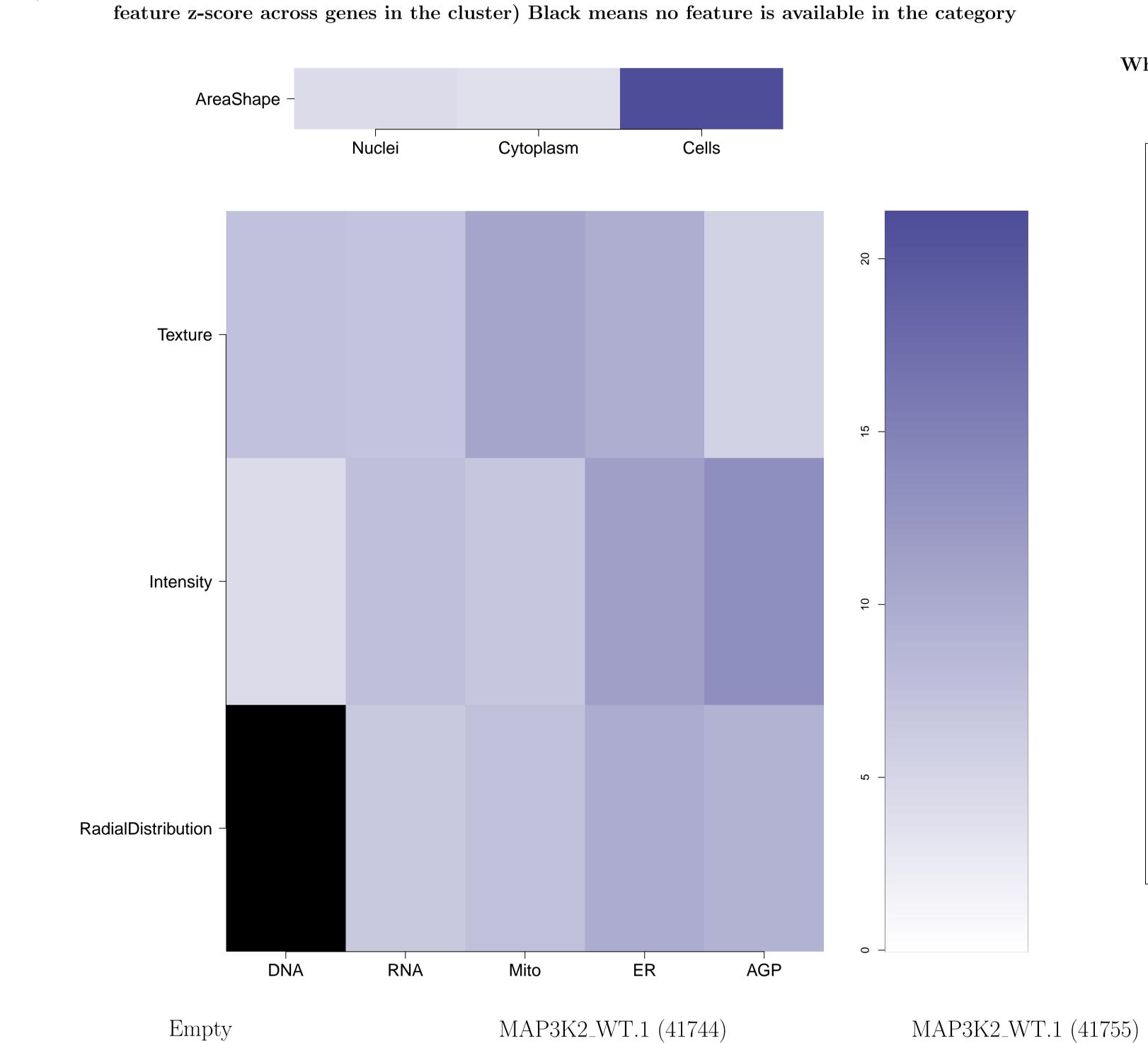


What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a



RNA

AGP

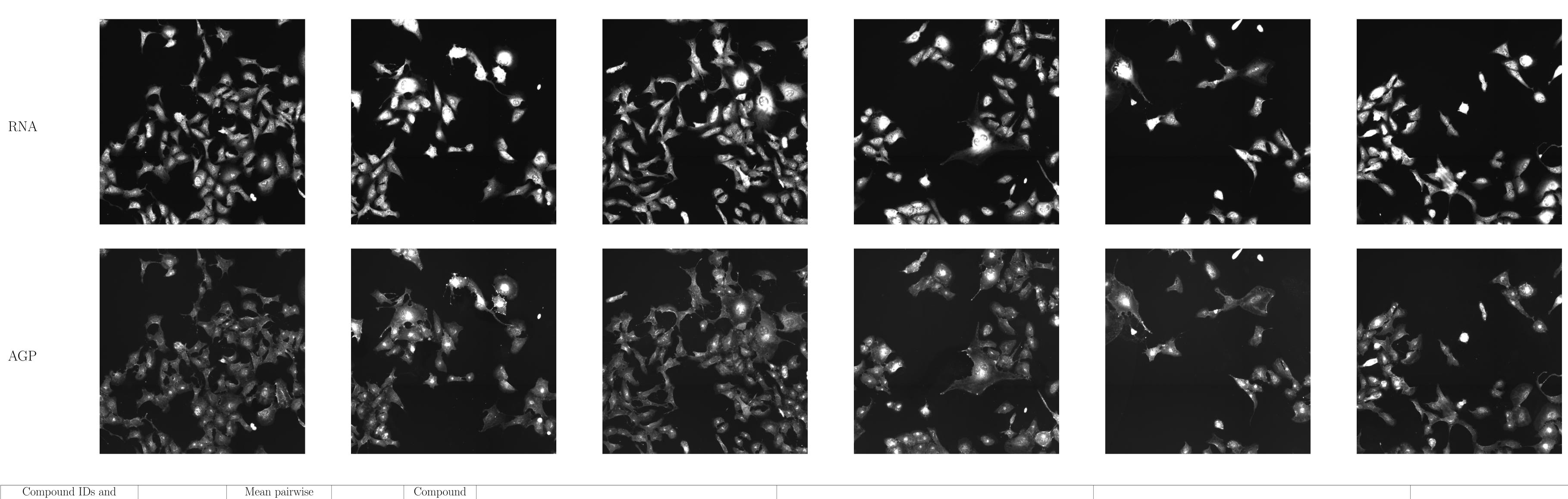
Which individual morphological features are distinguishing in the gene relative to the untreated samples? Blue/Red means the feature has a positive/negative z-score. Size is proportional to the z-score value.

MAP3K2_WT.1 (41757)

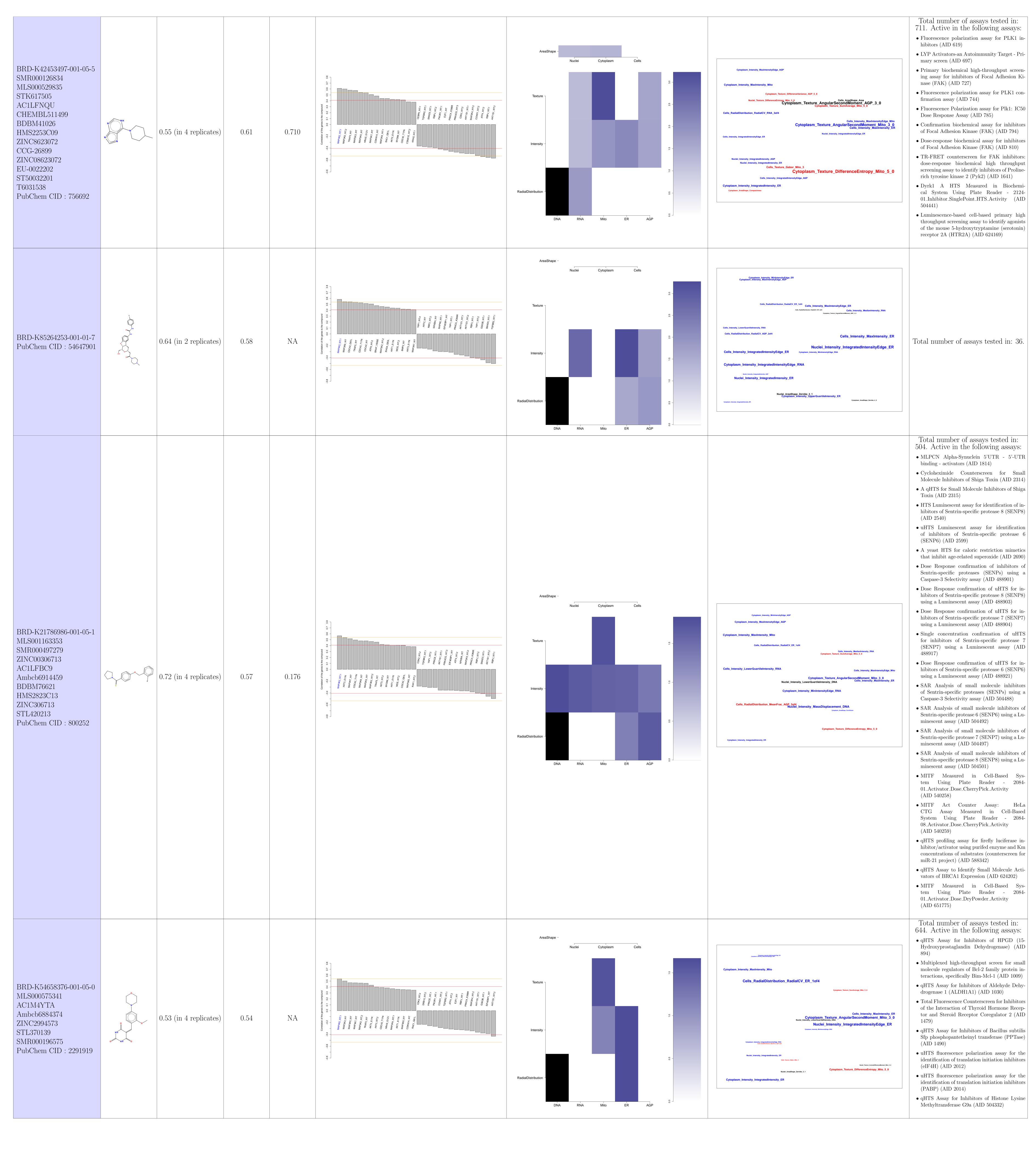
 $MAP3K2_WT.1 (41754)$

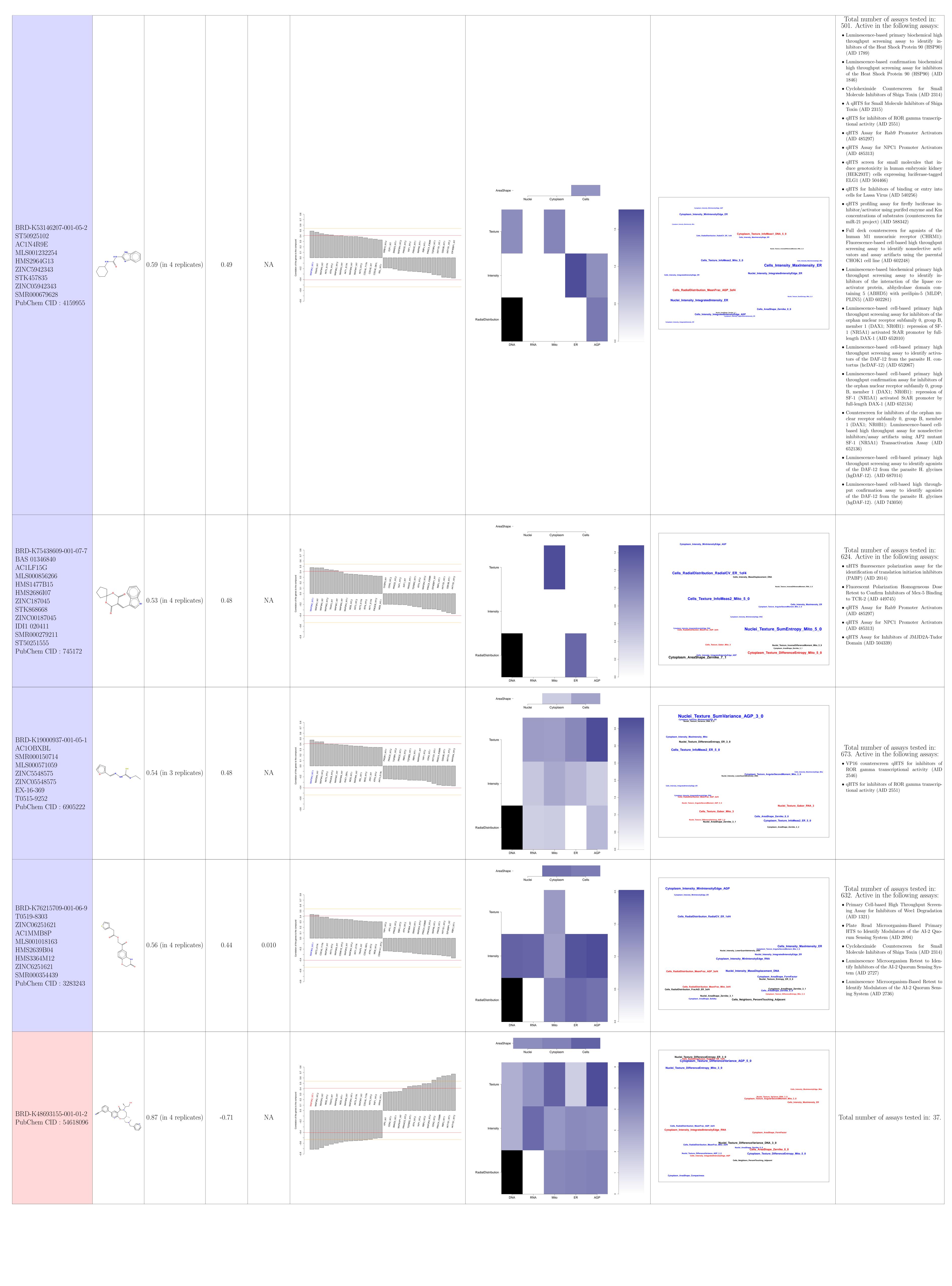


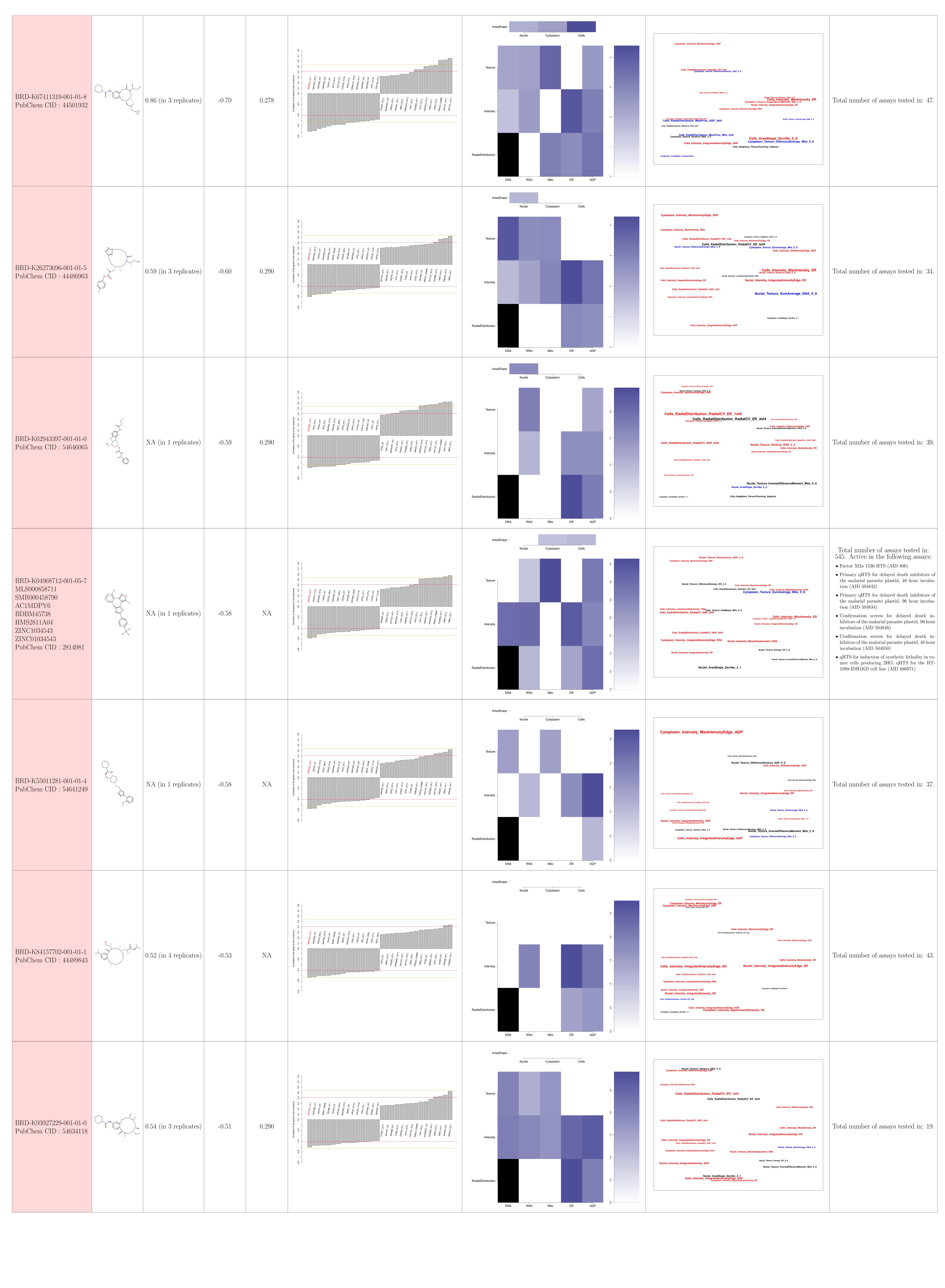
 $MAP3K2_WT.1 (41756)$



common names (where available); blue/red colored box means the matching compound is positively/negatively correlated with the cluster	Chemical structure	replicates correlation of the compound signature (95th DMSO replicate correlation is 0.52)	Correlation between compound the gene	against the	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)	LCamman distinsilishins teatiire catesaries in the compaiind 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	the compound was tested; assays in
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Cytoplasm Cells_RadialDistribution_RadialCV_ER_10f4
Nuclei_Texture_DifferenceEntropy_ER_3_0
Cells_RadialDistribution_RadialCV_ER_4of4 Cells_Intensity_MaxIntensityEdge_DNA BRD-K09160804-001-01-4 PubChem CID : 54619505 0.65 (in 4 replicates) Total number of assays tested in: 37. 0.892Cells_RadialDistribution_RadialCV_AGP_2of4 Cytoplasm_Texture_AngularSecondMoment_Mito_3_0 Intensity -Cells_RadialDistribution_RadialCV_AGP_3of4 Cells_RadialDistribution_FracAtD_ER_3of4
Cells_Intensity_IntegratedIntensityEdge_AGP
Cytoplasm_AreaShape_Solidity Cytoplasm_Texture_InfoMeas2_ER_5_0 RadialDistribution -Cytoplasm_AreaShape_Zernike_4_2 Mito ER AGP