

Figure S5: Quantified data (via Chenomx) from 1D ^1H NMR of willow stem tissue normalised to the extracted metabolite pool. Concentrations given in micrograms per mg of the extractable pool. Data points correspond to mean \pm standard deviation from two biological replicates. Red = Resolution; Blue = Tora. T = material harvested from the top of the plant. M = material harvested from the middle of the plant. B = material harvested from the bottom of the plant.

