

Model Type:simult_XSL_gmo

Fit results:

GMO rln = 0.057371 +- nan

M_4 posterior = -0.0511297 +- nan

Centroid posterior = 0.714612 +- nan

lam_E0= 0.743111 +- nan

1st e.s: 0.406889 +- nan

proton_E0= 0.604289 +- nan

1st e.s: 0.221707 +- nan

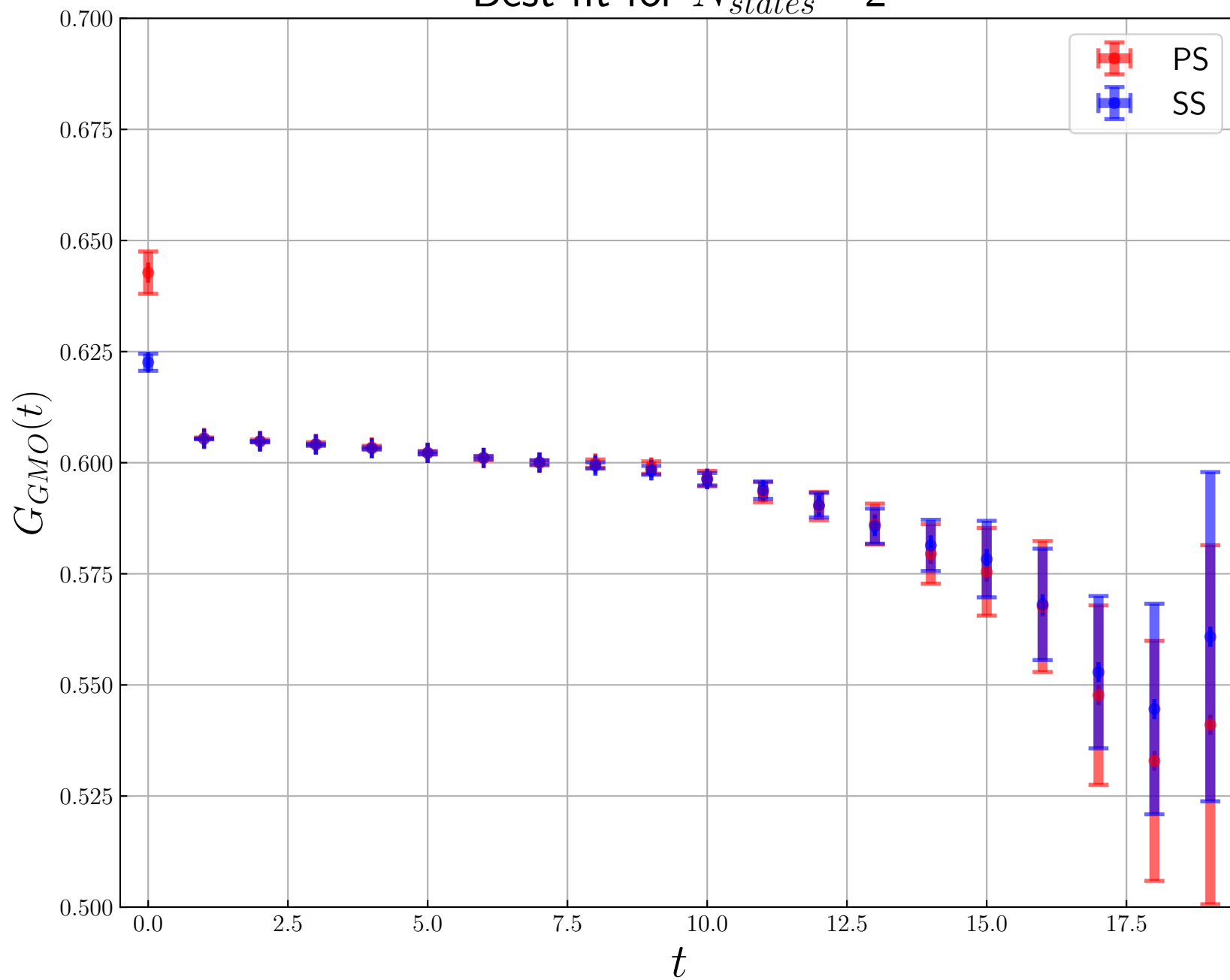
xi_E0= 0.788892 +- nan

1st e.s: 0.256778 +- nan

sigma_E0= 0.729141 +- nan

1st e.s: 0.132567 +- nan

Best fit for $N_{states} = 2$



gmo_ratio_eff for $N_{states}=2$

