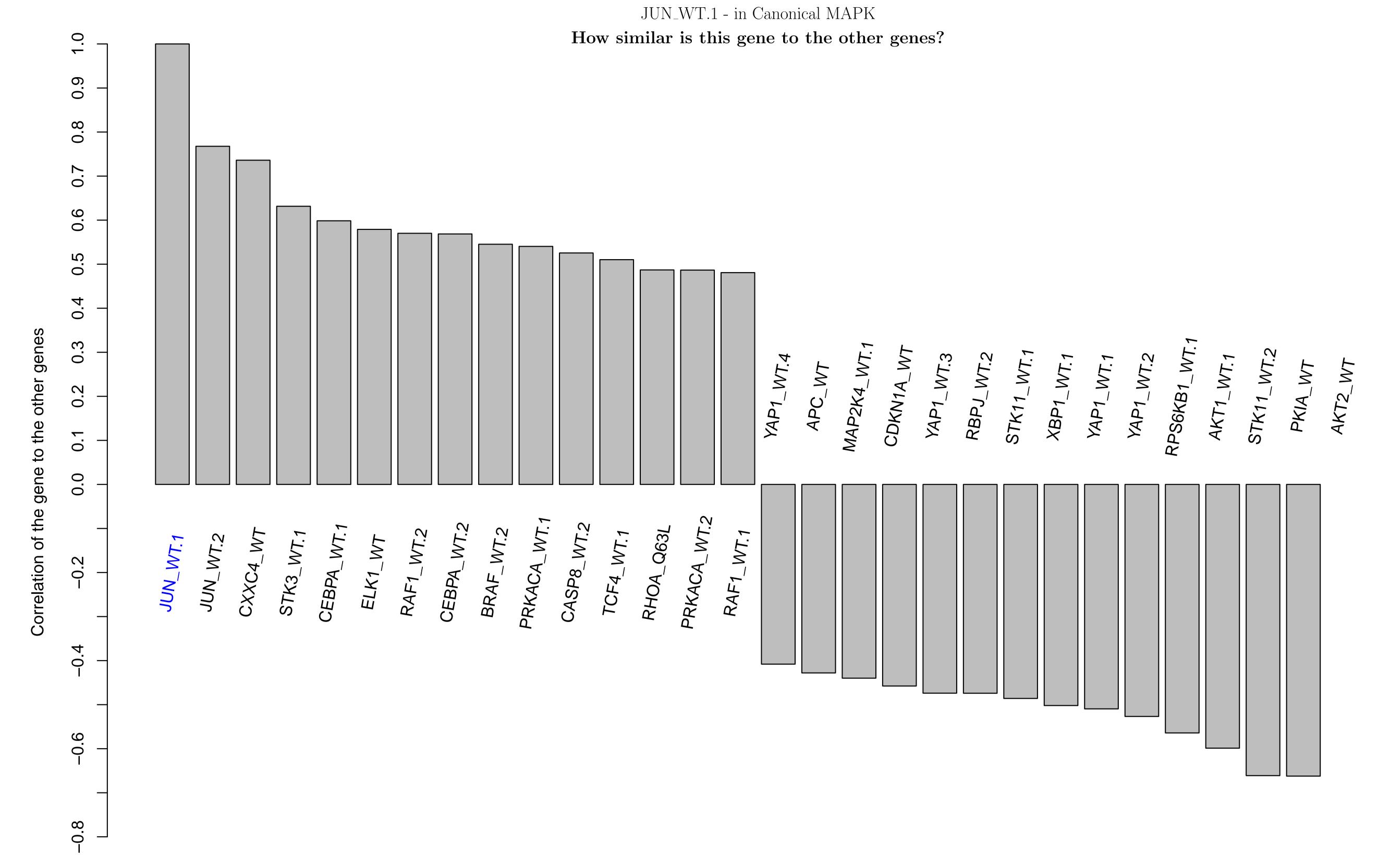
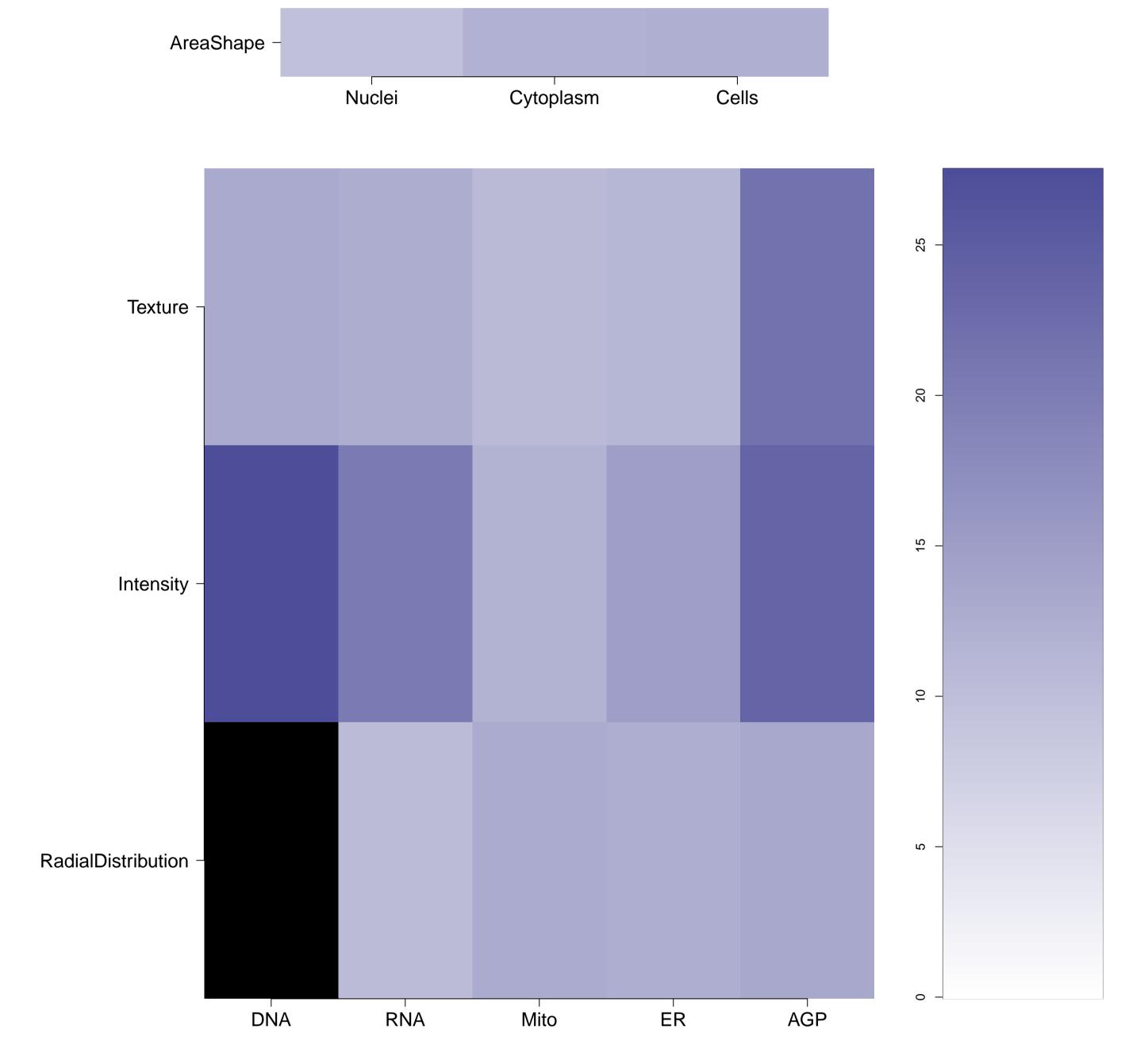
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein

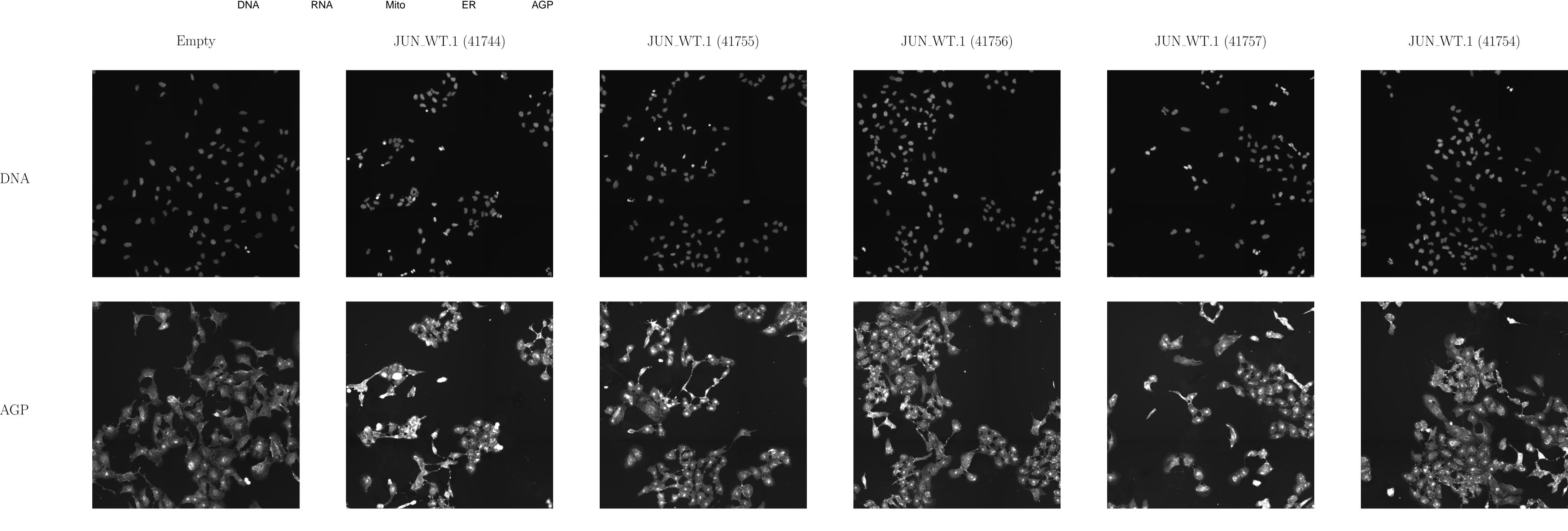


What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a feature z-score across genes in the cluster) Black means no feature is available in the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



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	d IDs and		Mean pairwise		Compound				
	mes (where	Chemical structure	replicates	the Correlation between	rank when	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 feature 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 (= which the compound was test and z-score in magnitude) or oppositely active in the gene	Number of PubChem assays in which
/ /	ie/red colored		correlation of the						the compound was tested; assays in
	he matching		compound signature	1	compound against the gene using				
compo	ound is		(95th DMSO	the gene					
positively/	'negatively		replicate correlation	elation L1000			sman z-score in magnitude, or oppositely active in the gene	Tocimized	
correlated wit	th the cluster		is 0.52)		profiling				

