

hsa-miR-6131 read count : 1  
remaining reads : 0

freq.

1

0.75

0.5

0.25

0

length

hsa-miR-6131

5' - ucccgcauucccucugcuuuggucaggugggugccuccuuccaugggguagagccagagauugguggguuucuggcuggucagauugggaguggacagagacccgggguccuc - 3'

reads mm sample

1 1 seq

hsa-miR-6131

[illegible]