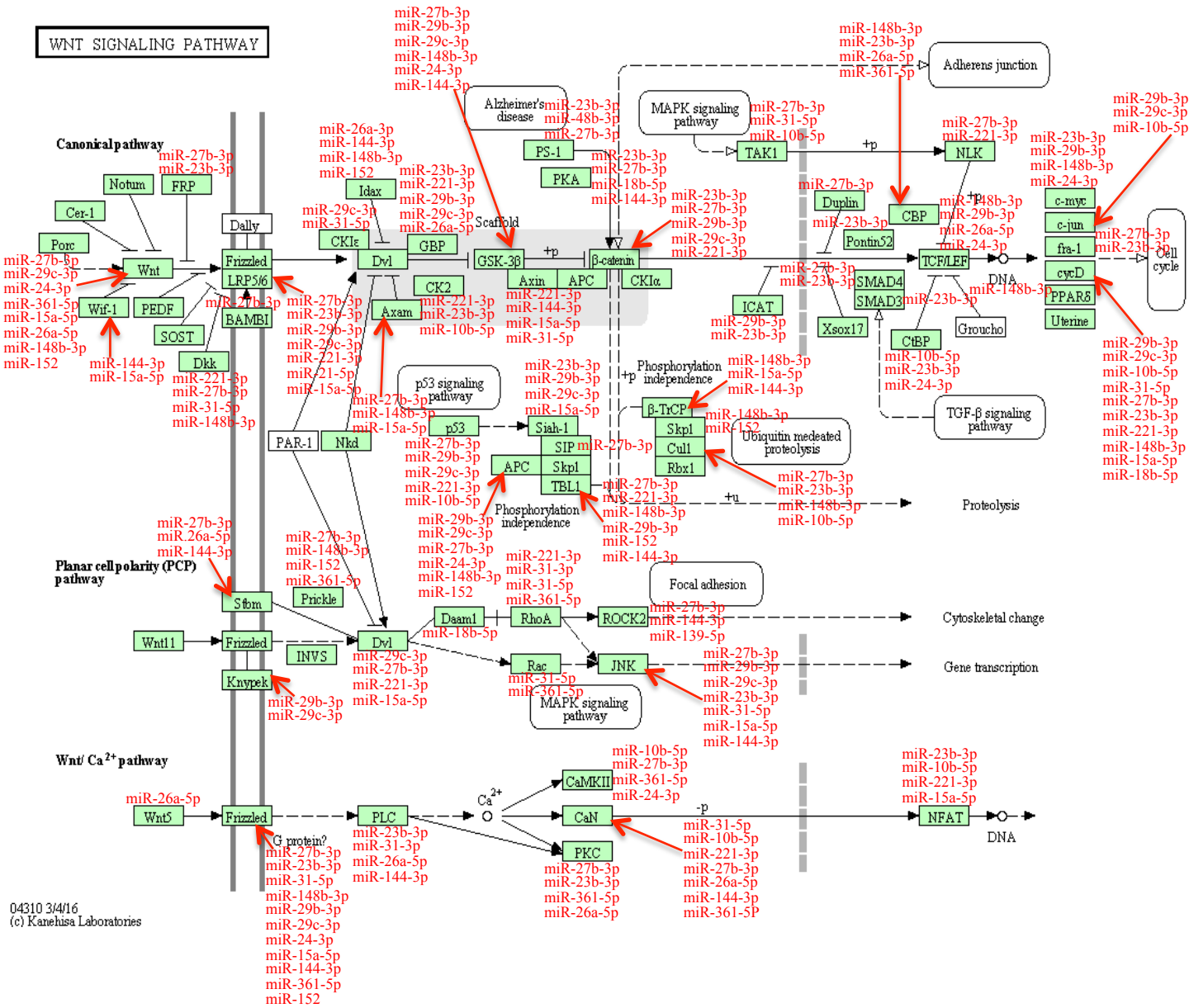

S3 Fig. Schematic diagram of Wnt signaling pathway enriched by differentially expressed microRNAs in Exo_D21 compared with Exo_P6 and Exo_D3. microRNAs, predicted and/or experimentally validated to participate in Wnt signaling pathway through regulating their potential target genes, are marked in red. The schematic Wnt signalling pathway was adopted from KEGG database (Kanehisa, M., *et al*., *Nucleic Acids Res*, 2017) and the analysis of microRNAs (in red) participating in the pathway was analyzed using DIANA-mirPath v.3.