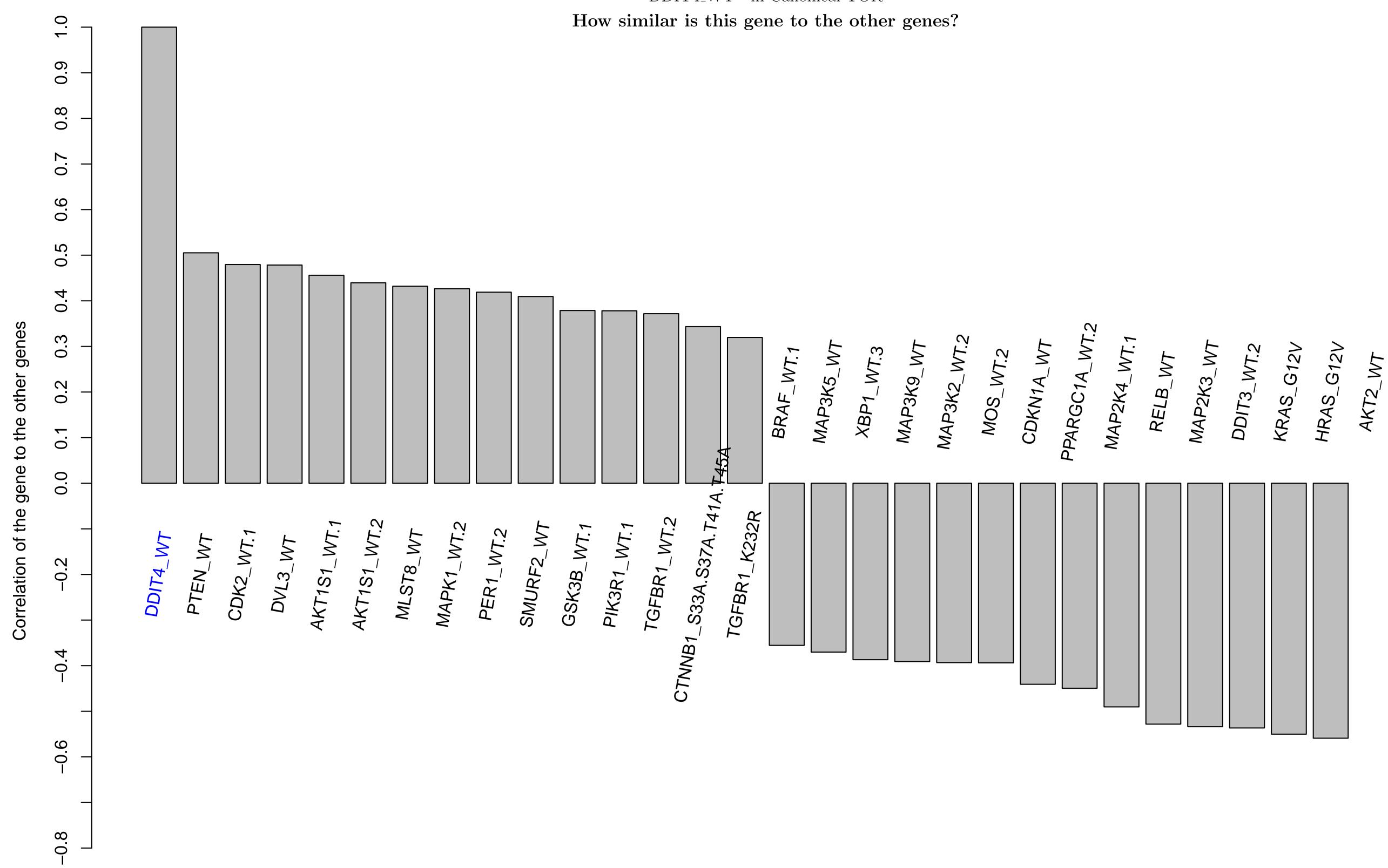
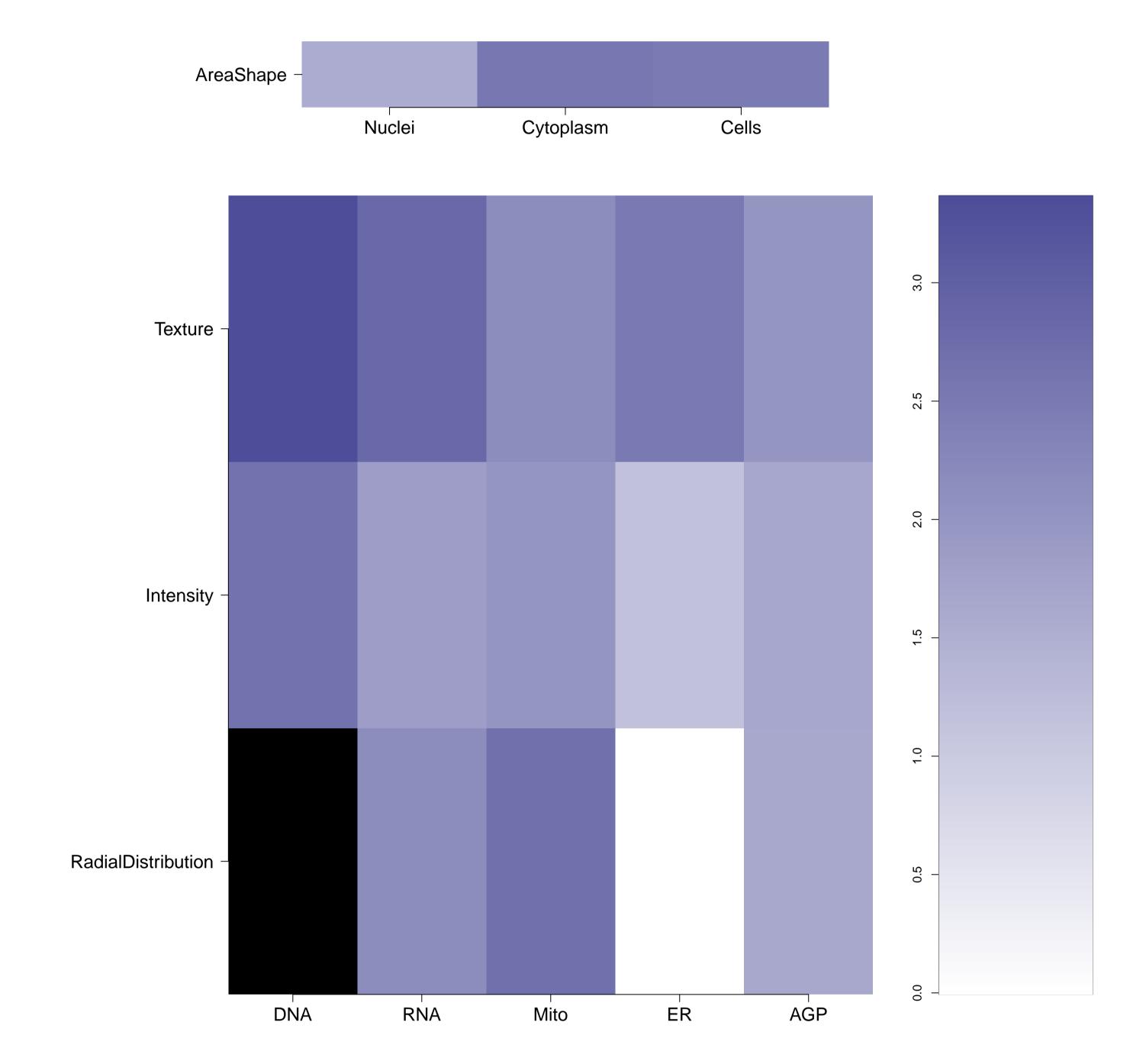
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein

DDIT4_WT - in Canonical TOR

How similar is this gene to the other genes?

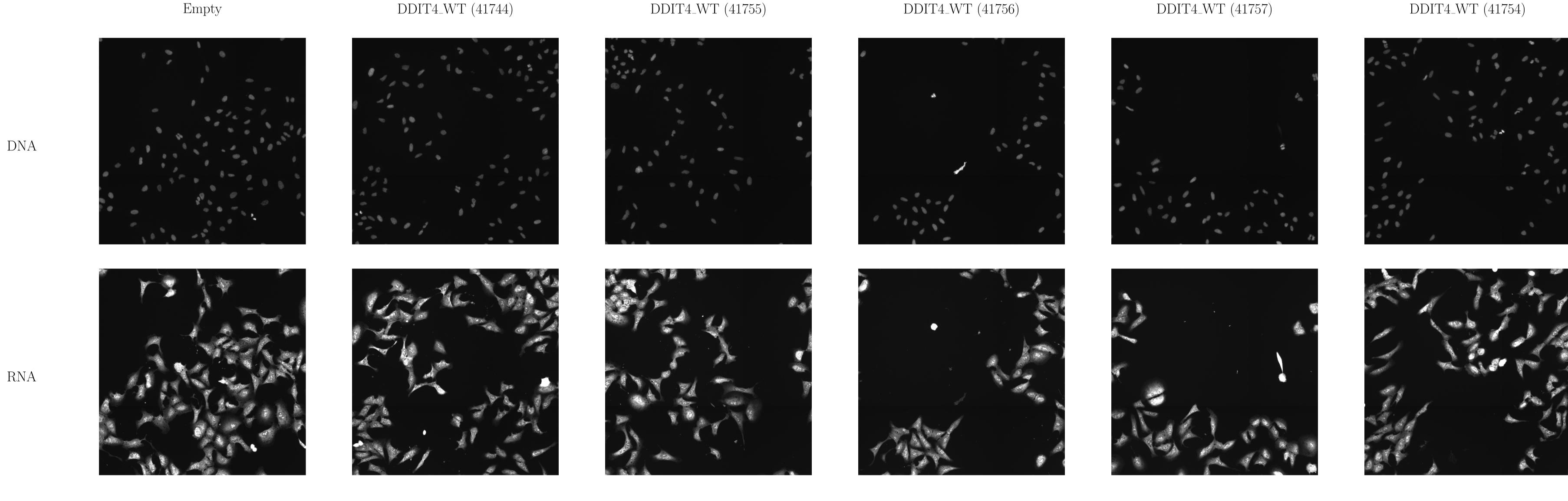


What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a feature z-score across genes in the cluster) Black means no feature is available in the 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 IDs and		Mean pairwise		Compound					
common names (where	Chemical structure	replicates	Correlation	rank when	How similar is the compound signature to the genes in this	Common distinguishing feature categories in the compound and	Distinguishing individual features for the compound relative to Number of PubChem assays in which		
available); blue/red colored		correlation of the		scored					
box means the matching		compound signature	between	against the			z-score in magnitude) in the compound, and either inactive (= mgn which the compound was tested, assays in small z-score in magnitude) or oppositely active in the gene itemized		
compound is		(95th DMSO	compound gene usin	gene using				-	
positively/negatively		replicate correlation	the gene	L1000					
correlated with the cluster		is 0.51)		profiling					

