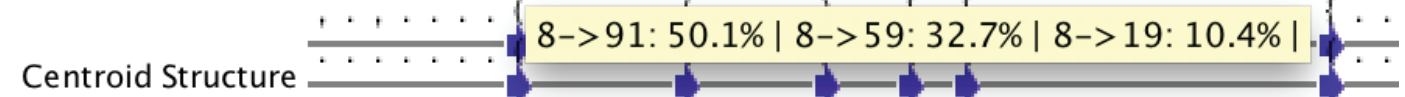


*R.norvegicus.2/1-64* G - UUACA UU - - GAU GAGAA CAGAAACA - UAAA - - CUAUG  
*R.norvegicus.1/1-61* A - UAUUU GUU - UAU GAUGG UCACAGUG - UAAA - - GUUCA  
*O.aries.1/1-68* G - ACGCU UC - - - AUGACAG GAAGGACU - GAAA - UGUCUC



RNAalifold Consensus G - UUUCAUUUU + AUGACGGCCUGUGCUCUAAA - CCCU + C  
 G \_UUUCAUU \_ \_ \_AUGACGGCCUGUGCU \_UAAA \_CCUCC



Sequence position 8 8->91: 50.1% | 8->59: 32.7% | 8->19: 10.4% |