

tf_4 gene_2

tf_2 gene_2

tf_3 gene_1

tf_1 gene_1

t1

t2

t3

t4

t5

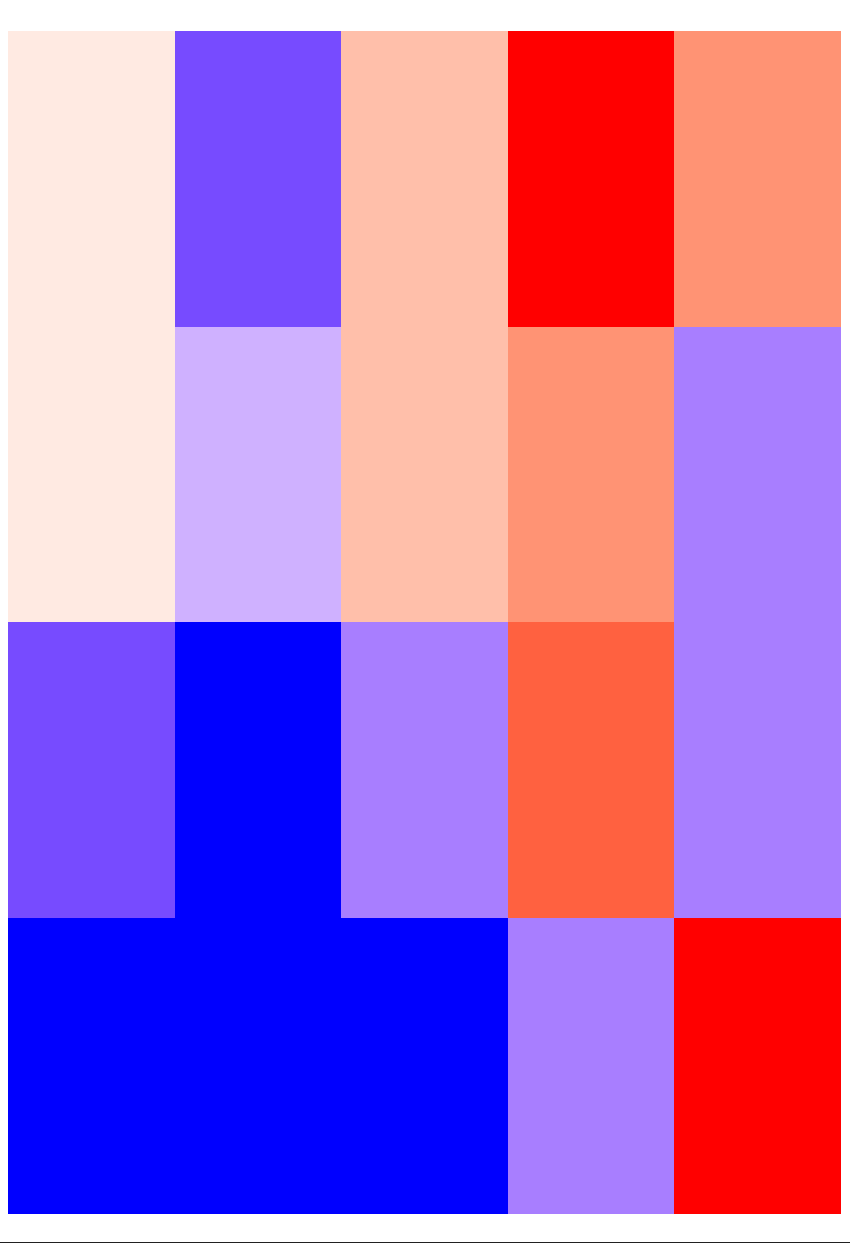
Prediction Score

7.5

5.0

2.5

0.0



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