

Kruskal–Wallis, $p = 2.4\text{e-}09$

Relative abundance of predicted phenotypes

compartment

- Mucus
- Tissue
- Skeleton

2

1

0

-1

M_above

M_neutral

M_below

T_above

T_neutral

T_below

S_above

S_neutral

S_below

Compartments and respective model fit

