

E. Quantitative characterization of the *lac* promoter

lac promoter of *E. coli*:

- best-studied system of molecular biology
 - all molecular components characterized
 - many mutants studied *in vivo*
 - most parameters measured *in vitro*
- exemplary model system of combinatorial gene regulation
 - involves activation, repression, and DNA looping

Quantitative confrontation of model and experiment

- applicability of the thermodynamic description of tsx control?
- can the *in vivo* behavior of a system be understood in terms of its molecular parts?

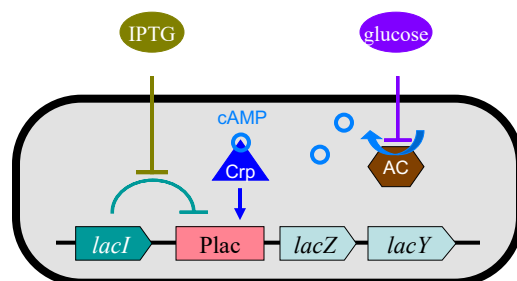
22

Review: regulation of the *lac*-operon of *E. coli*

Physiology:

- *lac*-operon: utilization of lactose
- repressed by the **Lac Repressor** (encoded by *lacI*)
- repression alleviated by allo-lactose (by-product of lactose metabolism) or the synthetic inducer **IPTG**
- activated by the global regulator **Crp**; requires the inducer **cAMP**
- cAMP synthesized endogenously by **Adenylate Cyclase** (encoded by *cyaA*)
- activity of **AC** repressed by **glucose** uptake

Function: expression ONLY in the presence of lactose **AND** absence of glucose



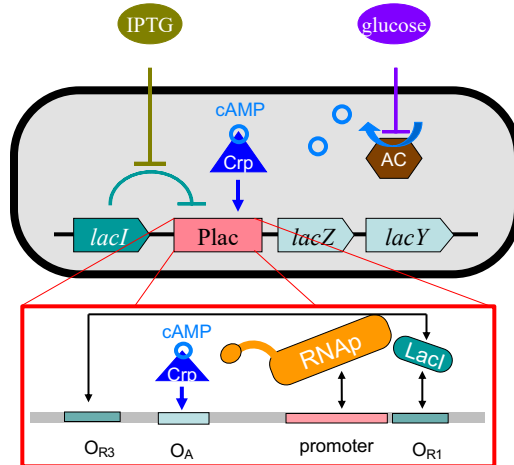
qualitative behavior:

IPTG	glucose	expression
low	high	OFF
low	low	OFF
high	high	OFF
high	low	ON

23

Review: regulation of the *lac*-operon of *E. coli*

Function: expression **ONLY** in the presence of lactose **AND** absence of glucose



qualitative behavior:

IPTG	glucose	expression
low	high	OFF
low	low	OFF
high	high	OFF
high	low	ON

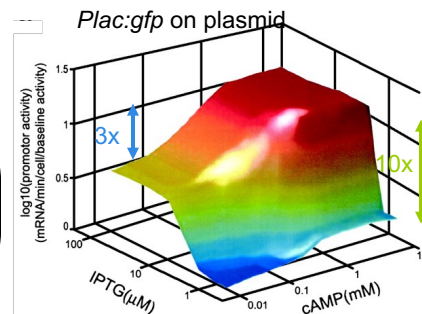
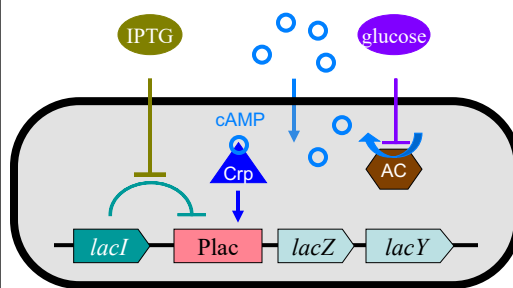
molecular ingredients:

- specific protein-DNA binding
- protein-protein interaction
- protein-mediated DNA looping

- theory: quantitative prediction of gene regulation by LacI, cAMP-Crp
 → expt: characterize **LacZ activity** for different levels of regulatory proteins
 -- control protein levels by varying the inducers (IPTG and cAMP)

24

Quantitative characterization



Previous expt: [Setty et al, PNAS, 2003]

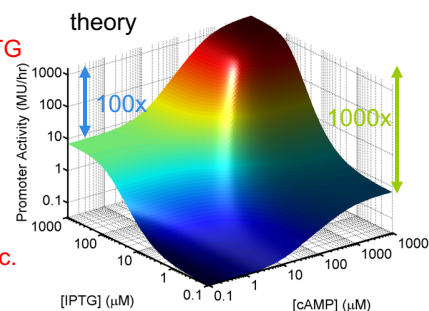
Grow cells in medium with glucose, cAMP, IPTG

- use glucose to suppress cAMP synthesis
- control cAMP-level extracellularly

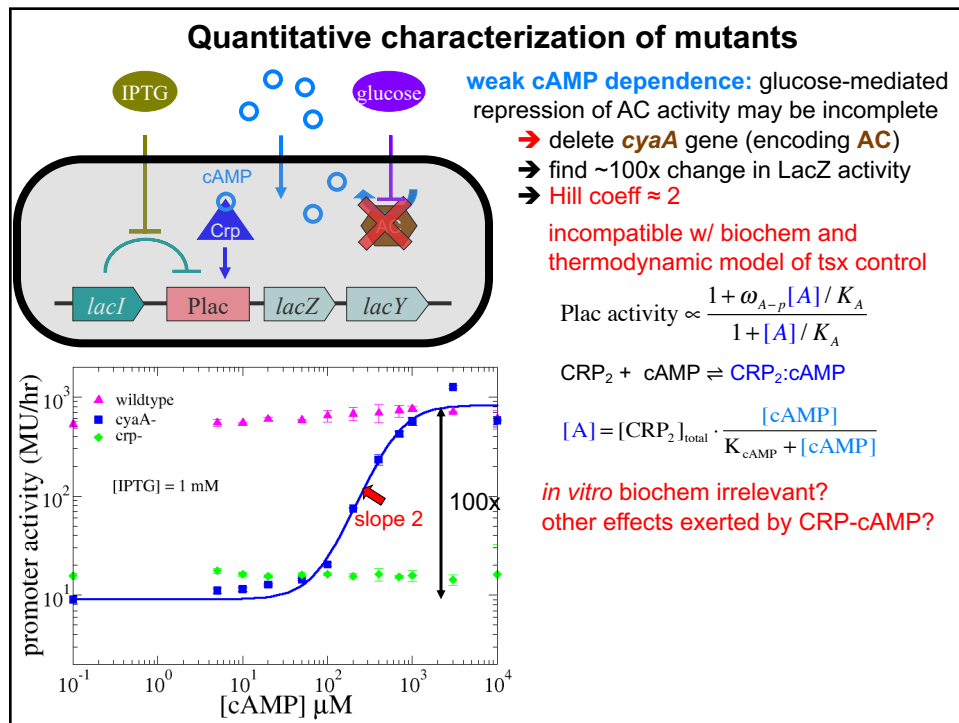
inconsistent with behavior of mutants:

$\Delta lacI$: > 1000x; Δcrp > 50x

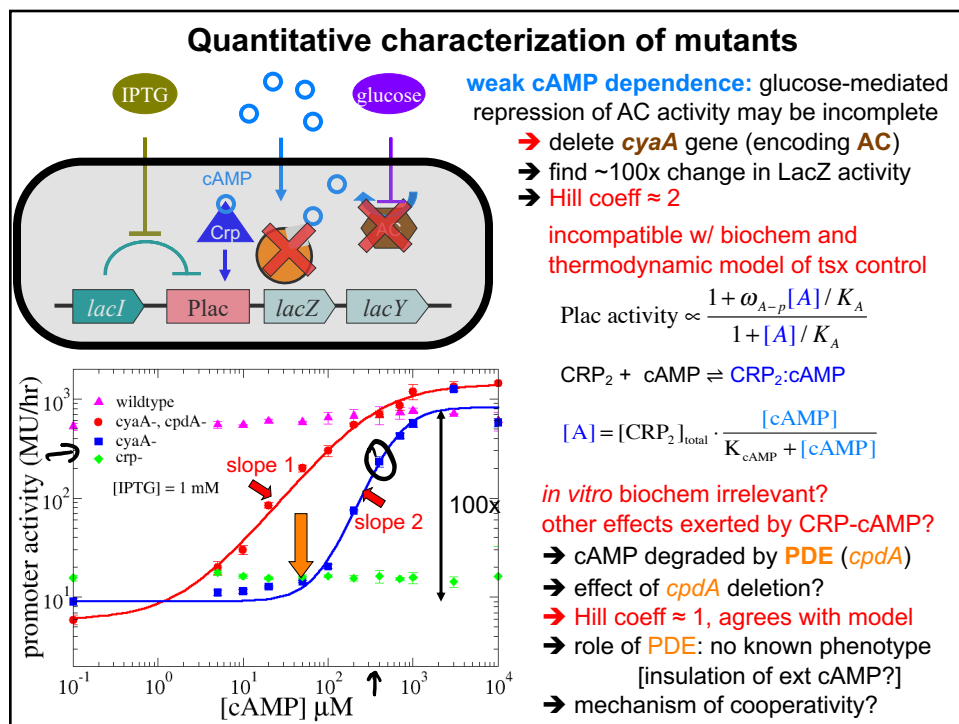
- possible problems: complex links between extracellular and intracellular inducer conc.



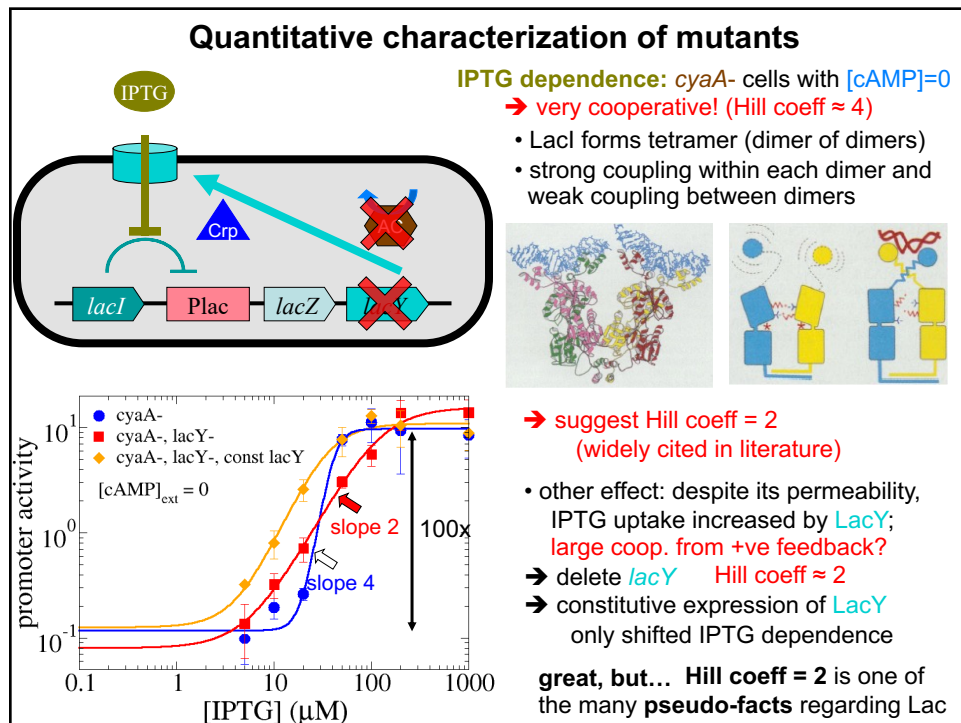
25



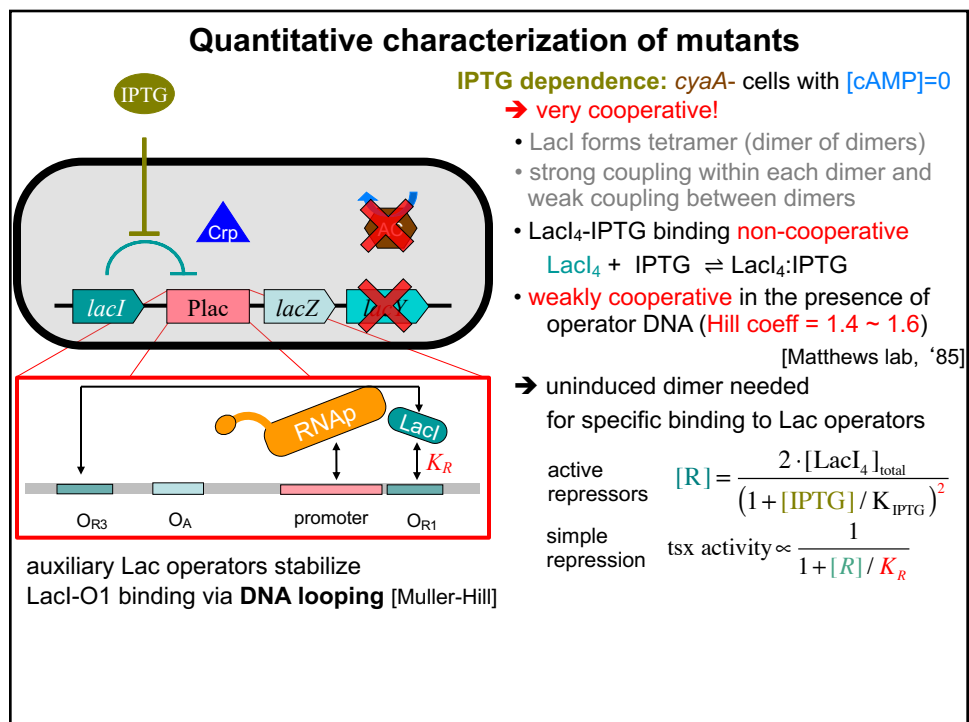
26



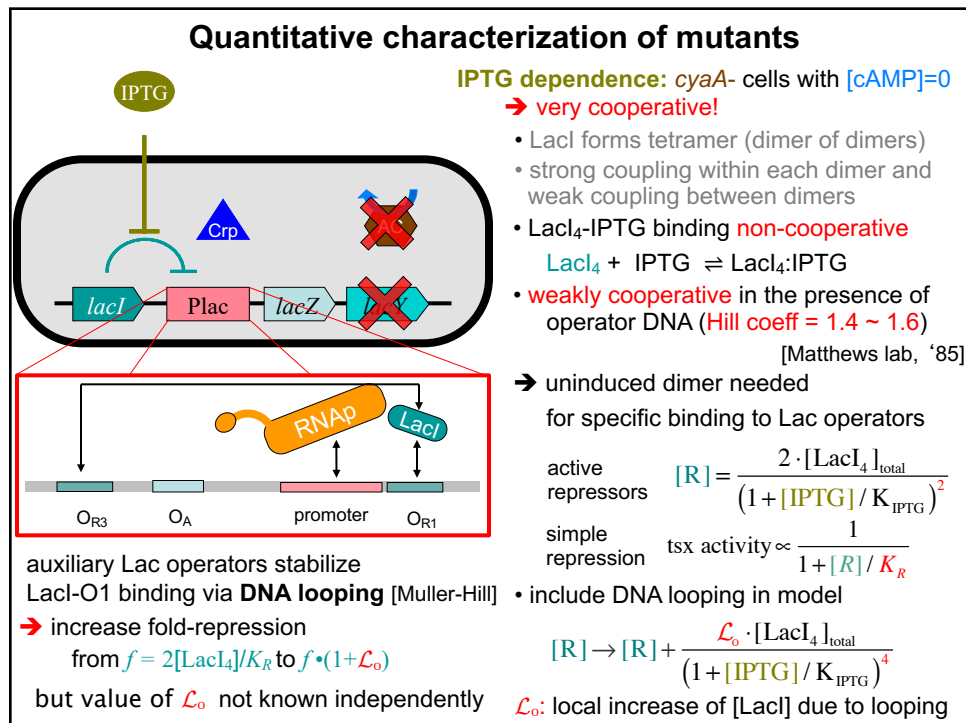
27



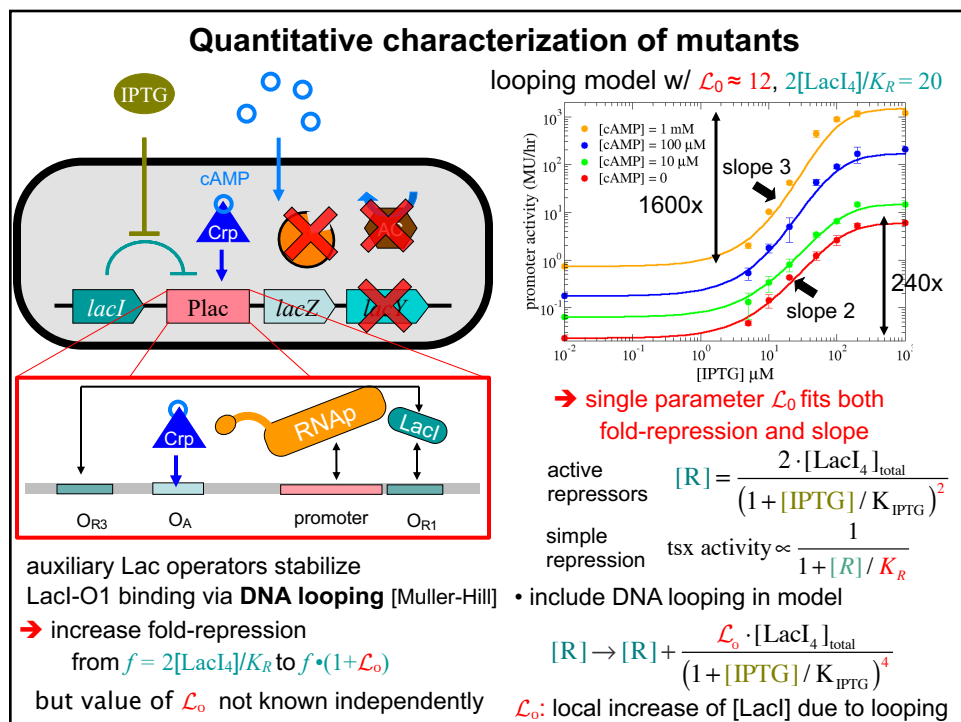
28



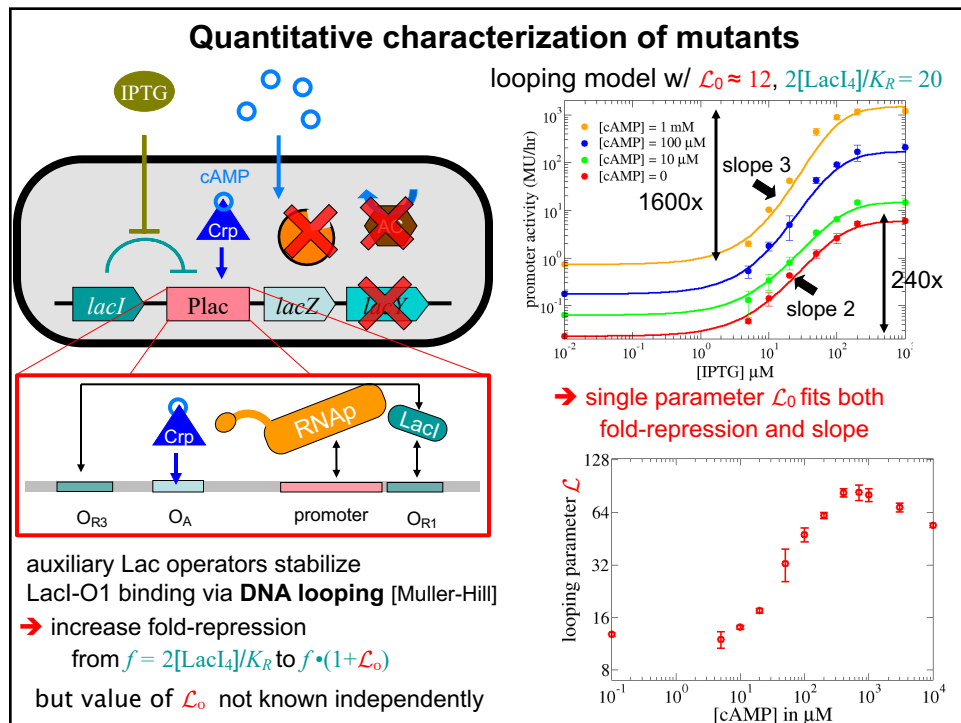
29



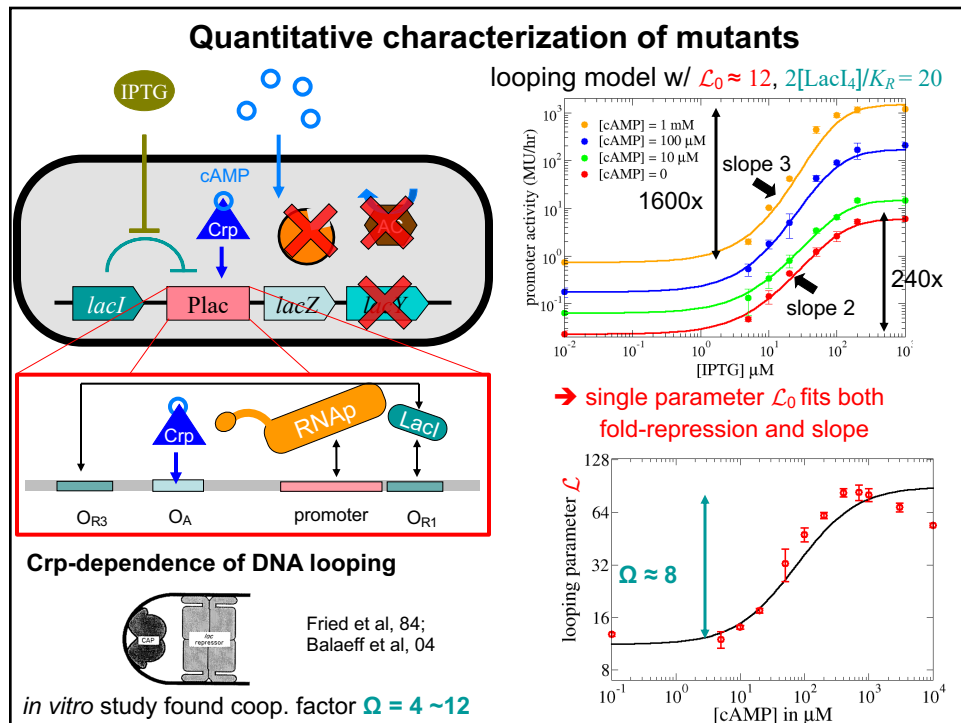
30



31



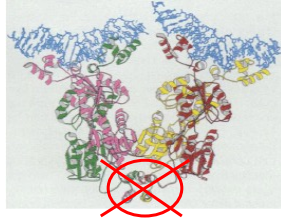
32



33

Direct probe of DNA looping *in vivo*

Use dimeric LacI mutant



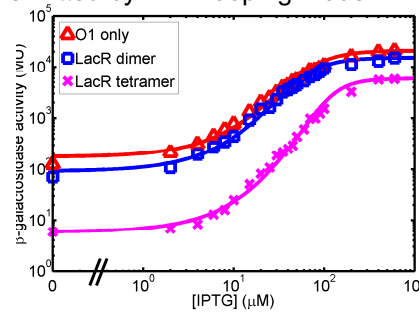
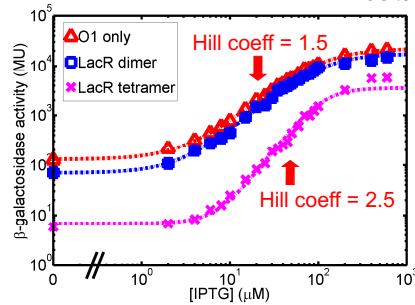
remove auxiliary operators



→ cooperativity in IPTG response requires DNA looping (Lac tetramer + auxiliary ops)

[Oehler & Muller-Hill, 06]

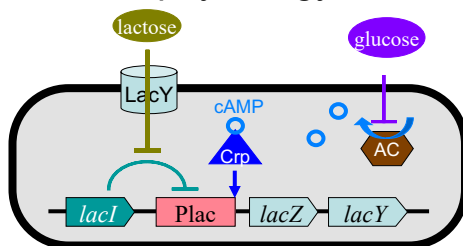
data well-fitted by DNA looping model



→ IPTG-LacI-operator interaction same as *in vitro*

34

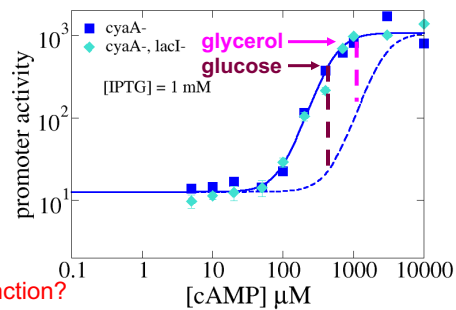
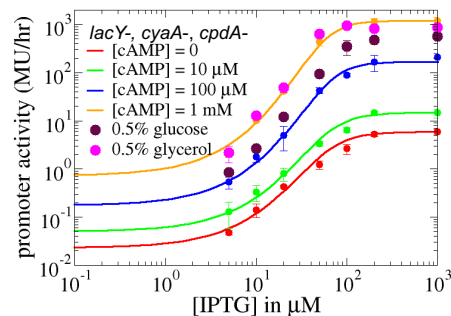
back to physiology



lactose	glucose	expression
low	high	OFF
low	low	OFF
high	high	OFF
high	low	ON

- only ~3x decrease from glucose to glycerol
- small fraction of dynamic range; (operating in saturation of cAMP-CRP)
- 10x change possible by reducing K_{crp}

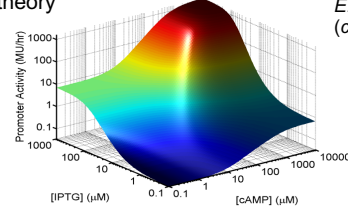
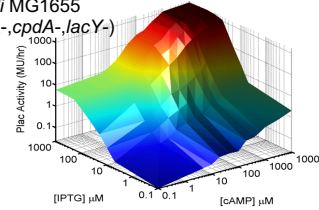
→ repression by glucose not the intended function?



35

Summary

theory


E. coli MG1655
(*cyaA*⁻, *cpdA*⁻, *lacY*⁻)


- **main findings for the *lac* promoter:**
 - Crp enhances DNA looping
 - abrupt IPTG response despite non-cooperative LacI-IPTG interaction;
 - **suggests physiological role of Crp-cAMP as enhancer of repression**
 - mechanism of Crp-LacI interaction?
 - coop cAMP response due to PDE; physiological function? mechanism?
- **general lessons for quantitative systems biology:**
 - hidden interaction and pseudo-facts abound even for the “best studied” system
 - quantitative description of *in vivo* biology is possible
 - need **solid, qualitative** knowledge of the components (e.g., Hill coeff)
(*in vitro* results surprisingly robust in this regard)
 - **(semi) quantitative** characterization generates spectrum of phenotypes
 - provides clues for identifying unknown components and mechanisms
 - provides **phenomenological description** of Plac for high-level studies