

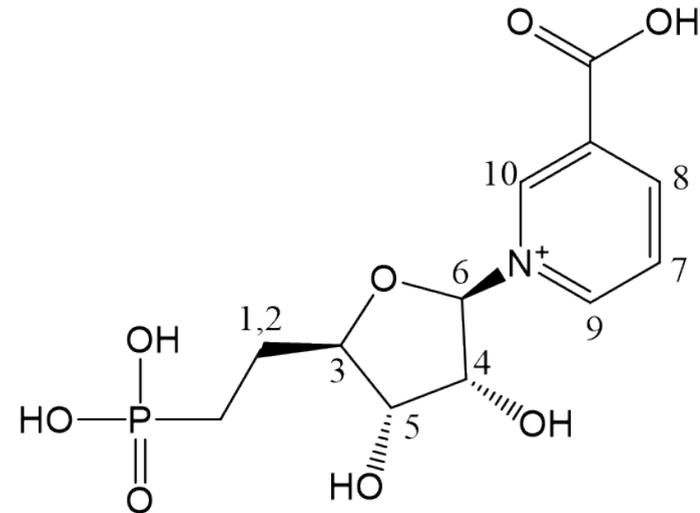
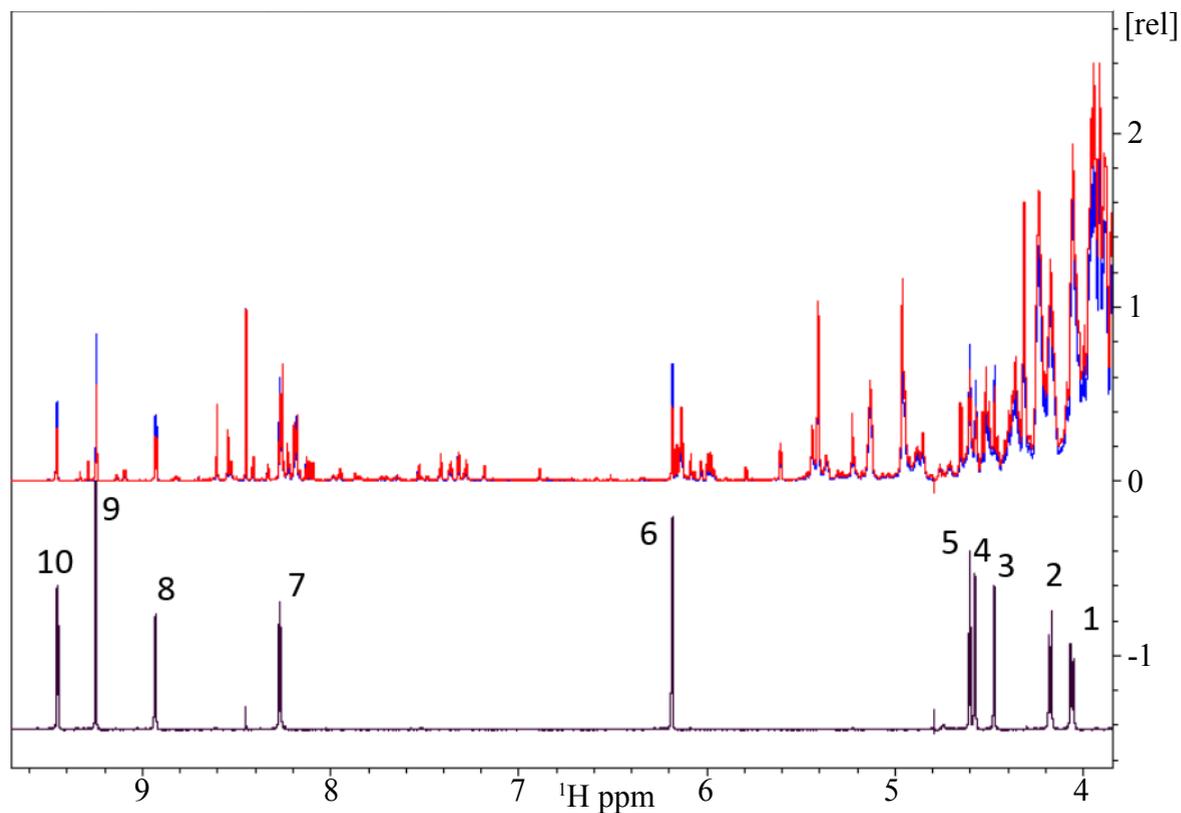
Supplementary Material

**Integrated metabolomics and transcriptomics suggest  
the global metabolic response to 2-aminoacrylate stress  
in *Salmonella enterica***

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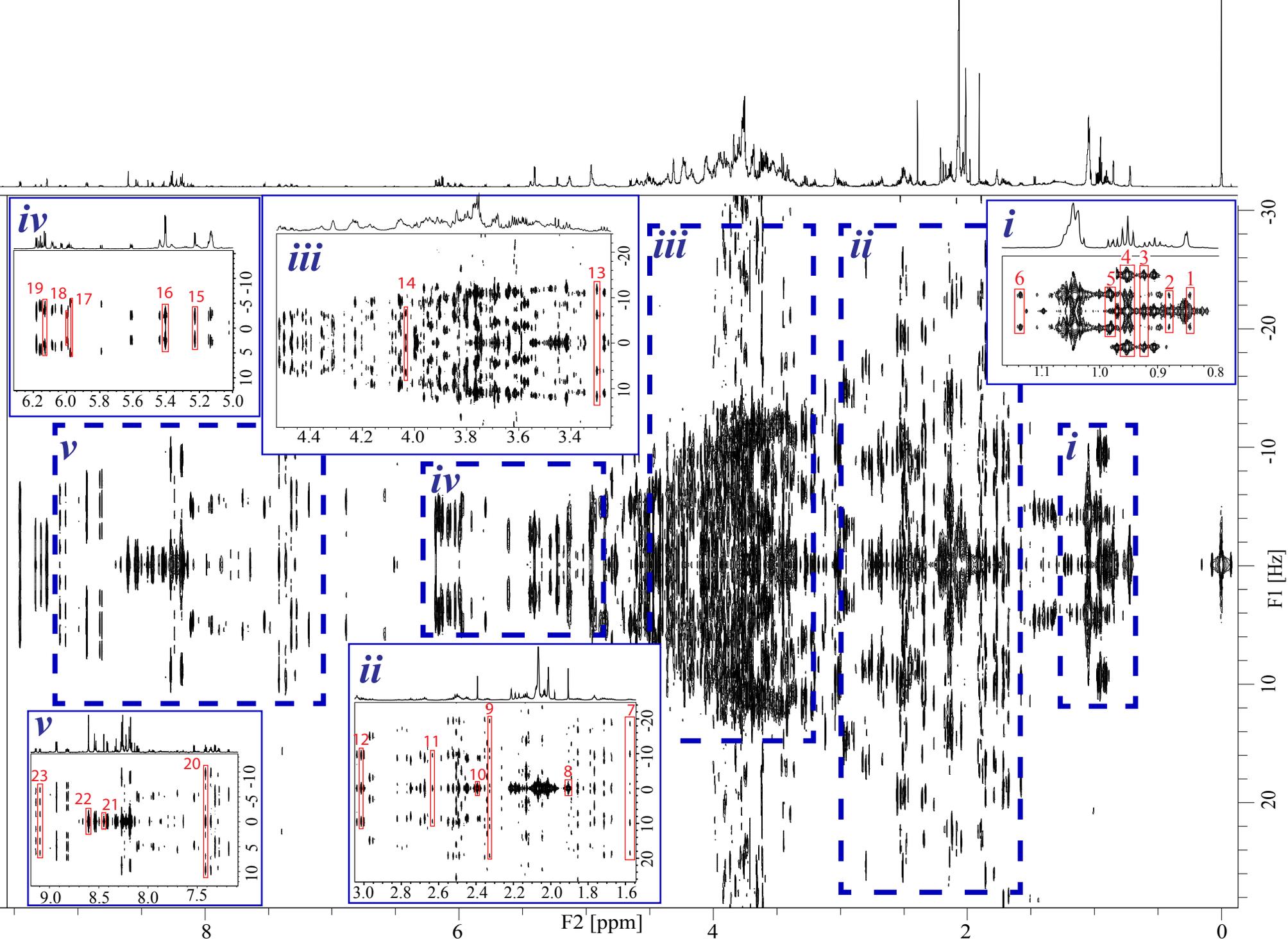
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Nicotinate mononucleotide (NMN)

Atom	<sup>1</sup> H peak	<sup>13</sup> C peak
1	4.05	65.87
2	4.17	65.87
3	4.47	73.59
4	4.57	90.34
5	4.6	80.36
6	6.18	102.75
7	8.27	131.26
8	8.93	149.55
9	9.25	144.39
10	9.45	143.93

**Figure S1. Nicotinate mononucleotide identification by sample spiking.** The <sup>1</sup>H NMR spectrum was obtained for the pooled metabolite extraction samples with (blue line) and without (red line) the inclusion of 150 μM pure NMN standard. The <sup>1</sup>H NMR spectra of NMN alone is provided below (black line). Labeled peaks correspond to labeled atoms on the structural diagram. Exact <sup>1</sup>H NMR and <sup>13</sup>C peak assignments are listed in the adjacent table.



**Figure S2. Representative 2D J-RES Spectrum.** Features outlined with red boxes were used for quantification: 1) 2-isopropylmalic acid, 2) Pantothenic acid, 3) Ile, 4) Leu, 5) Val, 6) Thr, 7) N-acetylputrescine, 8) Acetate, 9) Glu, 10) Succinate, 11) Met, 12) Lys, 13) GSSG, 14) Ser, 15) Glucose, 16) Maltose, 17) UTP, 18) CMP, 19) dCMP, 20) Phe, 21) Formate, 22) AMP, 23) NAD