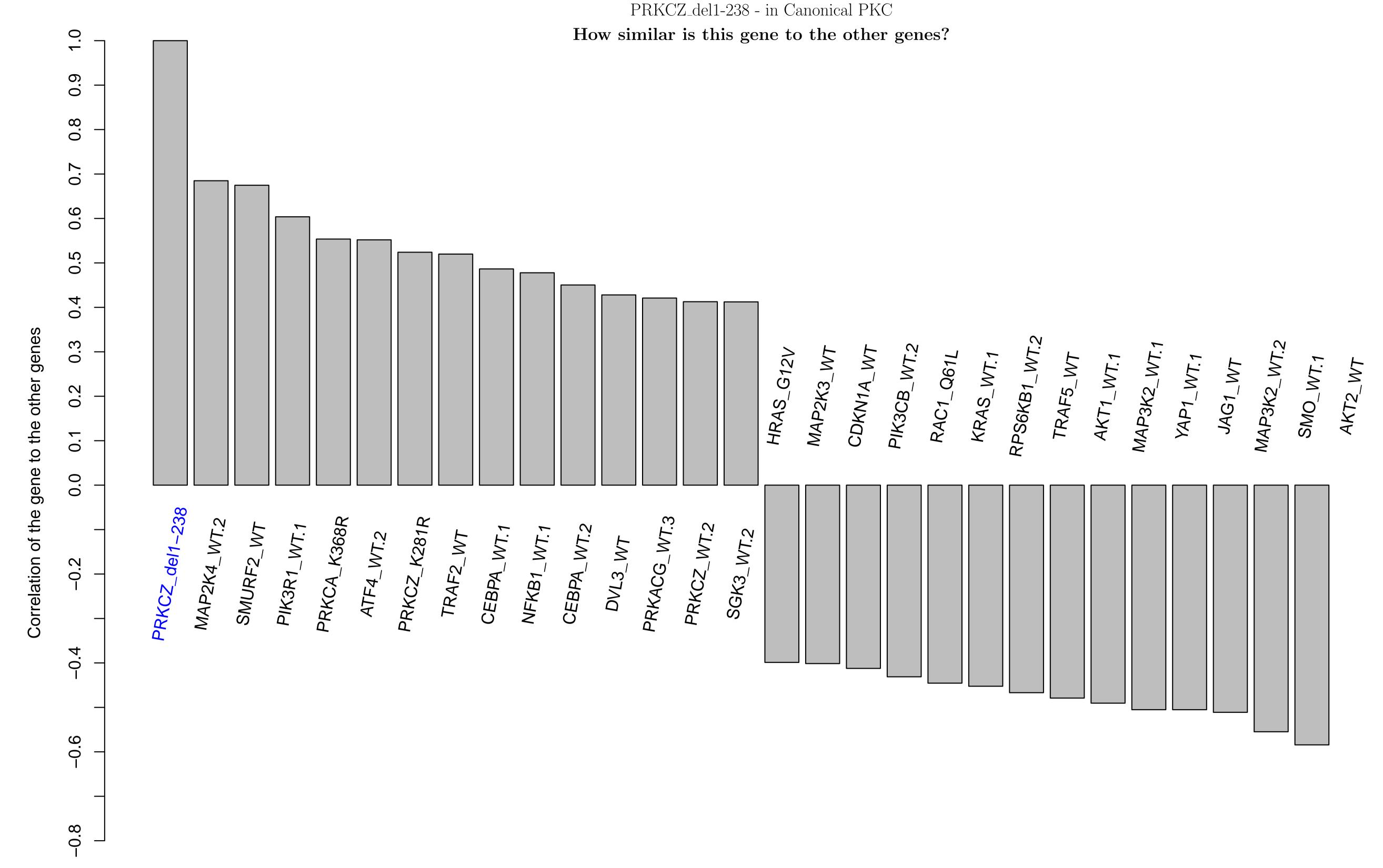
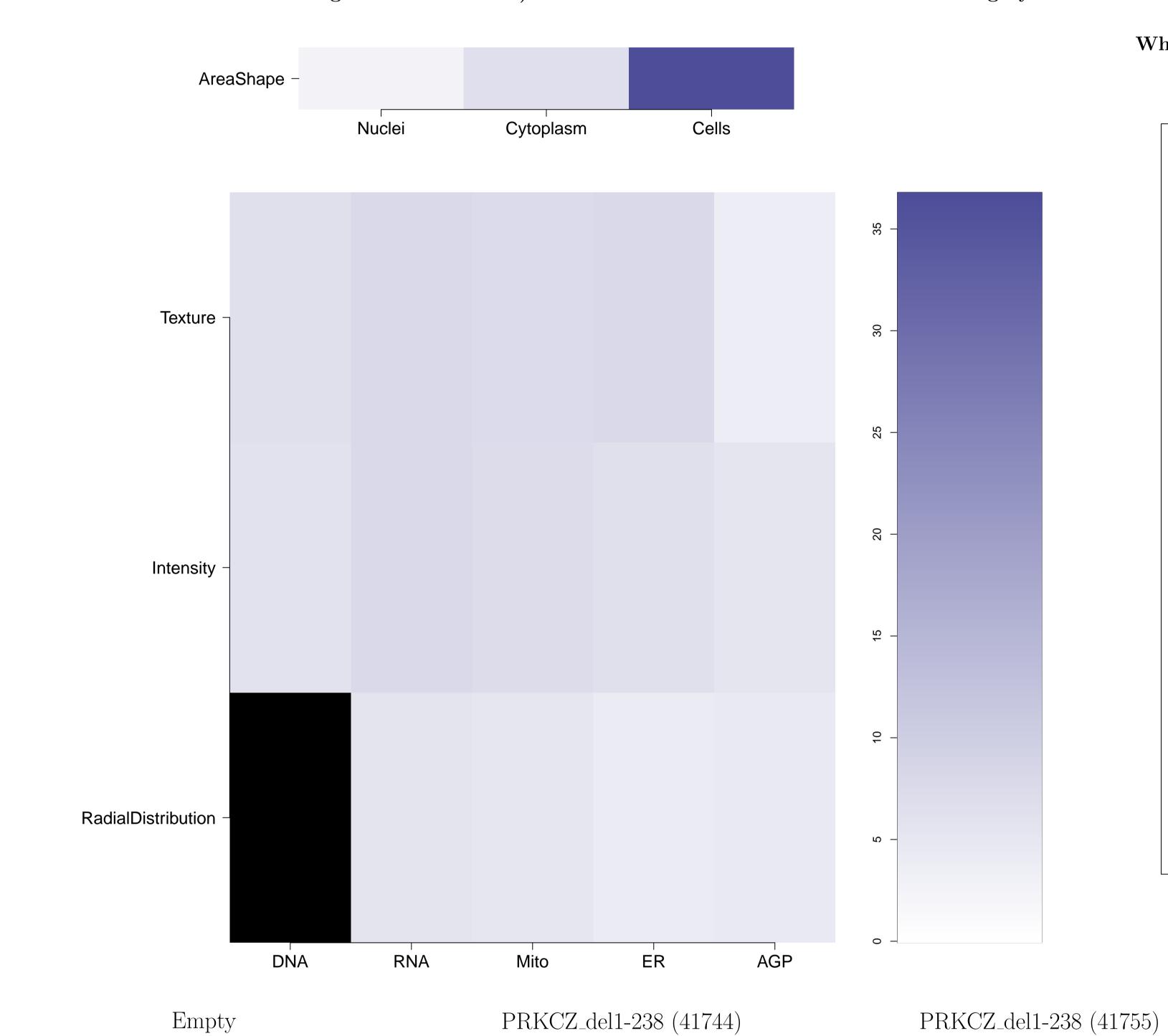
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein



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



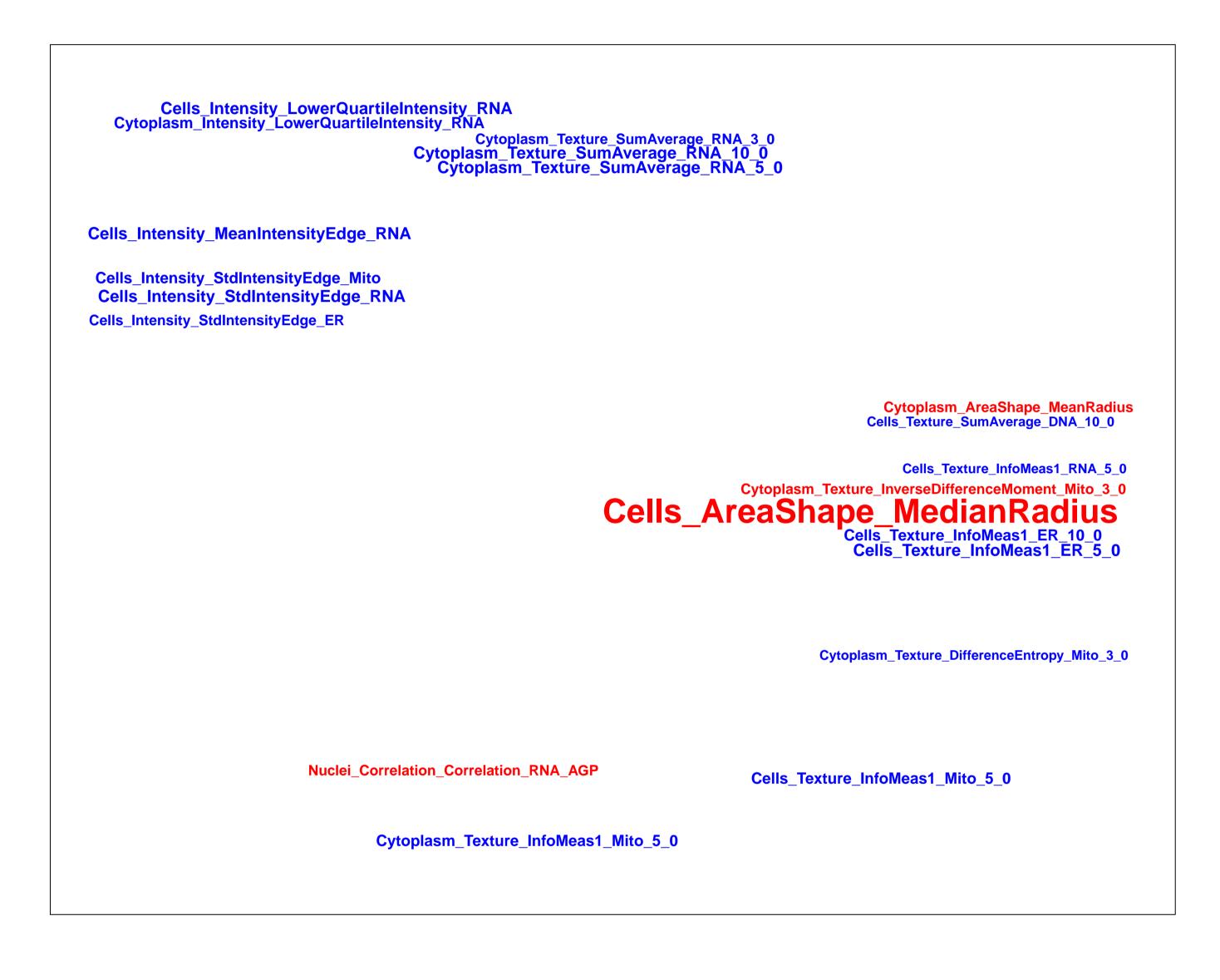
RNA

ER

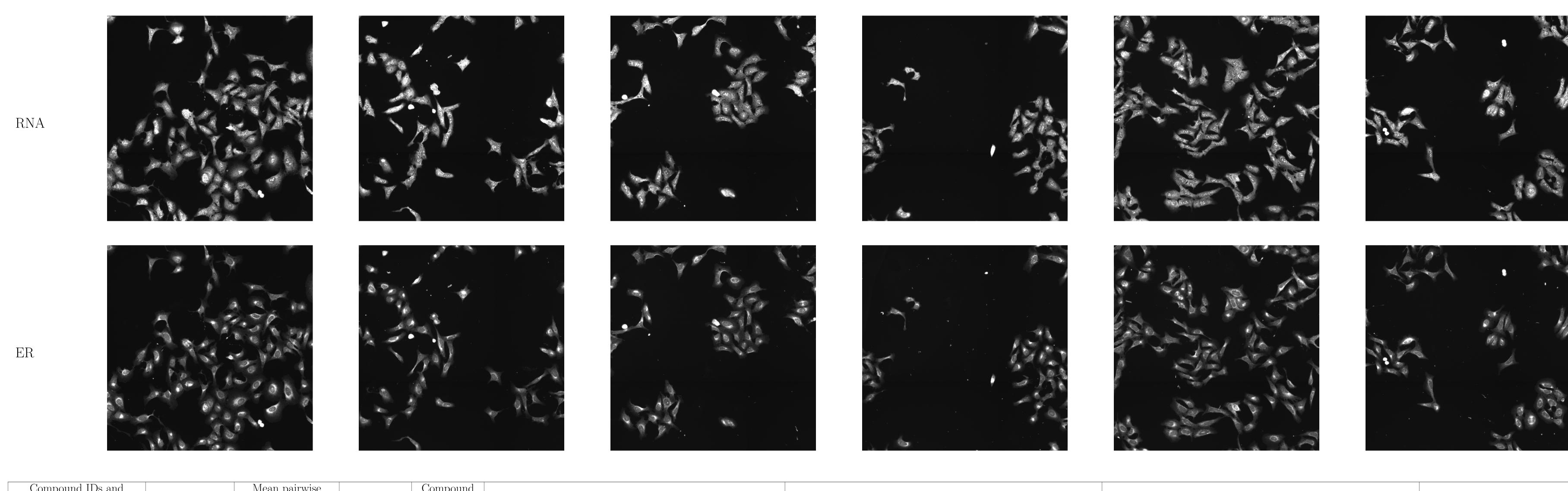
Which individual morphological features are distinguishing in the gene relative to the untreated samples? Blue/Red means the feature has a positive/negative z-score. Size is proportional to the z-score value.

PRKCZ_del1-238 (41757)

PRKCZ_del1-238 (41754)



PRKCZ_del1-238 (41756)



Compound IDs and		Mean pairwise		Compound			
common names (where		replicates	Correlation	rank when		Distinguishing individual features for the compound relative to	Number of PubChem aggreg in which
available); blue/red colored	Chemical structure	correlation of the	the ature compound	scored	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)	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
box means the matching		compound signature		against the			
compound is		(95th DMSO		gene using			
positively/negatively		replicate correlation	the gene	L1000		sman z-score in magnitude) of oppositely active in the gene	ıtemızed
correlated with the cluster		is 0.51)		profiling			

