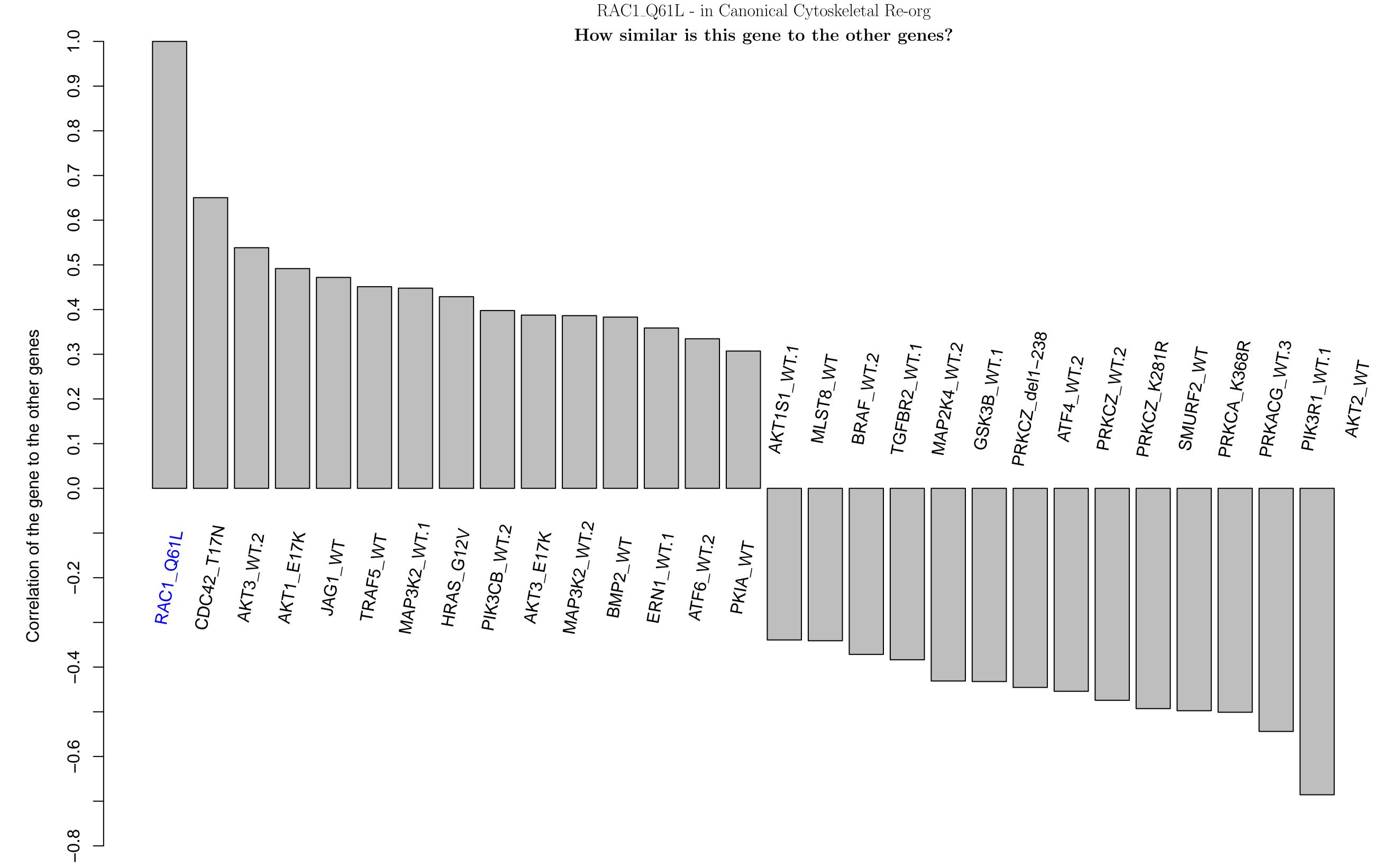
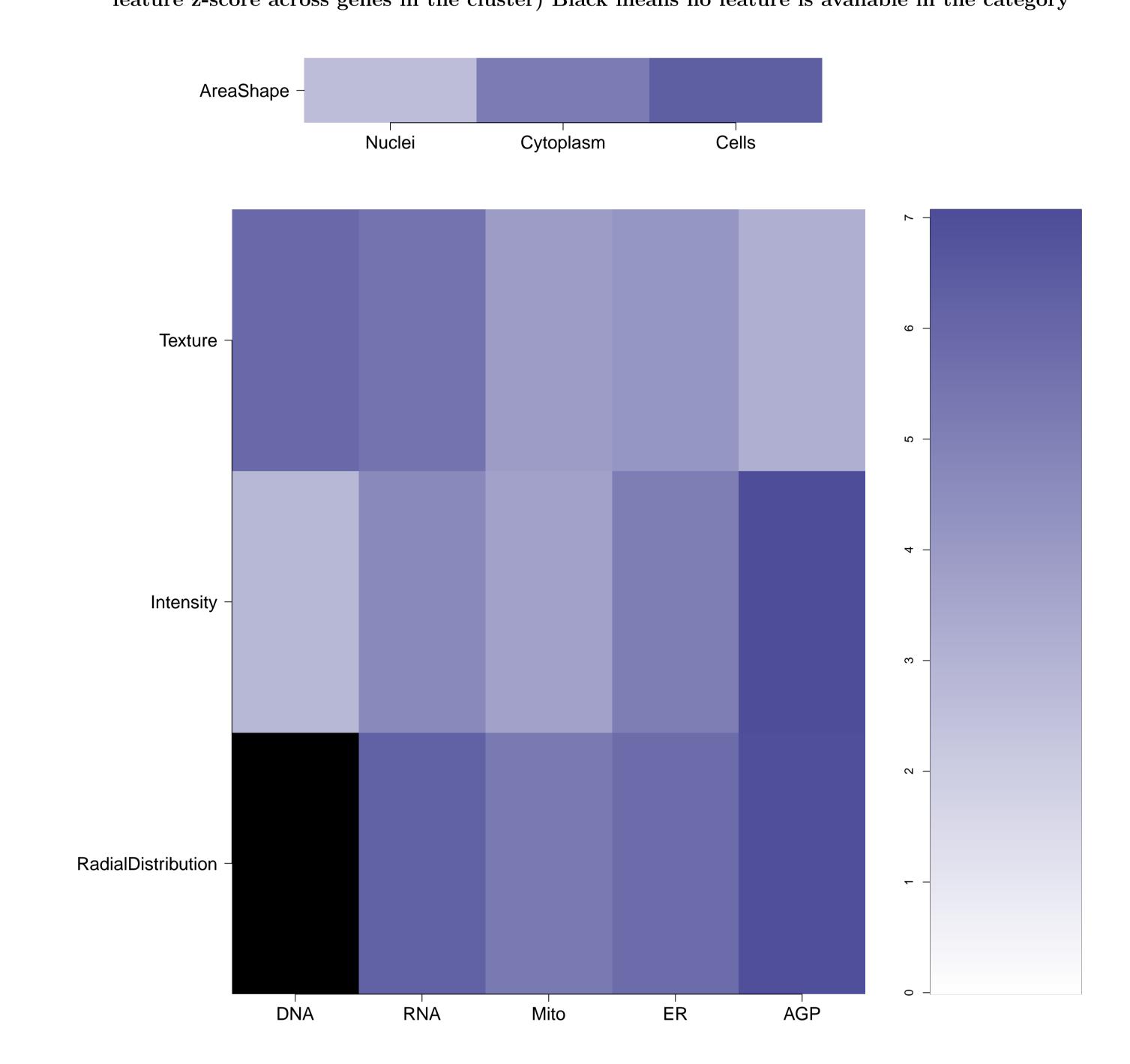
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein



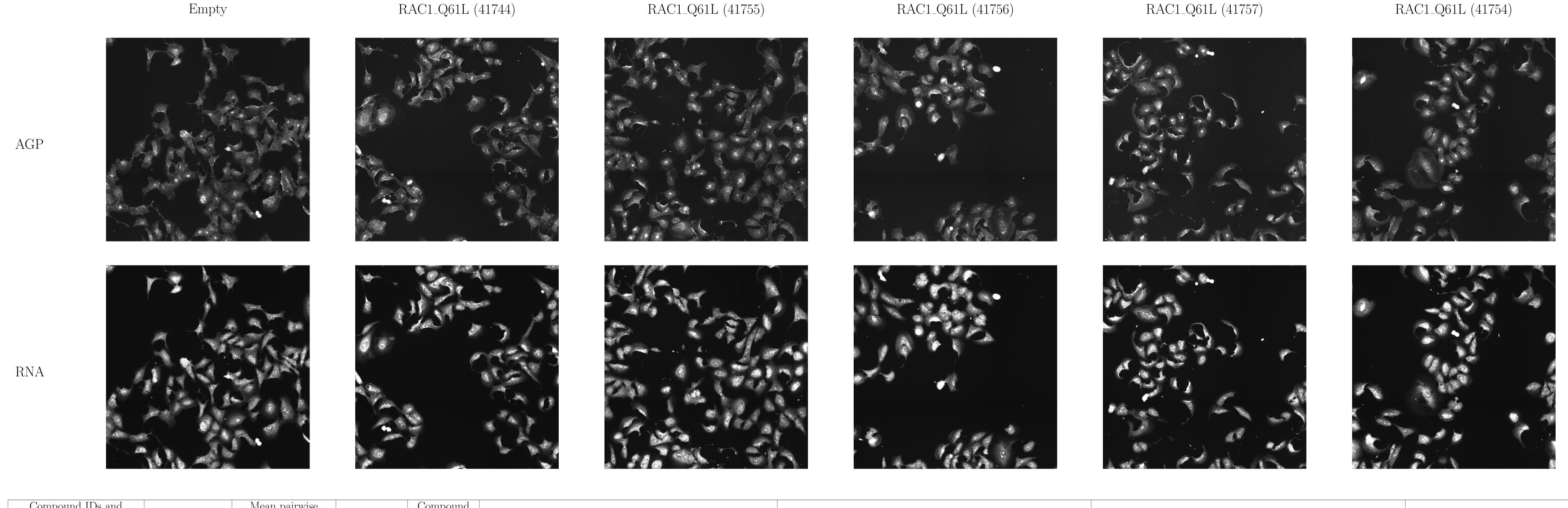
What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a feature z-score across genes in the cluster) Black means no feature is available in the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



RNA

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				
common names (where available); blue/red colored	Chemical structure	replicates correlation of the	of the gnature compound scored against the	scored	How similar is the compound signature to the genes in this	Lommon 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	
box means the matching compound is		compound signature (95th DMSO		experiment? (Yellow and red lines correspond to top/bottom 1st and 5th percentile DMSO correlation to all the genes)		z-score in magnitude) in the compound, and either inactive (= small z-score in magnitude) or oppositely active in the gene	, , ,	
positively/negatively correlated with the cluster		replicate correlation is 0.52)	one gene	L1000 profiling			sman z-score in magnitude, or oppositely active in the gene	Toermzea

