

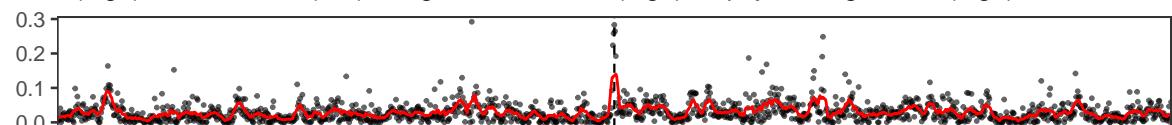
Peak: Chromosome\_3\_RagTag:36680000–36730000

Model: Overlapping models

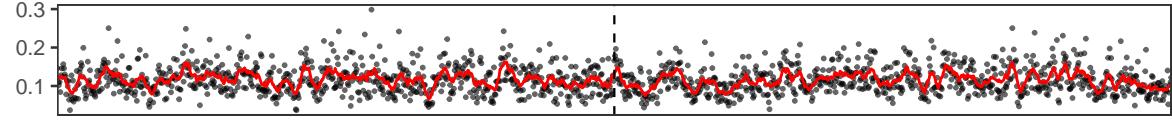
Overlapping models: Selection–bottleneck + Introgression + Deep lineage sorting

Overlap signals: Belem\_enrichment (high); Belem\_RTH (low); Xingu\_Belem\_RTH (high); Tapajos\_Xingu\_JCR (high)

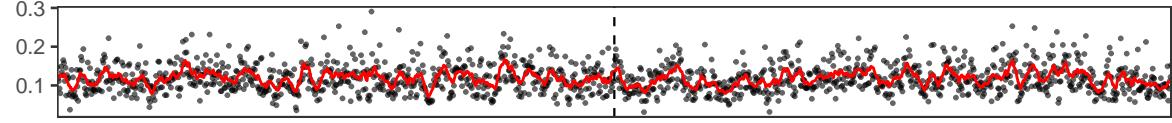
Xingu–Belem Fst



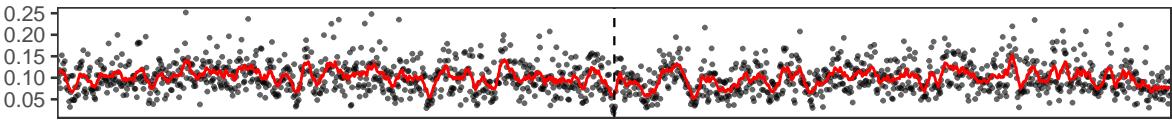
Xingu–Belem Dxy



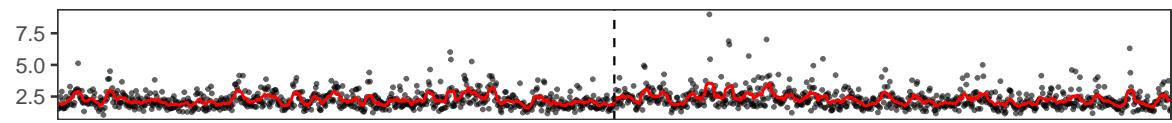
Xingu Pi



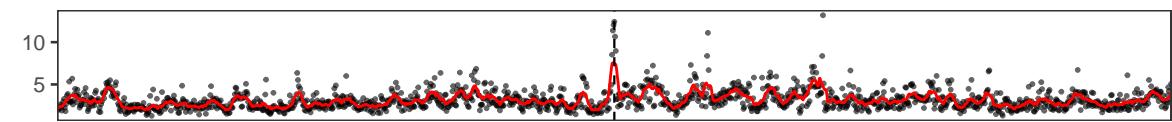
Belem Pi



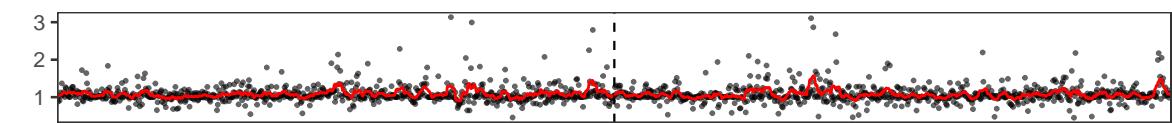
Xingu enrichment



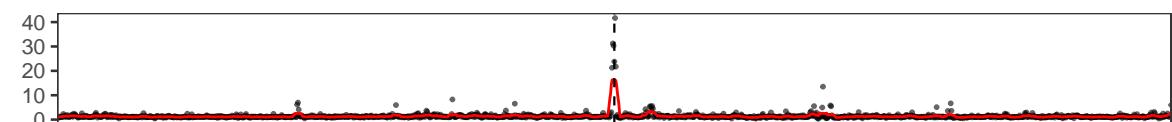
Belem enrichment



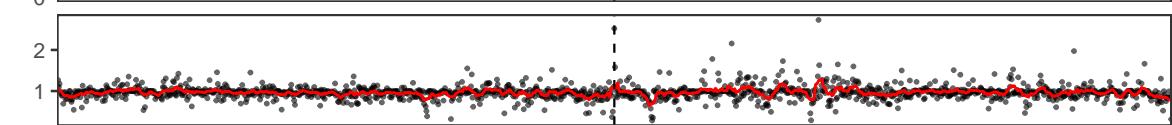
Xingu RTH (inverse)



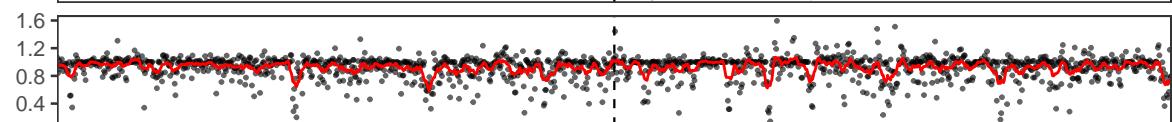
Belem RTH (inverse)



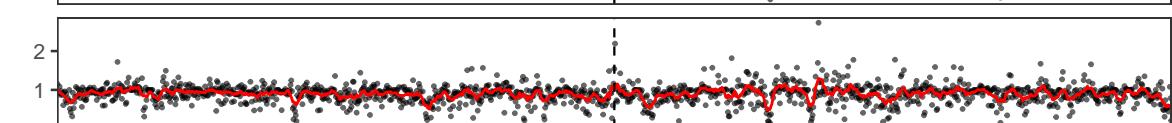
Xingu–Belem RTH



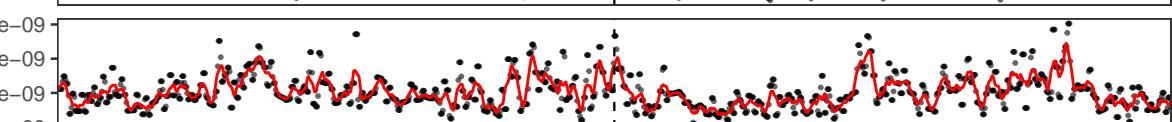
Tapajos–Xingu RTH (inverse)



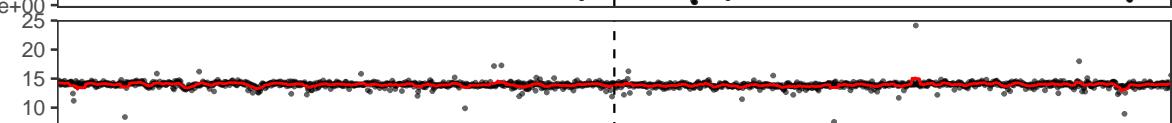
Tapajos–Xingu JCR



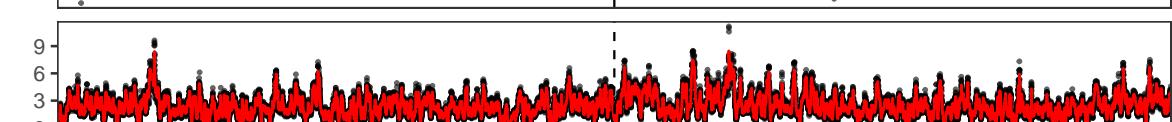
Recombination rate



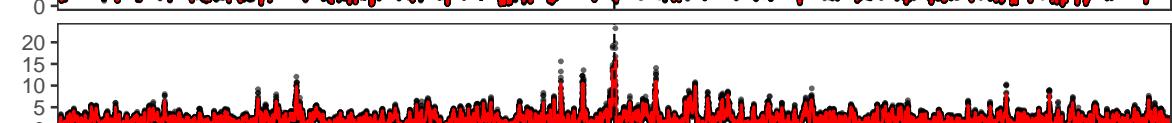
Average depth



Xingu RAiSD u



Belem RAiSD u



Position on Chromosome\_3\_RagTag (Mb)