

How similar is this gene to the other genes?

| Gene | Correlation |
|---------------|-------------|
| CASP8_WT.1 | 1.00 |
| CASP8_WT.2 | 0.72 |
| RHOA_Q63L | 0.64 |
| PRKACA_WT.2 | 0.61 |
| PRKACA_WT.1 | 0.58 |
| GLI1_WT | 0.55 |
| JUN_WT.2 | 0.54 |
| CXXC4_WT | 0.49 |
| CEBPA_WT.2 | 0.49 |
| CEBPA_WT.1 | 0.49 |
| BCL2L1_WT | 0.46 |
| JUN_WT.1 | 0.43 |
| PPARGC1A_WT.2 | 0.39 |
| RAF1_L613V | 0.38 |
| BRAF_WT.2 | 0.30 |
| PRKCZ_WT.2 | -0.31 |
| YAP1_WT.2 | -0.32 |
| PRKCZ_WT.1 | -0.32 |
| JAG1_WT | -0.33 |
| ERN1_WT.1 | -0.35 |
| AKT3_WT.3 | -0.37 |
| TGFBR1_K232R | -0.38 |
| SMAD3_WT.1 | -0.39 |
| DLL1_WT | -0.40 |
| RPS6KB1_WT.2 | -0.43 |
| CSNK1A1_WT.3 | -0.46 |
| MAPK1_WT.2 | -0.47 |
| RPS6KB1_WT.1 | -0.53 |
| AKT1_WT.1 | -0.58 |
| AKT2_WT | -0.65 |

AreaShape

Nuclei Cytoplasm Cells

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DNA RNA Mito ER AGP

0 10 20 30 40 50 60

Cells_Intensity_MedianIntensity_ER

Cytoplasm_Intensity_MeanIntensity_ER

Cells_Intensity_MinIntensity_ERNA
Cytoplasm_Intensity_MinIntensity_ERNA

Cells_Intensity_StdIntensity_AQP

Cells_Intensity_MeanIntensityEdge_ERNA

Cytoplasm_Intensity_MinIntensityEdge_ERNA

Cells_Intensity_MeanIntensityEdge_ER

Cells_Intensity_StdIntensityEdge_Mito

Cells_Intensity_UpperQuartileIntensity_DNA

Cells_Intensity_MaxIntensityEdge_DNA

Cells_Intensity_StdIntensityEdge_DNA

Cytoplasm_Intensity_MeanIntensity_DNA

Cytoplasm_Texture_SumVariance_DNA_10_0

Cells_Intensity_MeanIntensityEdge_DNA

Cytoplasm_Texture_Contrast_DNA_10_0

Cytoplasm_Intensity_UpperQuartileIntensity_DNA

Cytoplasm_Texture_Contrast_DNA_5_0

Nuclei_Texture_Gabor_ER_10

CASP8_WT.1 (41754)

