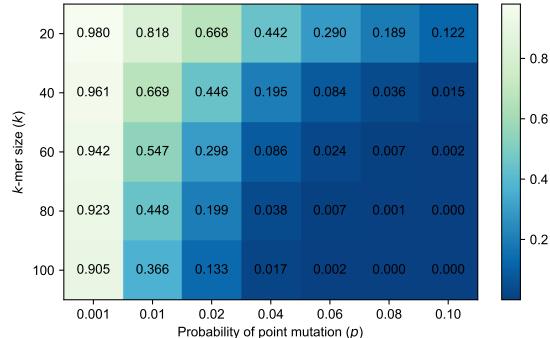
Expected containment across *k*-mer sizes and mutation rates



k-mers of in the original

Containment of mutated sequence