



Figure S5 Sequence logos of novel motifs in promoters of homologous recombination-related genes found by MEME. (a), E-value: 7.2×10^{-5} ; found within 11 promoters. (b), E-value: 2.9×10^1 ; found within ten promoters. (c), E-value: 3.7×10^1 ; found within 13 promoters.