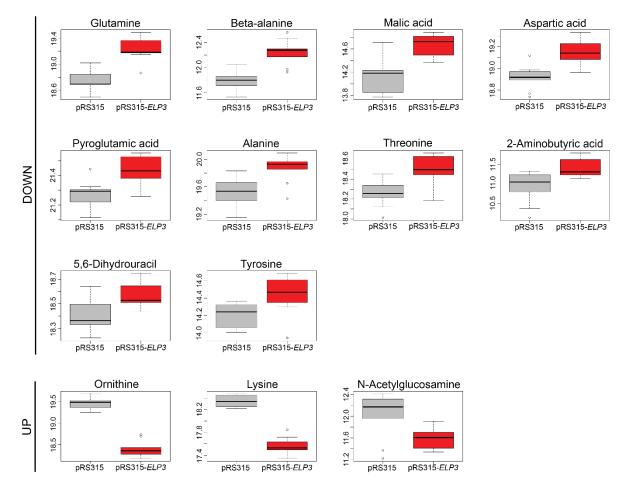
Loss of ncm⁵ and mcm⁵ wobble uridine side chains results in an altered metabolic profile

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Online Resource 3. Levels of identified metabolites with a VIP score above 1 in the PLS-DA comparing the $elp3\Delta$ -l.c.-empty and $elp3\Delta$ -l.c.-ELP3 strains grown at 30°C. The $elp3\Delta$ strain containing either an empty low copy pRS315 vector or a pRS315 vector carrying the wild-type ELP3 gene was grown logarithmically to an OD₆₀₀ of ~0.5 and harvested (see material and methods). Metabolites were extracted and then quantified using GC-TOF-MS. Values obtained were normalized and log2-transformed, and metabolic alterations were analyzed using PLS-DA. Metabolites shown have a VIP score >1. Metabolites were classified as either increasing (UP) or decreasing (DOWN) in the $elp3\Delta$ strain containing an empty low copy pRS315 vector when compared to the $elp3\Delta$ strain carrying the wild-type ELP3 gene. Boxplots were generated using R software with the Y-axis displaying the relative intensity in log2-scale.