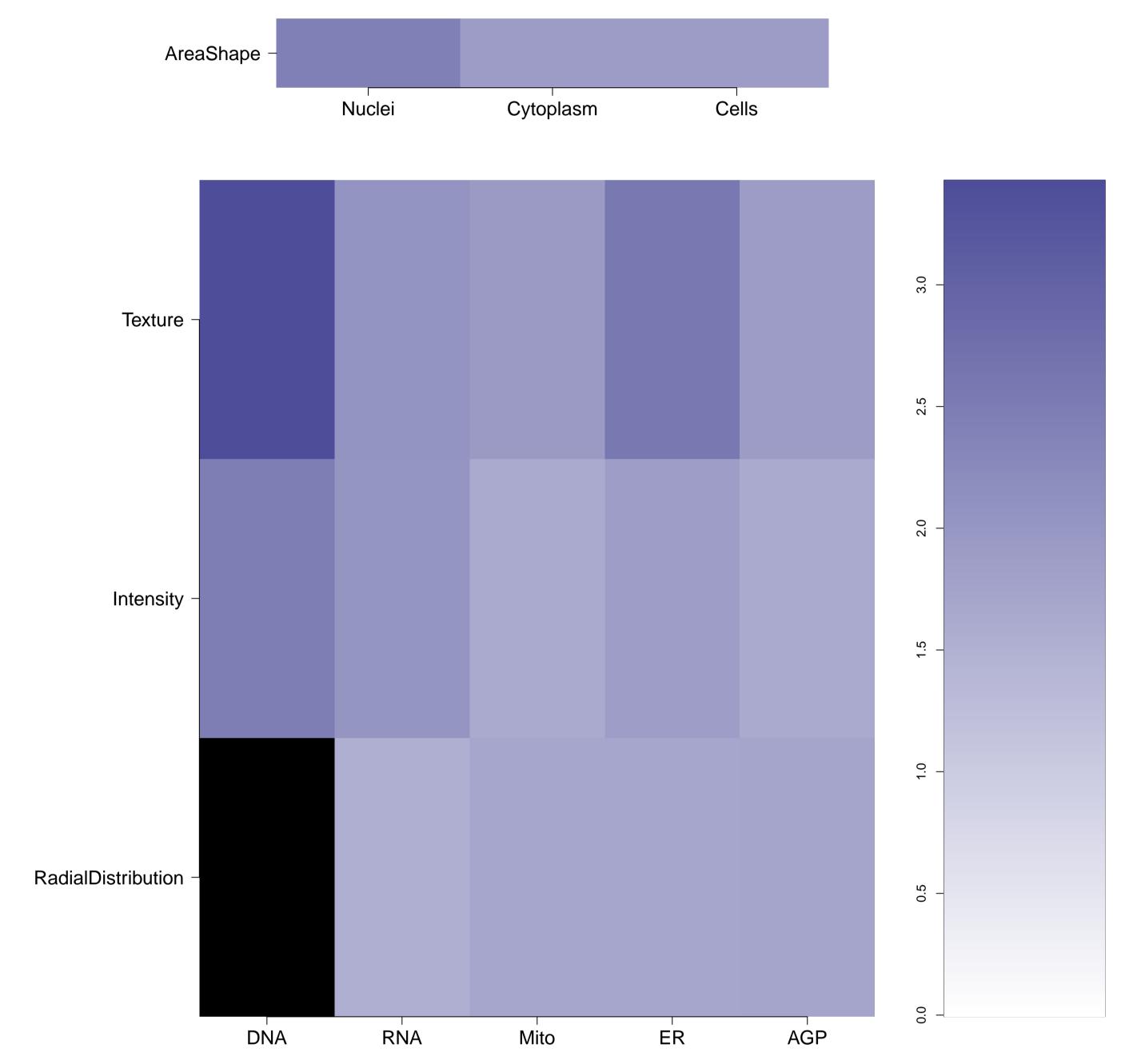
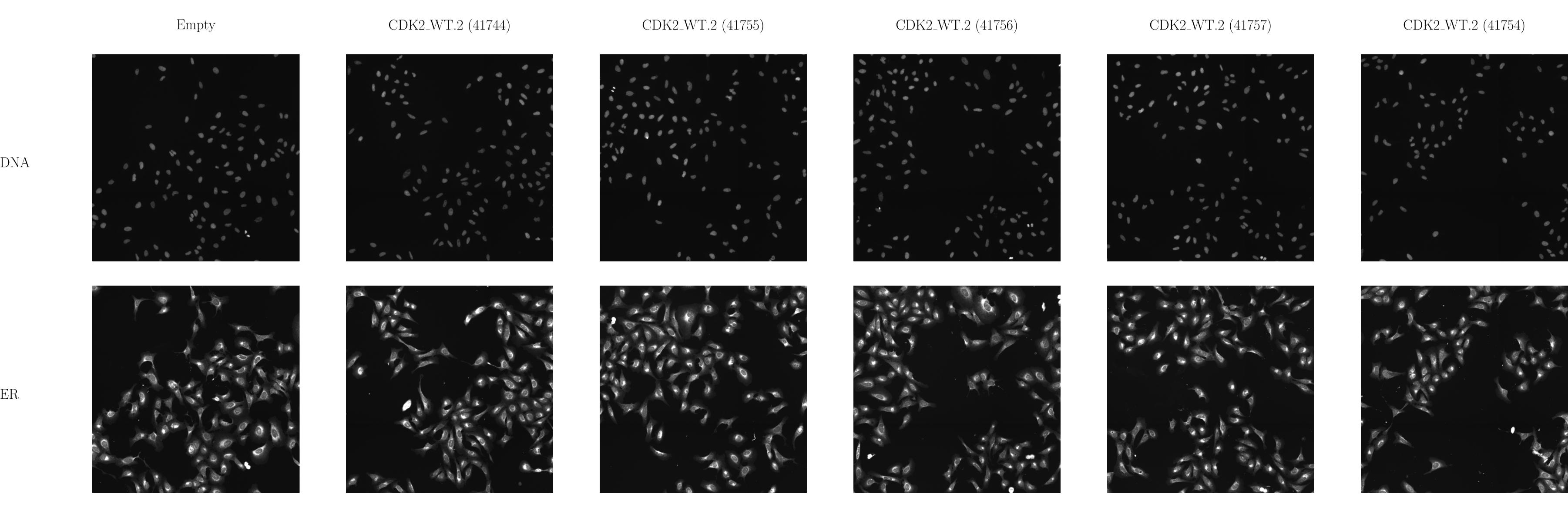


What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a feature z-score across genes in the cluster) Black means no feature is available in the 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	c between against the compound the gene using			Distinguishing individual features for the compound relative to Number of PubChem assays in	Number of PubChem aggree in which	
available); blue/red colored		correlation of the		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)	Common distinguishing feature categories in the compound and	d untreated samples. Black means a mismatch; i.e. active (= high   the com	
box means the matching		compound signature		against the				7
compound is		(95th DMSO		gene using				itemized
positively/negatively		replicate correlation				sman z-score in magnitude) of oppositely active in the gene	Itemized	
correlated with the cluster		is 0.52)		profiling				

