

**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





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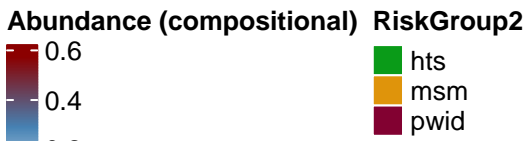
Otu\_308|Flavonifractor

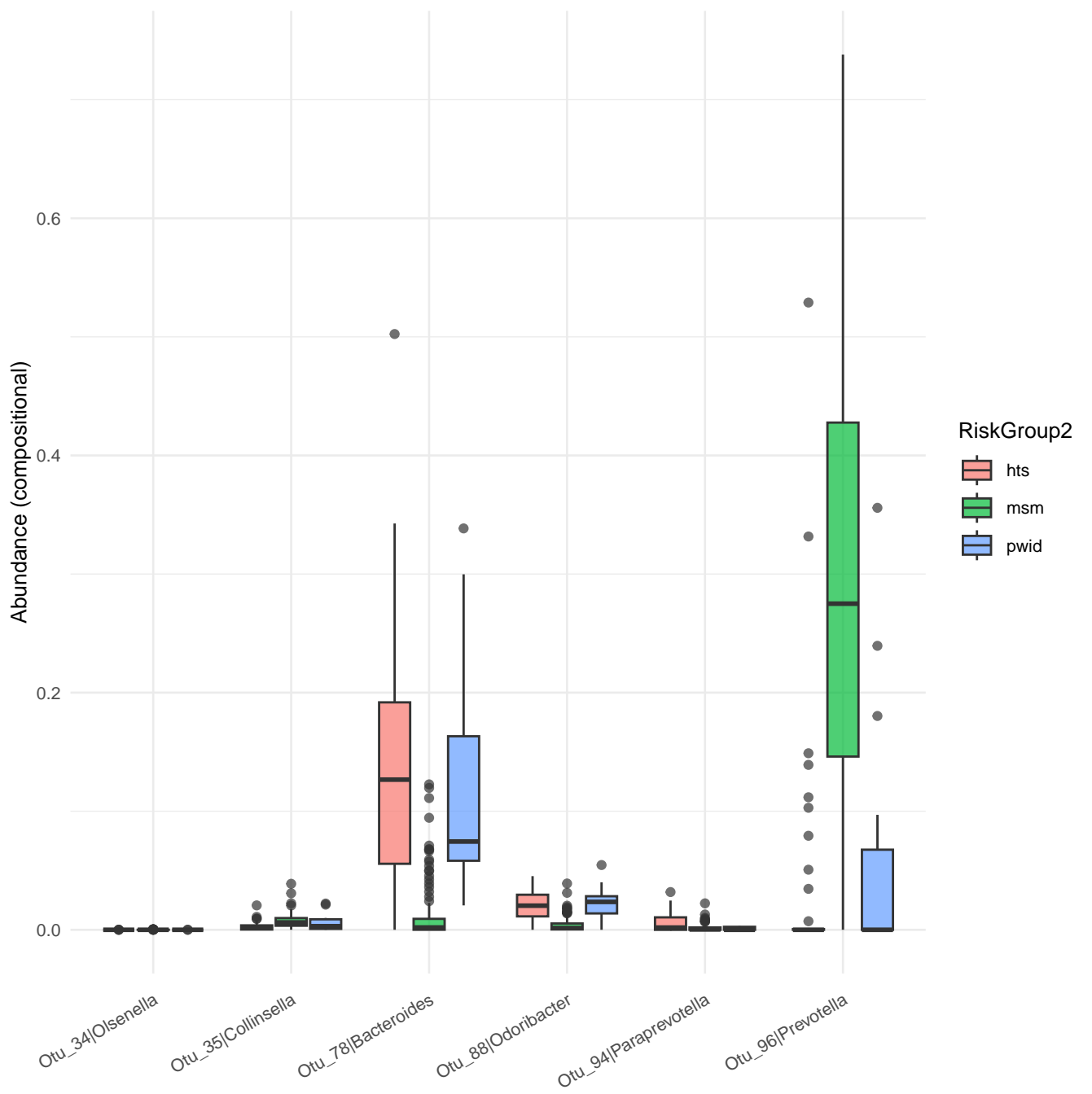
Otu\_330|Ruthenibacterium

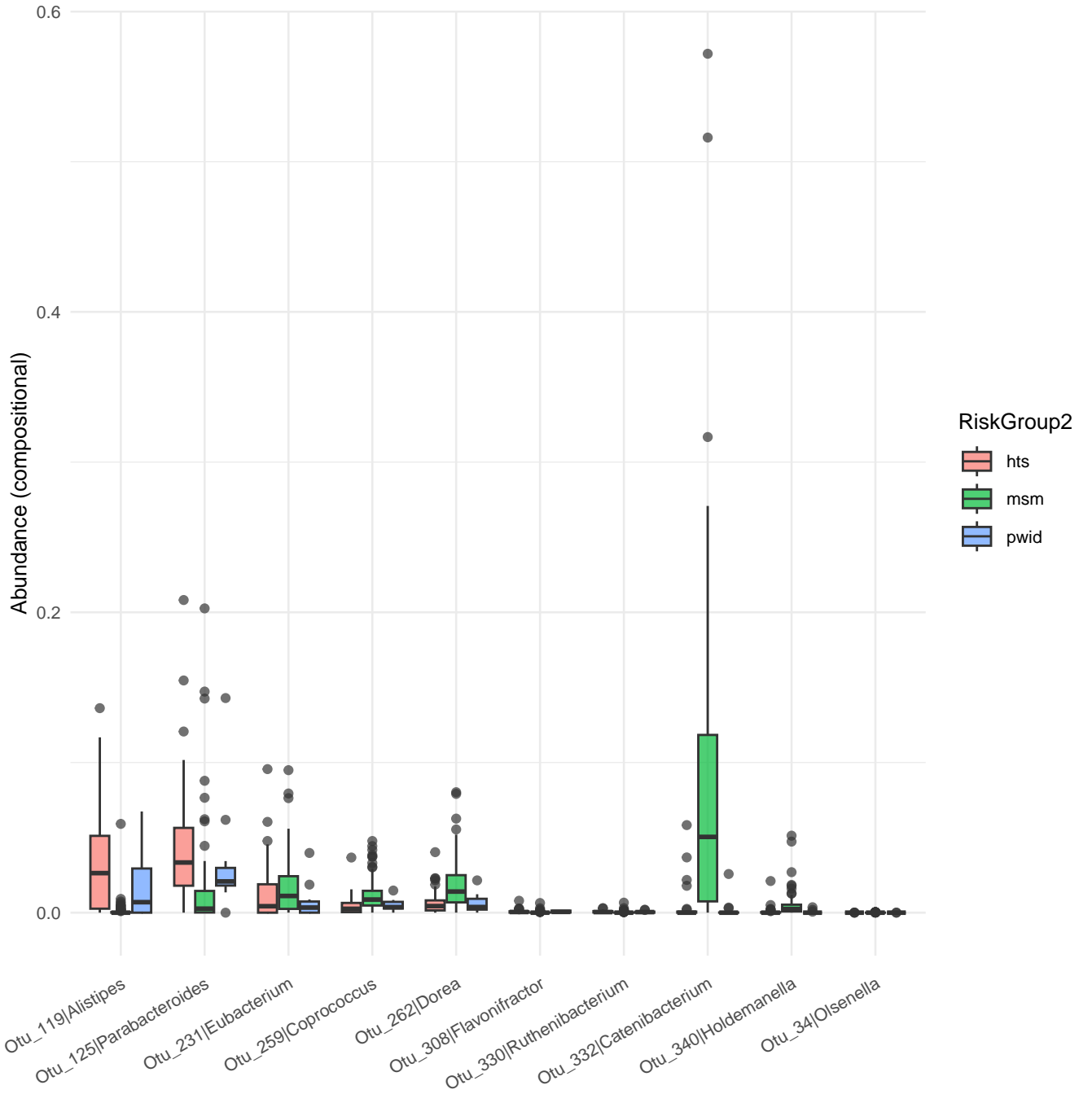
Otu\_365|Mitsuokella

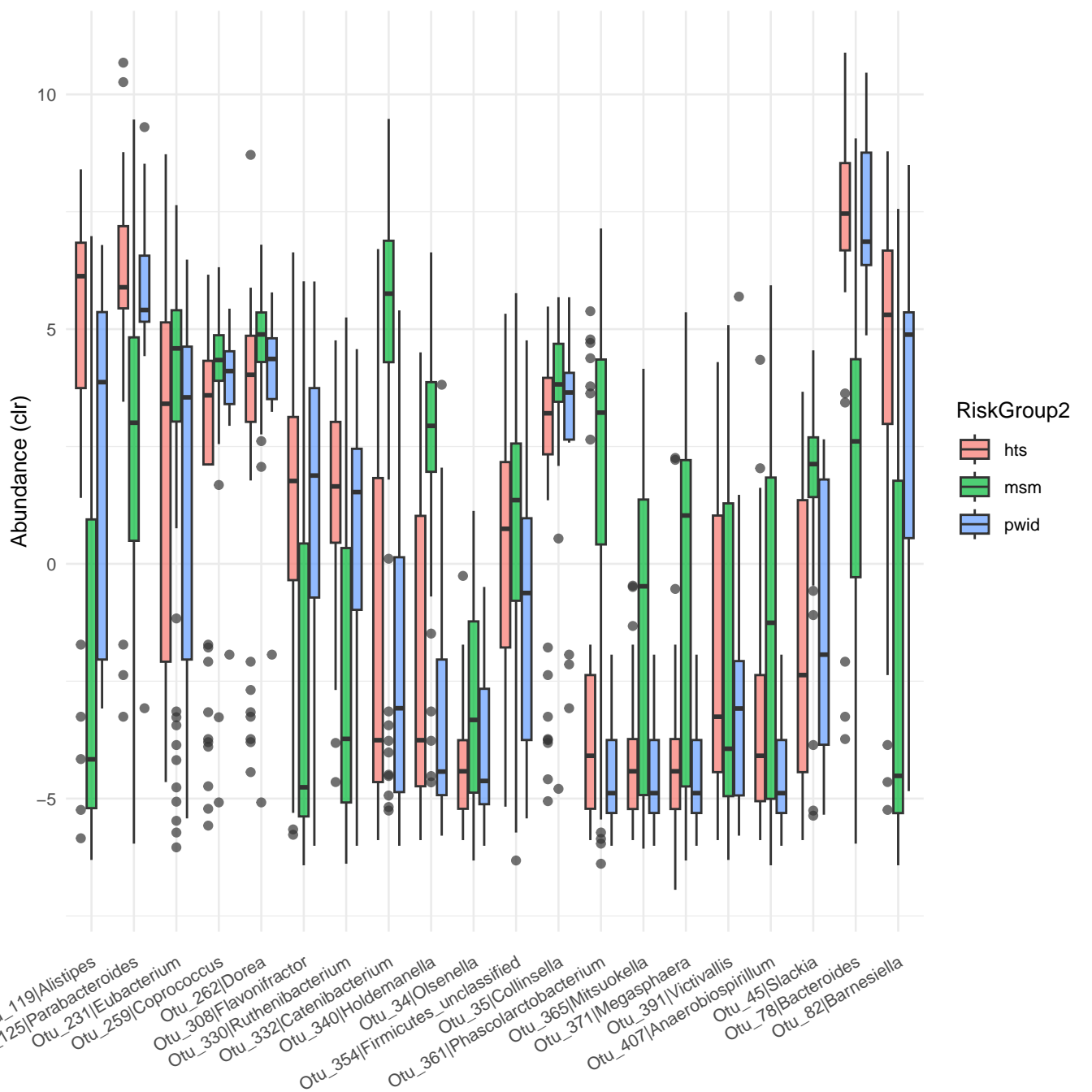
Otu\_34|Olsenella

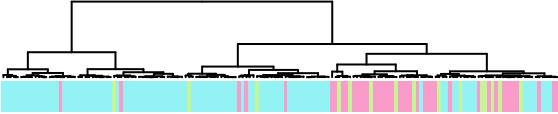
Otu\_371|Megasphaera











**RiskGroup2**



Otu\_96|Prevotella

Otu\_78|Bacteroides

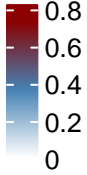
Otu\_88|Odoribacter

Otu\_35|Collinsella

Otu\_94|Paraprevotella

Otu\_34|Olsenella

**Abundance (compositional) RiskGroup2**

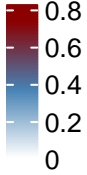




**RiskGroup2**



**Abundance (compositional) RiskGroup2**



hts  
msm  
pwid

