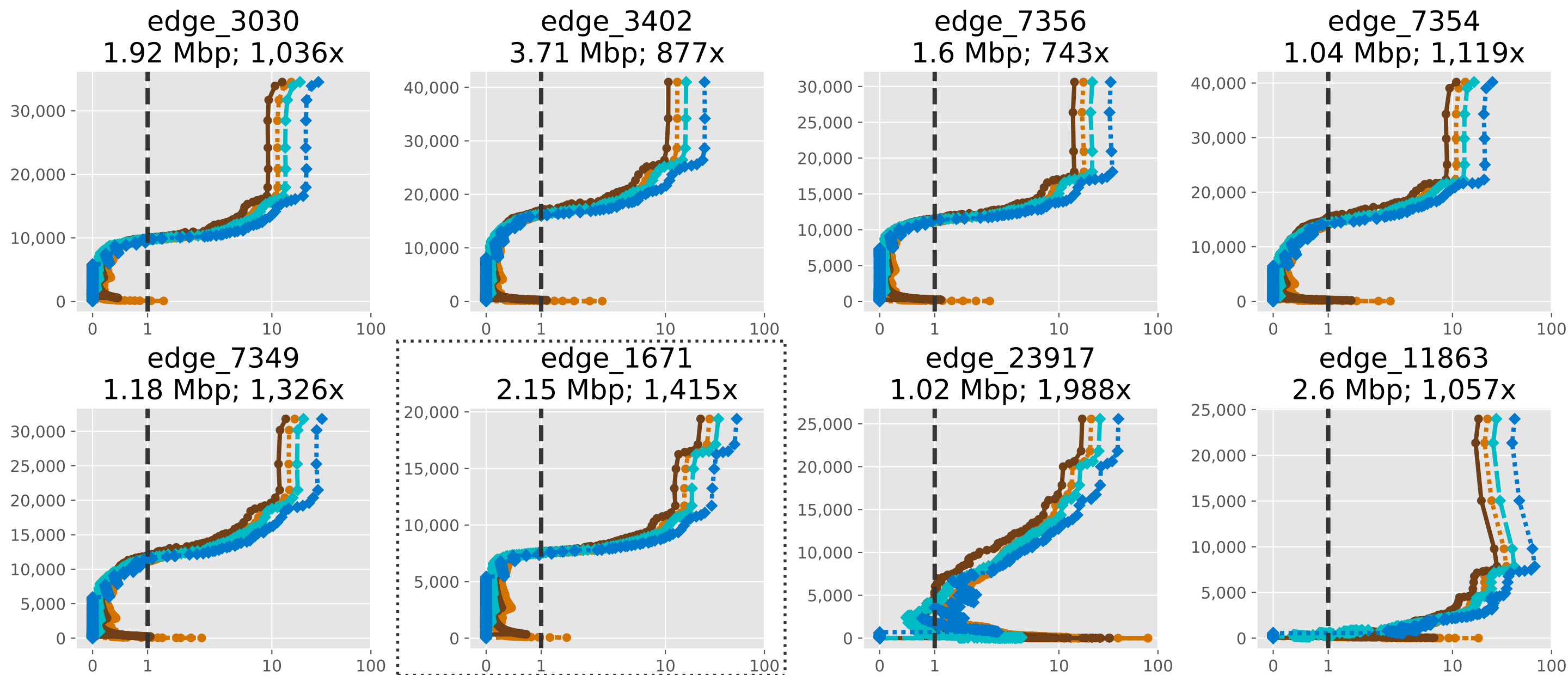
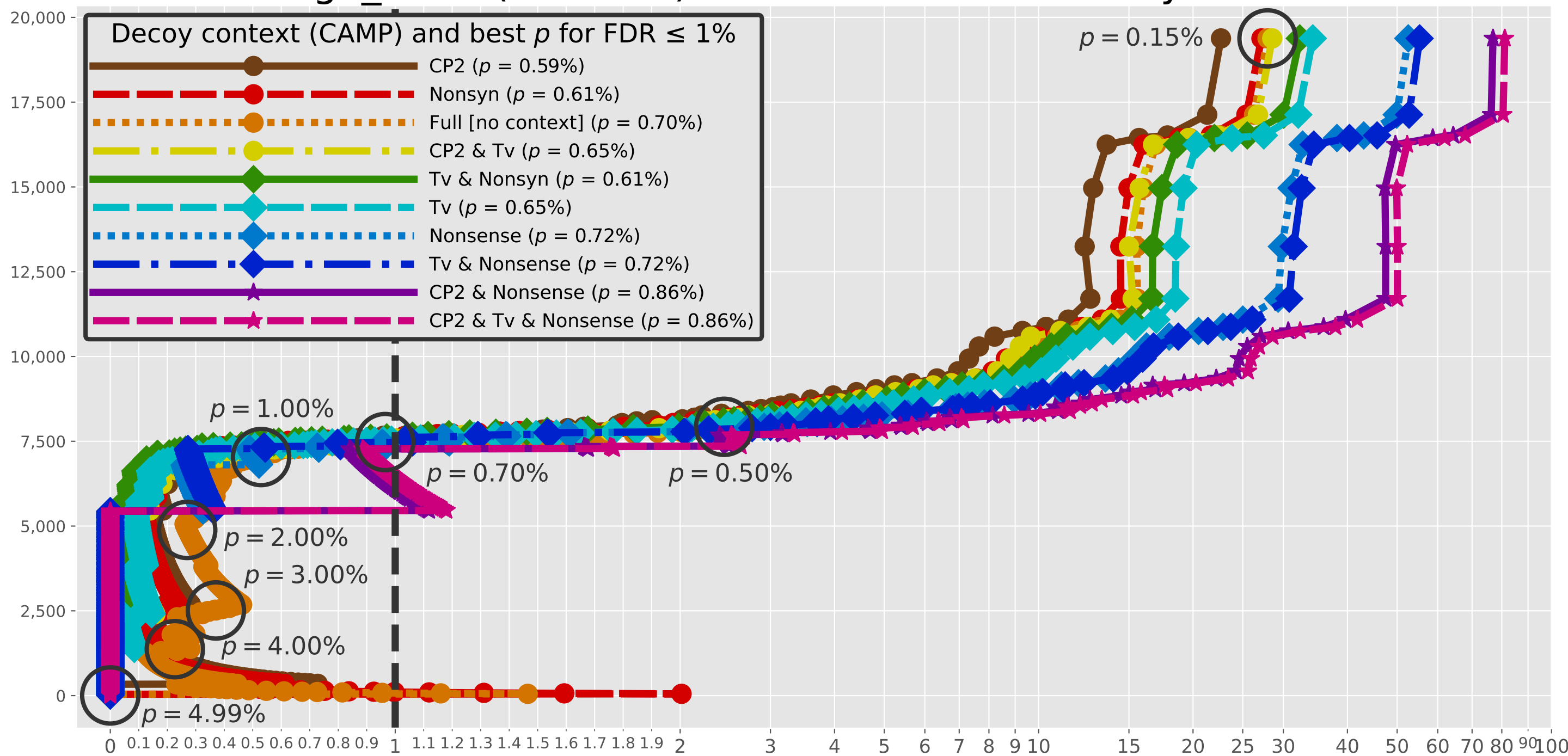


FDR curves for naïve p -mutation calling ($p = 4.99\%$ to $p = 0.15\%$),
using the 8 highest-diversity-index ($p = 0.5\%$) target contigs ≥ 1 Mbp

Number of called rare ($p < 5\%$) p -mutations per megabase



edge_1671 ("BACT1") FDR curves for 10 decoy contexts



Estimated FDR of called rare ($p < 5\%$) p -mutations (%),
using a \log_{10} scale to highlight order-of-magnitude differences