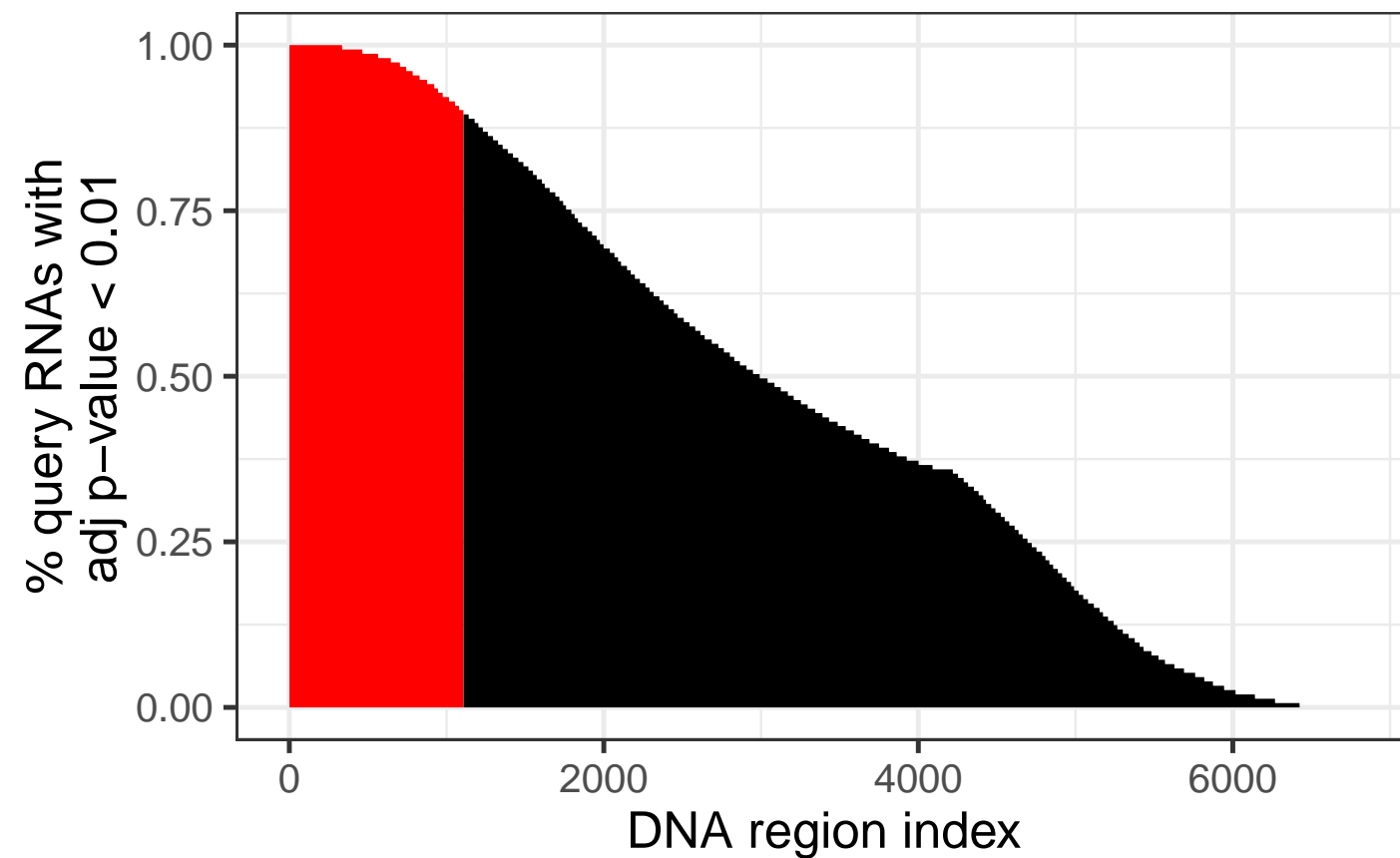


a

ChOP-seq peaks

Universal TTS
(n = 1107)

Other regions
(n = 5691)



b

Control DNA regions

Universal TTS
(n = 41)

Other regions
(n = 6757)

