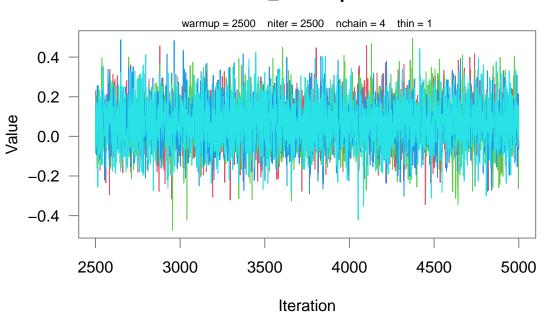
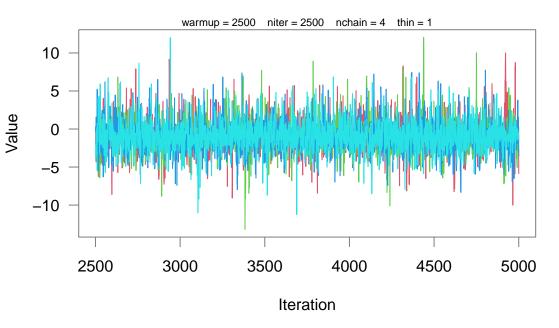
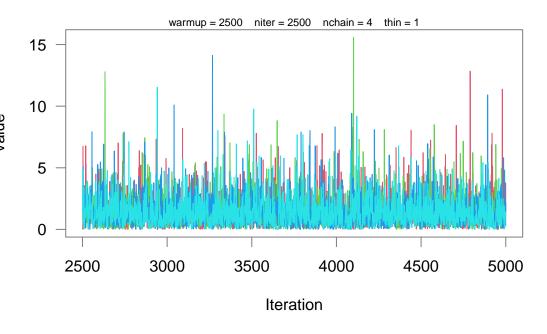
b_Intercept



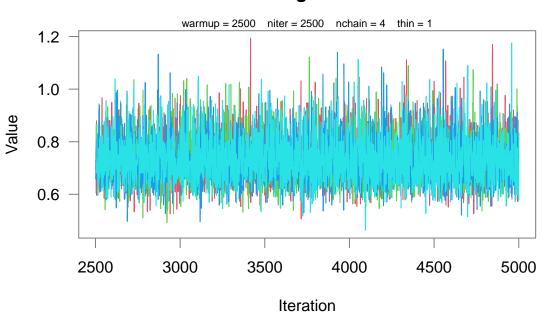
bs_smean.anom_1



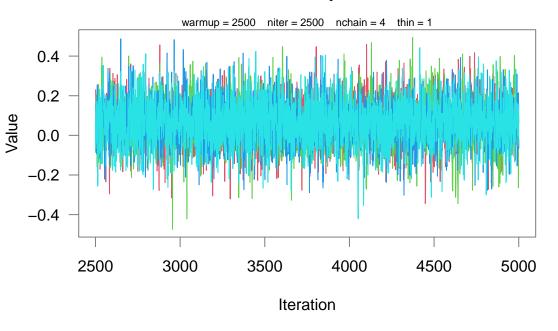
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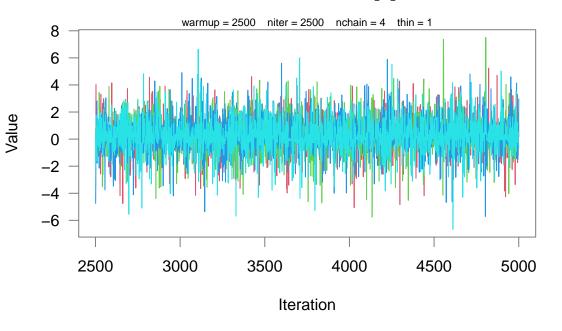
sigma



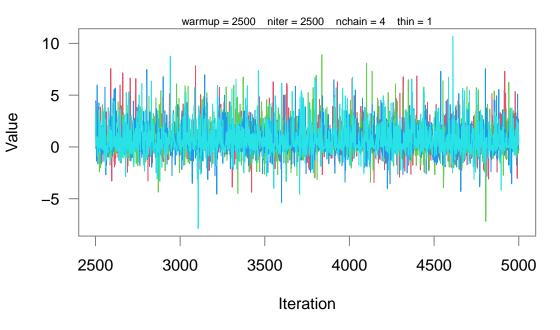
Intercept



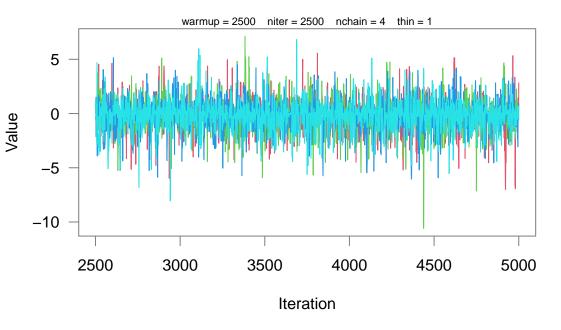
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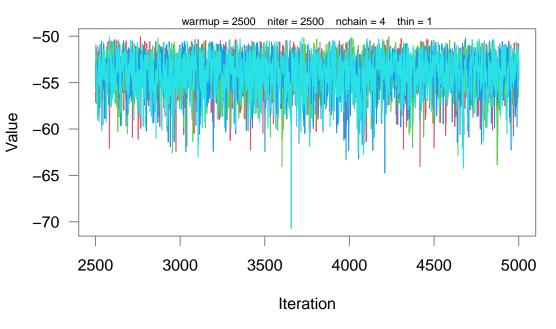
s_smean.anom_1[2]



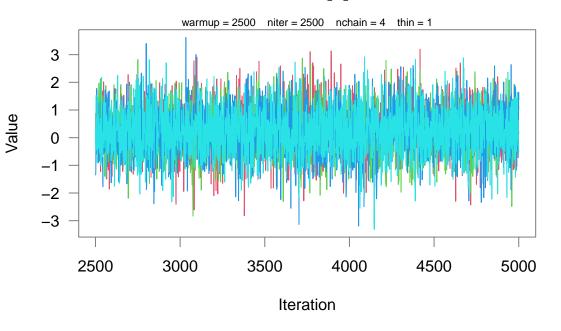
s_smean.anom_1[3]



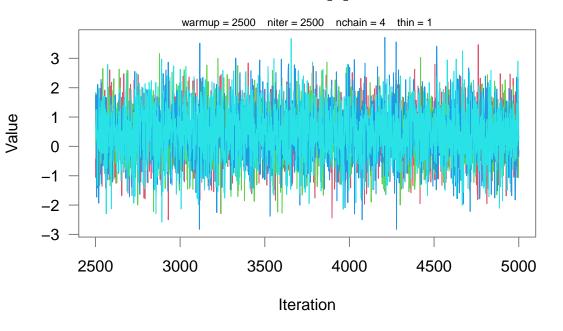




zs_1_1[1]



zs_1_1[2]



zs_1_1[3]

