The image contains a network of gene expression levels, with various genes and their interactions represented. The graph shows a complex network of connections between different gene identifiers, such as H-TGC_M-CAG72, H-TGC_M-CAG211, E-ACT_M-CTA104, and H-TGC_M-CAG112. The network appears to be dense, with many lines connecting different nodes, indicating interactions or expression correlations. The network is color-coded, likely to differentiate between different types of interactions or conditions. The vertical axis seems to represent some form of expression level, possibly ranging from 0 to 300, as inferred from the scale markers at the sides of the graph.