



C

	<i>S. cerevisiae</i> 0.005mM Spike
4-HBA	126070
VA	722906
SA	539391
CA	2525796
GGE	1262382

Fig. S2: A. Representative mass spectra (QToF-MS) for each uptake compound (4-HBA, VA, SA, CA, and GGE) in *S. cerevisiae* lysates. Integrated peak for internalization experiments is marked with a red arrow. DMSO vehicle control mass spectra are shown at the retention time for each compound of interest for peak comparison. B. Ion counts for each compound in *S. cerevisiae* cell lysates from 3 biological replicates. C. Ion counts for spiked *S. cerevisiae* lysate with indicated compounds at a concentration of 0.005mM. Spikes were used to calculate the relative intracellular concentrations in Fig. 1 from ion counts in Fig. S2C.