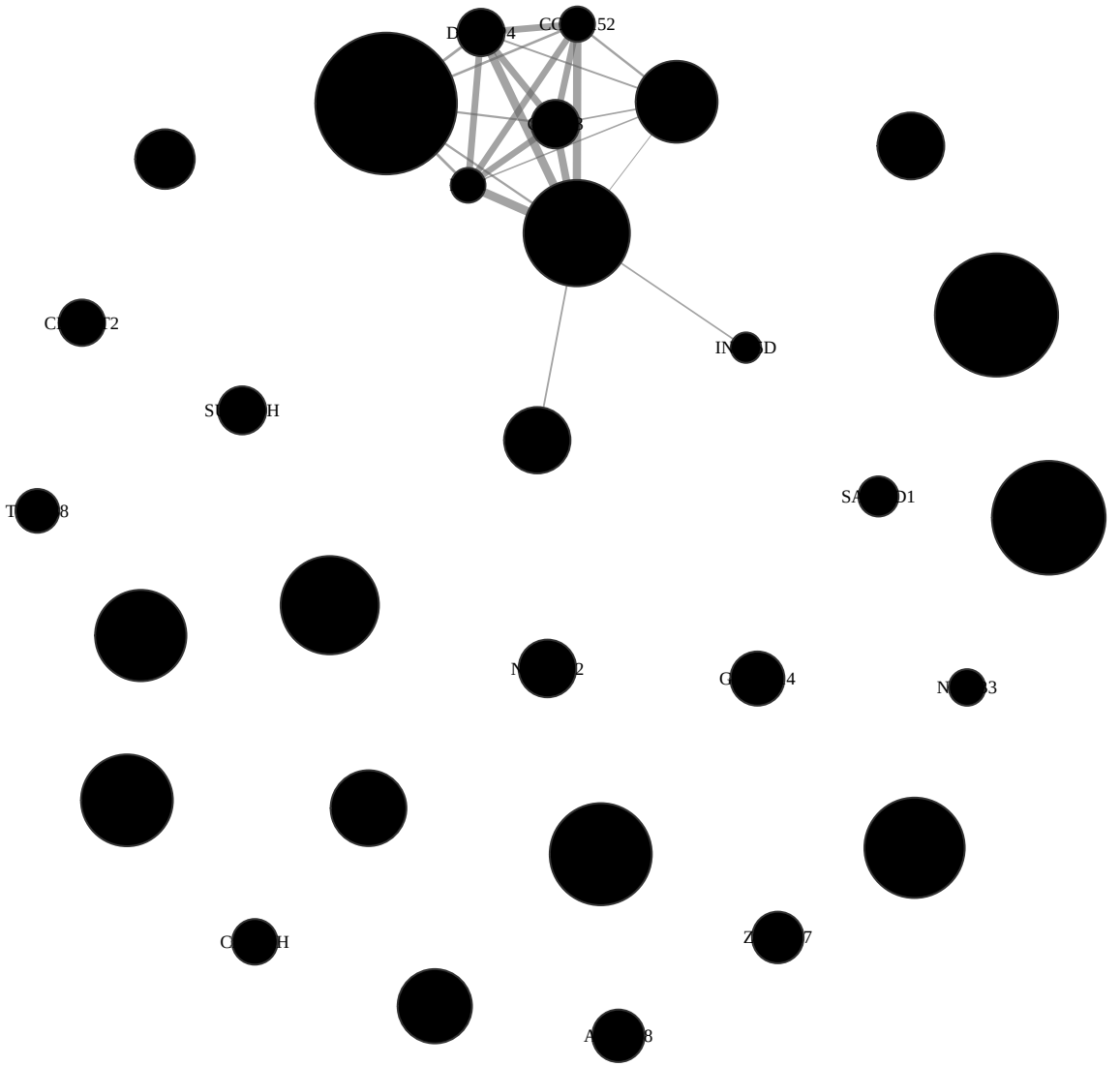
Module black hub network (n genes = 410, r = −0.32, p = 0.0425)



Top 30 genes by kME; edges: intramodular TOM >= 95th percentile (0.239)