

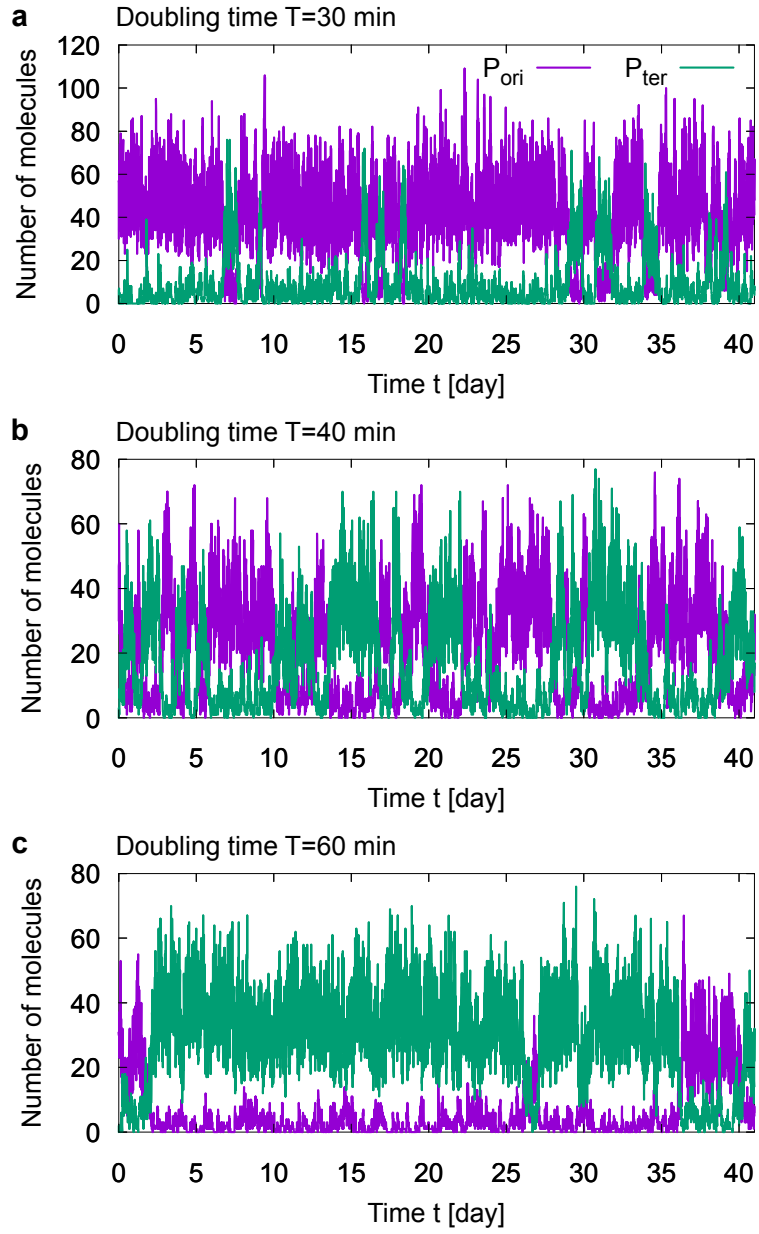


Figure S6: Number of protein monomers from sample stochastic trajectories computed for  $r_g = 0.002$  and for three doubling times: 30, 40, and 60 min.