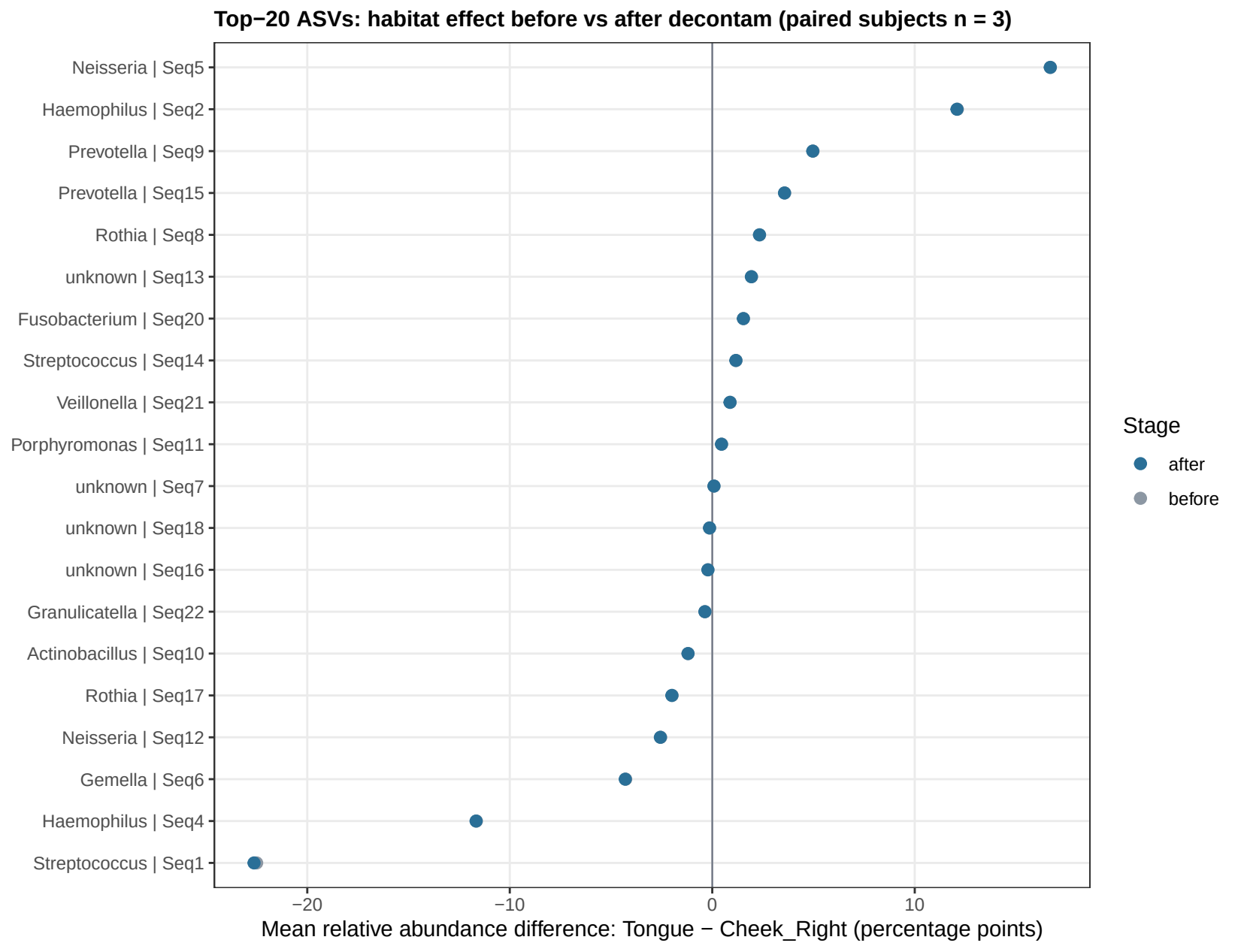
Top−20 ASVs: habitat effect before vs after decontam (paired subjects n = 3)
Neisseria | Seq5
Haemophilus | Seq2
Prevotella | Seq9
Prevotella | Seq15
Rothia | Seq8
unknown | Seq13
Fusobacterium | Seq20
Streptococcus | Seq14
Veillonella | Seq21
Porphyromonas | Seq11
unknown | Seq7
unknown | Seq18
unknown | Seq16
Granulicatella | Seq22
Actinobacillus | Seq10
Rothia | Seq17
Neisseria | Seq12
Gemella | Seq6
Haemophilus | Seq4
Streptococcus | Seq1
−20
−10
0
10
Mean relative abundance difference: Tongue − Cheek_Right (percentage points)
Stage
after
before