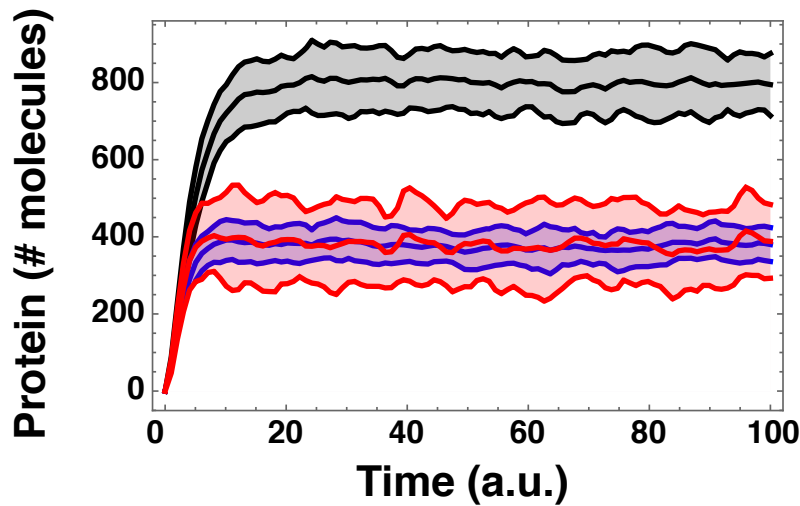


**A**

Model	Burst Freq.	$b_{P_{eff}}$	$b_{r_{eff}}$	T *	Analytical Prediction				From Simulations			
					<P>	Variance	Fano factor	CV <sup>2</sup>	<P>	Variance	Fano factor	CV <sup>2</sup>
Two-state Model (High mean)	0.091	100	20	0	18182	36363636	2000	0.11	17466 (±1060)	33912654 (±3230930)	1954 (±311)	0.11 (±0.02)
Two-state Model (one-half mean)	0.046	100	20	0	9091	18182000	2000	0.22	9614 (±27)	19601286 (±1077555)	2039 (±118)	0.21 (±0.01)
Auto-repression (one-half mean)	0.046	100	20	≤ 1	9091	9091000	≥ 1000	≥ 0.11	9049 (±89)	11189103 (±964941)	1237 (±114)	0.14 (±0.01)
Auto-depletion (one-half mean)	0.091	50	20	≤ 1	9091	6060606	≥ 500	≥ 0.055	9336 (±164)	5716108 (±381944)	612 (±30)	0.07 (±0.01)
Auto-repression (one-third mean)	0.030	100	20	≤ 2	6061	8661169	≥ 667	≥ 0.11	5248 (±219)	6611475 (±163289)	1260 (±21)	0.24 (±0.01)
Auto-depletion (one-third mean)	0.091	33.3	20	≤ 2	6061	2000130	≥ 222	≥ 0.037	5078 (±13)	1675023 (±126109)	330 (±26)	0.06 (±0.01)

\*For the analytical predictions the linear feedback approximation was used where  $|T| = \frac{\langle p \rangle_{NFB}}{\langle p \rangle_{FB}} - 1$ . The subscript NFB means without feedback and the subscript FB means with feedback. Since the linear feedback approximation does not saturate, it overestimates the magnitude of  $|T|$  for realistic repression relationships that do saturate. Accordingly, the analytical predictions for Fano factor and CV<sup>2</sup> should be interpreted as lower limits.

**B**



**C**

	Two State	Two State (lower mean)	Transcriptional Negative FB	Precursor Depletion FB	Simple RNA Depletion FB
Mean	800	400	381	376	381
CV <sub>2</sub>	0.01	0.025	0.07	0.009	0.008
Fano	8.5	9.5	30	5.6	3.2

**D**

