Biology of the *noisy* gene

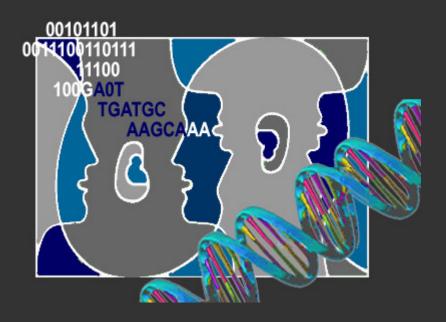
Universidad Autónoma de Madrid Jan 2008

Juan F. Poyatos

Logic of Genomic Systems laboratory

Spanish National Biotechnology Centre (CNB)

day III: noisy bacteria



- Regulation of noise (*B. subtilis*)
- Intrinsic/Extrinsic noise (*E. coli*)
- Noise time scales



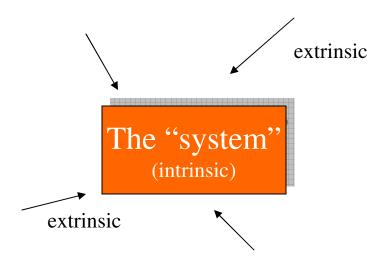
Stochastic dynamics of gene expression, experiments!

- Intrinsic noise in Bacillus subtilis

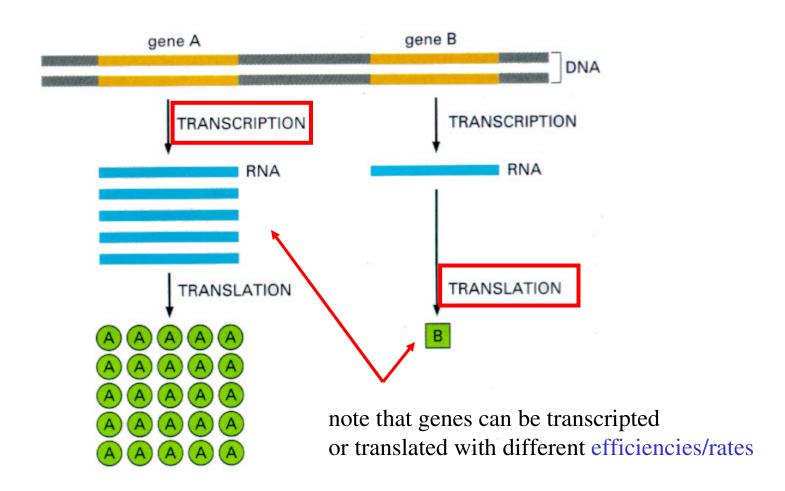
Molecular fluctuations within single cells (biochemical noise) → variability in a genetically identical population (phenotypic noise).

- Extrinsic vs. intrinsic noise in Escherichia coli

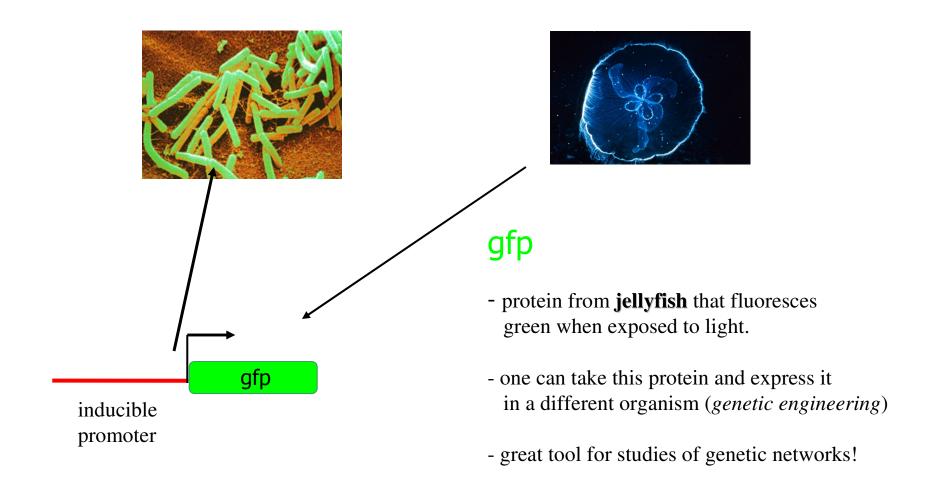
Detection of noise and discrimination between intrinsic and extrinsic noise

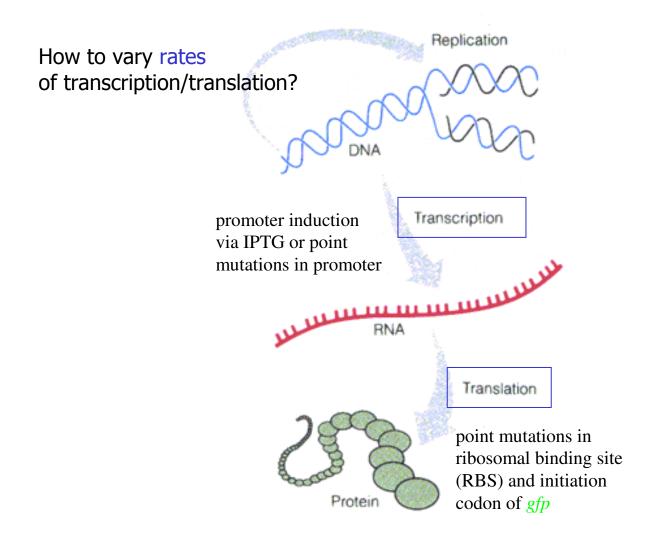


Gene expression in a nutshell



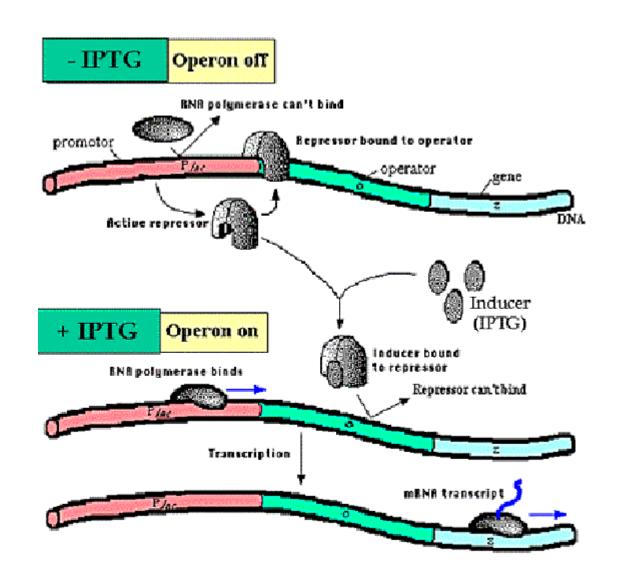
- A single-copy chromosomal gene with an inducible promoter was introduced in *B. subtilis*





Transcription efficiency

promoter induction via IPTG or ...



Induction of the *lac* Operon

Transcription efficiency

... or point mutations in promoter

 Transcriptional mutants: point mutations 	5
in the P _{spac} promoter	

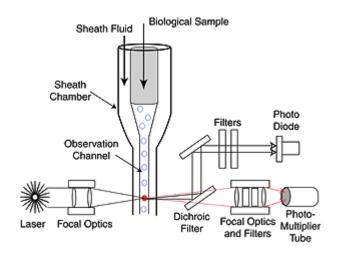
Strain	–10 regulatory region –10 +1	Transcriptional efficiency
ERT57	CAT AAT GTG TGT AAT	6.63
ERT25	CAT AAT GTG TGG AAT	1.00
ERT53	CAT AAT GTG TGC AAT	0.79
ERT51	CAT AAT GTG TGA AAT	0.76
ERT55	CAT AAT GTG TAA AAT	0.76

Translational efficiency

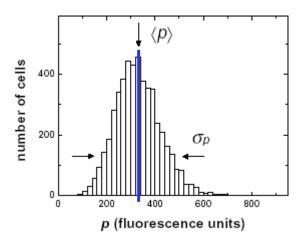
point mutations in ribosomal binding site (RBS) and initiation codon of *gfp*

• Translational mutants: point mutations in the RBS and initiation codon of *gfp*

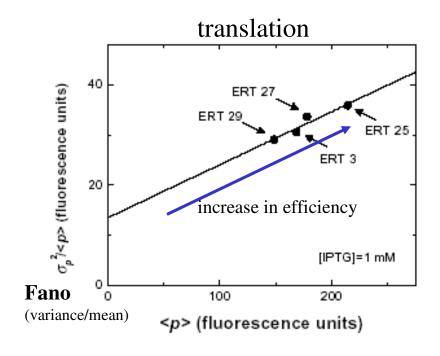
Strain	Ribosome binding site	Initiation codon	Translational efficiency
ERT25	GGG AAA AGG AGG TGA ACT	ACT ATG	1.00
ERT27	GGG AAA AGG AGG TGA ACT	ACT <u>T</u> TG	0.87
ERT3	GGG AAA AGG TGG TGA ACT	ACT ATG	0.84
ERT29	GGG AAA AGG AGG TGA ACT	ACT <u>G</u> TG	0.66



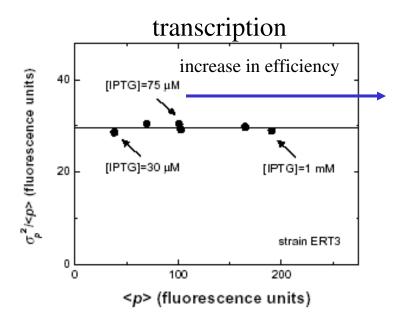
GFP expression level is measured for single cells in a bacterial population using flow cytometry



Expression level vary from cell to cell (phenotypic noise) as a consequence of molecular fluctuations within single cells (biochemical noise)



translational efficiency vs. transcriptional efficiency



Recall: gene expression model

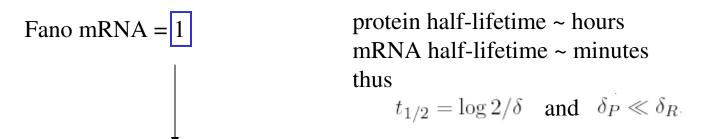
$$\frac{d[mRNA]}{dt} = k_R - \delta_R[mRNA] \qquad mRNA \xrightarrow{\delta_R} \emptyset,
\frac{d[P]}{dt} = k_P[mRNA] - \delta_P[P] \qquad mRNA \xrightarrow{k_P} P,
P \xrightarrow{\delta_P} \emptyset,$$

Master equation valid to ...

$$\frac{dp_{m,n}}{dt} = -p_{m,n}[m\delta_R + mk_P + k_R + n\delta_P]
+ p_{m,n+1}(n+1)\delta_P + p_{m+1,n}(m+1)\delta_R
+ p_{m,n-1}k_Pm + p_{m-1,n}k_R$$

... to get the final expressions for the macroscopic statistics

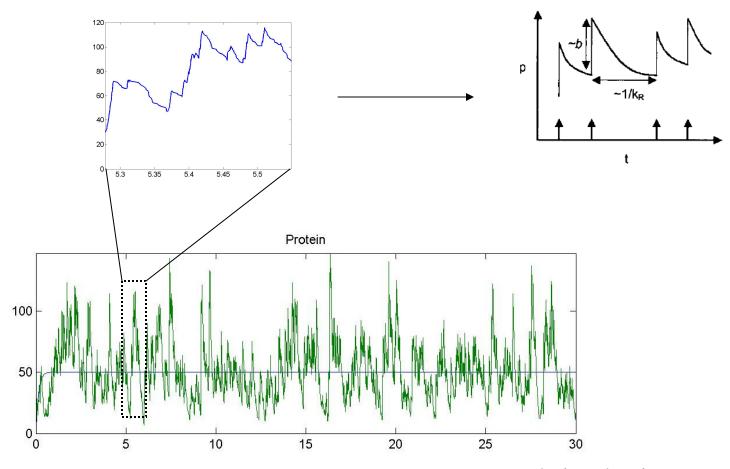
$$\text{Fano Protein} = \ \frac{\langle n^2 \rangle - \langle n \rangle^2}{\langle n \rangle} = 1 + \frac{k_P/\delta_R}{1 + \delta_P/\delta_R} \approx 1 + \frac{k_P}{\delta_R} \qquad \text{translation efficiency influences noise}$$



transcription efficiency does not influence noise

"Random bursts model"

$$b = \frac{k_P}{\delta_R}$$

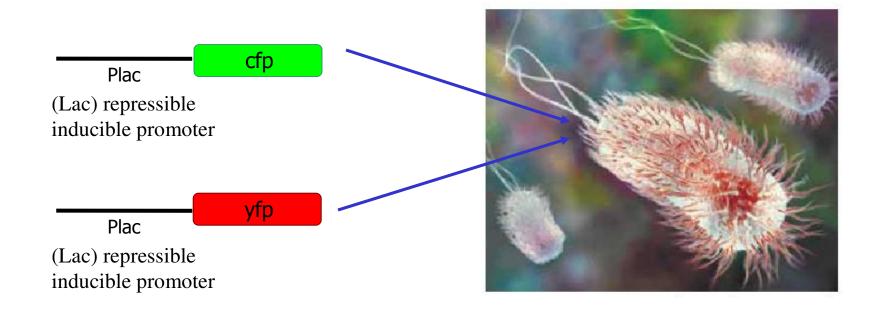


Translational noise control

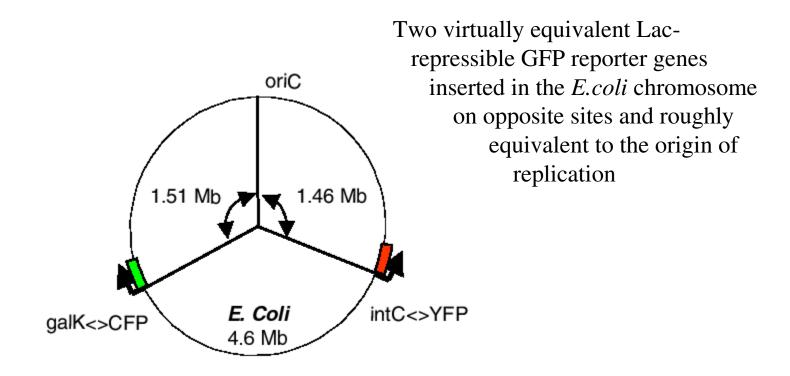
Intrinsic noise, even if all cellular conditions are equivalent for cells, we have seen that the reactions associated to transcription and translation originate noise Extrinsic noise, other molecular species (genes themselves too!), e.g., RNA polymerase, originate noise

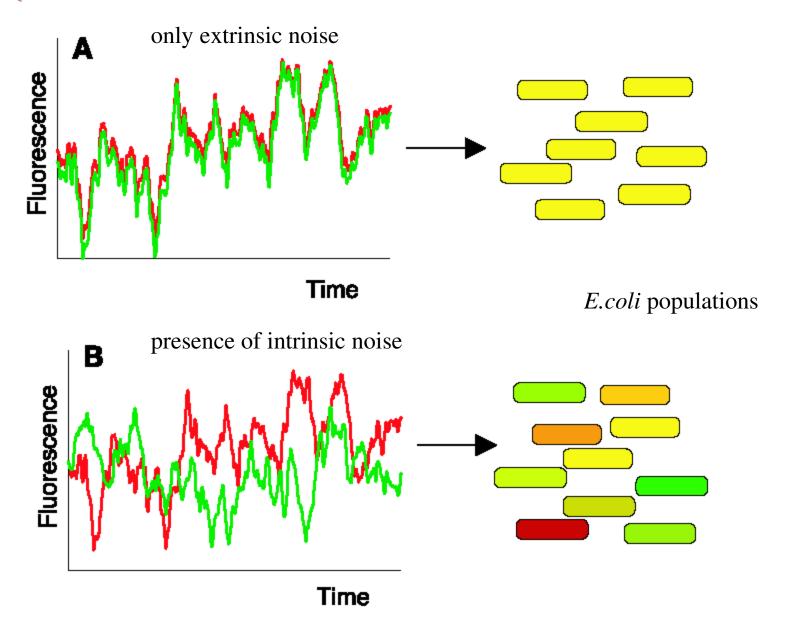
Can we discriminate both sources of noise?

Intrinsic noise:= Difference in gene expression that arises between two identical copies of a gene expressed under precisely the same conditions



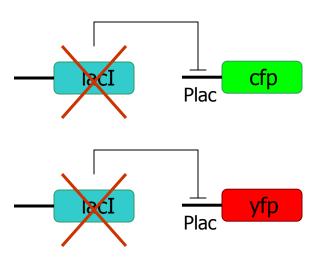
Intrinsic noise:= Difference in gene expression that arises between two identical copies of a gene expressed under precisely the same conditions

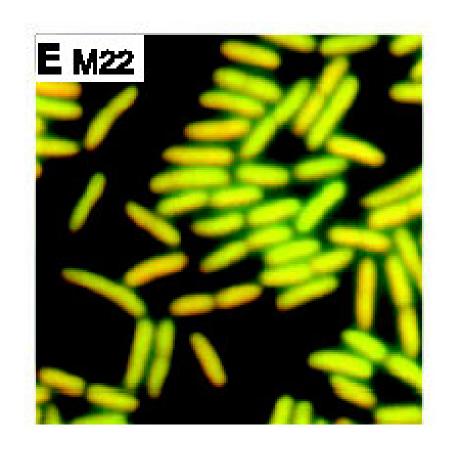




(lacl-cells)

strong **constitutive promoter** stable protein high transcription low noise

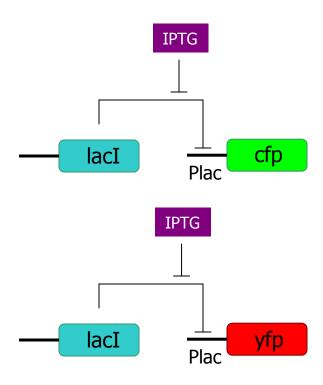


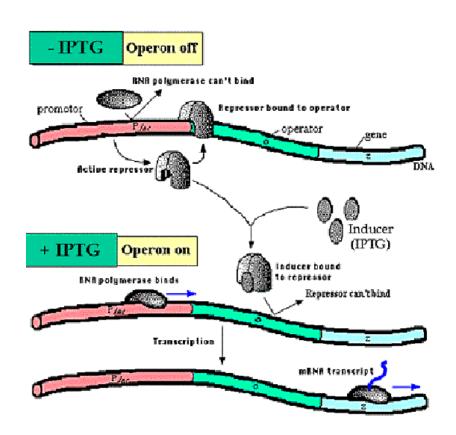




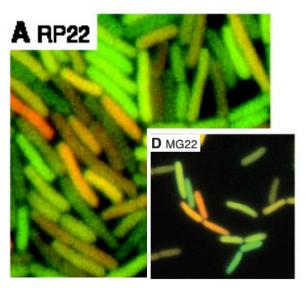
(lacl+cells)

inducible promoter



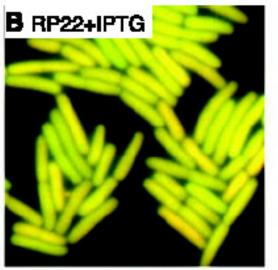


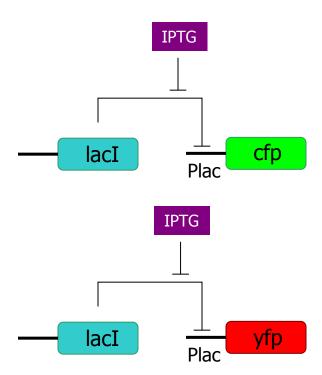
Induction of the lac Operon

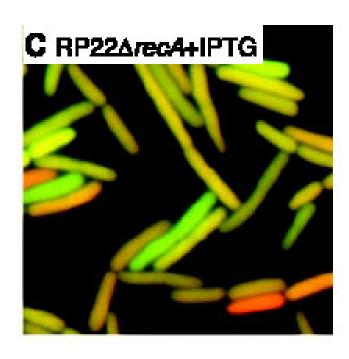


Promotors repressed by wild-type repressor (lacI) gene (-IPTG operon OFF) low transcription, high noise

Presence of inducer (+IPTG operon ON) high transcription, low noise

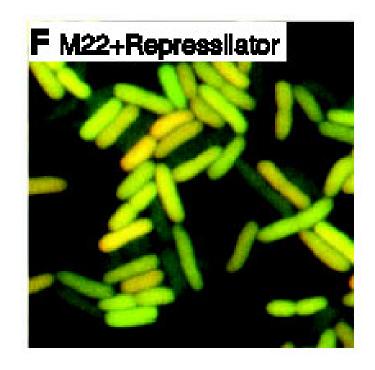


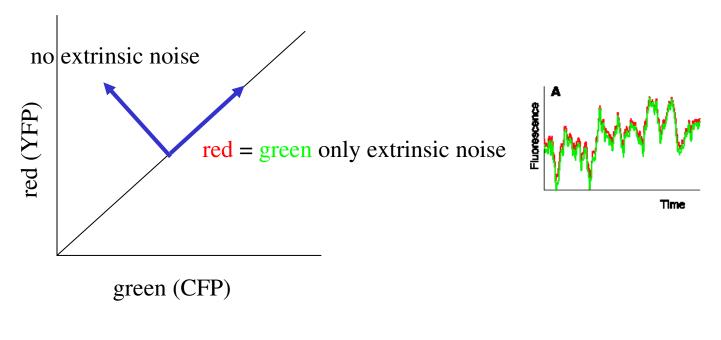




modified genetic background noisy

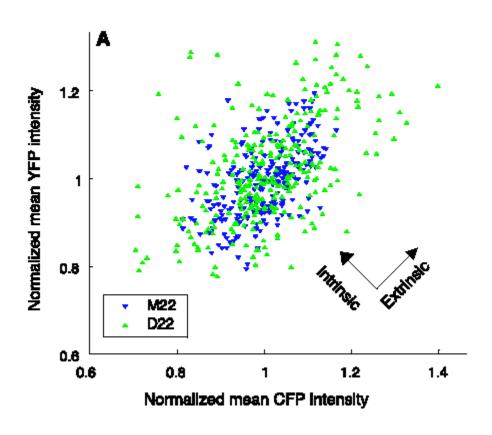
oscillating expression also noisy

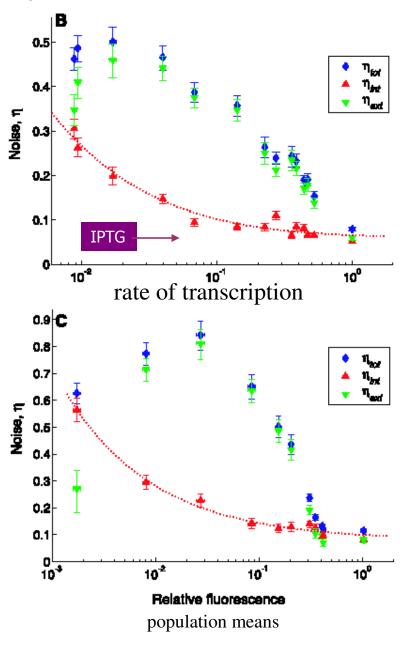




noise =
$$\frac{\text{variance }(\sigma^2)}{\text{mean}^2}$$
; $\text{noise}_{\text{total}}^2(\xi) = \text{noise}_{\text{intrinsic}}^2 + \text{noise}_{\text{extrinsic}}^2$

$$\rightarrow \text{ different to previous definition } n_2 = \frac{\sigma^2}{\langle n \rangle}$$





(recA+ lacI- cells)
LacI in plasmid

intrinsic noise decreases with rate of transcription (transcription in these experiments **does** have an effect on noise!)

extrinsic noise peaks at intermediate levels (fluctuations in Lac repressor proteins. At high or low IPTG concentrations fluctuations are buffered by excess IPTG or excess LacI, respectively)

Intrinsic/Extrinsic noise; time scales

