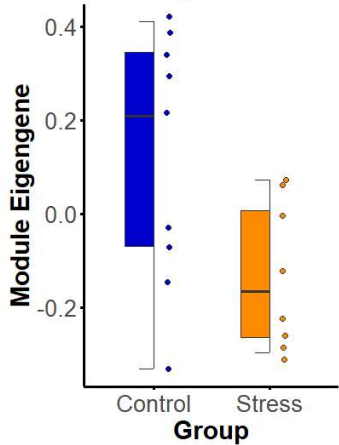
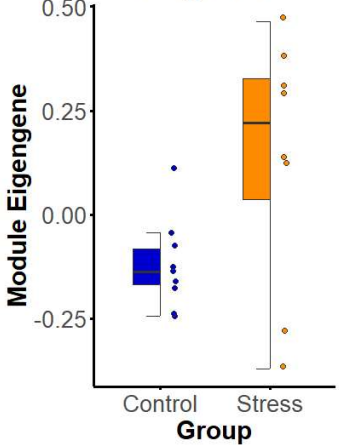
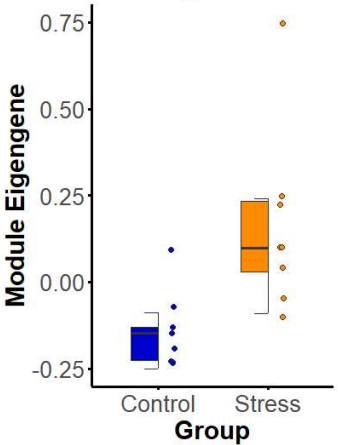


B. WGCNA: DMH

Lower in Stressed:	Higher in Stressed:	
<p>Darkred: 29 genes</p>  <p>Module Eigengene</p> <p>Control Stress</p> <p>Group</p>	<p>Skyblue: 21 genes</p>  <p>Module Eigengene</p> <p>Control Stress</p> <p>Group</p>	<p>Darkgrey: 26 genes</p>  <p>Module Eigengene</p> <p>Control Stress</p> <p>Group</p>
<p>Alcohol biosynthetic process</p> <p>L-glutamate transmembrane transport</p> <p>Positive regulation of cellular respiration</p>	<p>Forebrain cell migration</p> <p>Cellular response to interleukin-4</p> <p>Developmental maturation</p>	<p>Import across plasma membrane</p> <p>Positive regulation of mast cells</p> <p>Xenobiotic transport</p>
<p><i>Arhgdig</i>: 0.9355</p> <p><i>Rpl36</i>: 0.8992</p> <p><i>Ssu72</i>: 0.8873</p> <p><i>Pcbd1</i>: 0.8788</p> <p><i>Psm6</i>: 0.8243</p>	<p><i>Cntnap1</i>: 0.9521</p> <p><i>Fgfr1</i>: 0.9261</p> <p><i>Ogdh</i>: 0.8810</p> <p><i>Reln</i>: 0.8808</p> <p><i>Bap1</i>: 0.8695</p>	<p><i>Fcer1g</i>: 0.8937</p> <p><i>C1qc</i>: 0.8817</p> <p><i>Ak3</i>: 0.8746</p> <p><i>Ctsz</i>: 0.8528</p> <p><i>Slco2b1</i>: 0.8420</p>