Supplementary Figure 2. Protein–Protein Interaction (PPI) Network of Metastasis-Associated Genes

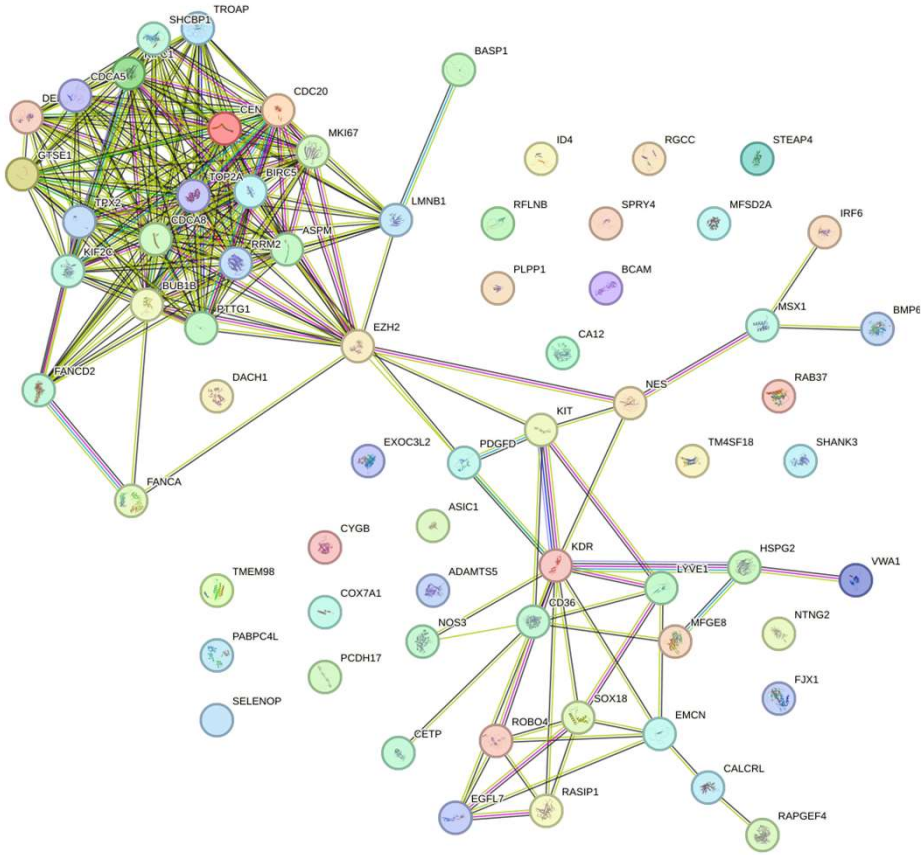


Figure S2. STRING-based protein–protein interaction (PPI) network constructed from 69 differentially expressed genes (DEGs) between M1 and M0 groups. Nodes represent proteins encoded by DEGs, and edges indicate predicted interactions with a confidence score > 0.4. A densely connected subnetwork enriched for mitotic and chromosomal regulation-related genes is visible in the upper left cluster, highlighting potential molecular drivers of metastatic progression.