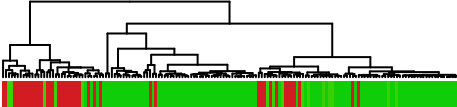




RiskGroup2

- Otu_78|Bacteroides
- Otu_332|Catenibacterium
- Otu_125|Parabacteroides
- Otu_82|Barnesiella
- Otu_119|Alistipes
- Otu_231|Eubacterium
- Otu_262|Dorea
- Otu_259|Coprococcus
- Otu_361|Phascolarctobacterium
- Otu_35|Collinsella
- Otu_340|Holdemanella
- Otu_391|Victivallis
- Otu_354|Firmicutes_unclassified
- Otu_45|Slackia
- Otu_407|Anaerobiospirillum
- Otu_308|Flavonifractor
- Otu_330|Ruthenibacterium
- Otu_365|Mitsuokella
- Otu_34|Olsenella
- Otu_371|Megasphaera





RiskGroup2



Otu_78|Bacteroides

Otu_332|Catenibacterium

Otu_125|Parabacteroides

Otu_82|Barnesiella

Otu_119|Alistipes

Otu_231|Eubacterium

Otu_262|Dorea

Otu_259|Coprococcus

Otu_361|Phascolarctobacterium

Otu_35|Collinsella

Otu_340|Holdemanella

Otu_391|Victivallis

Otu_354|Firmicutes_unclassified

Otu_45|Slackia

Otu_407|Anaerobiospirillum

Otu_308|Flavonifractor

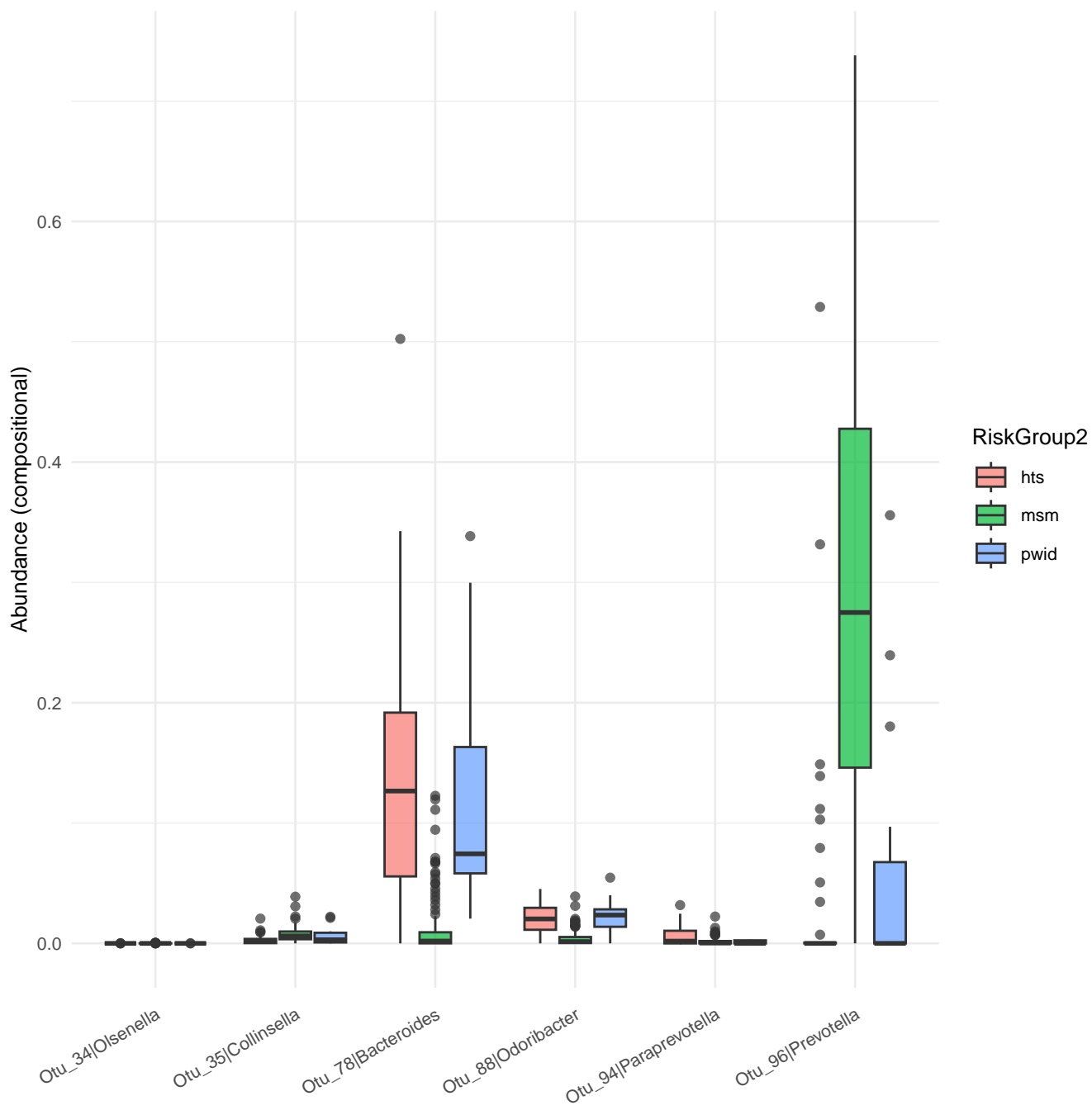
Otu_330|Ruthenibacterium

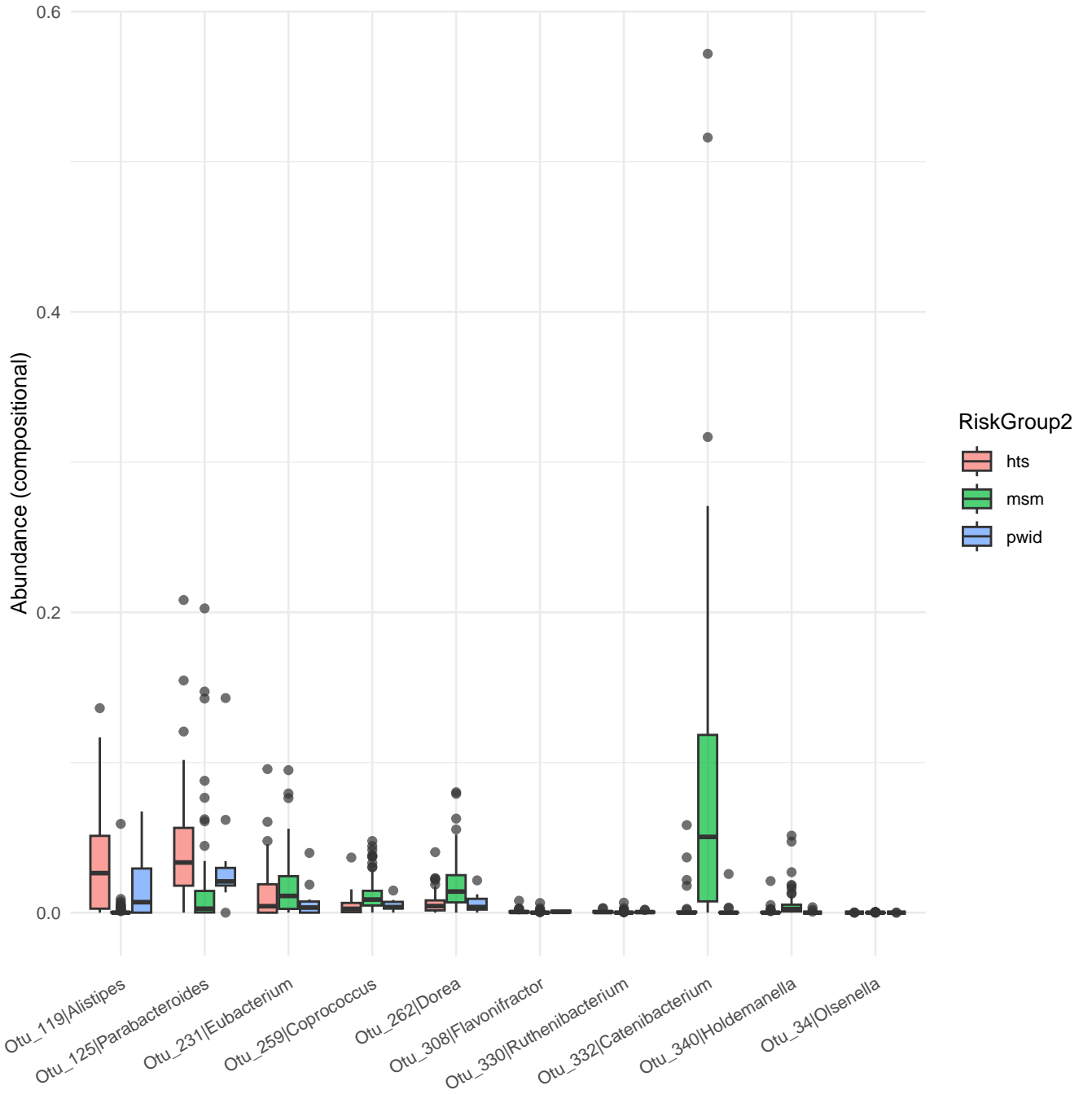
Otu_365|Mitsuokella

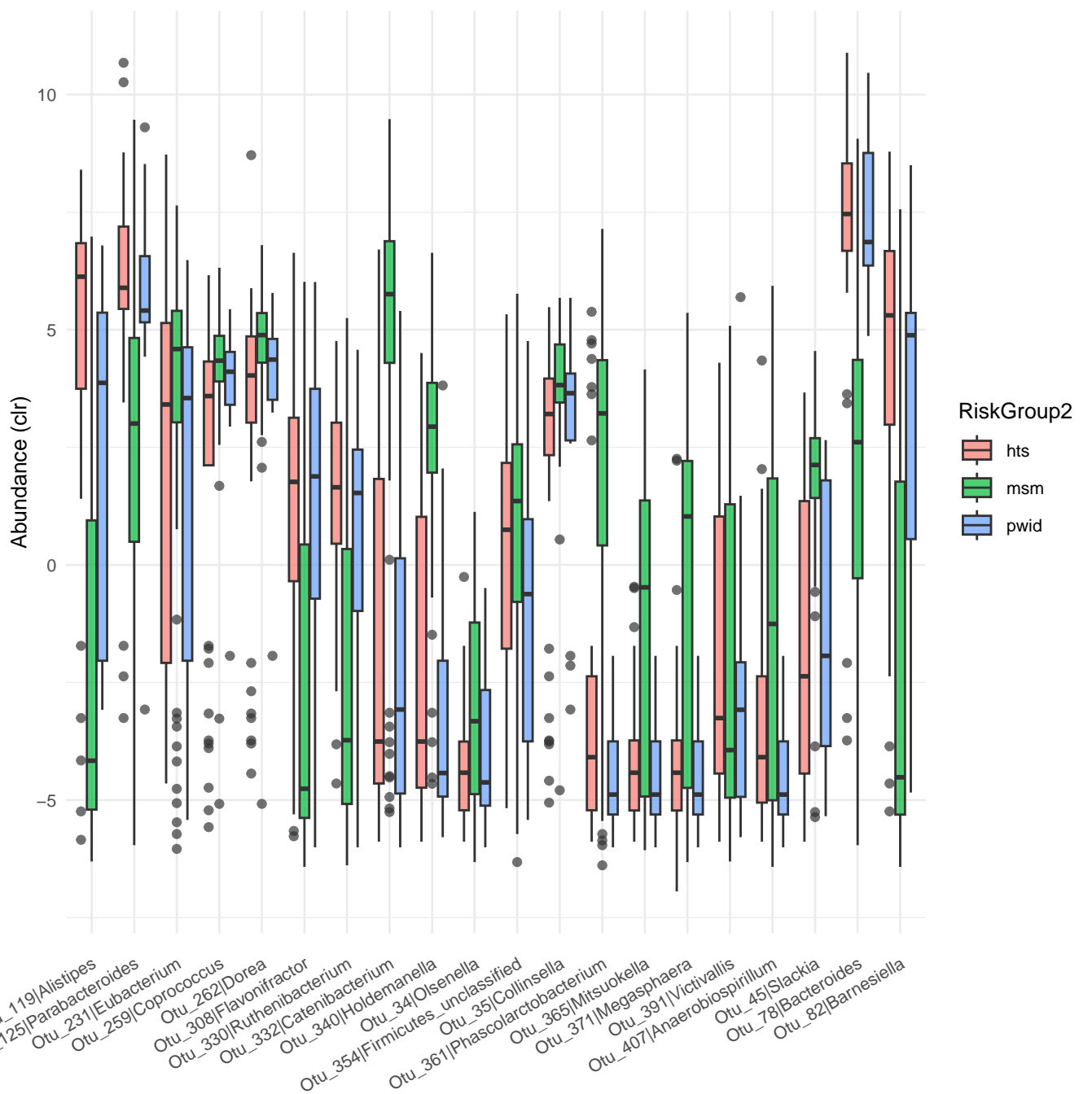
Otu_34|Olsenella

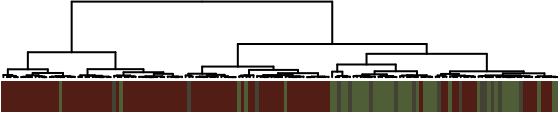
Otu_371|Megasphaera











RiskGroup2



Otu_96|Prevotella

Otu_78|Bacteroides

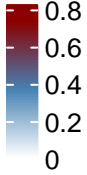
Otu_88|Odoribacter

Otu_35|Collinsella

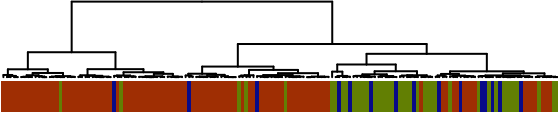
Otu_94|Paraprevotella

Otu_34|Olsenella

Abundance (compositional) RiskGroup2



hts
msm
pwid



RiskGroup2



Otu_96|Prevotella

Otu_78|Bacteroides

Otu_88|Odoribacter

Otu_35|Collinsella

Otu_94|Paraprevotella

Otu_34|Olsenella

Abundance (compositional) RiskGroup2

