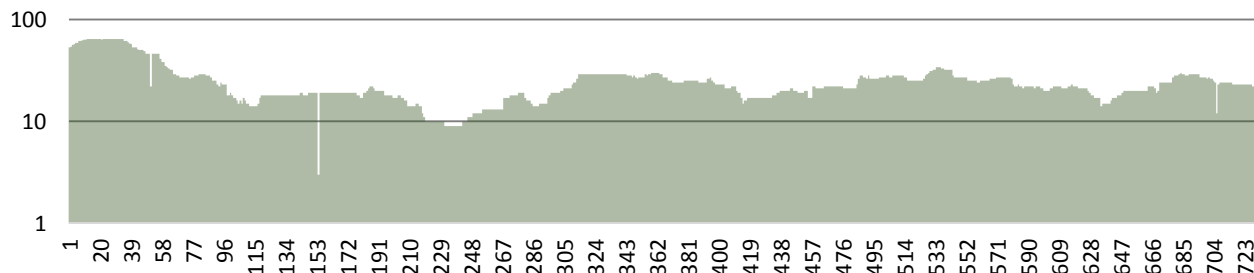


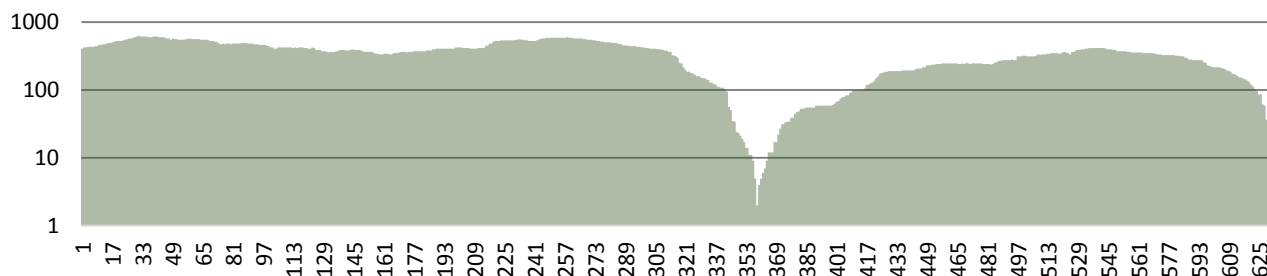
ITS-2 MK411275 *R. robustus* II vs. SRX1996481

1.6% <10 reads



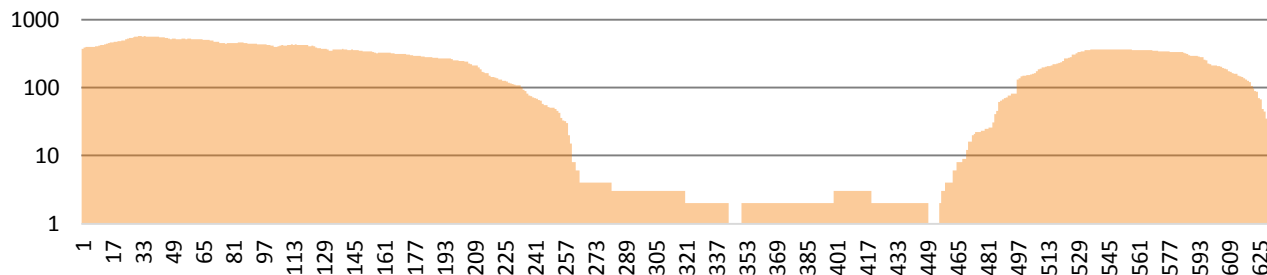
D2-28S AF435858.1 *R. robustus* II vs. SRX1996481

1.3% <10 reads



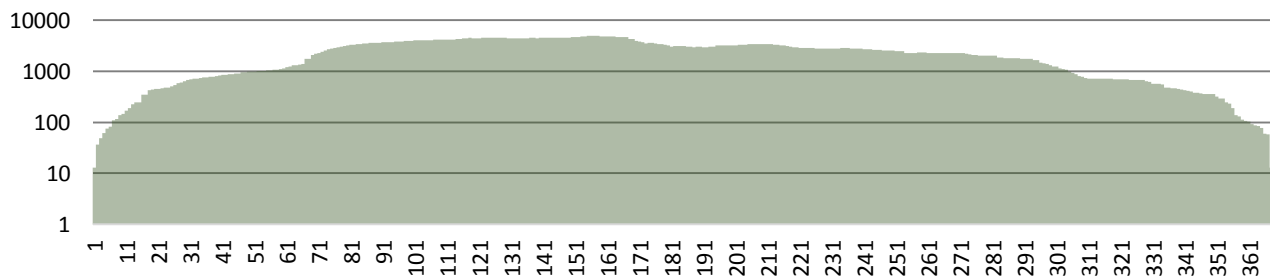
D2-28S AF435859.1 *R. robustus* IV vs. SRX1996481

33% <10 reads



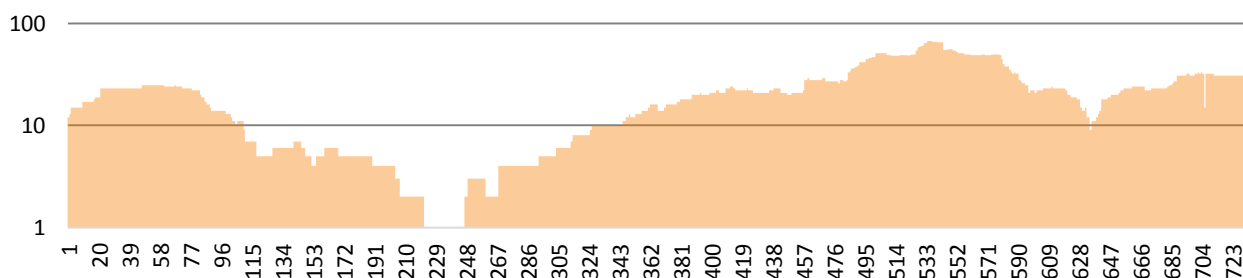
Cytb KR072682.1 *R. montenegrensis* vs. SRX1996482

0.3% <10 reads



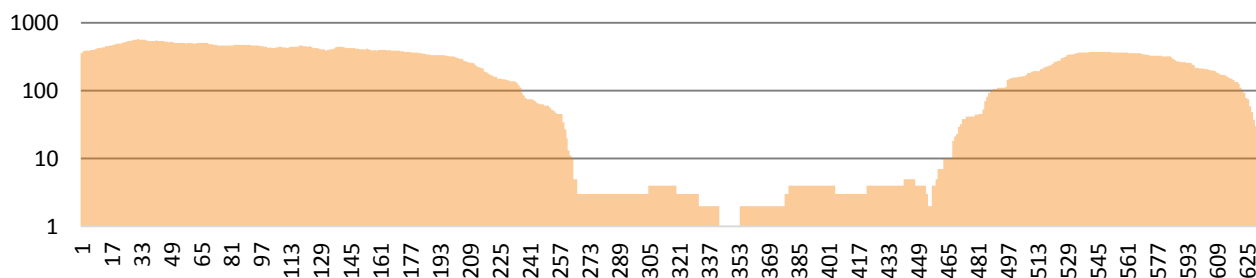
ITS-2 MK411274 *R. robustus* II vs. SRX1996482

29.6% <10 reads



D2-28S AF435859.1 *R. robustus* IV vs. SRX1996482

31.3% <10 reads

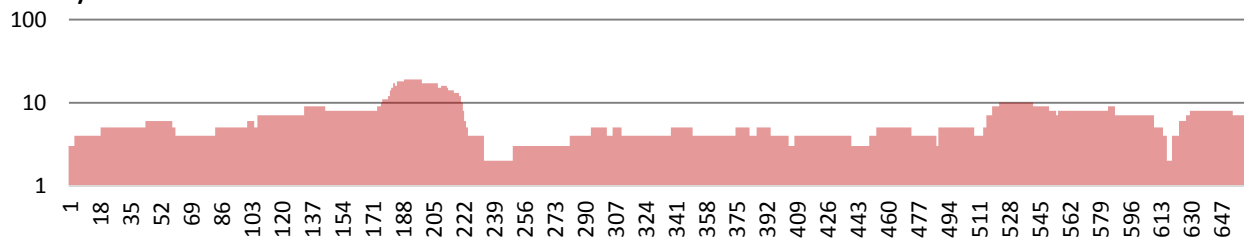


Nucleotide position

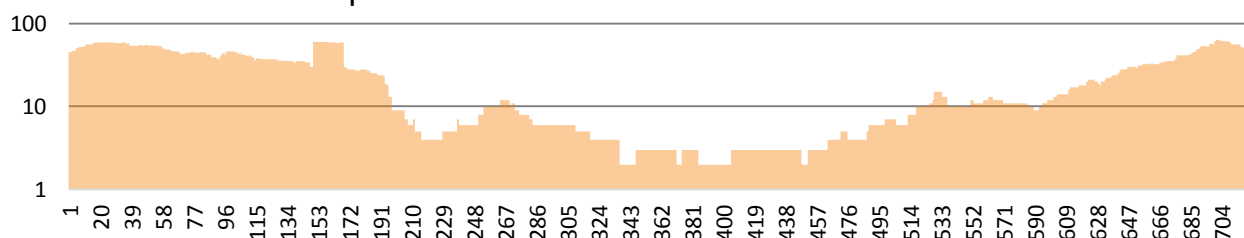
Depth-coverage (number of reads)

Cytb EF011724.1 *R. robustus* II vs. SRX1996483

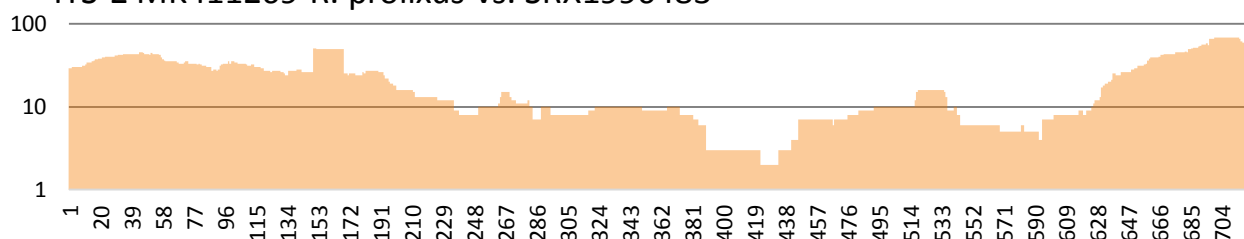
90.2% <10 reads

ITS-2 MK411270 *R. prolixus* vs. SRX1996483

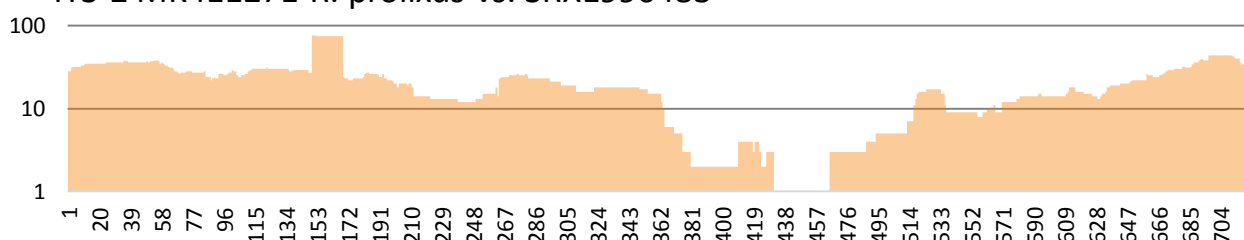
42.2% <10 reads

ITS-2 MK411269 *R. prolixus* vs. SRX1996483

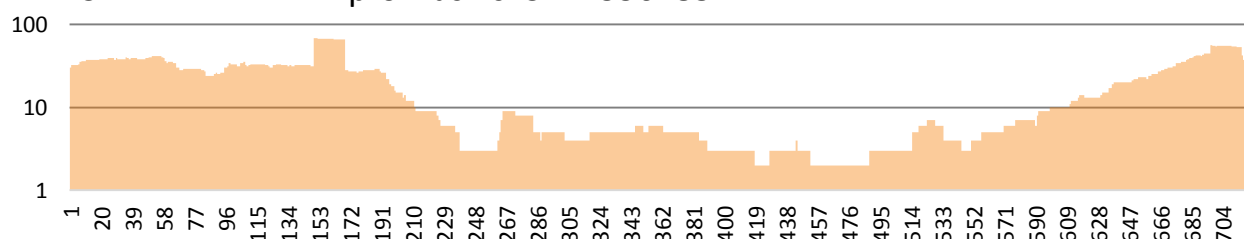
36.9% <10 reads

ITS-2 MK411271 *R. prolixus* vs. SRX1996483

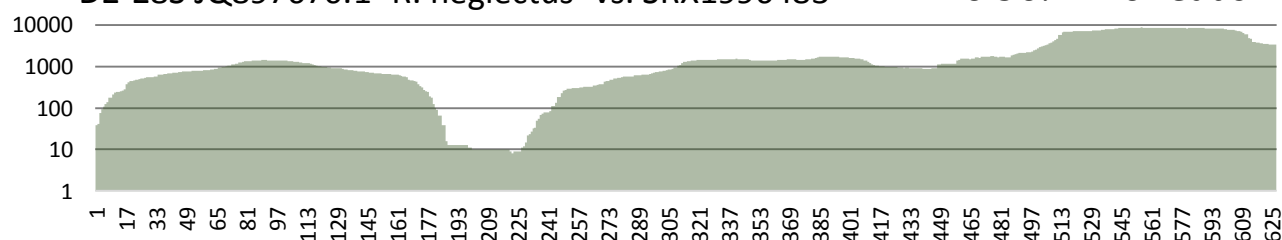
25.1% <10 reads

ITS-2 MK411272 *R. prolixus* vs. SRX1996483

53.7% <10 reads

D2-28S JQ897670.1 '*R. neglectus*' vs. SRX1996483

0.96% <10 reads



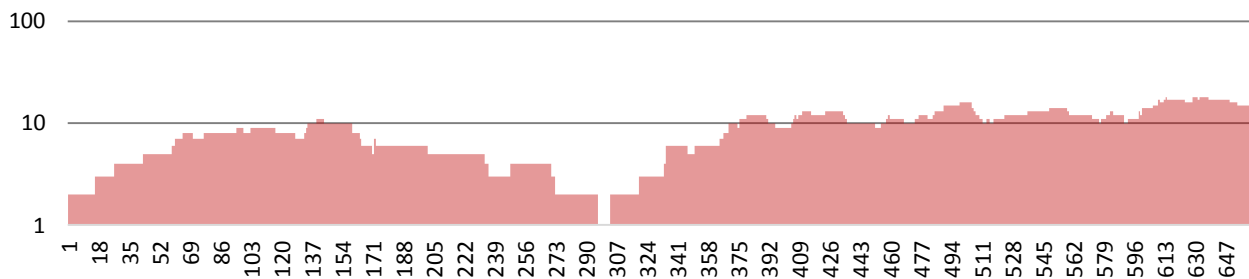
Depth-coverage (number of reads)

Nucleotide position

Cytb JX273156.1 *R. neglectus* vs. SRX1996484

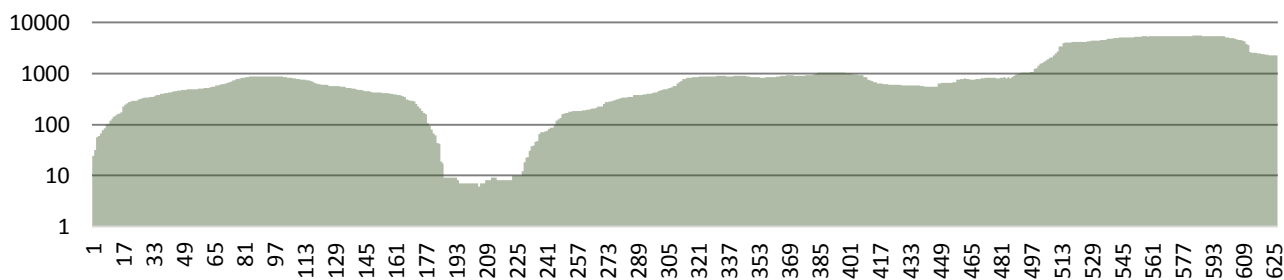
53.5% <10 reads

Depth-coverage (number of reads)



D2-28S JQ897670.1 '*R. neglectus*' vs. SRX1996484

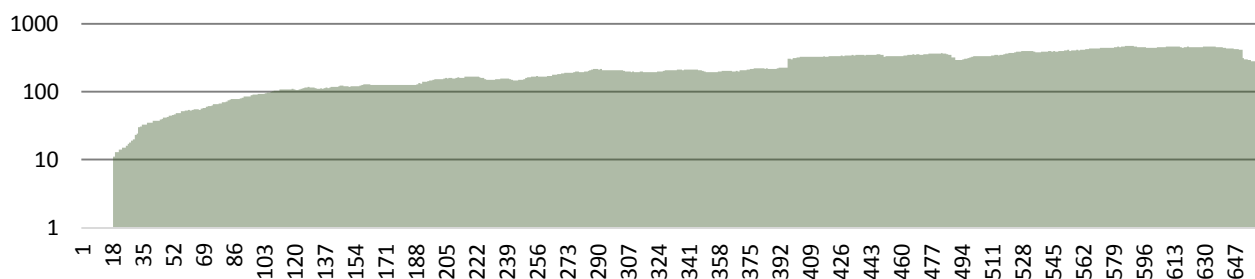
5.7% <10 reads



Nucleotide position

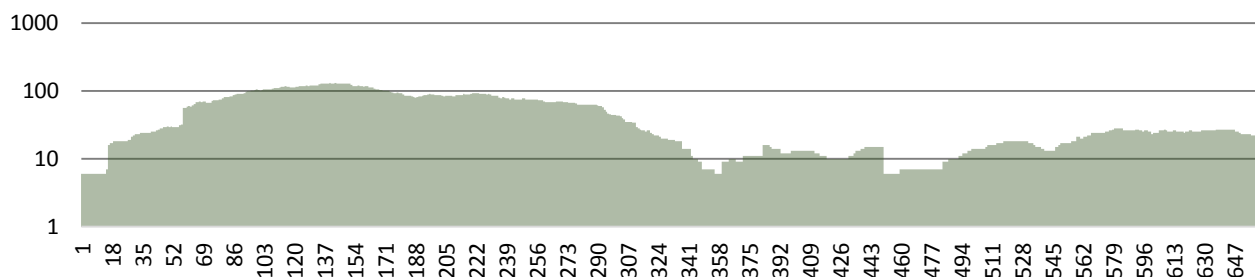
Cytb AF421339.1 *R. prolixus* vs. ERX1387159

2.7% <10 reads



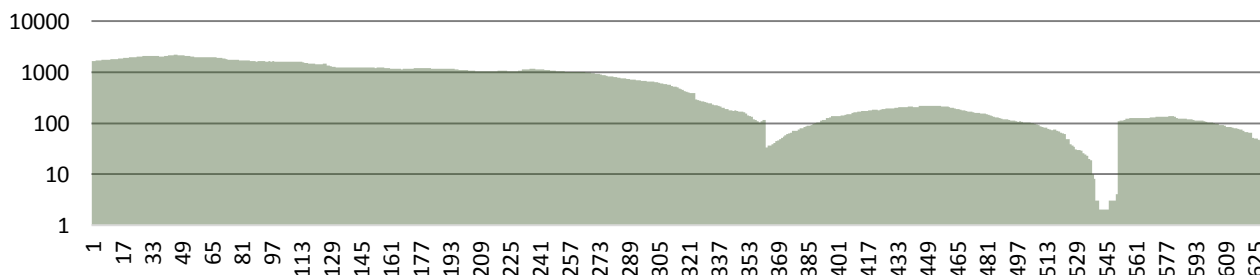
Cytb AF421341.1 *R. robustus* II vs. ERX1387159

10.9% <10 reads



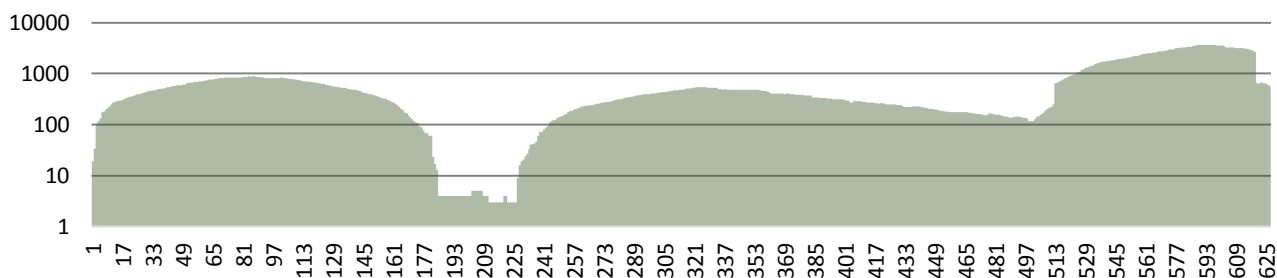
D2-28S AF435857.1 *R. robustus* III vs. ERX1387159

2.1% <10 reads



D2-28S JQ897670.1 '*R. neglectus*' vs. ERX1387159

6.9% <10 reads

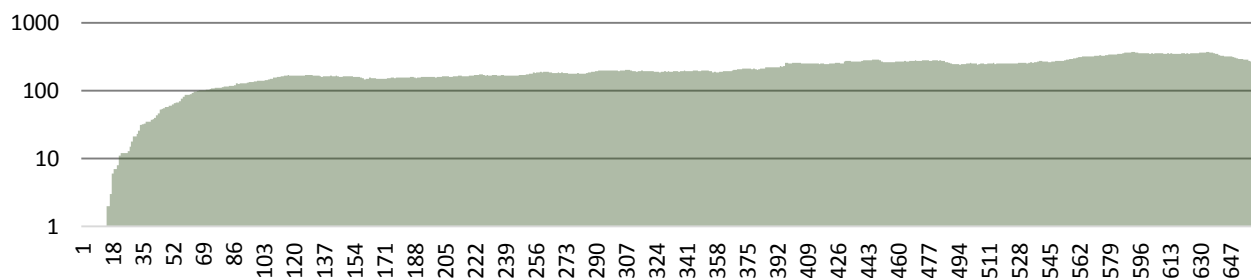


Nucleotide position

Depth-coverage (number of reads)

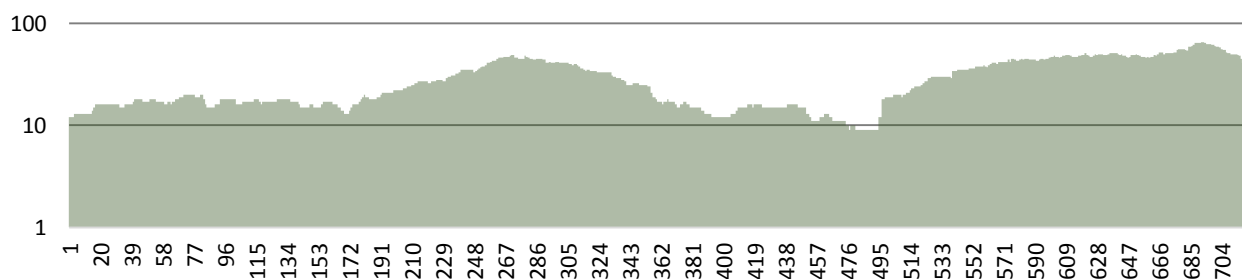
Cytb AF421339.1 *R. prolixus* vs. ERX1387160

3.2% <10 reads



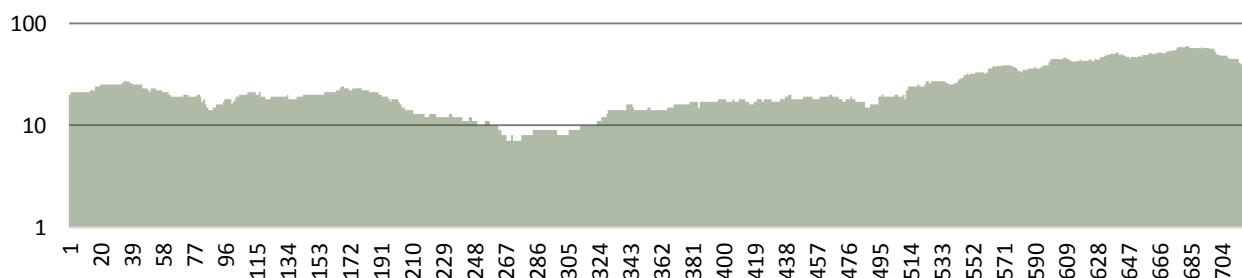
ITS-2 MK411271 *R. prolixus* vs. ERX1387160

2.1% <10 reads



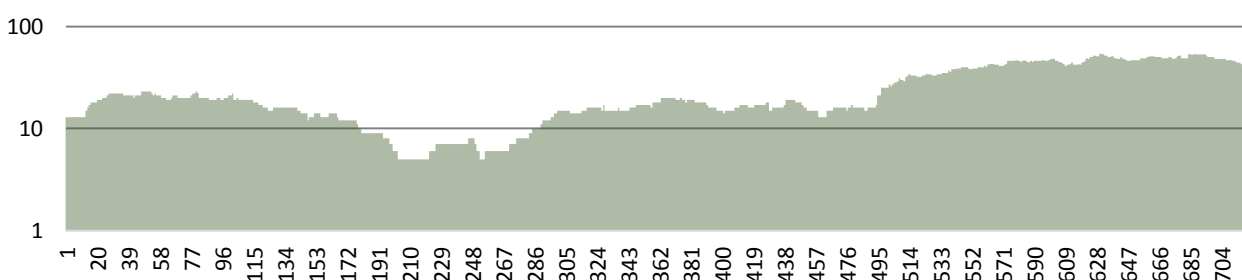
ITS-2 MK411269 *R. prolixus* vs. ERX1387160

6.9% <10 reads



ITS-2 MK411272 *R. prolixus* vs. ERX1387160

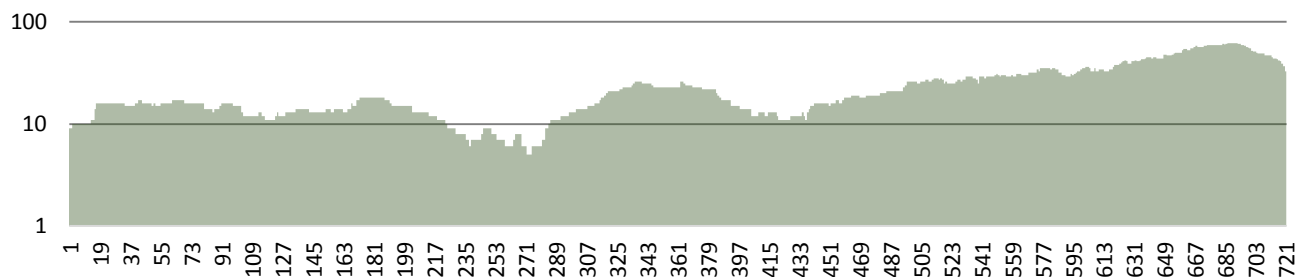
14.4% <10 reads



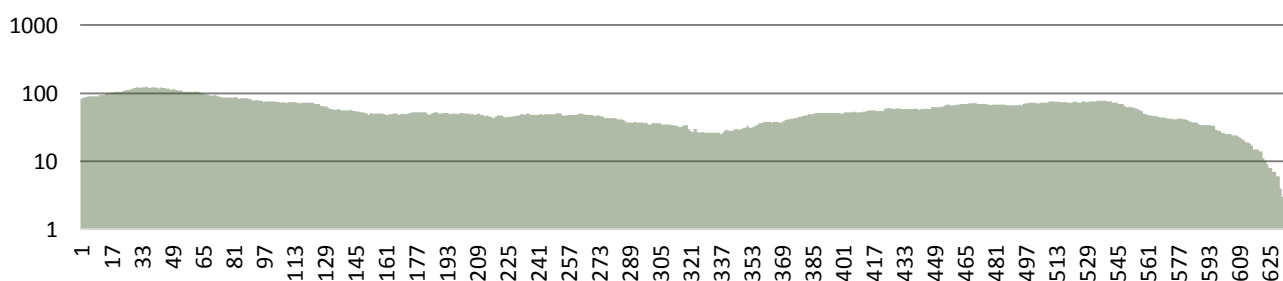
Nucleotide position

ITS-2 MK411270 *R. prolixus* vs. ERX1387160

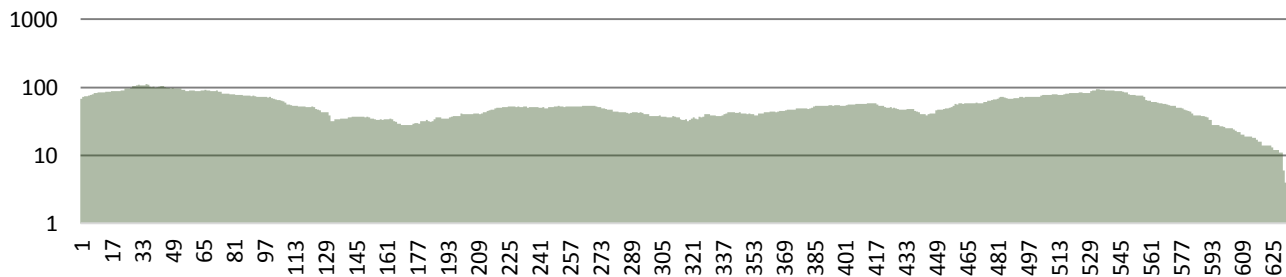
8.6% <10 reads

D2-28S AF435860.1 *R. prolixus* vs. ERX1387160

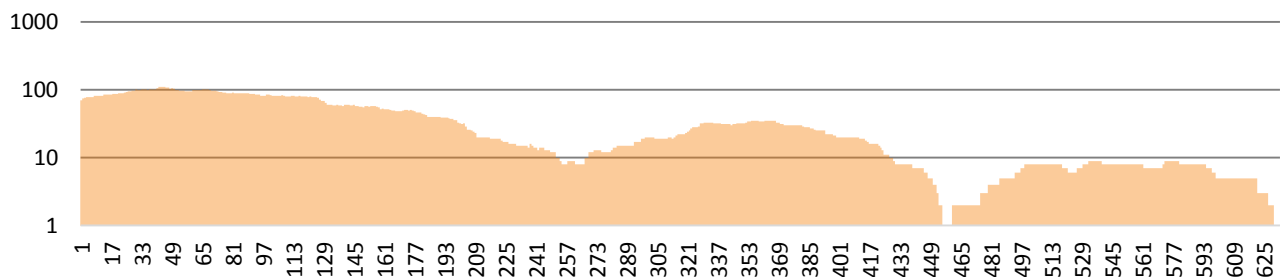
1.6% <10 reads

D2-28S AF435861.1 *R. robustus* I vs. ERX1387160

0.47% <10 reads

D2-28S AF435859.1 *R. robustus* IV vs. ERX1387160

34.3% <10 reads



Nucleotide position