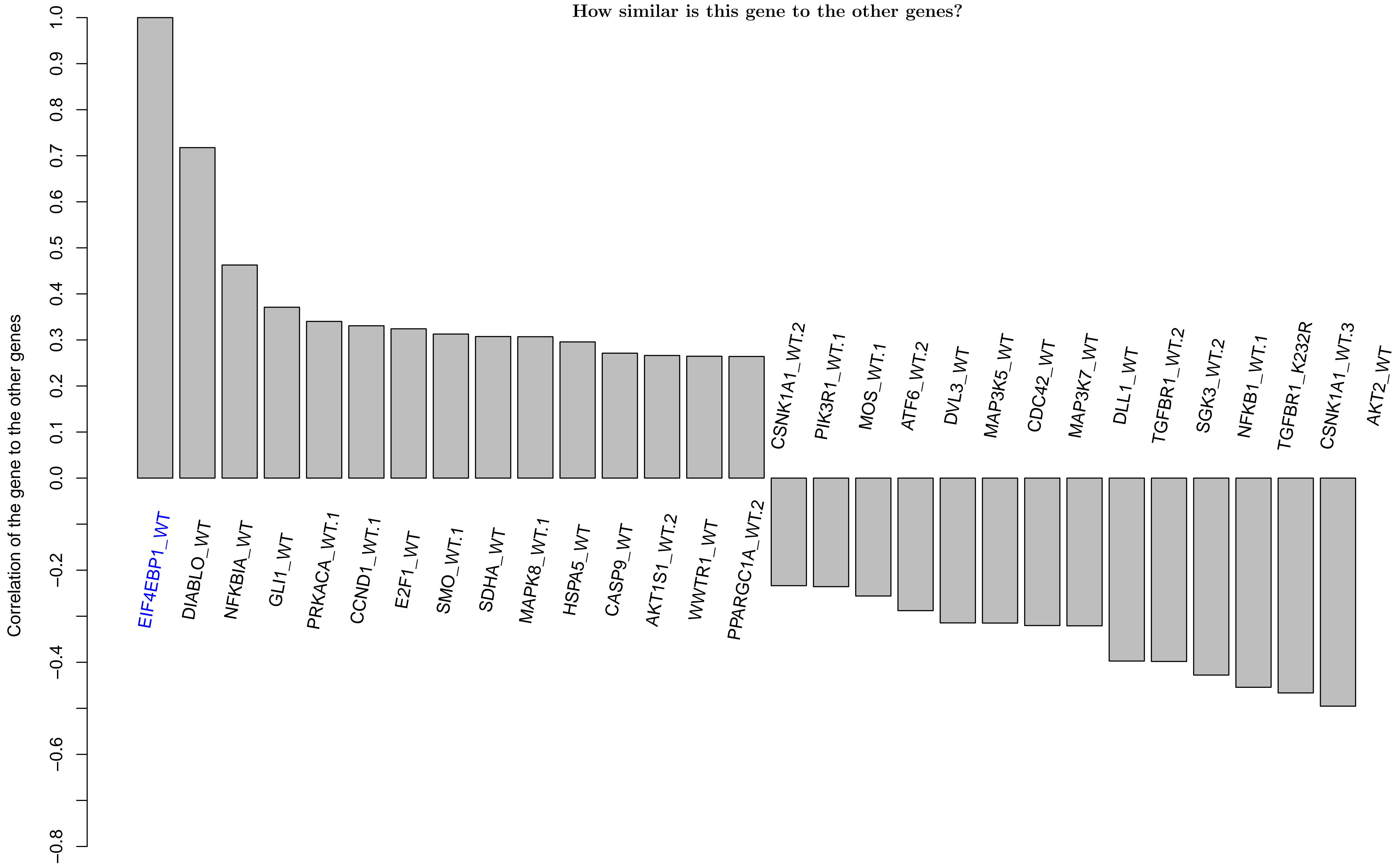
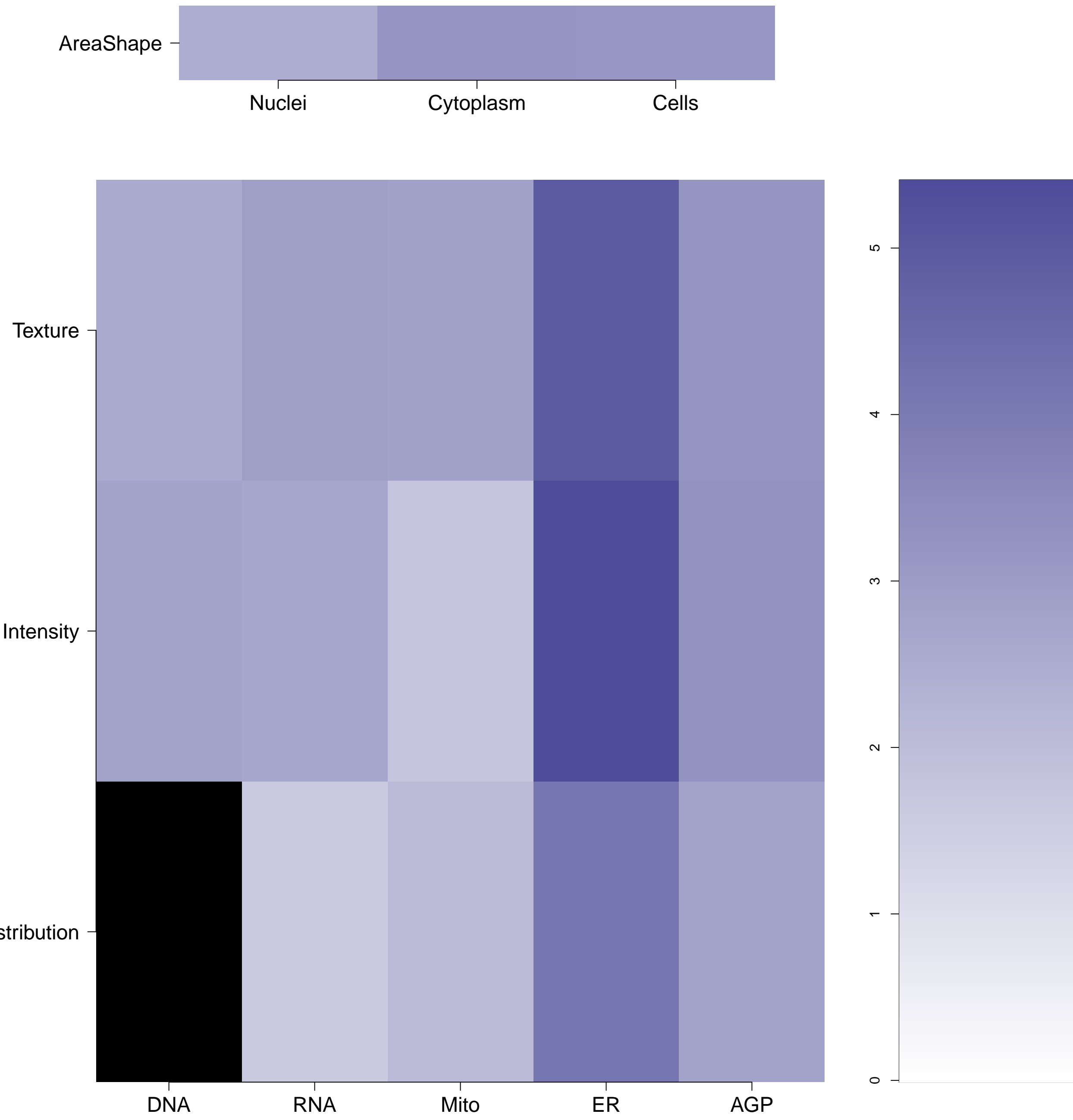


EIF4EBP1.WT - in Canonical TOR

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



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.



Empty

EIF4EBP1.WT (41744)

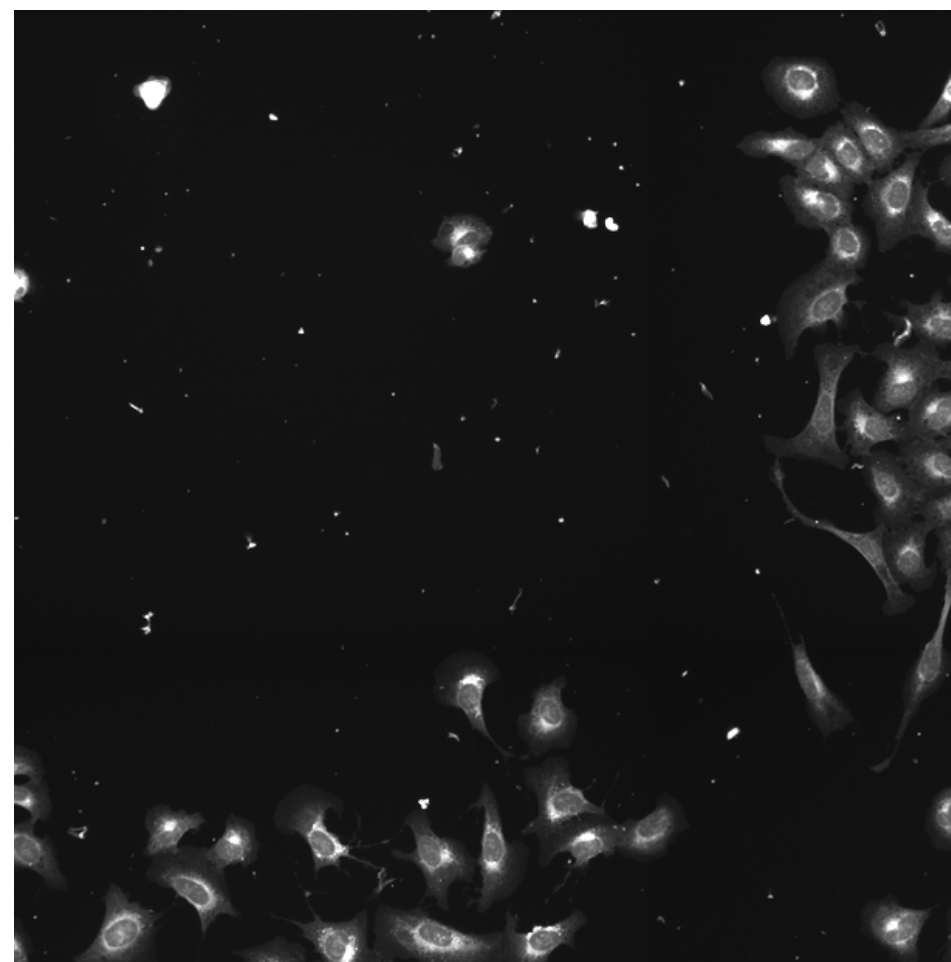
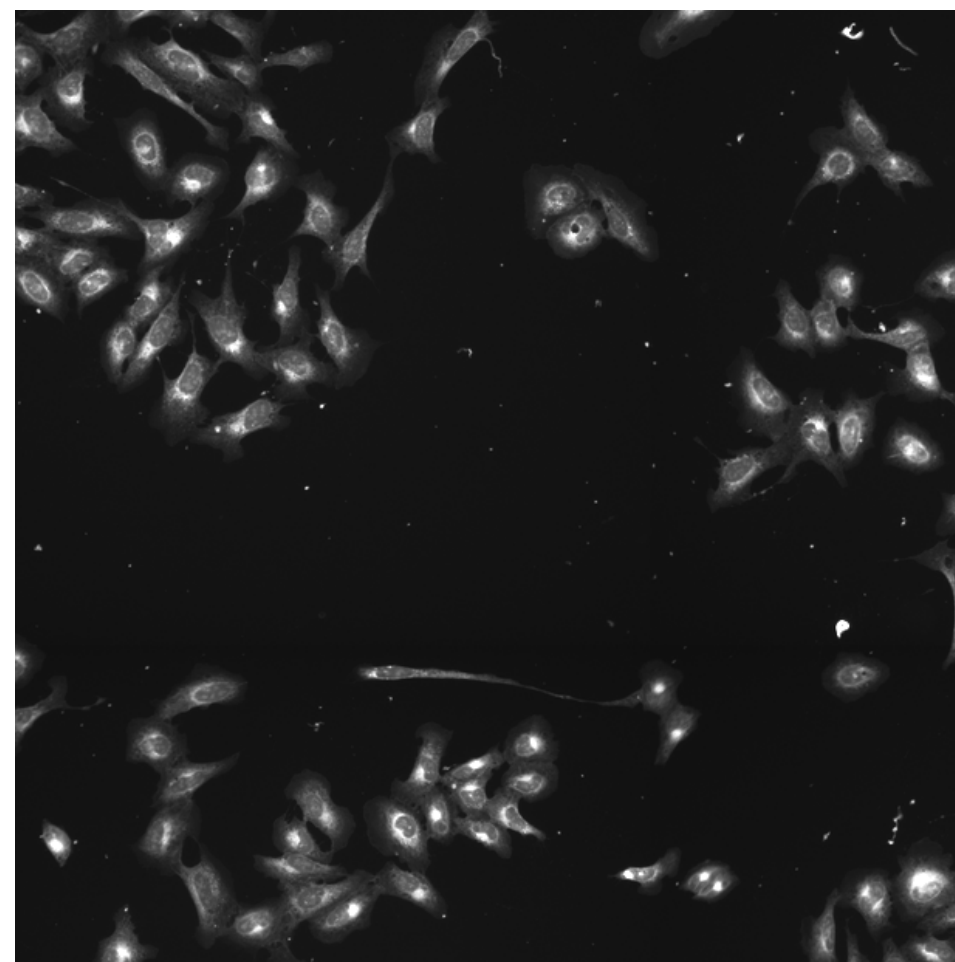
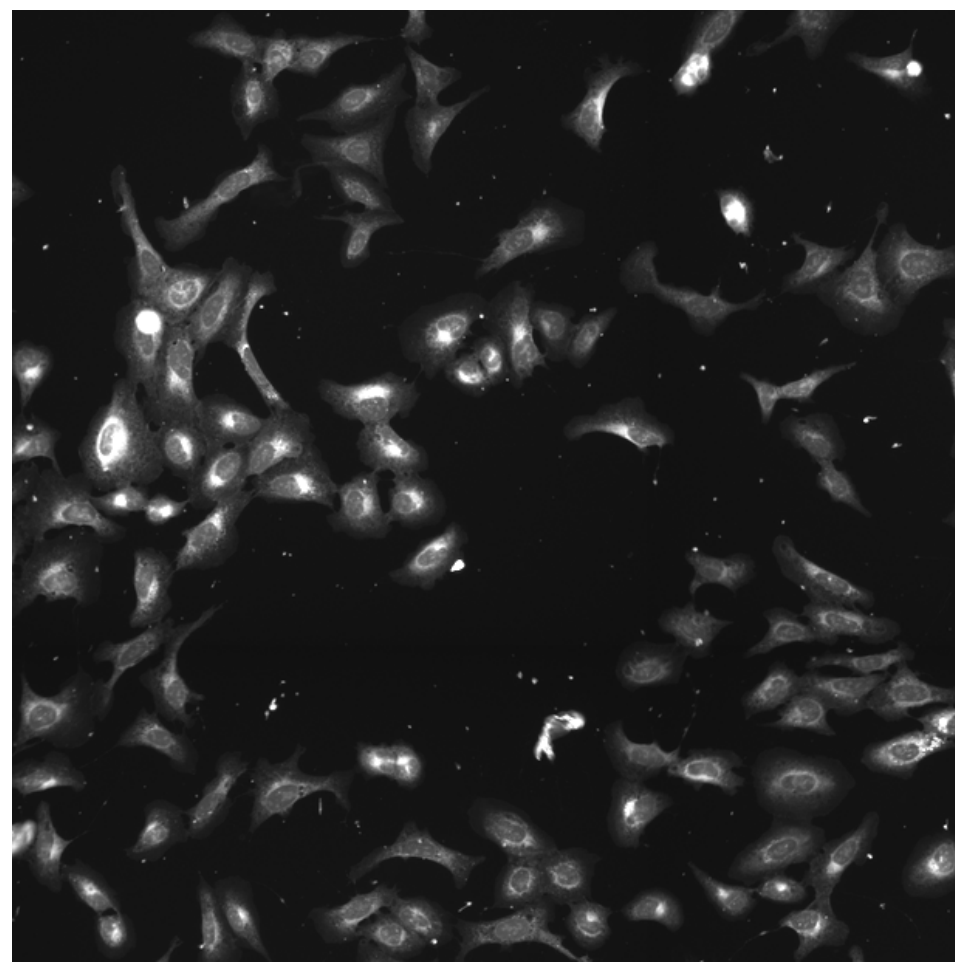
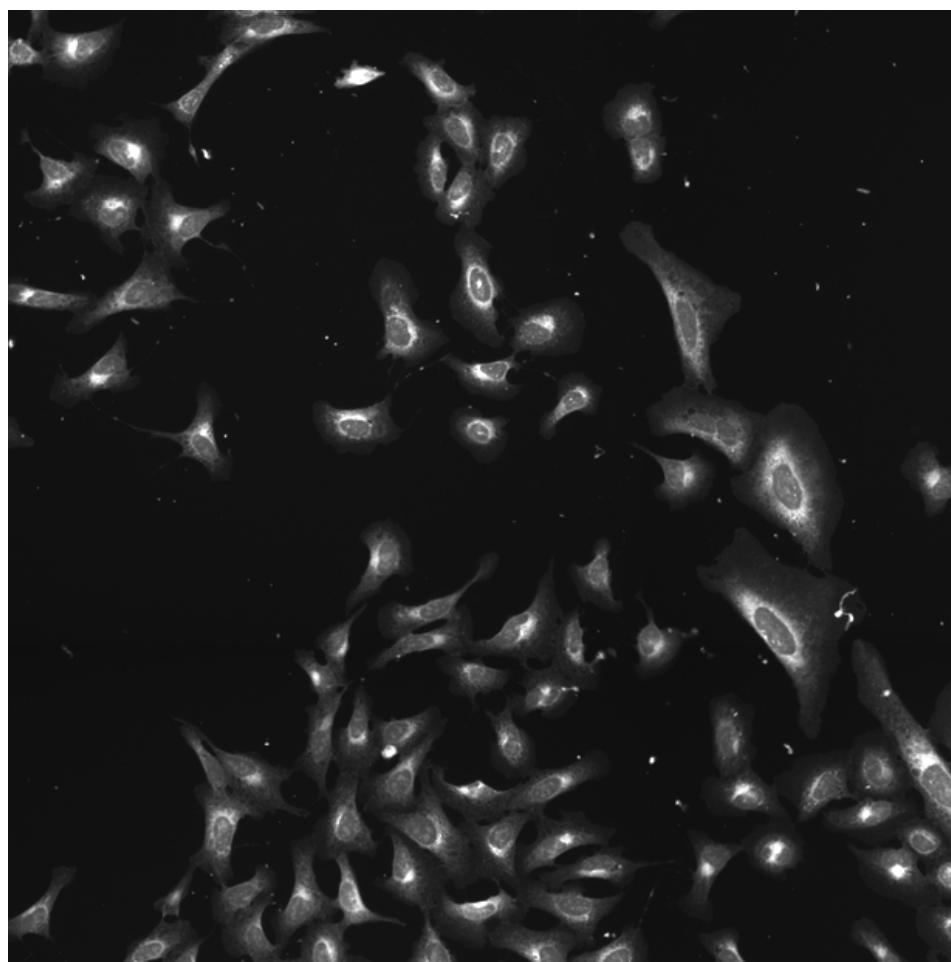
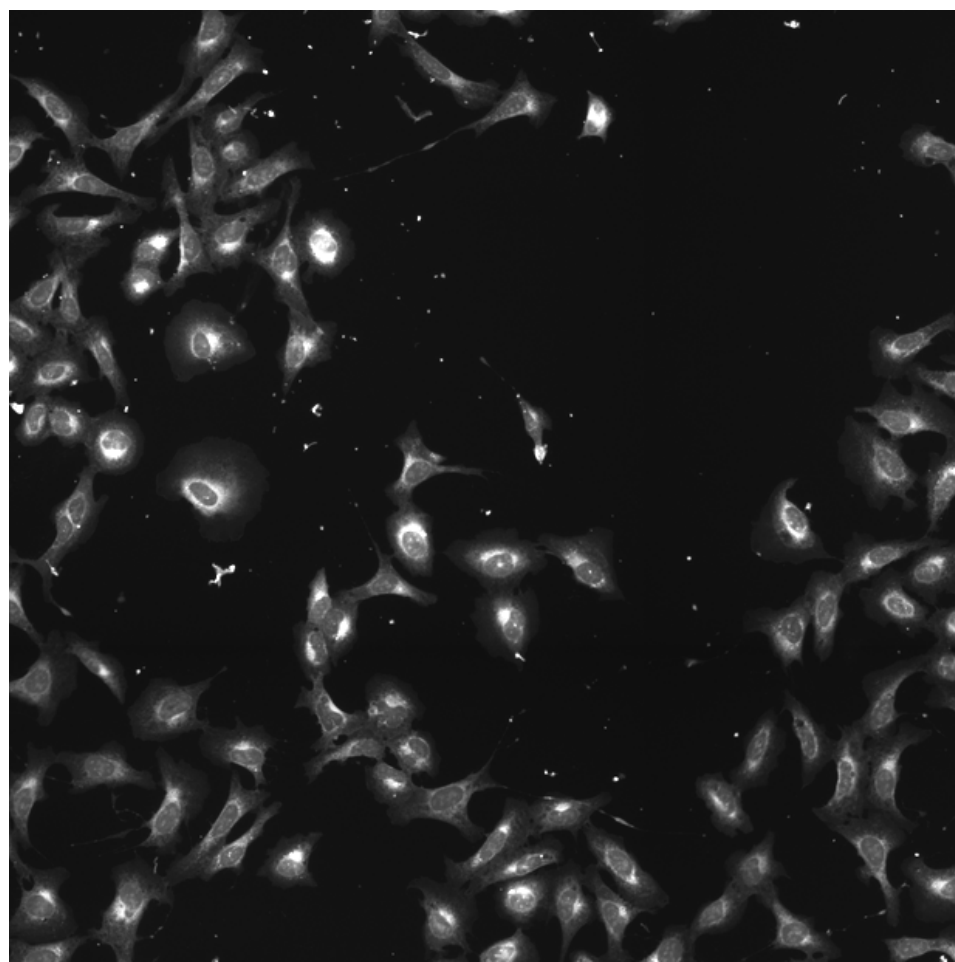
EIF4EBP1.WT (41755)

EIF4EBP1.WT (41756)

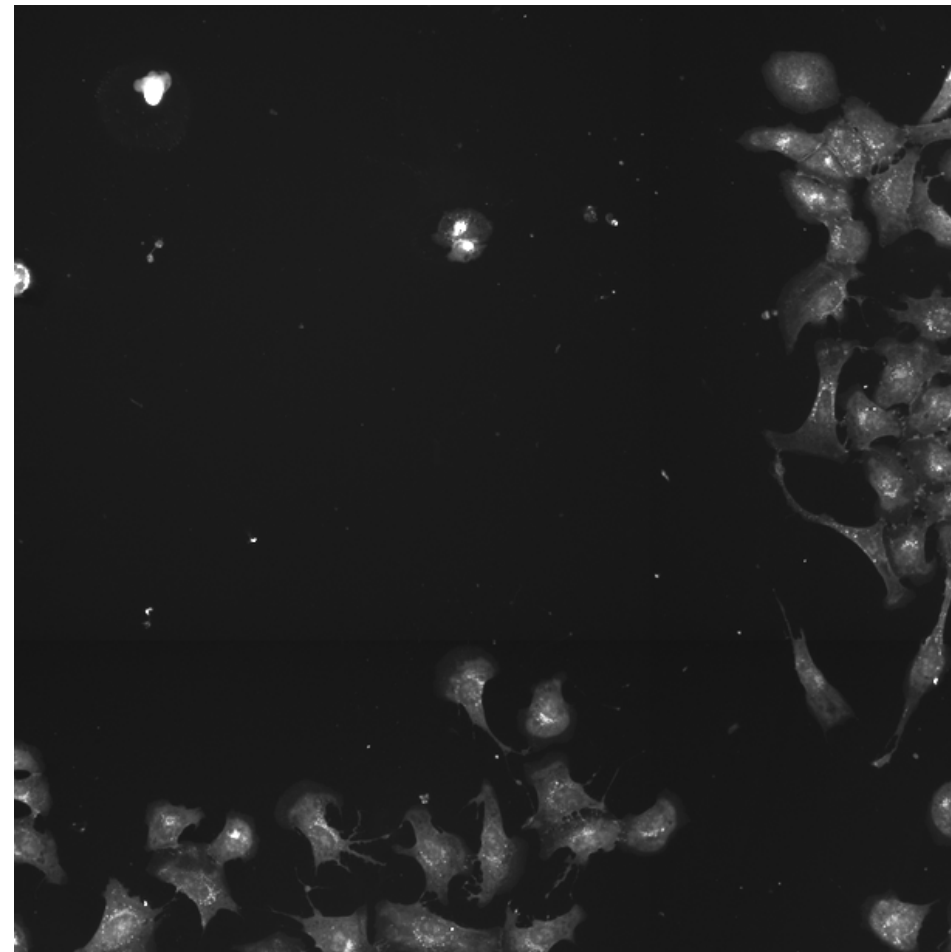
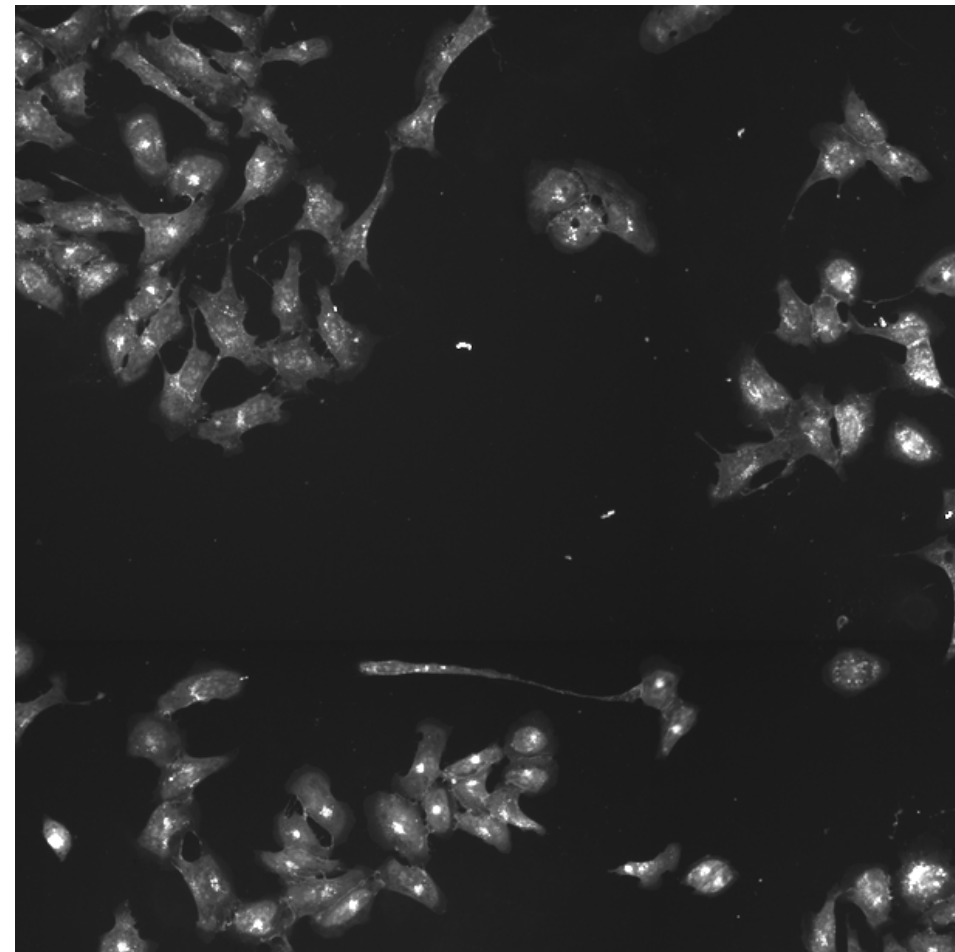
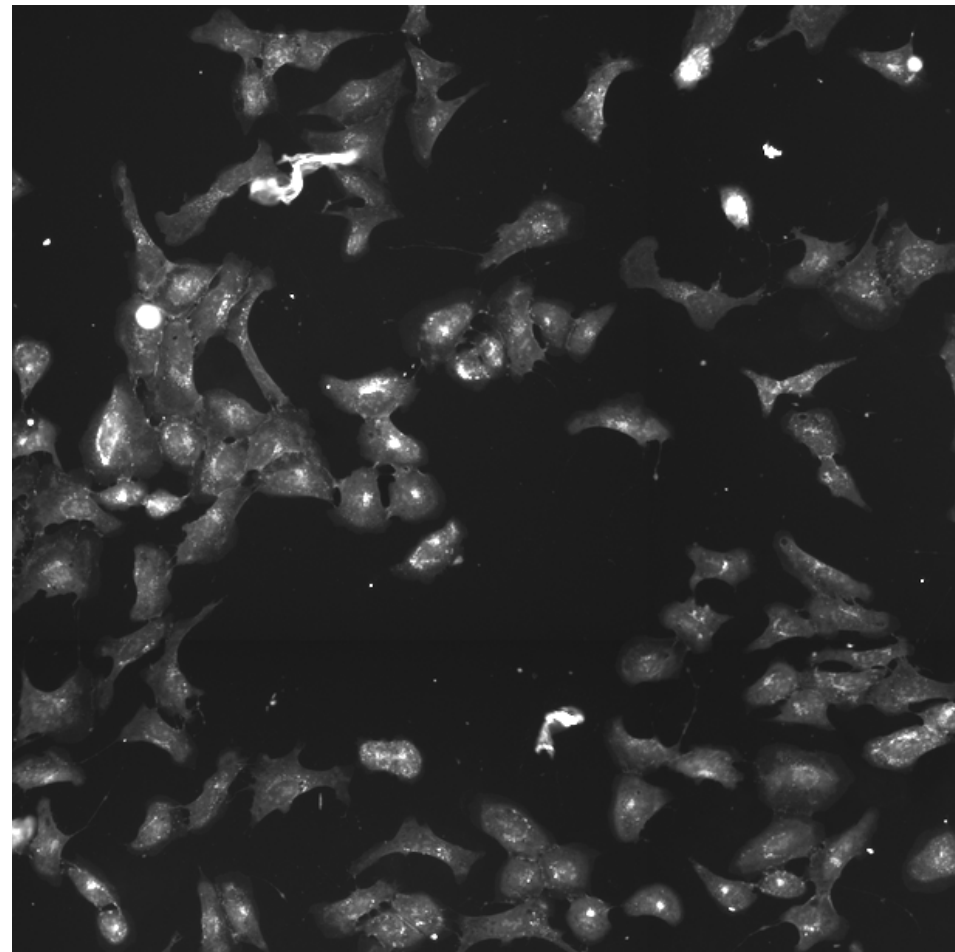
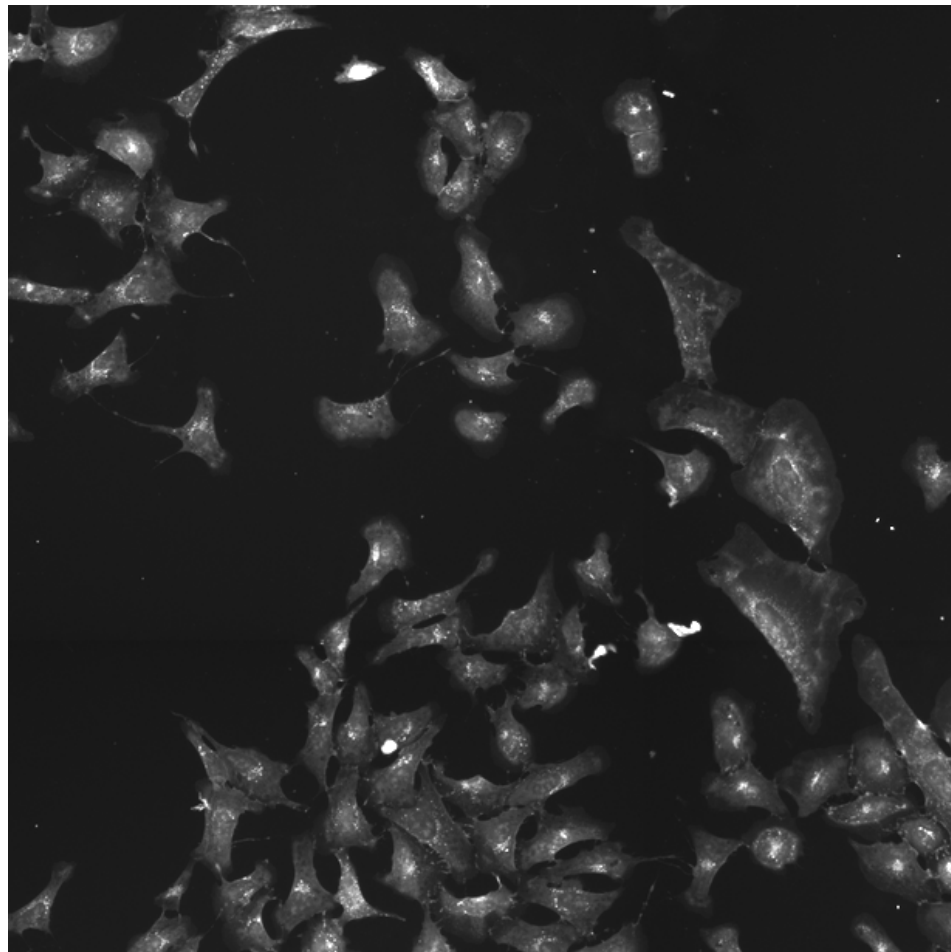
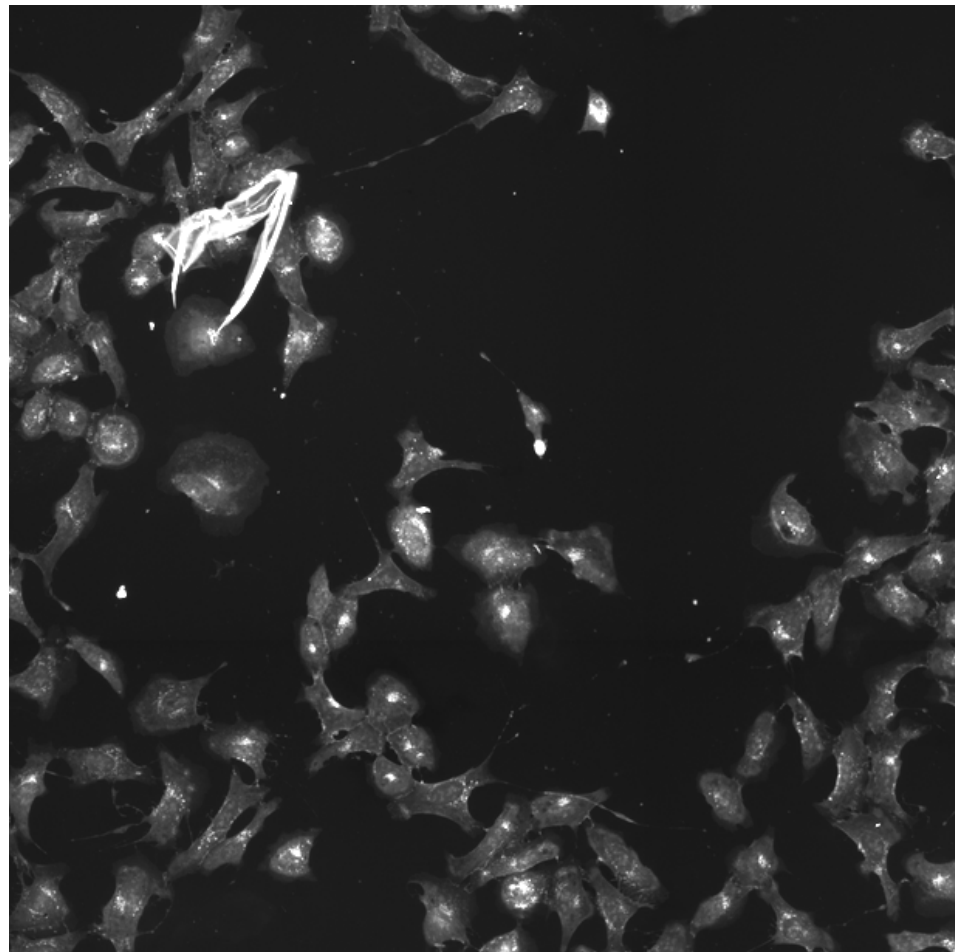
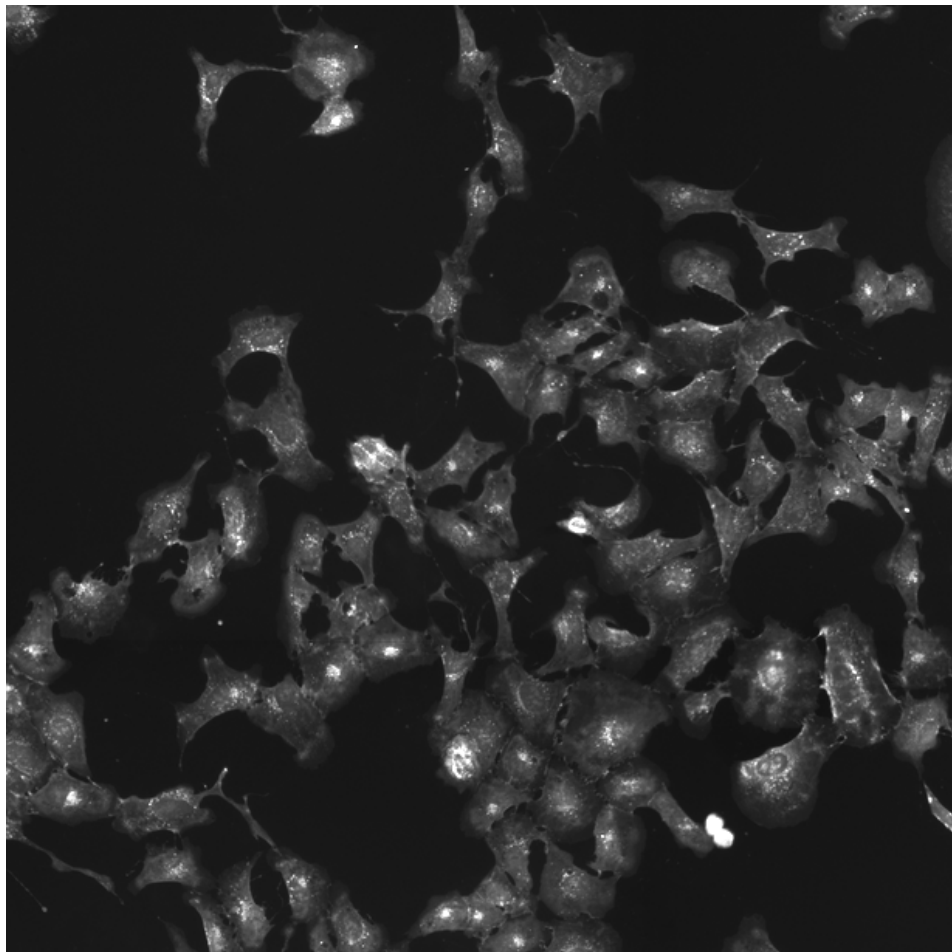
EIF4EBP1.WT (41757)

EIF4EBP1.WT (41754)

ER



AGP



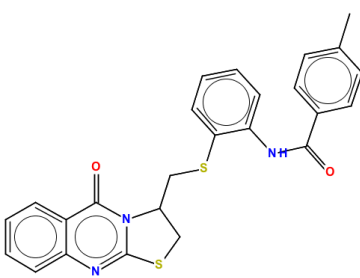
Compound IDs and common names (where available); blue/red colored box means the matching compound is positively/negatively correlated with the cluster	Chemical structure	Mean pairwise replicates correlation of the compound signature (95th DMSO replicate correlation is 0.52)	Correlation between compound the gene	Compound rank when scored against the gene using L1000 profiling	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 the gene relative to the untreated samples	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	Number of PubChem assays in which the compound was tested; assays in which the compound was active are itemized
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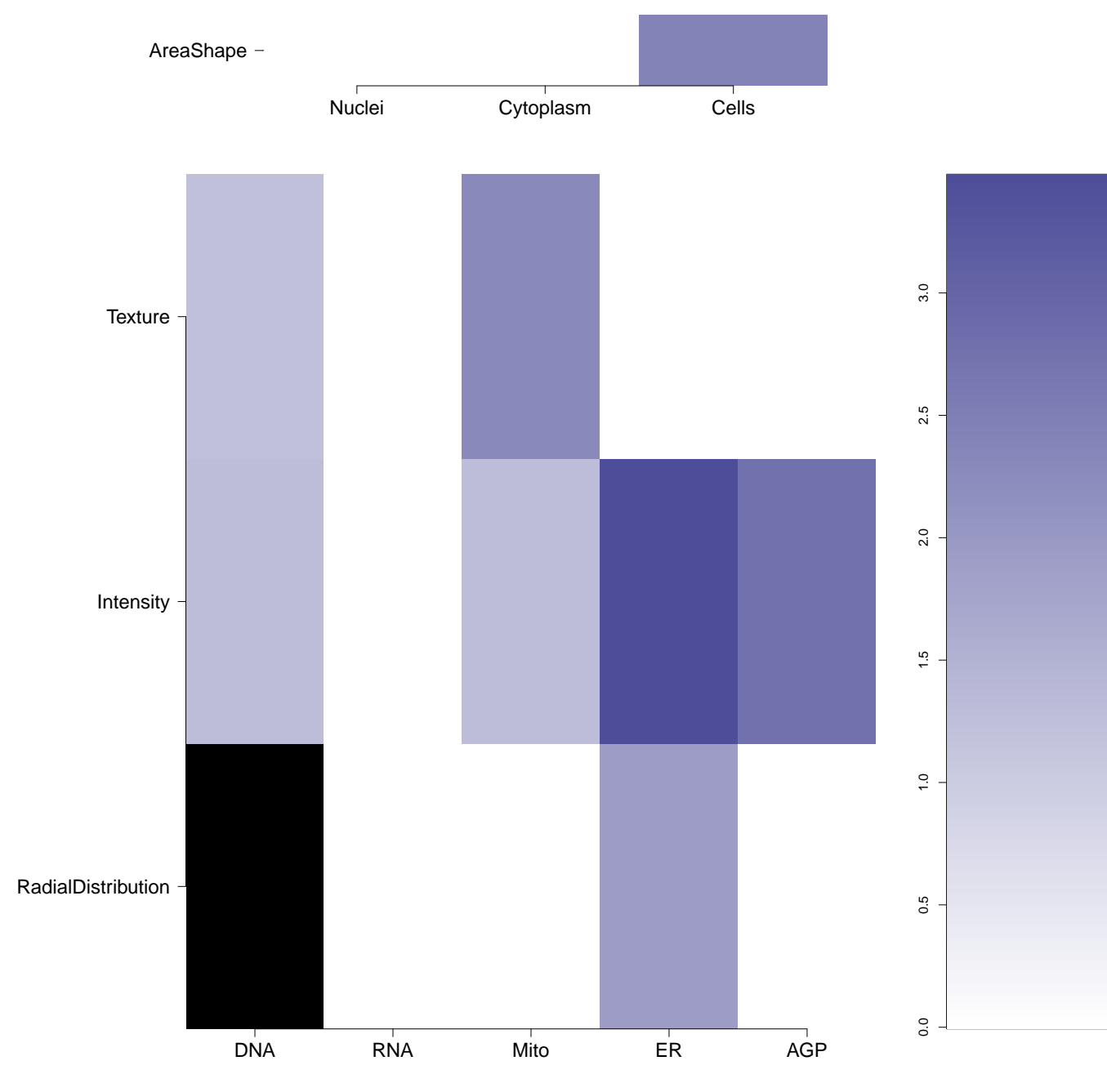
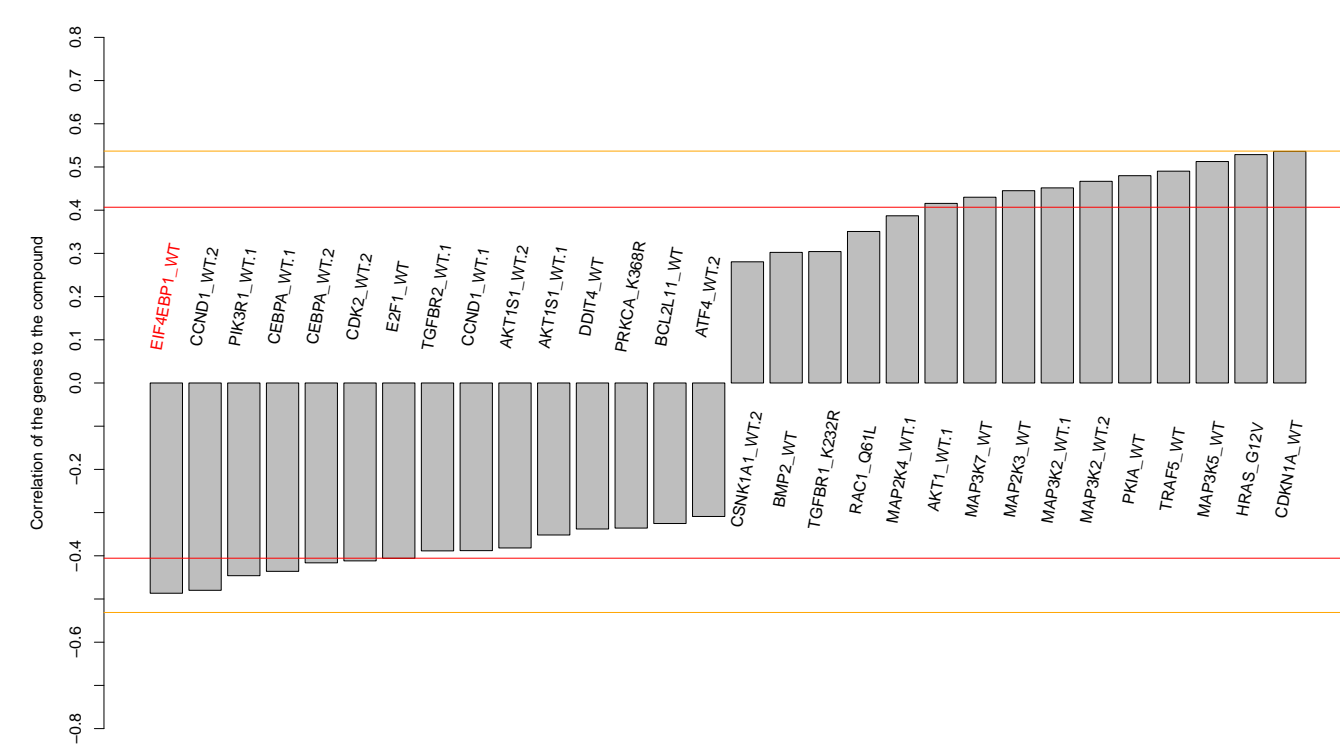
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0.66 (in 4 replicates)

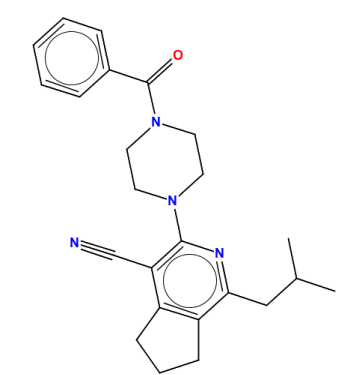
-0.49

0.302



- Total number of assays tested in: 666. Active in the following assays:
- Non-Nucleoside Inhibitor of Measles Virus RNA-Dependent RNA Polymerase Complex Activity HTS Single Point (MLSMR Library) (AID 841)
  - uHTS of Mcl-1/Bcl interaction inhibitors (AID 1021)
  - Leishmania major promastigote HTS (AID 1063)
  - Anti-Viral Drugs Against Arbovirus Infections, a Primary Screen (AID 1251)
  - Luminescence-based counterscreen assay for KLF5 inhibitors: cell-based high throughput screening assay to identify cytotoxic compounds using the IEC-6 intestinal epithelial cell line. (AID 1825)
  - Luminescence-based confirmation cell-based assay for cytotoxic compounds using the IEC-6 intestinal epithelial cell line. (AID 1907)
  - A Cell Based HTS Approach for the Discovery of New Inhibitors of Respiratory syncytial virus (RSV) (AID 2391)
  - An HTS Cytotoxicity Screen to evaluate New Inhibitors of Respiratory Syncytial Virus (RSV) (AID 2410)
  - qHTS for identification of Inhibitors of Mdm2/MdmX interaction in luminescent format. (AID 485346)
  - Single concentration confirmation of uHTS for Inhibitors of Mdm2/MdmX interaction in luminescent format. (AID 489028)
  - Single concentration confirmation of inhibitors of Mdm2/MdmX interaction using a Full-Length Luciferase Counterscreen assay (AID 504607)
  - Single concentration confirmation of inhibitors of Mdm2/MdmX interaction using a Breal/Bardil BiLC Counterscreen assay. (AID 504668)
  - Confirmation screen for delayed death inhibitors of the malarial parasite plastid, 96 hour incubation (AID 504848)
  - Confirmation screen for delayed death inhibitors of the malarial parasite plastid, 48 hour incubation (AID 504850)
  - qHTS for Inhibitors of binding or entry into cells for Lassa Virus (AID 540256)
  - qHTS for inhibitors of binding or entry into cells for Marburg Virus (AID 540270)
  - A Cell-Based Confirmatory Screen for Compounds that Inhibit VEEV, TC-83 (AID 588727)

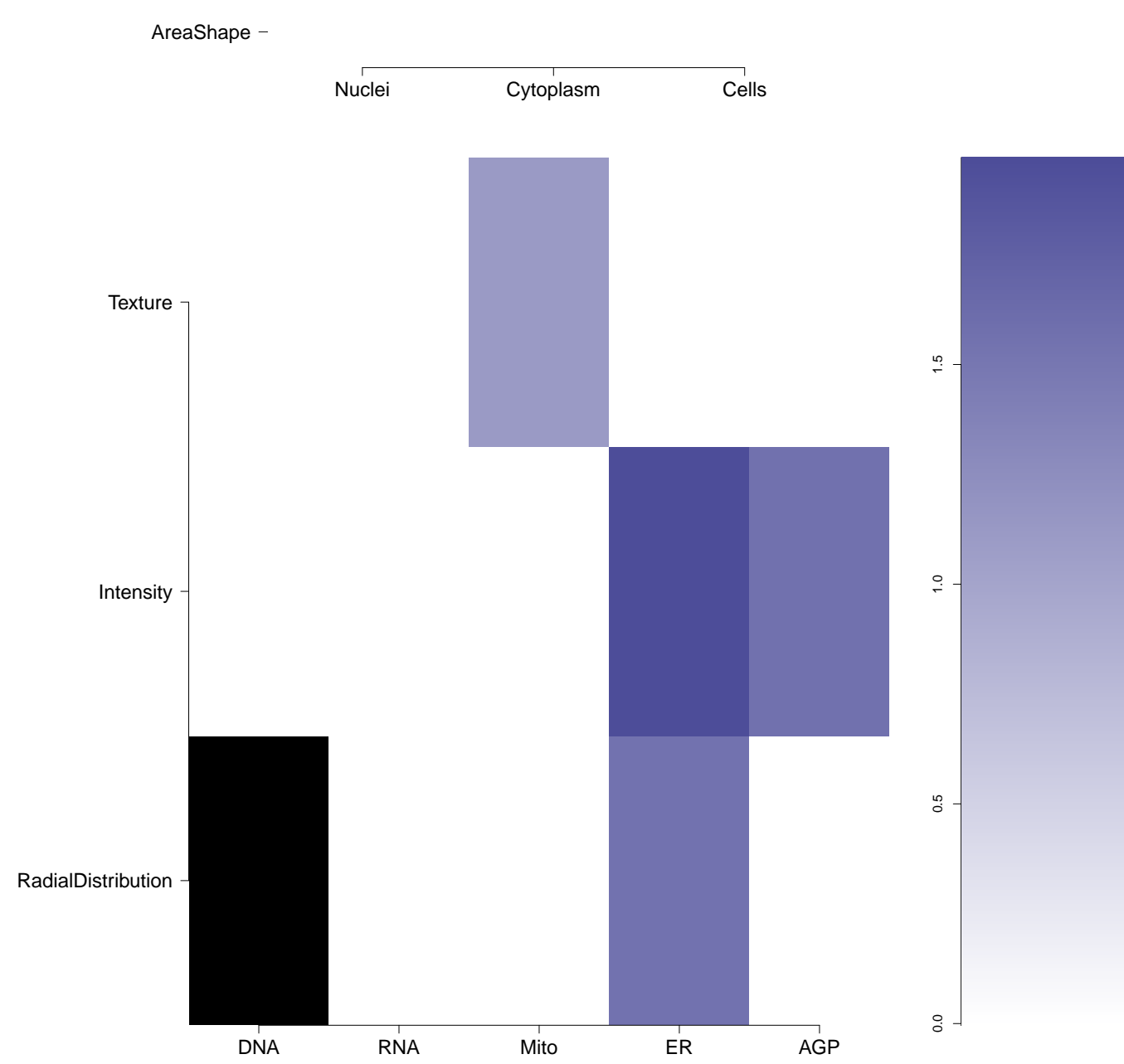
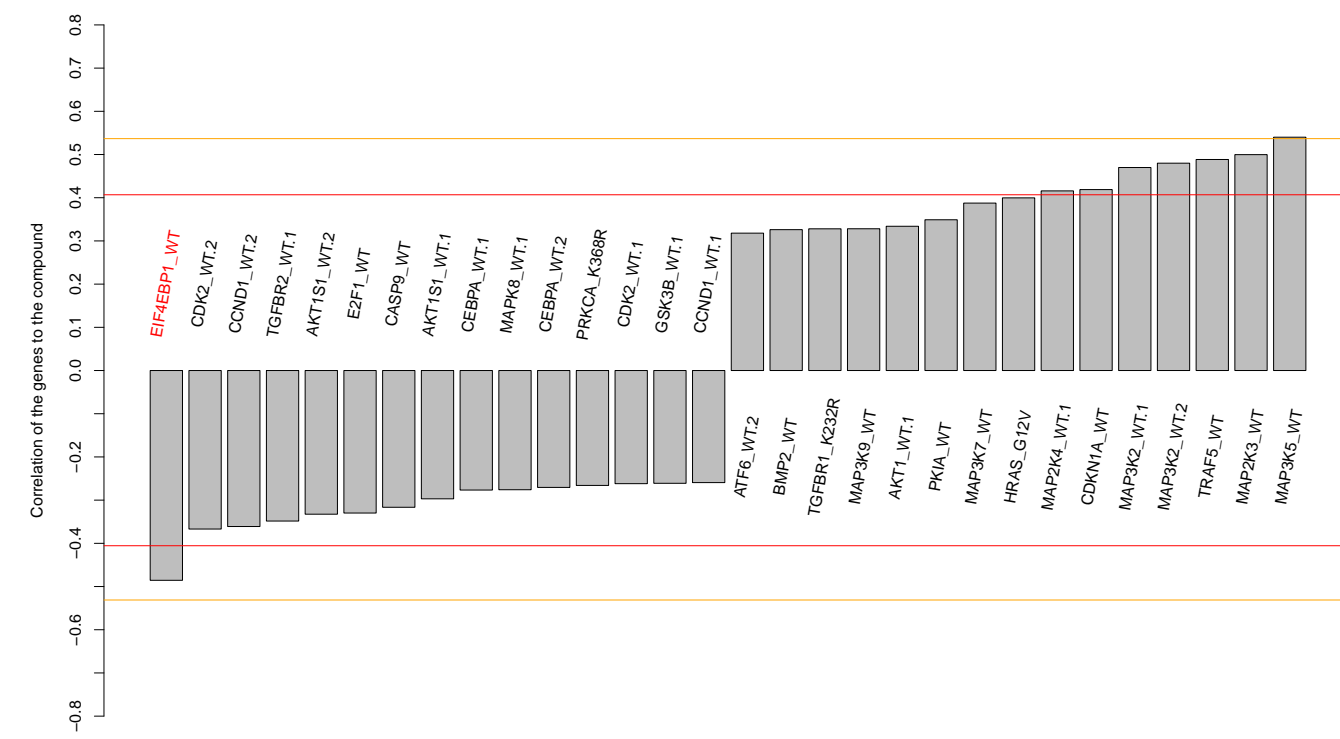
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0.60 (in 3 replicates)

-0.49

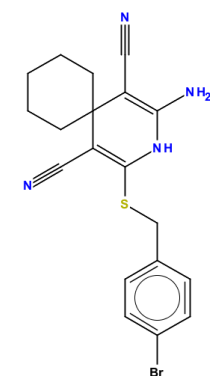
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- Total number of assays tested in: 683. Active in the following assays:
- CYP2C9 Assay (AID 777)
  - CYP2C19 Assay (AID 778)
  - Chemical Genetic Screen to Identify Inhibitors of Mitochondrial Fusion - Confirmatory Screen (AID 1361)
  - Chemical Genetic Screen to Identify Inhibitors of Mitochondrial Fusion - Primary Screen (AID 1362)
  - qHTS Assay for Antagonists of the Neuropeptide S Receptor: cAMP Signal Transduction (AID 1461)
  - Cytochrome panel assay with activity outcomes (AID 1851)
  - Primary cell-based high-throughput screening assay for identification of compounds that inhibit KCNQ2 potassium channels (AID 2156)
  - Primary cell-based high-throughput screening assay for identification of compounds that allosterically potentiate transient receptor potential cation channel C4 (TRPC4) (AID 2227)
  - VP16 counterscreen qHTS for inhibitors of ROR gamma transcriptional activity (AID 2546)
  - qHTS for inhibitors of ROR gamma transcriptional activity (AID 2551)
  - HTS for small molecule inhibitors of CHOP to regulate the unfolded protein response to ER stress (AID 2732)
  - Nrf2 qHTS screen for inhibitors (AID 504444)
  - qHTS for Inhibitors of binding or entry into cells for Lassa Virus (AID 540256)
  - qHTS Assay for Inhibitors of Hepatitis C Virus (HCV) (AID 651820)
  - qHTS Assay for Activators of ClpP (AID 651965)
  - qHTS of TDP-43 Inhibitors (AID 652104)



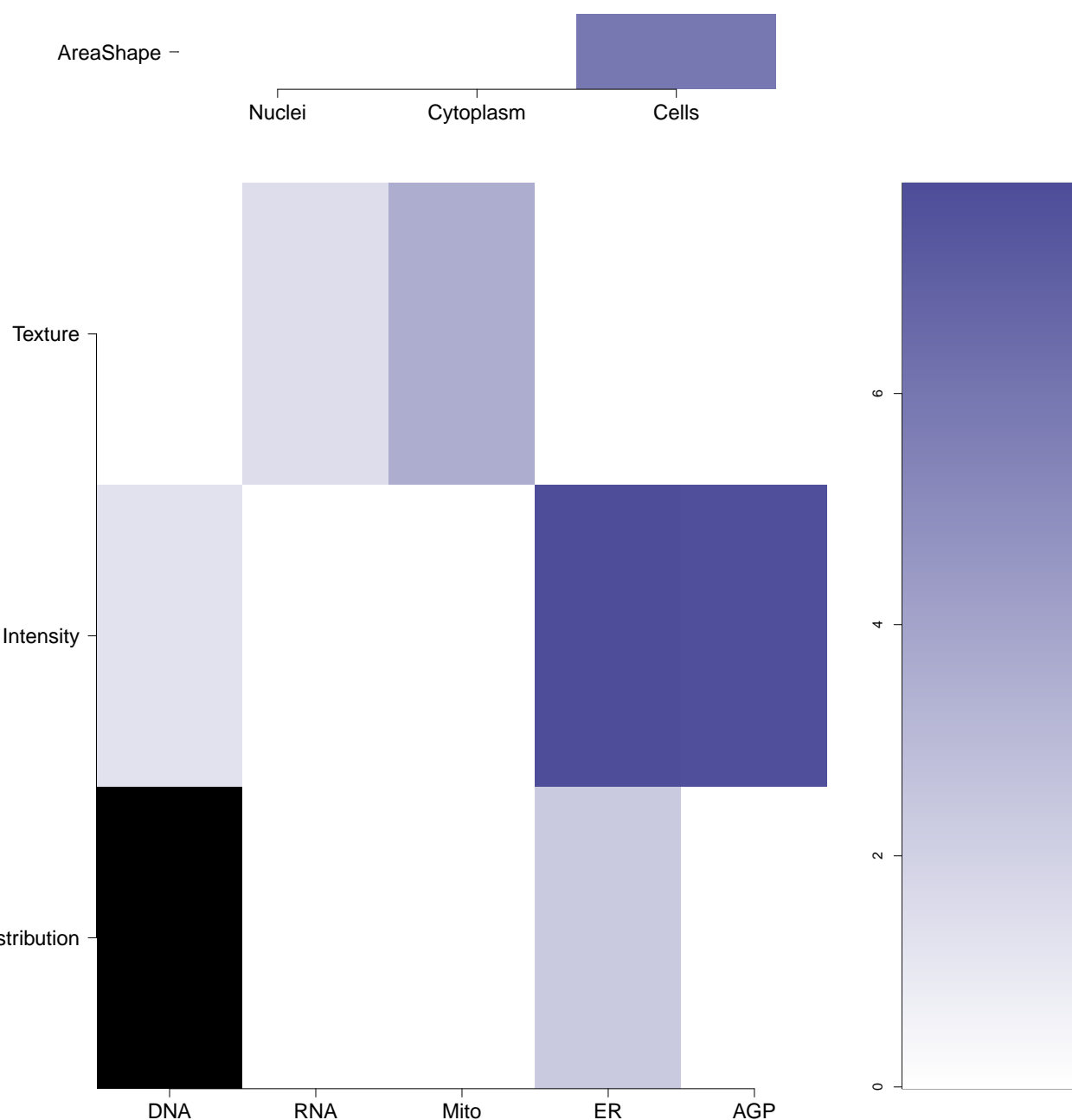
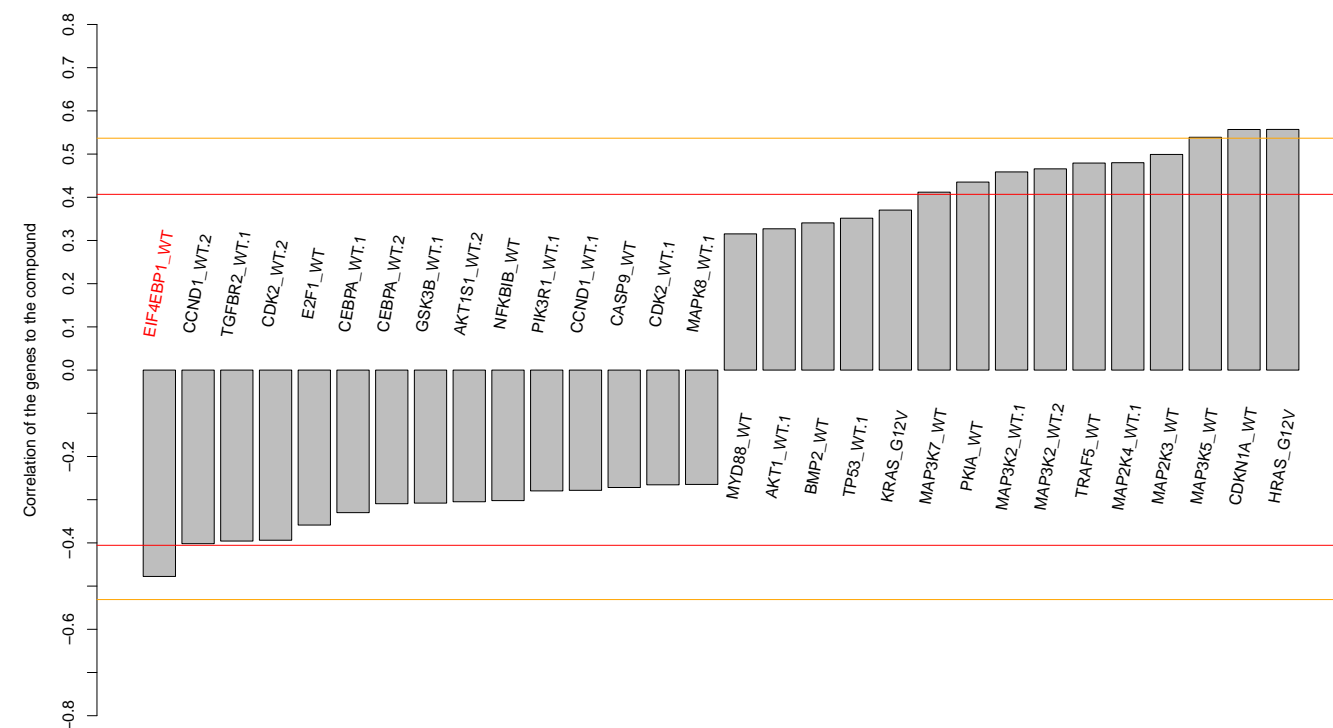
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0.52 (in 4 replicates)

-0.48

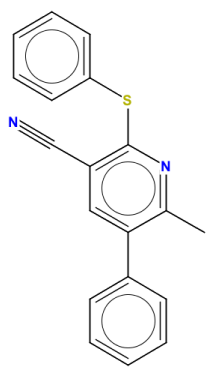
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total number of assays tested in: 674. Active in the following assays:

- Leishmania major promastigote HTS (AID 1063)
- Chemical Genetic Screen to Identify Inhibitors of Mitochondrial Fusion - Primary Screen (AID 1362)
- CounterScreen qHTS for Inhibitors of Tau Fibril Formation, Fluorescence Polarization (AID 1463)
- Luminescence Cell-Based/Microorganism Primary HTS to Identify Inhibitors of T.Cruzi Replication (AID 1885)
- High throughput discovery of novel modulators of ROMK K<sup>+</sup> channel activity: Primary Screen (AID 1918)
- Multiplex HTS Screen of TOR pathway components in *Saccharomyces cerevisiae*-specifically, MEPP2.MLPCN. (AID 2016)
- Multiplex HTS Screen of TOR pathway GTP-fusion proteins in *Saccharomyces cerevisiae*-specifically, RPL19A.MLPCN. (AID 2025)
- Luminescence Cell-Based/Microorganism Dose Confirmation HTS to Identify Inhibitors of T.Cruzi Replication. (AID 2044)
- Primary cell-based high-throughput screening assay for identification of compounds that inhibit KCNQ1 potassium channels (AID 2642)
- HTS for small molecule inhibitors of CHOP to regulate the unfolded protein response to ER stress (AID 2732)
- Fluorescence-based counter-screen for orexin 1 receptor antagonists in *Saccharomyces cerevisiae* to identify antagonists of the parental CHO cell line (AID 463079)
- High throughput fluorescence intensity-based biochemical assay to screen for small molecule inhibitors of Furin conducted by the Pittsburgh Molecular Library Screening Center. (AID 463115)
- Primary cell-based screen for identification of compounds that inhibit the two-pore domain potassium channel KCNK9 (AID 488922)
- Fluorescence polarization-based primary bioassay for high throughput screening assay to identify inhibitors of human platelet-activating factor acetylcholinesterase 1b, catalytic subunit 2 (PAPAHI2) (AID 492953)
- Fluorescence polarization-based biochemical high throughput confirmation assay for inhibitors of human platelet-activating factor acetylcholinesterase 1b, catalytic subunit 2 (PAPAHI2) (AID 493634)
- Validation (re-confirmation) assay for identification of compounds that inhibit KCNQ1 potassium channels (AID 585835)
- Counter screen assay for the parental CHO cells for identification of compounds that inhibit KCNQ1 potassium channels (AID 588366)
- Primary cell-based high-throughput screening for identification of compounds that inhibit bulk calcium-activated chloride channels (TMEM16A) (AID 588511)
- Full cell counter-screen for antagonists of the human M1 muscarinic receptor (CHRM1): Fluorescence-based cell-based high throughput screening assay to identify novel selective inhibitors and assay artifacts using the parental CHOx1 cell line (AID 602250)
- iHTS for antagonists of the Thyroid Stimulation Hormone Receptor: Hit Validation in Primary Screen (AID 602292)
- Primary cell-based screen for identification of compounds that inhibit the two-pore domain potassium channel KCNK3 (AID 602410)
- iHTS identification of small molecule inhibitors of the mitochondrial permeability transition pore via an absorbance assay (AID 602449)
- iHTS of GLP-1 Receptor Inverse Agonists (Inhibition Mode) (AID 62417)
- Single concentration confirmation of vHTS inhibitors of the mitochondrial permeability transition pore via a fluorescent based assay (AID 624504)
- Confirmation assay for identification of compounds that inhibit the two-pore domain potassium channel KCNK3 (Primary Screening) (AID 651638)
- Specificity screen against KCNQ1 for identification of compounds that inhibit KCNQ1 potassium channels (AID 651746)
- Counter screen assay for identification of compounds that inhibit the two-pore domain potassium channel KCNK3 (Primary Screening) in non-induced KCNK3 cells (AID 651747)
- Specificity screen against KCNQ1/KCNK1 for identification of compounds that inhibit KCNQ1 potassium channels (AID 652147)
- qPRET-based biochemical primary high throughput screening assay to identify exosite inhibitors of ADAM10. (AID 720582)

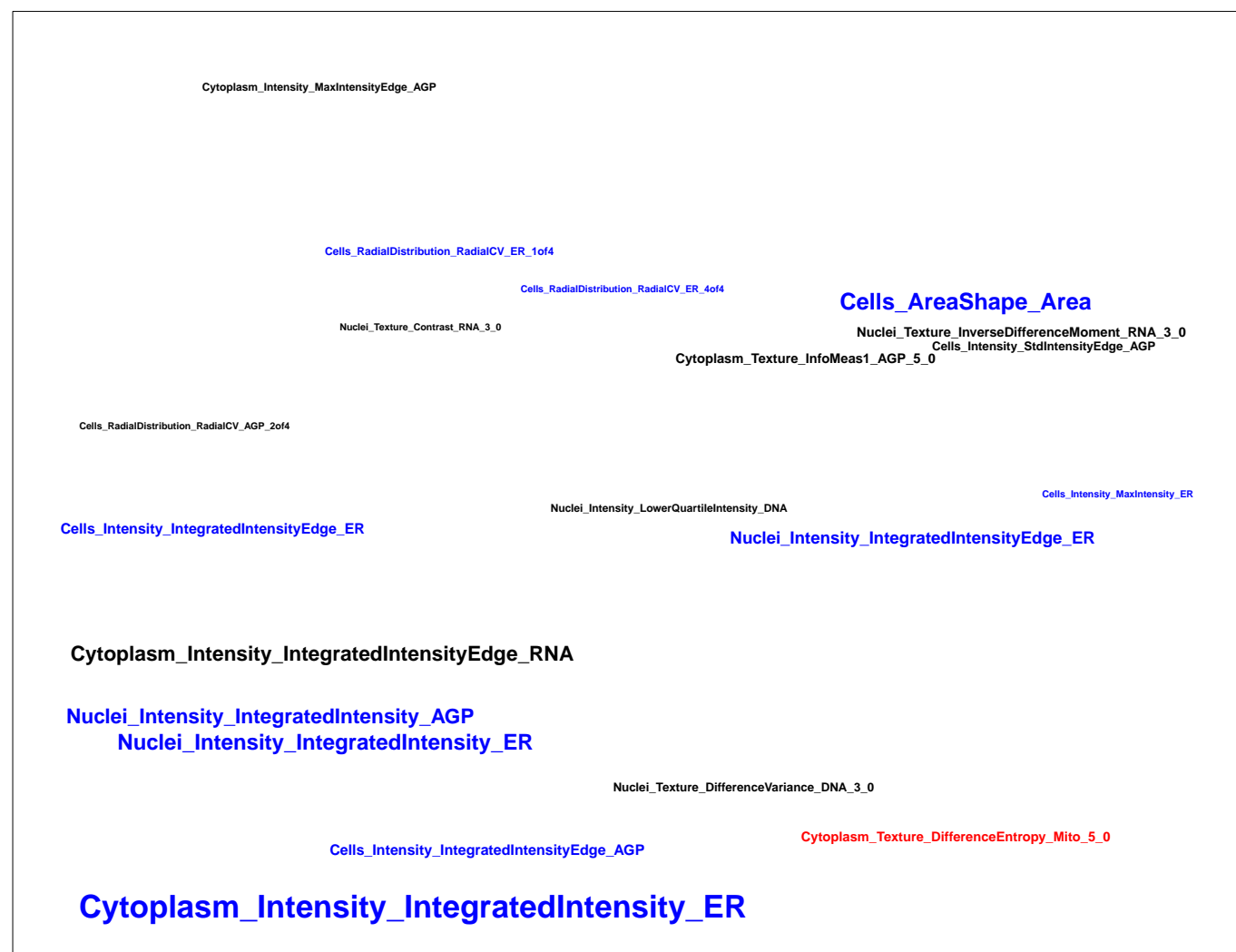
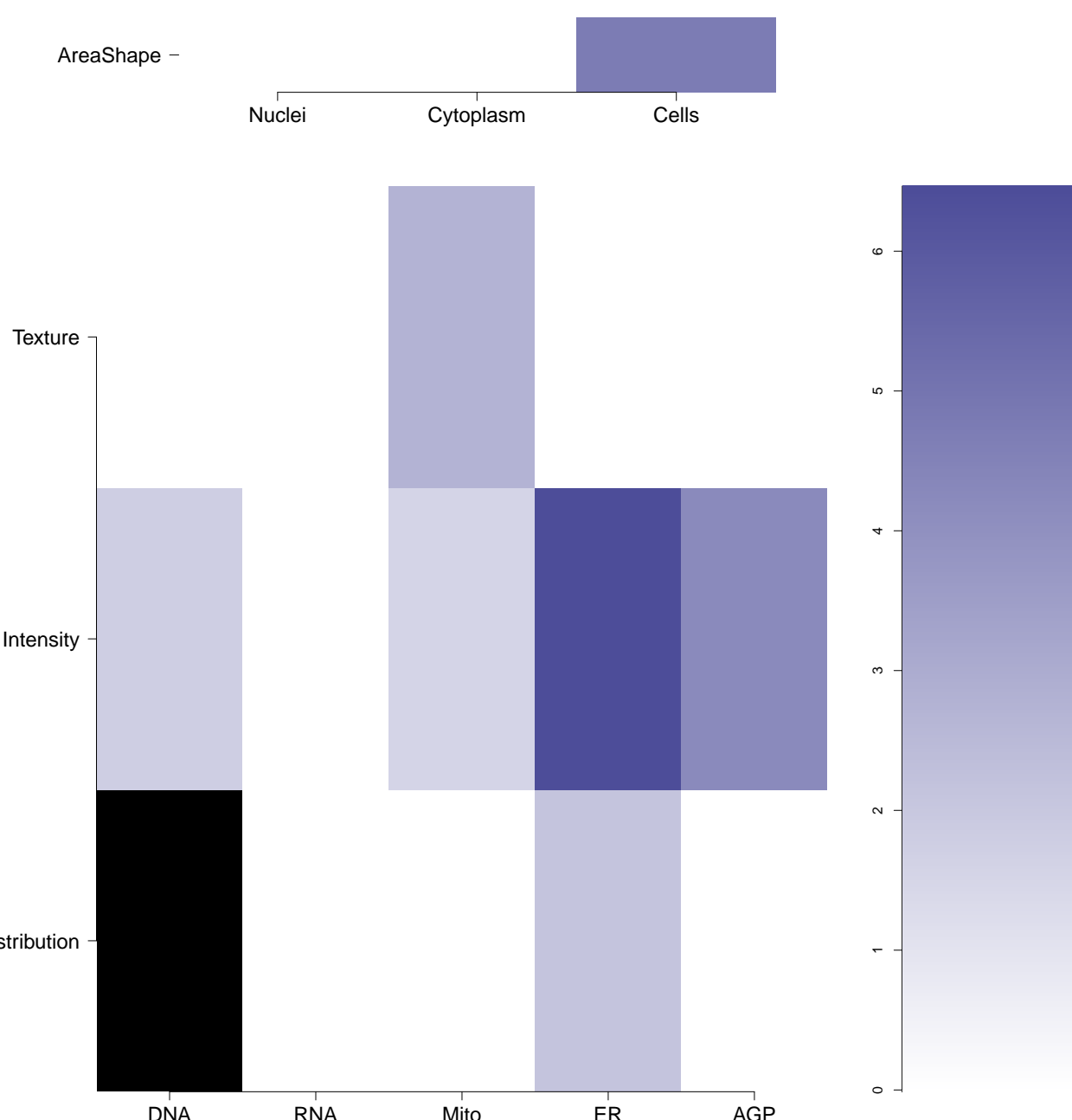
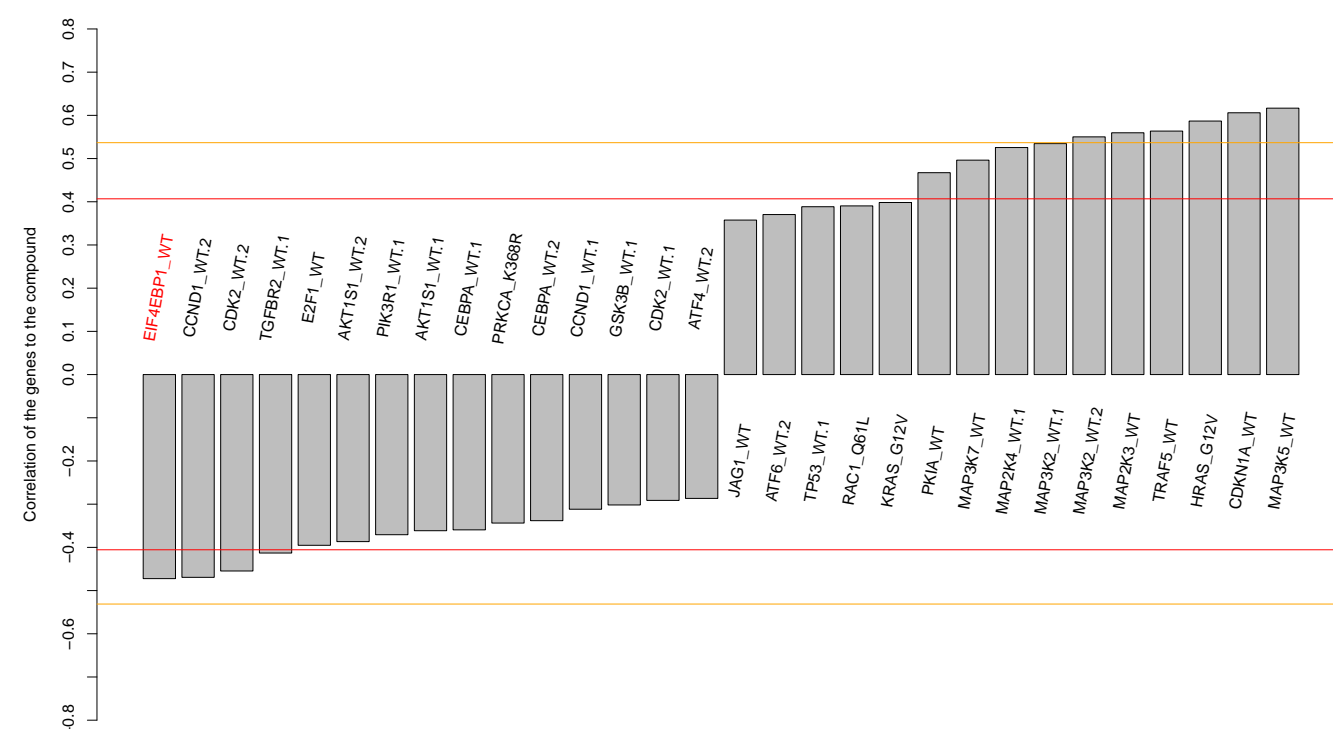
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ZINC1388577  
PubChem CID : 1476369



0.71 (in 4 replicates)


-0.47

NA



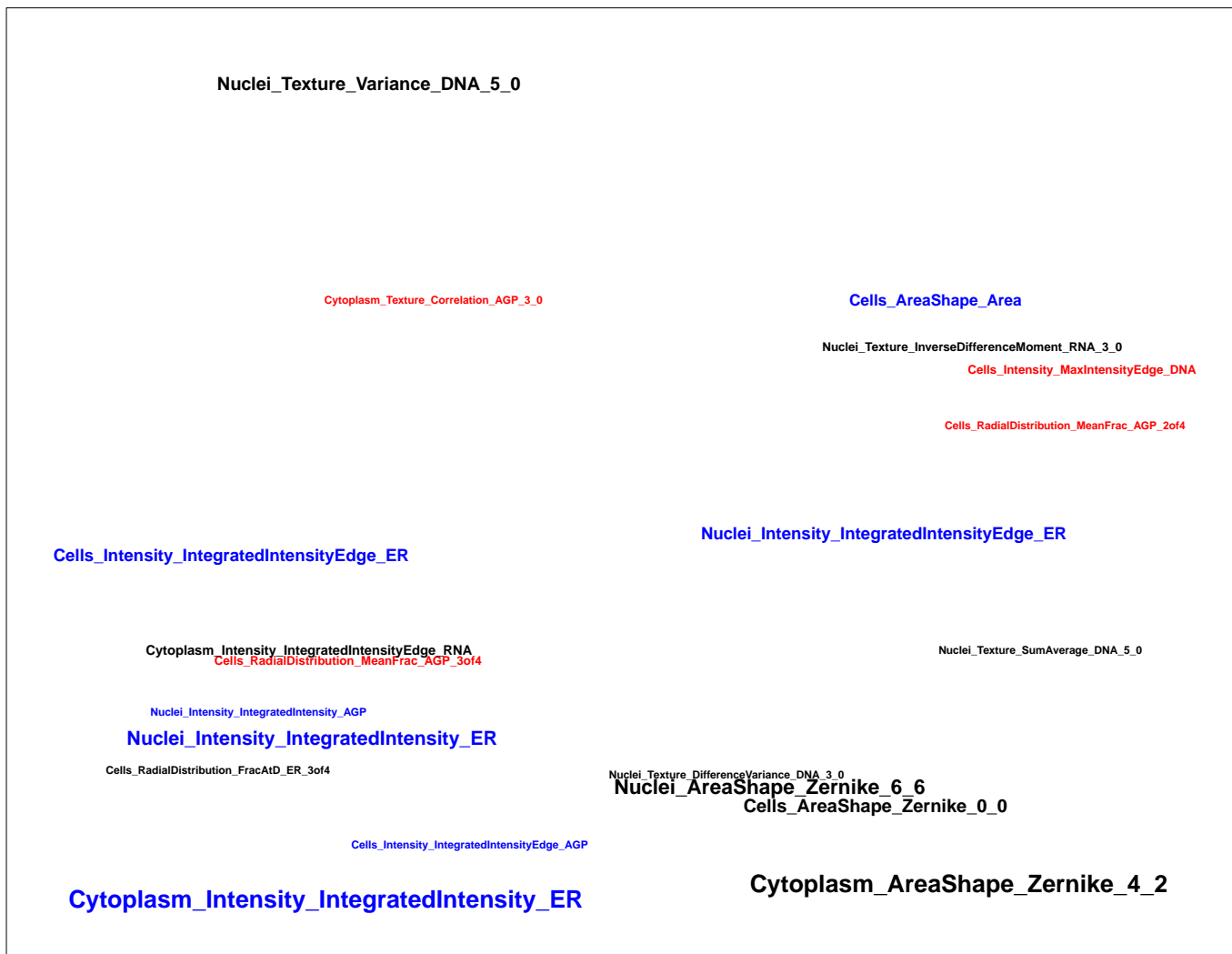
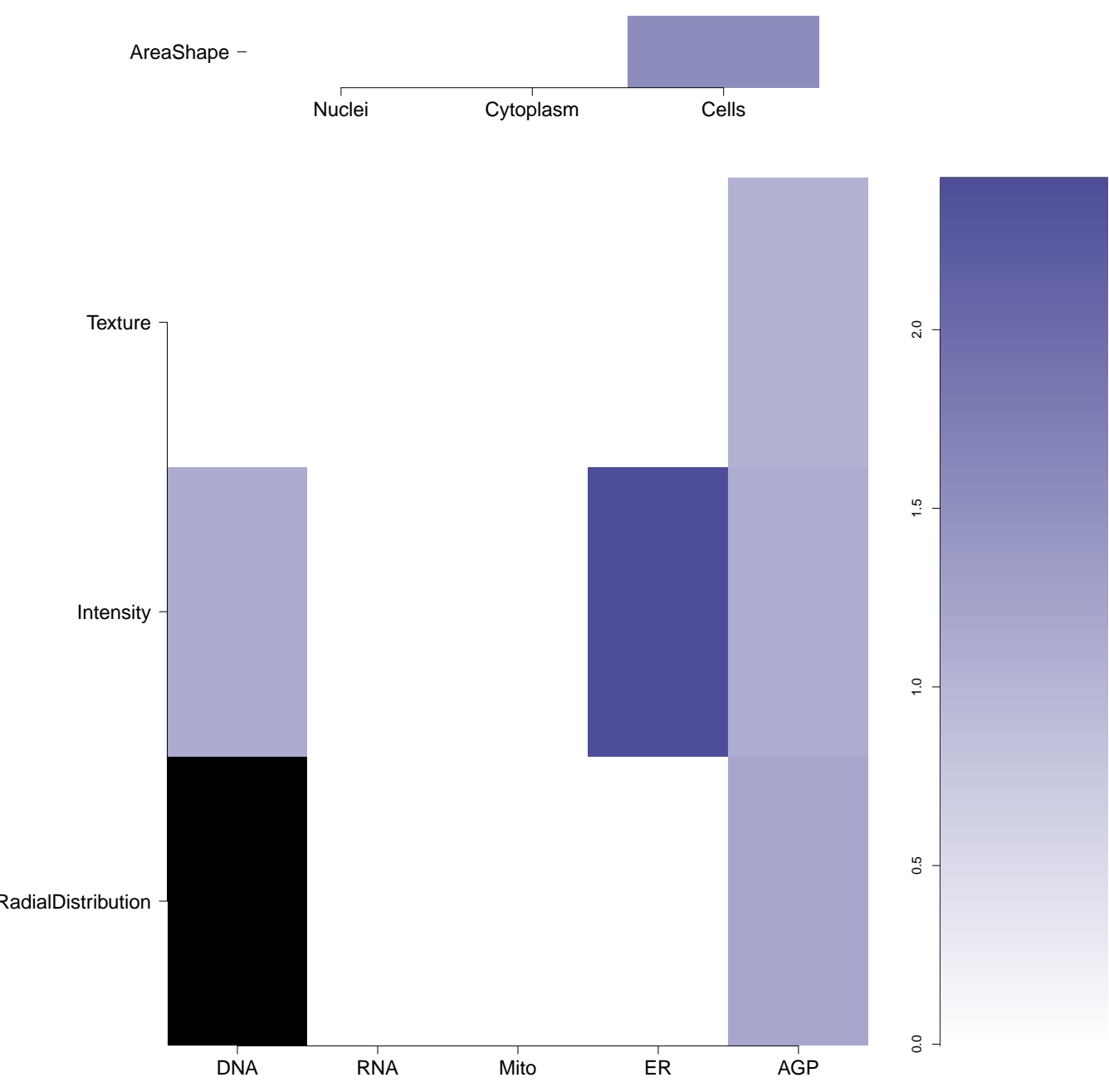
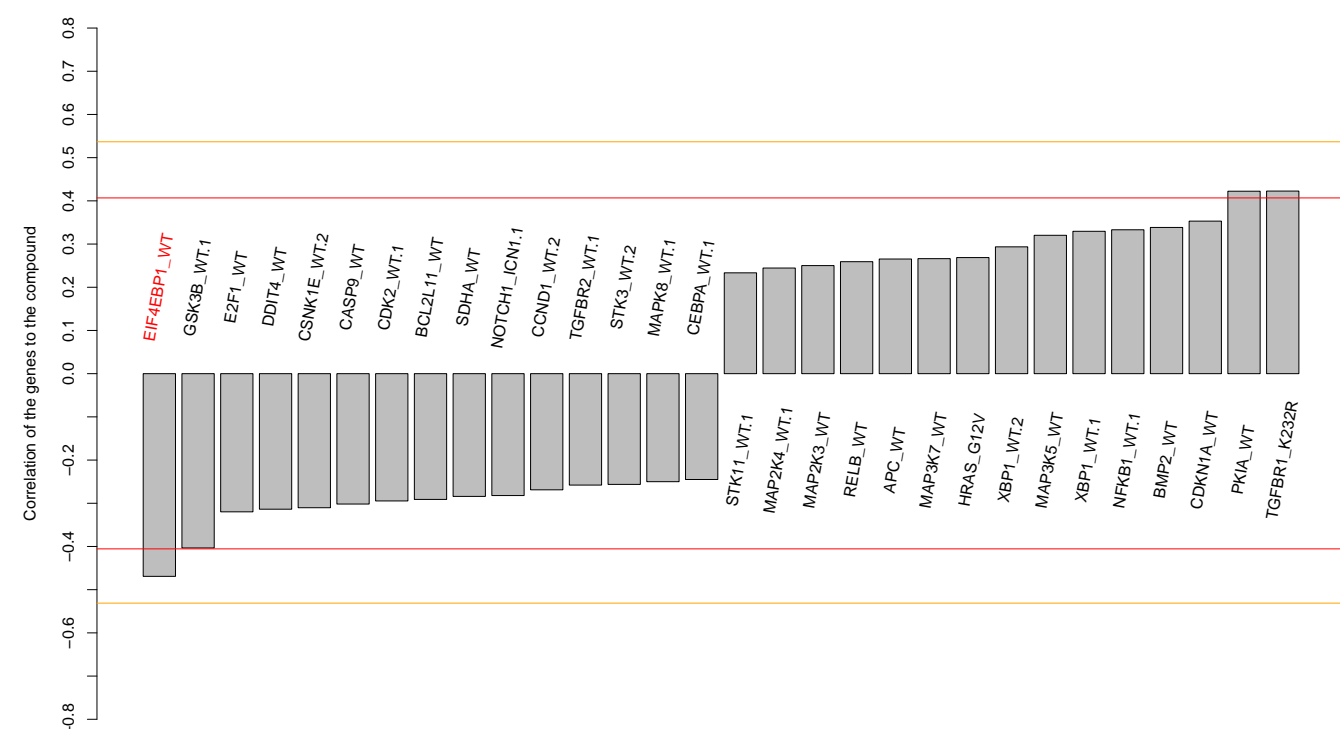
- Total number of assays tested in: 609. Active in the following assays:
  - CYP2C9 Assay (ADD 777)
  - CYP2C19 Assay (ADD 778)
  - Cytochrome panel assay with activity outcomes (ADD 1851)
  - Primary cell-based high-throughput screening assay for identification of compounds that inhibit KCNQ2 potassium channels (ADD 2156)
  - Cycloheximide Counterscreen for Small Molecule Inhibitors of Shiga Toxin (ADD 2314)
  - a HTS for Small Molecule Inhibitors of Shiga Toxin (ADD 2315)
  - Fluorescence Polarization Cell-Free Homogeneous Assay (HTS) to Identify Inhibitors of the ANA Histone H2A/H2B Interaction (ADD 2629)
  - Primary cell-based high-throughput screening assay for identification of compounds that inhibit KCNQ1 potassium channels (ADD 2642)
  - Primary cell-based high-throughput screening assay for identification of compounds that potentiate/activate regulator of G-protein signaling 4 (RGS4) (ADD 46311)
  - Validation (re-confirmation) assay for identification of compounds that inhibit KCNQ1 potassium channels (ADD 58553)
  - HTS identification of antagonists of the CRF-binding protein and CRF-R2 receptor complex (ADD 588475)
  - Primary cell-based high-throughput screening for identification of compounds that allosterically regulate MrxG1 receptor signaling (ADD 58875)
  - Dose Response confirmation of HTS hits for small molecule antagonists of the CRF-binding protein and CRF-R2 receptor complex (ADD 601280)
  - Validation assay for identification of compounds that activate the regulator of G-protein signaling 4 (RGS4) (ADD 602282)
  - Counter screen for identification of compounds that activate the regulator of G-protein signaling 4 (RGS4): Non-induced cells with the primary screen assay (ADD 602283)
  - Validation for compounds that inhibit KCNQ1 potassium channels on automated electrophysiology assay (ADD 621420)
  - Specificity screen against KCNQ2 for identification of compounds that inhibit KCNQ1 potassium channels (ADD 651746)
  - Specificity screen against KCNQ1/KCNEN1 for identification of compounds that inhibit KCNQ1 potassium channels (ADD 652147)




O=C(c1cc2ccccc2s1)c3ccccc3

-0.47

NA

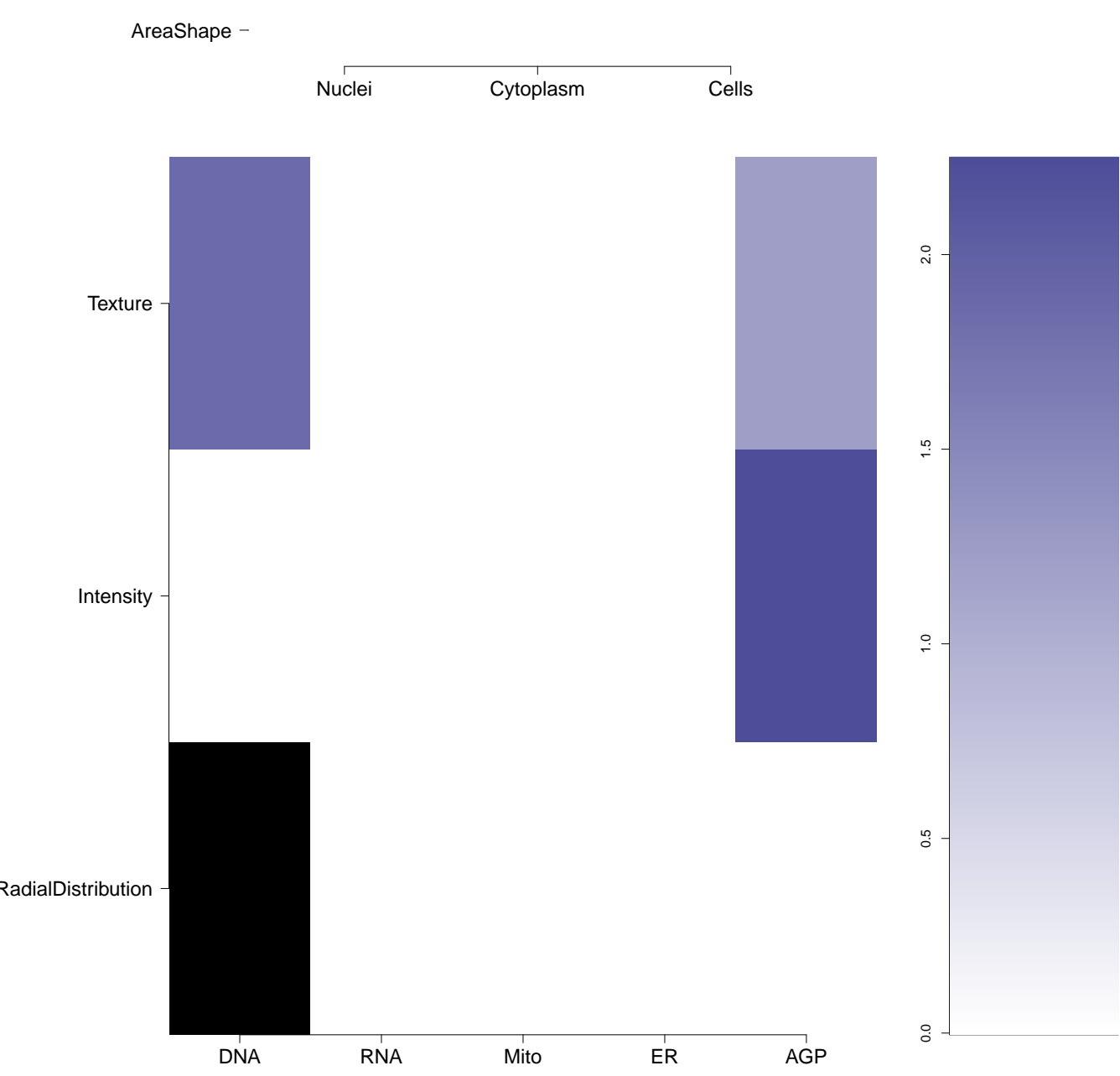
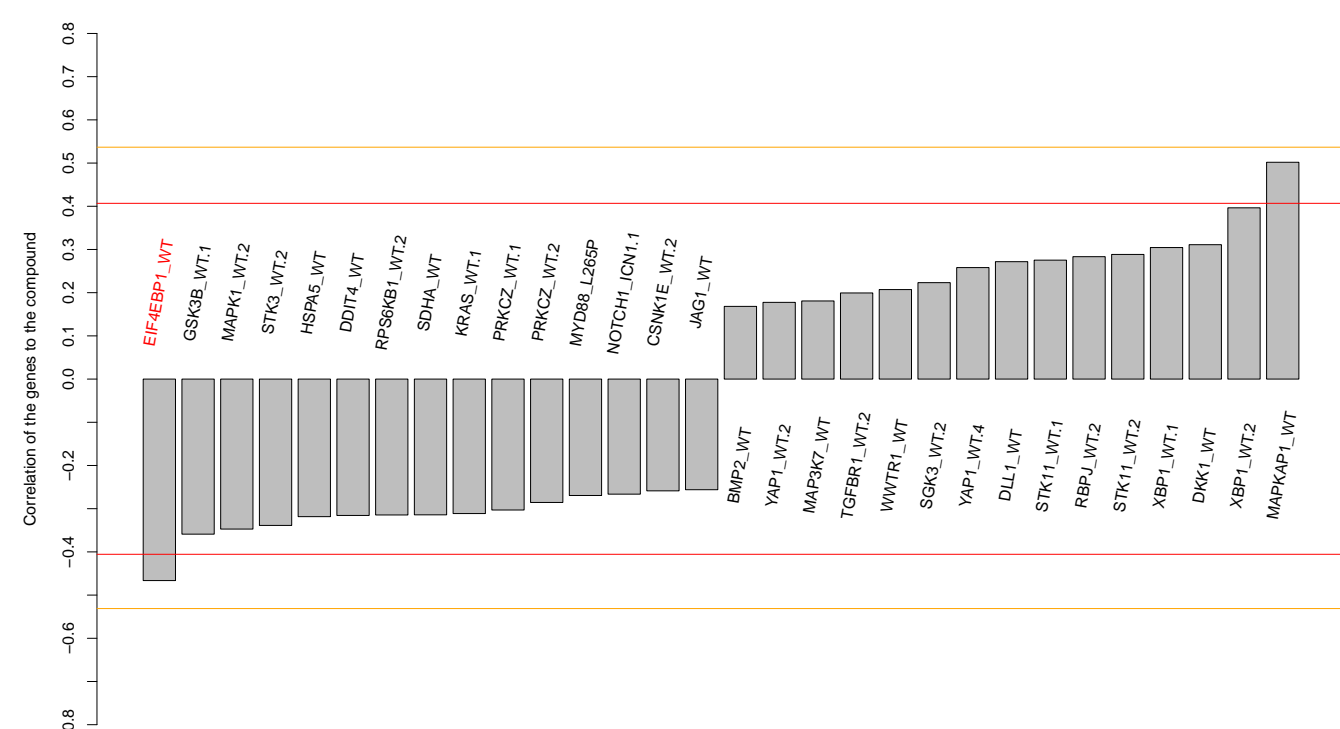


Total number of assays tested in: 38.

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-0.47

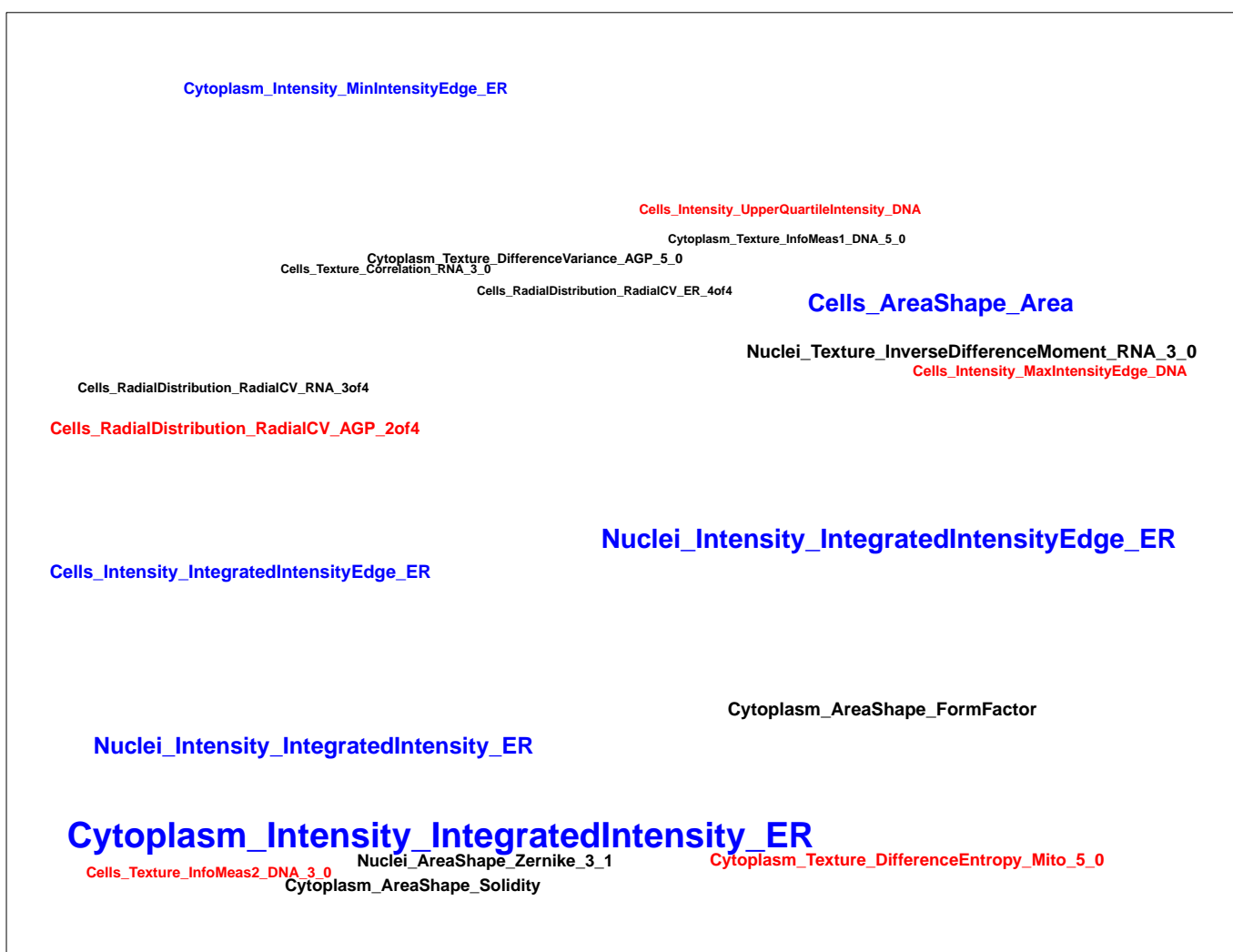
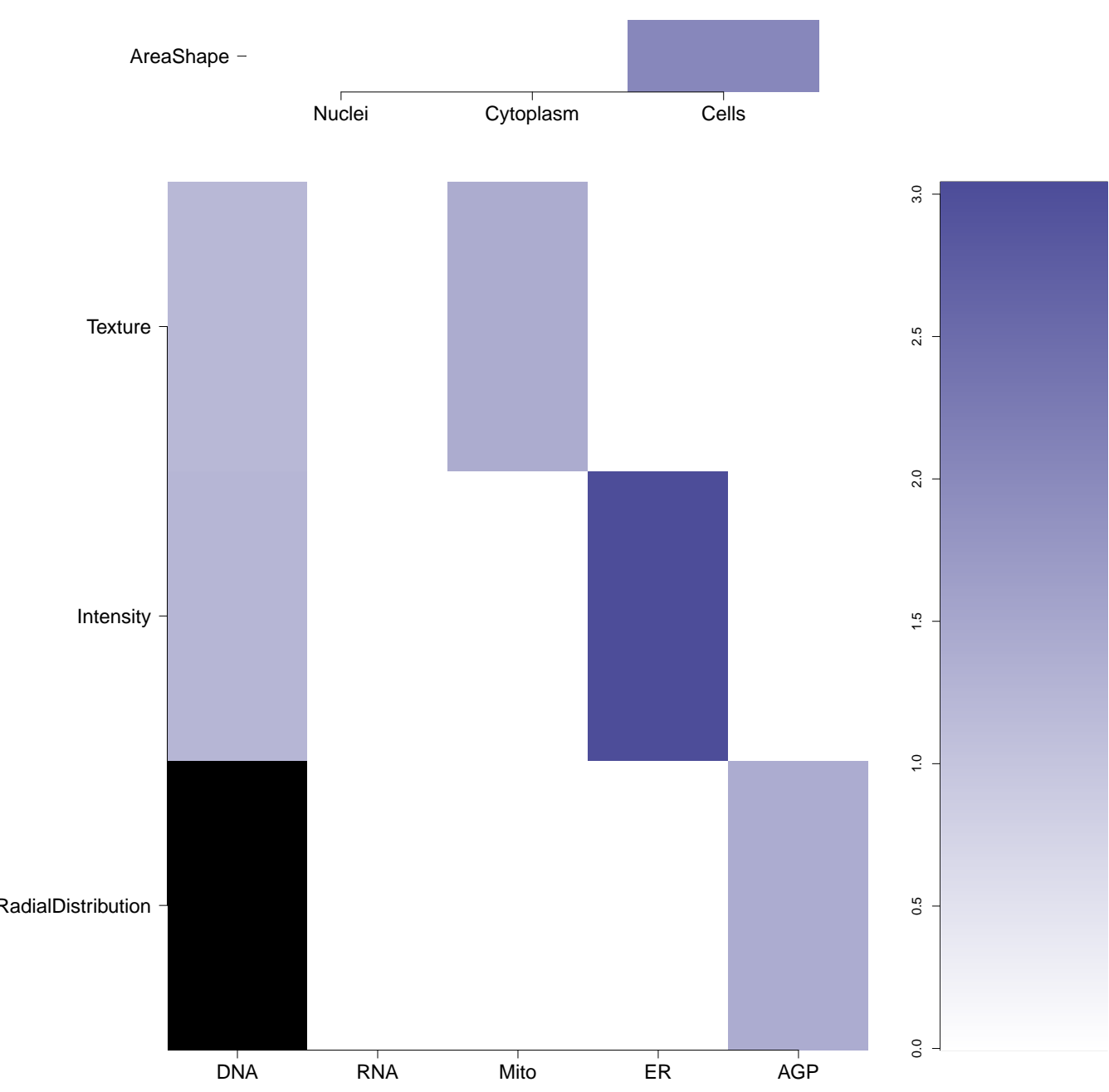
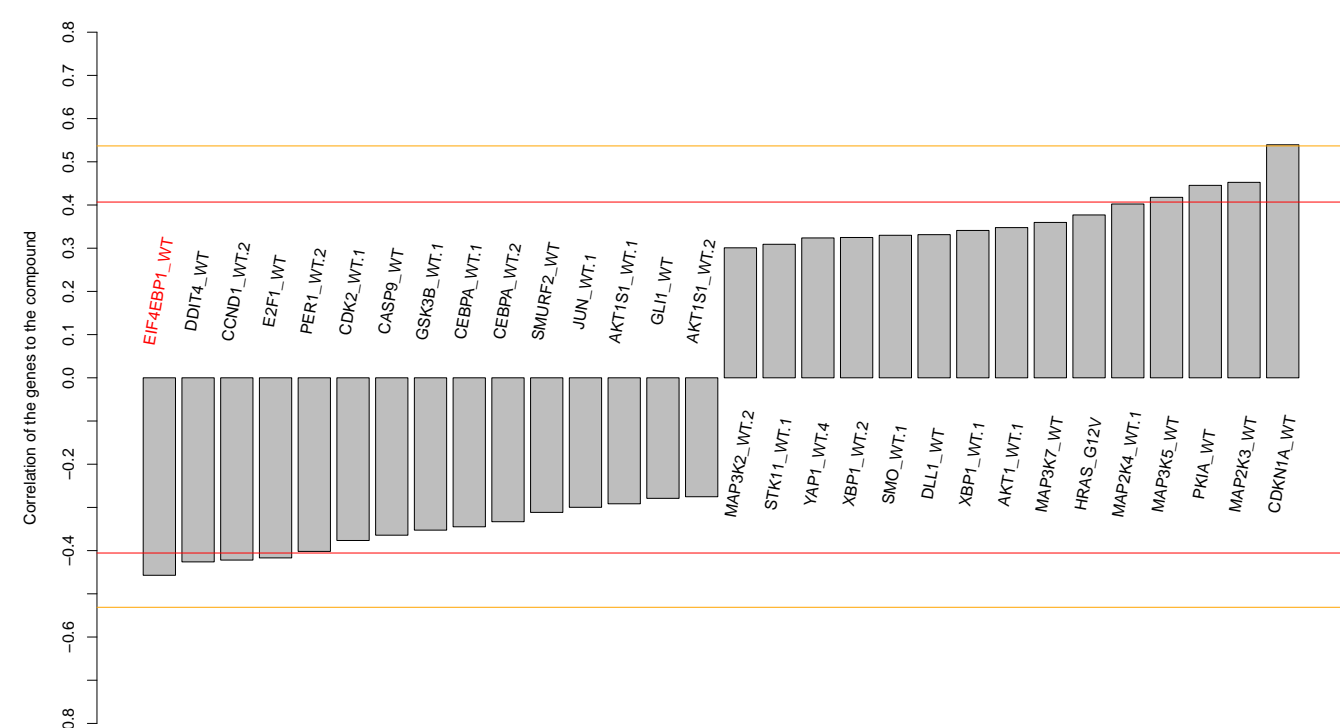
NA



Chemical structure of compound 10, a 12-membered macrocyclic lactone. It features a cyclopropylmethyl ketone side chain and a 2-hydroxyethyl side chain.

-0.46

0.302



Total number of assays tested in: 54.