

Additional file 3: **Figure S3.** (A) Gene levels of MYC-driven WNT signaling genes were compared between cancer tissues and normal tissues, and binary heatmaps were drawn to show the changes. Samples were sorted with p-value, and changes with p-value<0.05 were considered statistically significant. (B) Kaplan–Meier curves comparing survival of various cancer types correlated with LEF1. Comparisons between 30% highest and 30% lowest expression were generated using ONCLnc (<http://www.oncolnc.org/>).