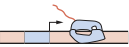


RNA synthesis



$$r_{\text{transcription}} \approx 40 \text{ nucleotides / (sec} \times \text{polymerase)}$$

BNID: 111871

RNA polymerase transcription rate

rRNA synthesis

$$L_{\text{rRNA genes}} \approx 4500 \text{ nucleotides}$$

BNID: 108093

summed nucleotide length of rRNA

$$r_{\text{RNAP loading}} \approx 1 / \text{sec}$$

BNID: 102362

promoter loading rate

$$L_{\text{RNAP spacing}} \approx \frac{r_{\text{transcription}}}{r_{\text{RNAP loading}}} \approx 40 \text{ nucleotides}$$

average spacing between RNA polymerase

$$N_{\text{RNAP}}^{(\text{rRNA})} \approx L_{\text{rRNA genes}} \times N_{\text{rRNA genes}} \times L_{\text{RNAP spacing}} \approx 1000 \text{ RNAP}$$

mRNA synthesis

$$N_{\text{mRNA}} \approx \frac{1 \text{ mRNA}}{1000 \text{ proteins}} \times 3 \times 10^6 \text{ proteins}$$

BNID: 104186; 106254

$$\approx 3000 \text{ mRNA (at steady state)}$$

number of mRNA molecules per cell

$$r_{\text{degradation}} \approx 1 \text{ mRNA / 300 s}$$

BNID: 111927

mRNA degradation rate

$$L_{\text{mRNA}} \approx 1000 \text{ nucleotides}$$

BNID: 100022

average length of mRNA

$$N_{\text{RNAP}}^{(\text{mRNA})} \approx \frac{N_{\text{mRNA}} \times r_{\text{degradation}} \times L_{\text{mRNA}}}{r_{\text{transcription}}} \approx 250 \text{ RNAP}$$

tRNA synthesis

$$N_{\text{tRNA}} \approx 4 \times 10^5 \text{ tRNA / cell}$$

BNID: 108611

number of tRNA molecules per cell

$$L_{\text{tRNA}} \approx 80 \text{ nucleotides / tRNA}$$

BNID: 102340

average length of tRNA

$$N_{\text{RNAP}}^{(\text{tRNA})} \approx \frac{N_{\text{tRNA}} \times L_{\text{tRNA}}}{r_{\text{transcription}} \times t_{\text{division}}} \approx 150 \text{ RNAP}$$

$$N_{\text{RNAP}} \approx N_{\text{RNAP}}^{(\text{rRNA})} + N_{\text{RNAP}}^{(\text{mRNA})} + N_{\text{RNAP}}^{(\text{tRNA})} \approx 1500 \text{ RNAP}$$

total number of RNAP complexes