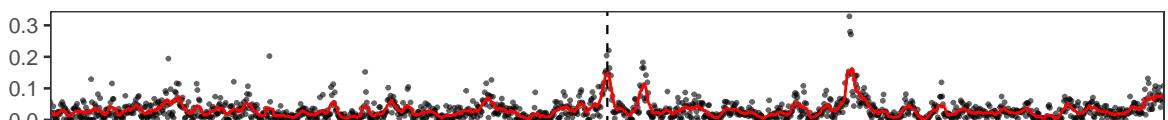


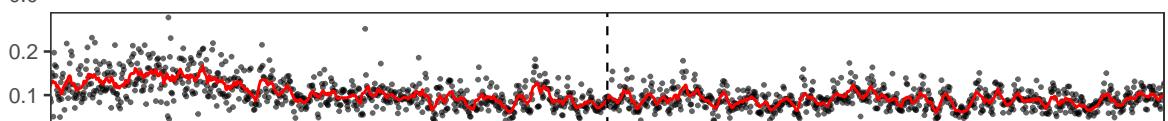
Peak: Chromosome_1_RagTag:119380000–119450000

Model: Selection–bottleneck: Selection

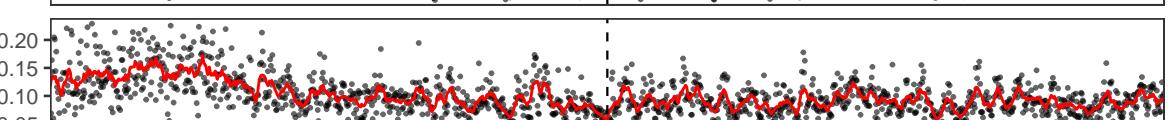
Xingu–Belem Fst



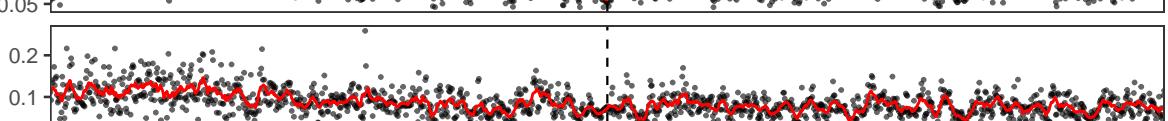
Xingu–Belem Dxy



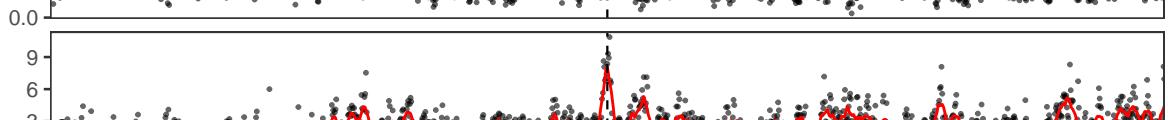
Xingu Pi



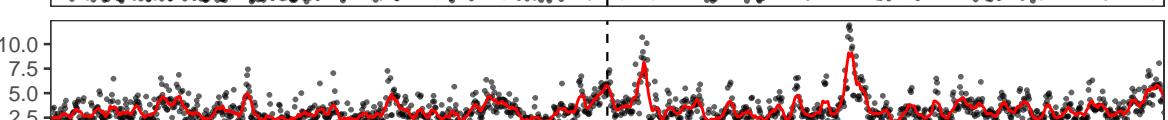
Belem Pi



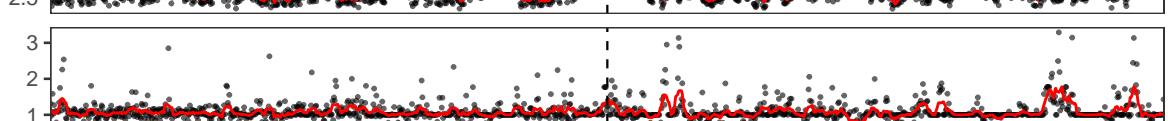
Xingu enrichment



Belem enrichment



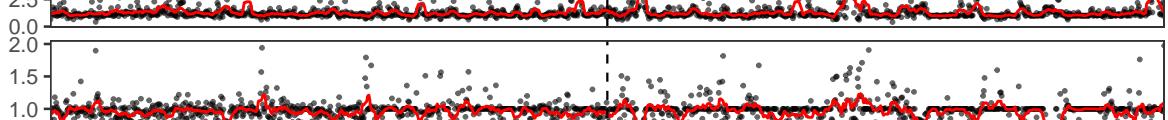
Xingu RTH (inverse)



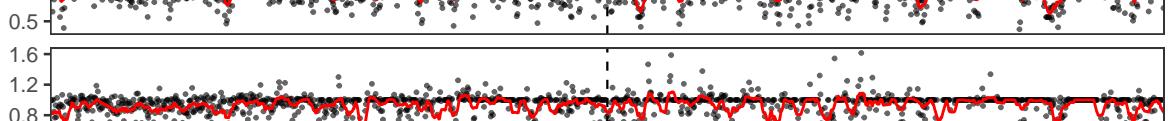
Belem RTH (inverse)



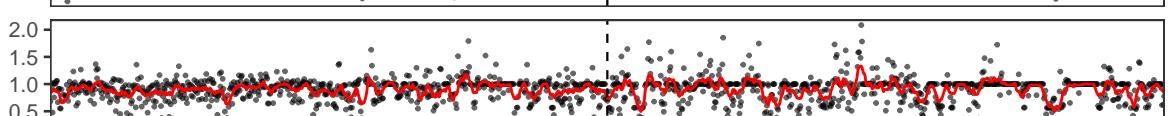
Xingu–Belem RTH



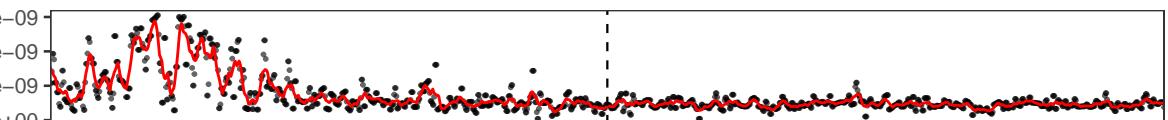
Tapajos–Xingu RTH (inverse)



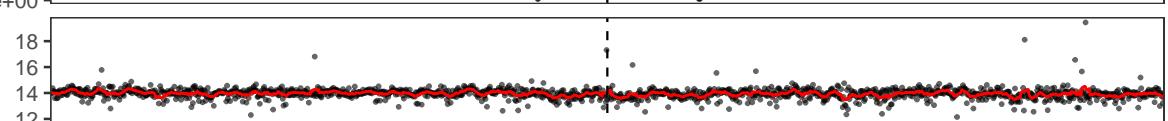
Tapajos–Xingu JCR



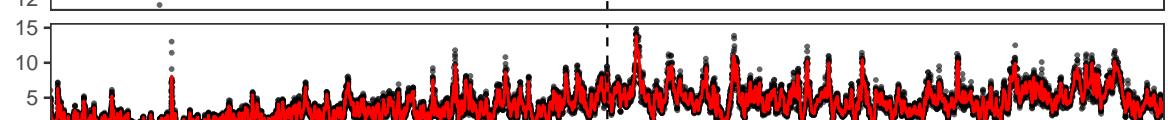
Recombination rate



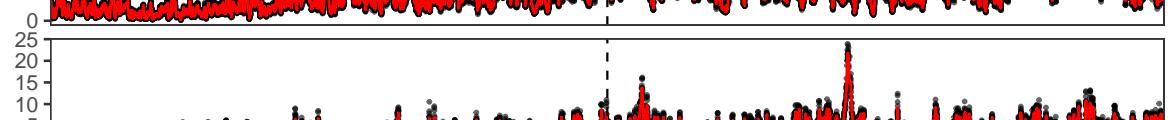
Average depth



Xingu RAiSD u



Belem RAiSD u



112 114 116 118 120 122 124 126
Position on Chromosome_1_RagTag (Mb)