**KEGG Pathway Enrichment (Rich Factor): AAB\_vs\_DGB** Neuroactive ligand-receptor interaction Cytoskeleton in muscle cells Neuroactive ligand signaling Gene Count 10 Cytokine-cytokine receptor interaction 20 30 ECM-receptor interaction 40 Pathway Glycine, serine and threonine metabolism -log10(adj. p-value) 3.5 Biosynthesis of amino acids 3.0 2.5 One carbon pool by folate 2.0 Arginine and proline metabolism Cysteine and methionine metabolism Alanine, aspartate and glutamate metabolism 0.004 0.002 0.006 0.008 Rich Factor