

Figure S1. Reverse transcription-quantitative PCR analysis of circ-CCT2, circ-TRIM33, circ-EIF4G3 and circ-LRP6 expression levels in osteosarcoma (tumor) and normal paracancerous tissues. **P<0.01, ***P<0.001. CCT2, chaperonin-containing TCP1 subunit 2; circ, circular RNA; EIF4G3, circ-eukaryotic translation initiation factor 4 γ 3; LRP, lipoprotein receptor 6; TRIM33, tripartite motif-containing 33.

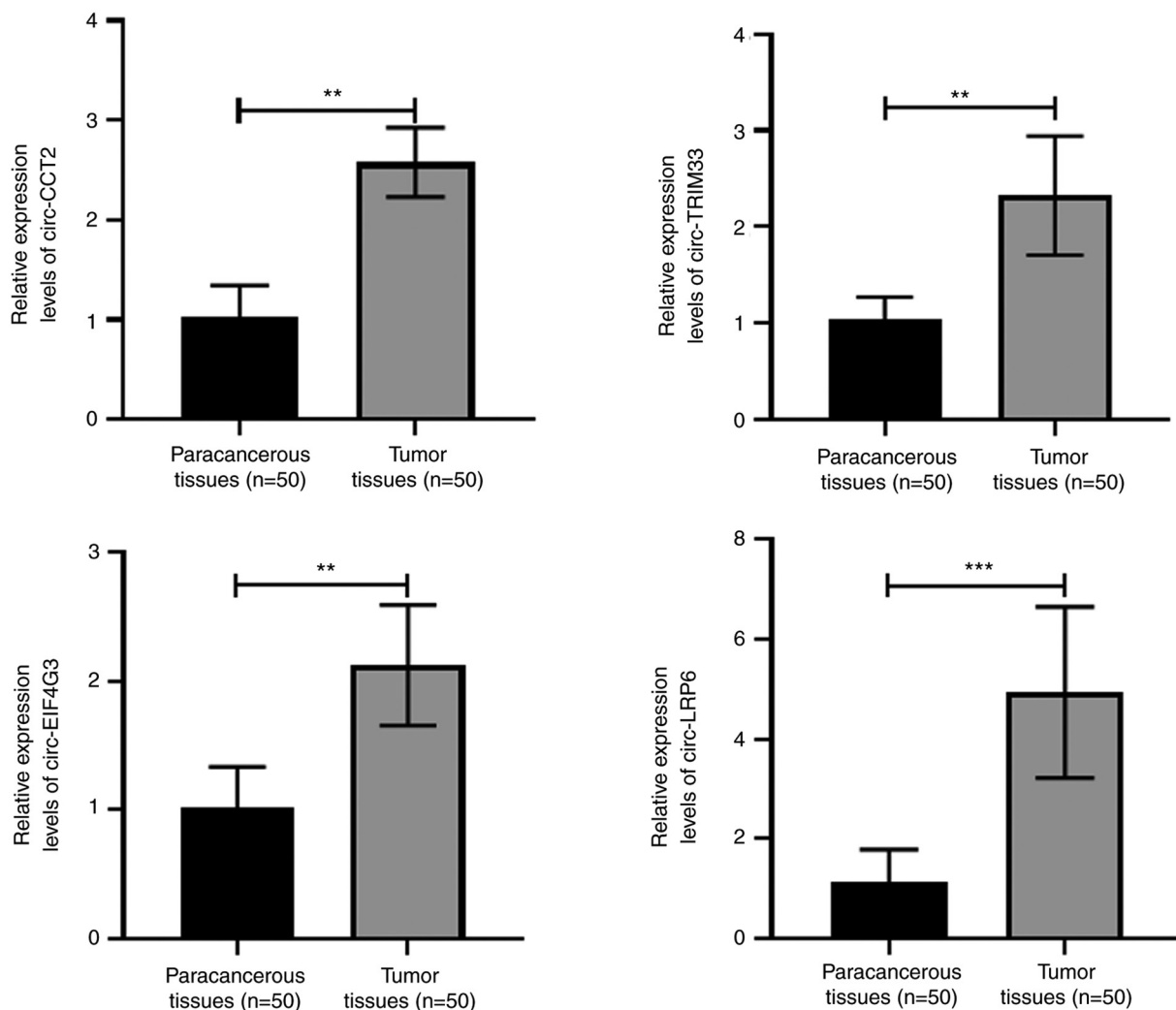


Figure S2. Circular RNA Interactome online tool prediction of four of the top five miRNAs that might interact with circ-LRP6. (A) Binding sites between circ-LRP6 and miR-1208. (B) Binding sites between circ-LRP6 and miR-326. (C) Binding sites between circ-LRP6 and miR-330-5p. (D) Binding sites between circ-LRP6 and miR-513a-3p. circ, circular RNA; LRP, lipoprotein receptor 6; miR, microRNA.

