

Figure S2. The enrichments of DE lncRNAs in Min dead (MD) vs. Min control (MC) groups. (A) KEGG pathway enriched by DE lncRNAs with significant up-regulation expression in MD vs. MC. (B) GO entries enriched by DE lncRNAs with significant up-regulation expression in MD vs. MC. (C) KEGG pathway enriched by DE lncRNAs with strong down-regulation expression in MD vs. MC. (D) GO entries enriched by DE lncRNAs with strong down-regulation expression in MD vs. MC.