

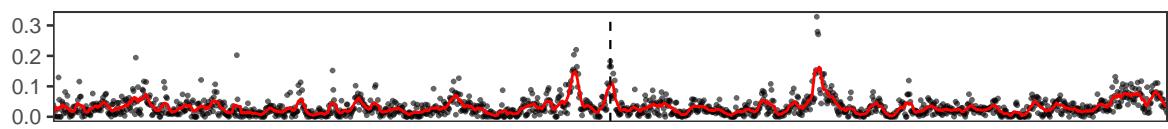
Peak: Chromosome_1_RagTag:119880000–119910000

Model: Overlapping models

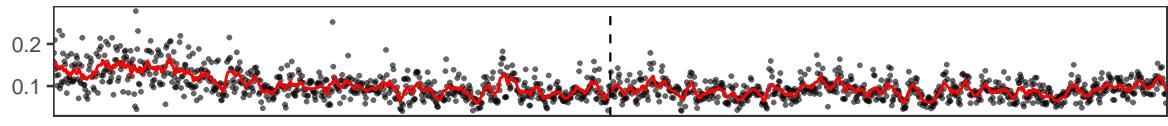
Overlapping models: Selection–bottleneck + Genomic architecture

Overlap signals: Xingu_enrichment (high); Belem_enrichment (high)

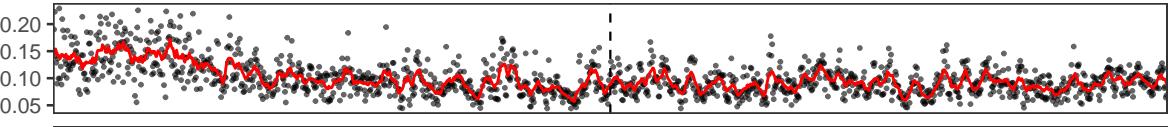
Xingu–Belem Fst



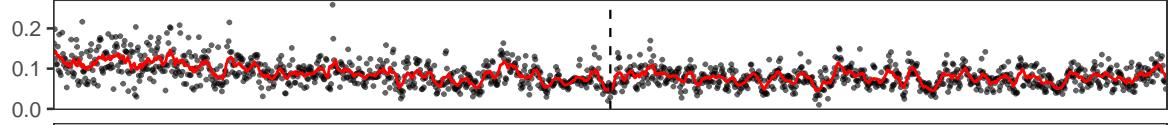
Xingu–Belem Dxy



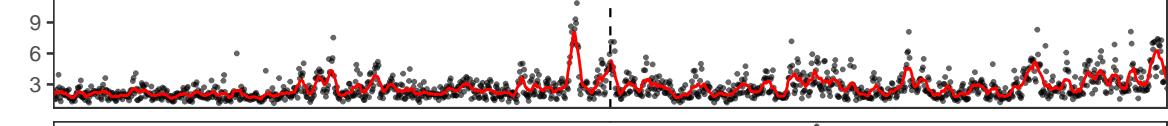
Xingu Pi



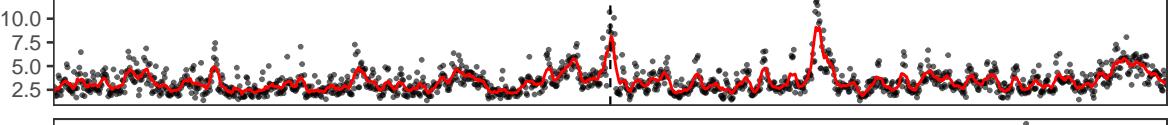
Belem Pi



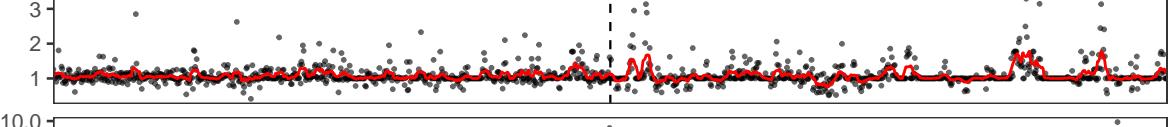
Xingu enrichment



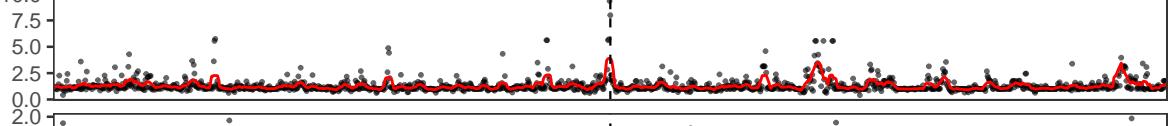
Belem enrichment



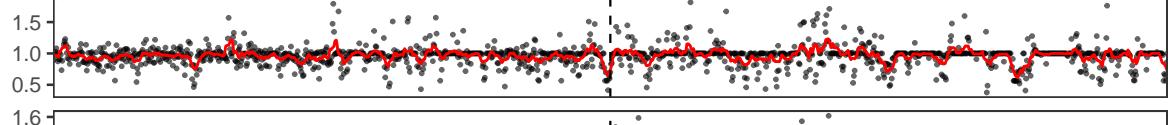
Xingu RTH (inverse)



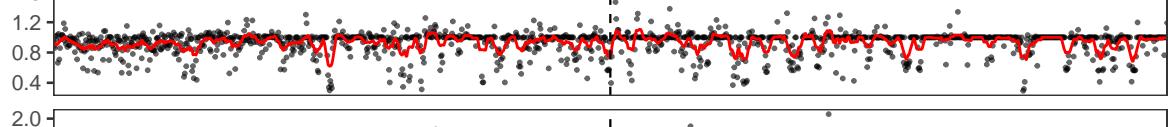
Belem RTH (inverse)



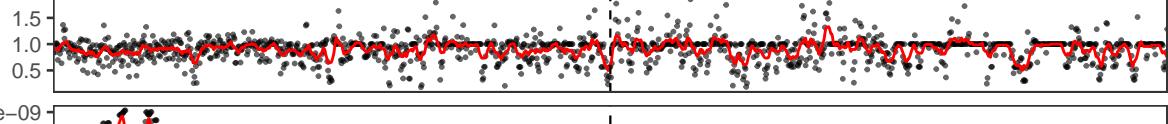
Xingu–Belem RTH



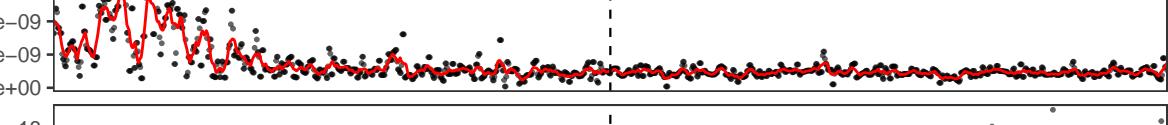
Tapajos–Xingu RTH (inverse)



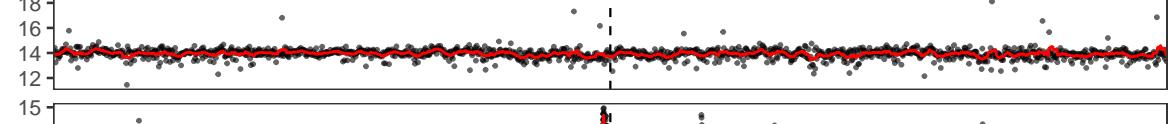
Tapajos–Xingu JCR



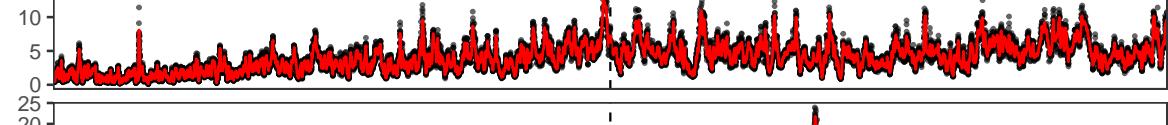
Recombination rate



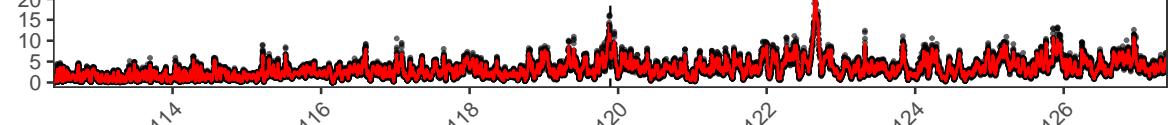
Average depth



Xingu RAiSD u



Belem RAiSD u



114 116 118 120 122 124 126 128

Position on Chromosome_1_RagTag (Mb)