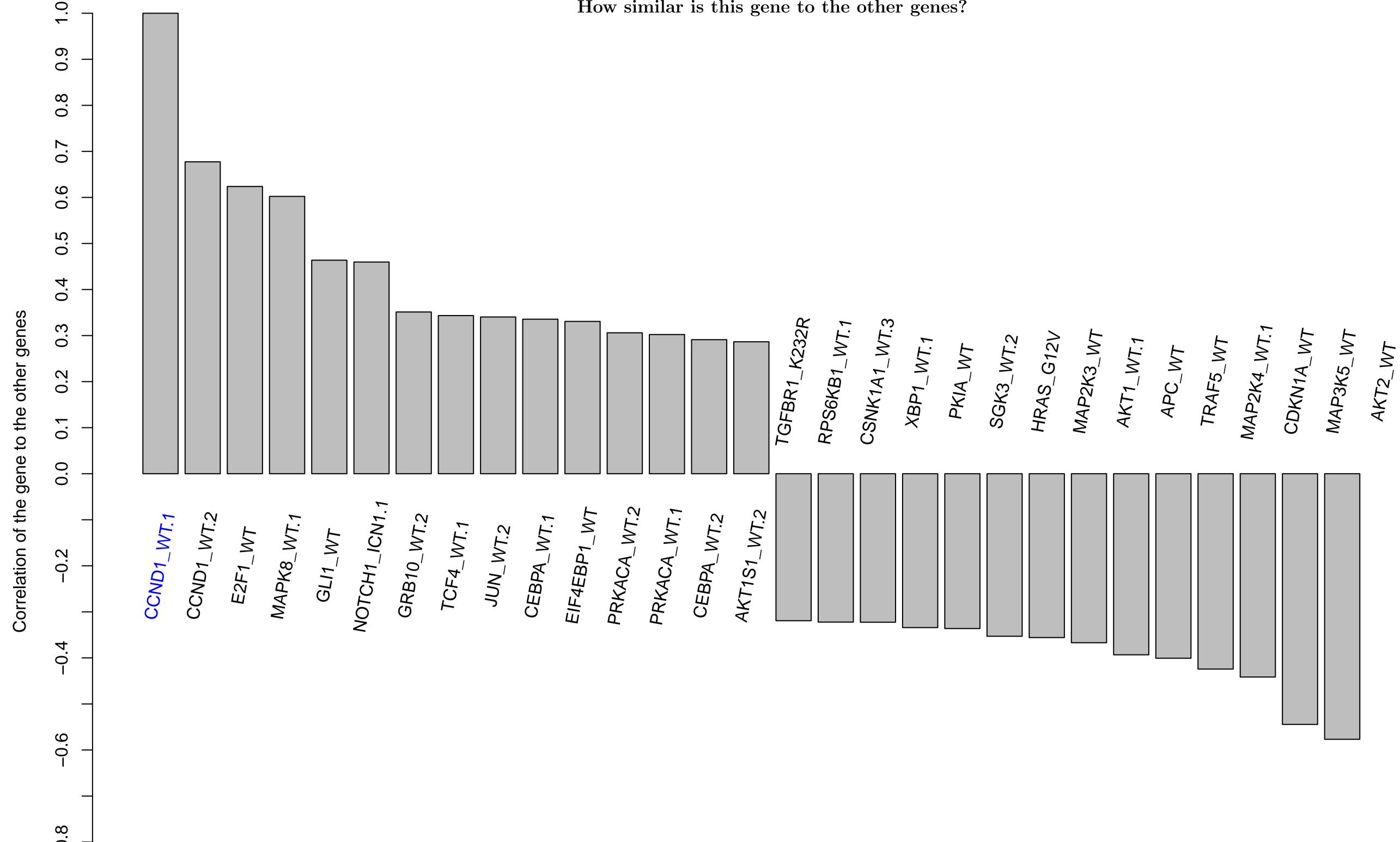
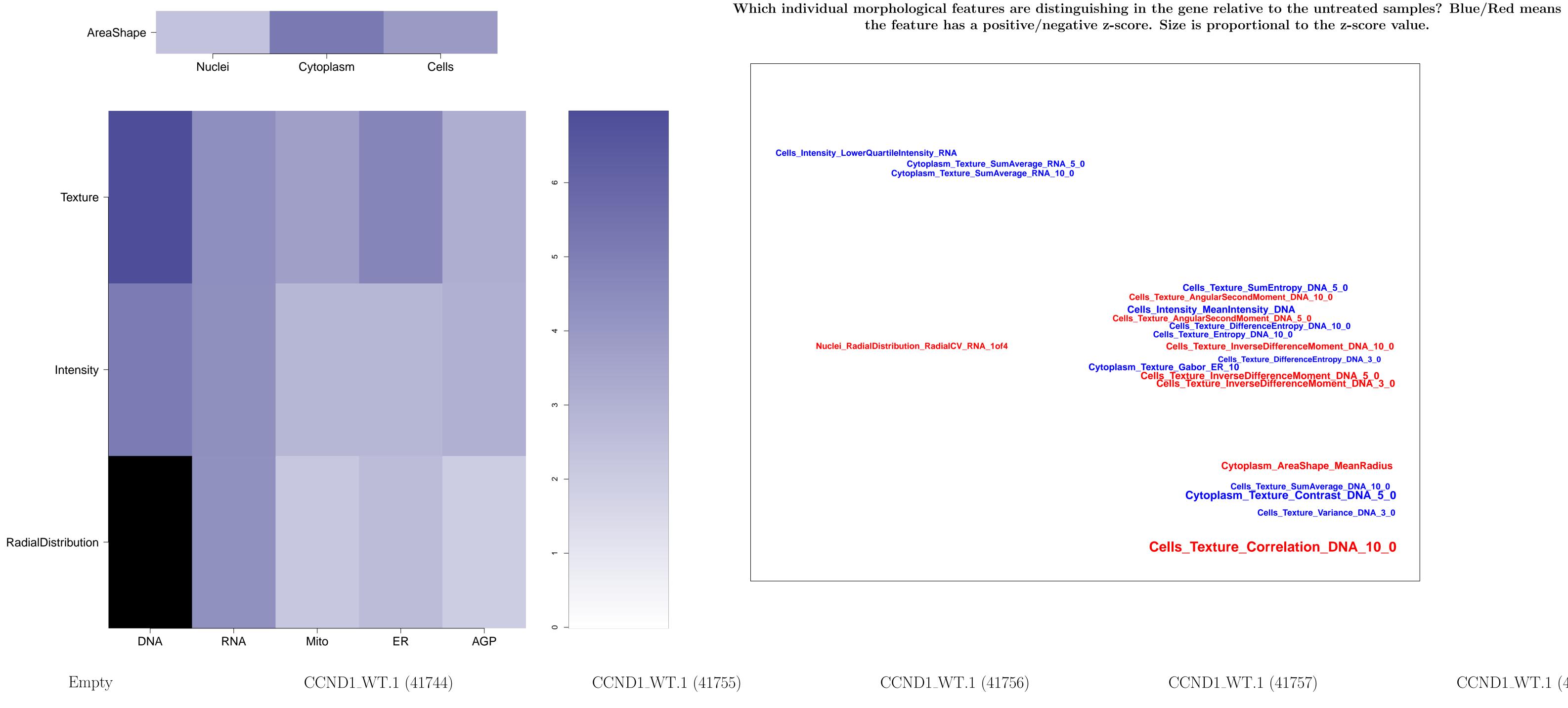
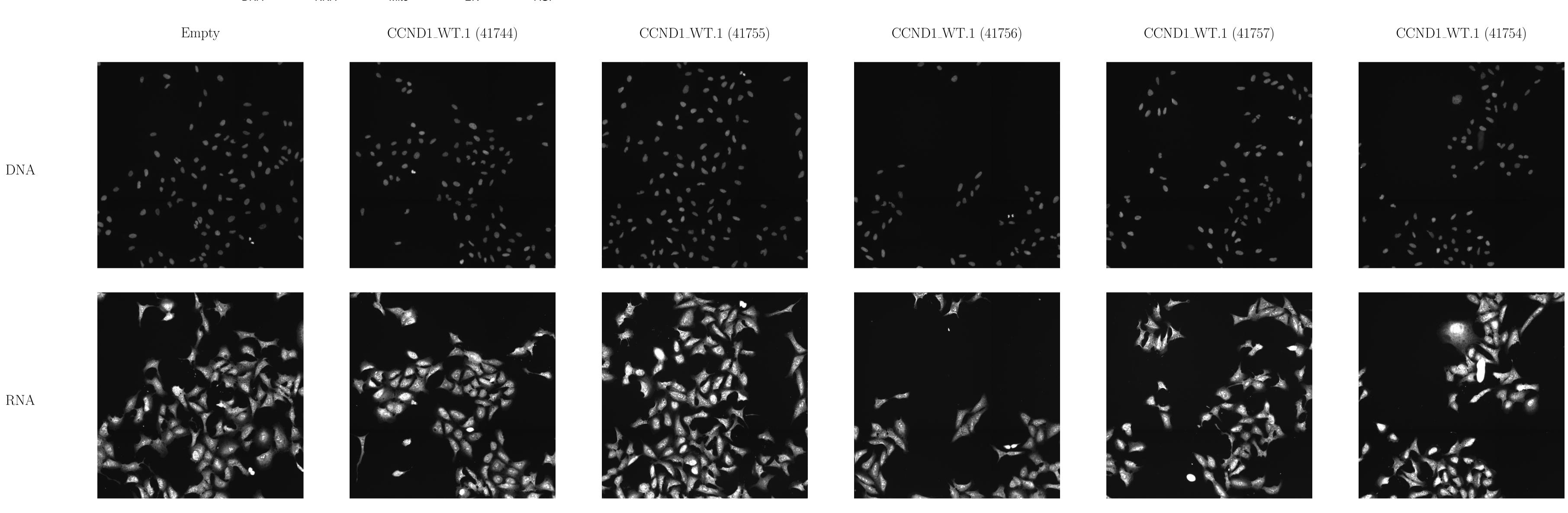
CONFIDENTIAL, contact the Imaging Platform to collaborate on the findings herein CCND1\_WT.1 - in Canonical Cell Cycle How similar is this gene to the other genes?



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





Compound IDs and Compound Mean pairwise rank when replicates common names (where Correlation Distinguishing individual features for the compound relative to Number of PubChem assays in which How similar is the compound signature to the genes in this available); blue/red colored correlation of the scored Chemical Common distinguishing feature categories in the compound and the compound was tested; assays in between untreated samples. Black means a mismatch; i.e. active (= high experiment? (Yellow and red lines correspond to top/bottom box means the matching against the compound signature compound the gene relative to the untreated samples z-score in magnitude) in the compound, and either inactive (= which the compound was active are structure compound is (95th DMSO 1st and 5th percentile DMSO correlation to all the genes) gene using the gene small z-score in magnitude) or oppositely active in the gene itemized positively/negatively replicate correlation L1000 correlated with the cluster is 0.52) profiling

