

## Supporting Information

# Enhanced Metabolome Coverage and Evaluation of Matrix Effects by the Use of Experimental-Condition-Matched $^{13}\text{C}$ -Labeled Biological Samples in Isotope-Assisted LC-HRMS Metabolomics

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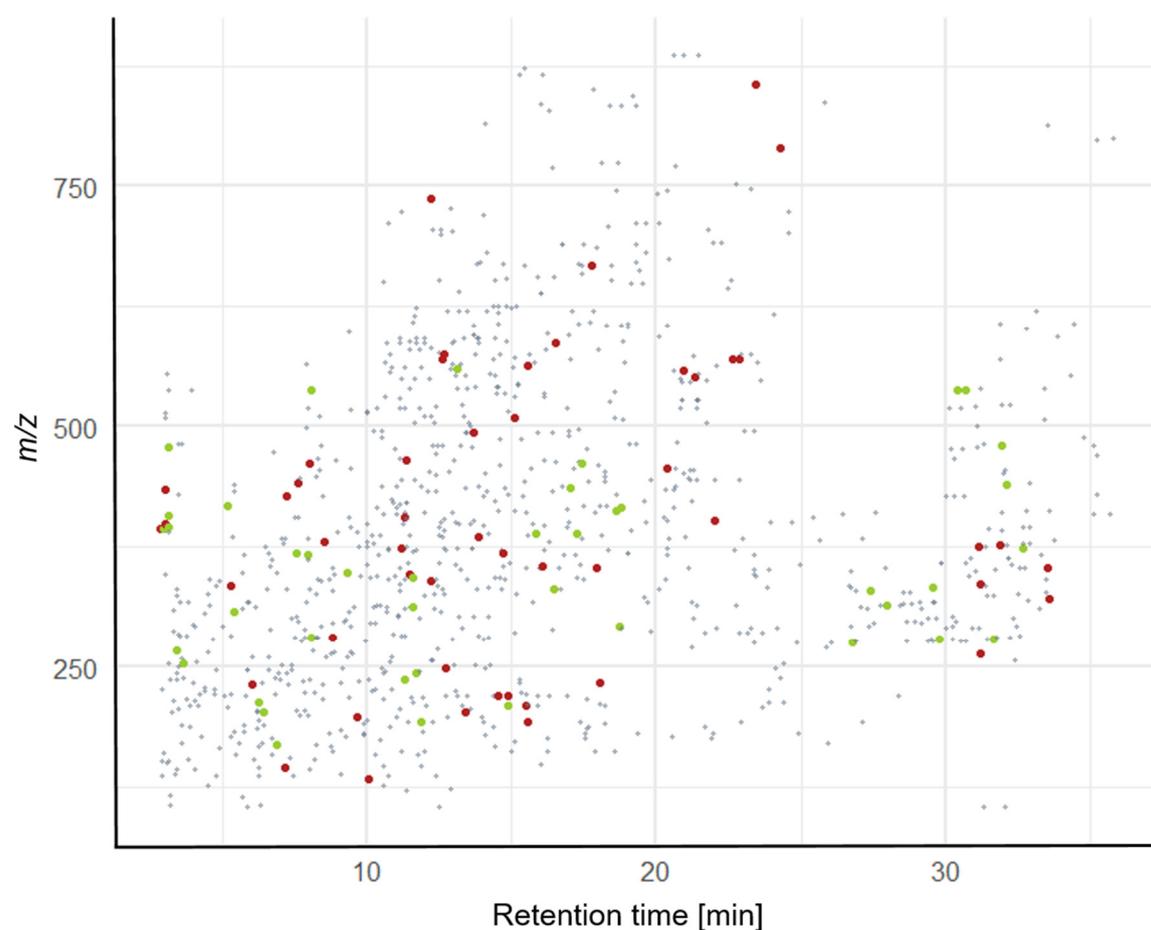
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**Figure S1.** Feature map of detected metabolites. Different color code represents metabolites that have shown different results for the significantly differing metabolites in the comparative quantification test (Toxin versus Control). In red are highlighted those which are significantly different (between

Toxin and Control) only if absolute native abundances are used, and in green only those obtained after normalized (native to labeled) abundances are applied.