



Heatmap illustrating the expression of a representative gene (eigengene) for the 38 co-expression modules defined by WGCNA\_global network. Modules are represented as columns, with the dendrogram illustrating eigengene relatedness. Each row represents one sample; coloured bars to the left indicate the high-level tissue, high-level age and high-level stress/disease of the sample. DESeq2 normalised expression levels are shown.