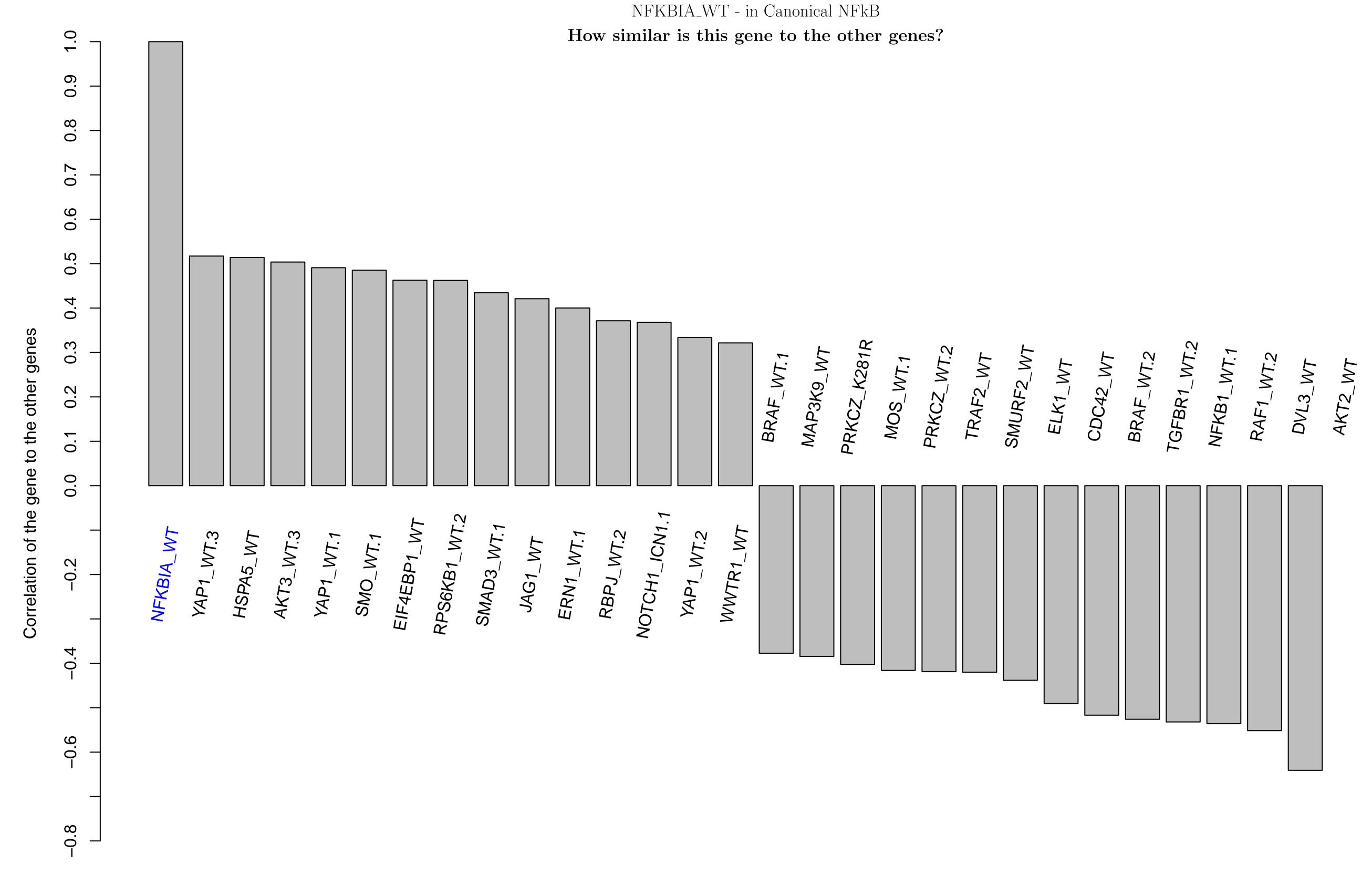
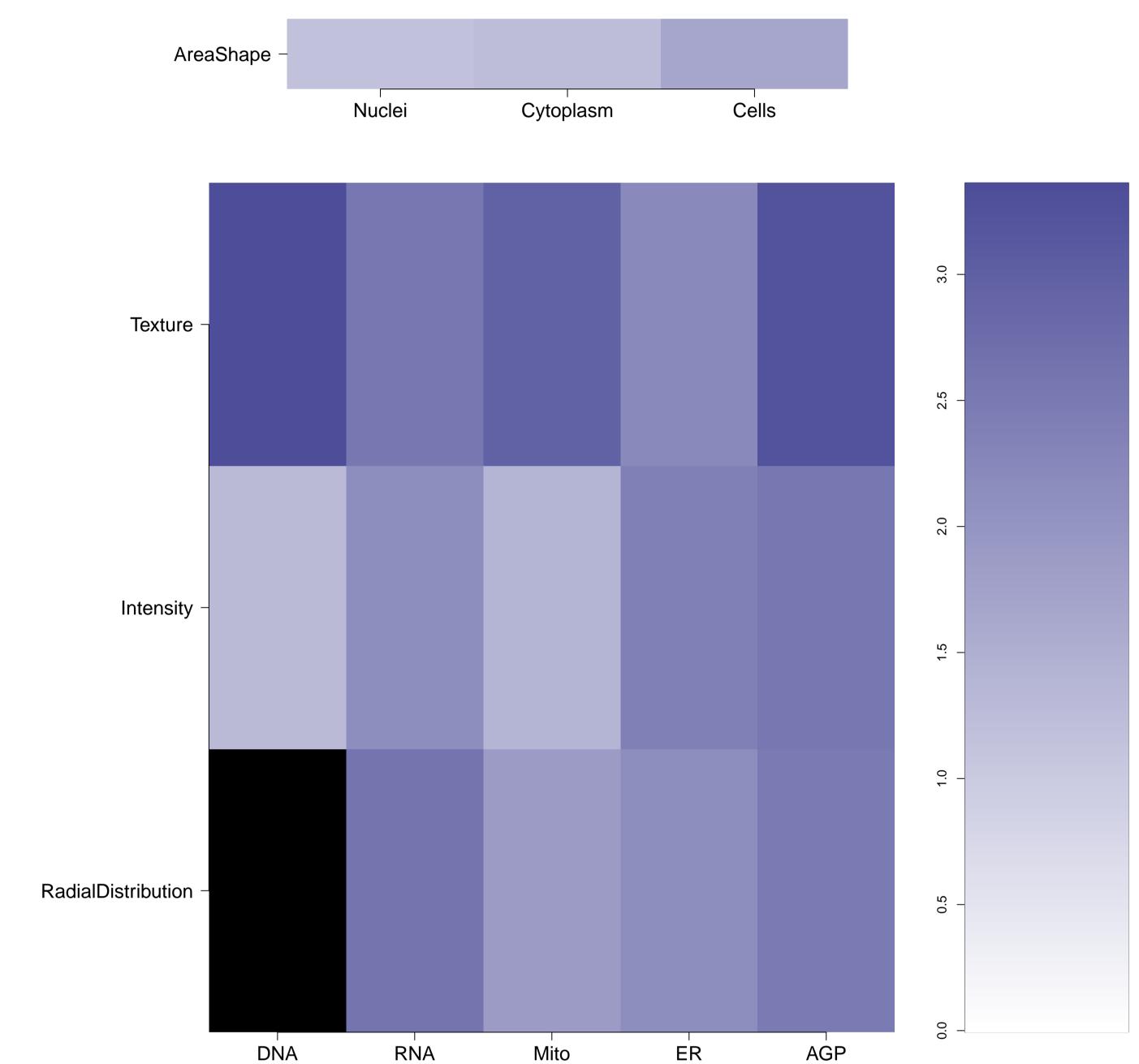
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein

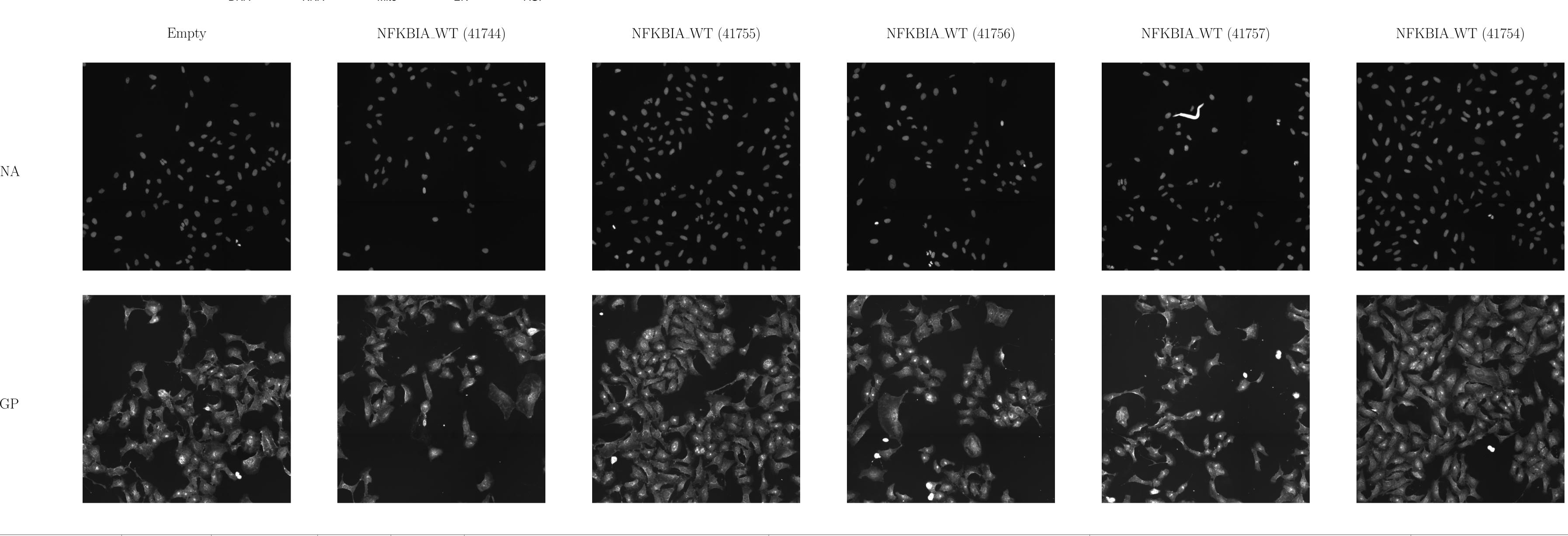


What groups of morphological features are distinguishing in the cluster relative to the untreated samples? (maximum of absolute m-score for the features belonging to the same category; m-score defined as median of a feature z-score across genes in the cluster) Black means no feature is available in the 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 common names (where available); blue/red colored box means the matching compound is positively/negatively correlated with the cluster Chemical structure Chemical structure Chemical structure (95th DMSO replicate correlation is 0.52) Mean pairwise replicates correlation between compound signature (95th DMSO replicate correlation is 0.52) Compound rank when scored against the gene using L1000 L1000 profiling Compound rank when scored against the gene using L1000 The profiling is the compound signature to the genes in this experiment? (Yellow and red lines correspond to top/botto 1st and 5th percentile DMSO correlation to all the genes)	± 1 Common distinguishing teature categories in the compound and \pm intreated samples. Black means a mismatch, i.e. active $1-$ high \pm the compound was tested, assays in
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