

The diagram illustrates a complex network of gene-gene interactions, represented by a large, irregular shape composed of several overlapping colored polygons. The genes are categorized by their markers: red circles, blue squares, green triangles, and purple crosses. The network is divided into several distinct clusters or modules, each highlighted by a different color. The green cluster on the left includes genes like CTDSP2, ESYT2, CXCL12, and INTS5. The red cluster at the top includes SZRD1, PCMTD1, SLC35A4, and COQ2. The blue cluster in the center includes KLF10, SLC29A1, and PNC1. The purple cluster on the right includes SLC38A4, ADH1B, and ECHDC3. The orange cluster at the bottom includes C1RL, CEBPB, and FPG5. The network also shows interactions between these clusters, with genes like MYC, PELO, and GNG5 acting as hubs. The background is a light gray grid.

