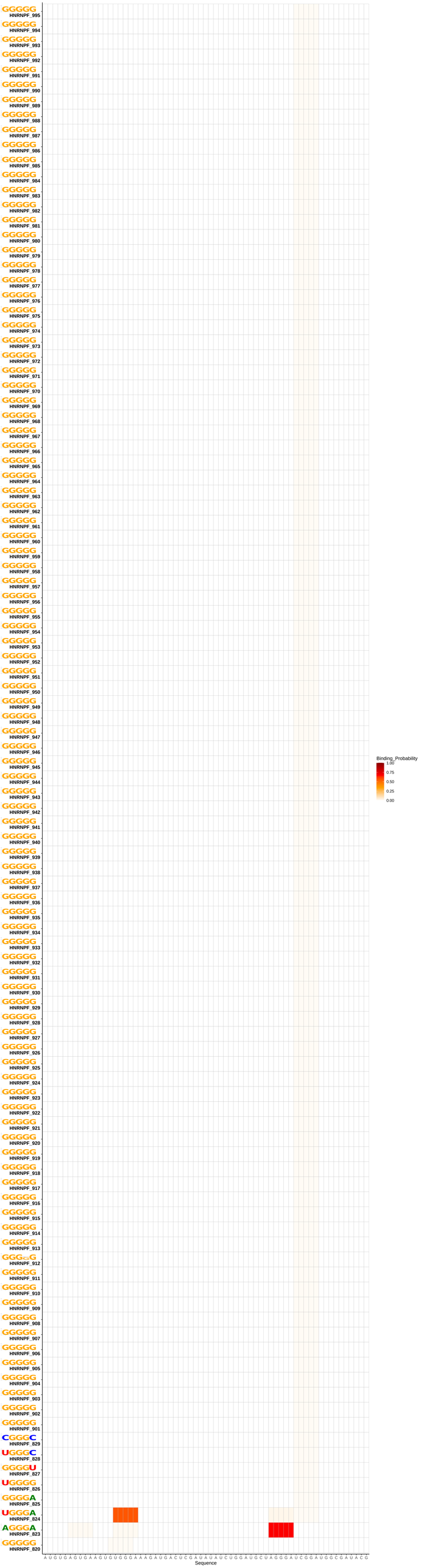
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Sequence
Binding_Probability
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0.25
0.00