Supplementary Materials: Comparing Simplification Strategies for the Skeletal Muscle Proteome

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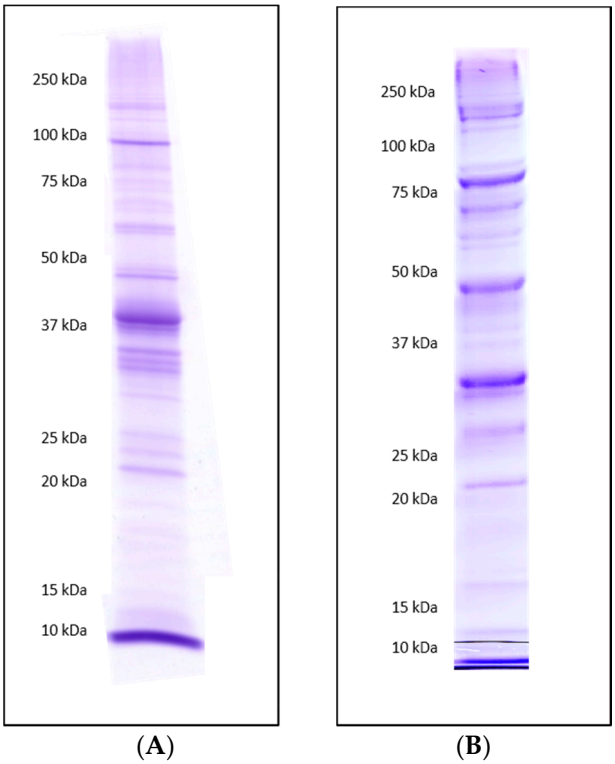

Figure S1. Visualisation of ProteoMiner Bead Equalisation. Proteins were subjected to equalisation using bead technology in order to reduce the dominance of highly abundant proteins. Panel **A** shows the pre-fractionated skeletal muscle soluble proteome with the simplified proteome in Panel **B**.

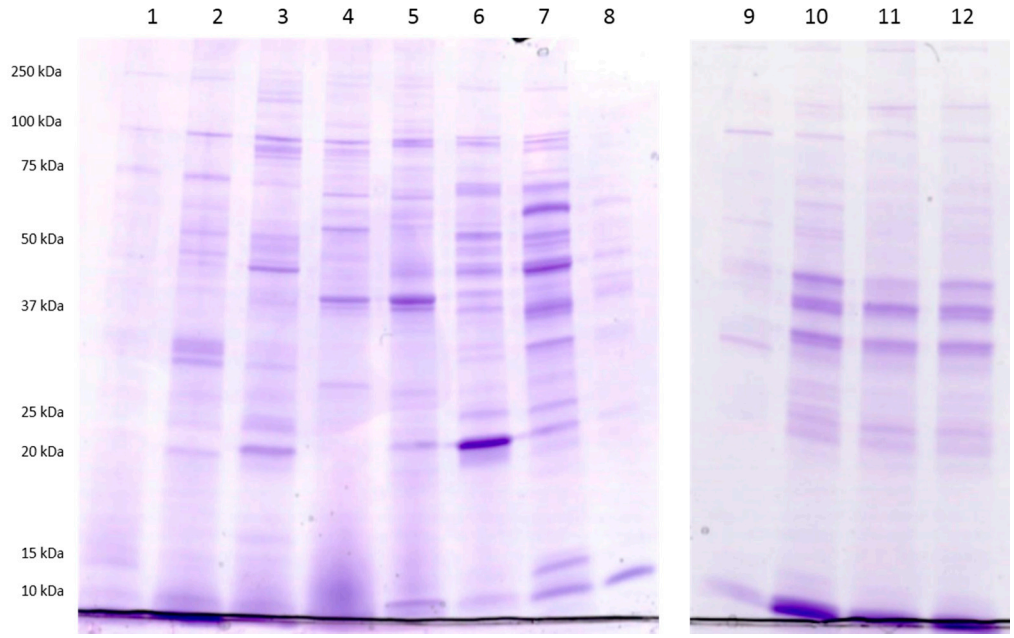

Figure S2. In-Solution Isoelectric Focussing of the Skeletal Muscle Proteome. Proteins were separated according to isoelectric point and each fraction further separated by 1D-SDS-PAGE.

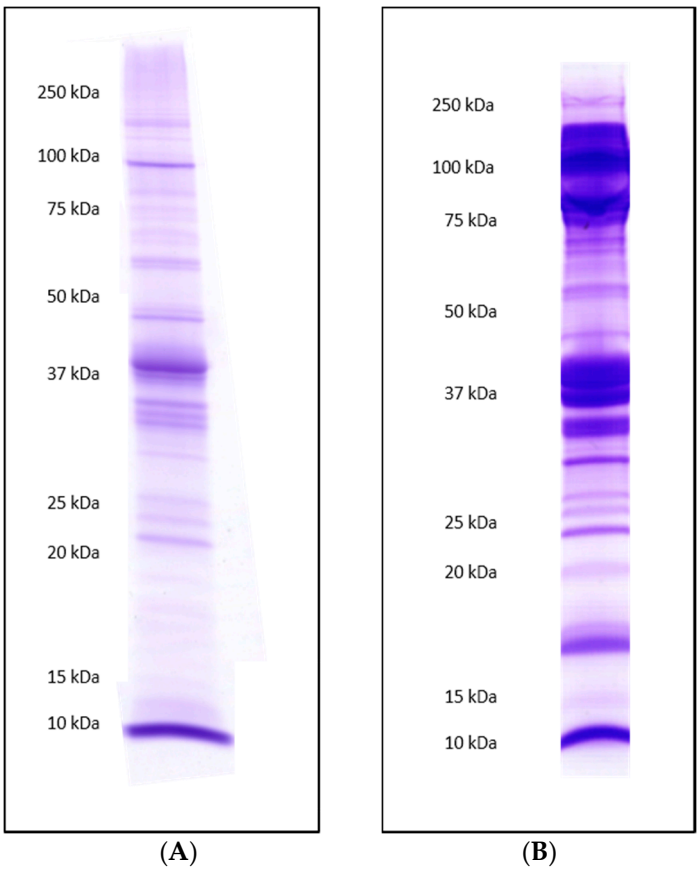

Figure S3. Protein Extraction for FASP Analysis. Proteins were solubilised prior to FASP analysis. Panel **A** shows the skeletal muscle soluble proteome used for standard 1D-SDS-PAGE with the extracted proteome for FASP analysis shown in Panel **B**.