



**S2. Fig: Correlation of log(2) expression values of transcriptomic and proteomic of two biological replicates under the three experimental conditions.**

<sup>a</sup>. Sum of total 100-mer alignment per gene; Panel A/B: Waste fatty acids exponential phase culture biological replicate 1 and 2; Panel C/D: Waste glycerol exponential phase culture biological replicate 1 and 2; Panel E/F: Waste glycerol stationary phase culture biological replicate 1 and 2.