

The figure displays a large, complex network graph representing protein-protein interactions. The nodes are colored based on their functional clusters, including shades of blue, green, orange, purple, pink, and red. Each node is labeled with a protein name and its corresponding UniProt accession number (e.g., P12345, Q6789). The graph is highly interconnected, with a large central cluster and several smaller peripheral clusters. The edges represent interactions between the proteins, forming a dense web of connections.