Mean Error for inference of s s= 0.7 (S=100%) s= 2.4 (S=1000%) s= 4.6 (S=10000%) 0.75 0.2 1.0 0.50 0.1 0.25 -0.5 0.0 0.00 0.0 method Mean Error birthDeathMCMC s= 6.9 (S=100000%) s= 9.2 (S=1000000%) s= 11.5 (S=10000000%) cloneRate - phylofit 1.5 1.0 0.5 0.0 11 32 58 75 91 32 58 75 11 32 58 75 91 n(Sampled Mutant Clade Size)