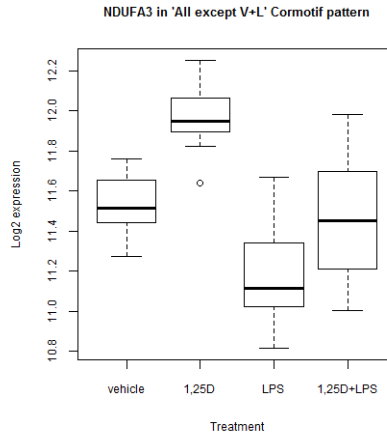
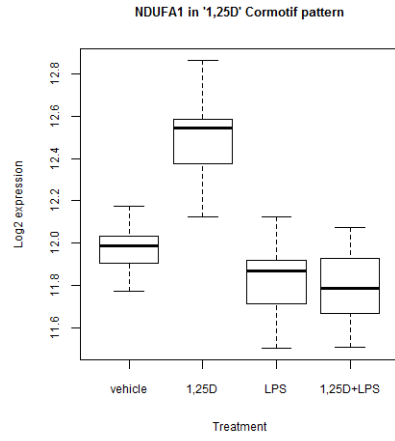
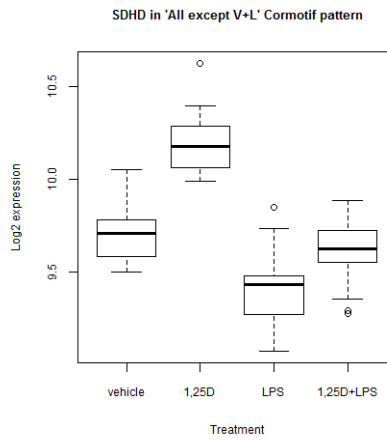
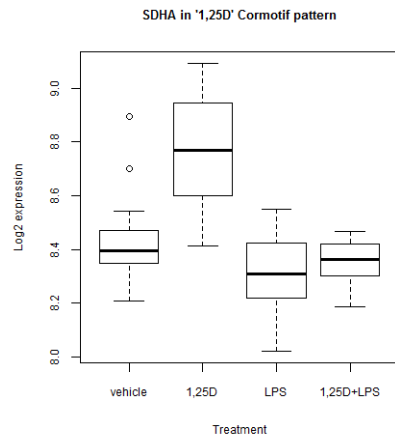


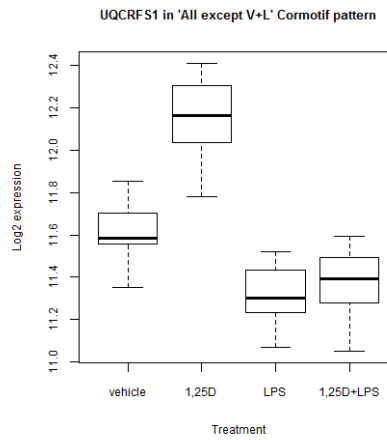
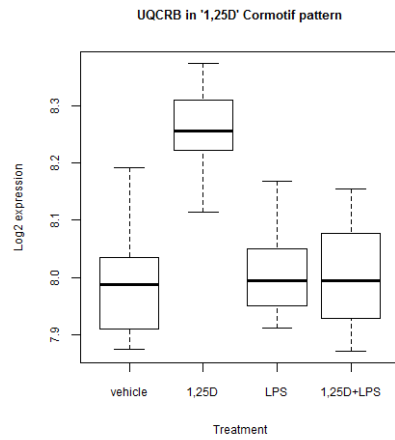
A: Complex I



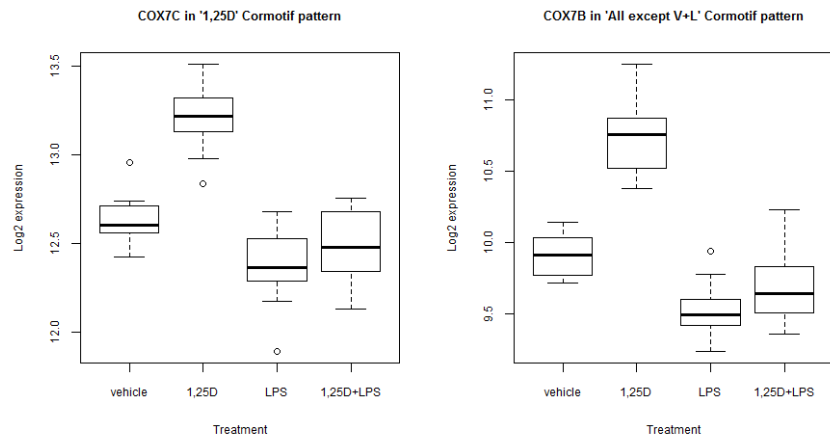
B: Complex II



C: Complex III



D: Complex IV



E: Complex V

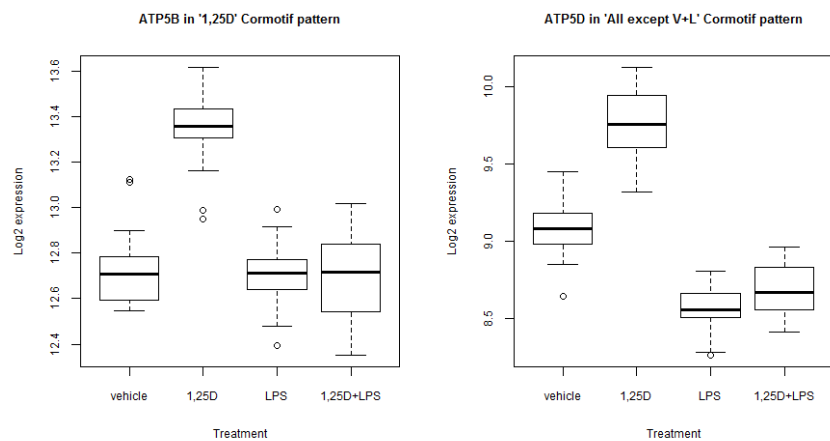


Figure S6: Boxplots of genes in the Oxidative Phosphorylation pathway clustered in the “1,25D” and “All except V+L” Cormotif patterns with treatment-specific response patterns. Genes representing the five respiratory complexes are shown in **A-E**. Oxidative phosphorylation pathway genes in the “1,25D” pattern were responsive only to 1,25D, while genes in the “All except V+L” pattern were responsive to both 1,25D and LPS in opposite directions.