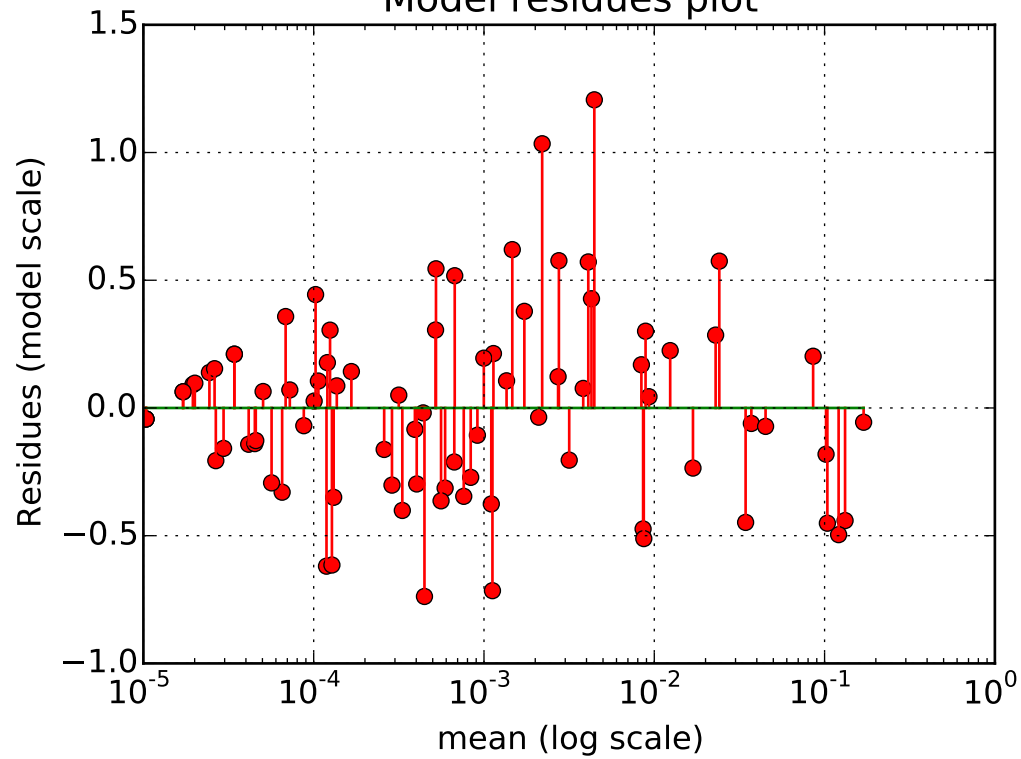
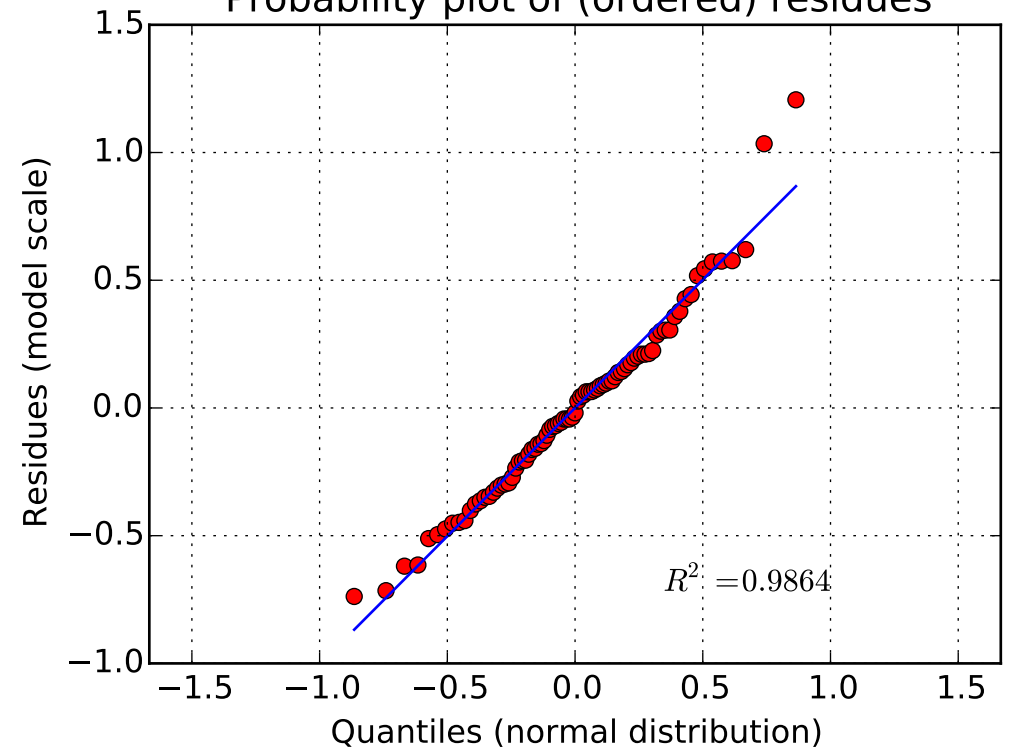


IBS_P1_metatranscriptomes_family: residues analysis of LLR-model fit

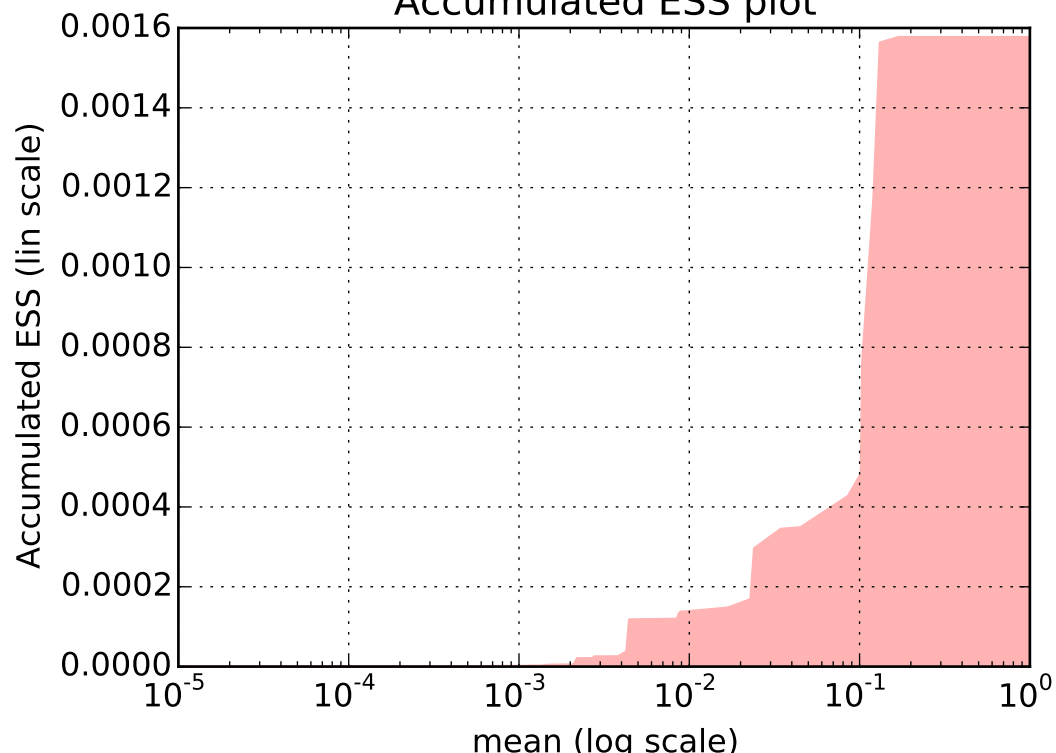
Model residues plot



Probability plot of (ordered) residues



Accumulated ESS plot



Normality plot of the residues

