RNA SYNTHESIS ≈ 40 nucleotides / sec BNID: 111871 (A) rRNA synthesis mRNA synthesis tRNA synthesis BNID: 104186;106254 $L_{rRNA\ genes} \approx 4500\ nucleotides$ BNID: 108093 $N_{tRNA} \approx \frac{3000}{\text{amino acid}} \times 20 \text{ amino acids}$ $N_{mRNA} \approx \frac{1 \text{ mRNA}}{1000 \text{ proteins}} \times 3 \times 10^6 \text{ proteins}$ $r_{\text{RNAP loading}} \approx 1 \text{ / sec}$ BNID: 111997 $\approx 6 \times 10^4 \text{ tRNA}$ ≈3000 mRNA (at steady state) BNID: 107873 $L_{RNAP footprint} \approx 40 \text{ nucleotides}$ $r_{degradation} \approx 1 \text{ mRNA} / 300 \text{ s}$ BNID:111927 L_{RNA} ≈ 80 nucleotides BNID:102340 80 nucleotides $L_{mRNA} \approx 1000 \text{ nucleotides}$ BNID:100022 $N_{RNAP}^{(mRNA)} \approx \frac{N_{mRNA} \times r_{degradation} \times L_{mRNA}}{r} \approx 250 \text{ RNAP}$ $N_{RNAP} \approx N_{RNAP}^{\text{(riftNA)}} + N_{RNAP}^{\text{(miRNA)}} + N_{RNAP}^{\text{(miRNA)}} \approx 700 \text{ RNAP}$ (B) (C) 105 10^{4} RNA Polymerases per cell 0 70 (RpoD) per cell 10^{4} 0 replication fork scaling replication fork scaling estimated value estimated value Li et al. 2014 Li et al. 2014 Peebo et al. 2015 Peebo et al. 2015 Schmidt et al. 2016 Schmidt et al. 2016 Valgepea et al. 2013 Valgepea et al. 2013

 10^{2}

0.00

0.25

0.50

1.00

growth rate [hr⁻¹]

1.25

1.50

1.75

 10^{2}

0.00

0.25

0.50

0.75

1.00

growth rate [hr-1]

1.25

1.50

1.75

2.00