

# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc473

Average coverage per 5'UTR

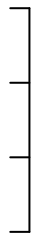
0.0020 0.0024

-500bp

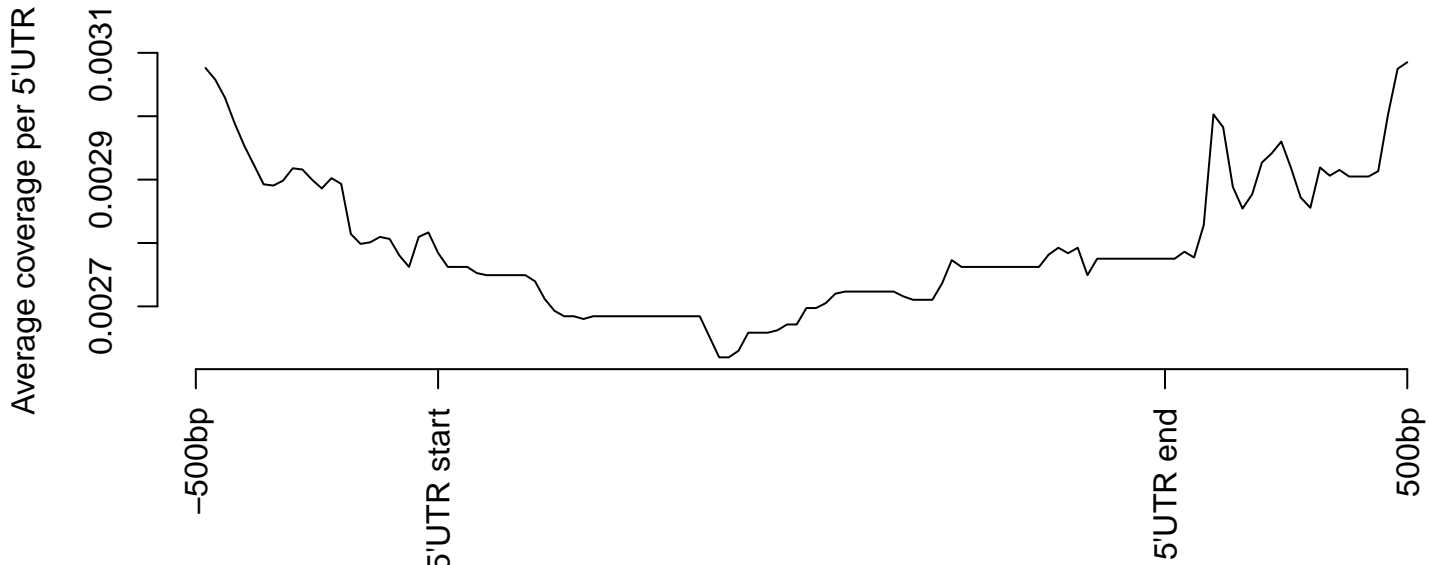
5'UTR start

5'UTR end

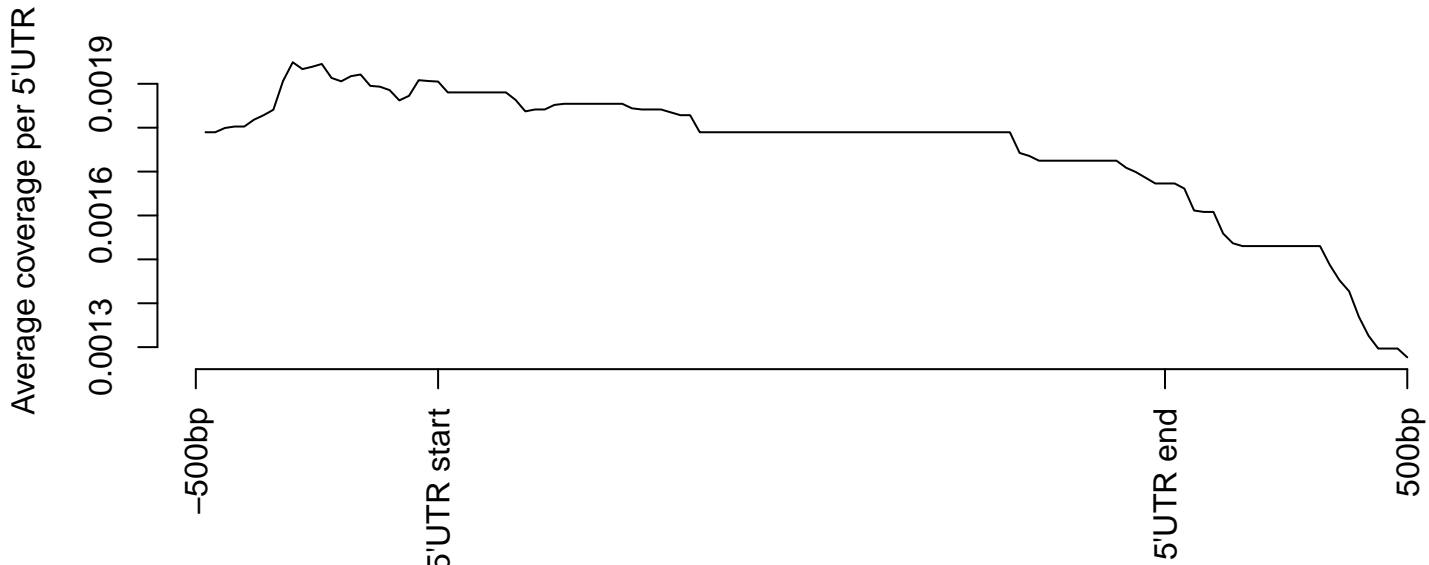
500bp



# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc474



# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc475



# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc476

Average coverage per 5'UTR

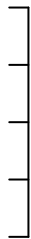
0.0014 0.0020

-500bp

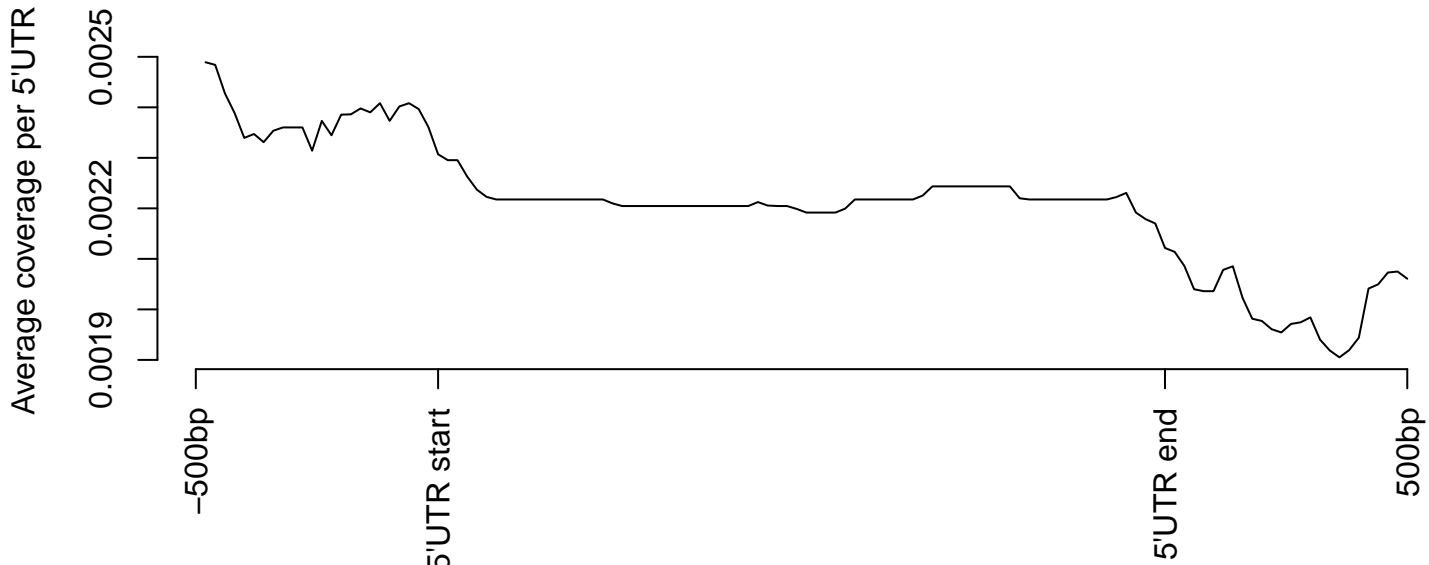
5'UTR start

5'UTR end

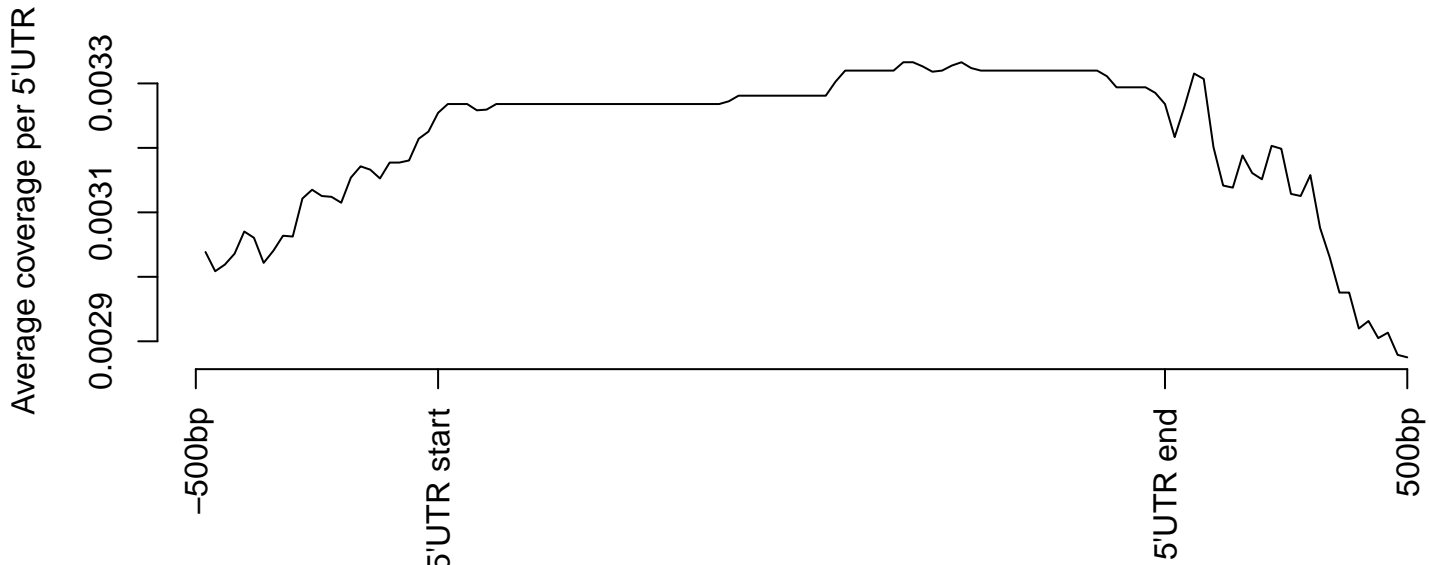
500bp



# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc477

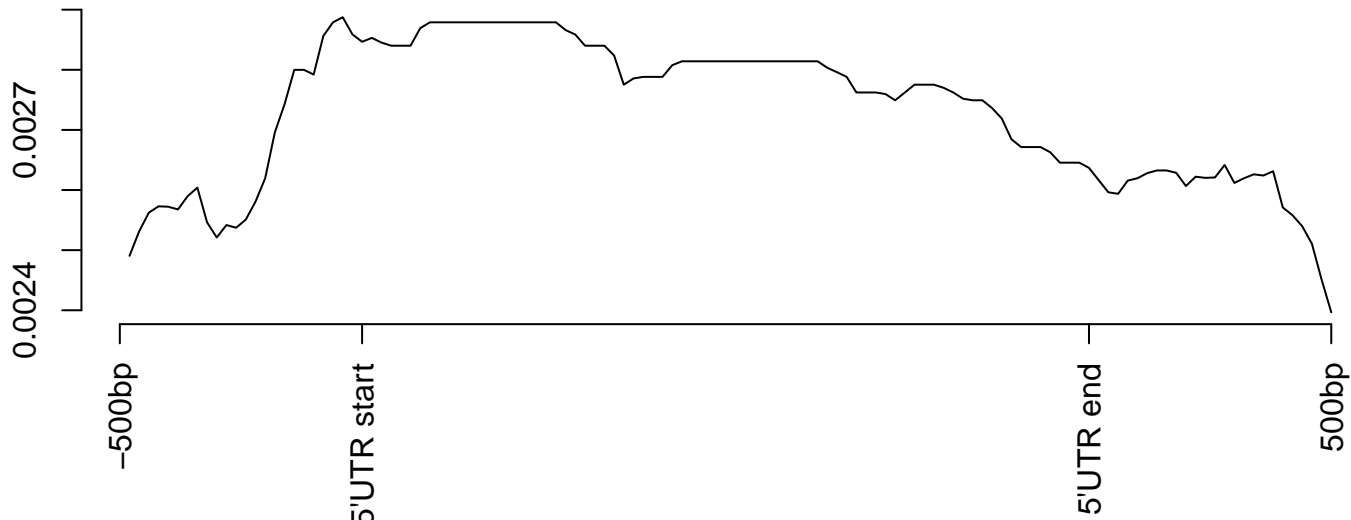


# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc478



# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc480

Average coverage per 5'UTR



# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc481

Average coverage per 5'UTR

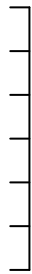
0.00195 0.00215

-500bp

5'UTR start

5'UTR end

500bp





# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc482

Average coverage per 5'UTR

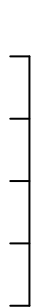
0.0028 0.0032 0.0036

-500bp

5'UTR start

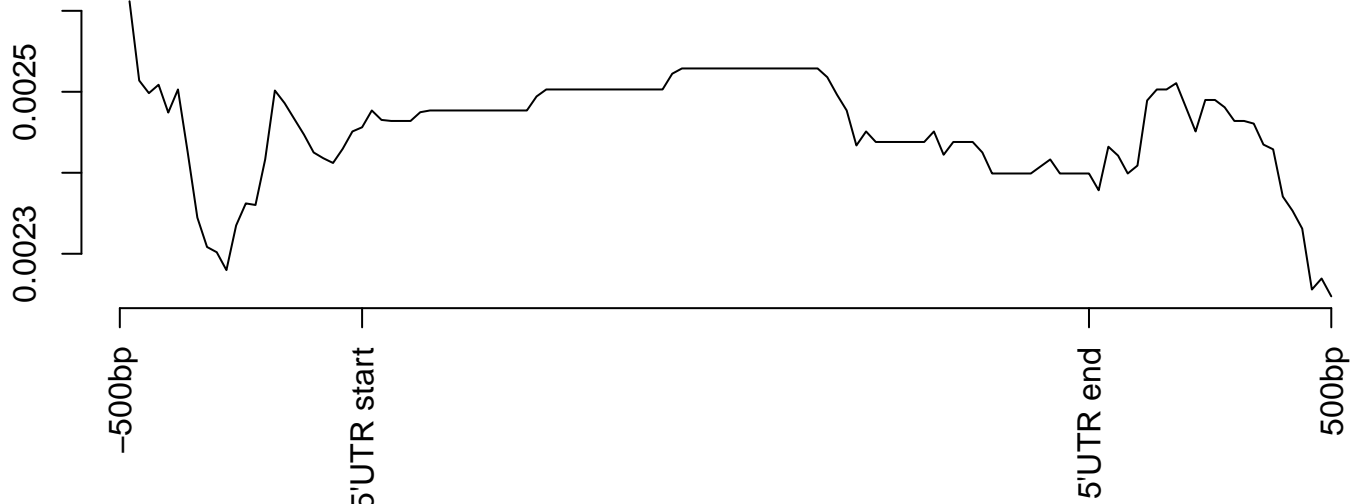
5'UTR end

500bp

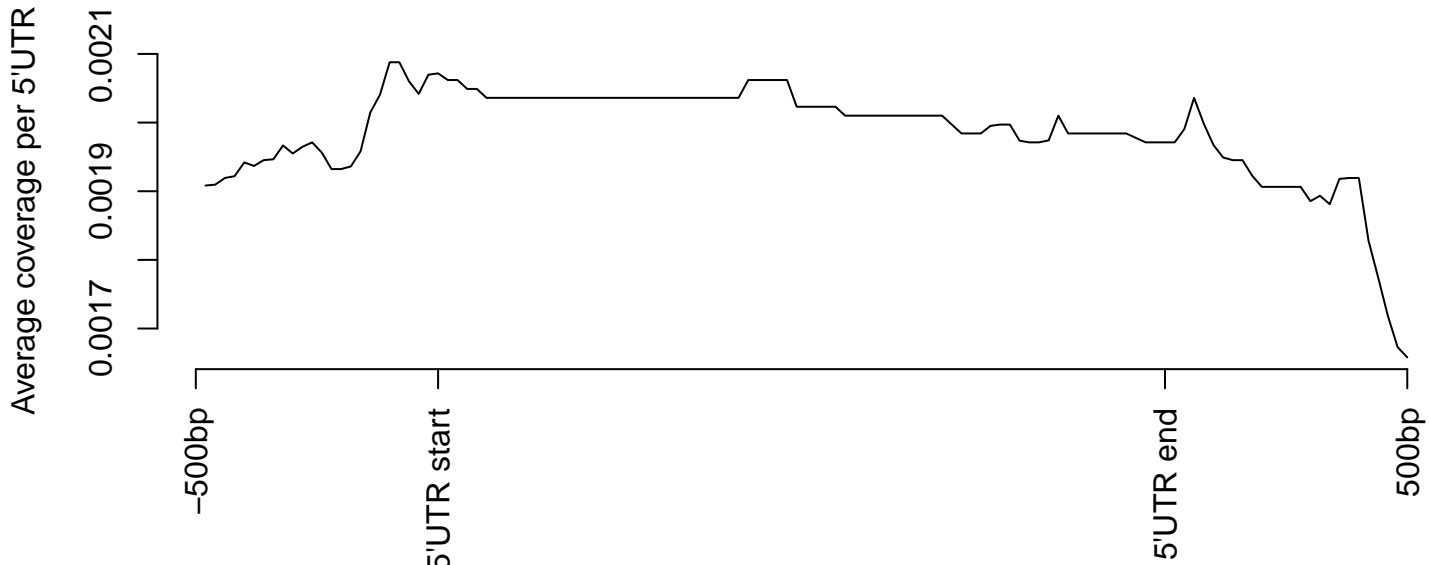


# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc483

Average coverage per 5'UTR



# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc484



# HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc485

Average coverage per 5'UTR

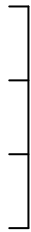
0.0021 0.0023

-500bp

5'UTR start

5'UTR end

500bp



# HumanNeuronCulture PositiveMerged E.– HumanNeuronCulturePositiveMerged

Average coverage per 5'UTR

0.026 0.028 0.030

-500bp

5'UTR start

5'UTR end

500bp

