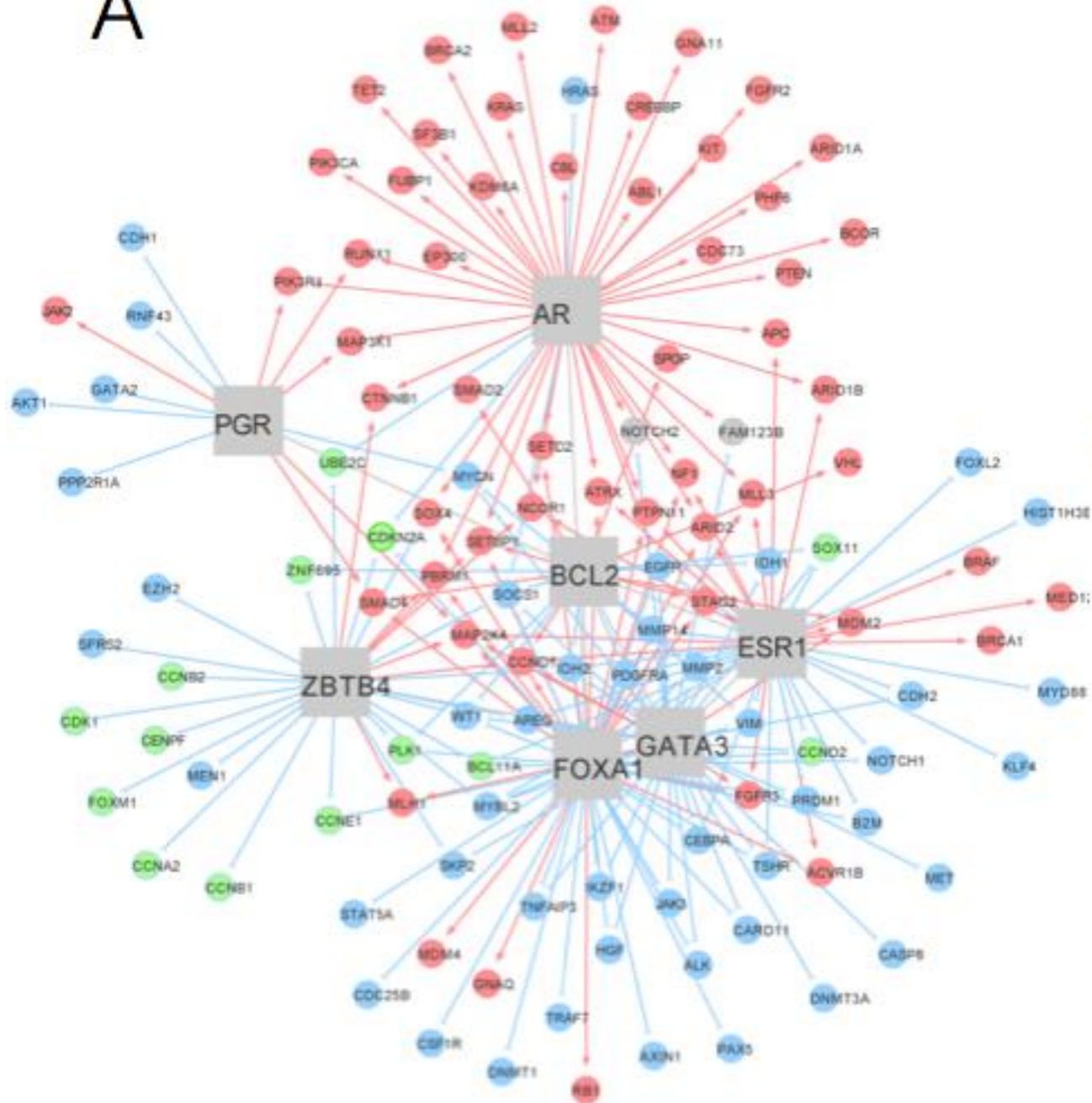


Figure S8. Subtype-specific gene regulatory networks in the TCGA cohort. We used two subtypes (luminal B and basal-like) to illustrate Subtype-specific gene networks. The circular nodes represent regulons and the edges connecting the MRs and the regulons indicate interactions between them. Red and blue colors denote up- and down-regulation, respectively. **A:** A network specific to the luminal B subtype. Green nodes are highlighted for either cell cycle genes or Group 2 MRs. Group 2 MRs, such as *ZNF695* and *SOX11*, and cell cycle genes, such as *CCNB1* and *CDK1*, are down-regulated by the central Group 1 MRs. **B:** A network specific to the basal-like subtype. The cell cycle genes, such as *PLK1* and *CDK1*, are up-regulated by the central Group 2 MRs.

Figure S8

A



B

