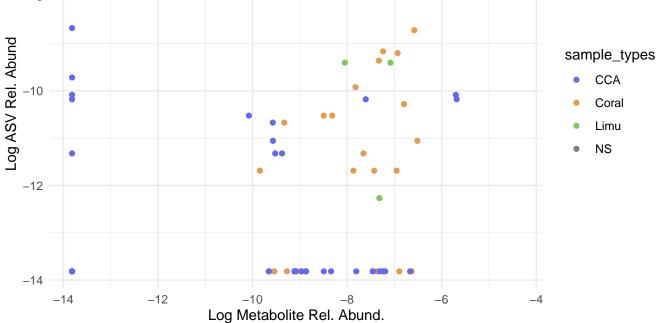
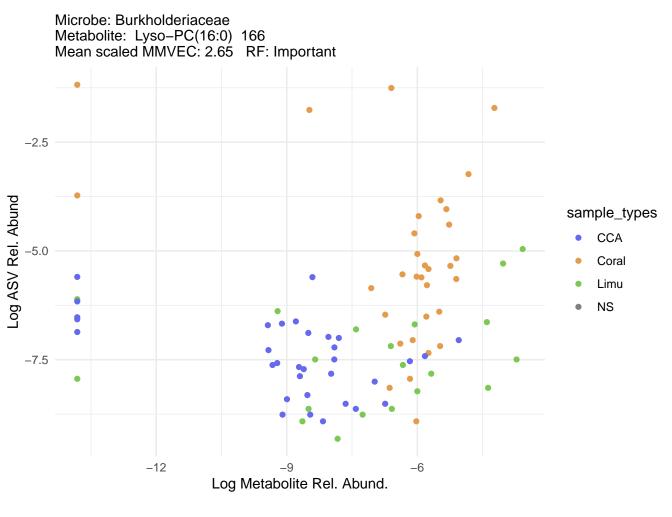


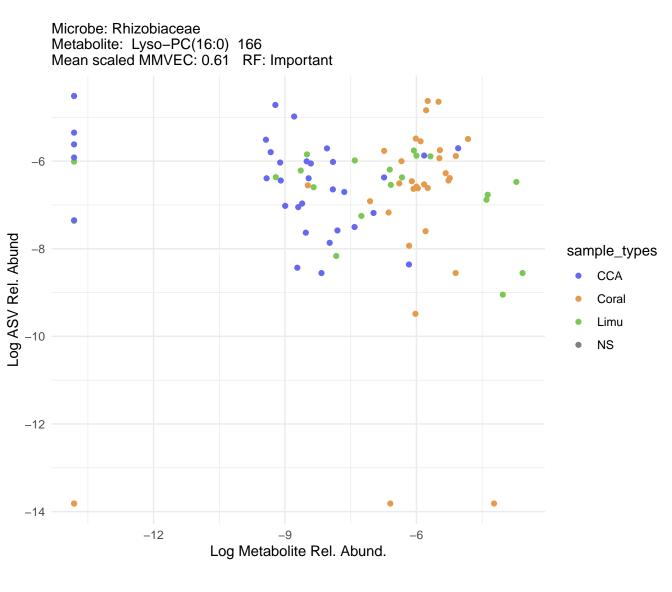
Microbe: Saprospiraceae
Metabolite: missing 146
Mean scaled MMVEC: 1.46 RF: Important

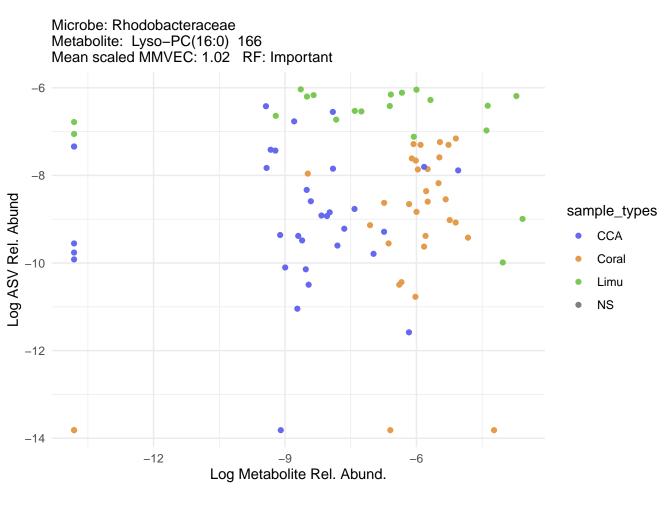
-8

sample\_types





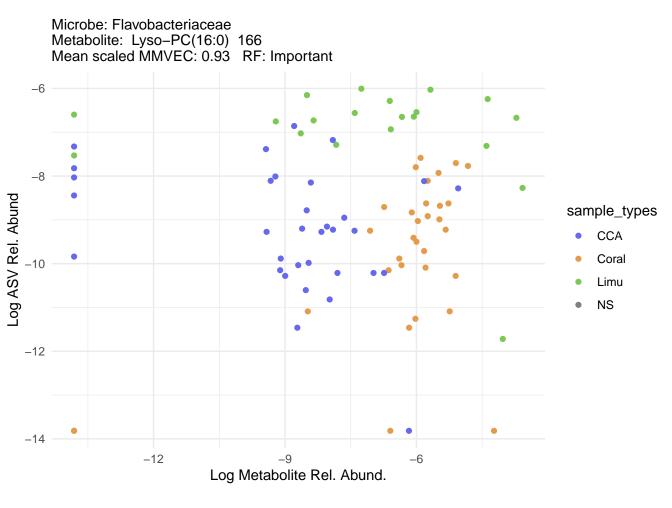


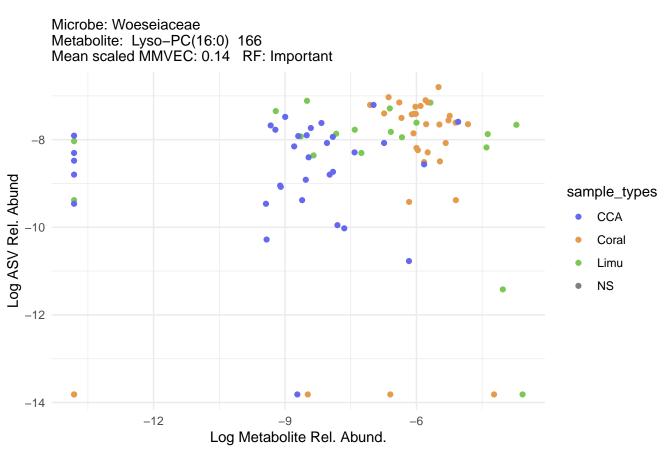


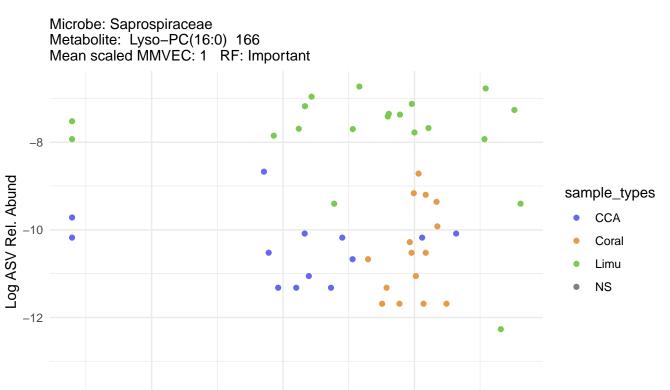
Microbe: Kiloniellaceae

Metabolite: Lyso-PC(16:0) 166 Mean scaled MMVEC: 0.27 RF: Important







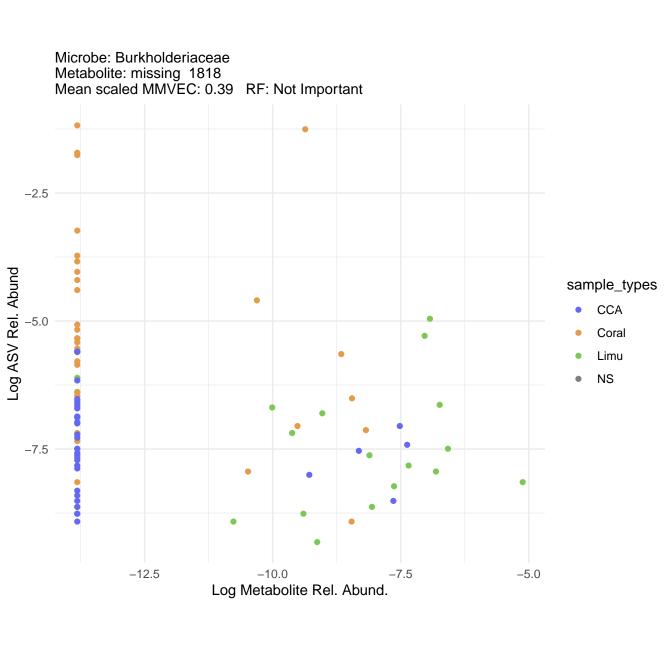


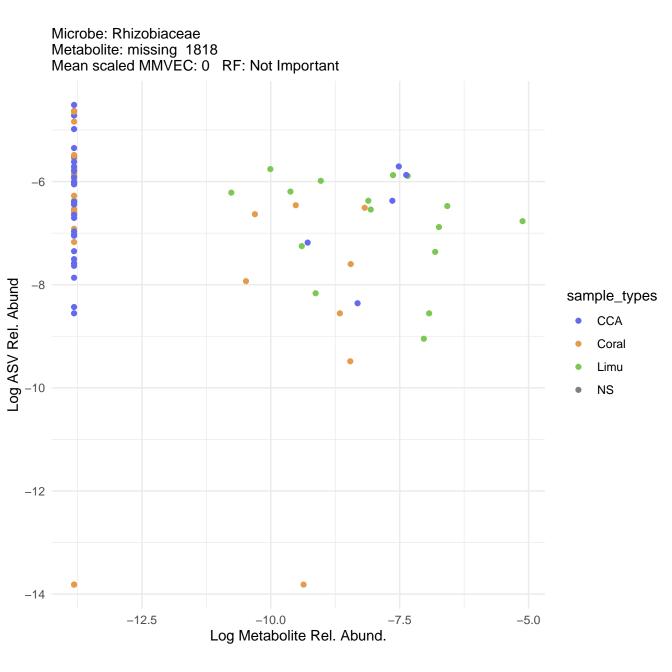
Log Metabolite Rel. Abund.

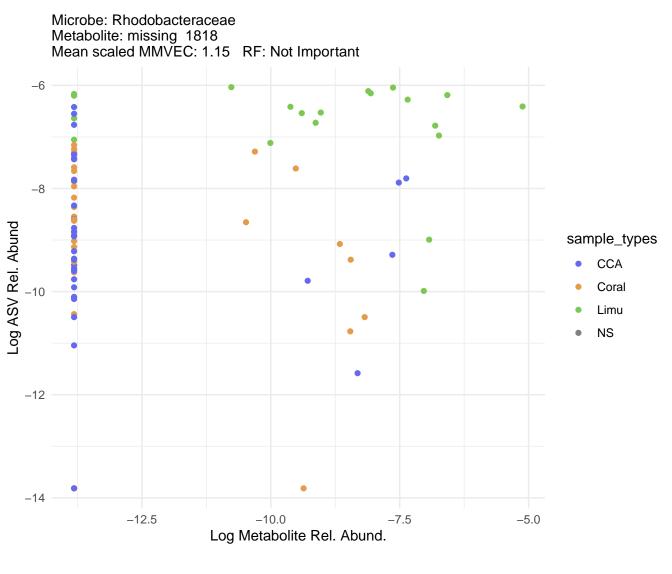
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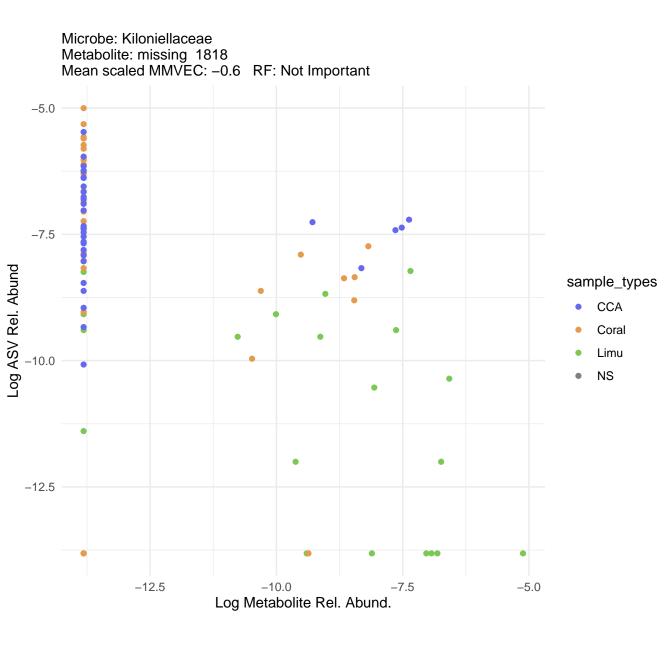
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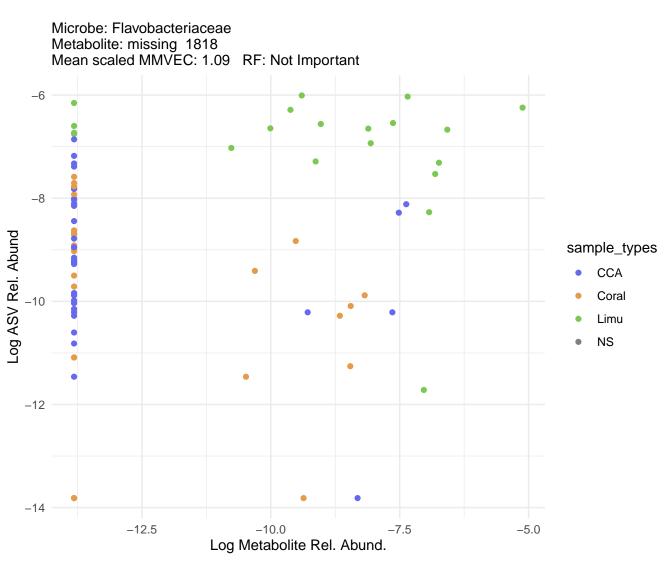
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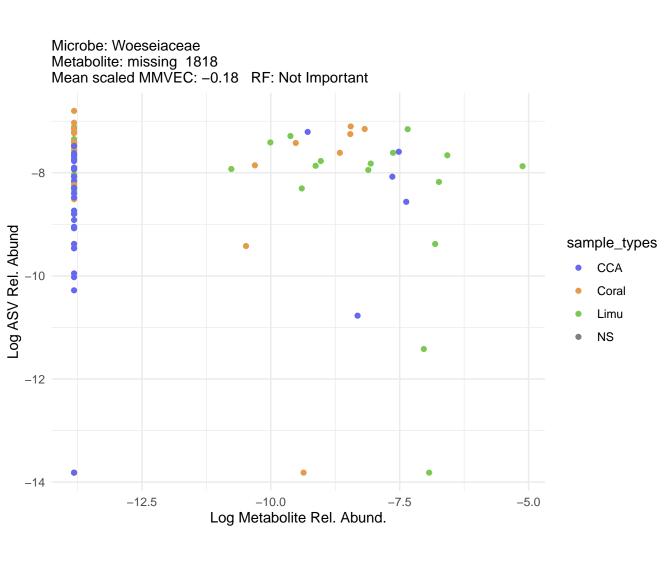


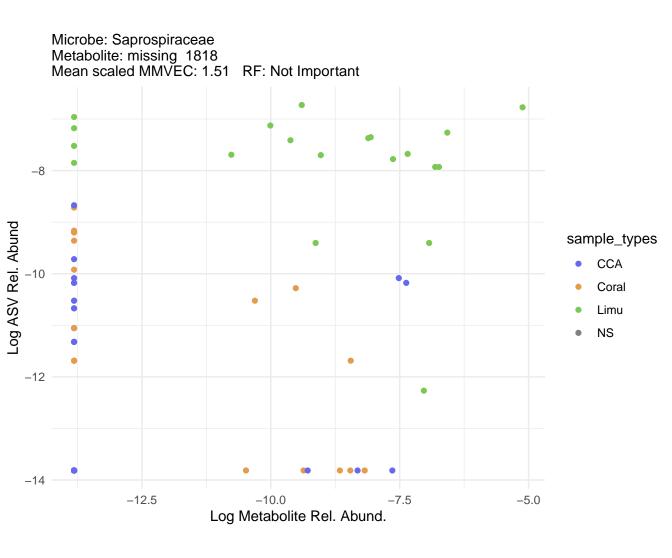


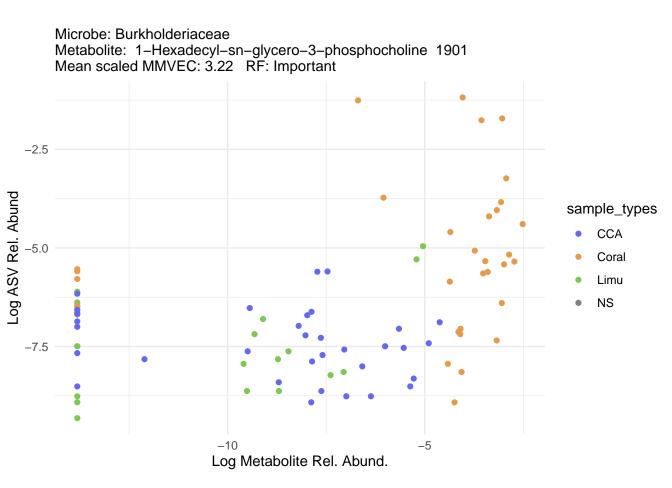


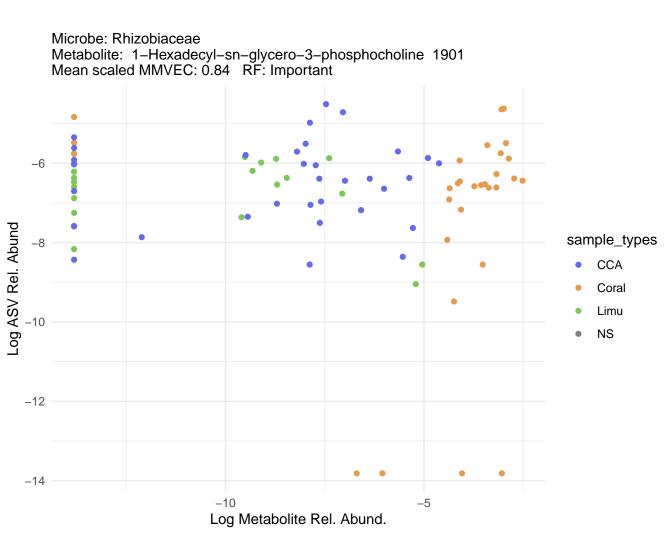


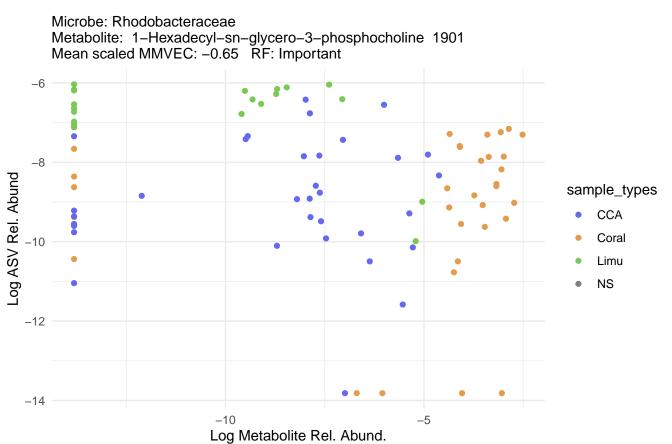


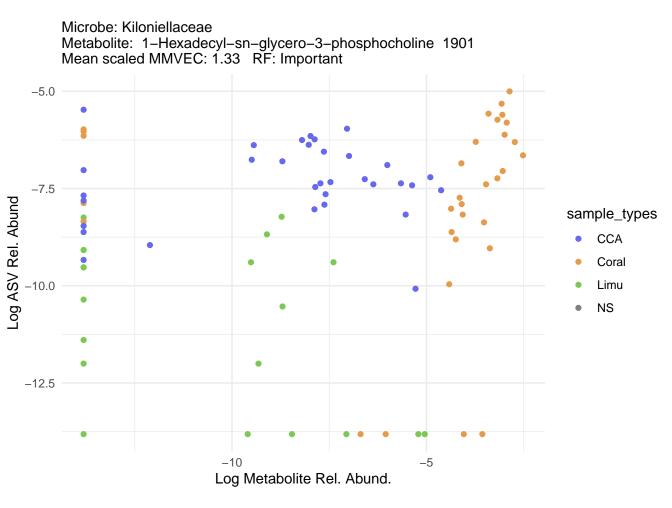


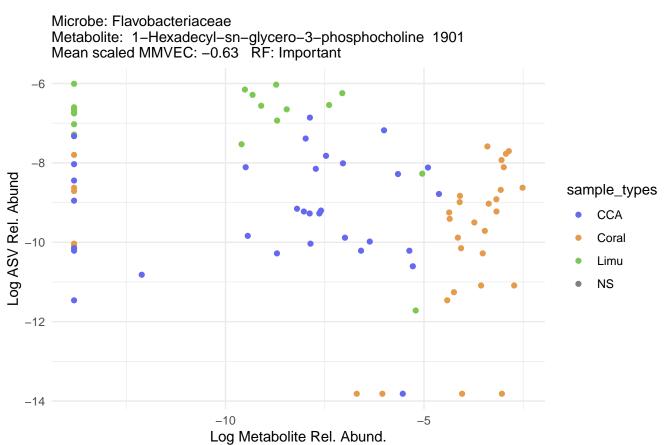


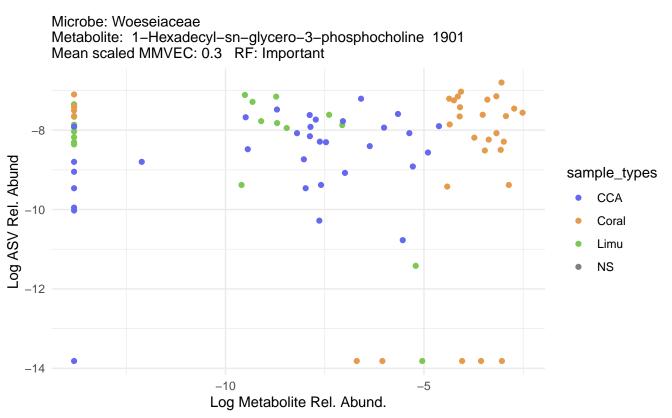


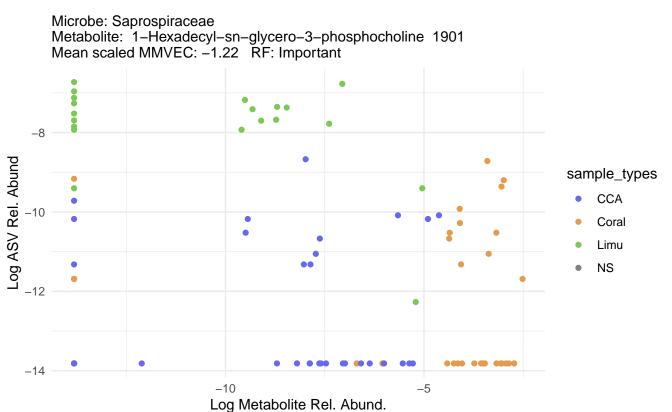


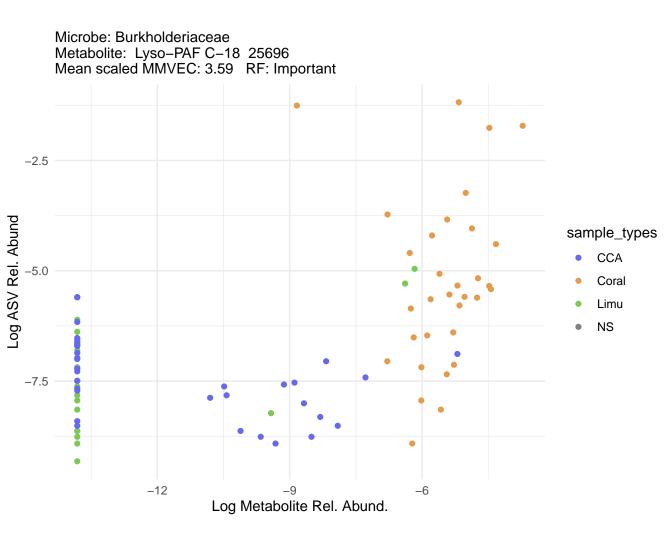


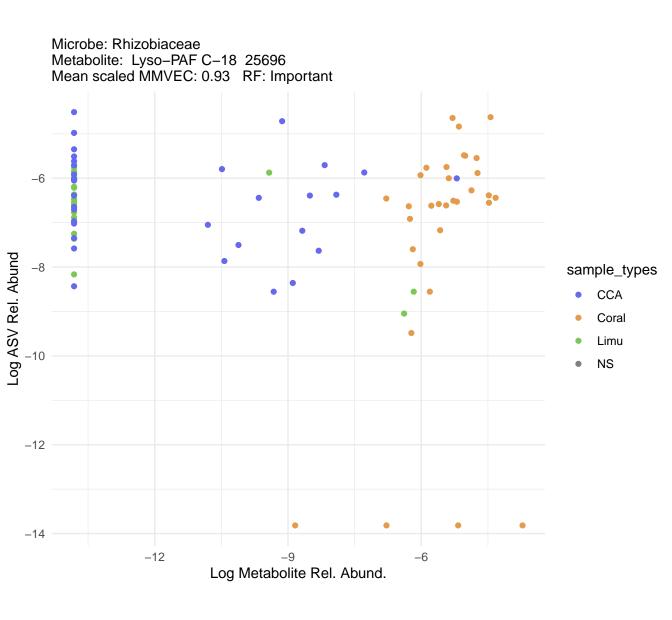


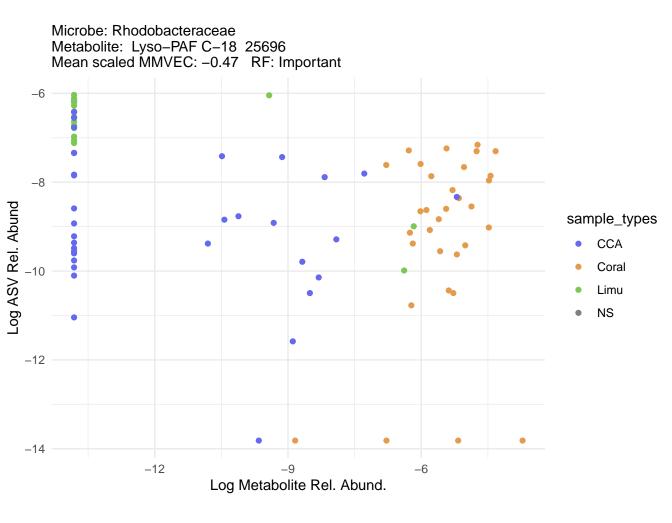


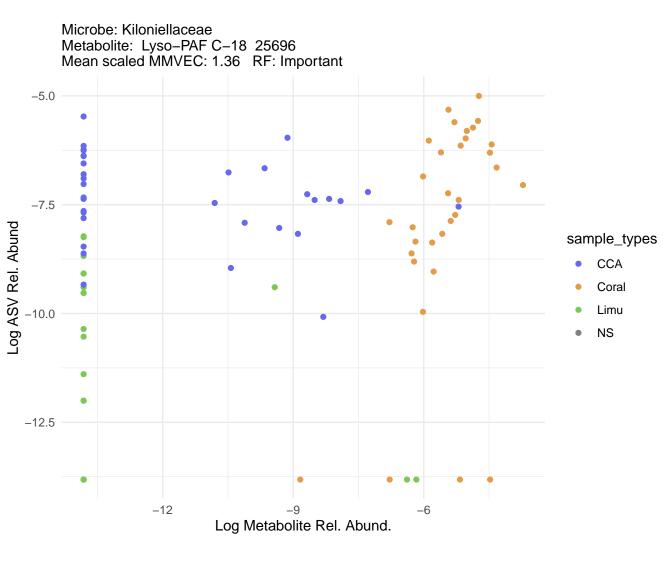


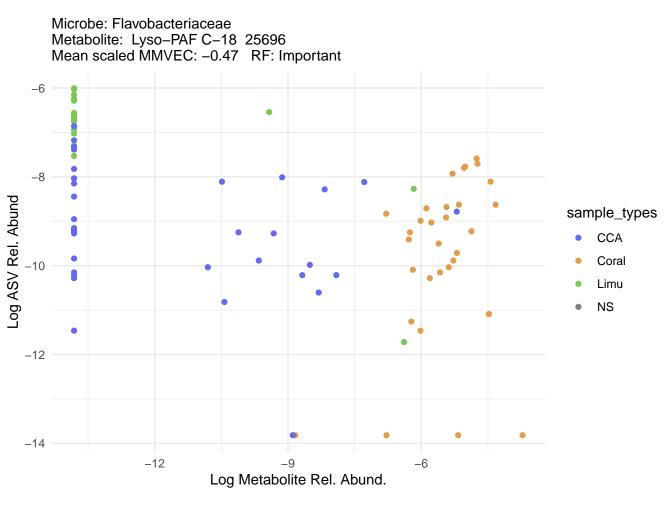


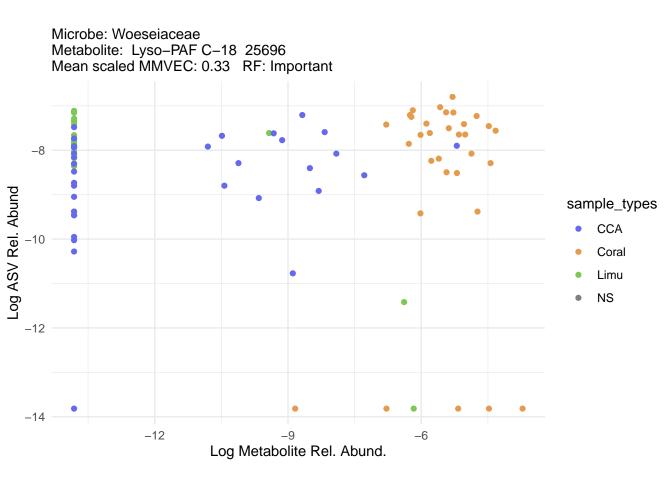


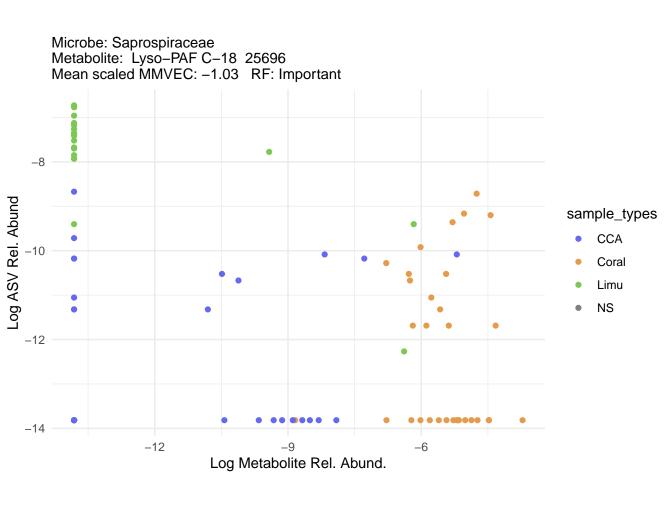


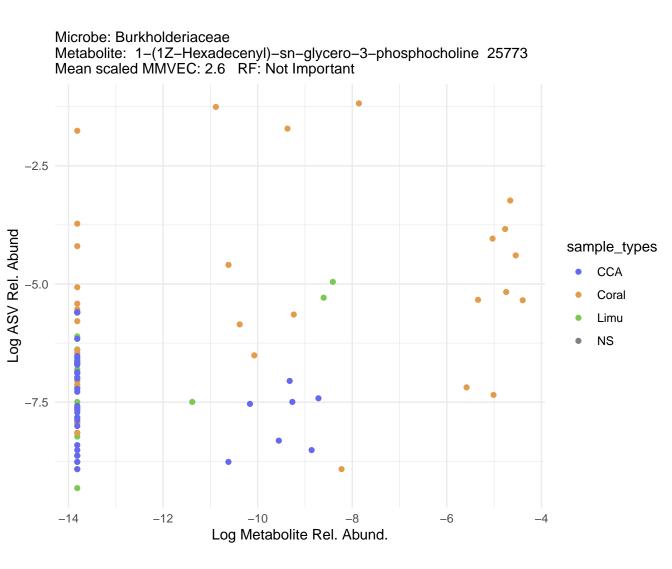


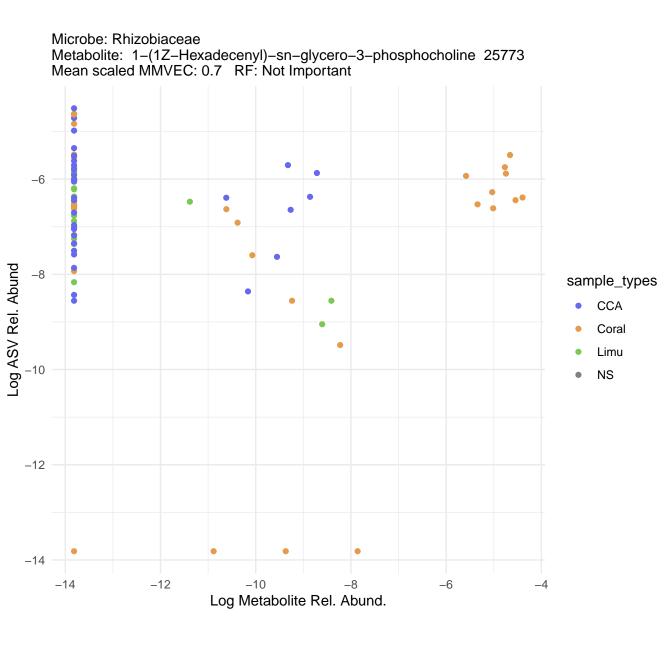


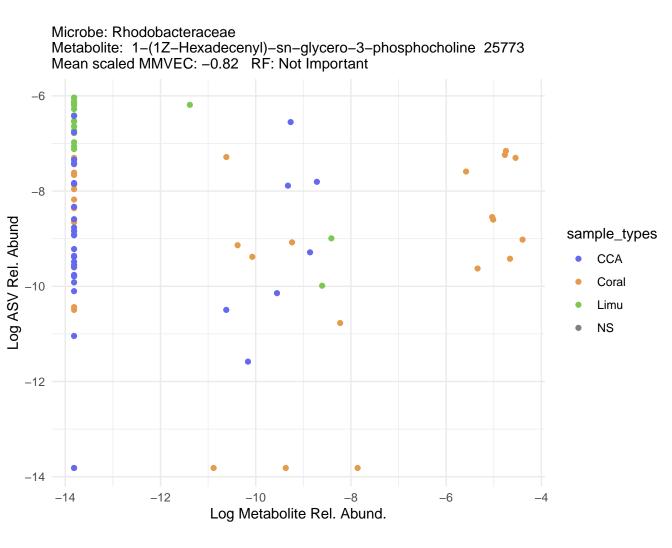


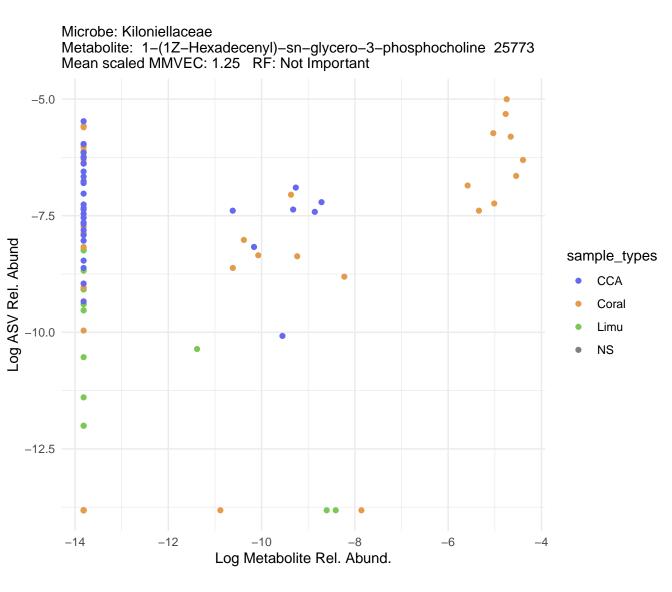






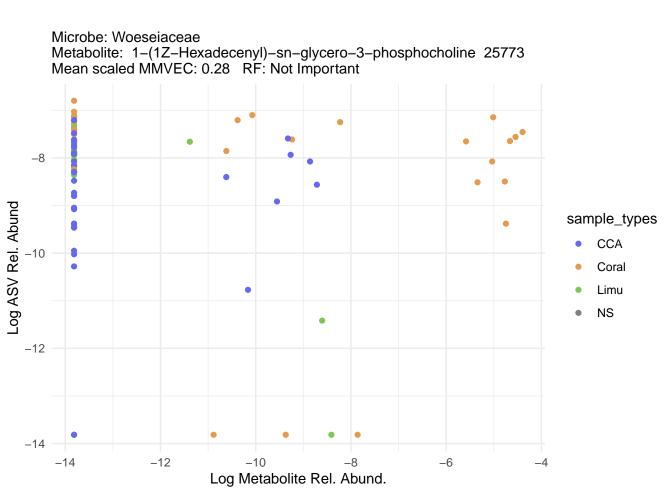


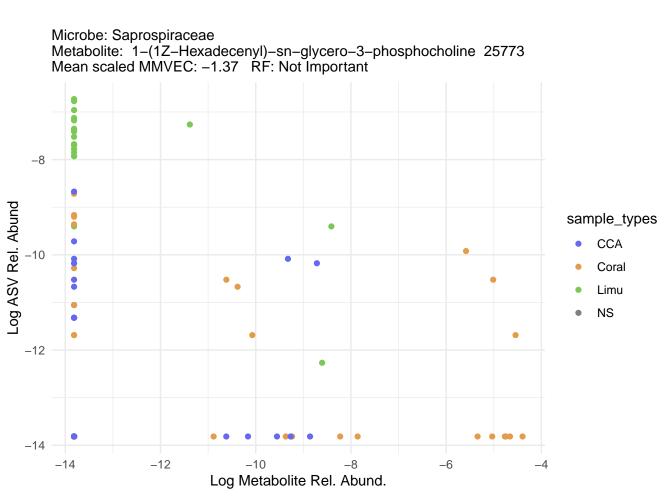


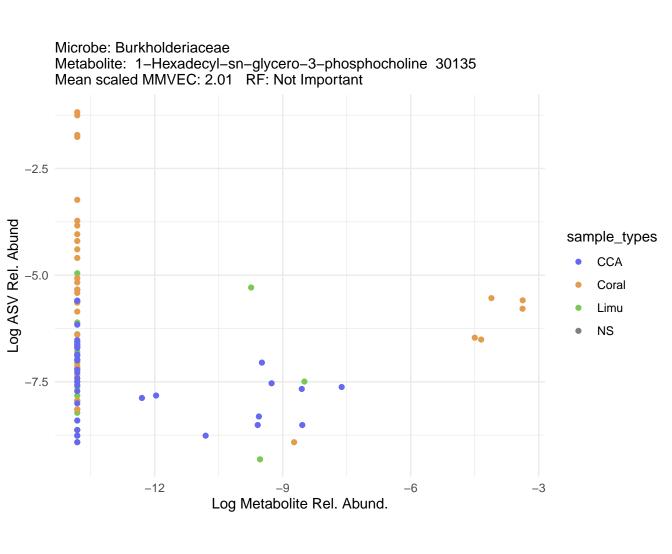


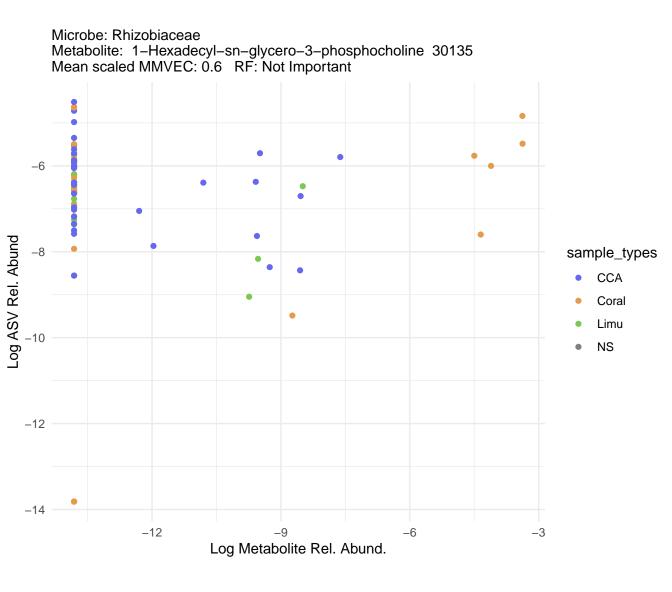
Microbe: Flavobacteriaceae Metabolite: 1–(1Z–Hexadecenyl)–sn–glycero–3–phosphocholine 25773 Mean scaled MMVEC: –0.79 RF: Not Important -6 -8 Log ASV Rel. Abund sample\_types CCA Coral Limu NS -12 -14 -14-12 -10 -6 -4

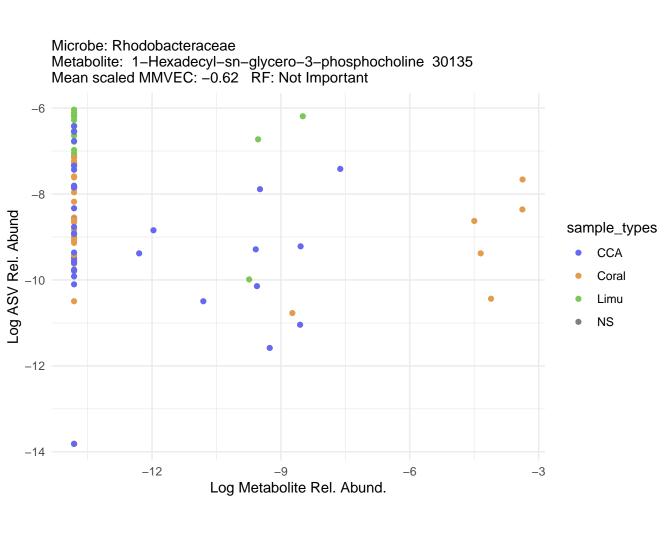
Log Metabolite Rel. Abund.

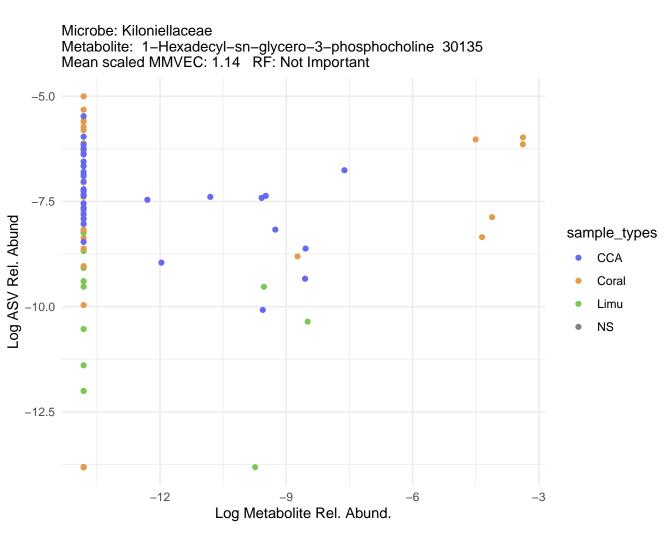


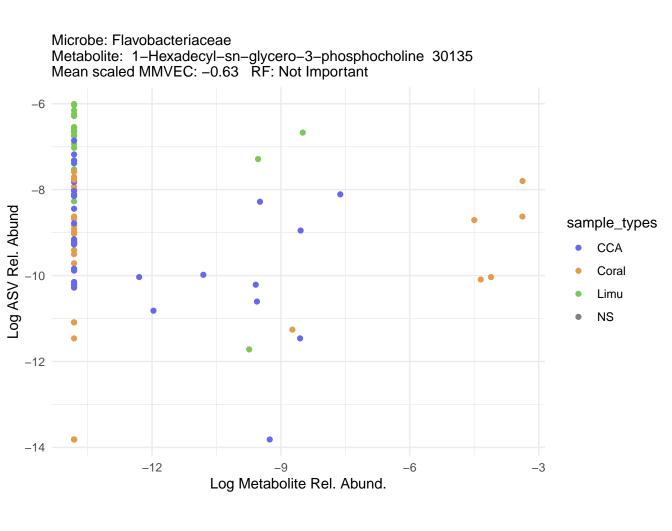


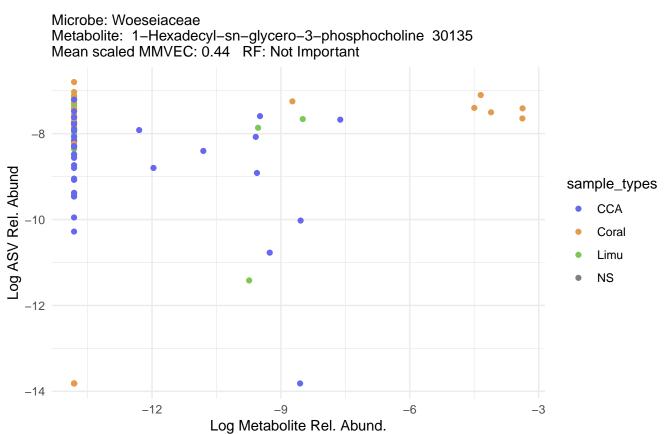


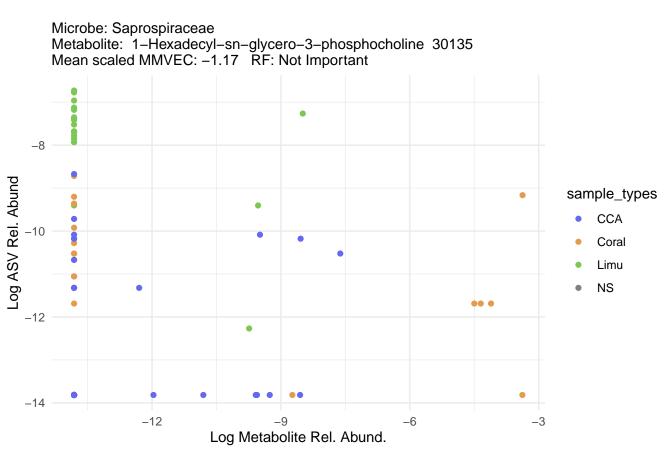


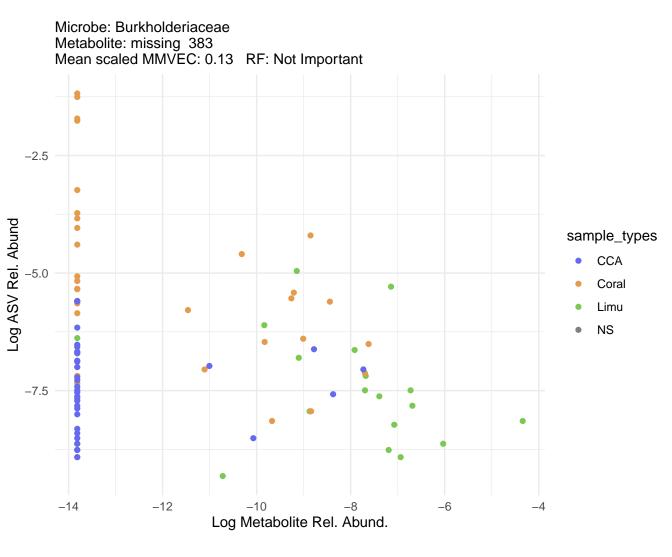


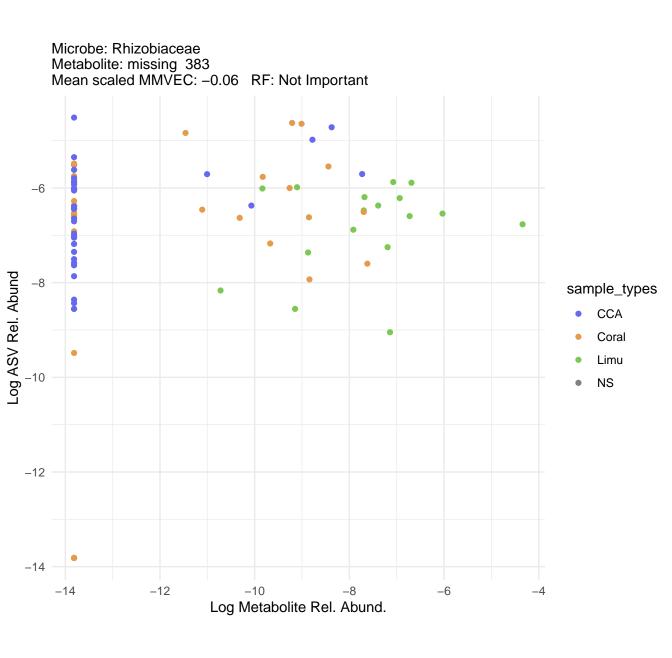




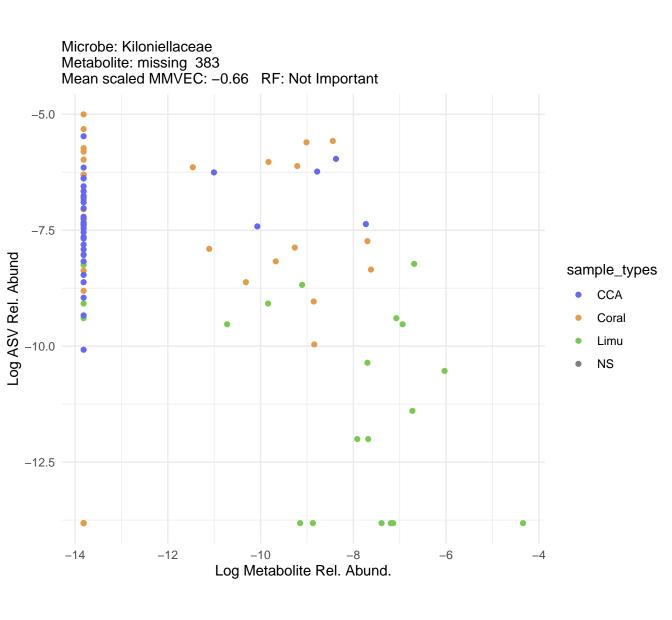


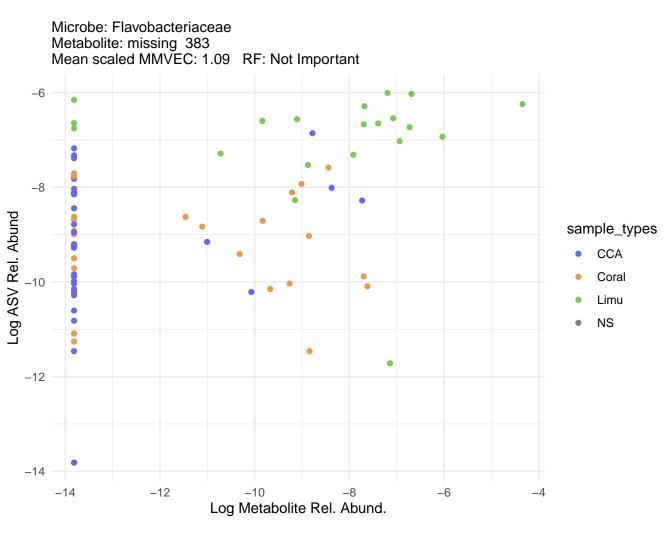


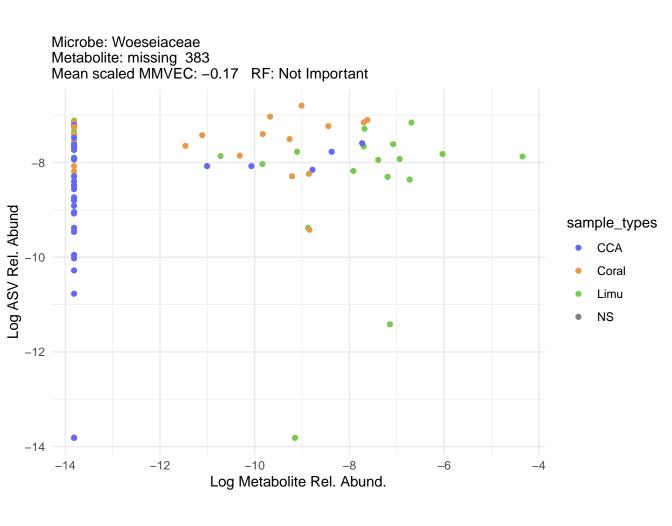


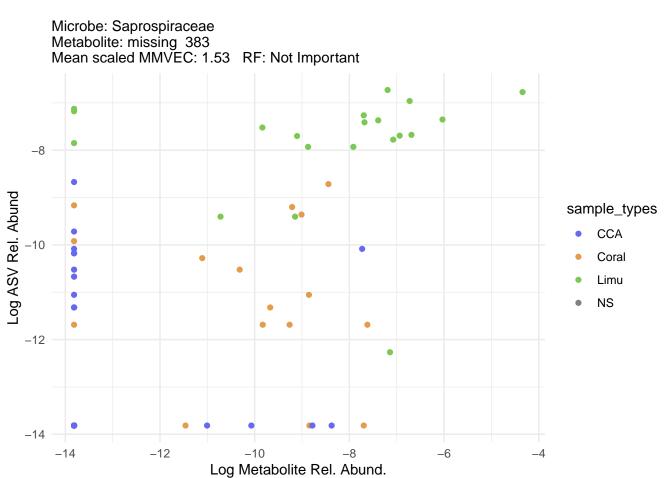


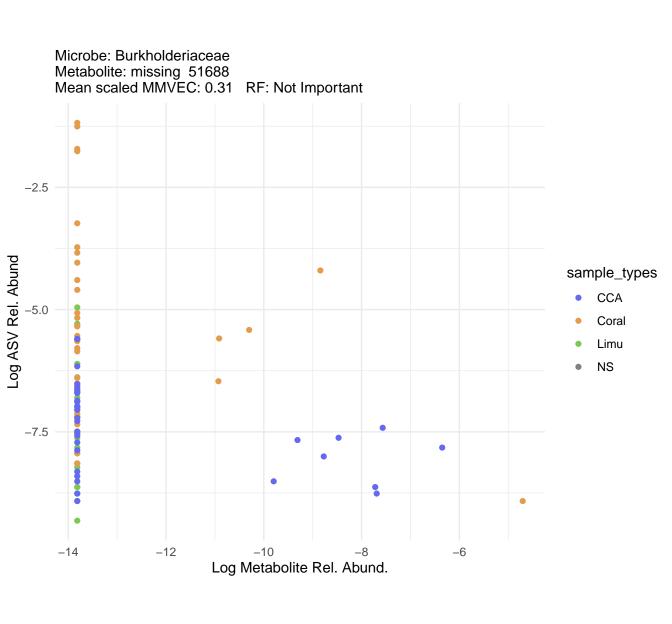
Microbe: Rhodobacteraceae Metabolite: missing 383
Mean scaled MMVEC: 1.15 RF: Not Important -6 -8 Log ASV Rel. Abund sample\_types CCA Coral -10 Limu • NS -12 -14 -14 -12 -10 -6 Log Metabolite Rel. Abund.

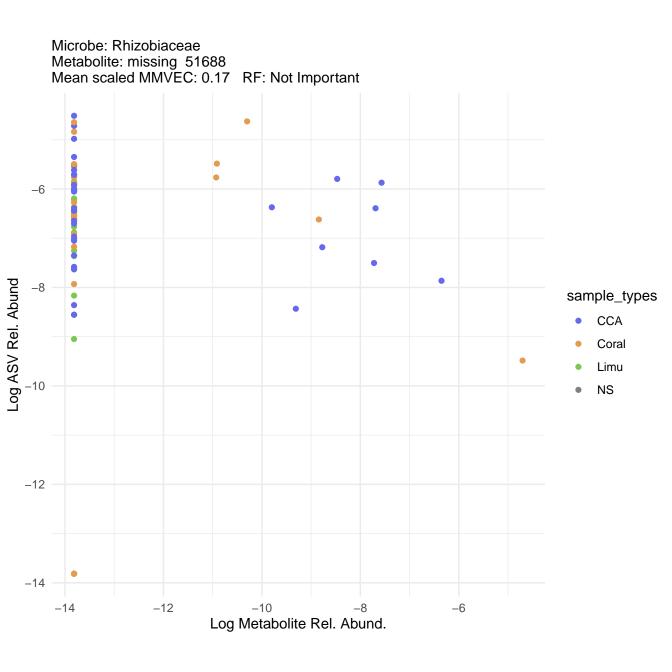


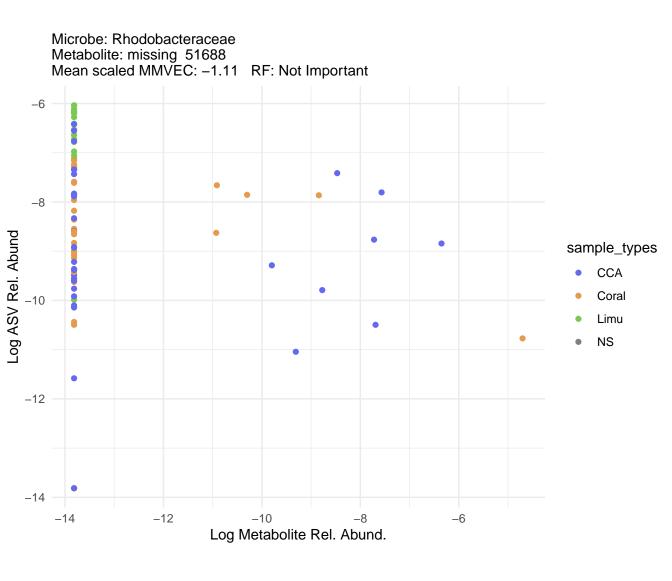


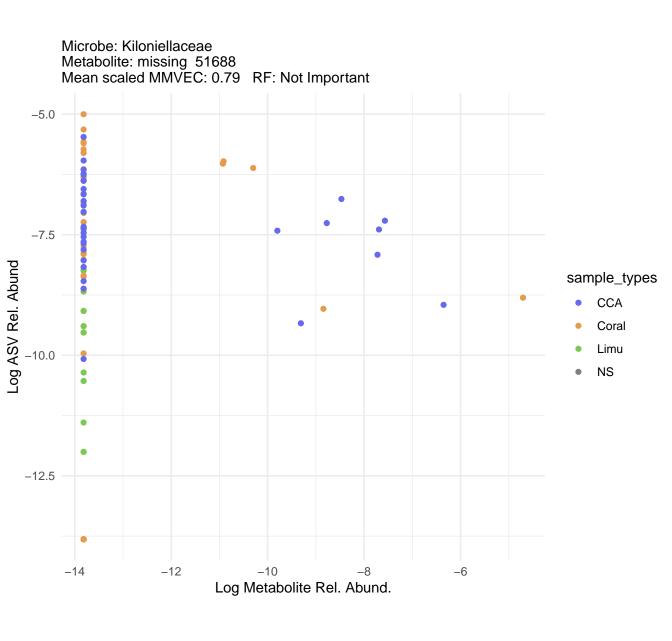


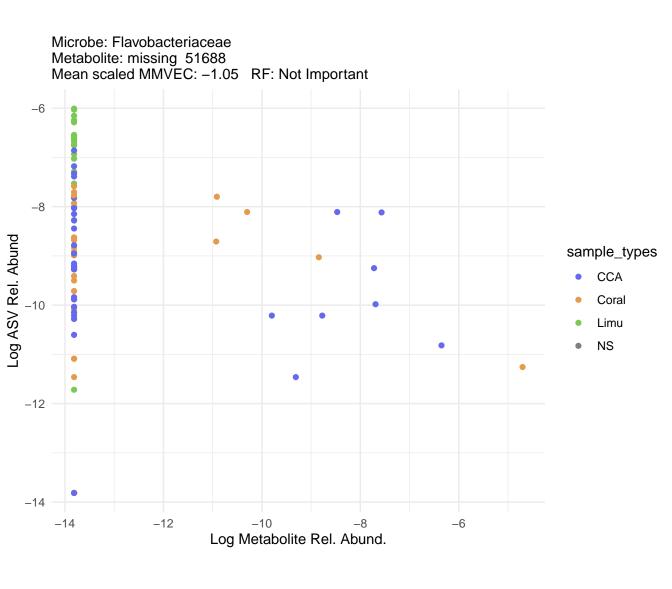


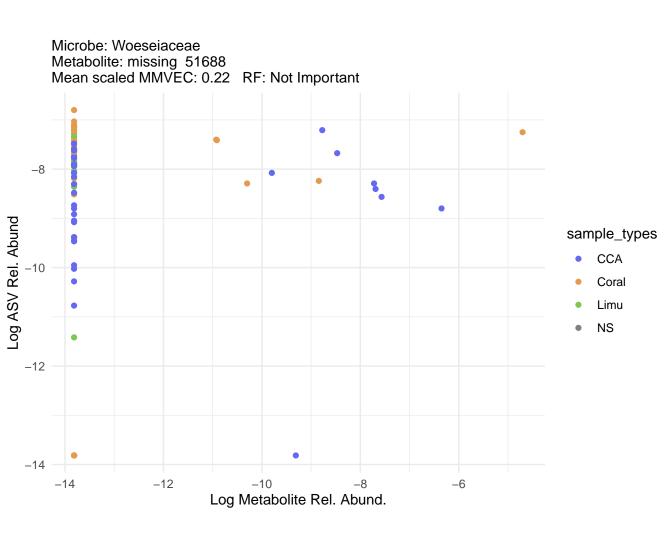


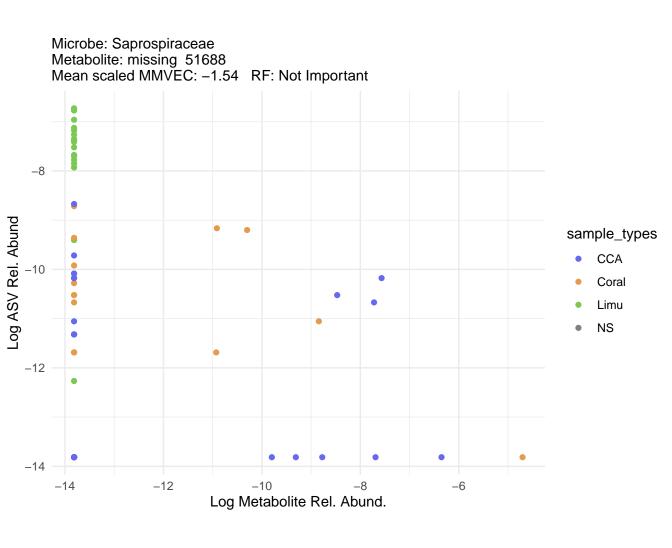


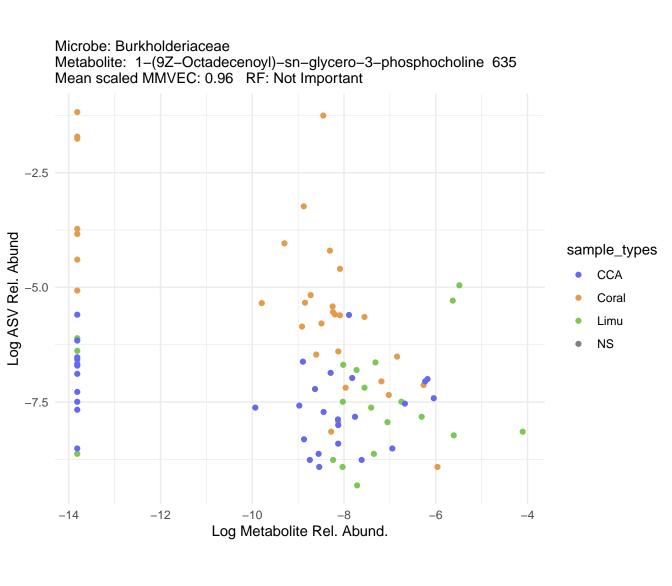


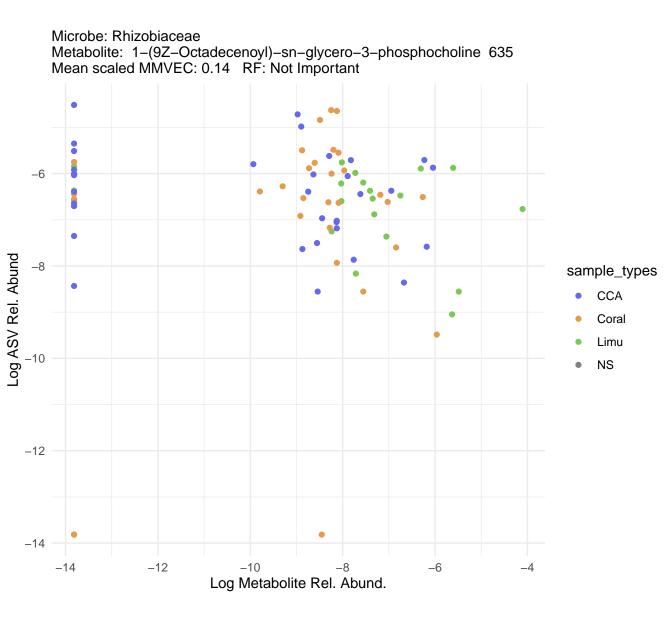


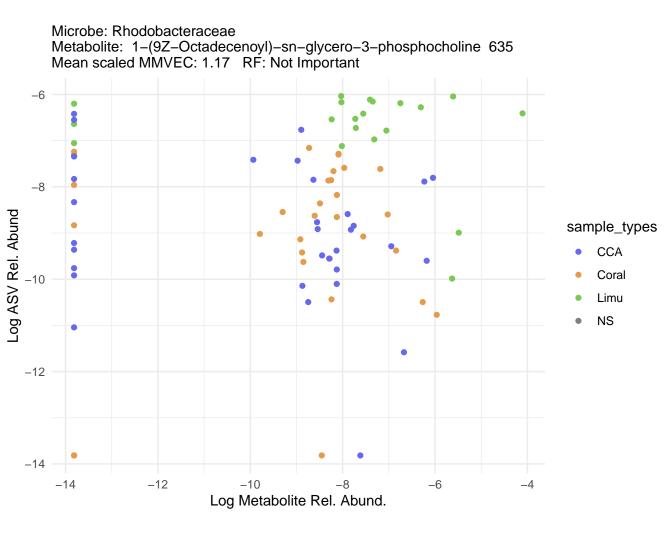


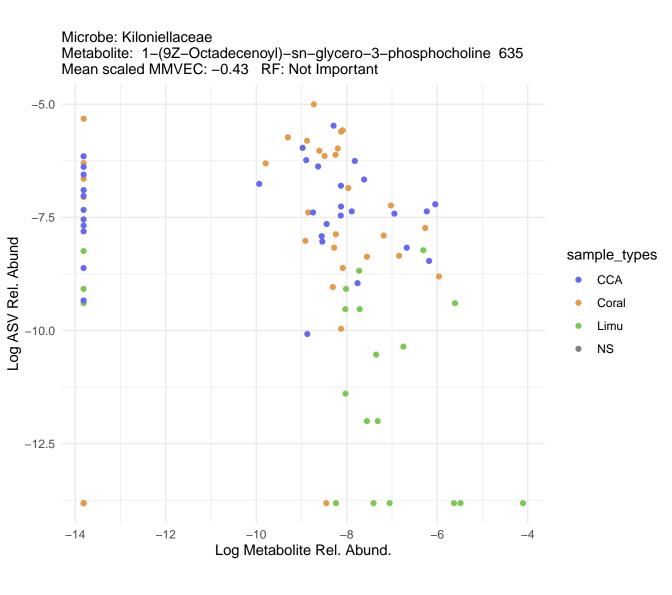


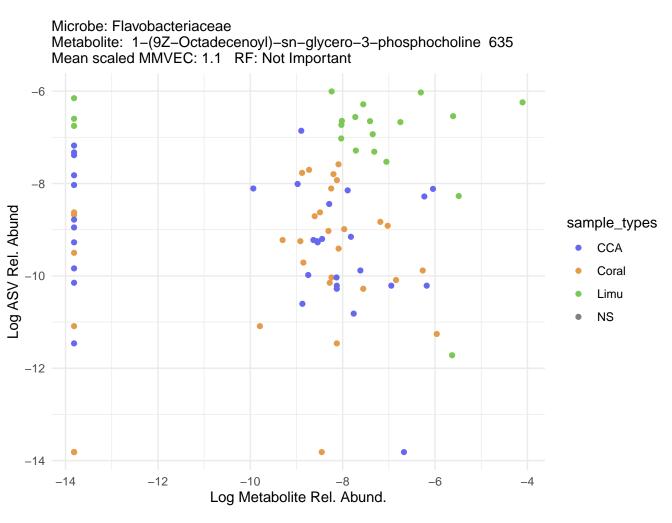


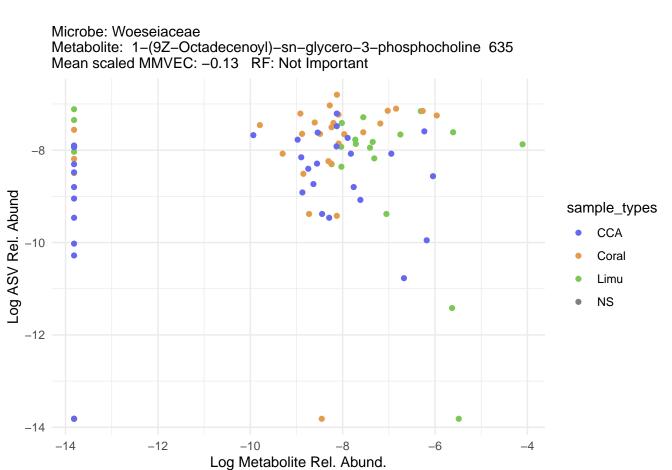






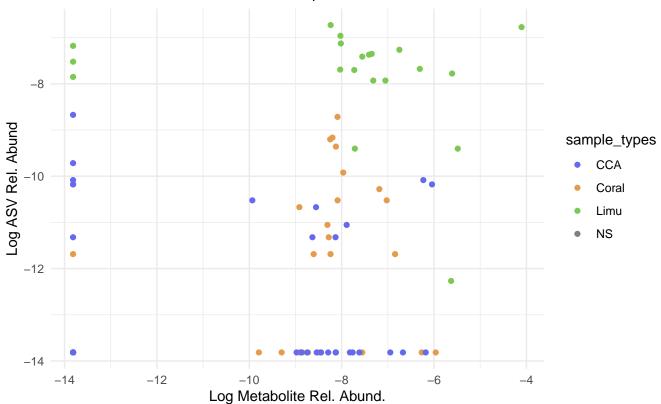


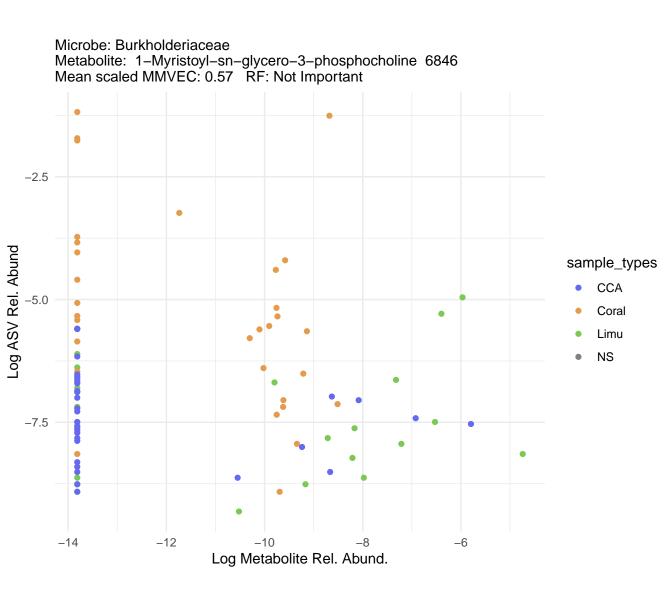


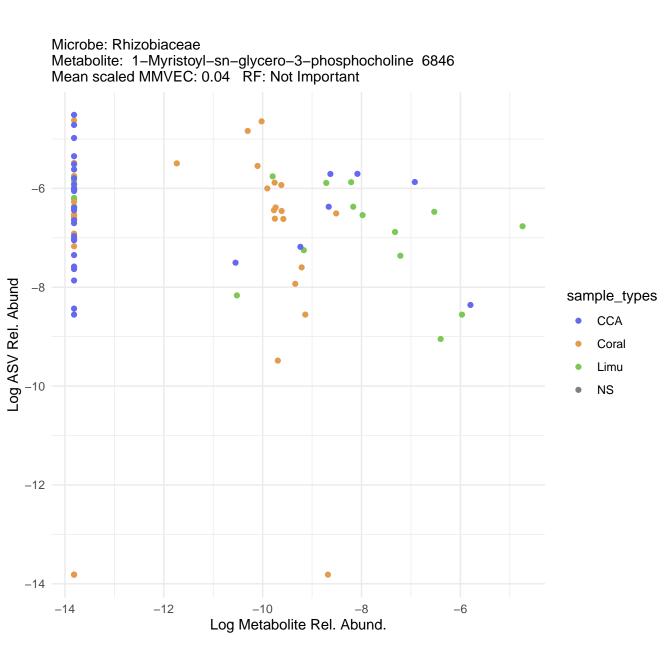


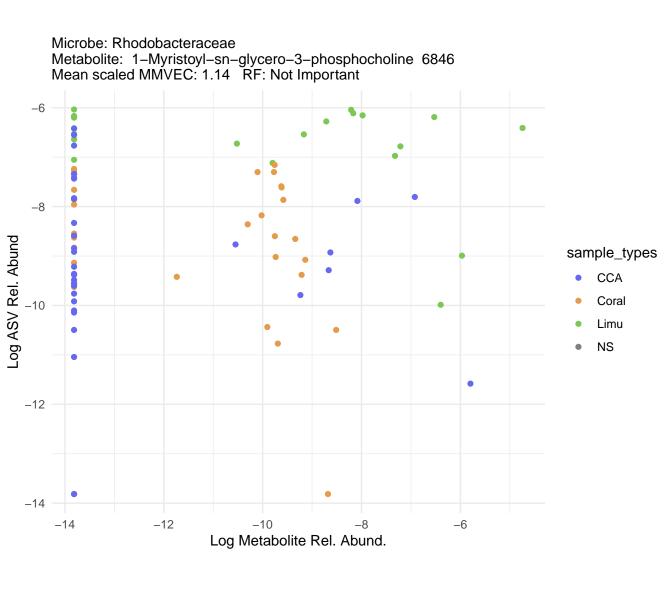
Microbe: Saprospiraceae

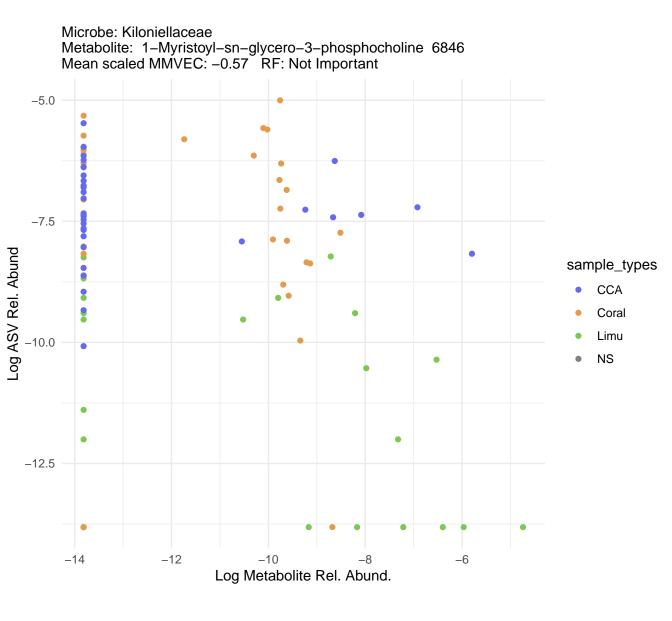
Metabolite: 1–(9Z–Octadecenoyl)–sn–glycero–3–phosphocholine 635 Mean scaled MMVEC: 1.47 RF: Not Important

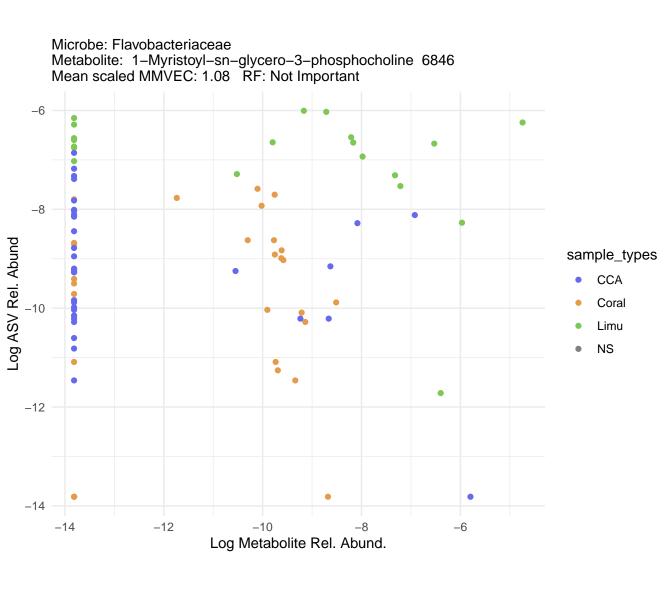


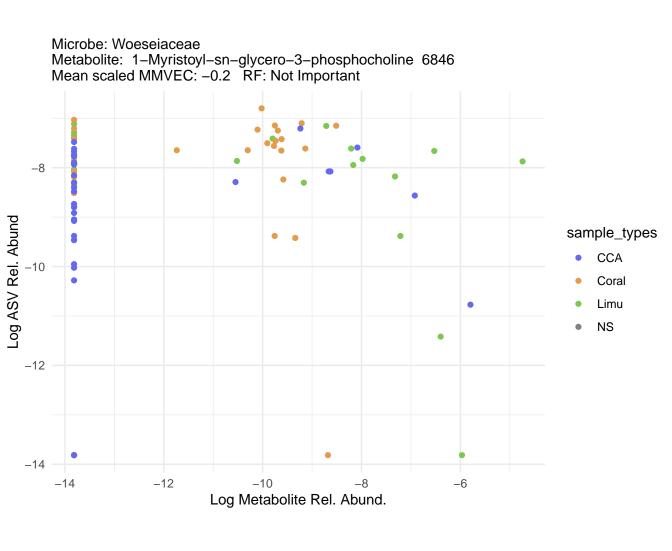












Microbe: Saprospiraceae
Metabolite: 1-Myristoyl-sn-glycero-3-phosphocholine 6846
Mean scaled MMVEC: 1.5 RF: Not Important

