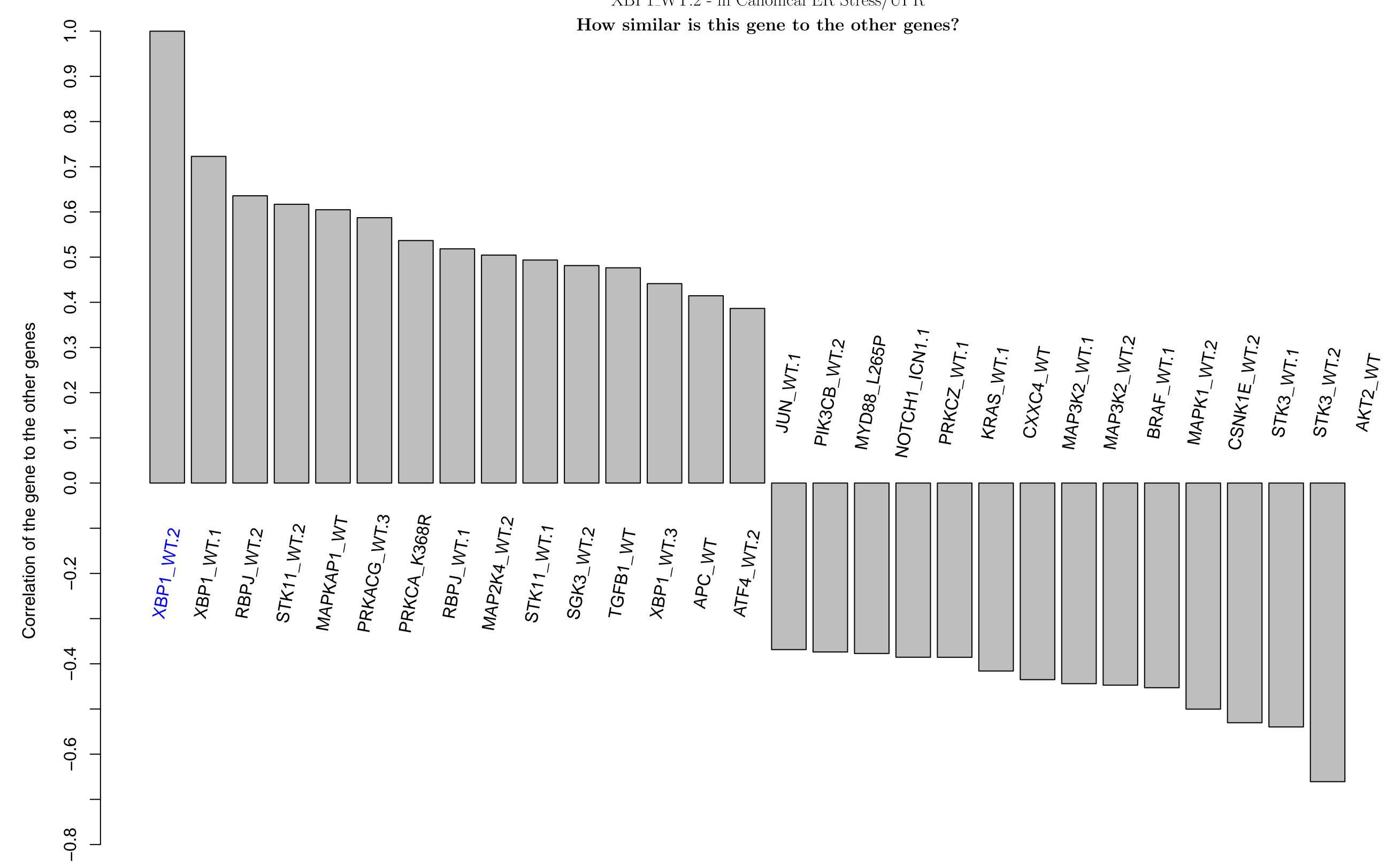
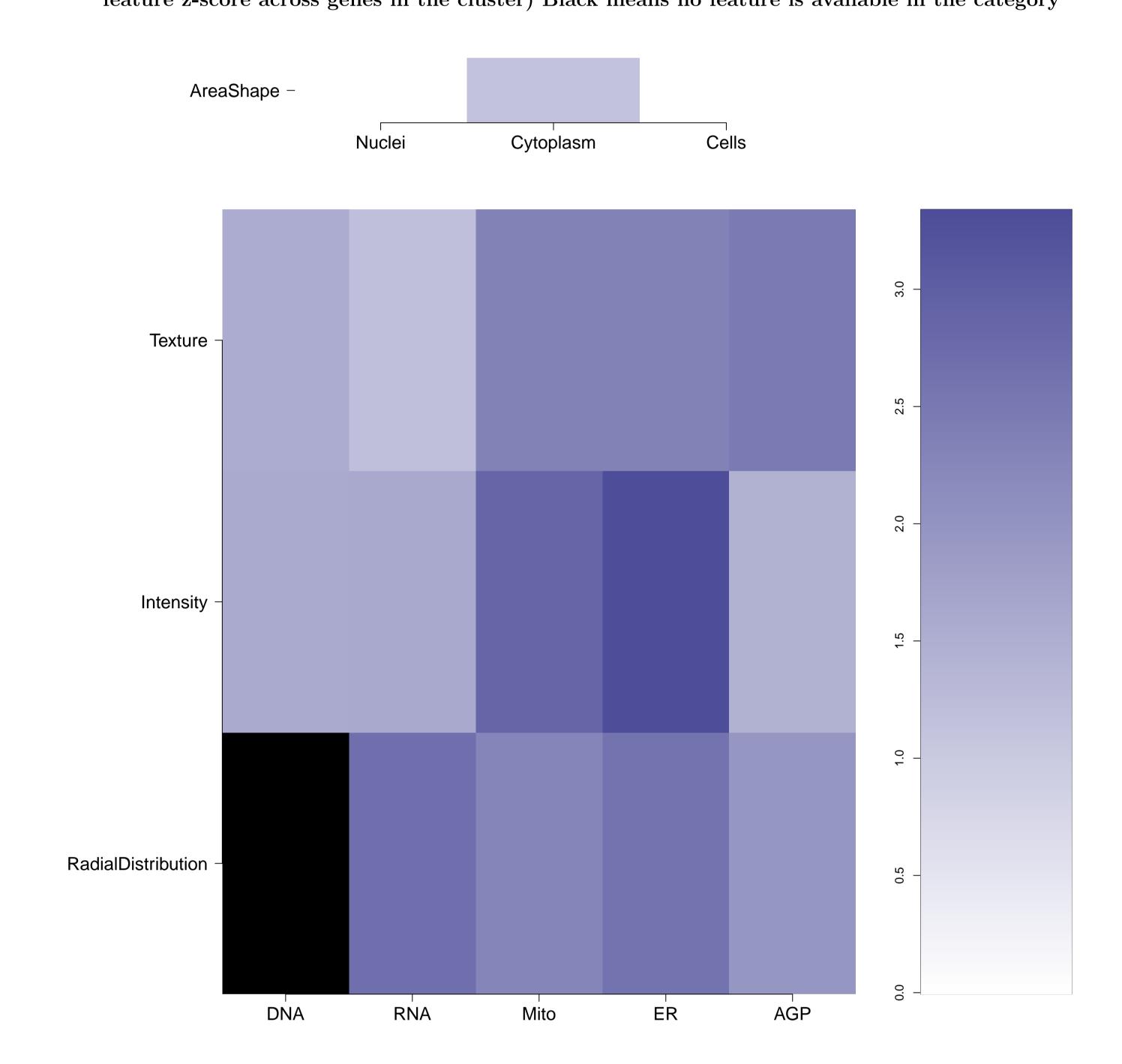
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein XBP1_WT.2 - in Canonical ER Stress/UPR

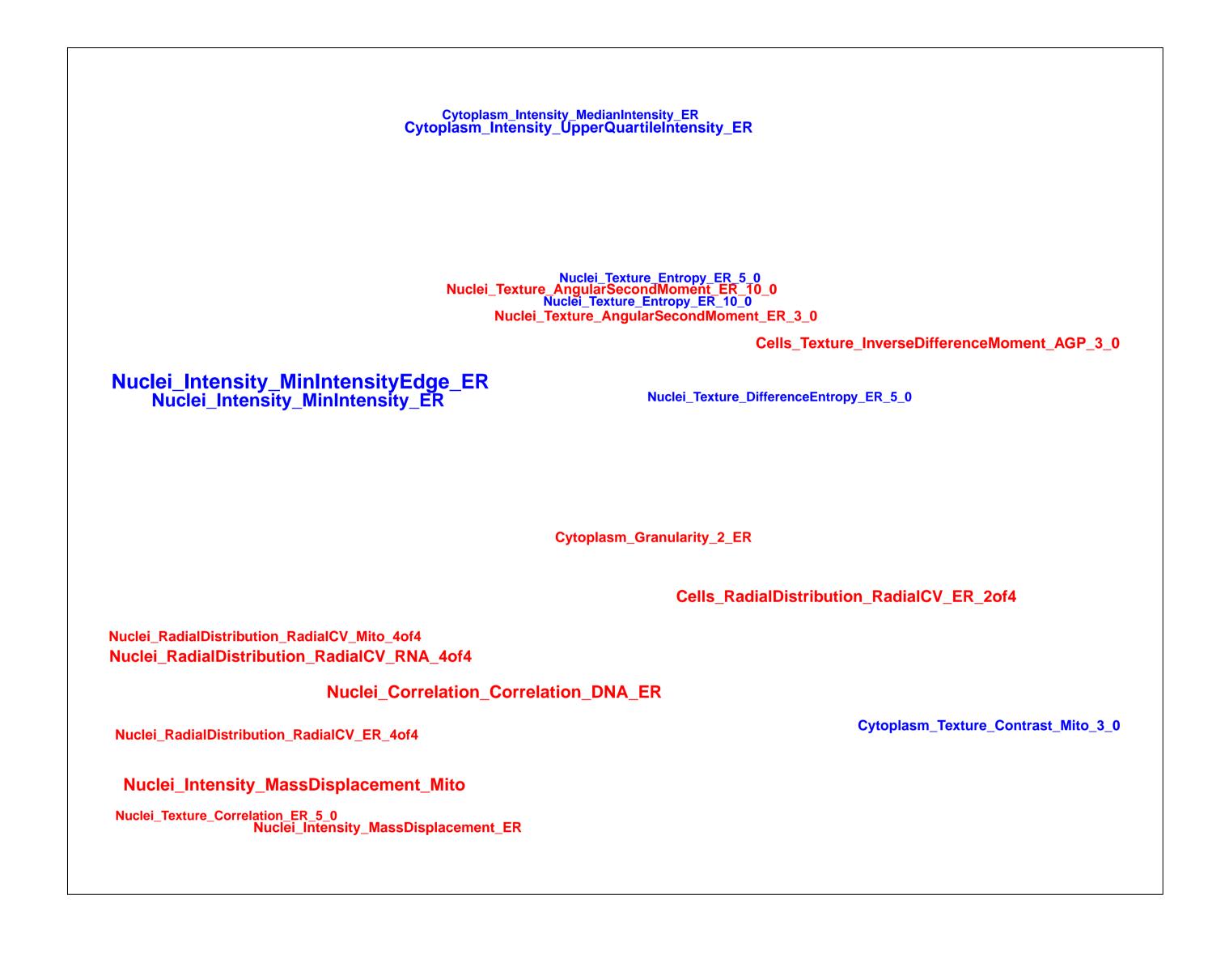


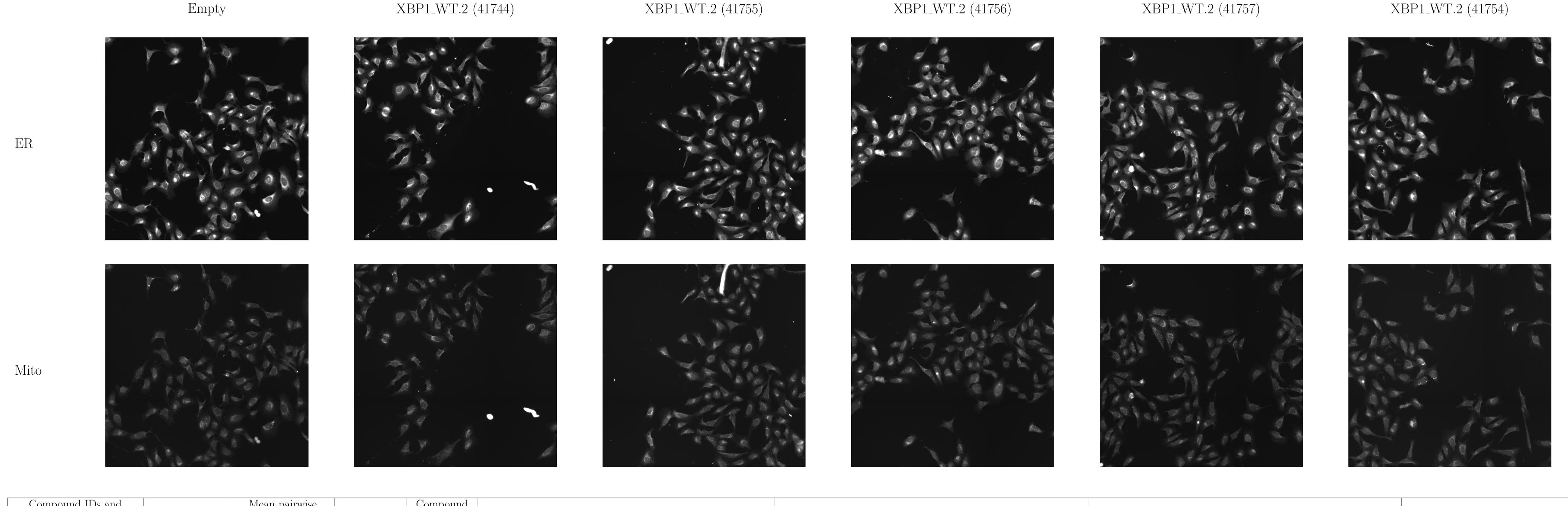
What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a feature z-score across genes in the cluster) Black means no feature is available in the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



Mito

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 box means the matching		replicates correlation of the compound signature	compound the gene using		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 (= small z-score in magnitude) or oppositely active in the gene Number of PubChe the compound was which the compound was which the compound item.	the compound was tested; assays in
compound is positively/negatively		(95th DMSO replicate correlation		gene using				which the compound was active are itemized
correlated with the cluster		is 0.52)		profiling				

