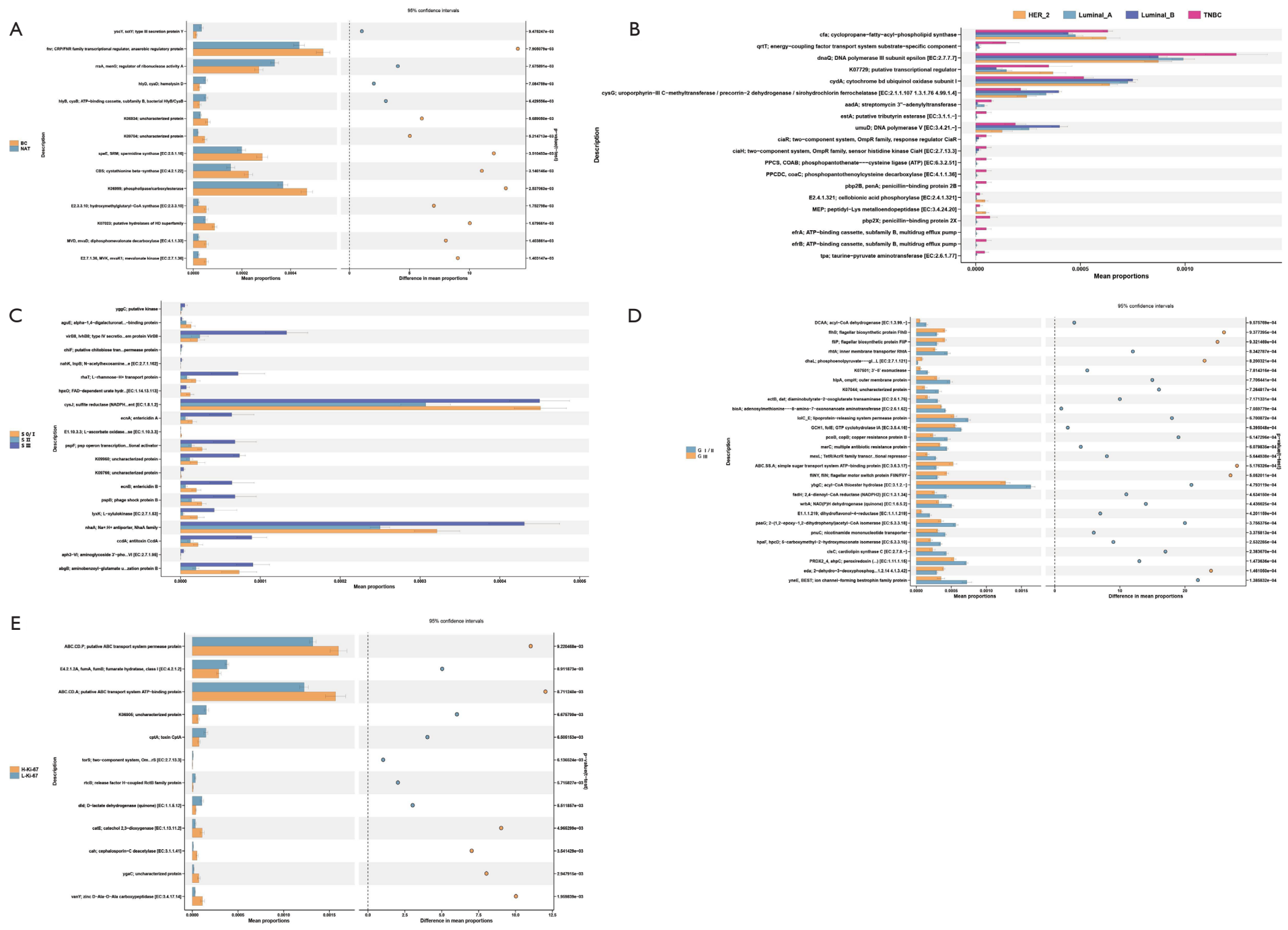

Figure S1 Differential KO gene analysis based on microbial functional prediction between BC and NAT, and among BC clinical and pathological subgroups. (A) The KO genes that differed significantly between the BC and NAT groups (P≤0.01). (B) The KO genes that differed significantly among different molecular subtypes (P<0.01). (C) The KO genes that differed significantly across clinical stages (P<0.01). (D) The KO genes that differed significantly between different histological grade groups (P<0.01). (E) The KO genes that differed significantly between different Ki-67 groups (P≤0.01). BC, breast cancer; KEGG, Kyoto Encyclopedia of Genes and Genomes; KO, KEGG Orthology; NAT, normal adjacent tissue.

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