**KEGG Pathway Enrichment (Rich Factor): AAB\_vs\_AAL** Cytoskeleton in muscle cells Glycolysis / Gluconeogenesis Gene Count Carbon metabolism 16 20 Pathway Biosynthesis of amino acids -log10(adj. p-value) 5 Starch and sucrose metabolism Glycine, serine and threonine metabolism Pentose phosphate pathway 0.0015 0.0020 0.0010 0.0025 0.0030 Rich Factor