



Figure S2. Patterns of base-substitution mutation (bpsm) rates at various size intervals extending clockwise from the origin of replication (*oriC*), in MMR-deficient mutation accumulation lineages of *Vibrio fischeri* (A) and *Vibrio cholerae* (B) on chromosome 2. All interval breakpoints are plotted relative to the initiation of replication of *oriC* so that the boundaries of the intervals are at identical locations.