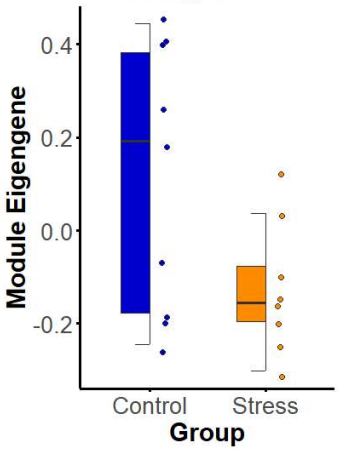
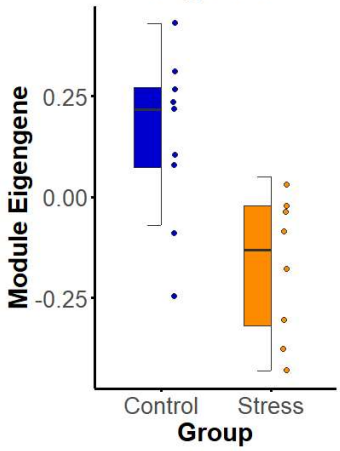
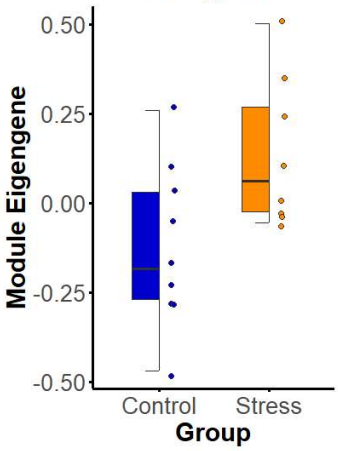
C. WGCNA: ARC

Lower in Stressed:		Higher in Stressed:
Yellow: 365 genes 	Purple: 66 genes 	Blue: 1528 genes 
Translation at synapse Mitochondrial gene expression Regulation of protein stability	Cardiac epithelial to mesenchymal transition Negative regulation of DNA binding Blood vessel endothelial cell differentiation	Chromatin remodeling mRNA processing Ribonucleoprotein complex biogenesis
Rpsa: 0.9375 *Uba52*: 0.9340 *Rpl34*: 0.9189 *Rps16*: 0.9126 *Rpl29*: 0.9074	*Trip6*: 0.8998 *Phgdh*: 0.8841 *Mrpl27*: 0.8768 *Selenom*: 0.8762 *Timm13*: 0.8407	*Eif3a*: 0.9613 *Atrx*: 0.9574 *Eea1*: 0.9514 *Gcc2*: 0.9512 *Suz12*: 0.9503