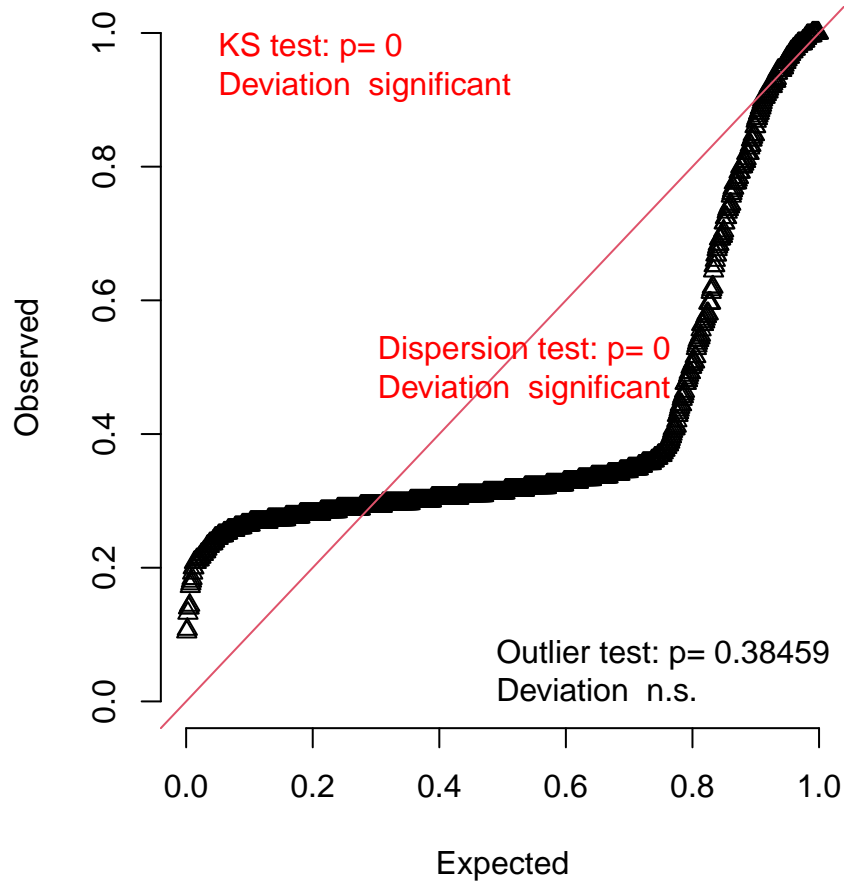DHARMa residual

Citrobacter_freundii (SGB10083_group)Beta model

QQ plot residuals

DHARMa residual vs. predicted
Quantile deviations detected (red curves)
Combined adjusted quantile test significant

KS test: p= 0
Deviation significant

Dispersion test: p= 0
Deviation significant

Outlier test: p= 0.38459
Deviation n.s.

Observed

Expected

DHARMa residual

Model predictions (rank transformed)