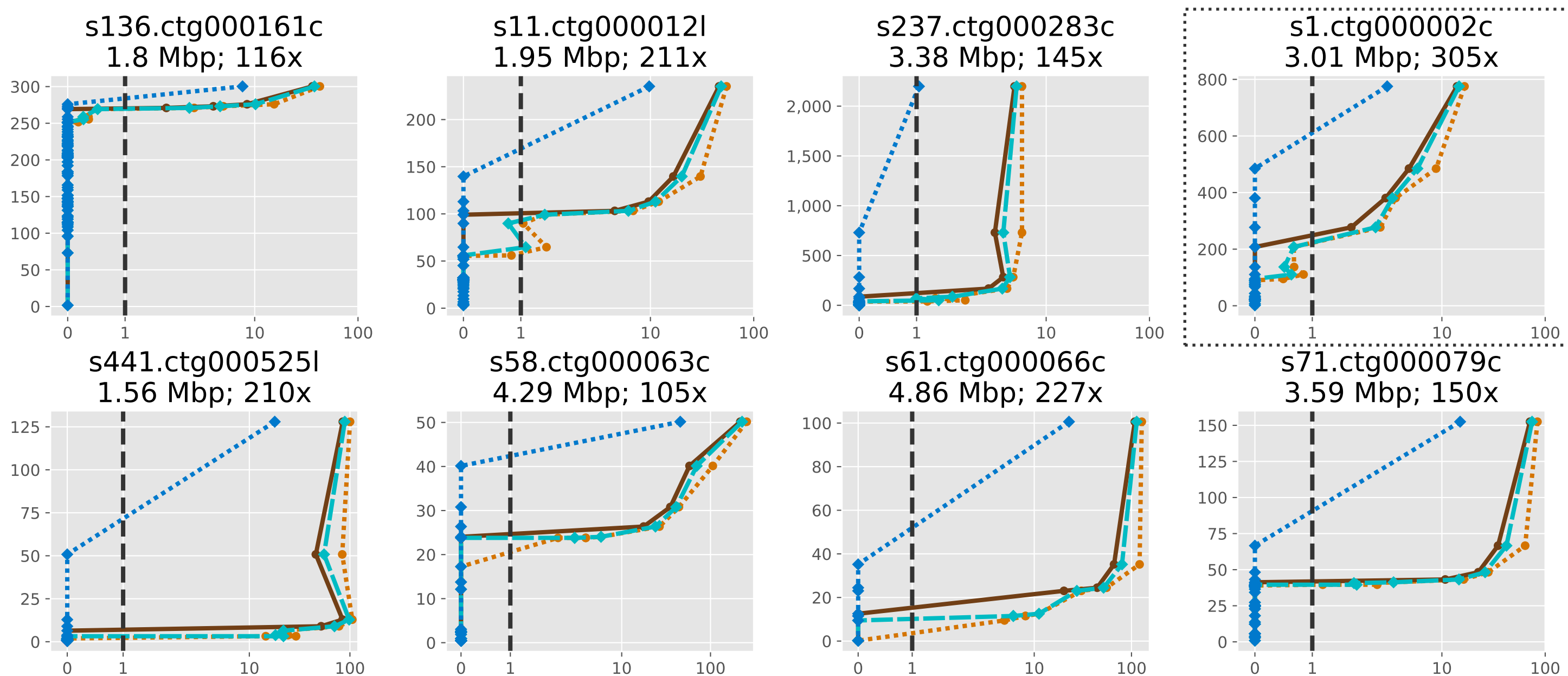
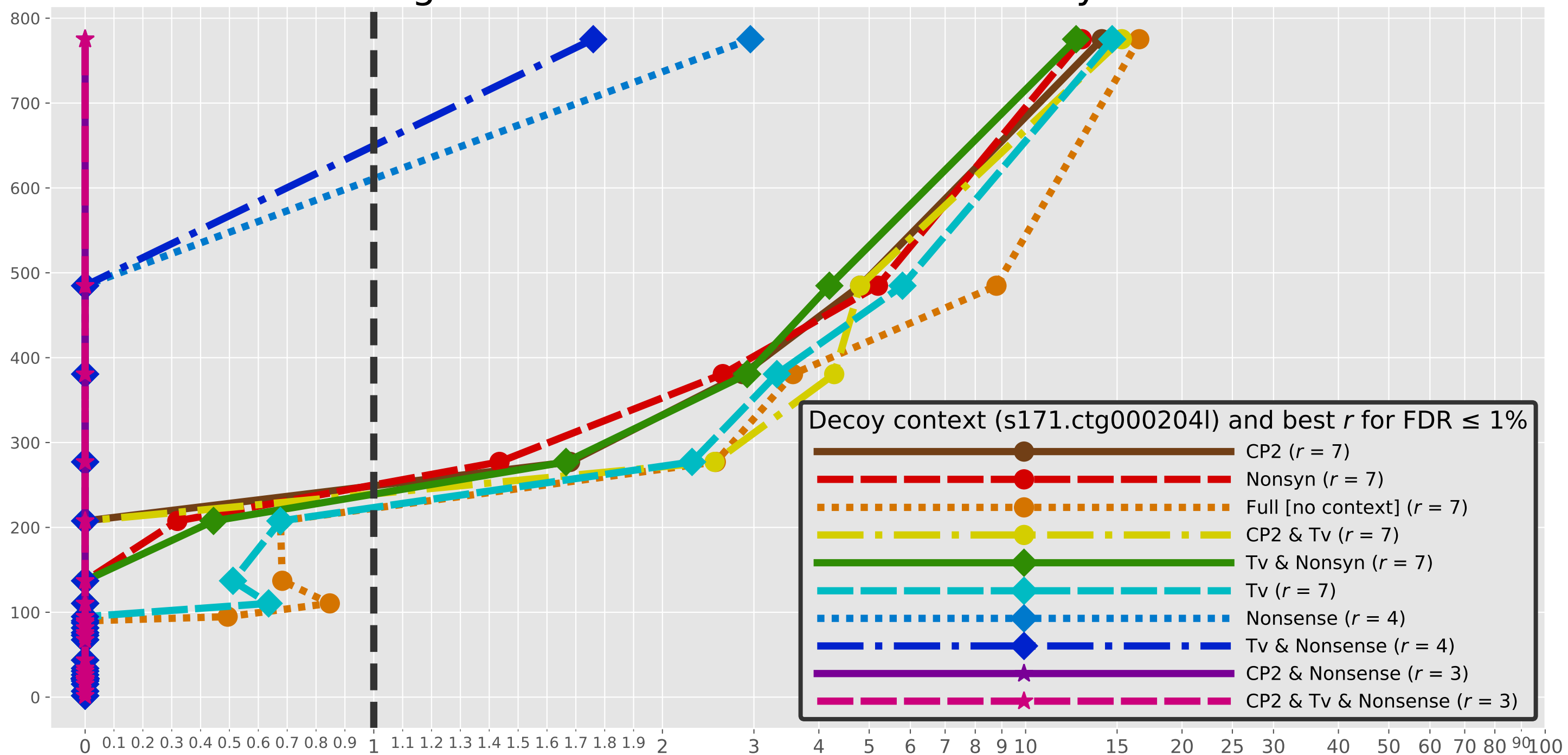


FDR curves for naïve r -mutation calling ($r = 99$ to $r = 3$), using the 8 highest-diversity-index ($r = 50$) target contigs ≥ 1 Mbp

Number of called rare ($r < 100$) r -mutations per megabase



s1.ctg000002c FDR curves for 10 decoy contexts



Estimated FDR of called rare ($r < 100$) r -mutations (%),
using a \log_{10} scale to highlight order-of-magnitude differences