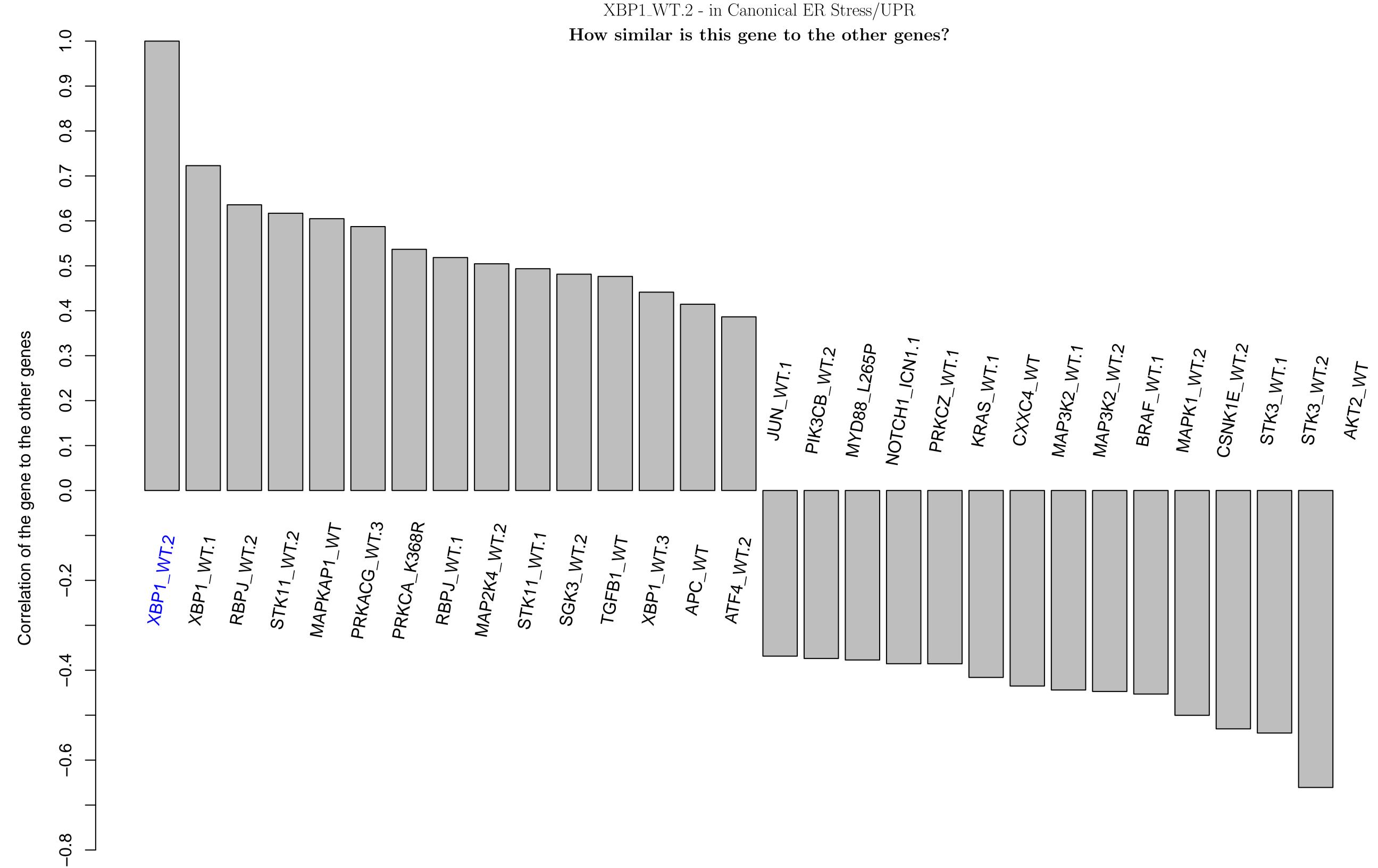
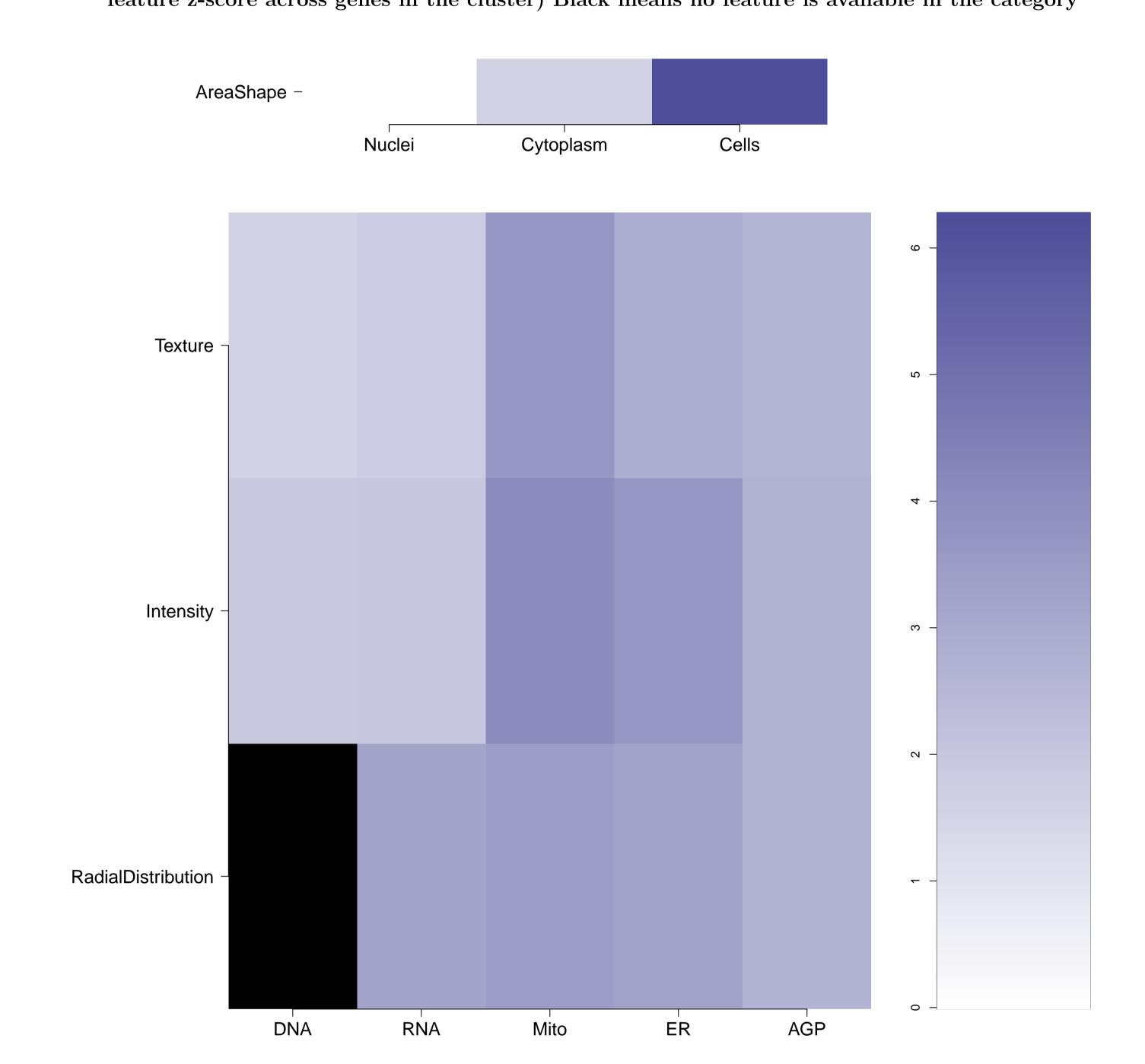
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein



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



XBP1_WT.2 (41744)

Empty

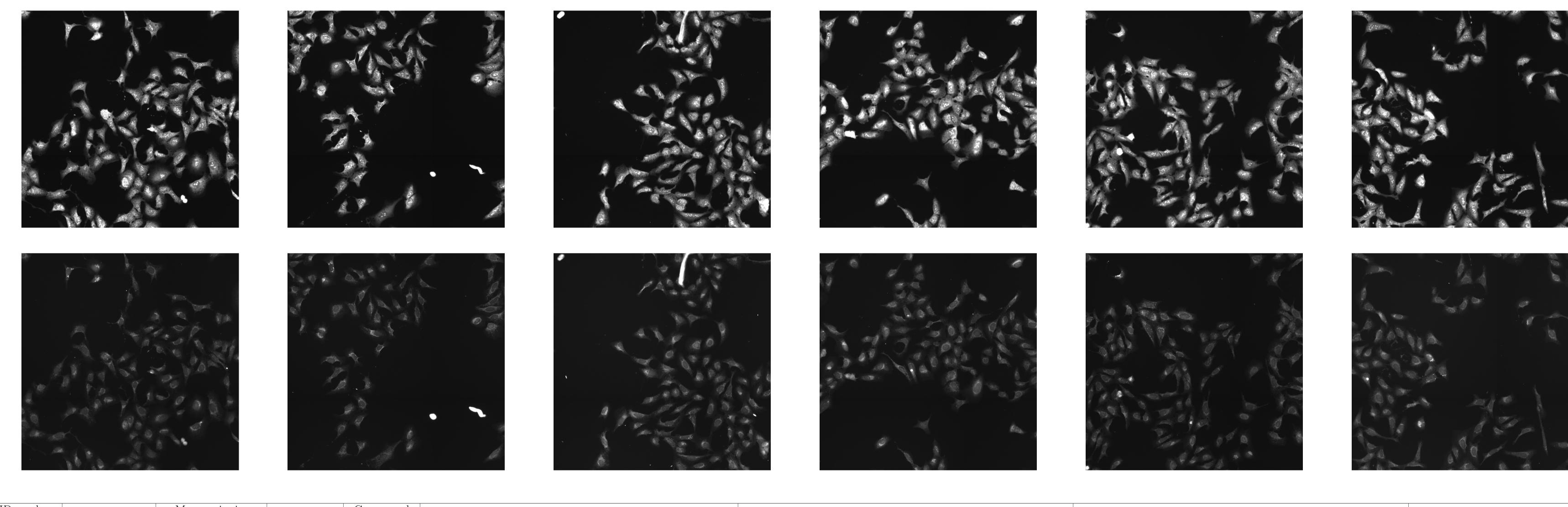
Which individual morphological features are distinguishing in the gene relative to the untreated samples? Blue/Red means the feature has a positive/negative z-score. Size is proportional to the z-score value.

XBP1_WT.2 (41757)

XBP1_WT.2 (41754)



XBP1_WT.2 (41756)



XBP1_WT.2 (41755)

Compound IDs and		Mean pairwise		Compound				
common names (where		replicates	Correlation	rank when			Distinguishing individual features for the compound relative to	Number of PubChem aggregate which
available); blue/red colored	Chemical structure	correlation of the	between 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)	Common distinguishing feature categories in the compound and the gene relative to the untreated samples	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 active a small z-score in magnitude) or expectative in the general property of the compound was active a small z-score in magnitude) or expectative in the general property of the compound was active a small z-score in magnitude) or expectative in the general property of the compound was active a small z-score in magnitude) or expectative in the general property of the compound was active a small z-score in magnitude) or expectative in the general property of the compound was active a small z-score in magnitude).	·
box means the matching		compound signature		against the				
compound is		(95th DMSO		gene using				
positively/negatively		replicate correlation	the gene	L1000			small z-score in magnitude) or oppositely active in the gene	itemized
correlated with the cluster		is 0.51)		profiling				

