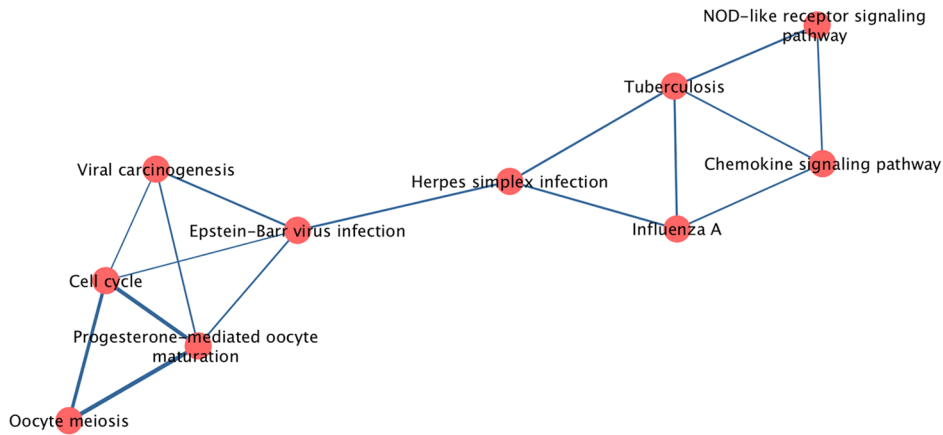
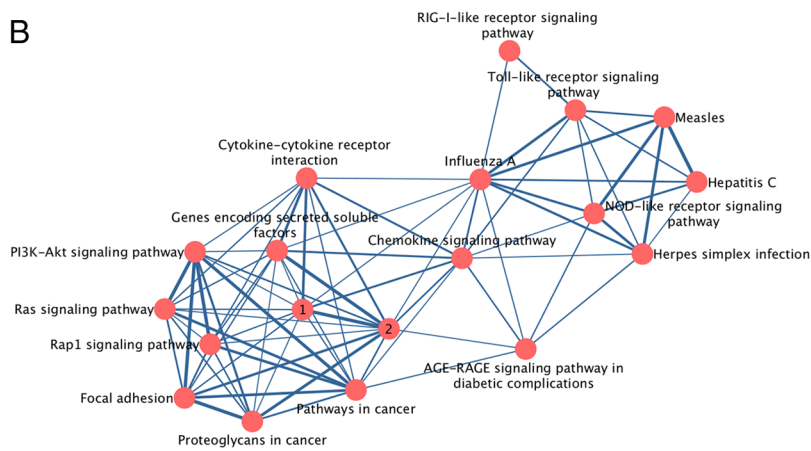


Figure S1. Pathway crosstalk analysis of phases I-IV. (A) Phase I, (B) phase II, (C) phase III and (D) phase IV. Nodes represent pathways and edge-width corresponds to the score of the specific pathway pair. Larger edge-width indicates higher score. In (B) 1 represents ‘Ensemble of genes encoding ECM-associated proteins including ECM-affiliated proteins, ECM regulators and secreted factors’ and 2 represent ‘Ensemble of gene encoding extracellular matrix and extracellular matrix-associated proteins’.

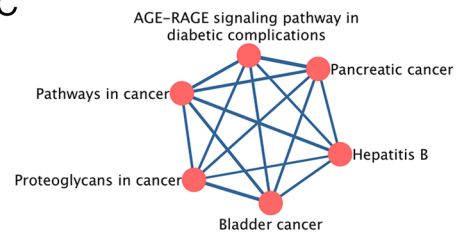
A



B



C



D

