

Metabolite Spectral Accuracy on Orbitraps

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SUPPORT INFORMATION

Figure S1. Spectral accuracy of glutathione (m/z 307) and serine (m/z 104)

Figure S2. Extracted ion chromatograms of NAD at high and low mass resolution

Figure S3. Mass spectra of NAD at different resolutions

Figure S4. Spectral accuracy of glutathione and serine as a function of mass scan range

Figure S5. The extracted ion chromatogram of NAD under full scan or smaller m/z scan range (500-1000)

Figure S6. Zoomed mass spectra of M+1 peaks reveals no evidence of contaminating species

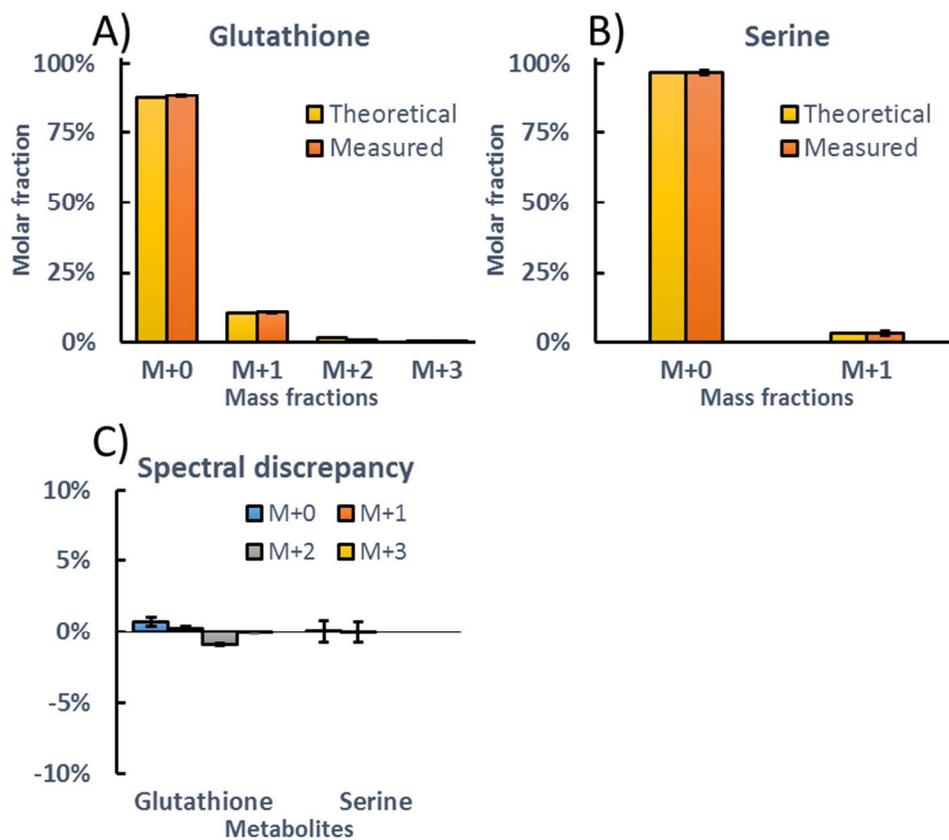


Figure S1. Spectral accuracy of glutathione (m/z 307) and serine (m/z 104). Measured and theoretical mass distribution of unlabeled A) glutathione and B) serine. C) Spectral discrepancies.

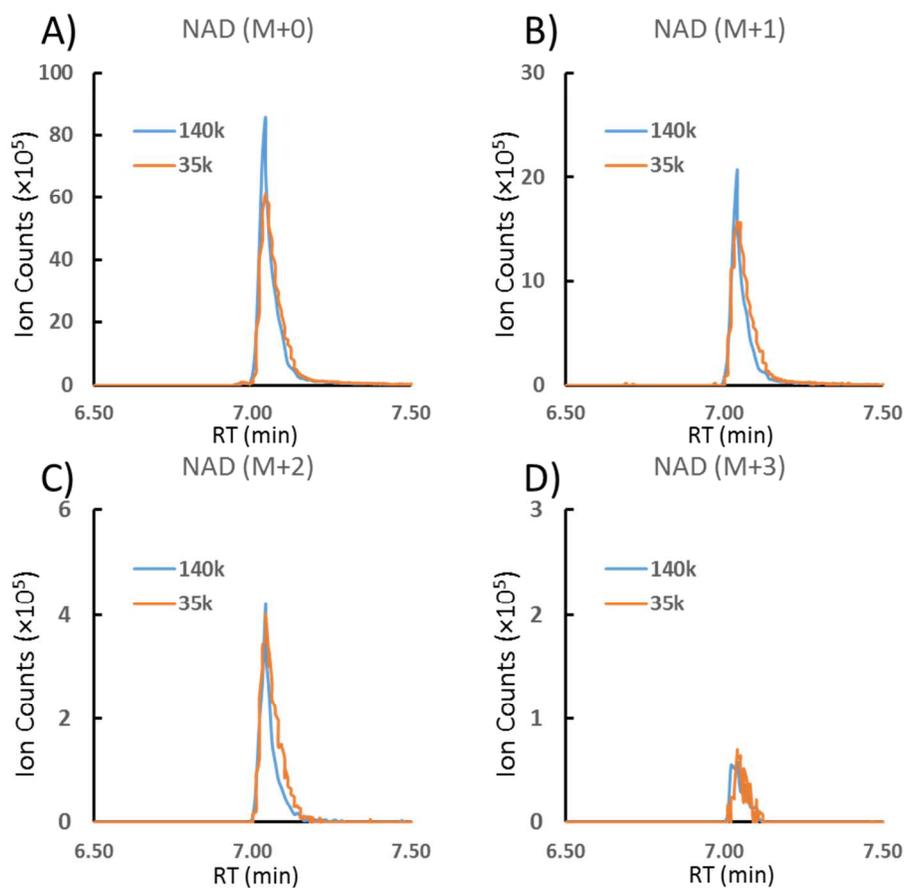


Figure S2. Extracted ion chromatograms of NAD at high and low mass resolution. A) M+0, B) M+1, C) M+2, D) M+3.

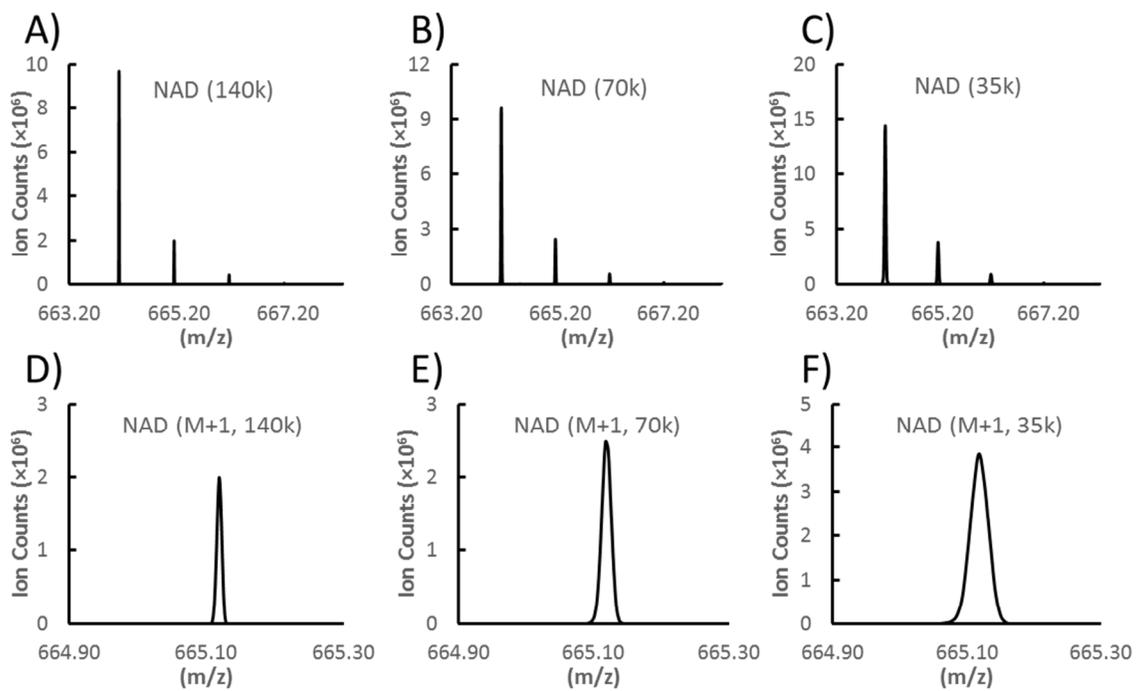


Figure S3. Mass spectra of NAD at different resolutions. A-C) Resolution 140k, 70k and 35k. D-F) Zoomed NAD M+1 peak.

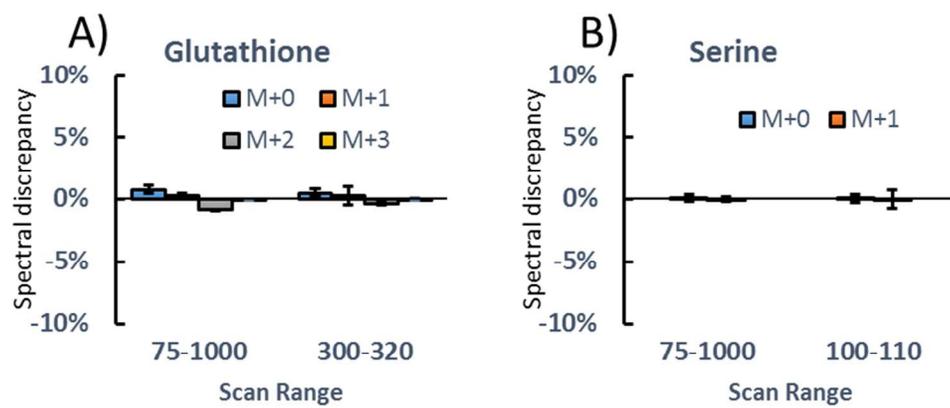


Figure S4. Spectral accuracy of A) glutathione and B) serine as a function of mass scan range.

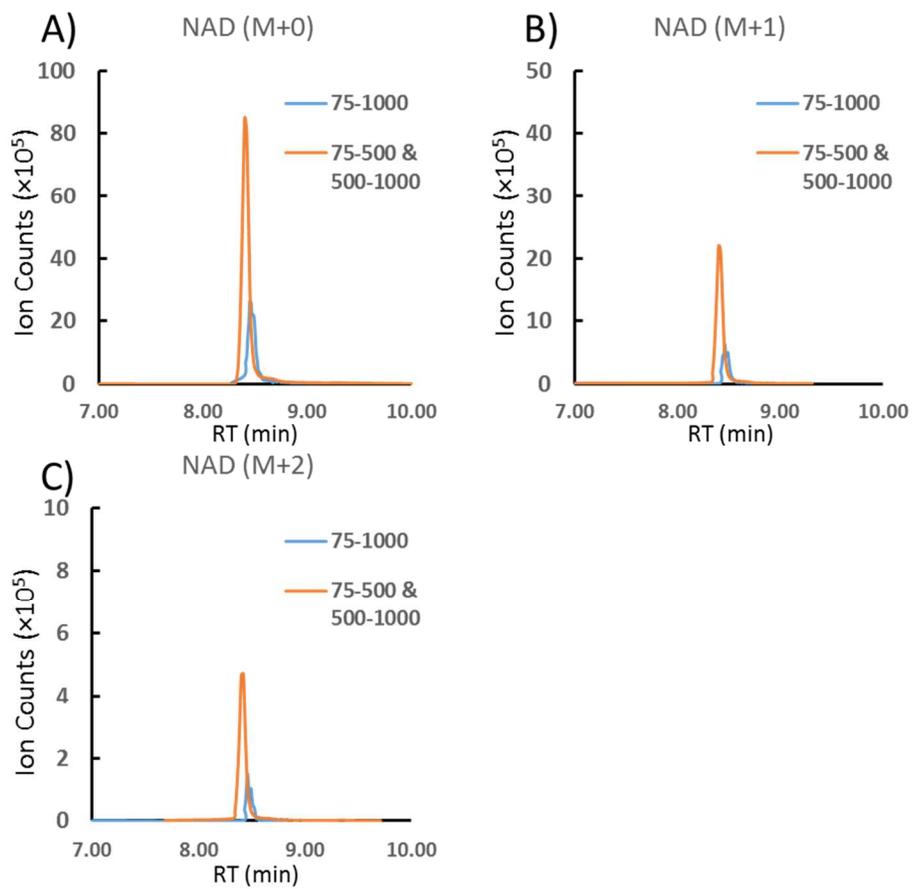


Figure S5. The extracted ion chromatogram of NAD under full scan or smaller m/z scan range (500-1000).

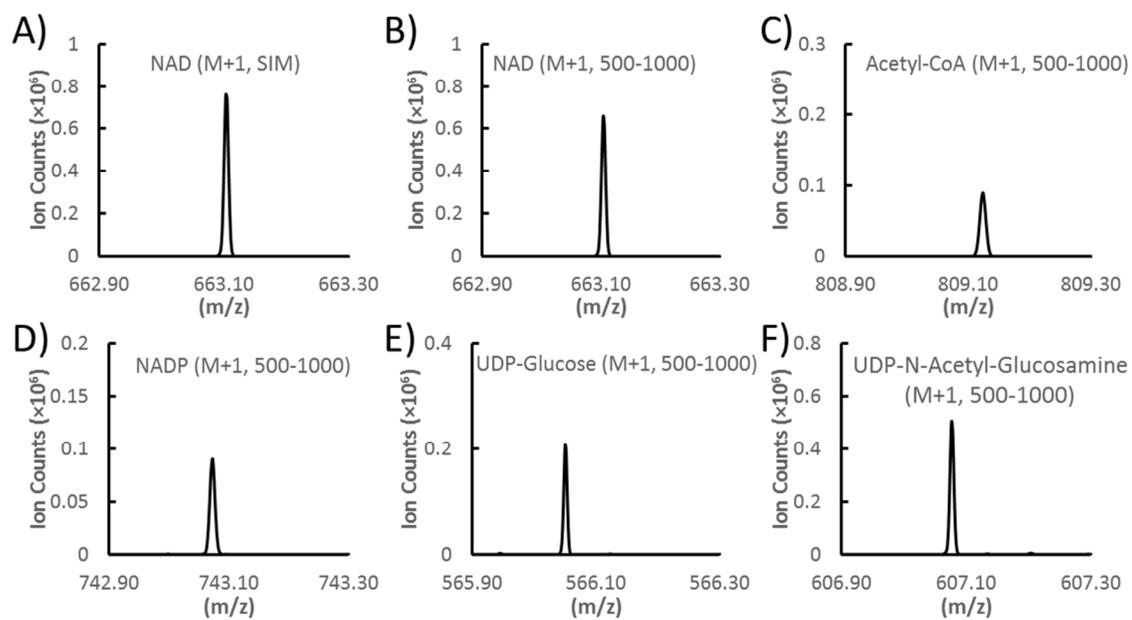


Figure S6. Zoomed mass spectra of M+1 peaks reveals no evidence of contaminating species. Values in parentheses show the scan range. A, B) NAD, C) Acetyl-CoA, D), NADP, E) UDP-glucose, F) UDP-N-acetyl-glucosamine.