

Figure S2

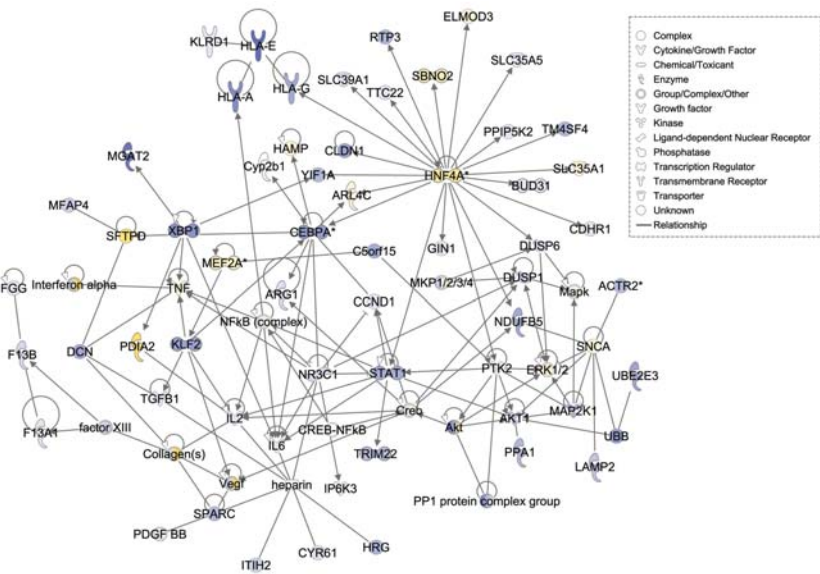


Figure S2. Magnification of Figure 4C. The cellular network differentially expressed between joint G2345 and joint G345 responsible for the G2345 to G345 transition (orange edge), merged with the G345e time-specific analysis gene sets, and built based on direct interactions between molecules. Data overlaid is the averaged G345e/G2 log₂ ratio expression data. Upregulated genes are shaded in yellow, and downregulated genes are shaded in blue.