

PROTEOMICS

Supporting Information for Proteomics

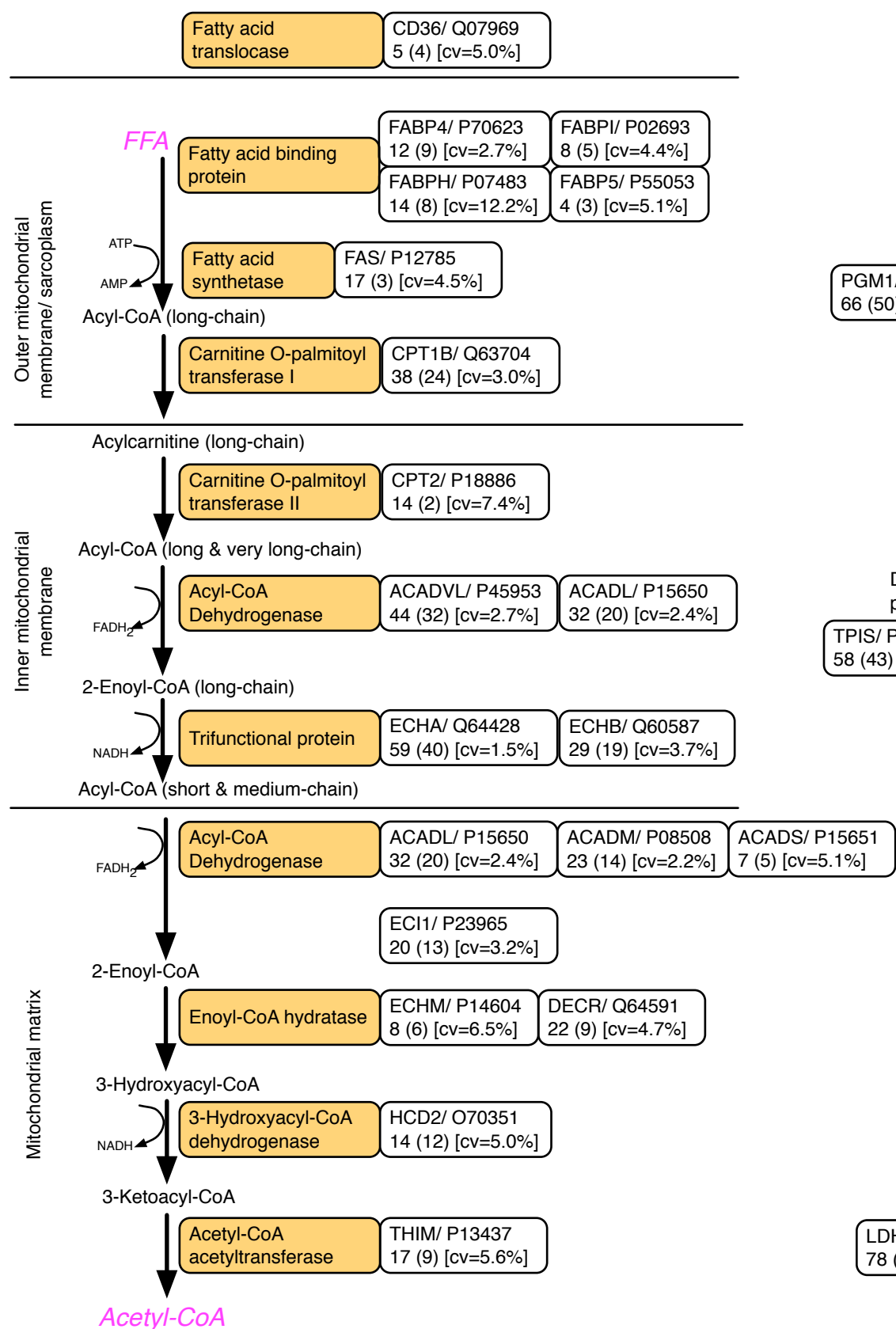
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Label-free profiling of skeletal muscle using high-definition mass spectrometry

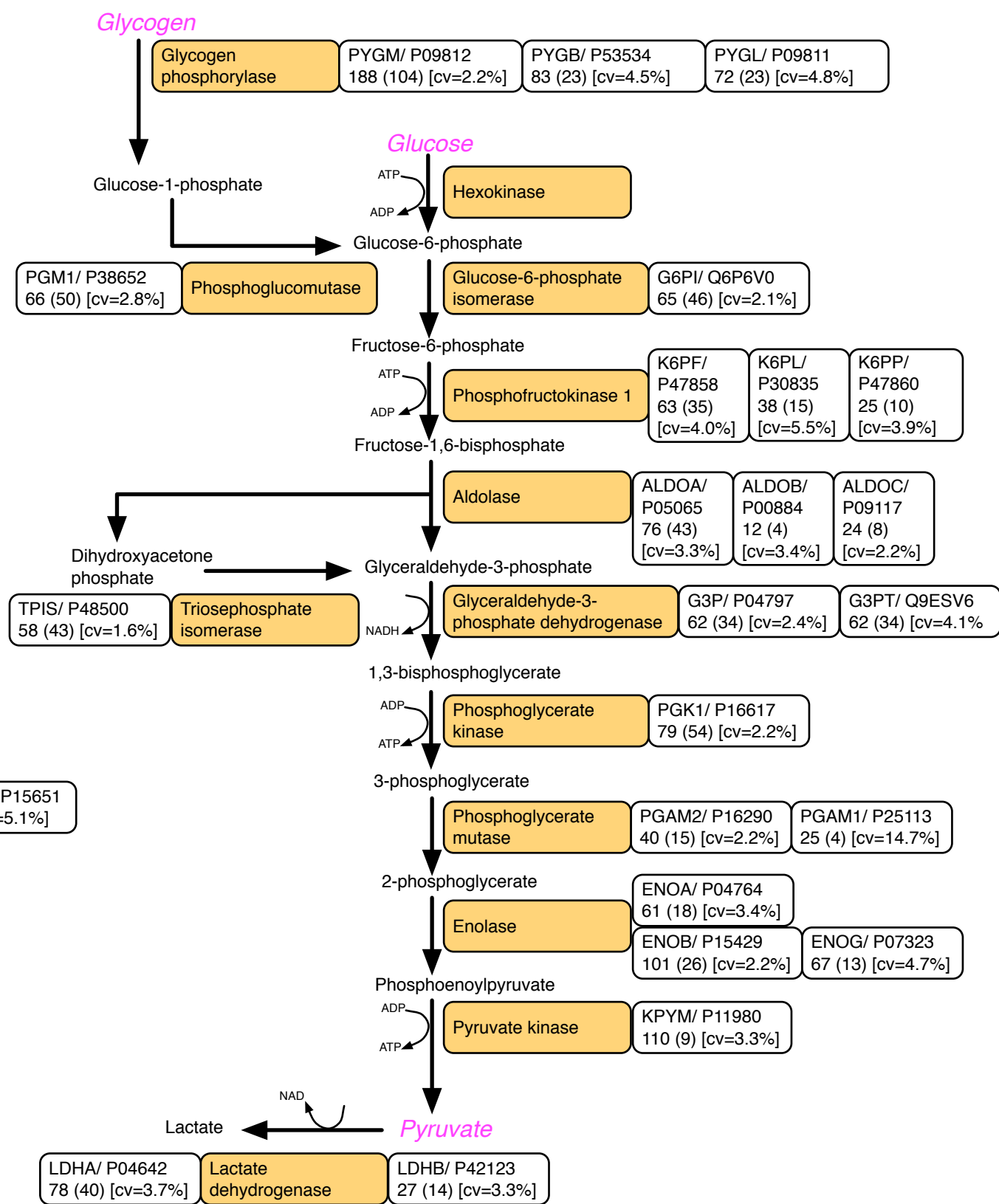
Fatty acid oxidation

(average coefficient of variation = 4.47 %)



Glycolysis

(average coefficient of variation = 3.76 %)



TCA cycle

(average coefficient of variation = 4.26 %)

