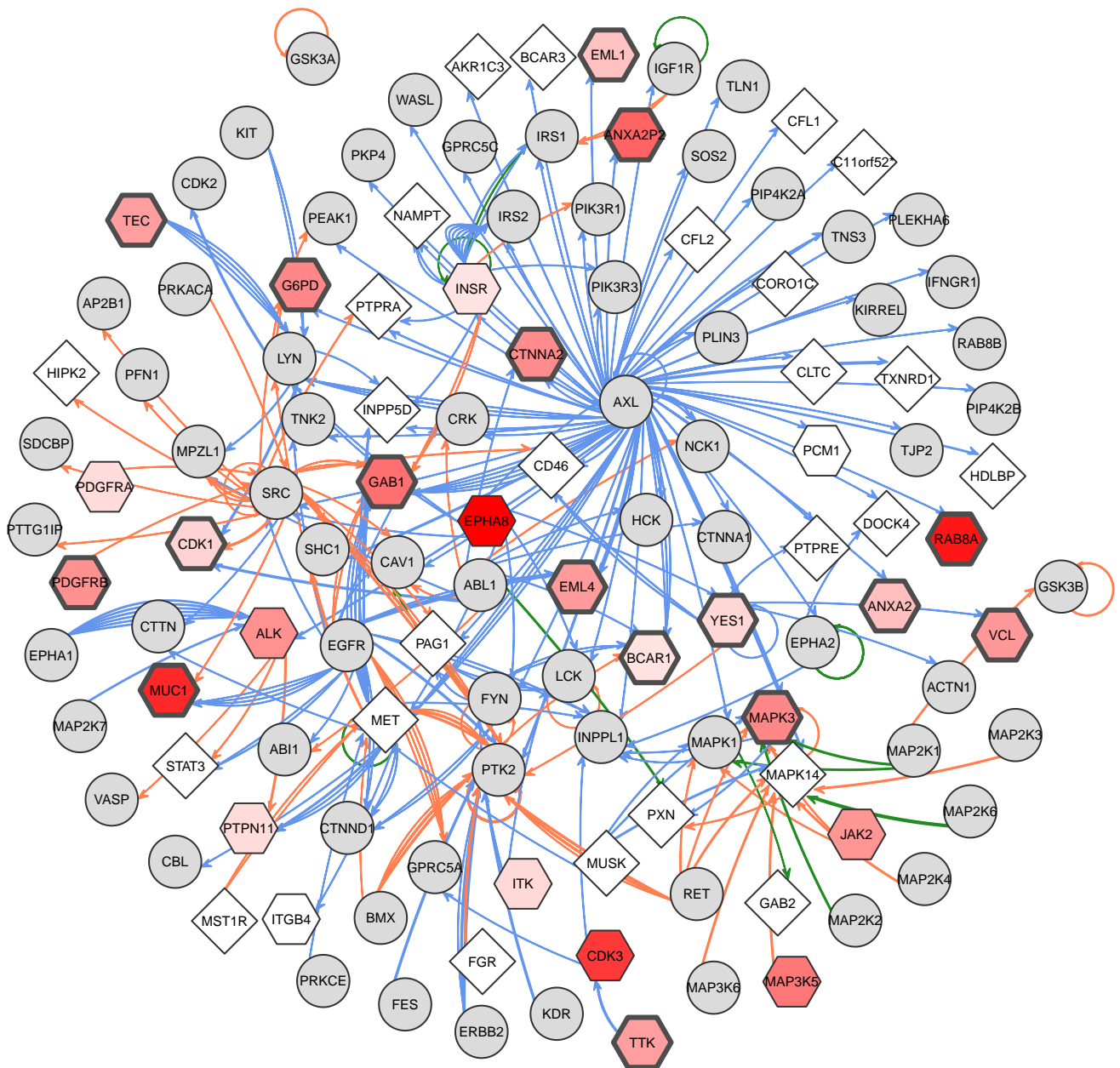
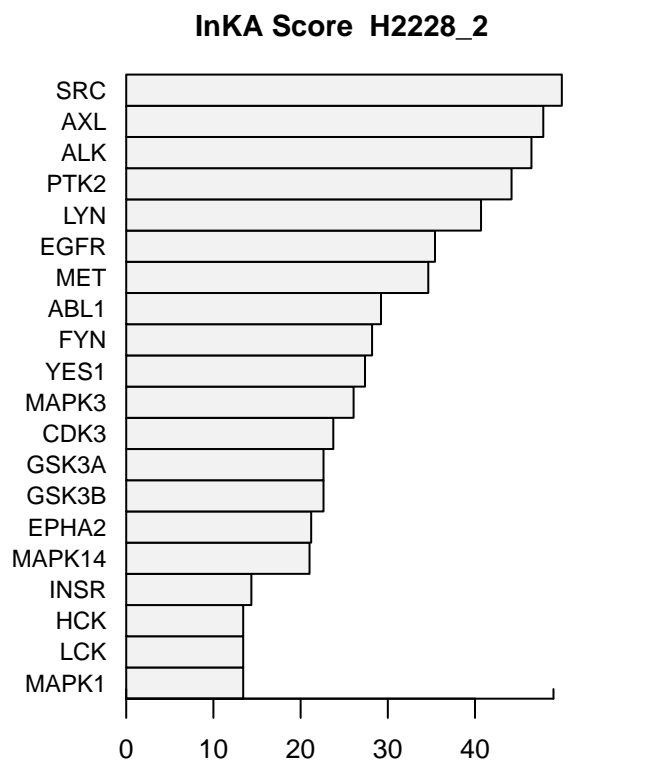
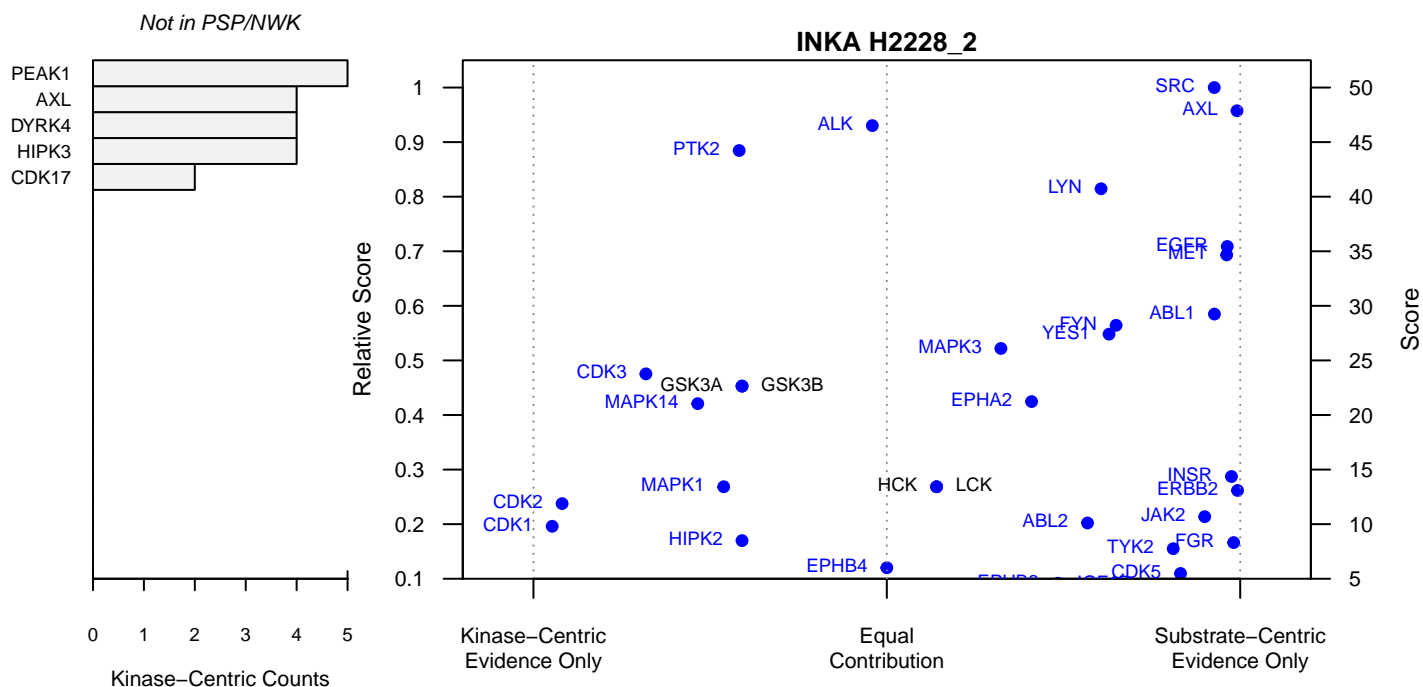
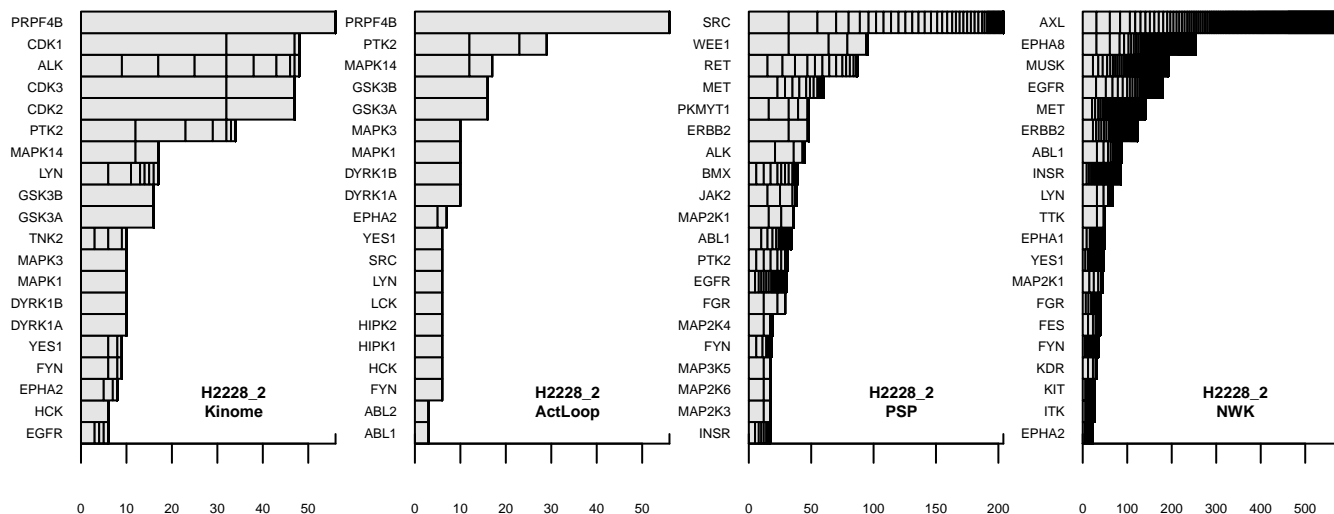
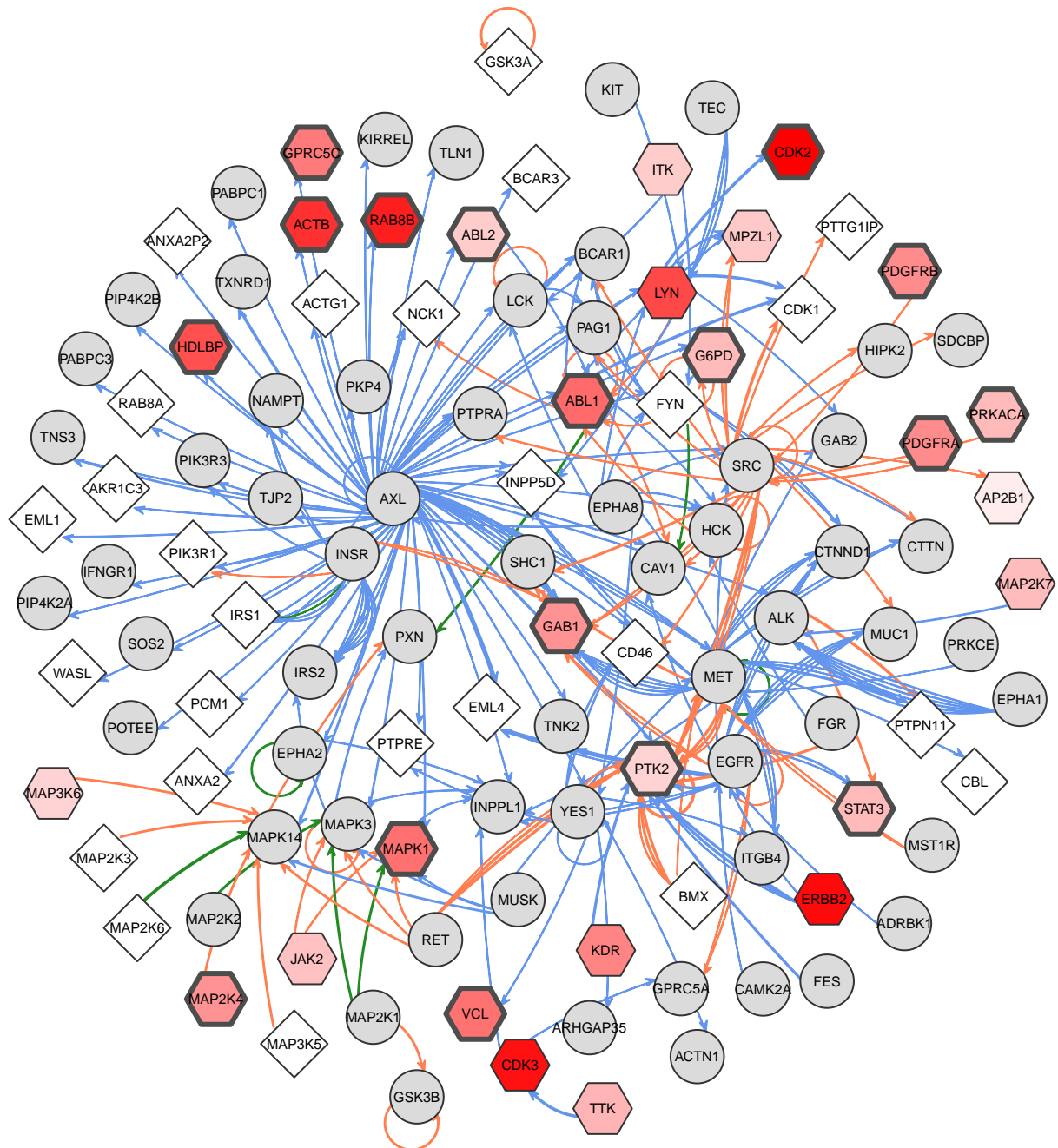


H2228_1

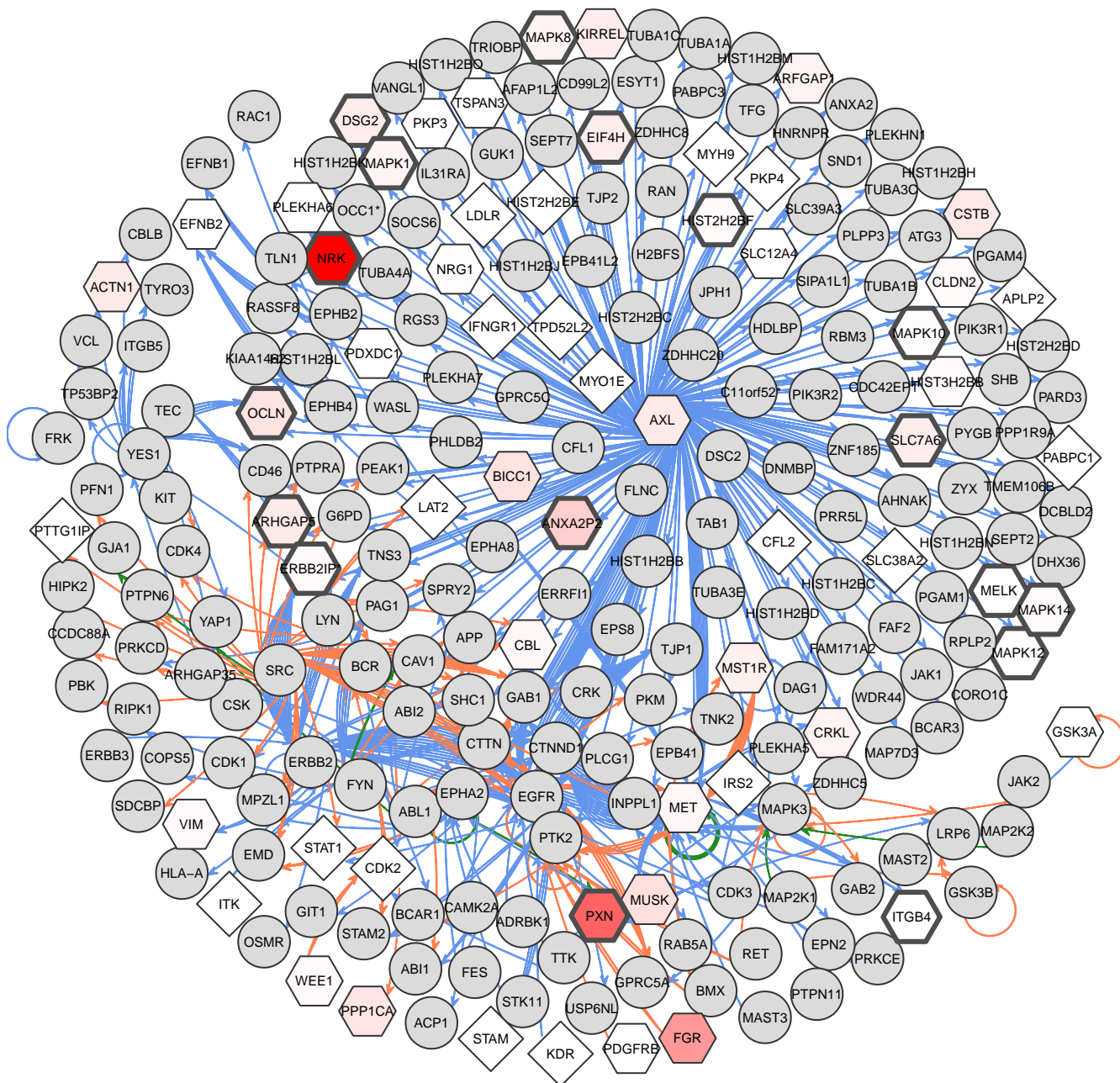


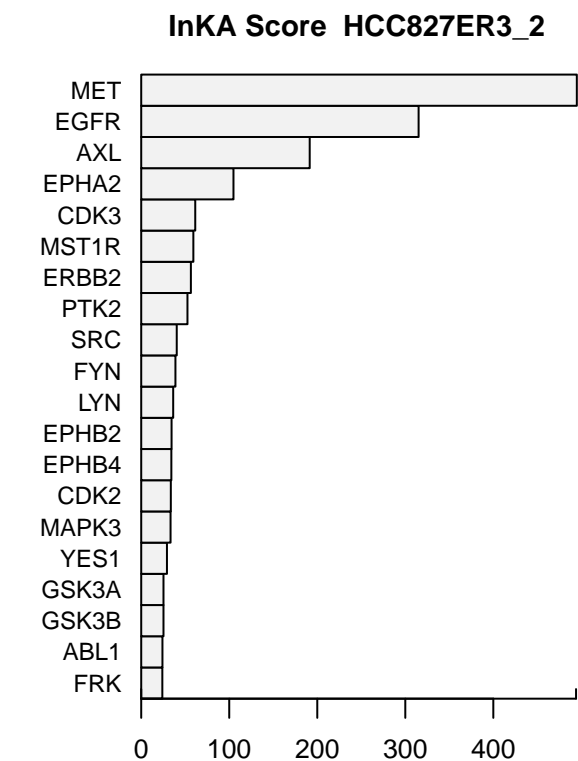
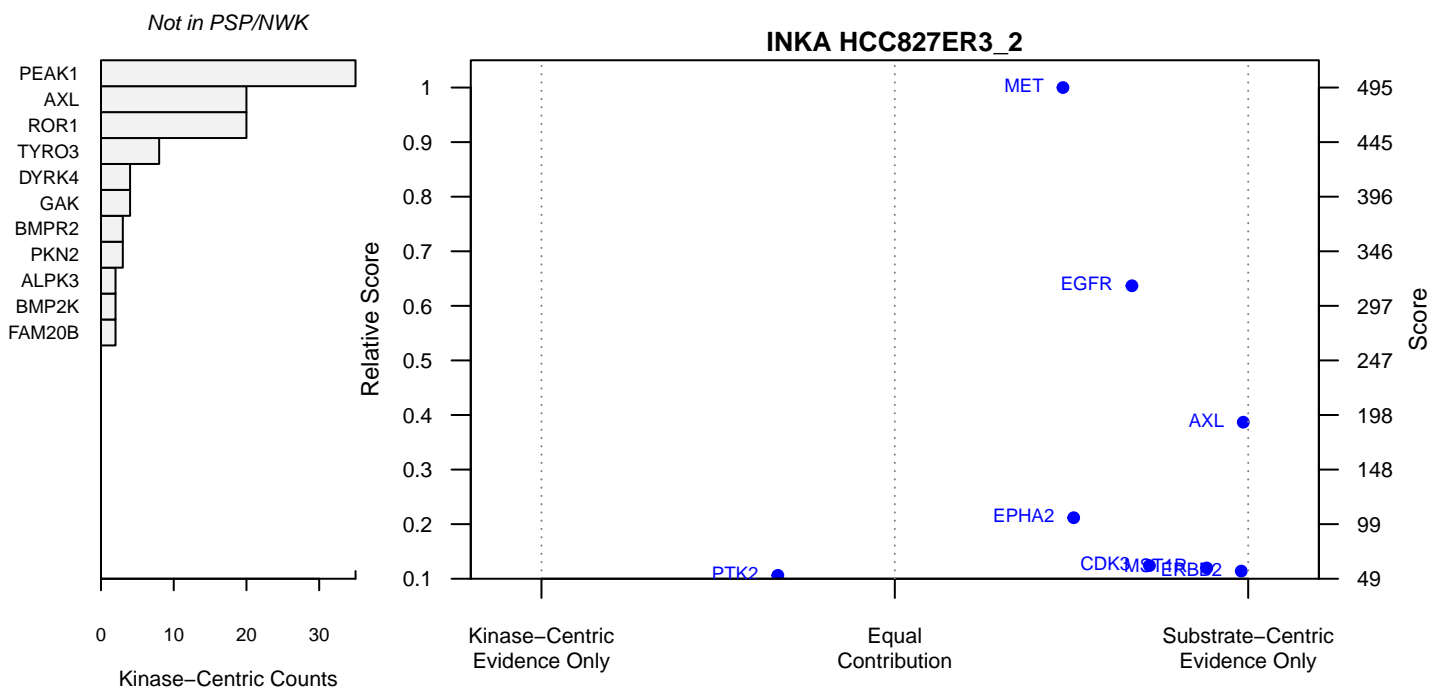
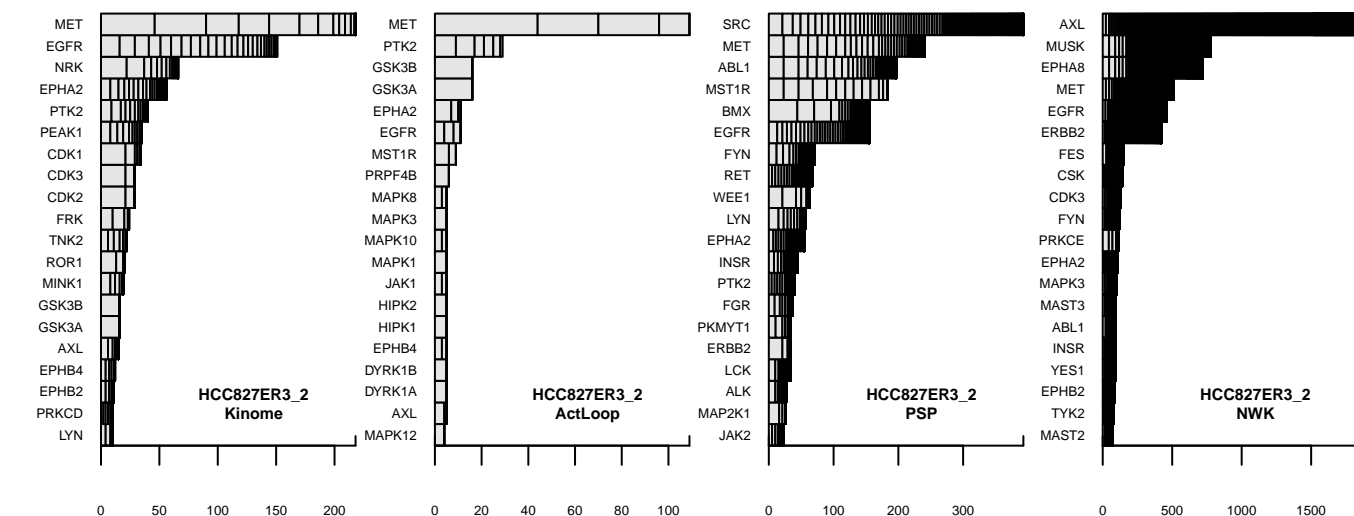


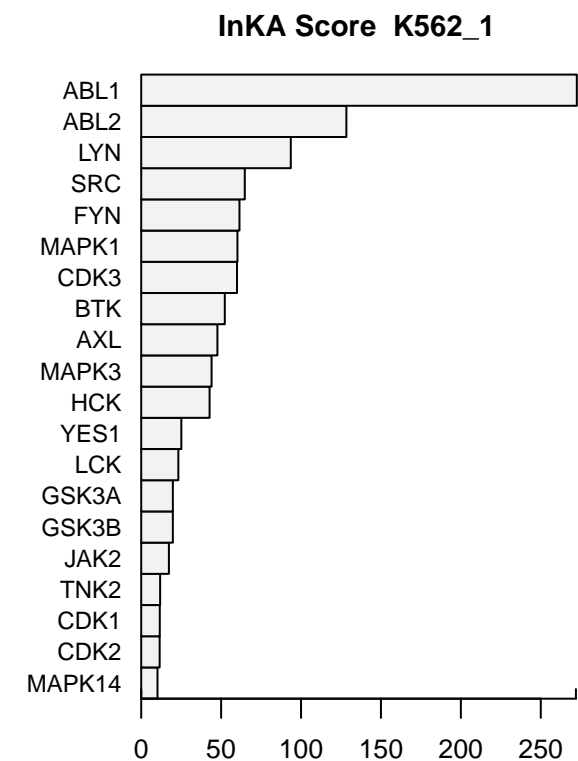
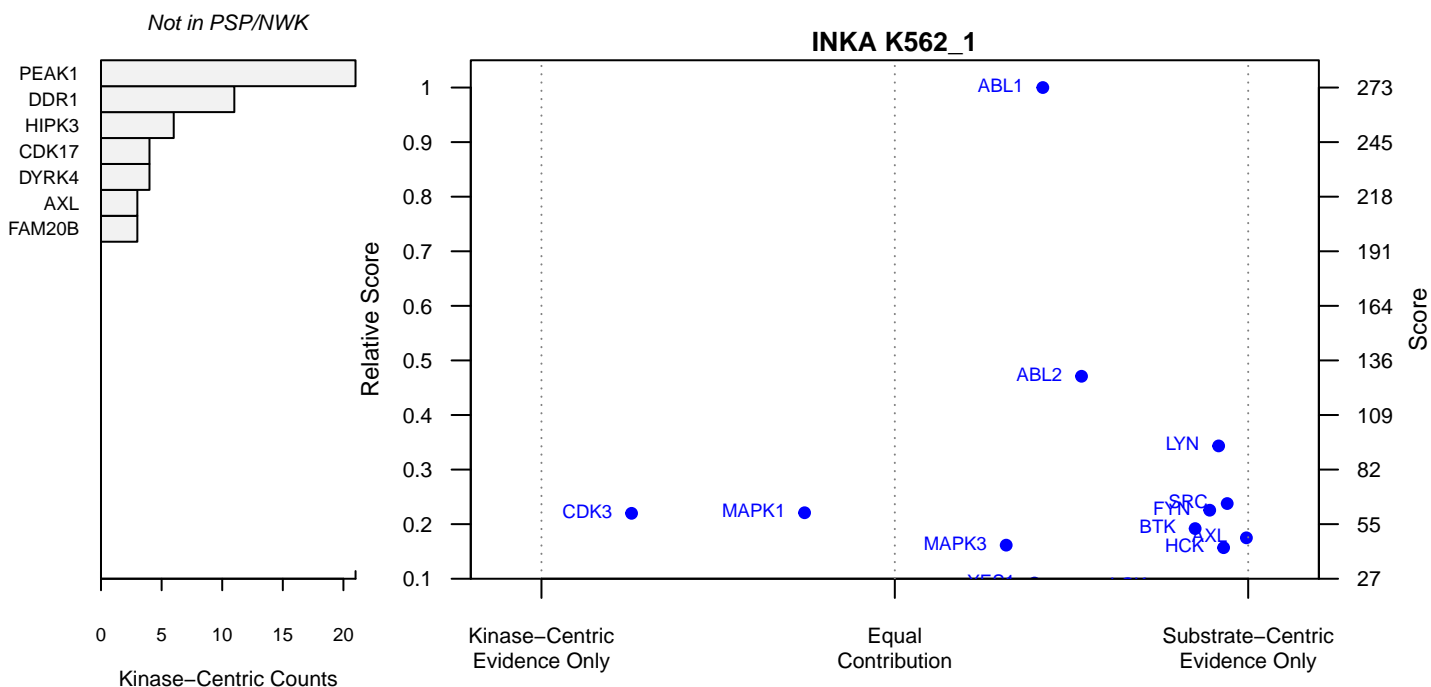
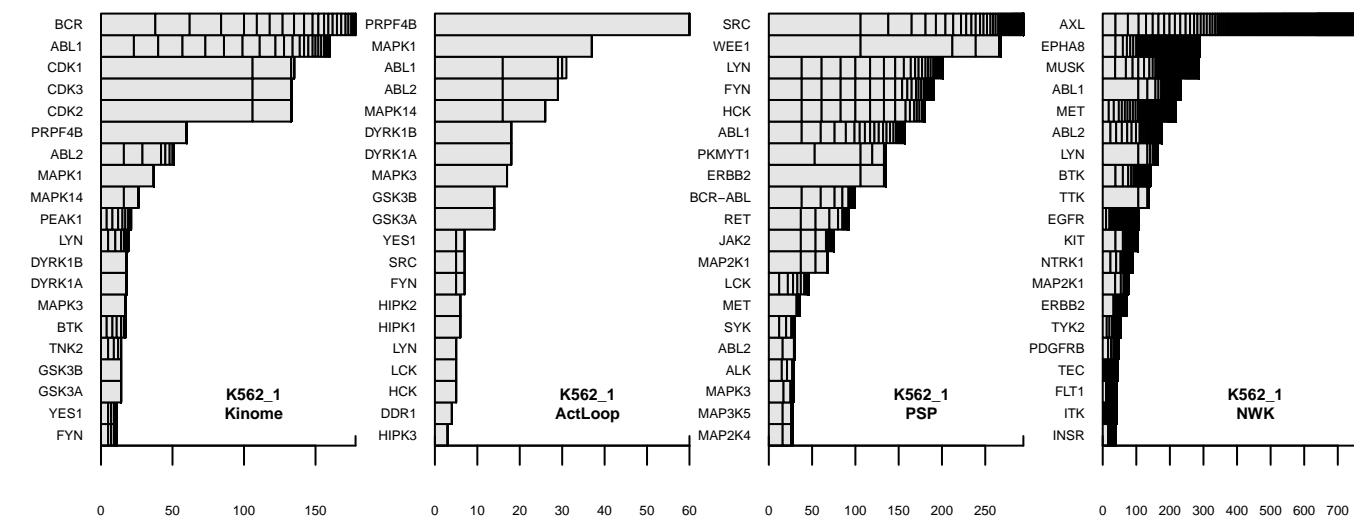
H2228_2



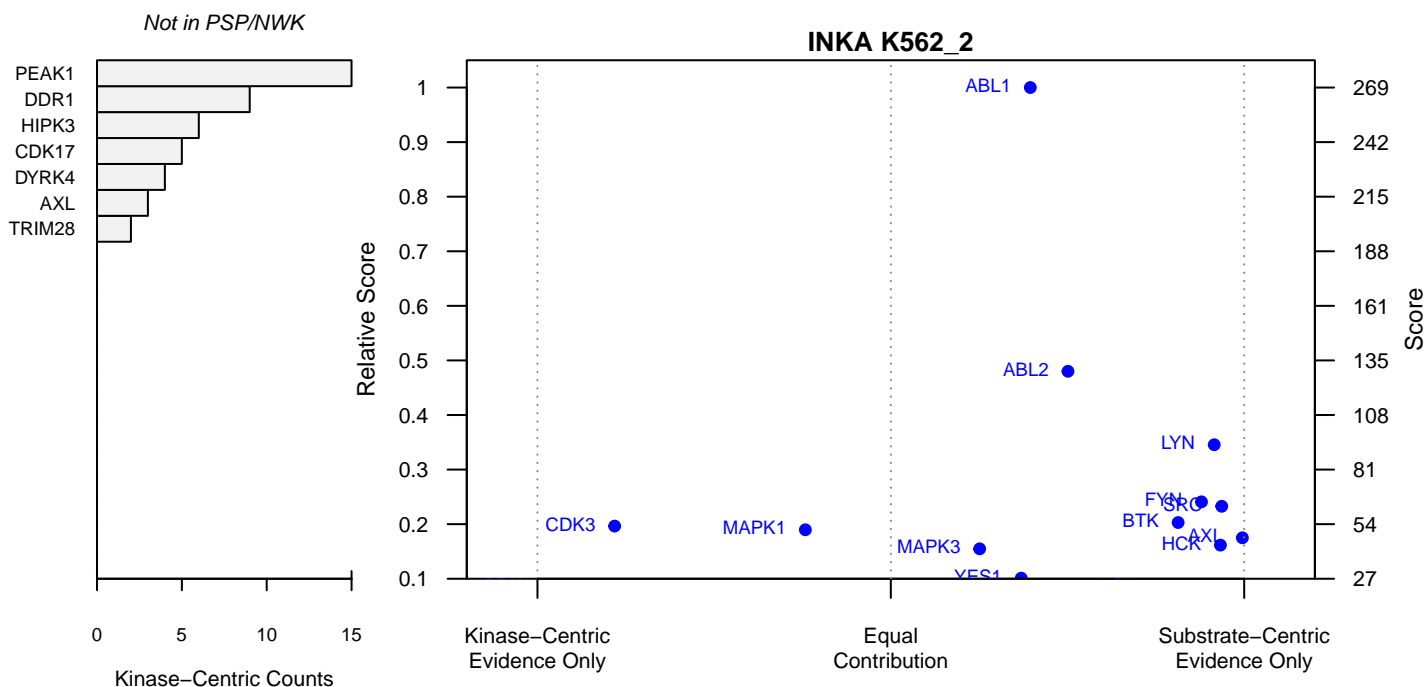
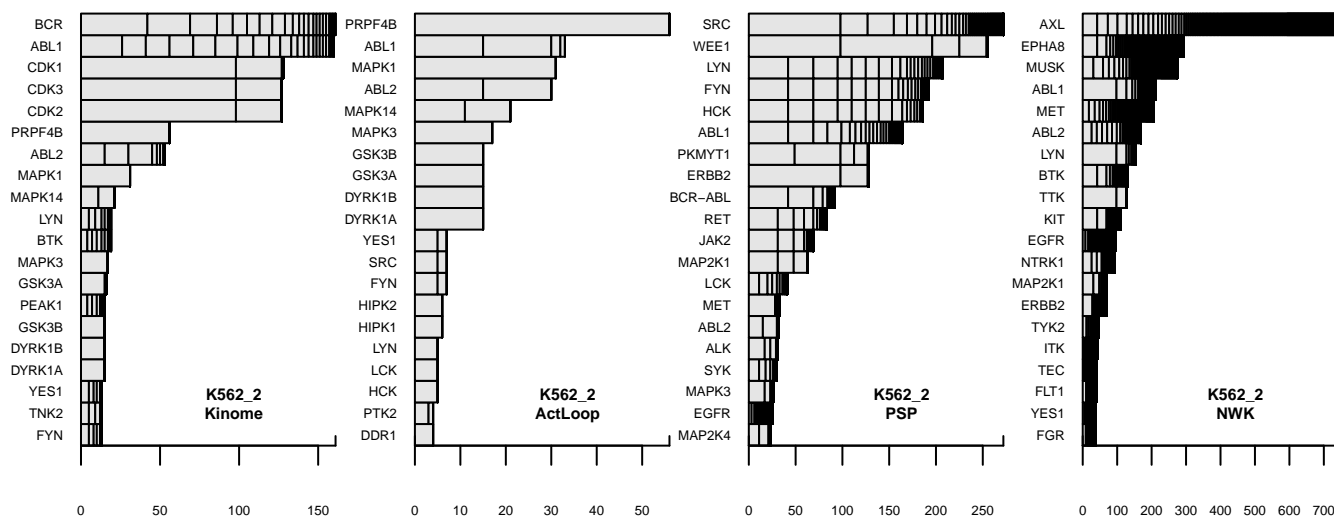
HCC827ER3_1



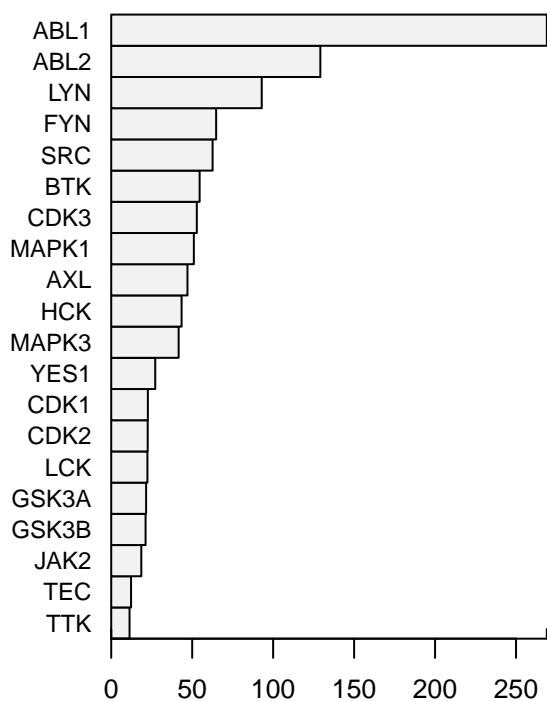


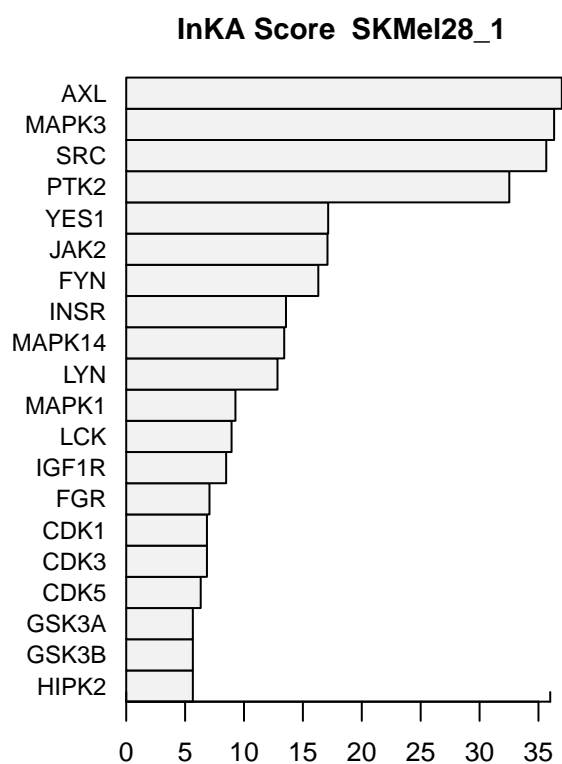
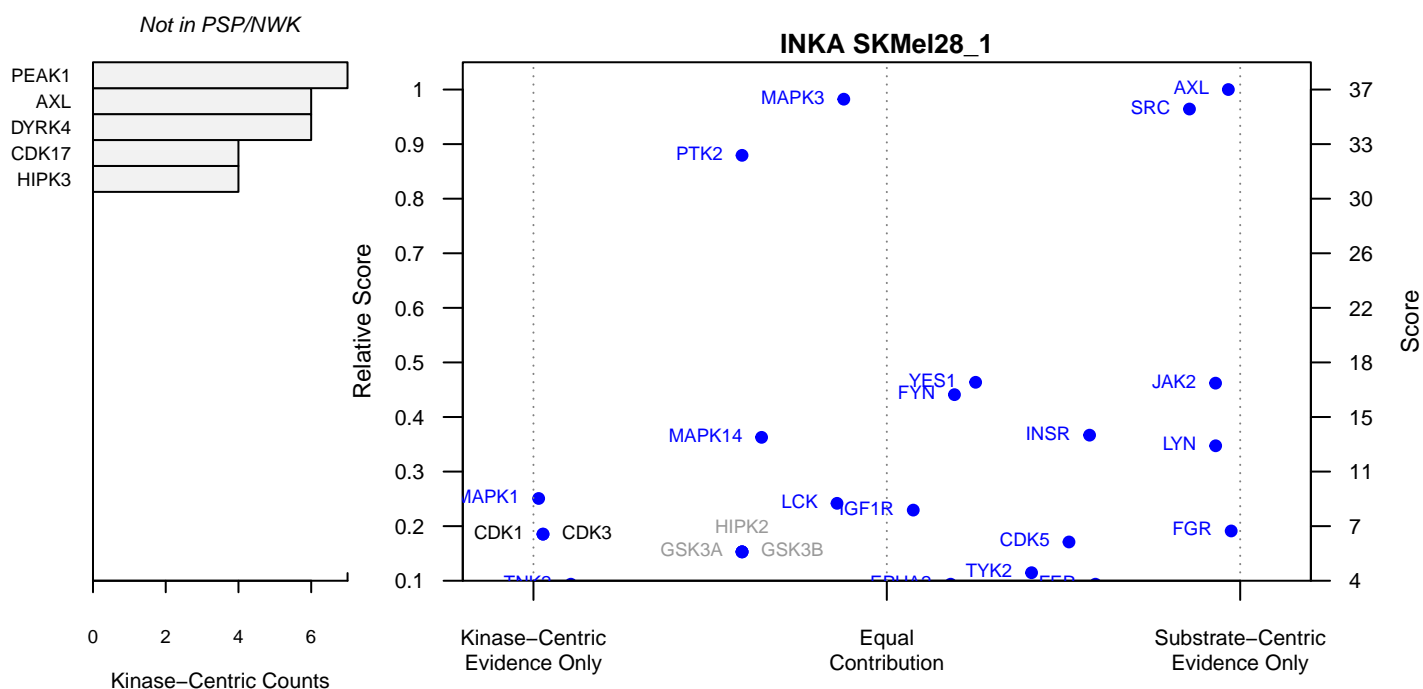
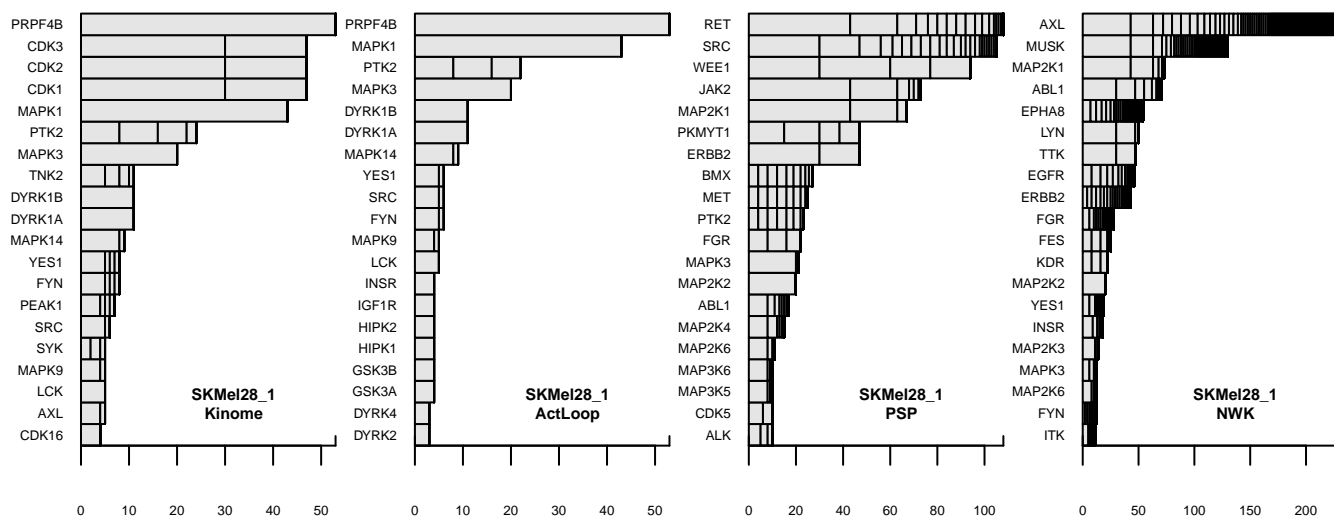


The diagram illustrates a complex network of protein-protein interactions. The nodes are represented by various shapes (circles, hexagons, diamonds) and colors (grey, red, black). The edges are colored lines (blue, orange, green) representing different types of interactions. The network is highly interconnected, with many nodes having multiple connections. Key nodes include SRC, ABL2, ABL1, BCR, YES1, and AXL, which are central to the network.

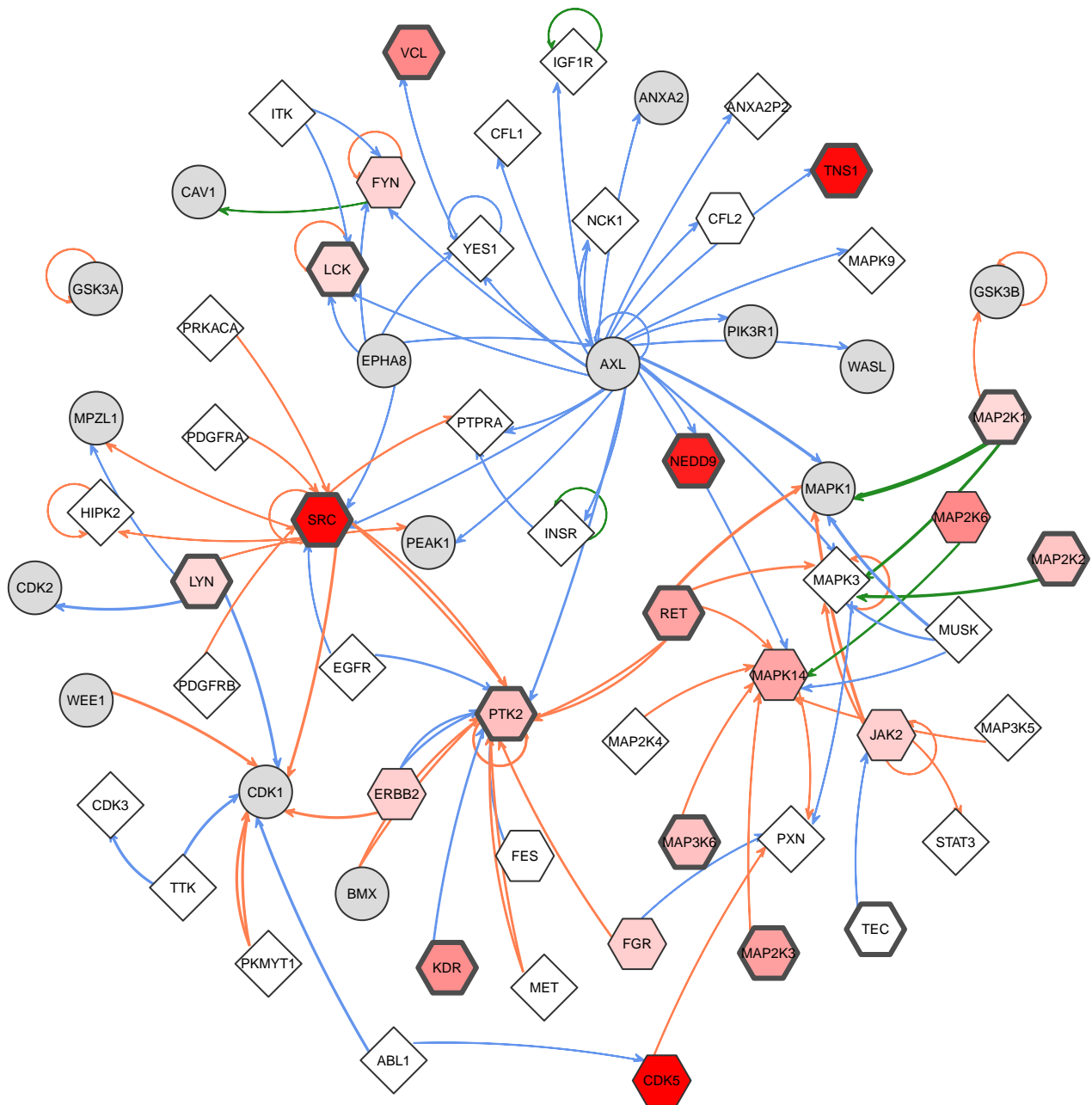


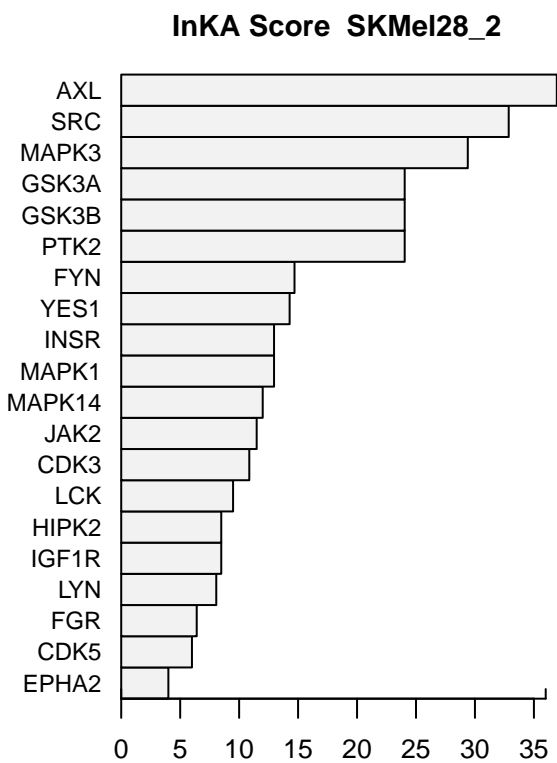
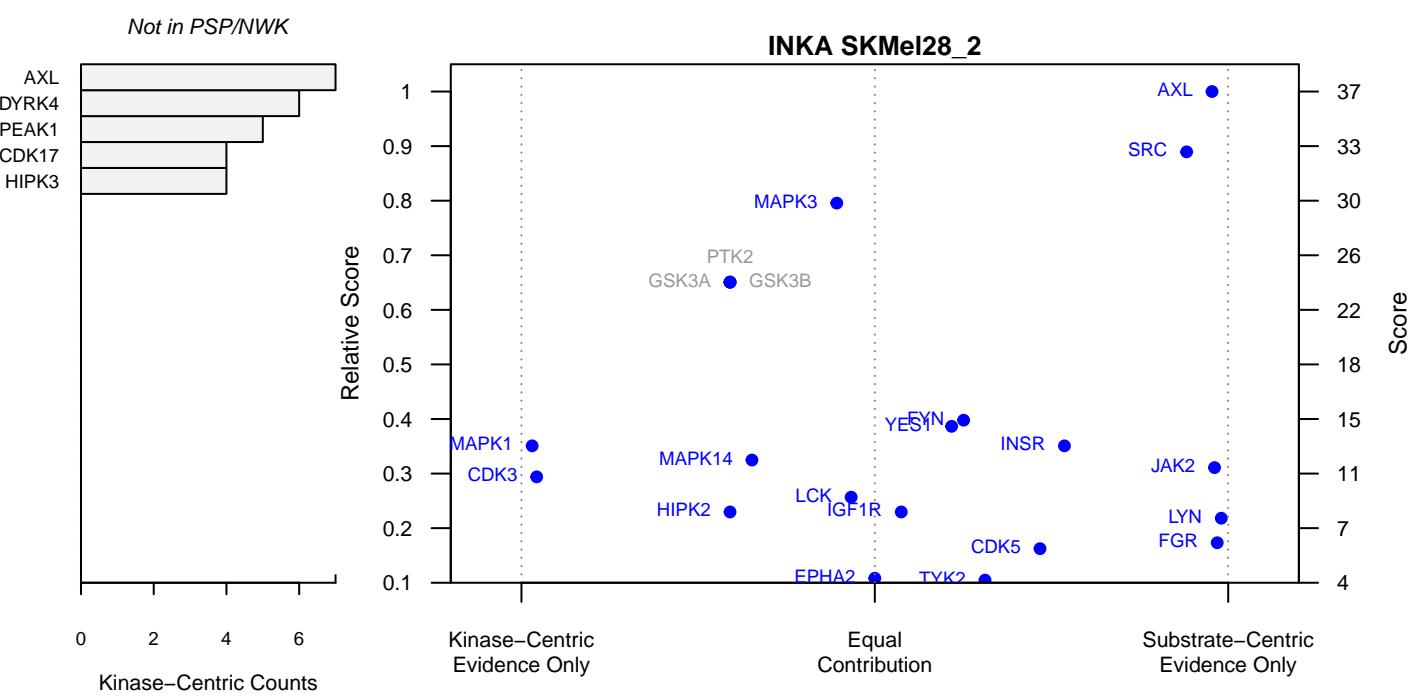
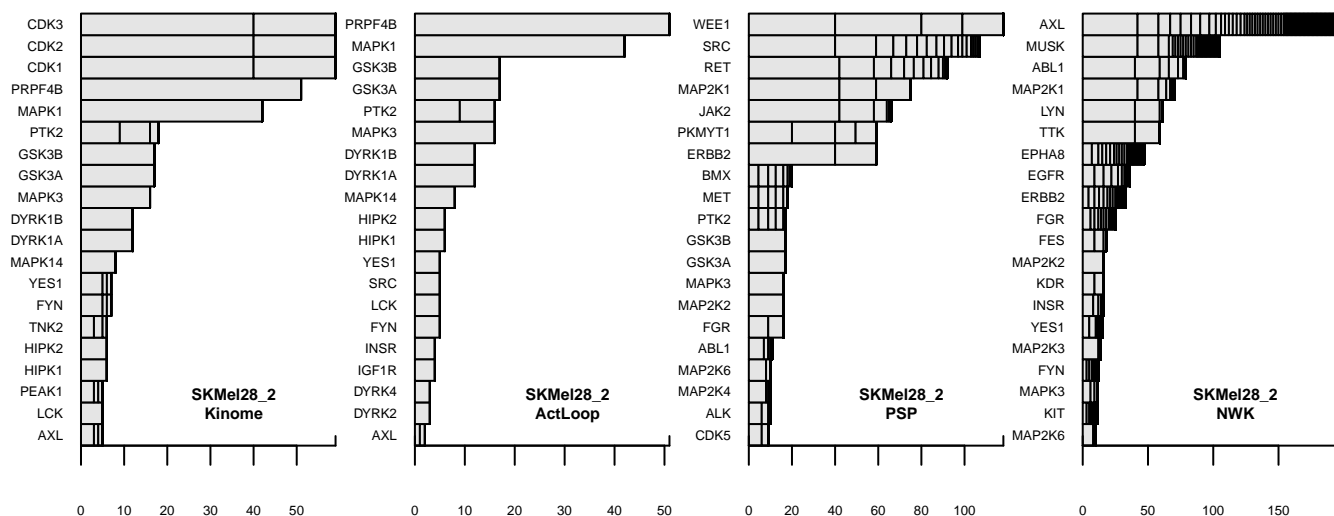
InKA Score K562_2





SKMel28_1





SKMel28_2

