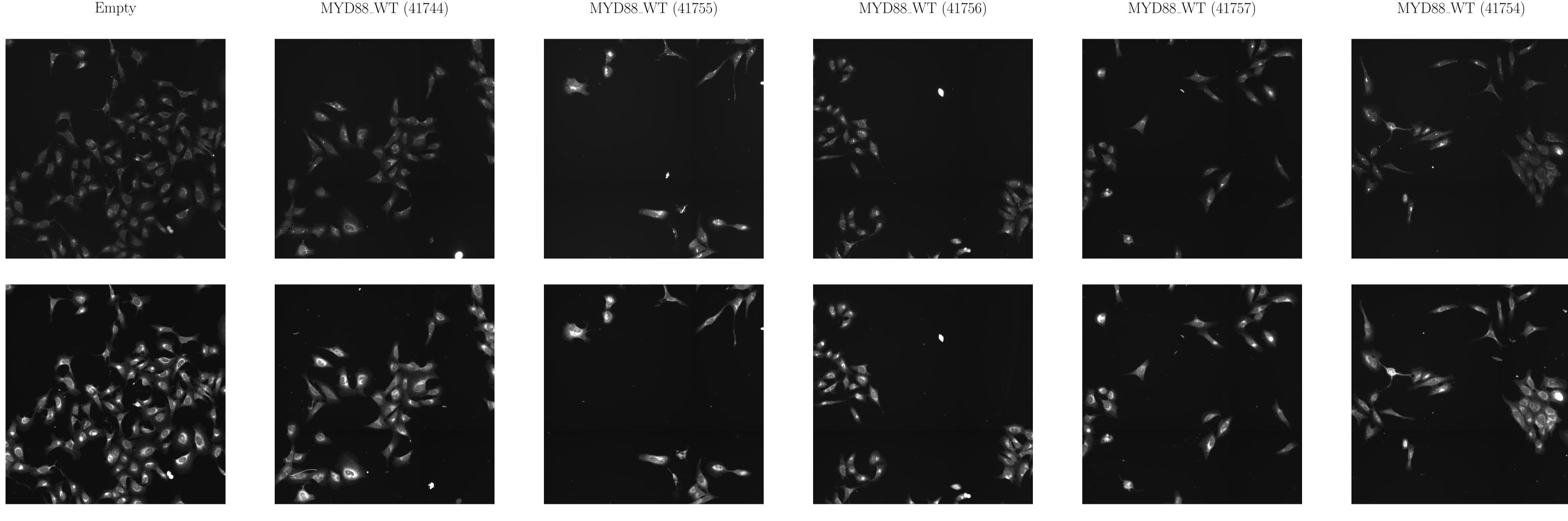


Mito

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.





Compound IDs and Mean pairwise Compound rank when common names (where replicates Correlation Distinguishing individual features for the compound relative to Number of PubChem assays in which How similar is the compound signature to the genes in this correlation of the available); blue/red colored scored Chemical Common distinguishing feature categories in the compound and untreated samples. Black means a mismatch; i.e. active (= high between the compound was tested; assays in experiment? (Yellow and red lines correspond to top/bottom against the box means the matching compound signature the gene relative to the untreated samples which the compound was active are compound z-score in magnitude) in the compound, and either inactive (= structure compound is (95th DMSO 1st and 5th percentile DMSO correlation to all the genes) gene using small z-score in magnitude) or oppositely active in the gene the gene itemized replicate correlation positively/negatively L1000 correlated with the cluster is 0.52) profiling

