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Applications to serum, urine and tissue extracts.
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Abstract
The effective analysis of polar ionic metabolites by LC-MS, such as those encountered in
central carbon metabolism, represents a major problem for metabolic profiling that is not
adequately addressed using strategies based on either reversed-phase or HILIC methods. Here
we have compared analysis of central carbon metabolites on optimized methods using HILIC,
porous graphitic carbon or ion pair chromatography (IPC) using tributyl ammonium as IP

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