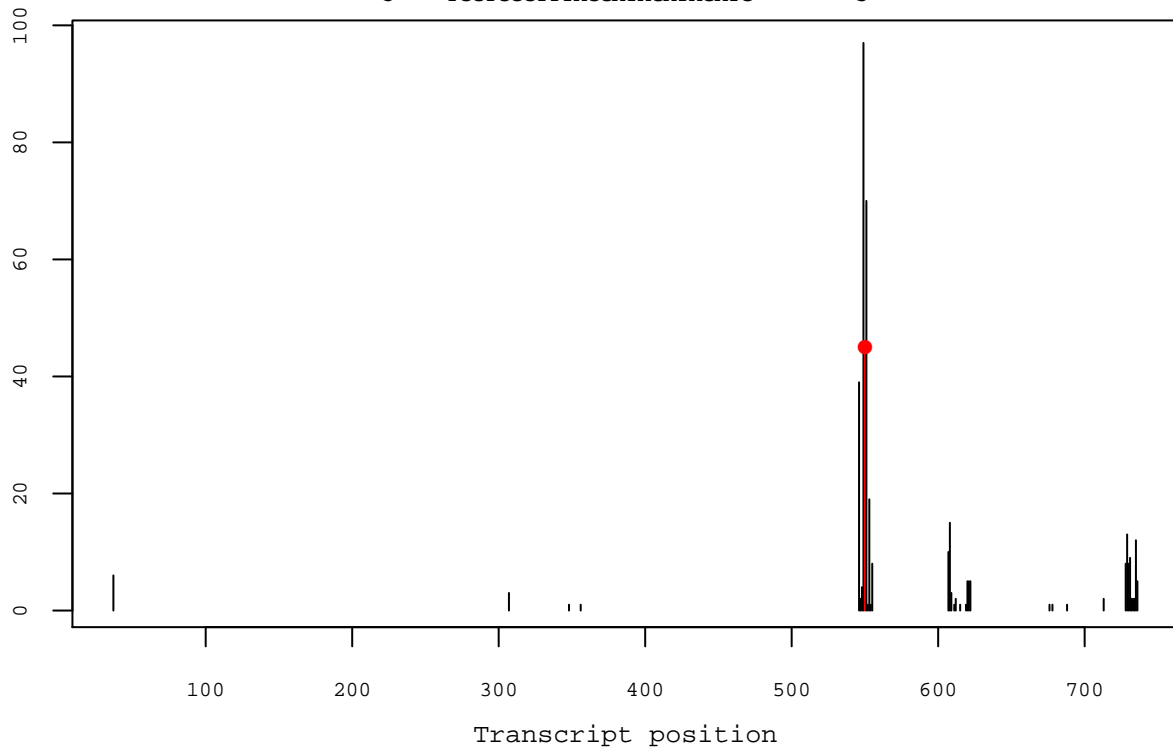


5' GTACCACTCAAATCCTATGTATTACATATCT '3
|||||o|||||||o|||
3' TGGTGGGTTTAGGATACATAGATG '5

Fragment Abundance



Cleavage site: 550 Tag abundance: 45 Weighted abundance: 1.318 Category: 2
sRNA abundance: 8 Alignment score: 1.5 MFE ratio: 0.866 p-value: 0.013