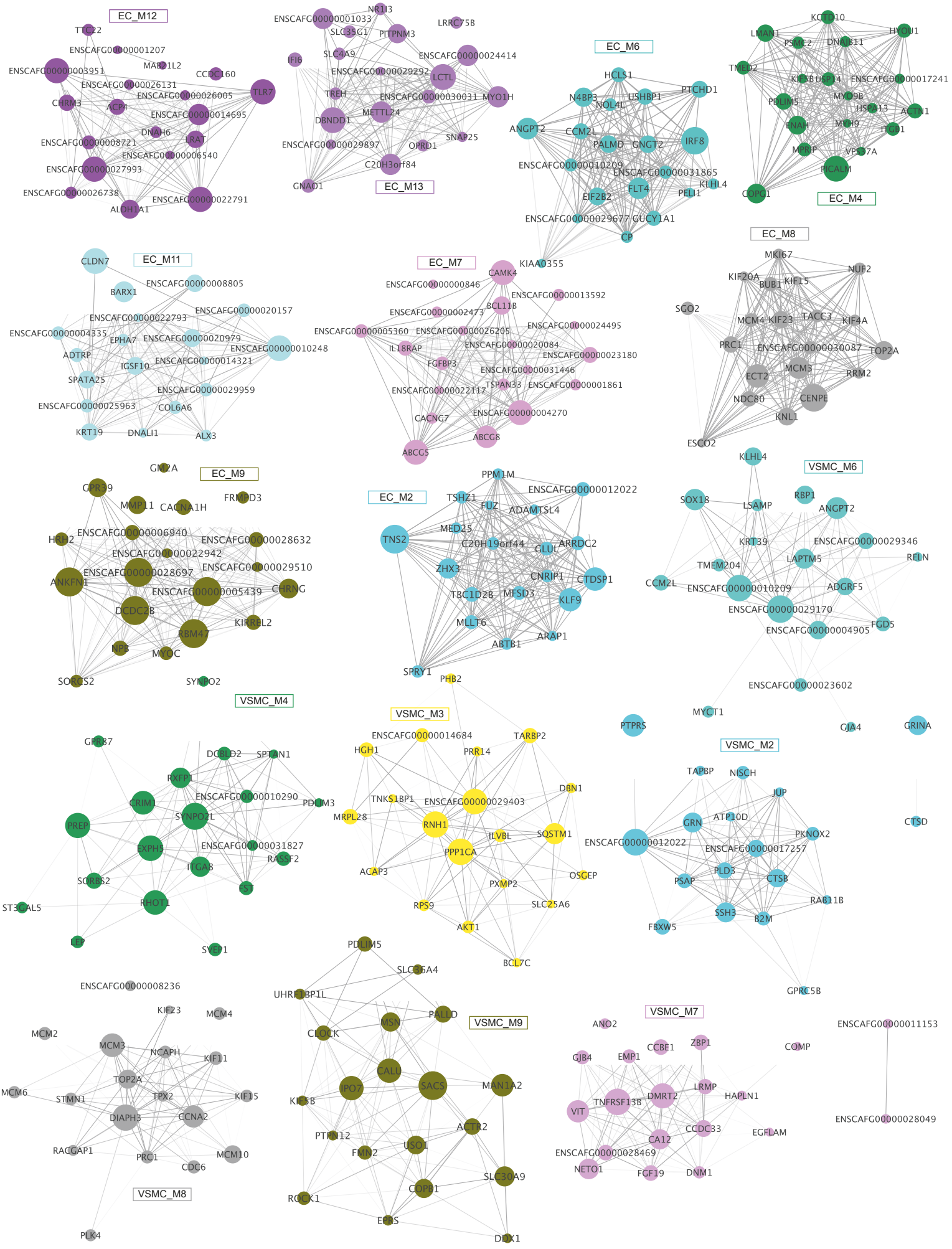

Supplemental Figure 5. Hub gene networks generated by Cytoscape.
Hub gene networks for a selection of modules significantly correlated to one or more vessel types in ECs or VSMCs. Each network consists of 20 genes with the highest intramodular connectivity in the module. Node size was scaled to intramodular connectivity and line transparency to edge weight.