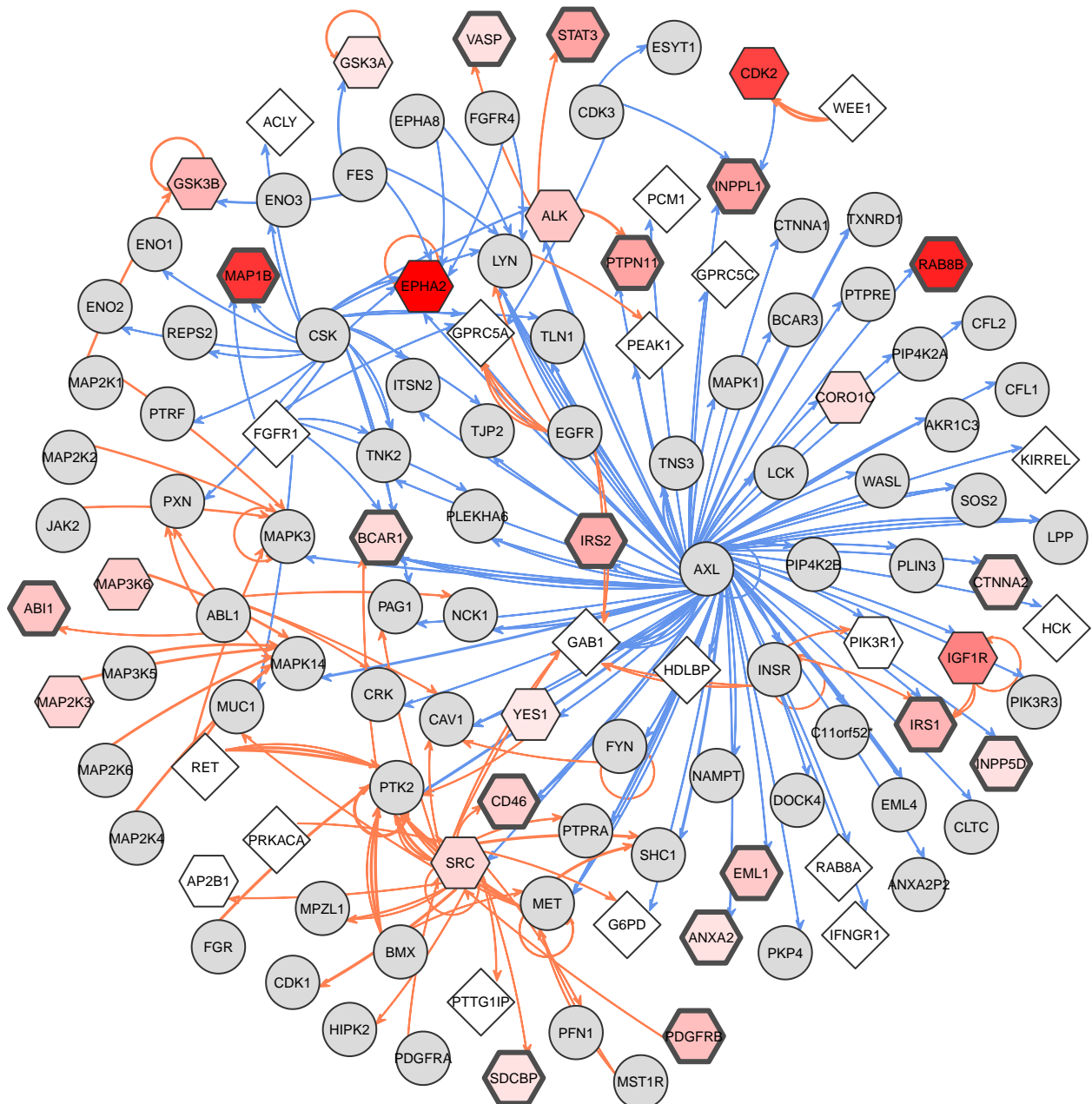
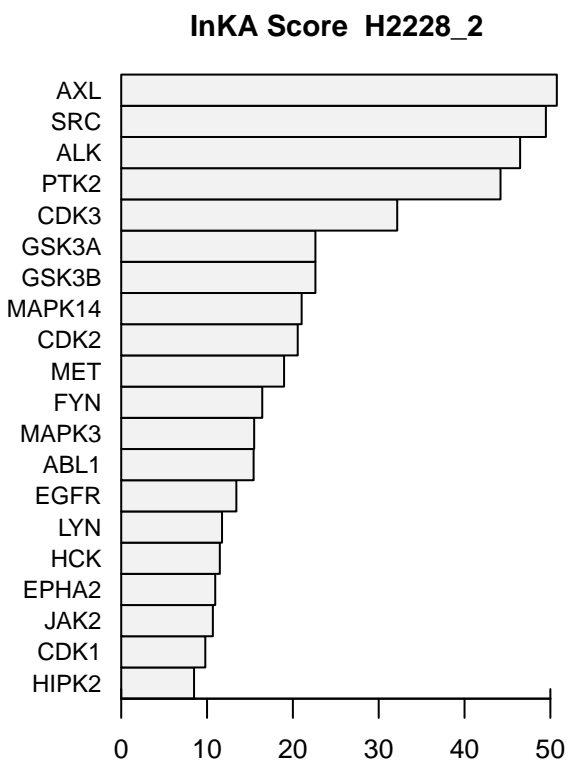
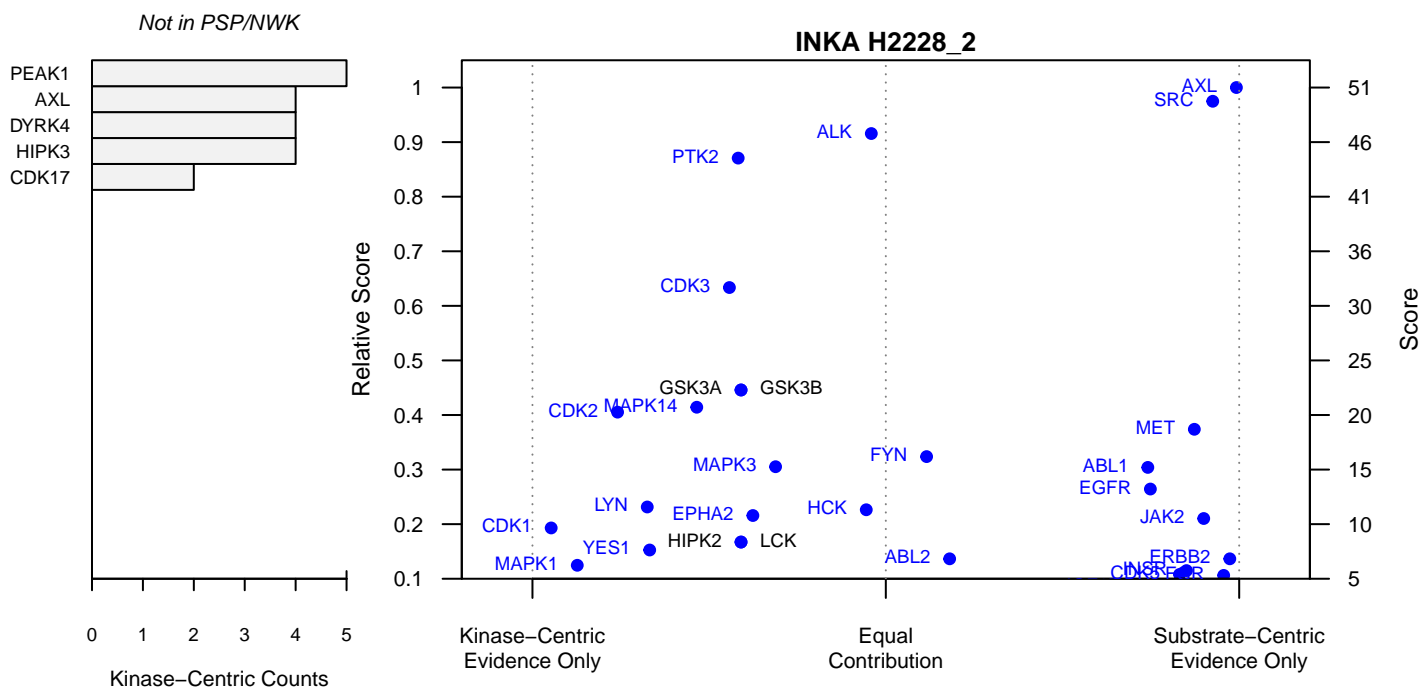
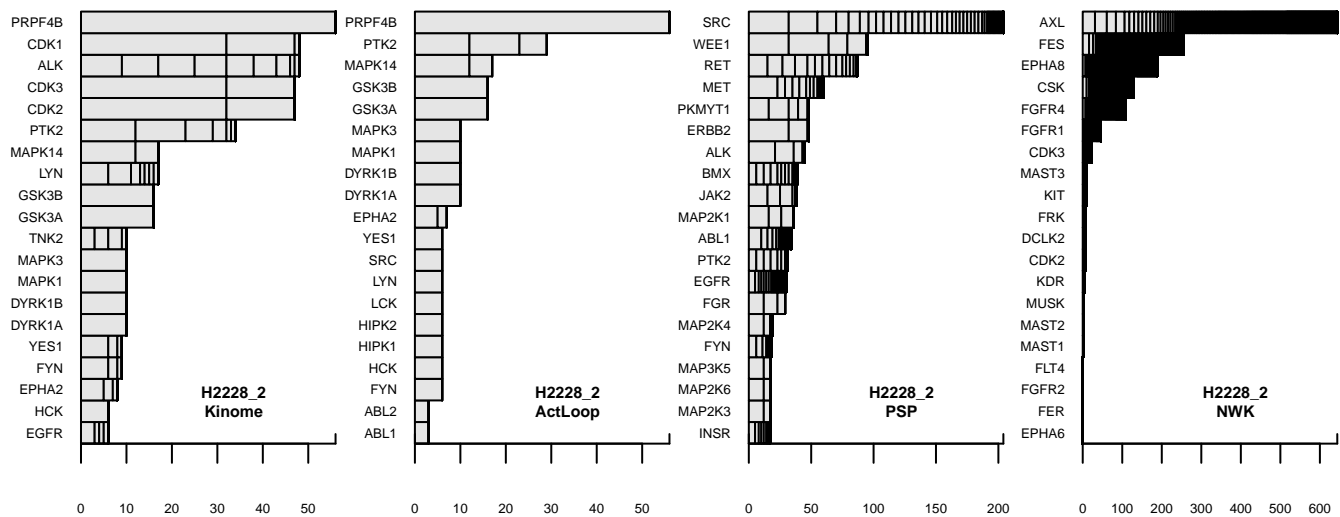
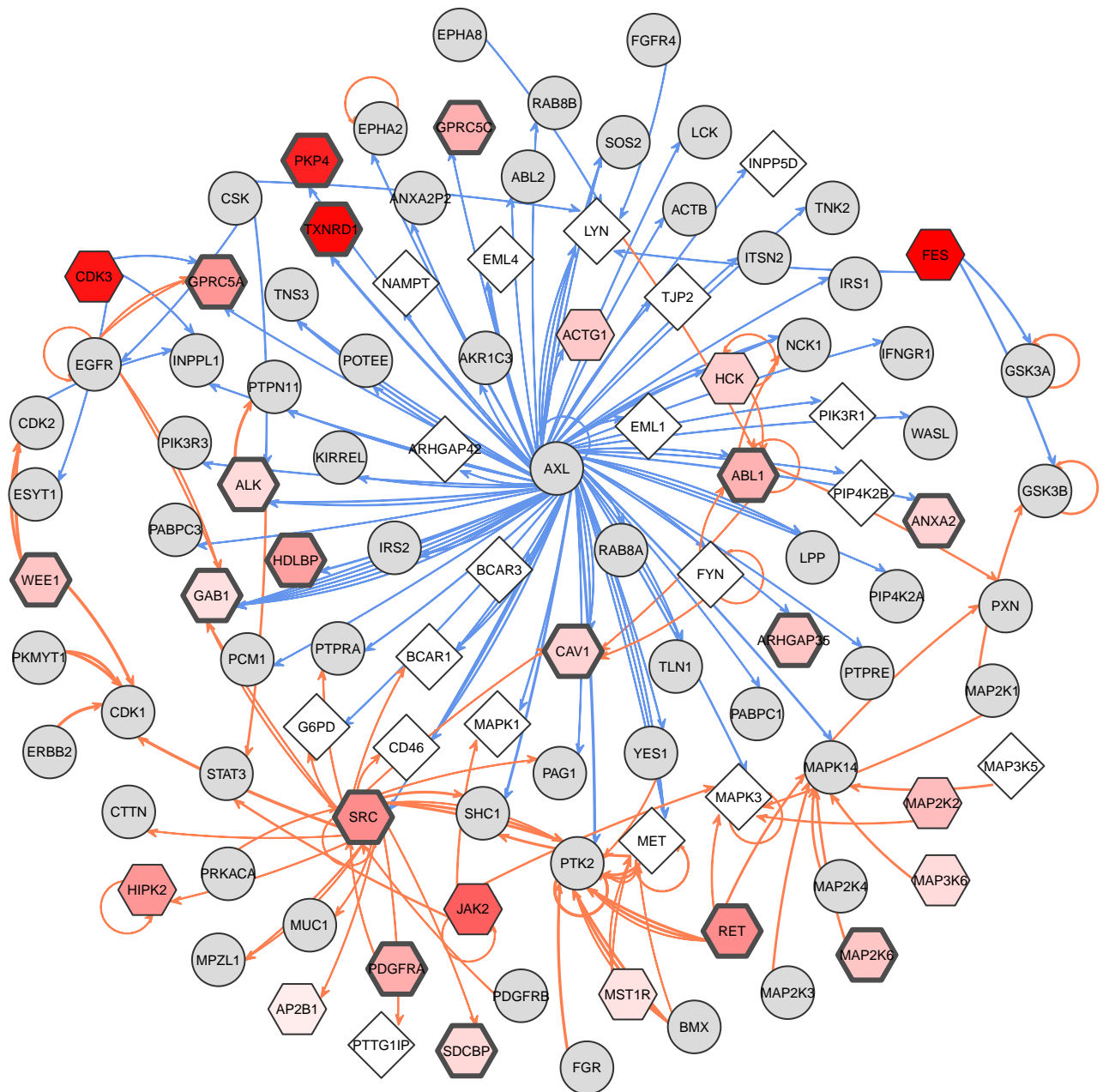


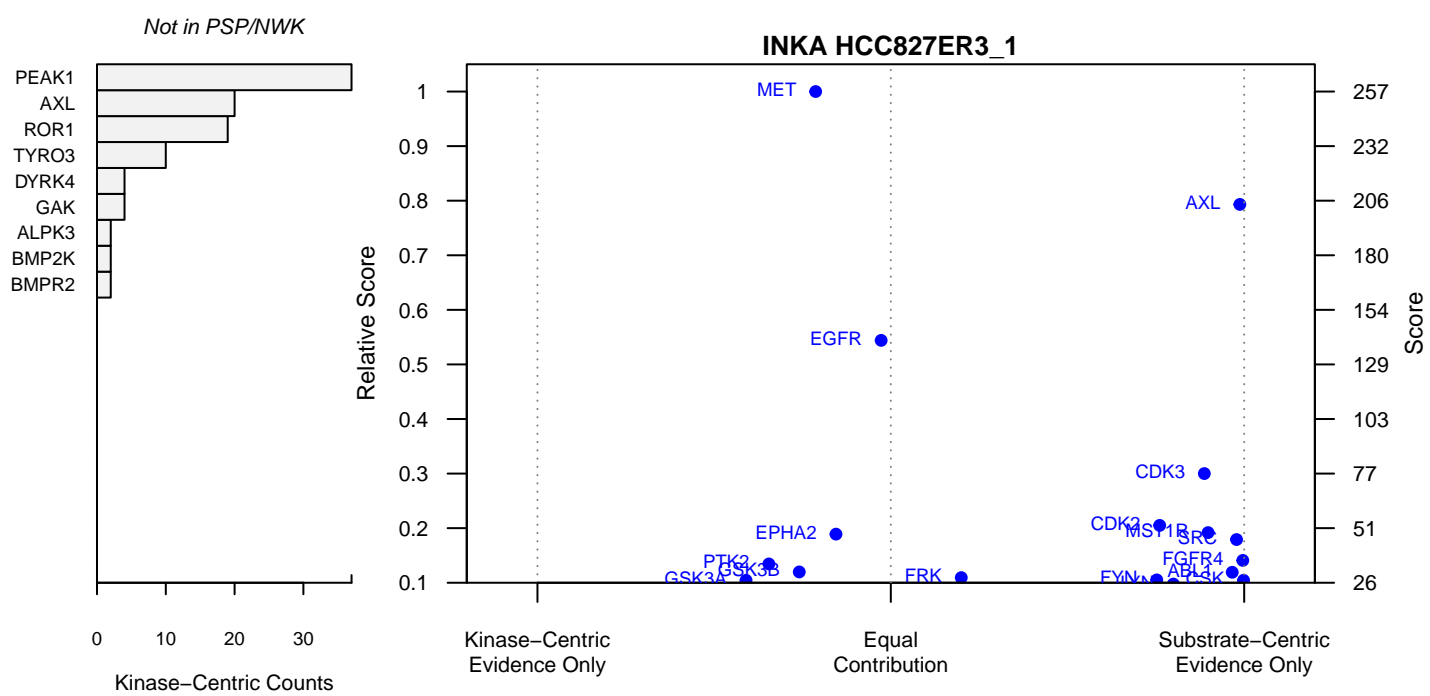
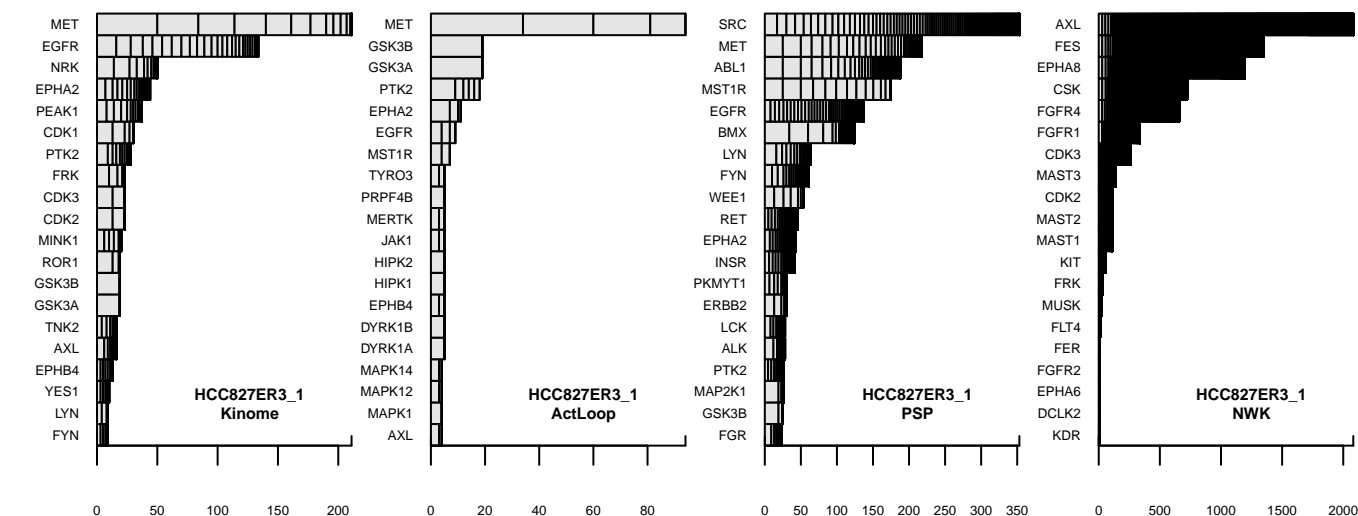
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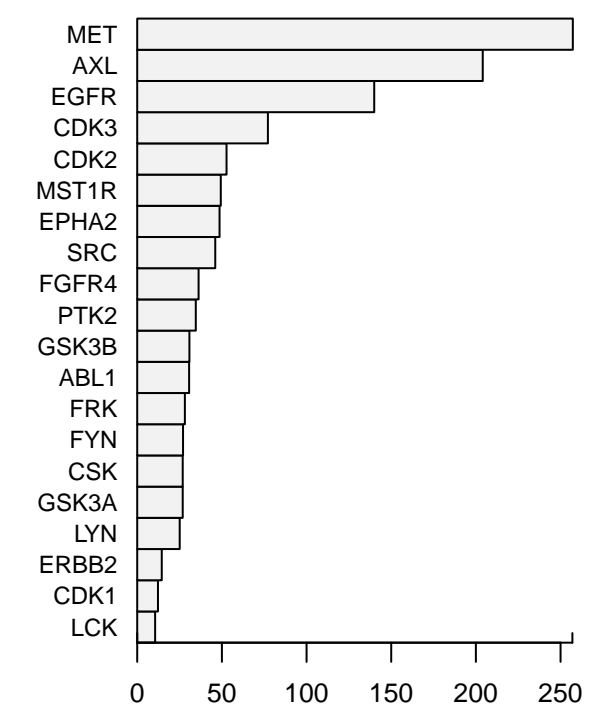


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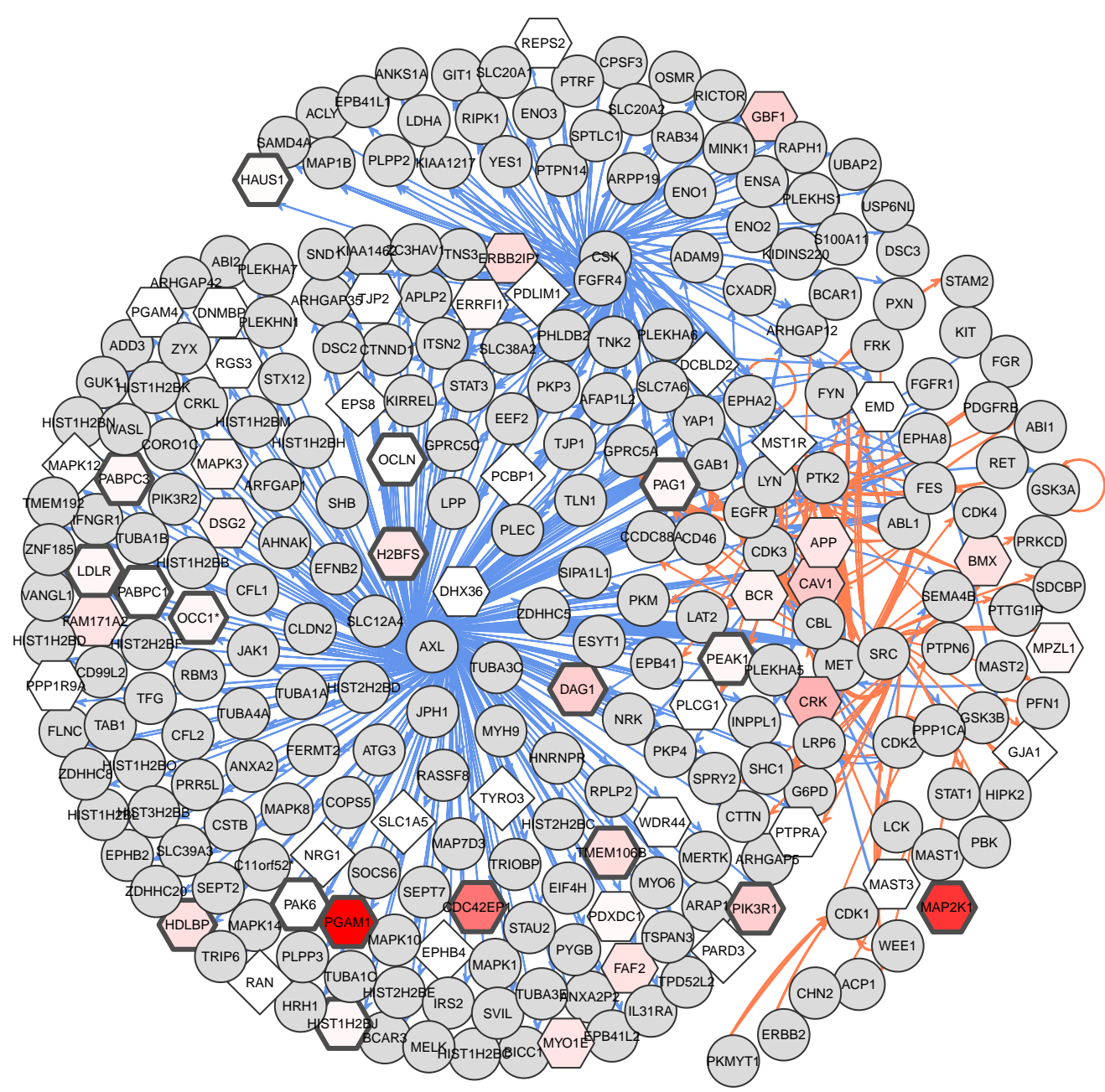


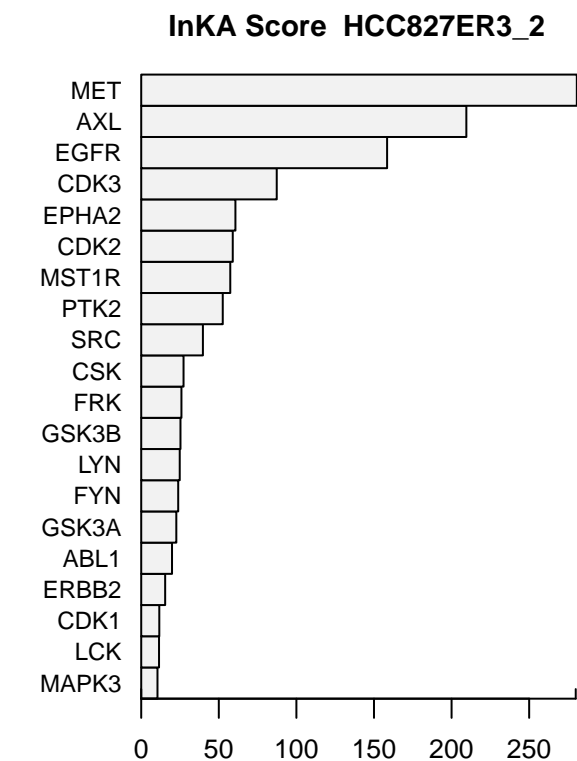
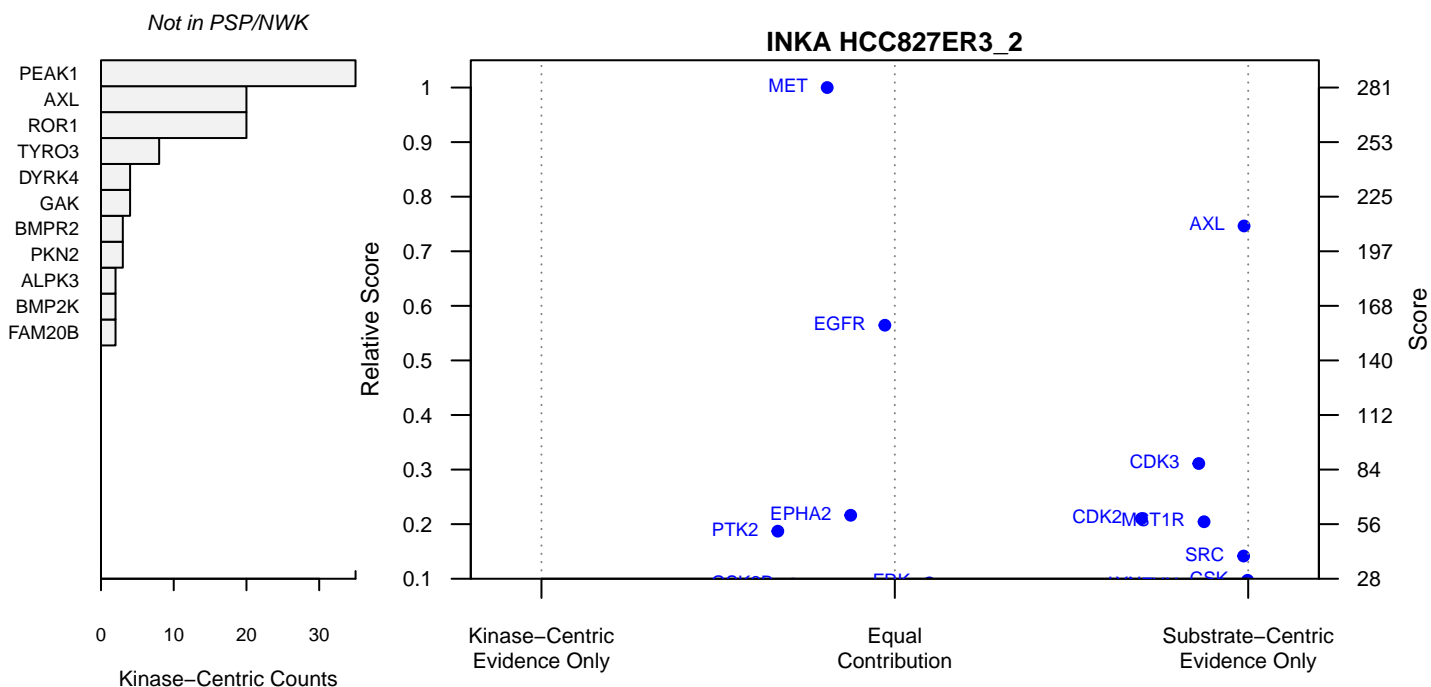
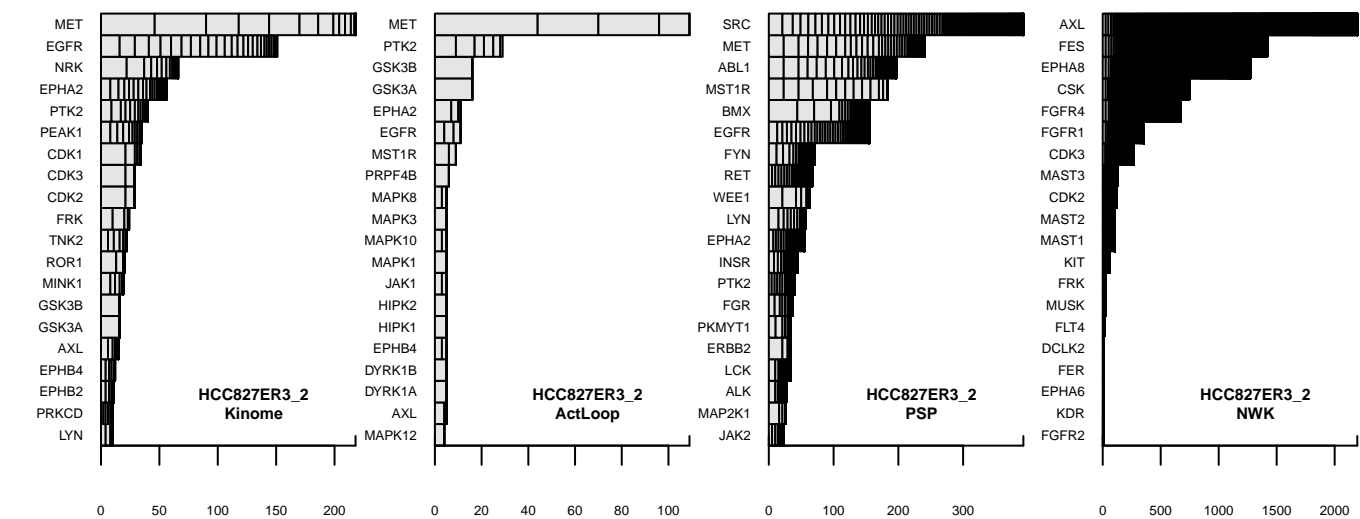


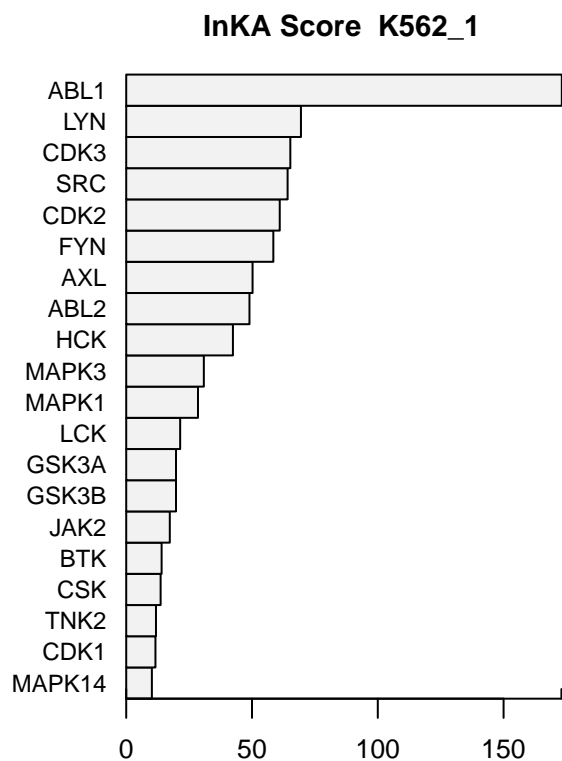
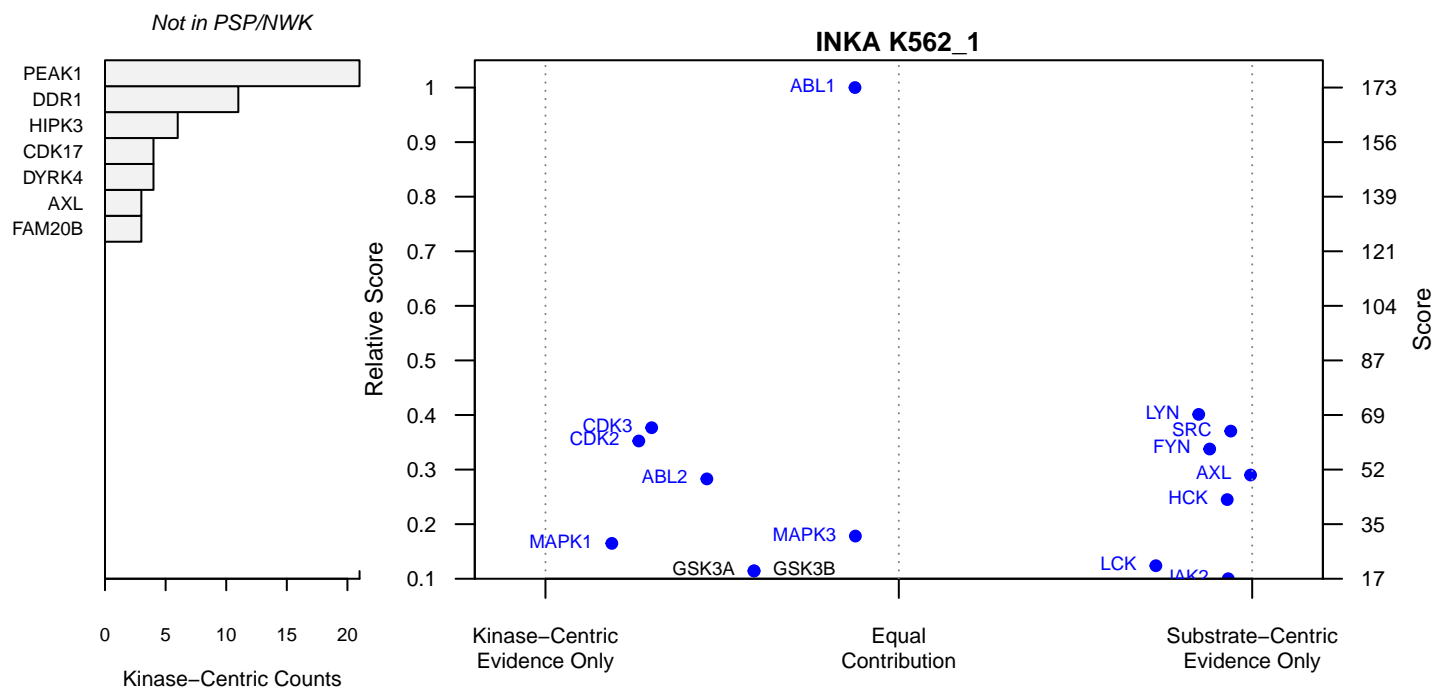
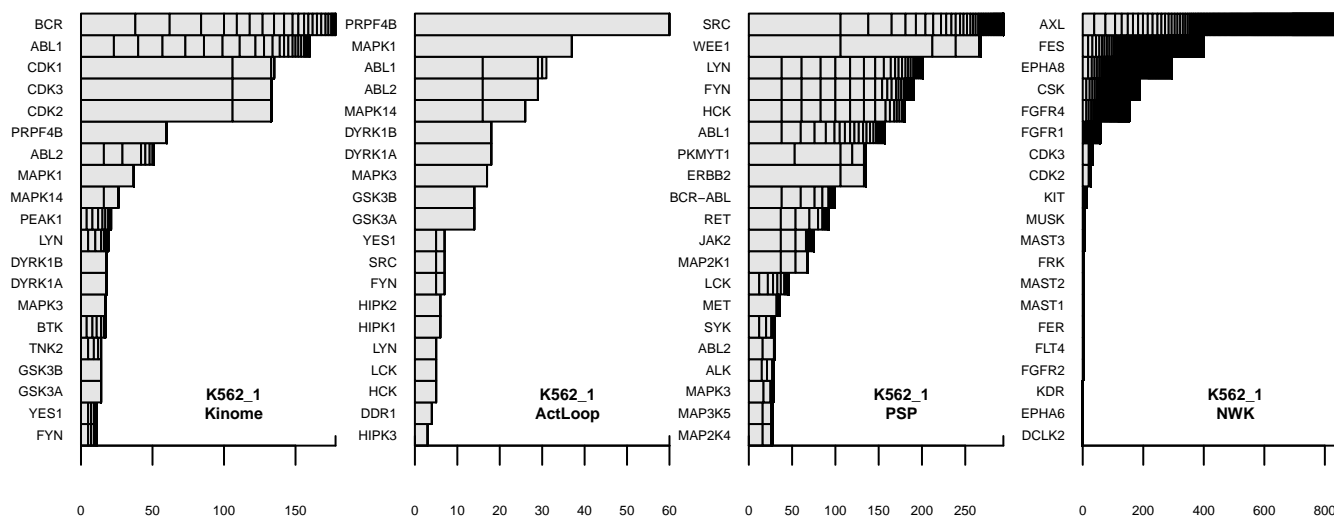
InKA Score HCC827ER3_1



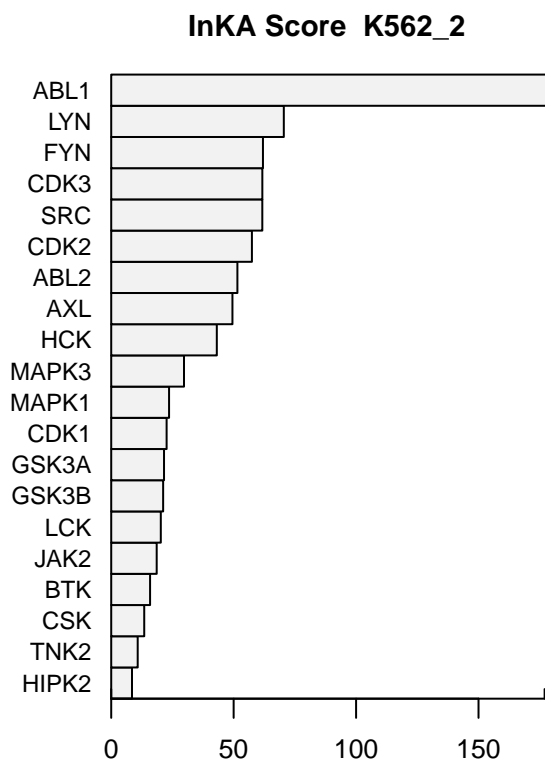
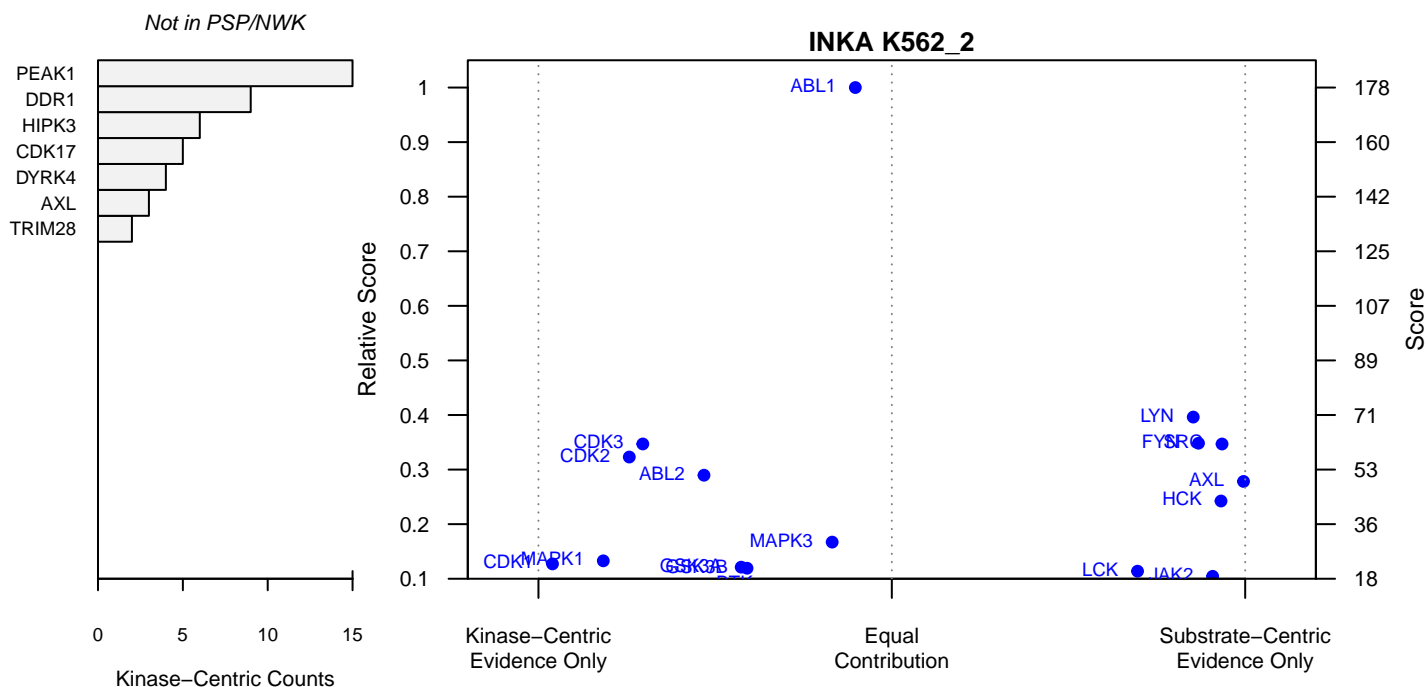
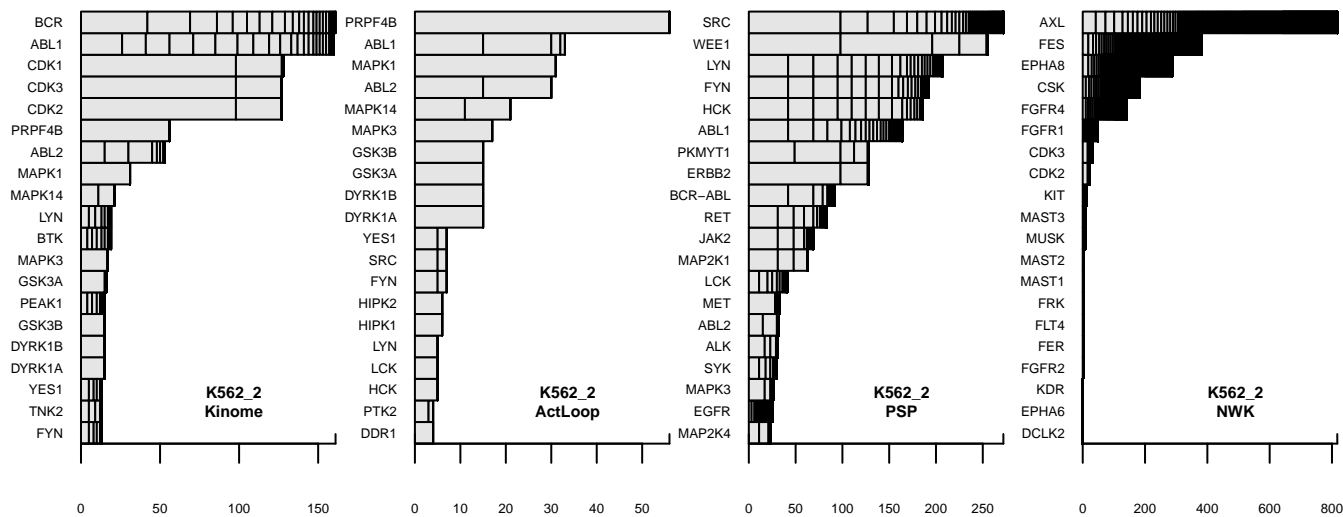
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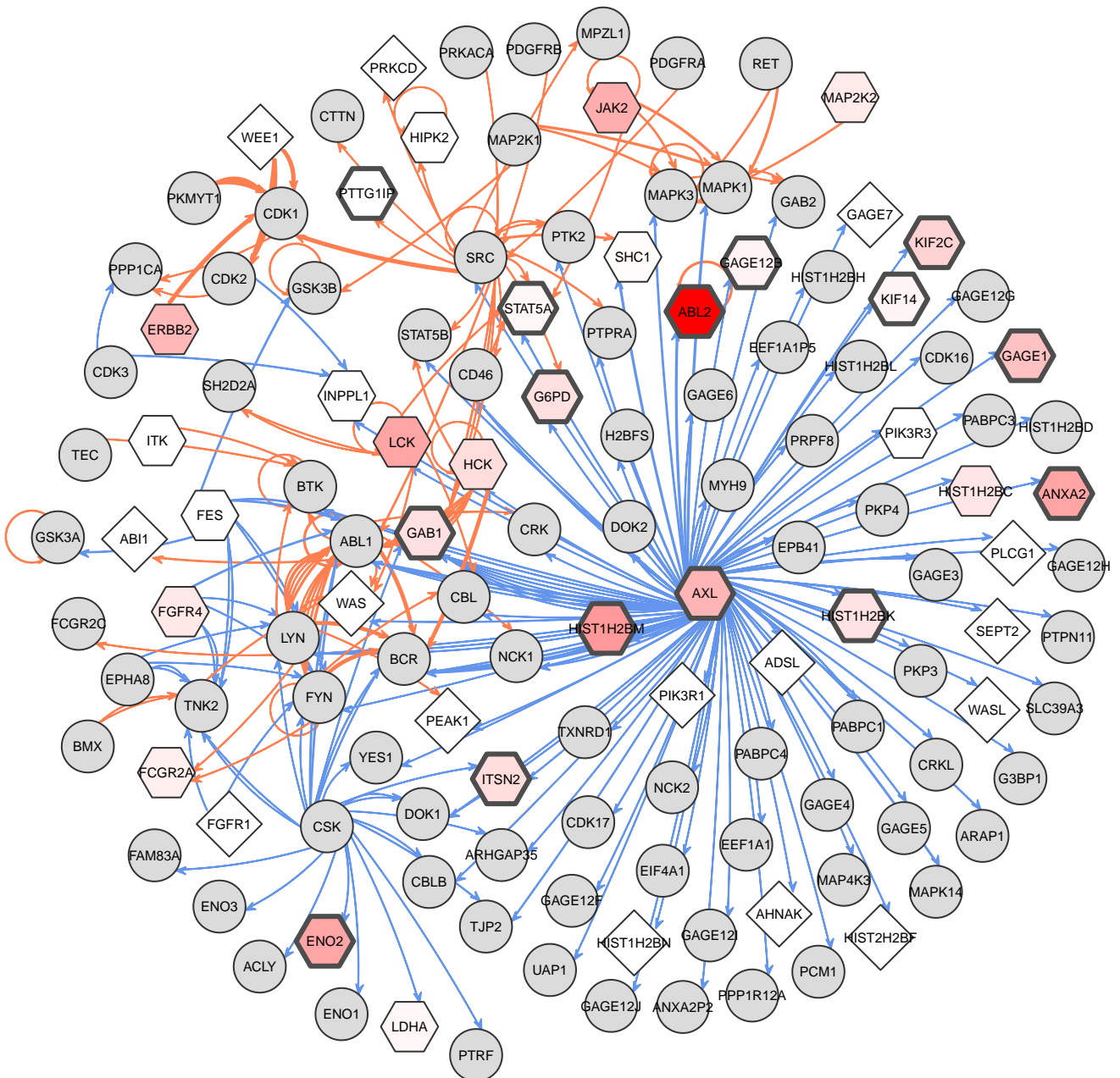


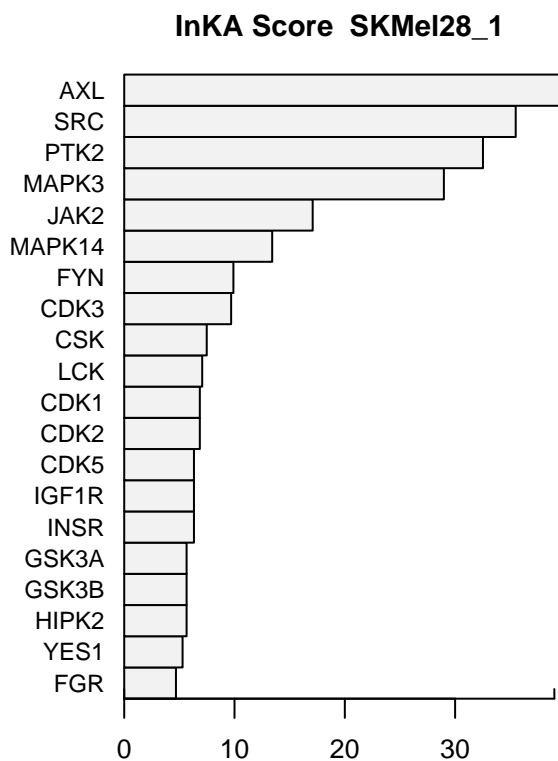
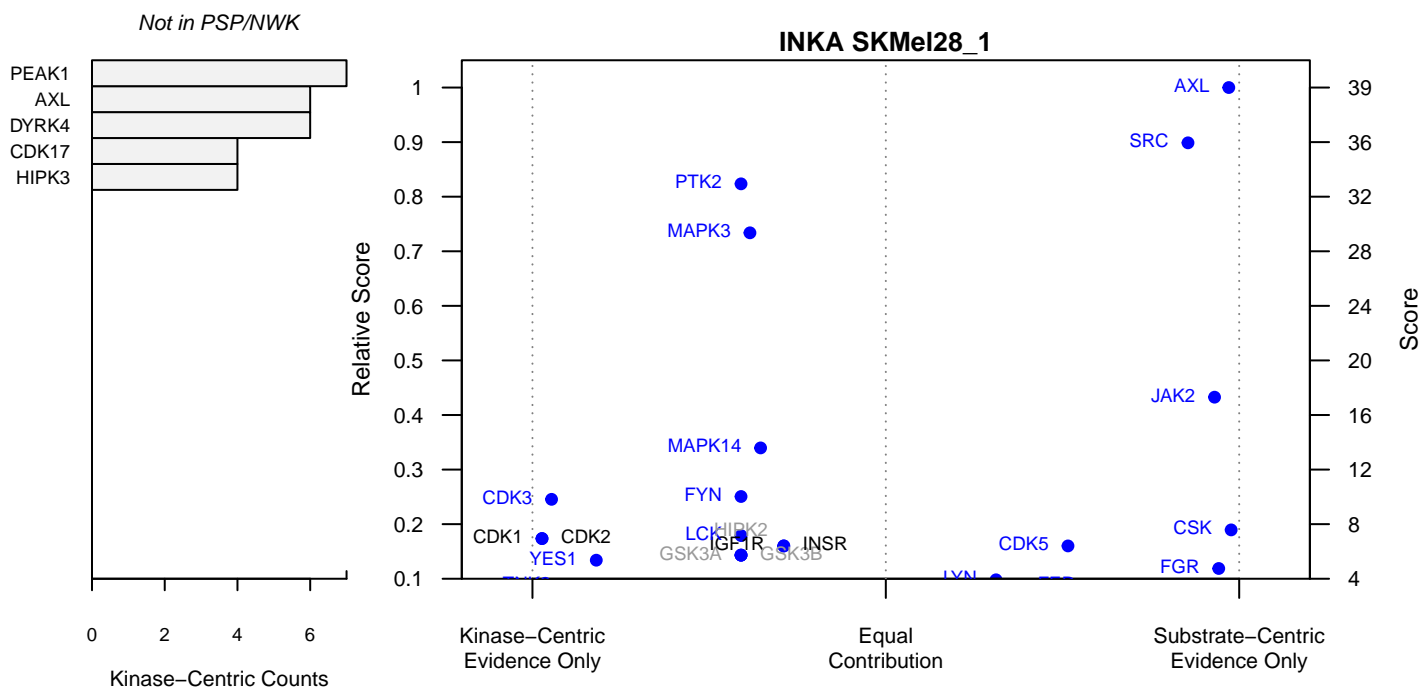
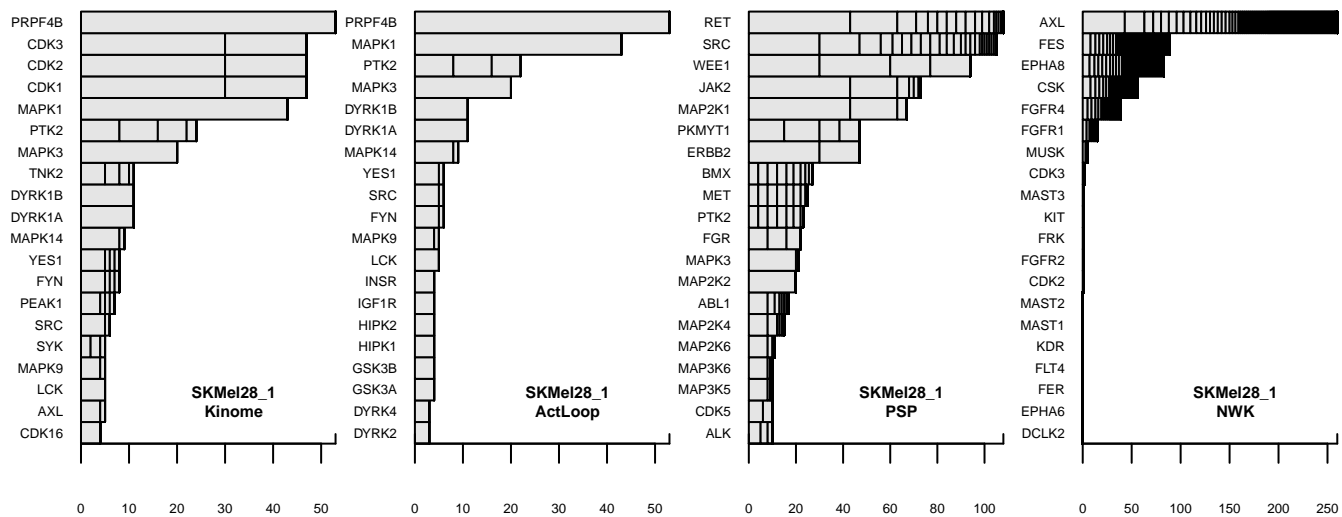


The diagram illustrates a complex network of protein-protein interactions. The nodes are represented by different shapes and colors, likely indicating different protein families or functional categories. The connections are represented by blue and orange lines, suggesting different types of interactions or signaling pathways. The network is highly interconnected, with many nodes having multiple incoming and outgoing connections. The central part of the network is particularly dense with many overlapping connections.

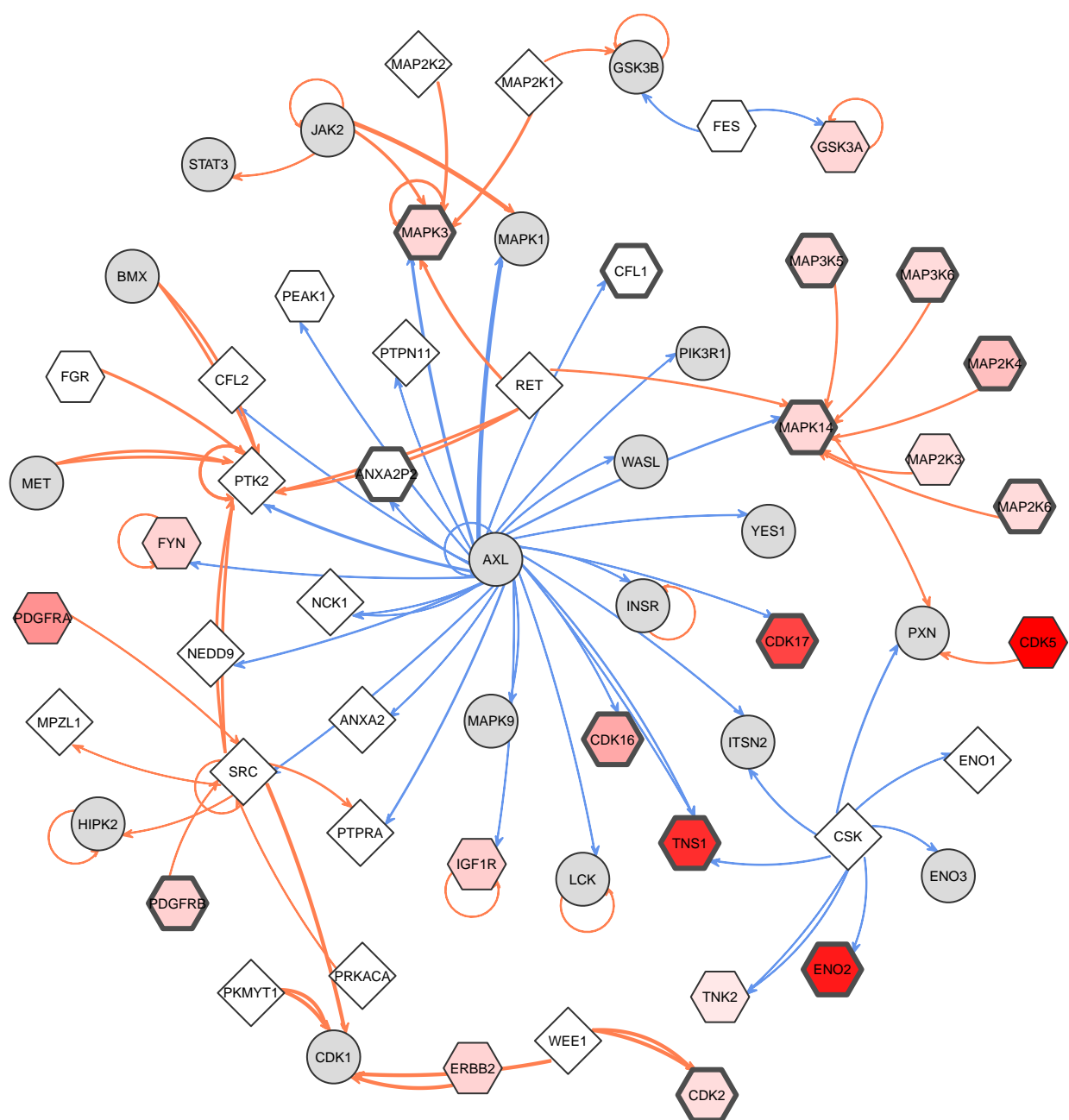


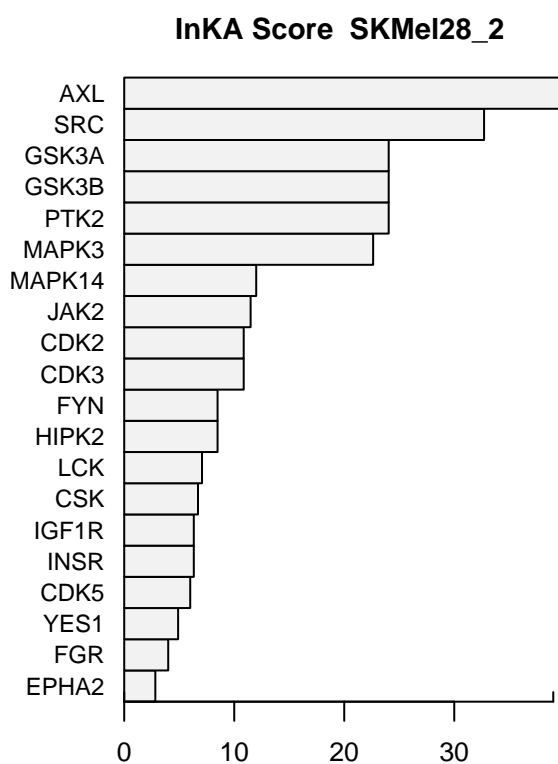
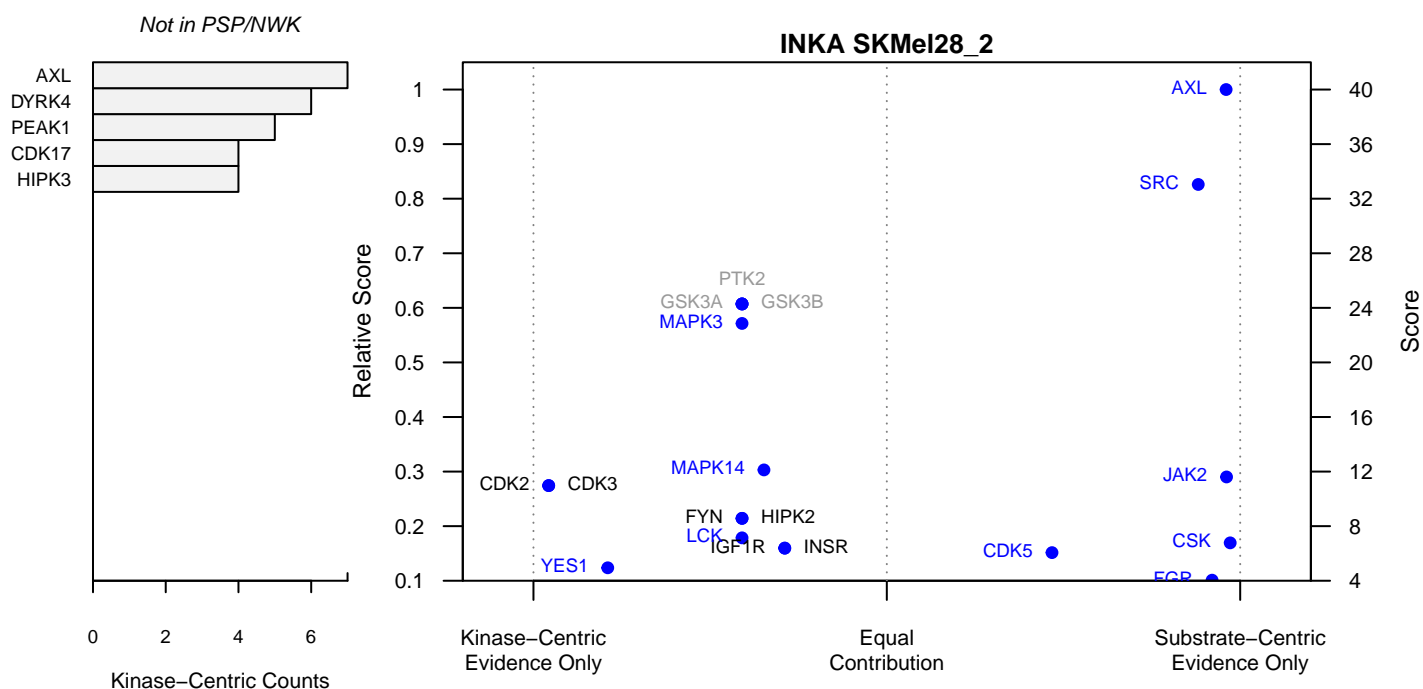
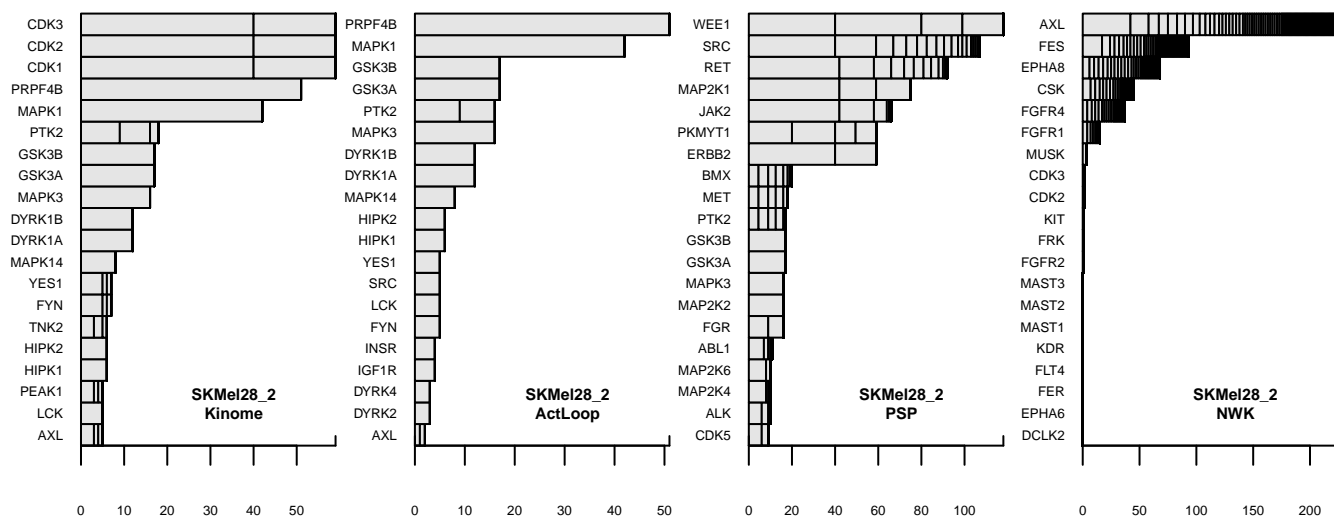
K562_2





SKMel28_1





SKMel28_2

