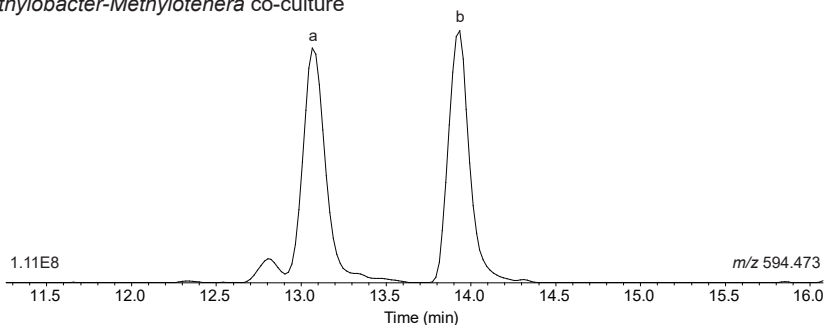
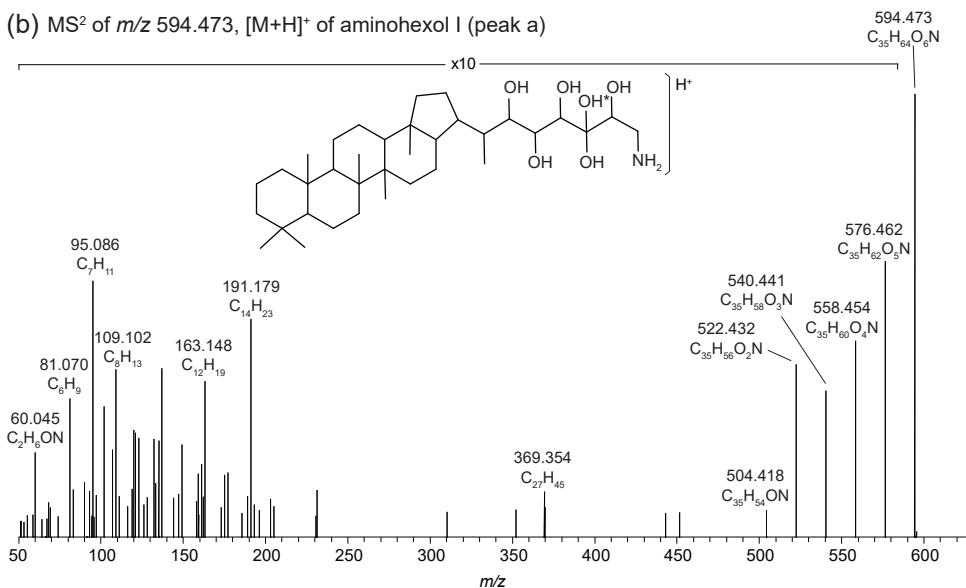


(a) *Methylobacter-Methylotenera* co-culture



(b) MS² of *m/z* 594.473, [M+H]⁺ of aminohexol I (peak a)



(c) MS² of *m/z* 594.473, [M+H]⁺ of aminohexol II (peak b)

