Supplementary Information (SI) for Molecular Omics.

Supplementary Information

Proteomic and metabolic profiling reveals molecular phenotype associated with chemotrophic growth of *Rubrivivax benzoatilyticus* JA2 on L-tryptophan

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Supplementary Information

Supporting Information Figures …… No S1 – S5

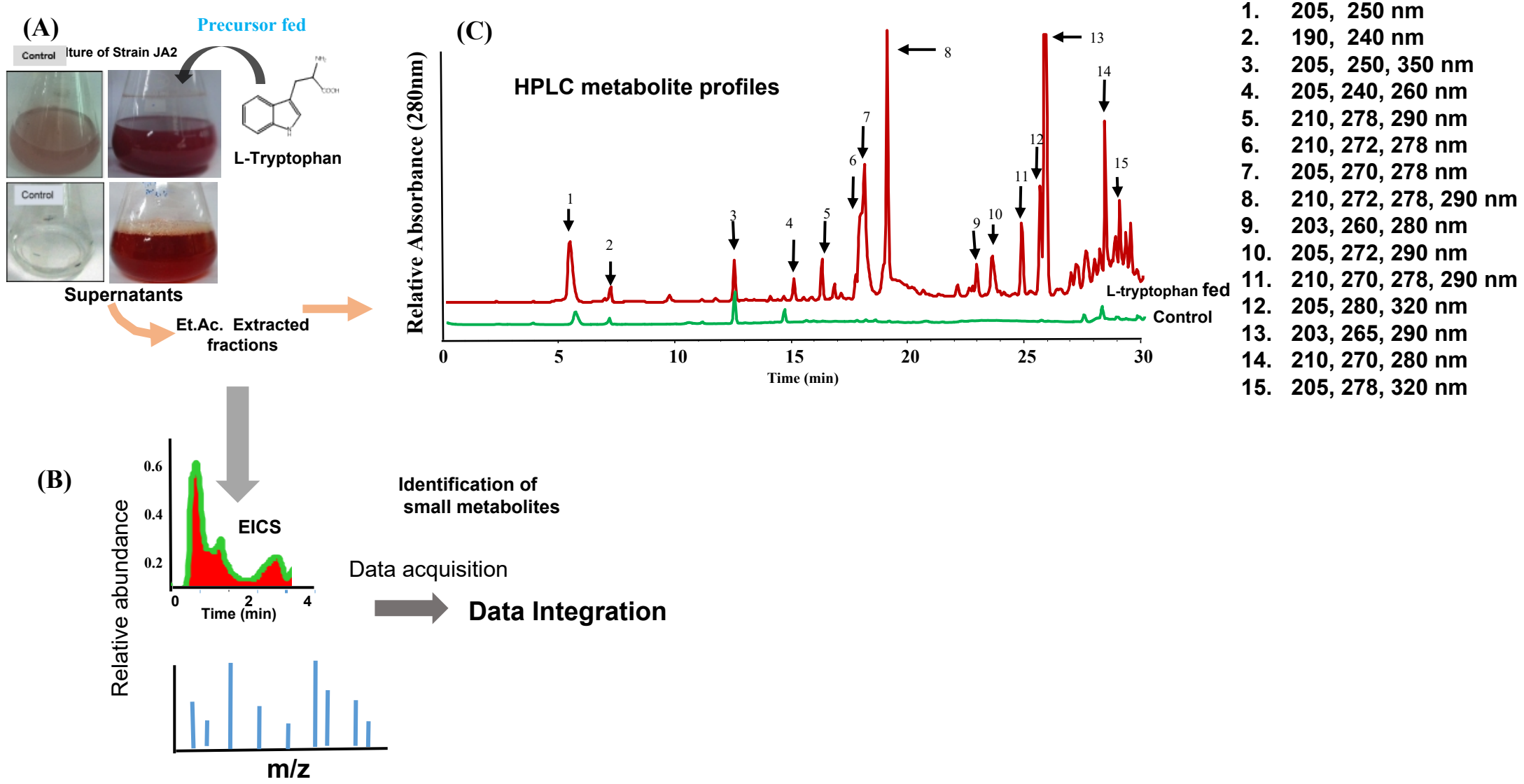

Fig. S1: Workflow of metabolic profiling of L-tryptophan fed culture supernatant and identification of small indole derivatives metabolites.

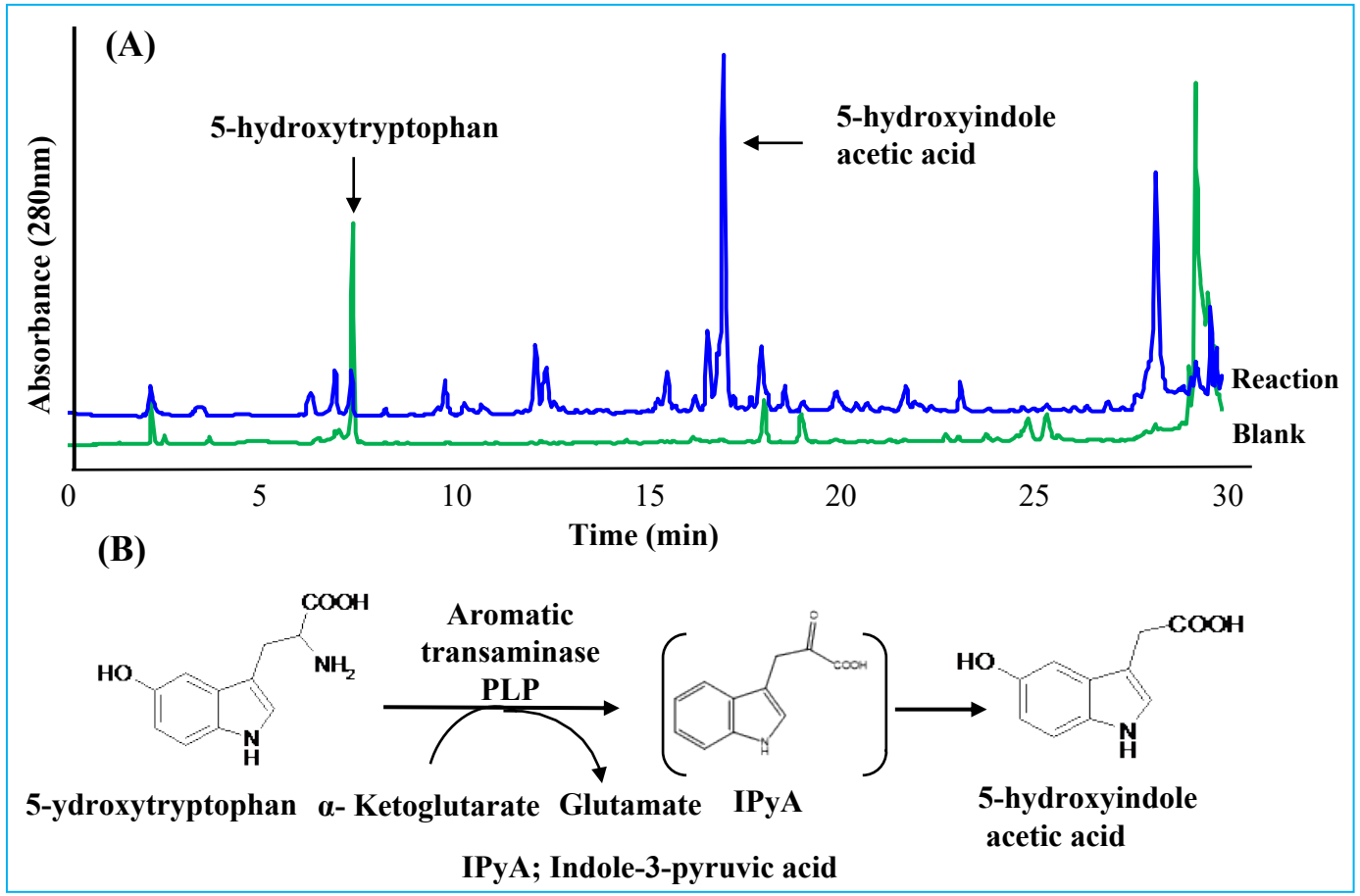

Fig. S2: HPLC chromatogram showing the aromatic aminotransferase enzyme activity **(A)** Transamination mediated catabolism of hydroxytryptophan in strain JA2 **(B)**.

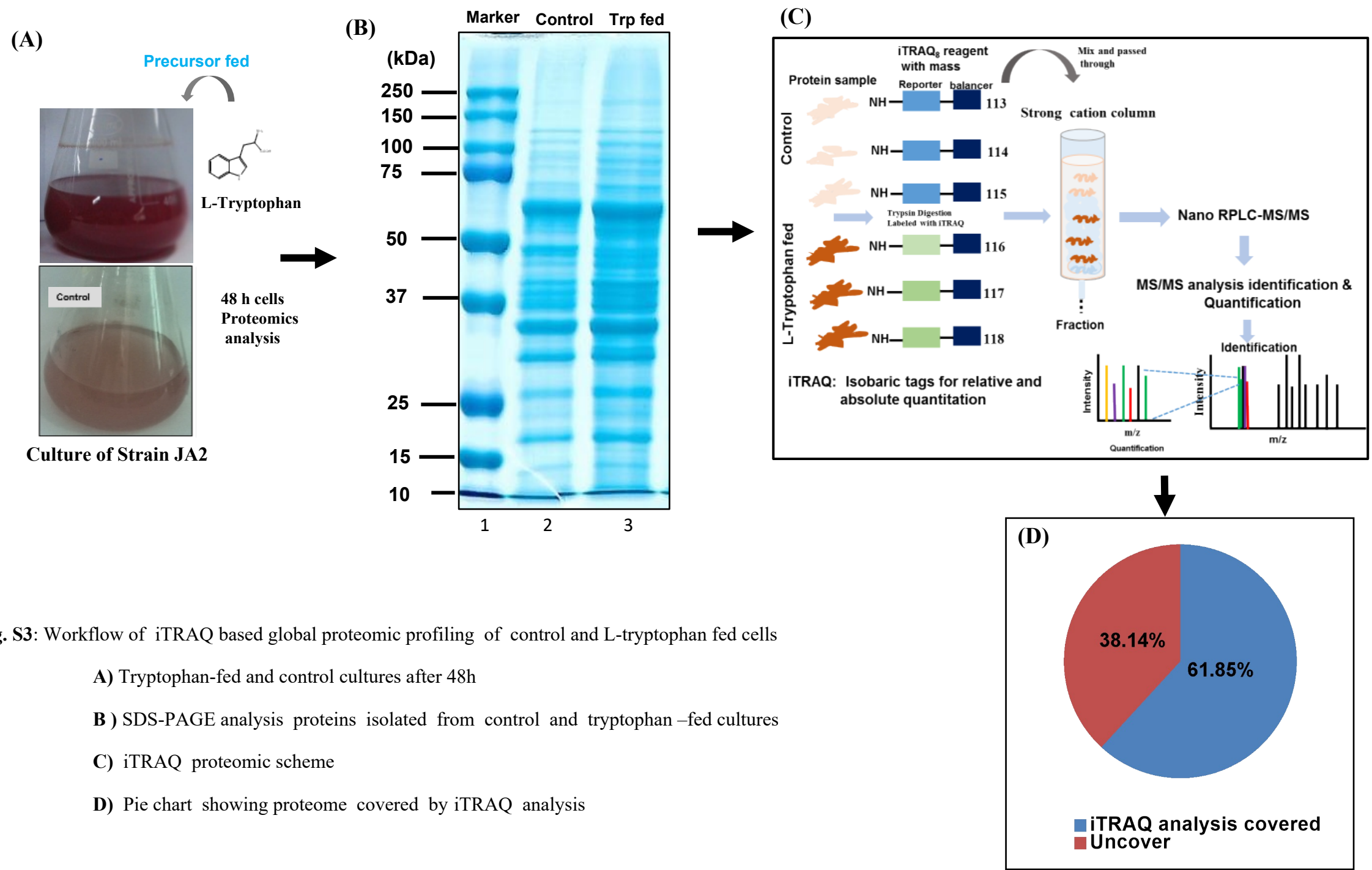

Fig. S3: Workflow of iTRAQ based global proteomic profiling of control and L-tryptophan fed cells

A) Tryptophan-fed and control cultures after 48h

B) SDS-PAGE analysis proteins isolated from control and tryptophan –fed cultures

C) iTRAQ proteomic scheme

D) Pie chart showing proteome covered by iTRAQ analysis

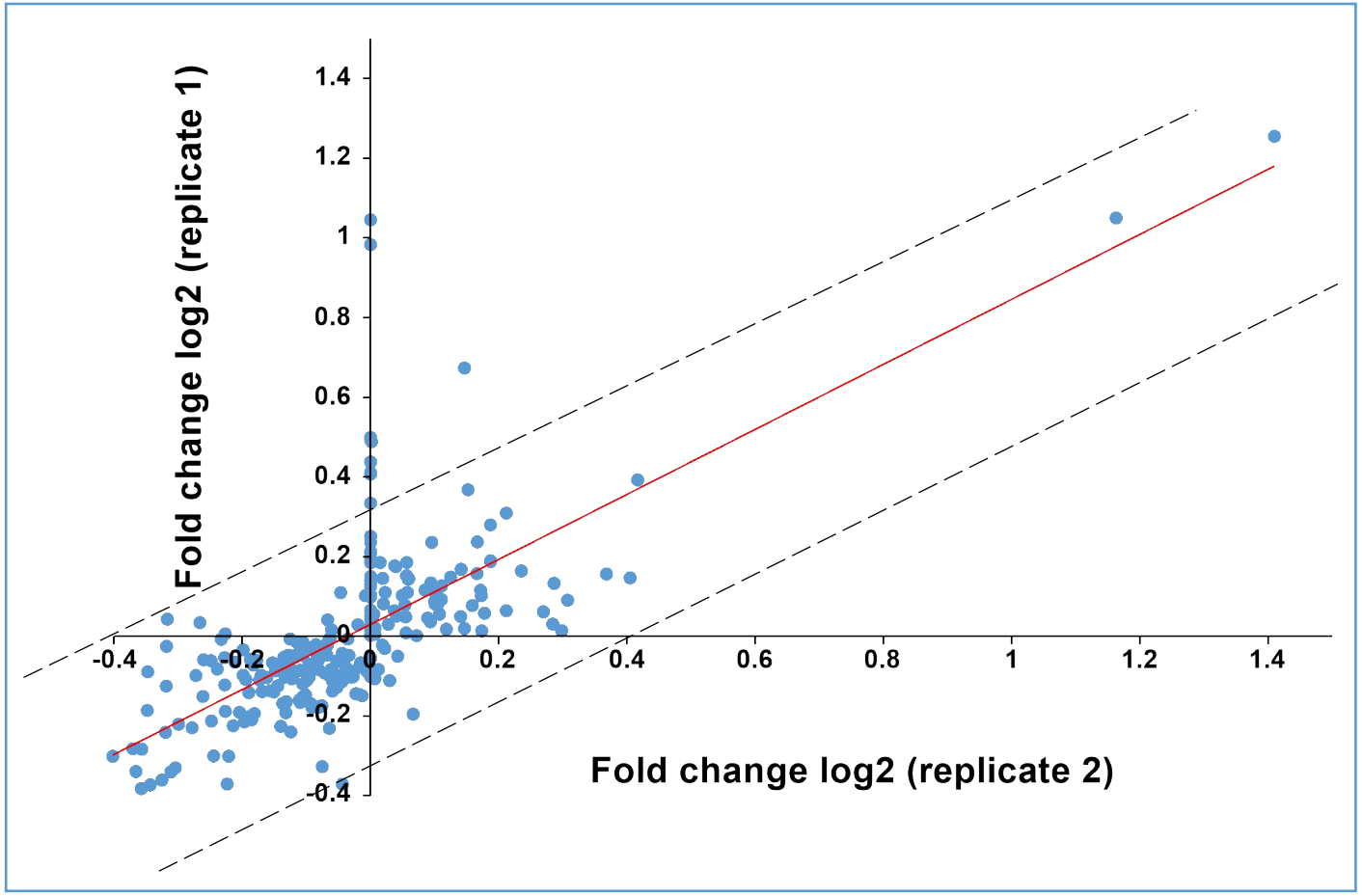

Fig. S4: Linear regression analysis deferential regulated proteins of two replicates

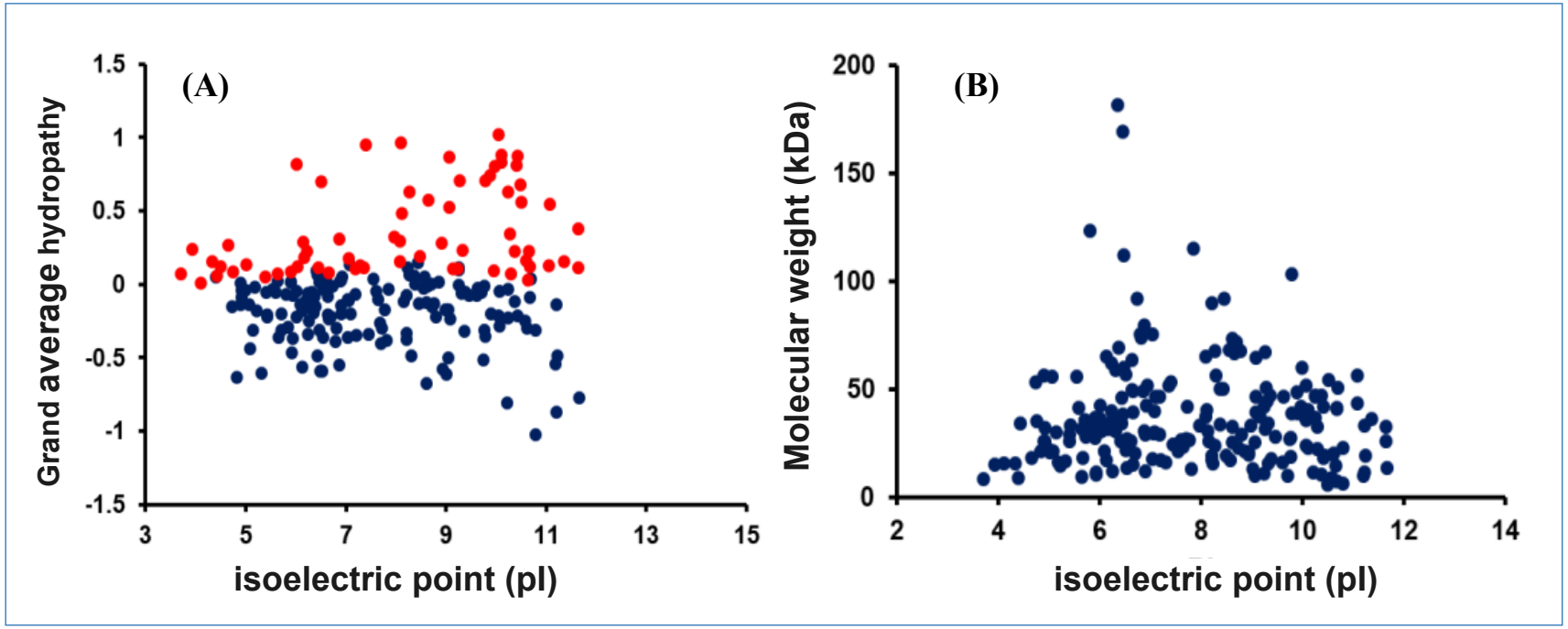

Fig. S5: *In-silico* characterization of differential regulated proteins of L-tryptophan fed chemotrophic conditions by proteomic Expasy tool. (A) Hydropathy of proteins, (B) Molecular weight *vs* pI plot