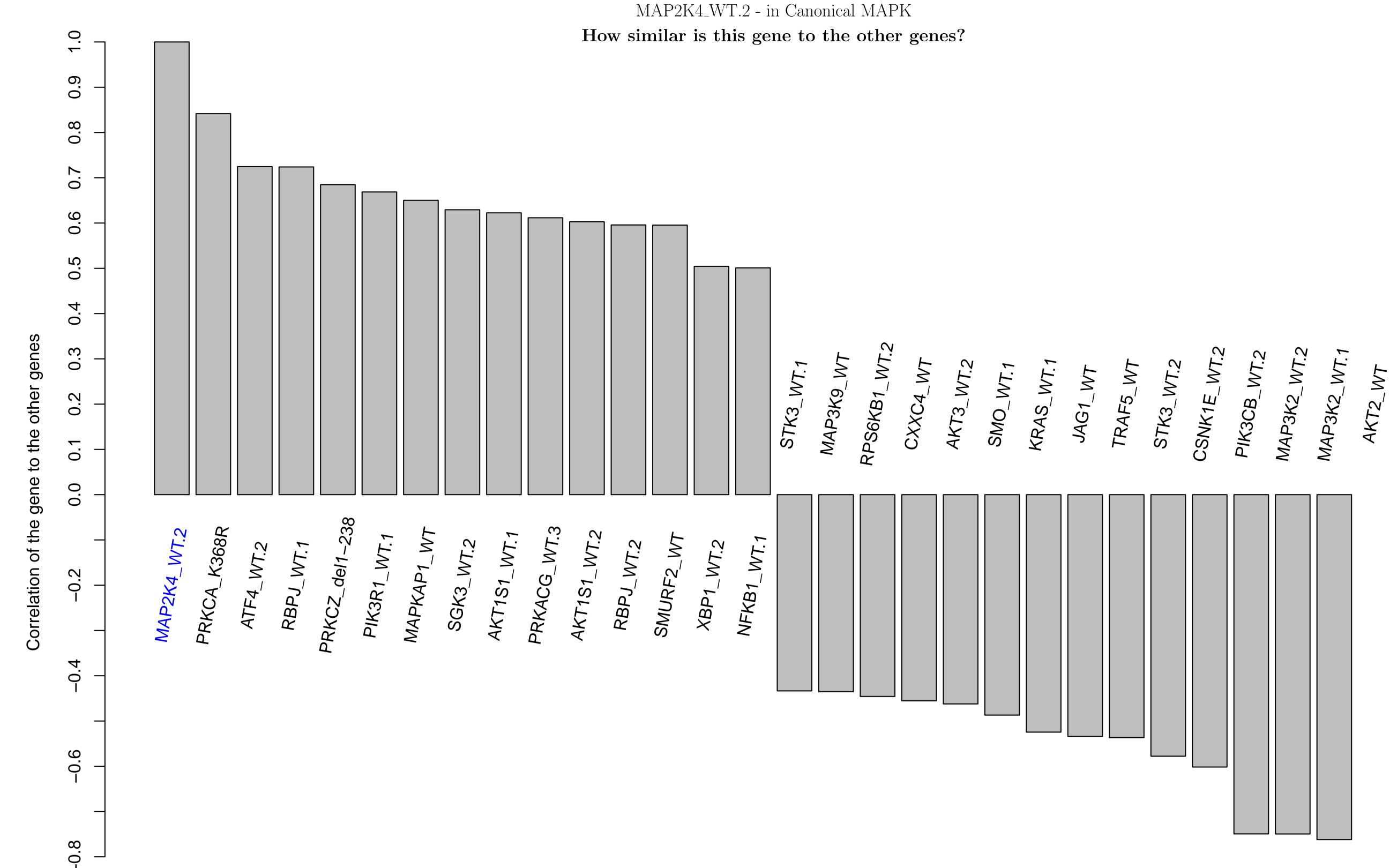
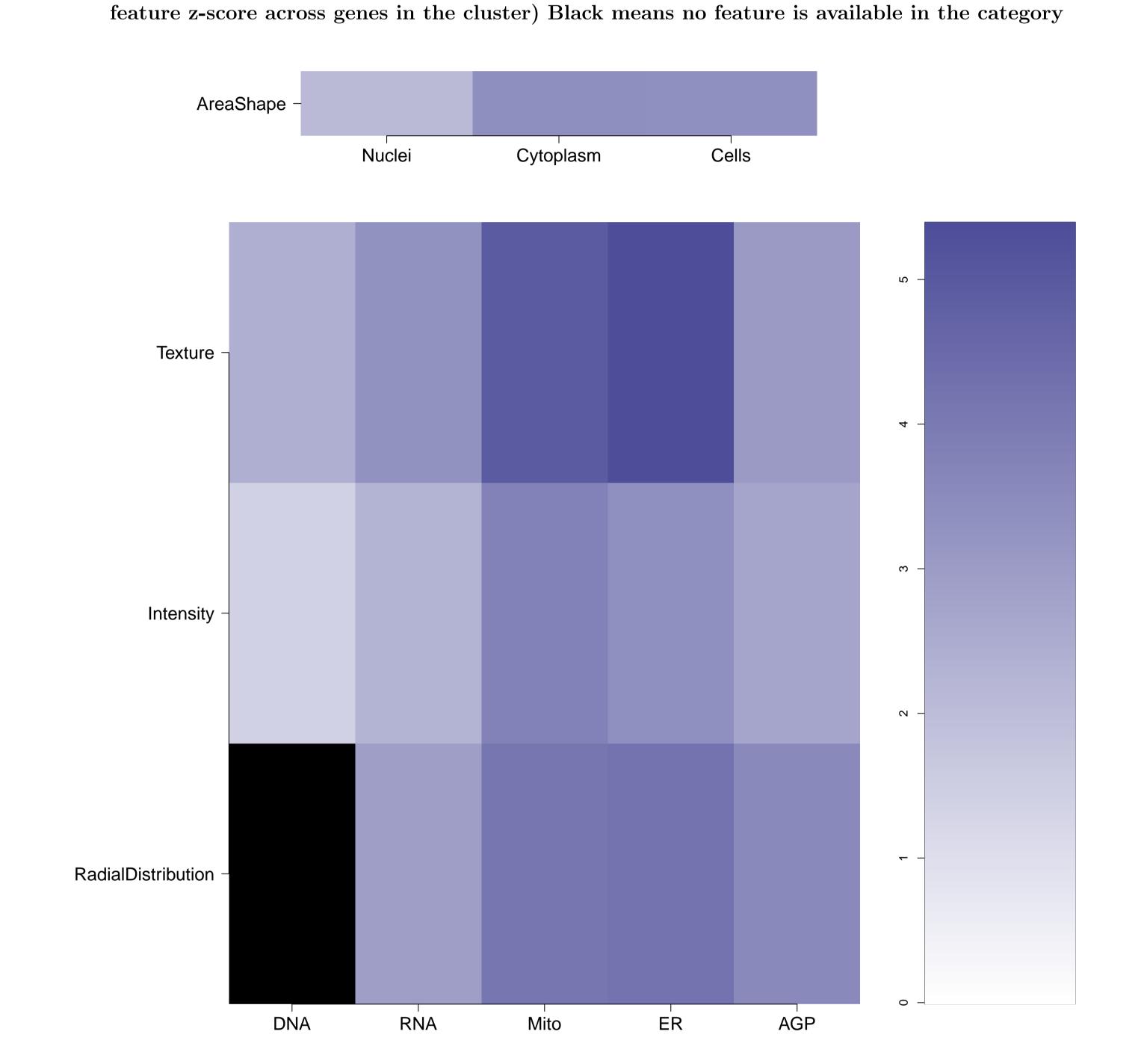
## CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein



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



Correlation

between

compound

the gene

scored

against the

gene using

L1000

profiling

correlation of the

compound signature

(95th DMSO

replicate correlation

is 0.52)

Chemical

structure

ER

Mito

available); blue/red colored

box means the matching

compound is

positively/negatively

correlated with the cluster

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.

Distinguishing individual features for the compound relative to Number of PubChem assays in which

untreated samples. Black means a mismatch; i.e. active (= high

z-score in magnitude) in the compound, and either inactive (=

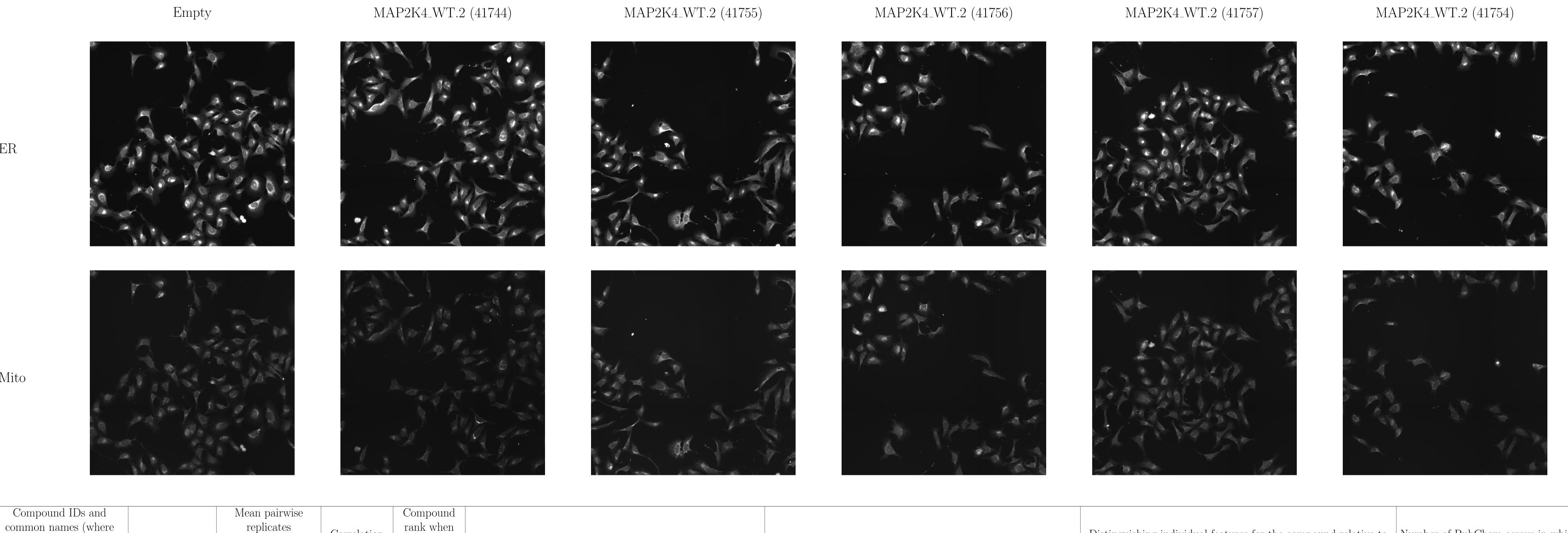
small z-score in magnitude) or oppositely active in the gene

the compound was tested; assays in

which the compound was active are

itemized





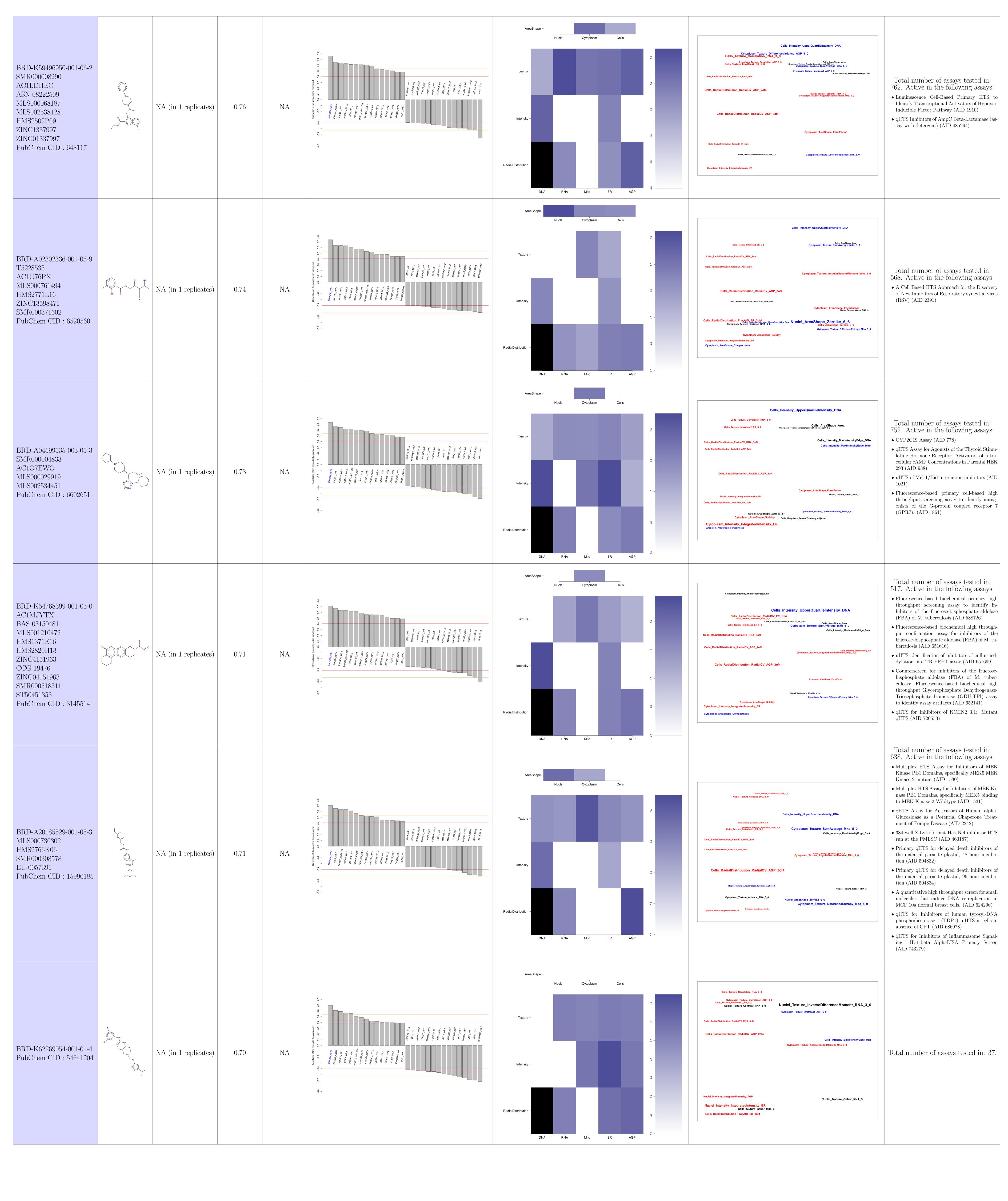
Common distinguishing feature categories in the compound and

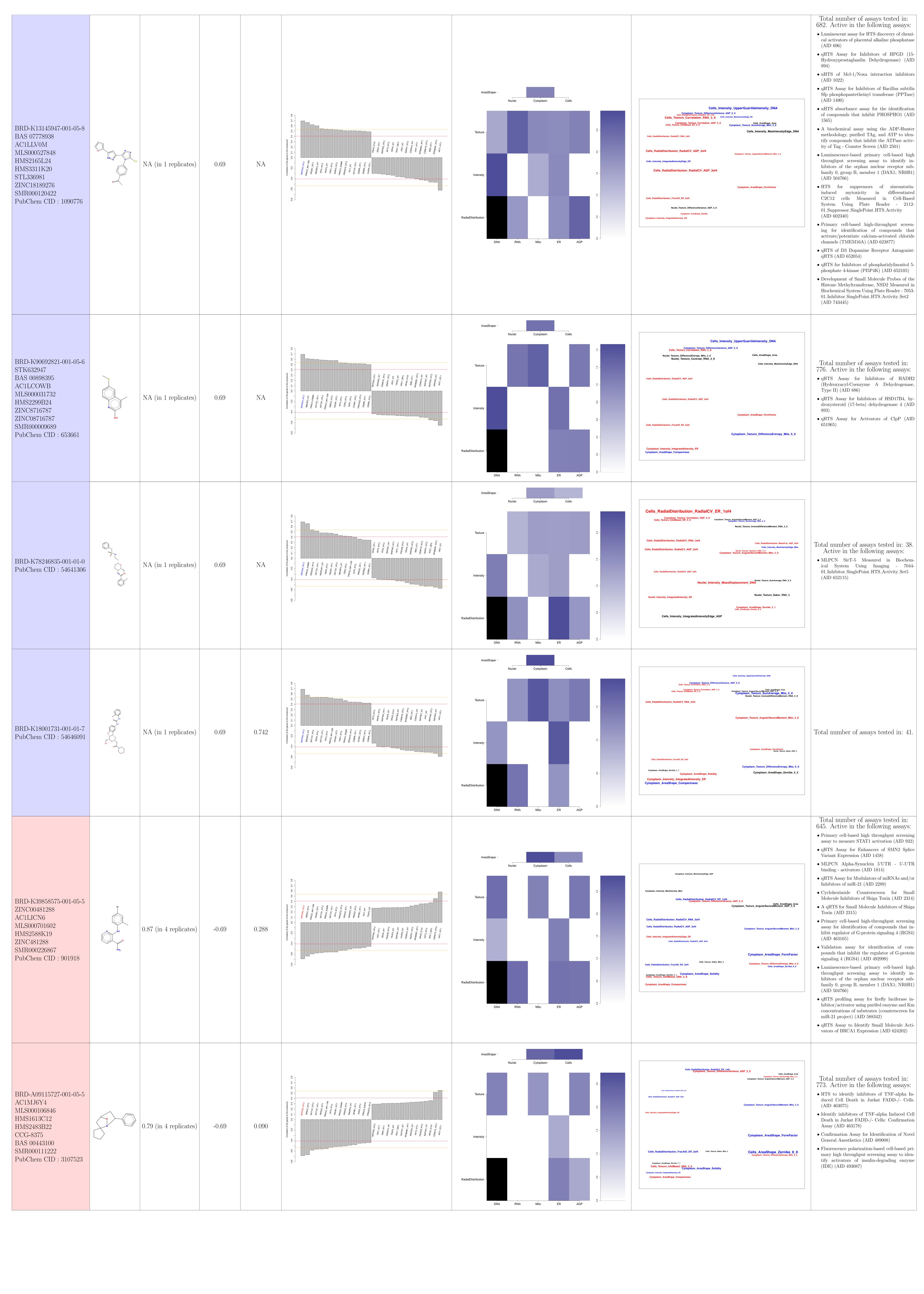
the gene relative to the untreated samples

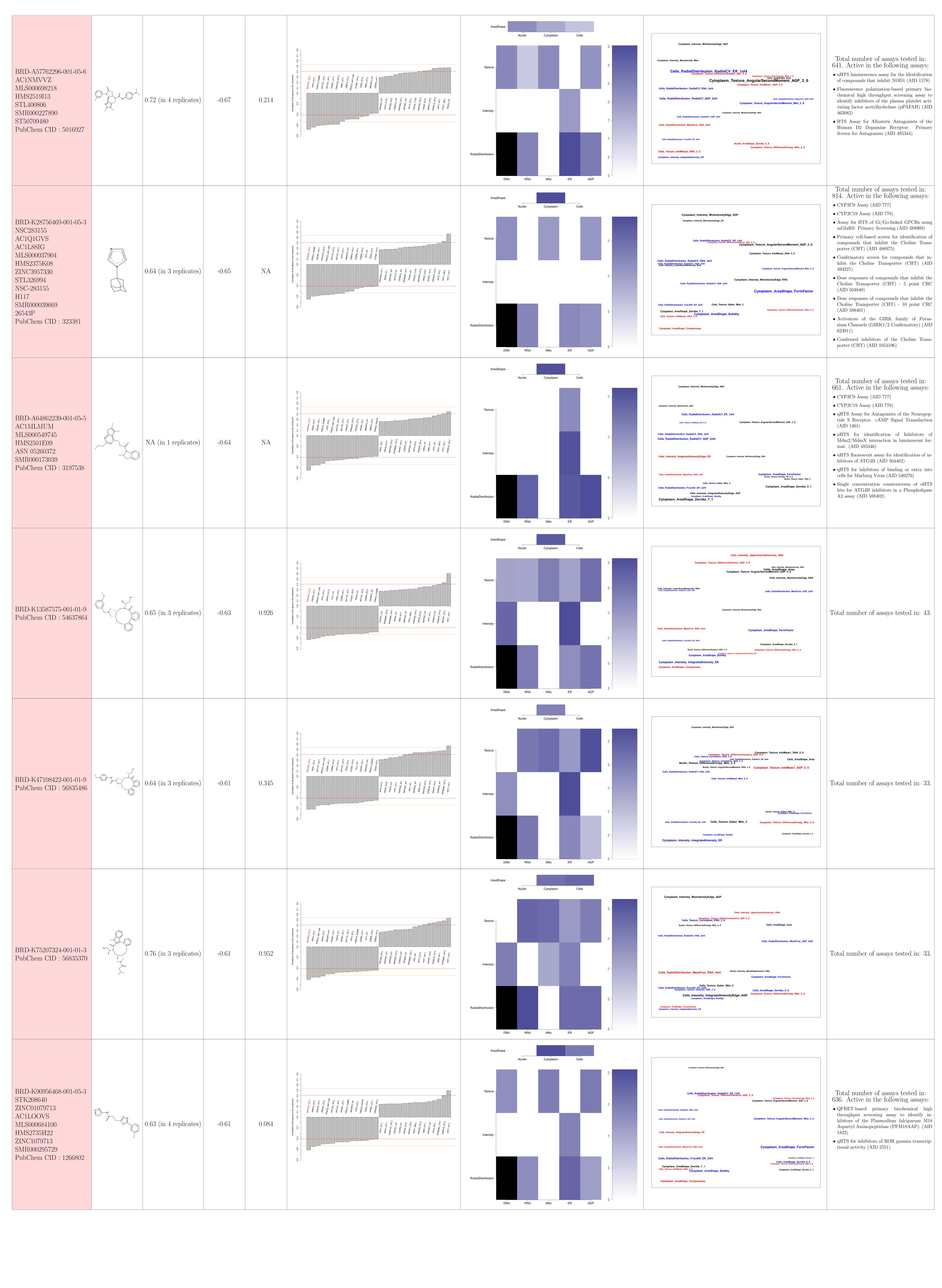
How similar is the compound signature to the genes in this

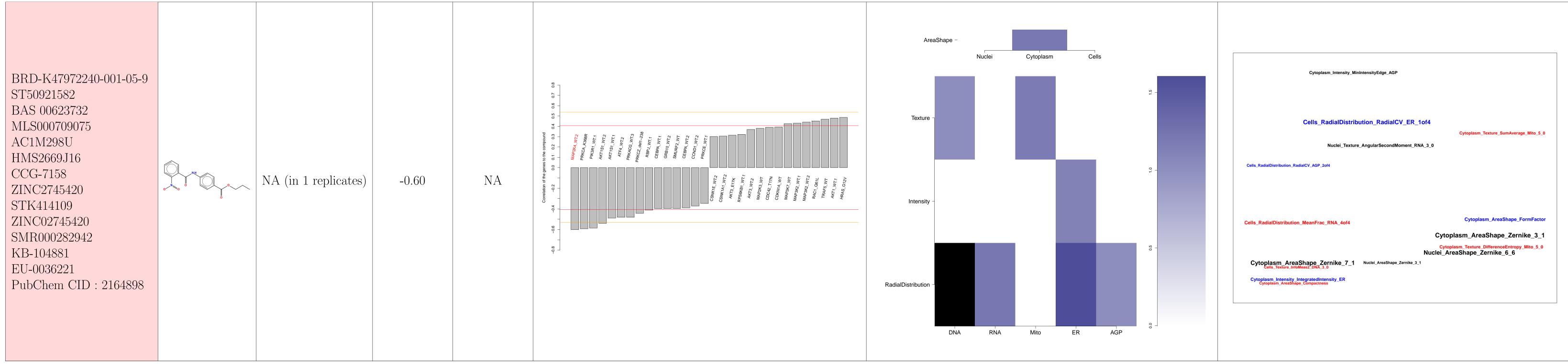
experiment? (Yellow and red lines correspond to top/bottom

1st and 5th percentile DMSO correlation to all the genes)









- Total number of assays tested in: 632. Active in the following assays: • Aqueous Solubility from MLSMR Stock Solutions (AID 1996) • Cycloheximide Counterscreen for Small Molecule Inhibitors of Shiga Toxin (AID 2314) • A qHTS for Small Molecule Inhibitors of Shiga
- Toxin (AID 2315) • Primary cell-based screen for identification of compounds that inhibit the two-pore domain
- potassium channel KCNK9 (AID 488922) • Confirmatory screen for identification of compounds that inhibit the two-pore domain potassium channel (KCNK9) (AID 492992)
- Primary cell-based screen for identification of
- compounds that inhibit the two-pore domain potassium channel KCNK3 (AID 602410) • Confirmation assay for identification of compounds that inhibit the two-pore domain potassium channel KCNK3 [Primary Screening]
- (AID 651638)• Small Molecule Inhibitors of FGF22-Mediated Excitatory Synaptogenesis and Epilepsy Measured in Biochemical System Using RT-PCR - 7012-01\_Inhibitor\_SinglePoint\_HTS\_Activity (AID 651658)