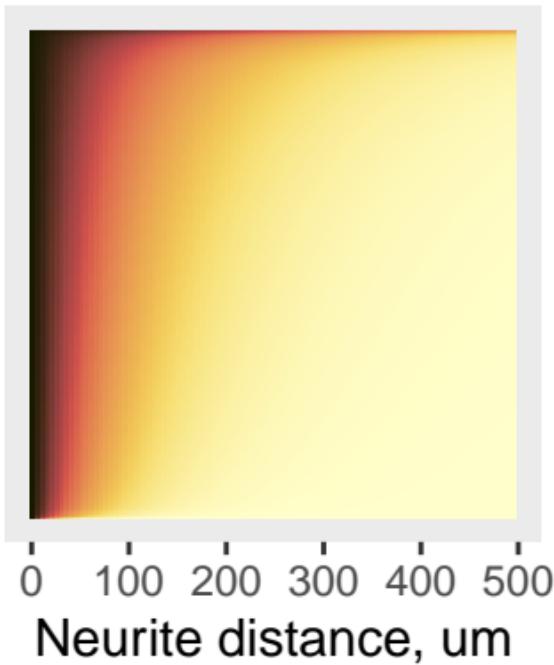


transcripts arranged by hl, 11042

GFP

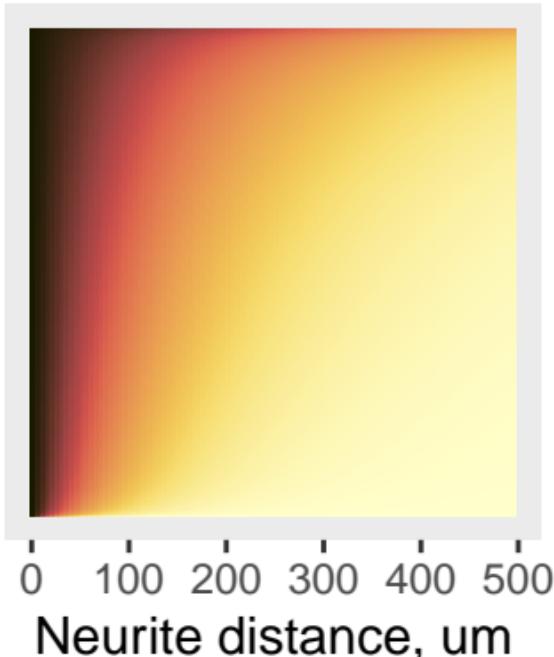


Probability:

1.00  
0.75  
0.50  
0.25

transcripts arranged by hl, 10368

dnCaf1



Probability:

1.00  
0.75  
0.50  
0.25