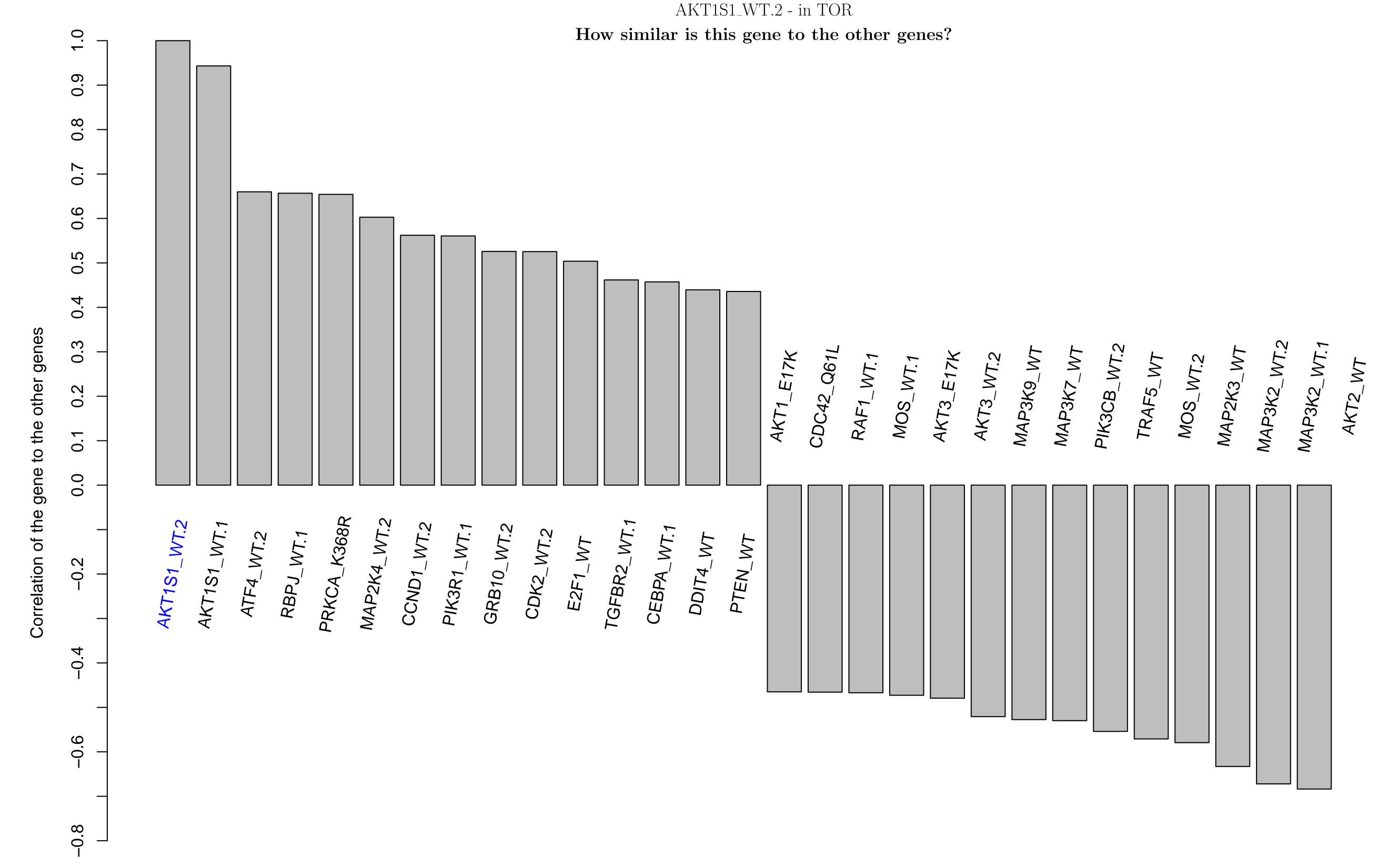
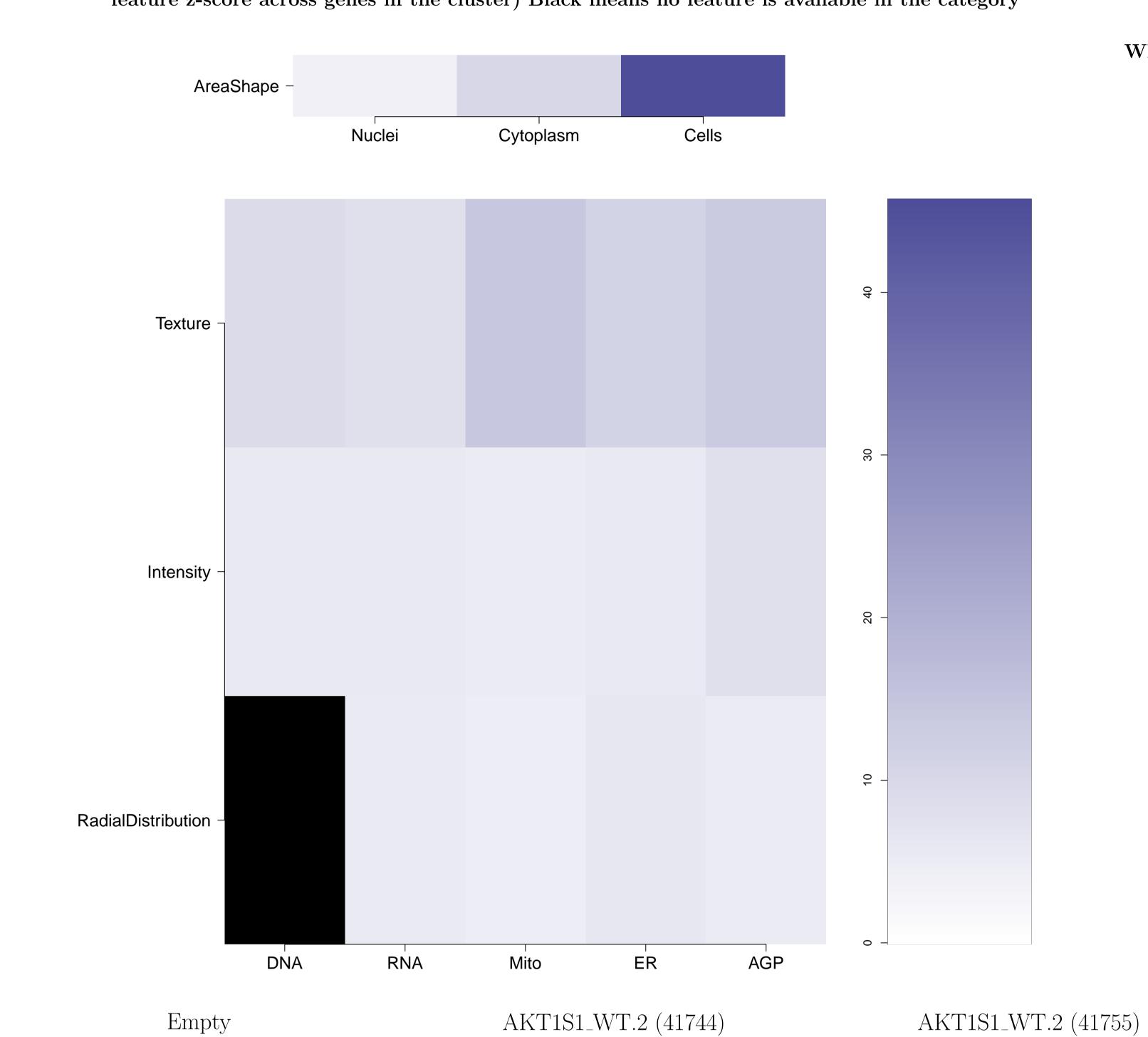
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein

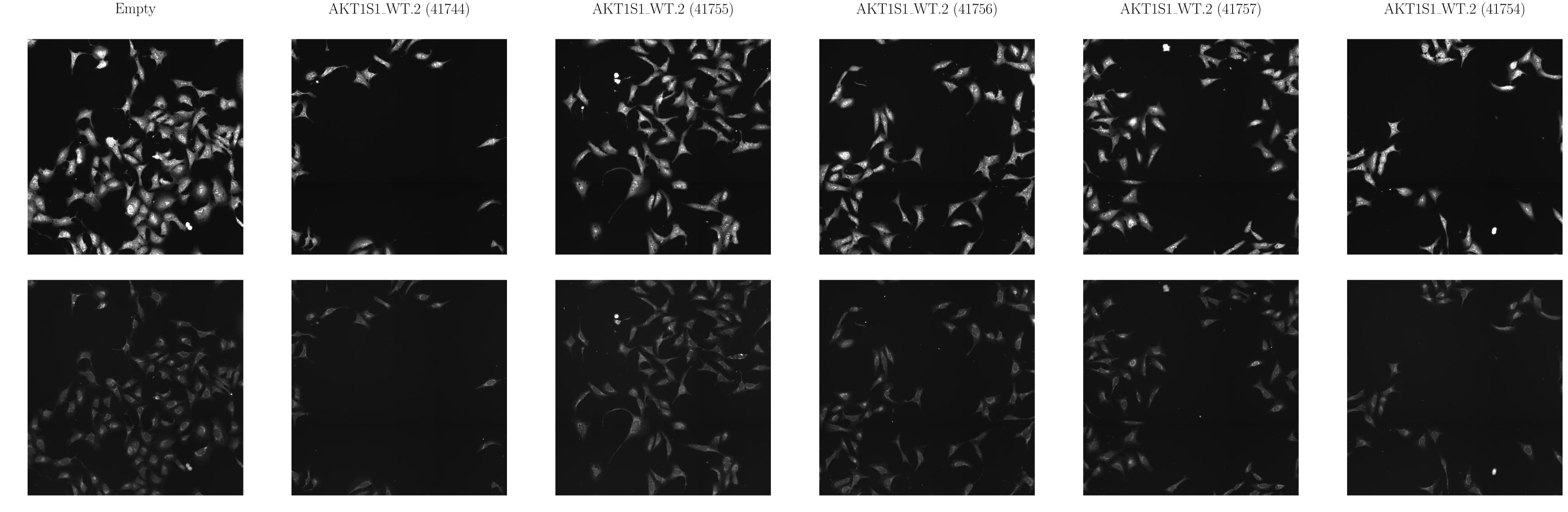


What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a feature z-score across genes in the cluster) Black means no feature is available in the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



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 compound is positively/negatively correlated with the cluster | Chemical<br>structure | replicates correlation of the compound signature (95th DMSO replicate correlation is 0.51) | Correlation between compound the gene | rank when 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 | 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 |
|   |                       | / /  |                                       | 1                |   |  |   |                                    |

