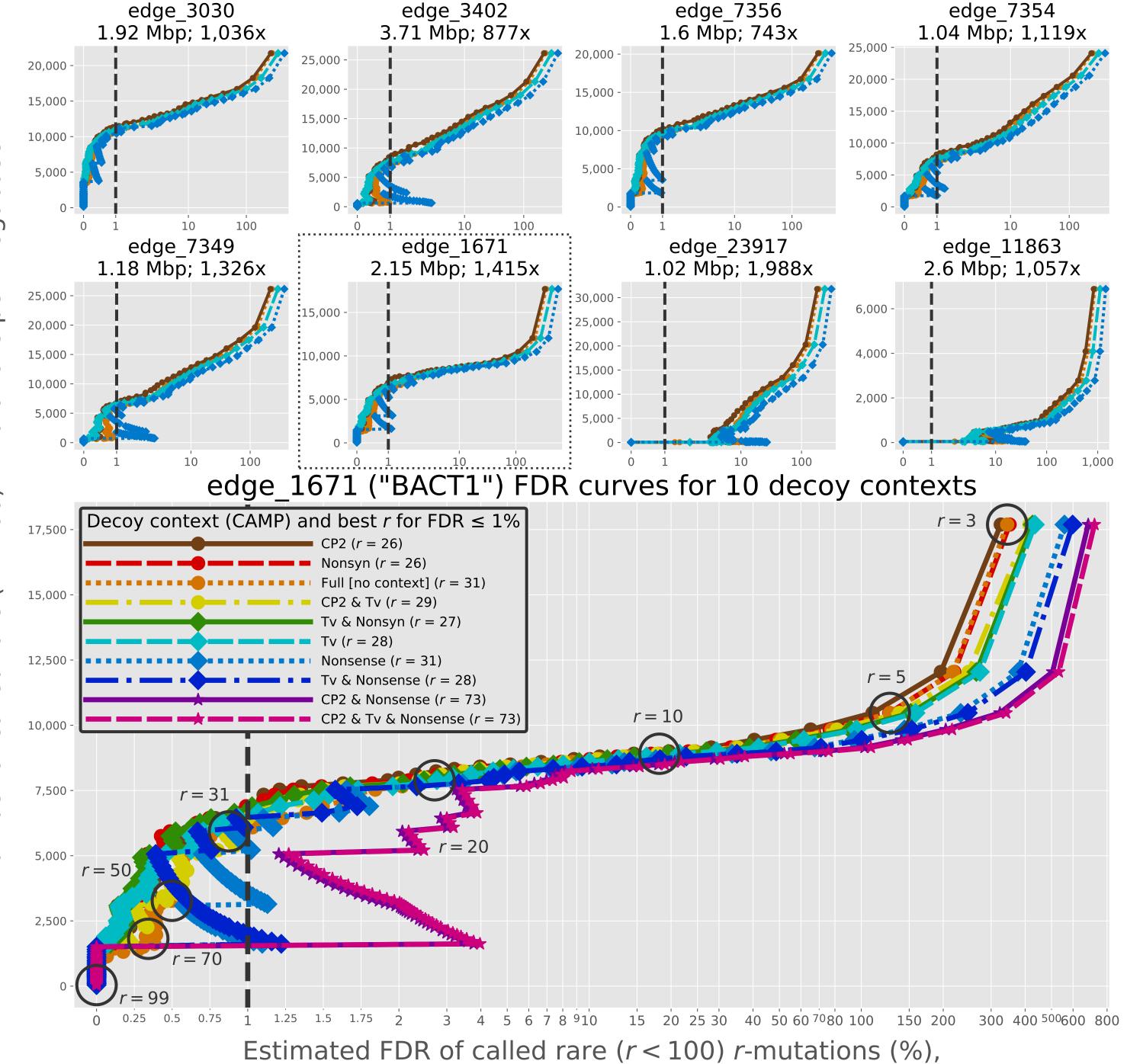
FDR curves for naïve r-mutation calling (r = 99 to r = 3), using the 8 highest-diversity-index (p = 0.5%) target contigs  $\geq 1$  Mbp



using a log<sub>10</sub> scale to highlight order-of-magnitude differences