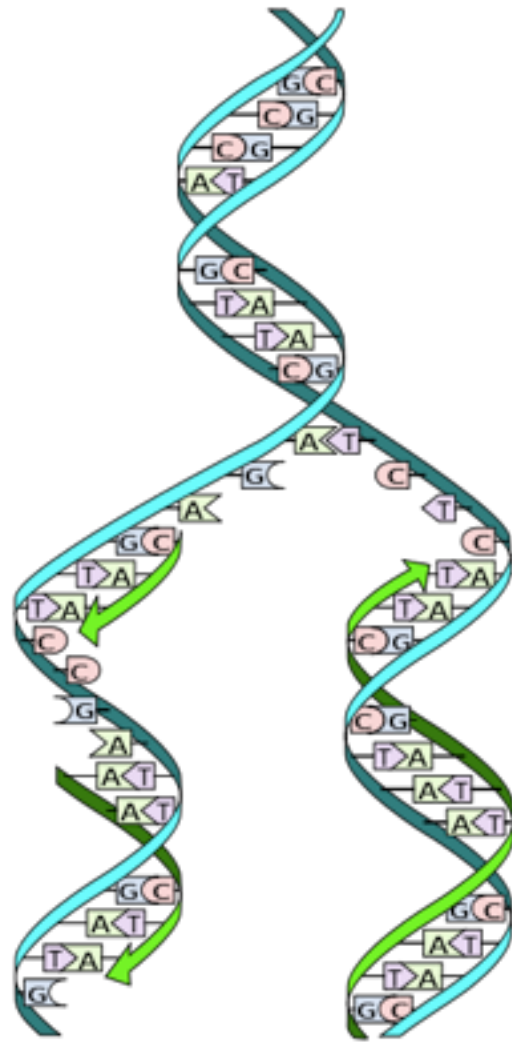


Thermodynamics of biological copying

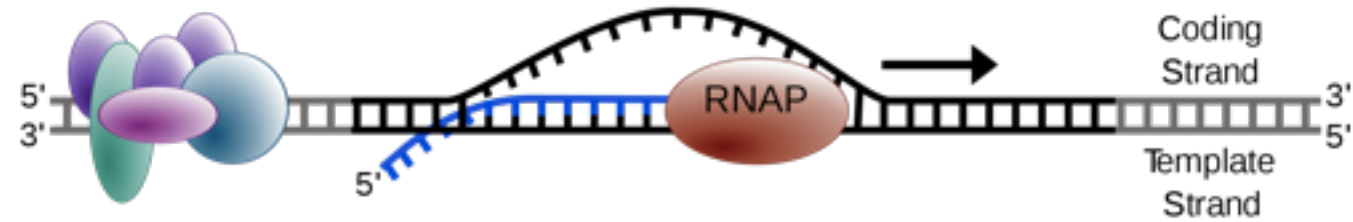
Simone Pigolotti (UPC, Barcelona)
Roma, 24/9/2013

with: Pablo Sartori (MPIPKS, Dresden)

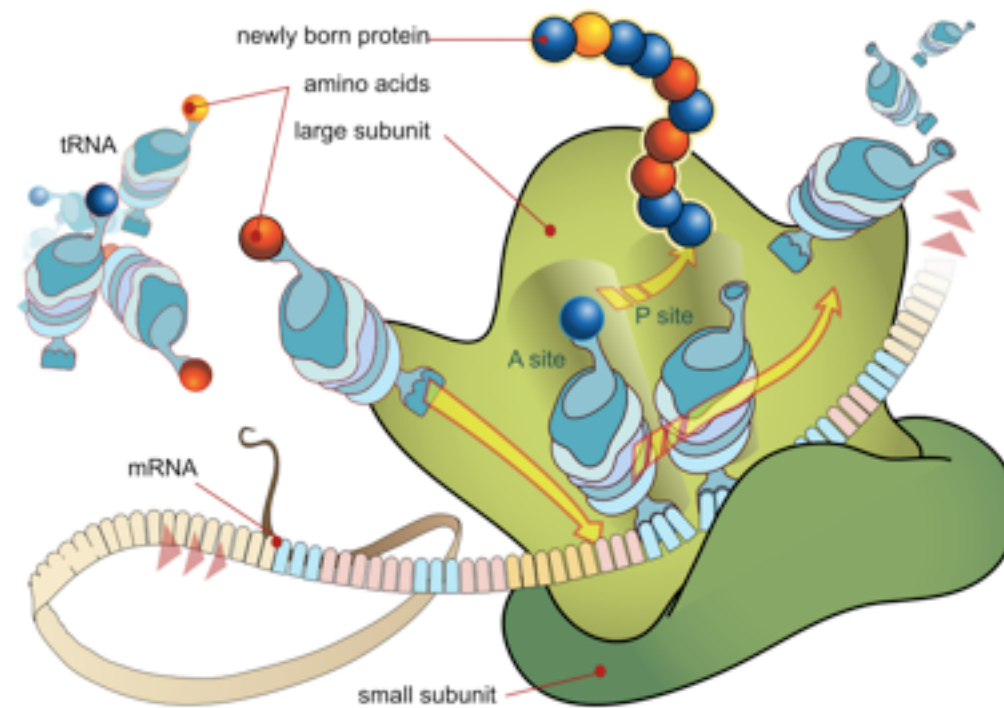
Biological copying



DNA duplication
error rate $\sim 10^{-9}$

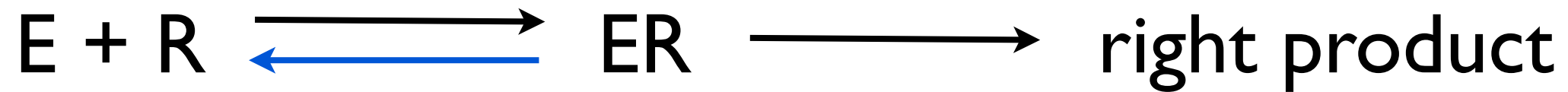
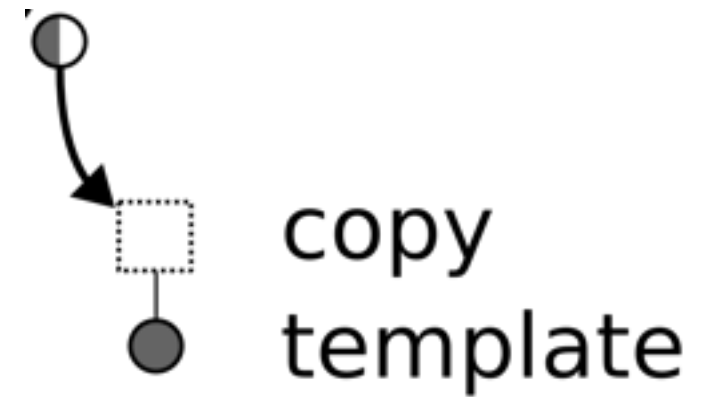


transcription
error rate $\sim 10^{-4}$



translation
error rate $\sim 10^{-4}$

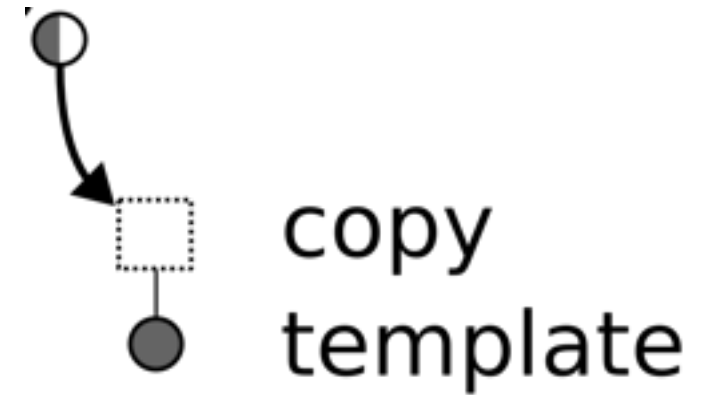
Enzymatic copying model



error rate $> e^{-\beta \Delta G}$ where $\Delta G = G_{EW} - G_{ER}$

minimum error achieved in quasi-equilibrium $\rightarrow$ very slow reaction

Enzymatic copying model



$$\text{error rate} > e^{-\beta \Delta G} \quad \text{where} \quad \Delta G = G_{EW} - G_{ER}$$

minimum error achieved in quasi-equilibrium $\rightarrow$ very slow reaction

DNA duplication:

$$kT \sim 2.479 \text{ kJ/mol}$$

$$\Delta G \sim 1 \text{ -- } 16 \text{ kJ/mol}$$

minimum error from binding energies > 0.01

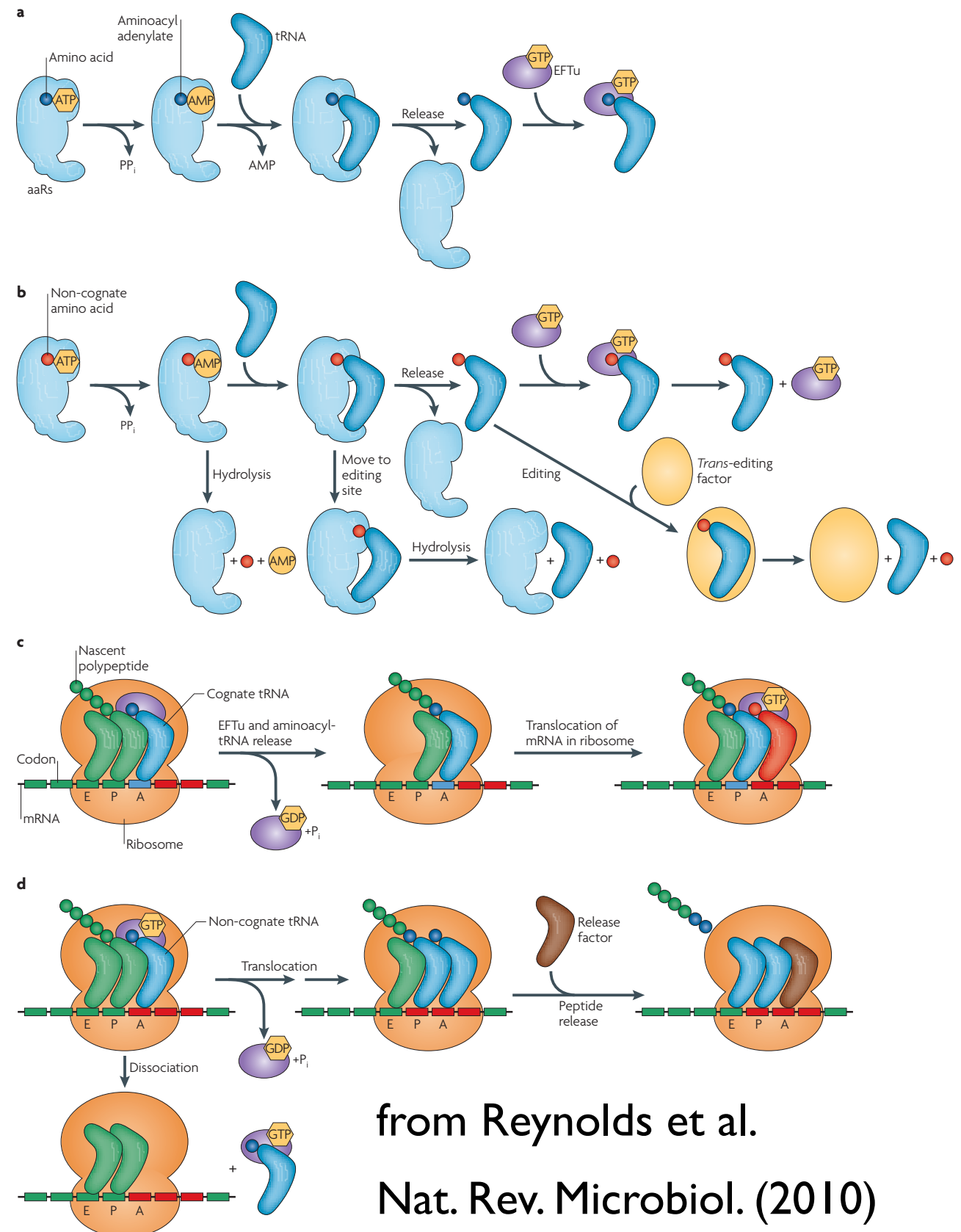
incompatible with observed value $= 10^{-9}$

Proofreading



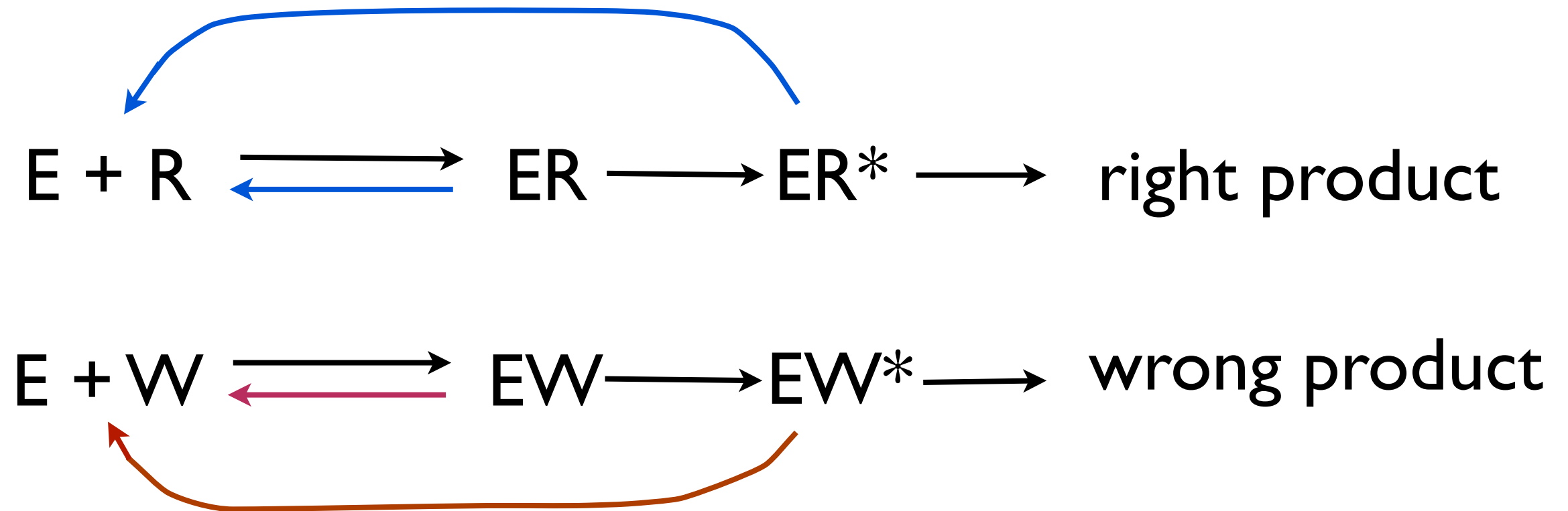
high fidelity is achieved via
complex multi-step reactions

example: translation



from Reynolds et al.
Nat. Rev. Microbiol. (2010)

Kinetic Proofreading



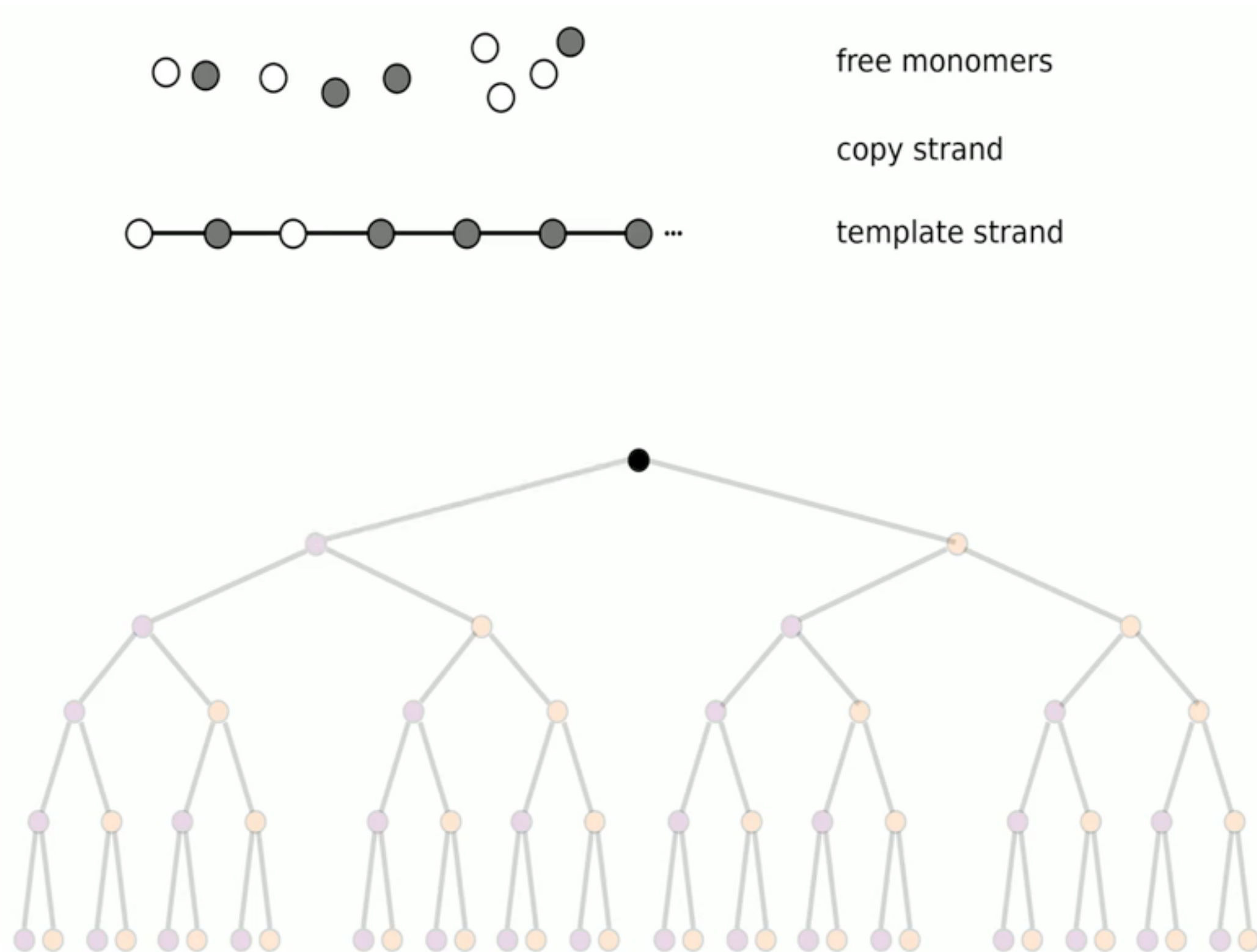
proofreading step is non-discriminating but reduces the error

$$\text{error rate} > \left(e^{-\beta\Delta G}\right)^2$$

proofreading is intrinsically dissipative

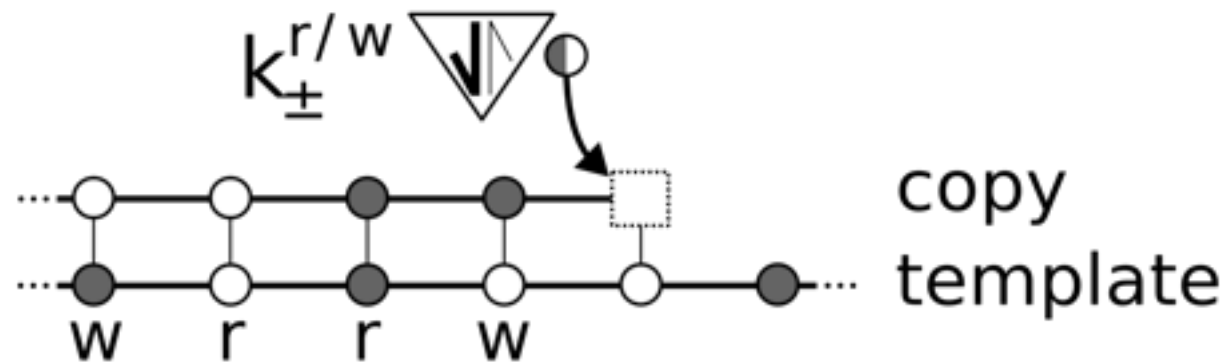
Hopfield, *PNAS* (1974)

Bennett's copolymerization model



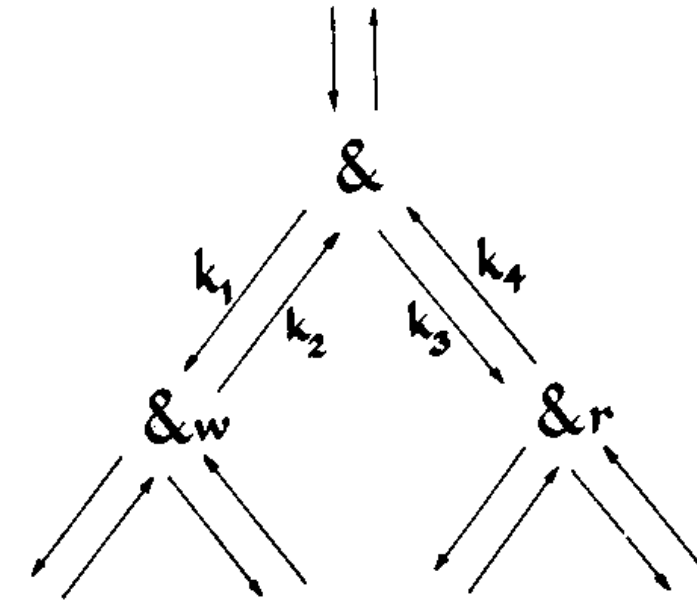
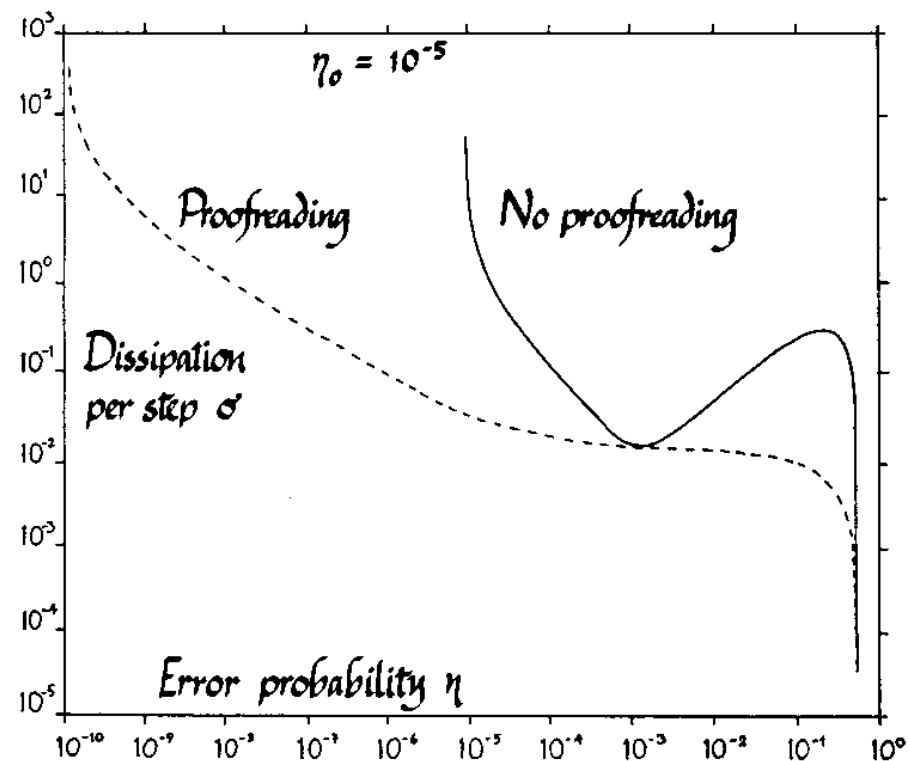
Bennett, *Biosystems* (1979)

Copolymerization model

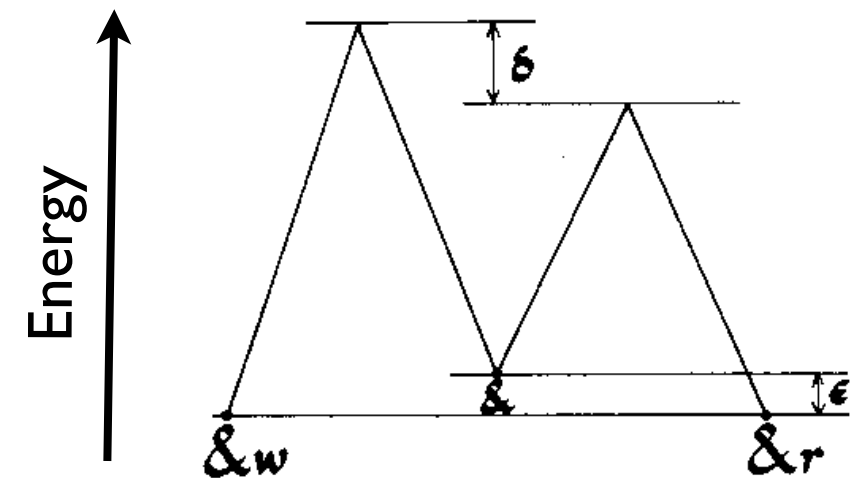


Copying of a long polymer

$k_{\pm}^{r/w}$ rates of binding/unbinding of right/wrong monomers



transition diagram



energy diagram - isoenergetic strains

Bennett, *Biosystems* (1979)
Andrieux and Gaspard, PNAS (2008), Esposito et al. JSTAT (2010)

Enzymatic copy vs. copolymerization

	enzymatic copy	copolymerization
ΔG	yes, determines minimum error	no, copies are isoenergetic
minimum error limit	adiabatic	$\dot{S} \rightarrow \infty$
proofreading	only dissipative step	less dissipation with proofreading than without

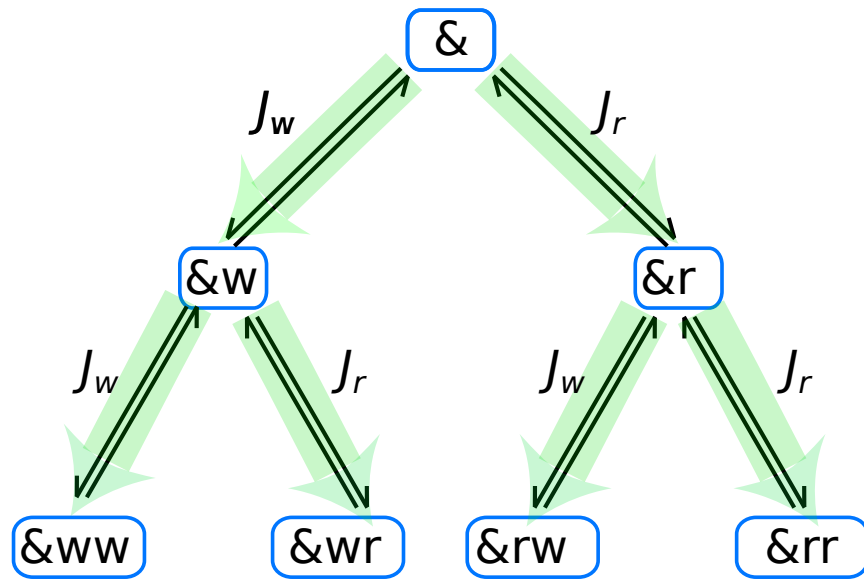
The bigger picture

I want a copying pathway with

- low error
- low dissipation
- high copying velocity

How should I design it?

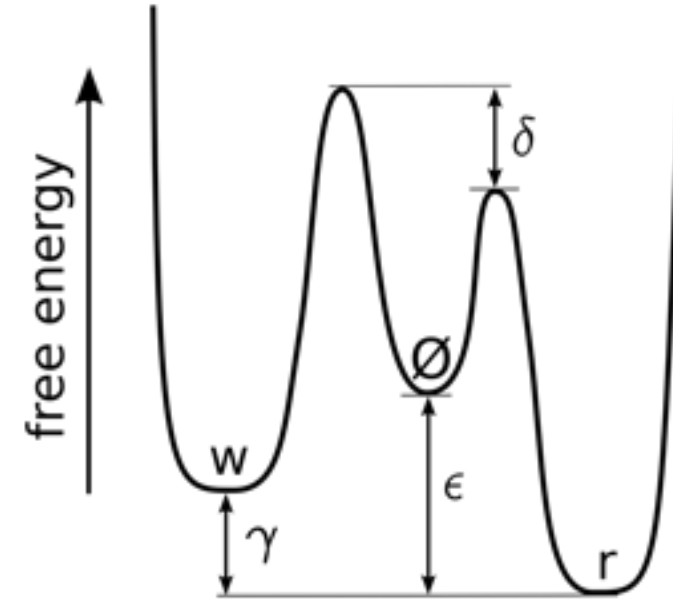
Copolymerization



write rates as Kramers rates:

$$k_+^r = \omega e^{\epsilon + \delta} \quad k_-^r = \omega e^{\delta}$$

$$k_+^w = \omega e^{\epsilon} \quad k_-^w = \omega e^{\gamma}$$



master equation:

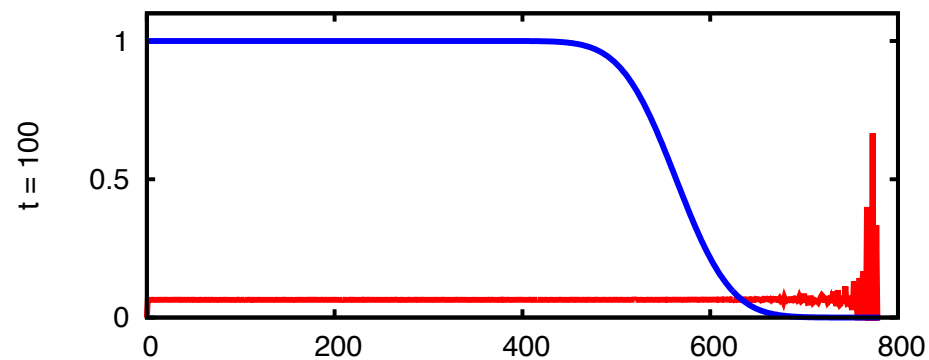
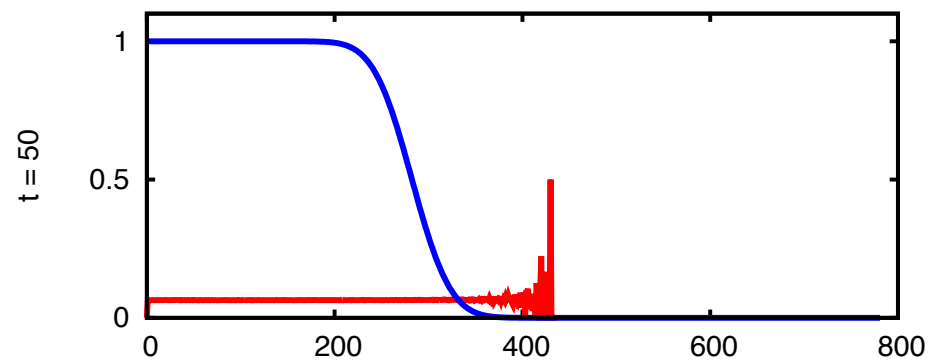
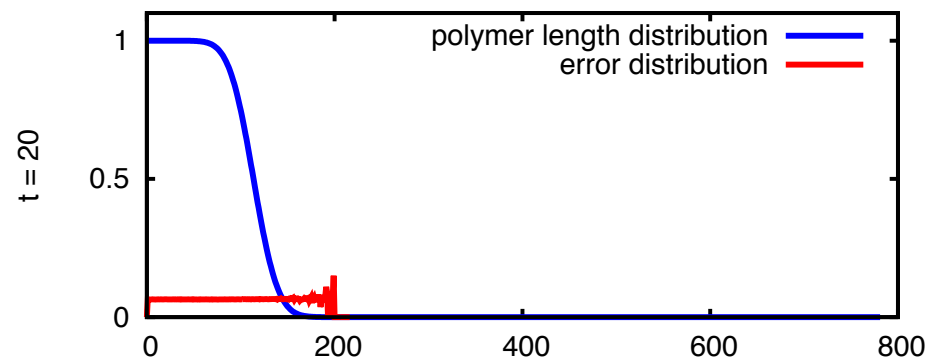
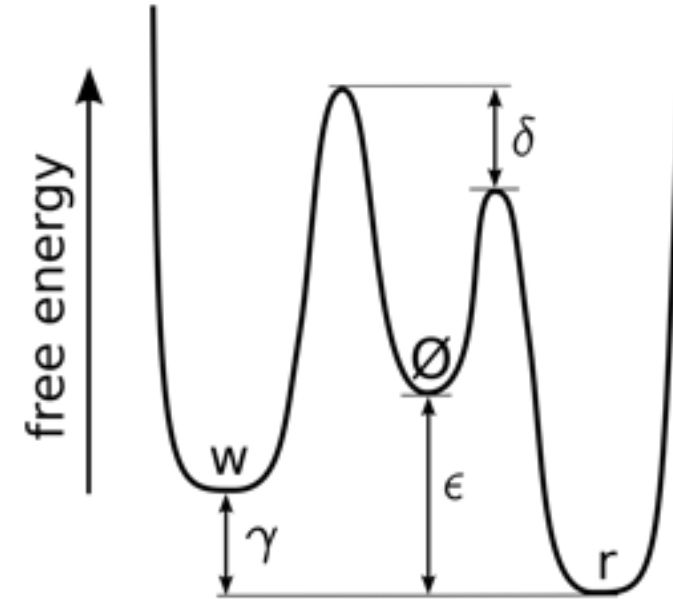
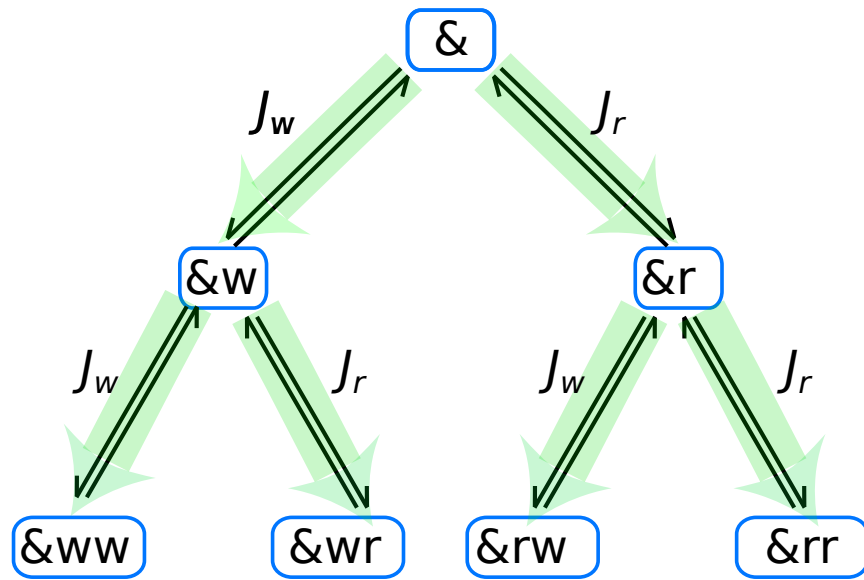
$$\frac{d}{dt}P(\cdots r) = k_+^r P(\cdots) + k_-^w P(\cdots rw) + k_-^r P(\cdots rr) - (k_-^r + k_+^w + k_+^r)P(\cdots r)$$

$$\frac{d}{dt}P(\cdots w) = k_+^w P(\cdots) + k_-^w P(\cdots ww) + k_-^r P(\cdots wr) - (k_-^w + k_+^w + k_+^r)P(\cdots w)$$

mean field solution:

$$\frac{\eta}{1 - \eta} = \frac{\text{net increase wrong monomers}}{\text{net increase right monomers}} = \frac{k_+^w - k_-^w \eta}{k_+^r - k_-^r (1 - \eta)}$$

Copolymerization



the ansatz

$$P(rwwrw \dots) \propto (1 - \eta)^{N_r} \eta^{N_w}$$

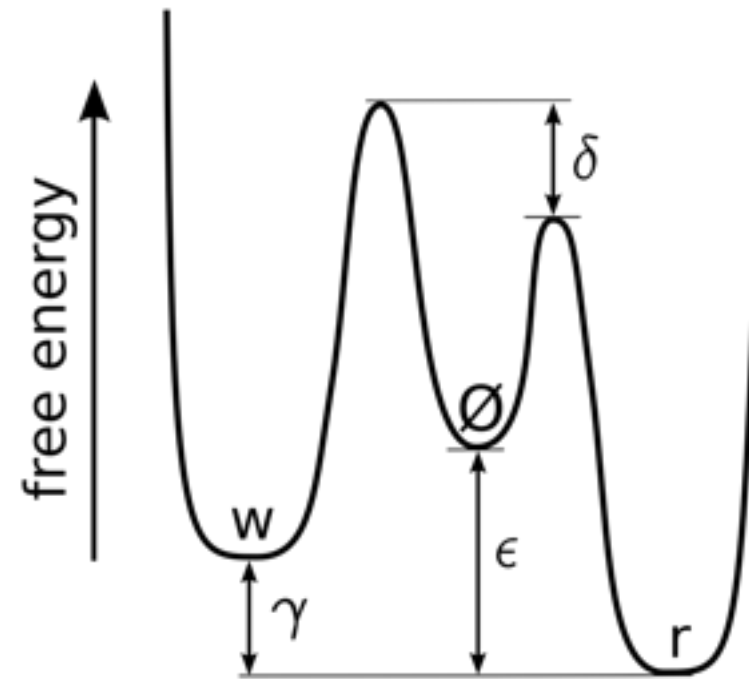
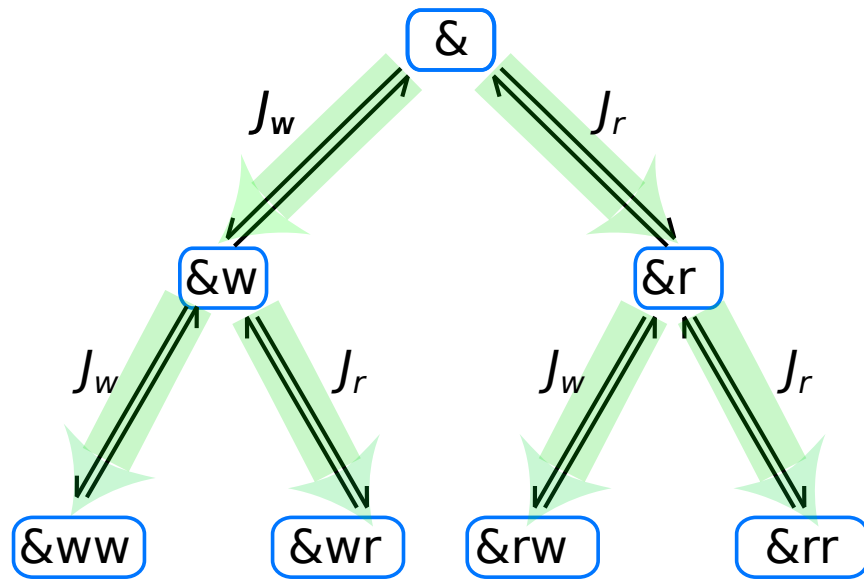
solves the model

-> mean field solution is exact

stochastic simulations

$$\delta = 2 \quad \gamma = 3 \quad \epsilon = 0.5$$

Results



$$\dot{S} = v\Delta S = v[(1 - \eta)\epsilon + \eta(\epsilon - \gamma) + H(\eta)]$$

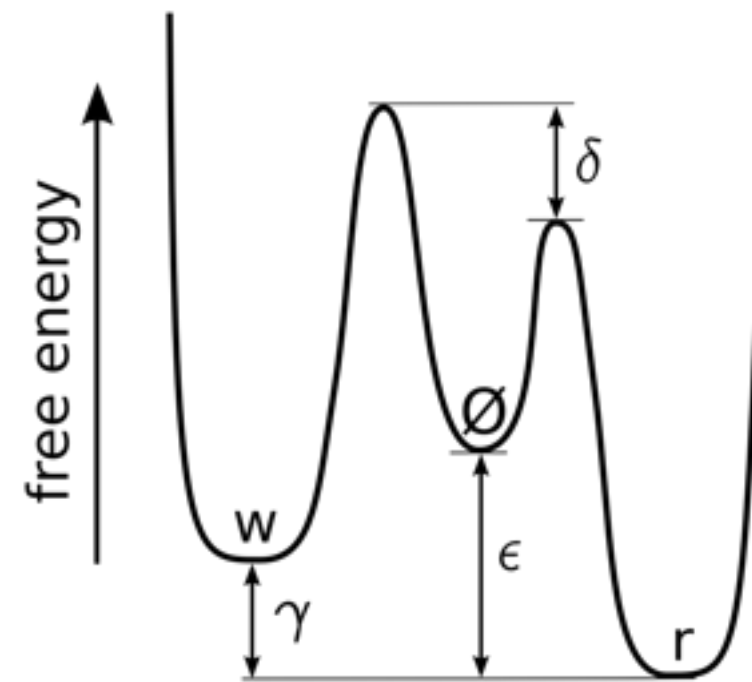
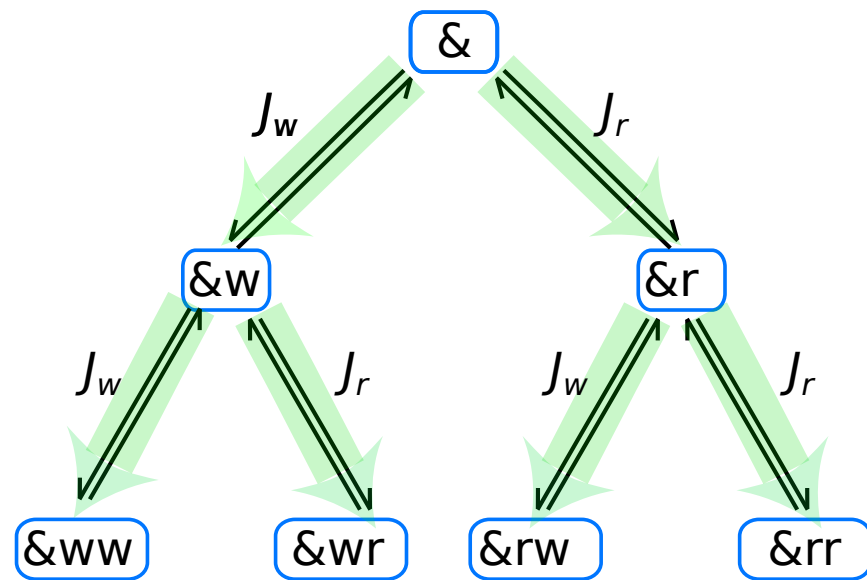
$\dot{S}$ entropy production

ΔS entropy production per copied base

v average copying velocity

$$H(\eta) = -\eta \ln(\eta) - (1 - \eta) \ln(1 - \eta)$$

Results



adiabatic limit:

$$\eta \rightarrow e^{-\gamma}$$

dissipation per step and velocity vanish

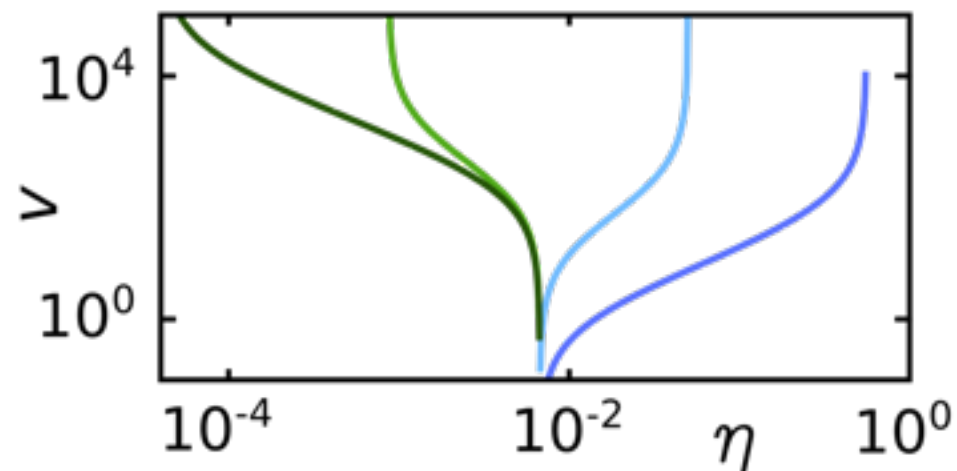
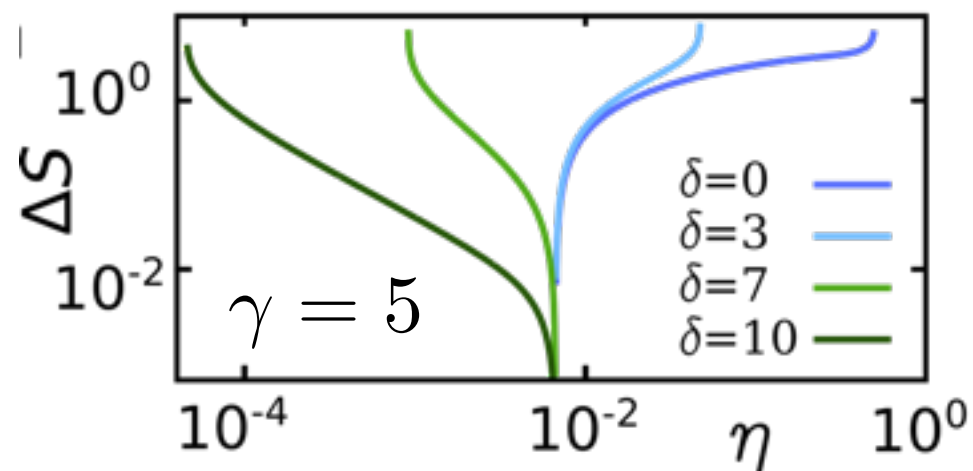
highly dissipative limit:

$$\eta \rightarrow e^{-\delta}$$

dissipation per step and velocity diverge

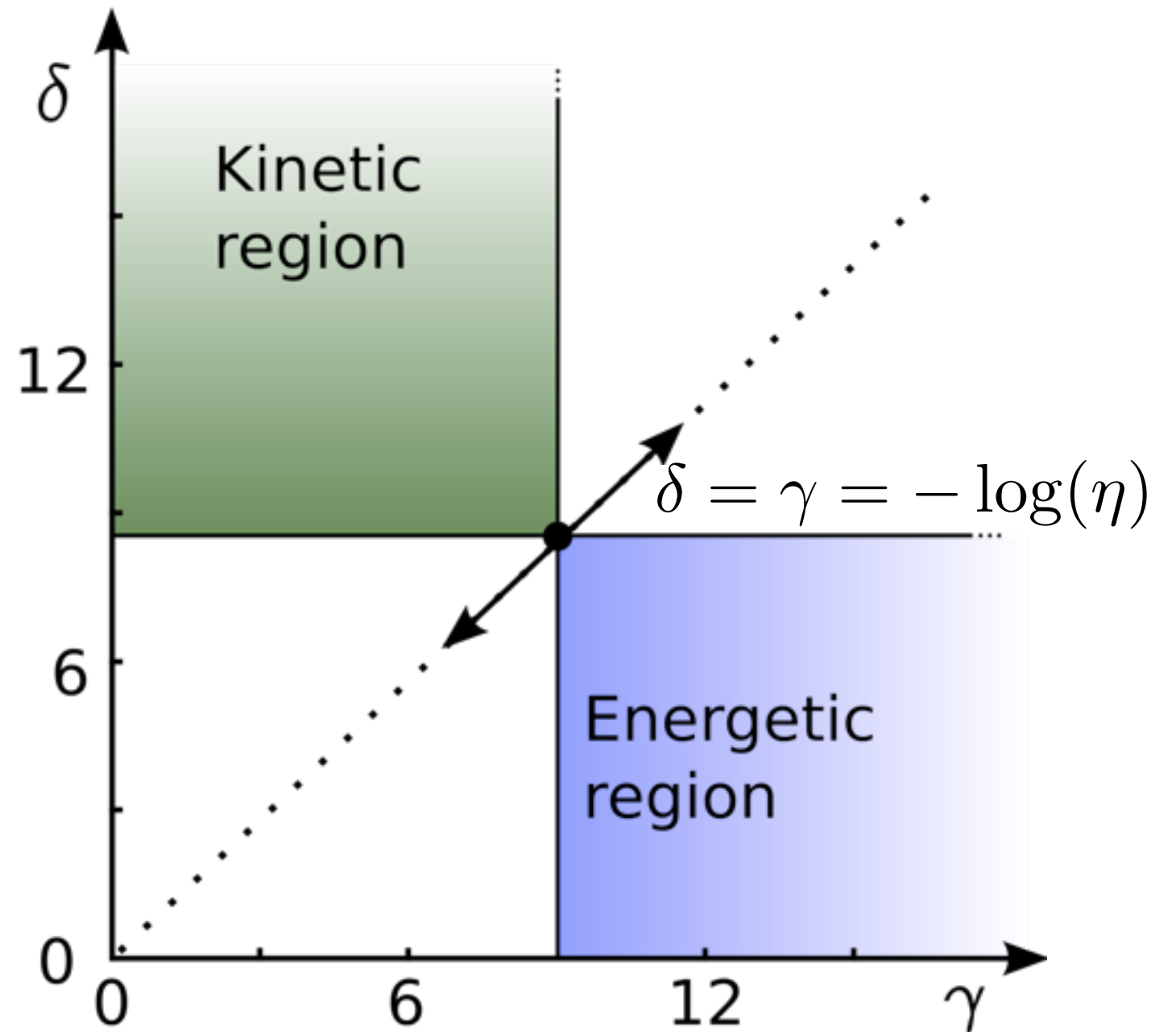
physical region:

$$\min[e^{-\delta}, e^{-\gamma}] < \eta < \max[e^{-\delta}, e^{-\gamma}]$$



Two distinct copying strategies

	Kinetic region	Energetic region
η	↓	↓
ϵ	↑	↓
ΔS	↑	↓
v	↑	↓



$$\min[e^{-\delta}, e^{-\gamma}] < \eta < \max[e^{-\delta}, e^{-\gamma}]$$

different tradeoffs in the two regions

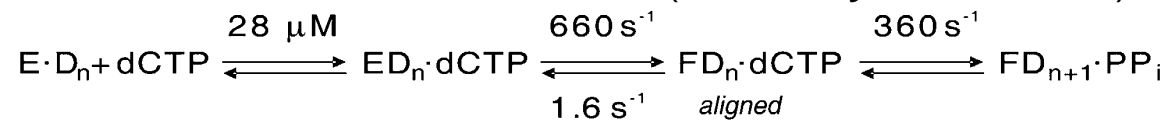
Experimental comparison

Pol γ human DNA polymerase
studied in (Andrieux and Gaspard, 2008)
assuming isoenergetic strains ($\gamma = 0$)
result: $\delta \approx 11$

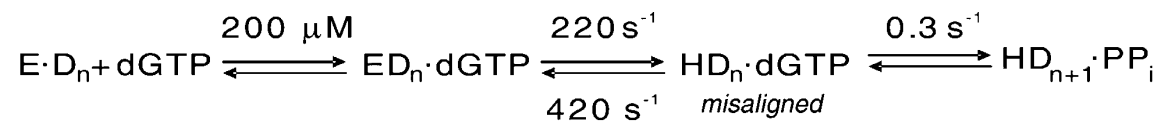
T7 phage DNA polymerase

Correct nucleotide - dCTP

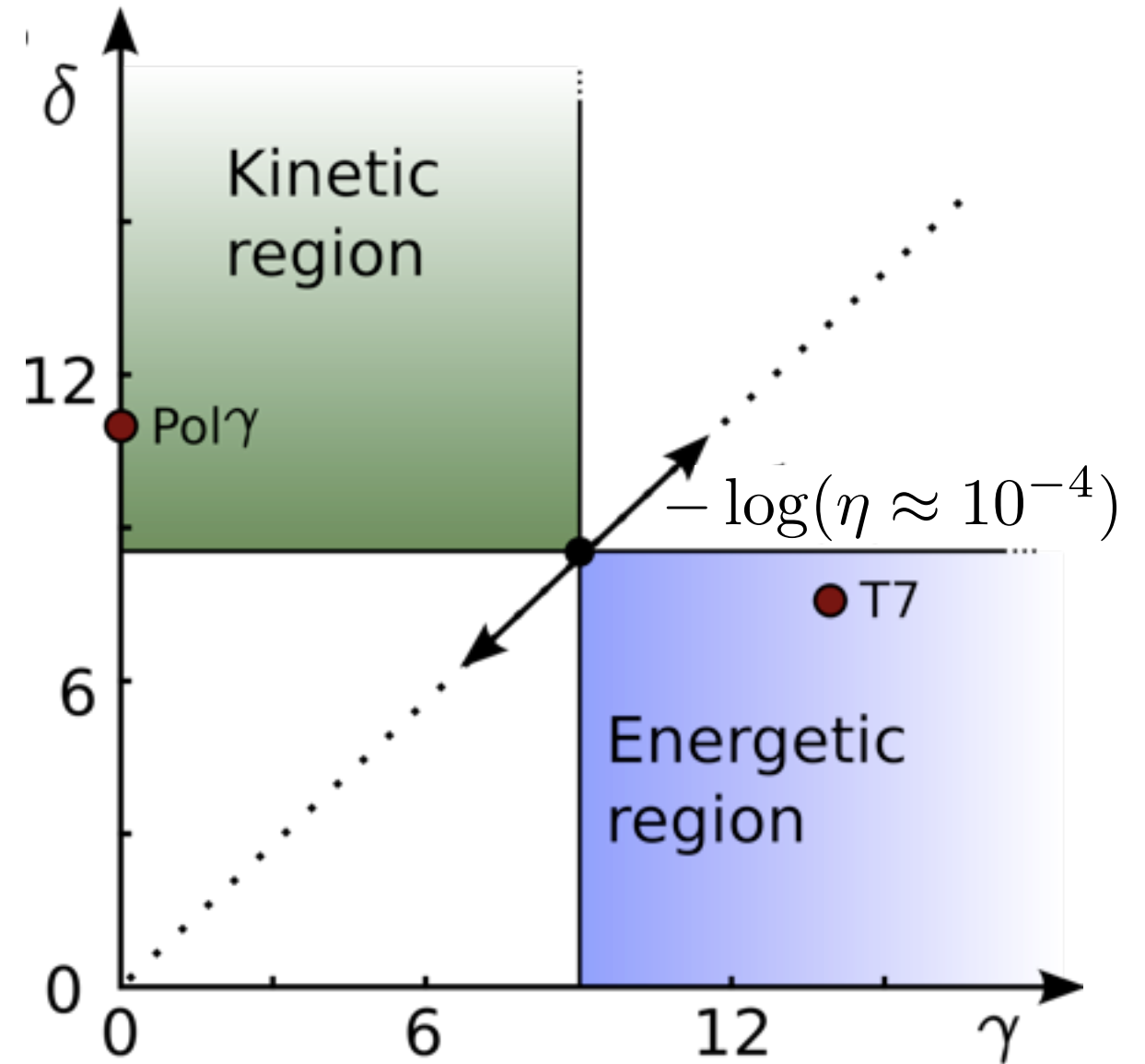
(Tsai and Johnson 2006)



Incorrect nucleotide - dGTP

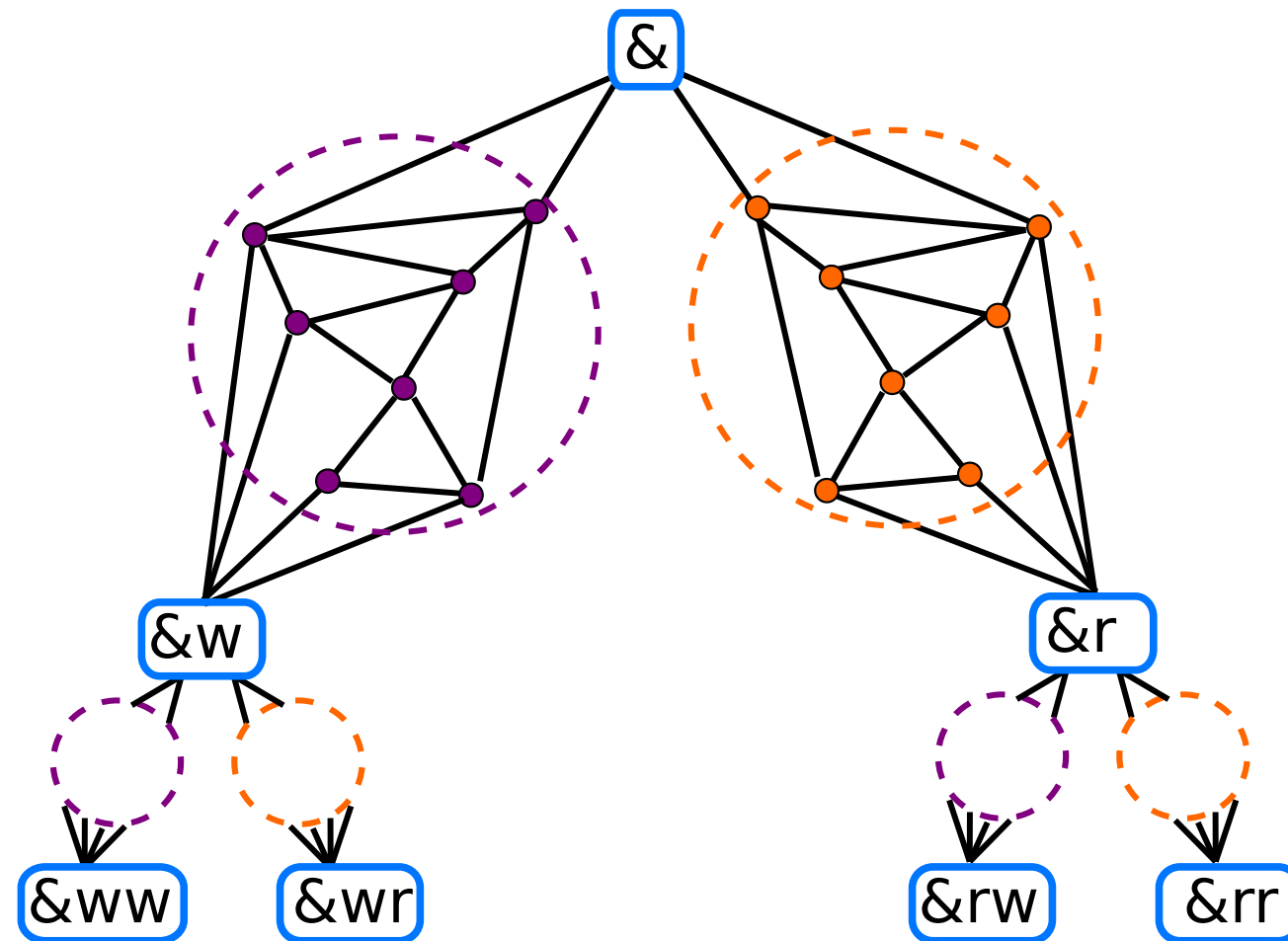


$$\gamma \approx 14 \quad \delta \approx 8$$



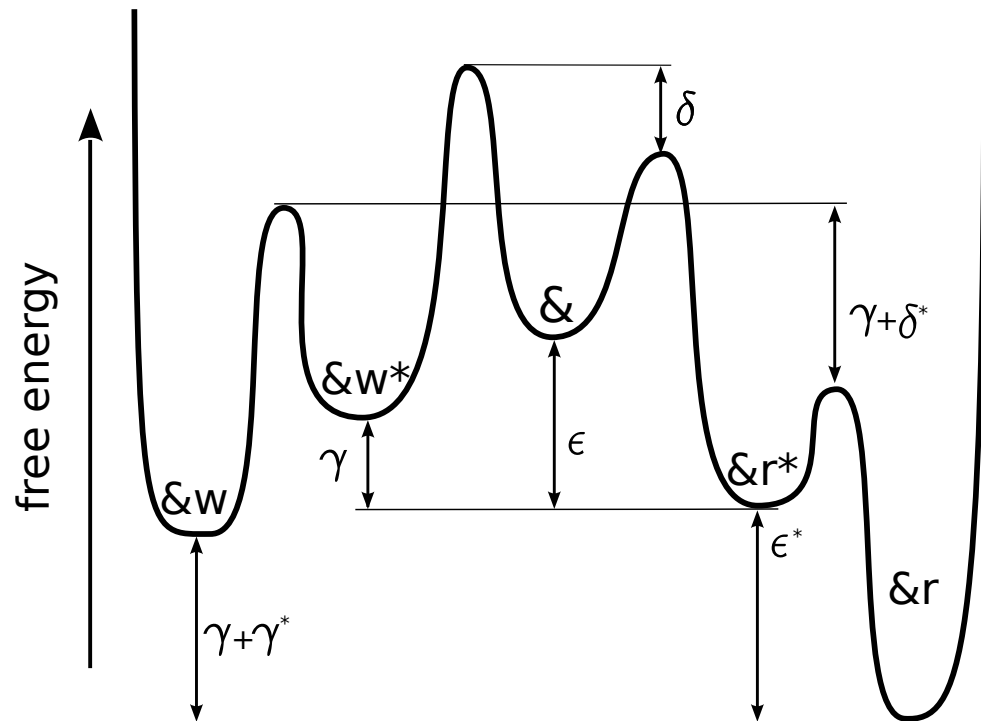
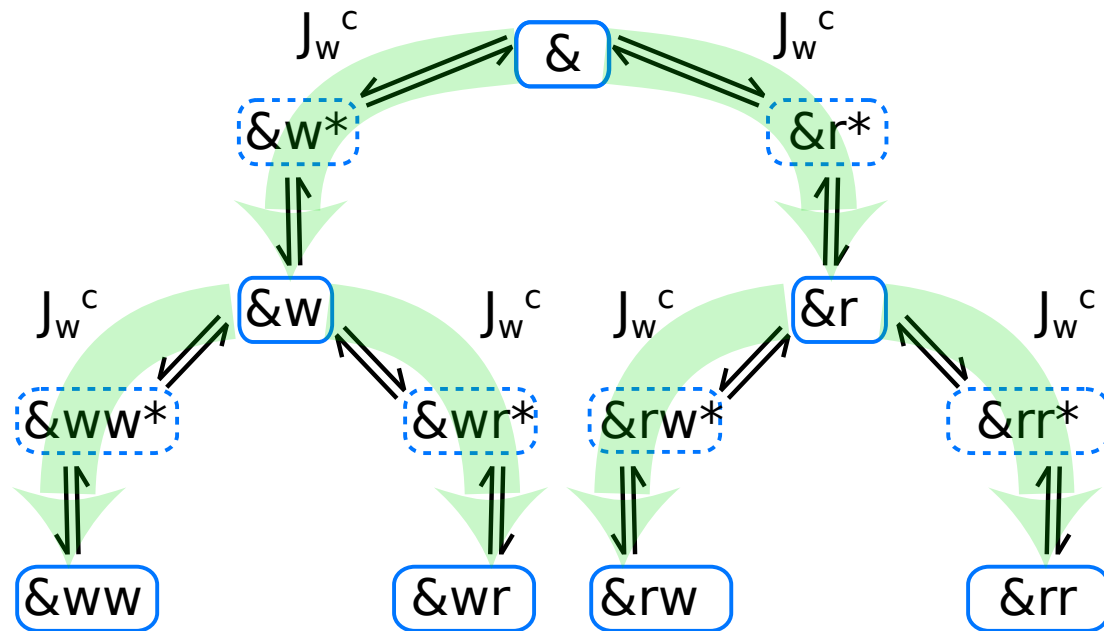
the two copying machines operate in different regions

Intermediate states



- 1) eliminate intermediate states via steady-state condition
- 2) same ansatz as before

Example: two-steps copy

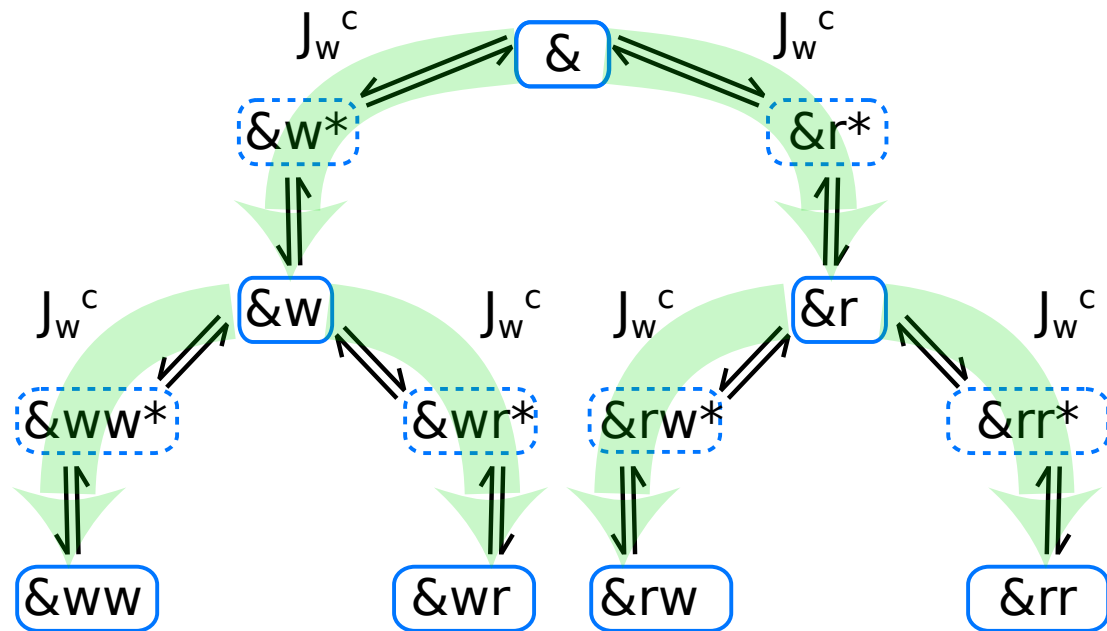


Solution:

$$\frac{\eta}{1 - \eta} = \frac{(e^{\delta} + \omega e^{\bar{\epsilon} + \bar{\delta}})(e^{\epsilon + \bar{\epsilon}} - \eta e^{\gamma + \bar{\gamma}})}{(e^{\gamma} + \omega e^{\bar{\epsilon}})[e^{\epsilon + \delta + \bar{\epsilon} + \bar{\delta}} - (1 - \eta)e^{\delta + \bar{\delta}}]}$$

P. Sartori and SP, in preparation

Example: two-steps copy

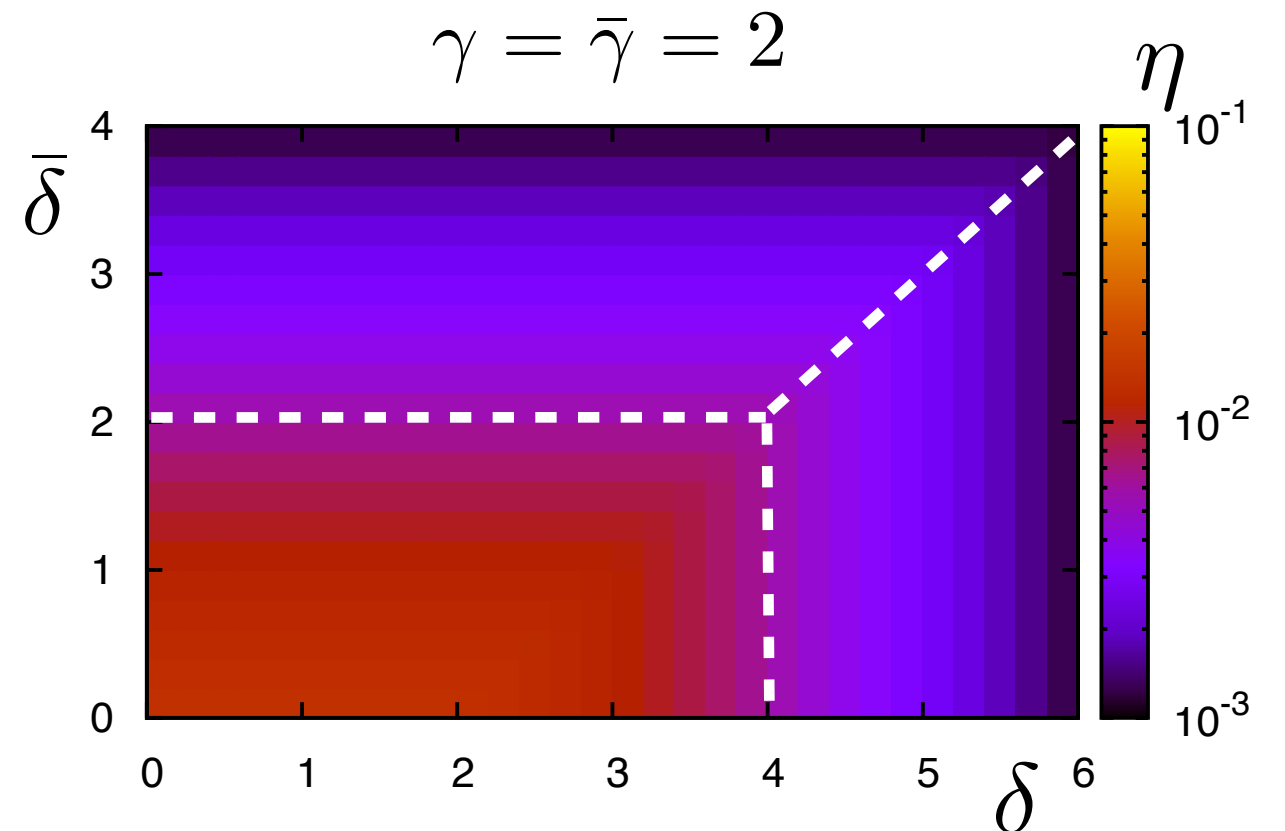
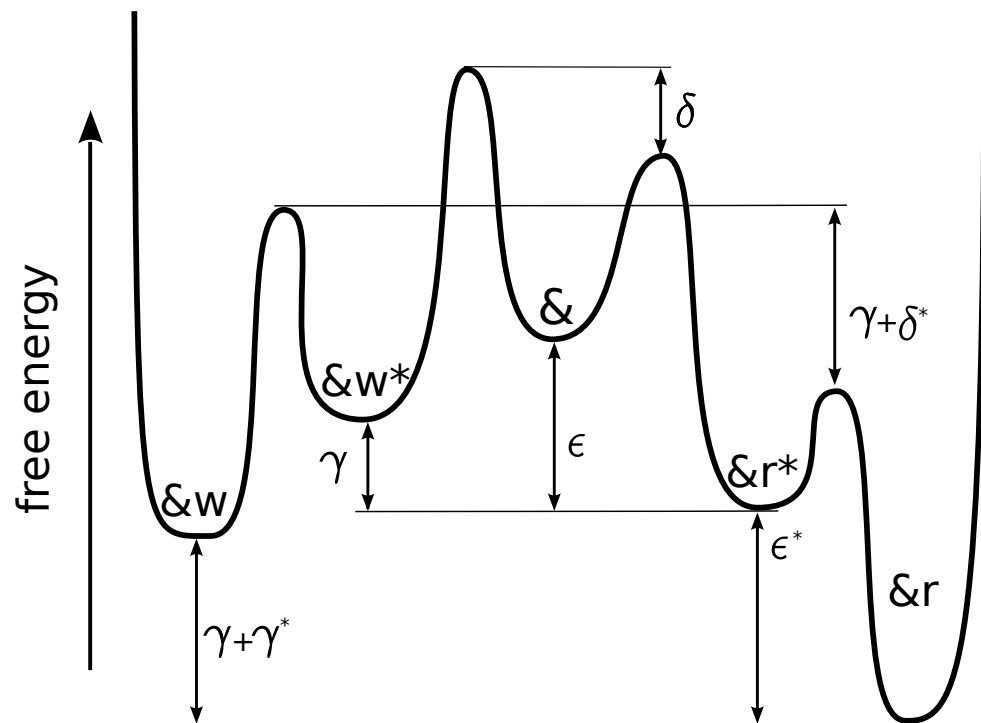


3 regions:

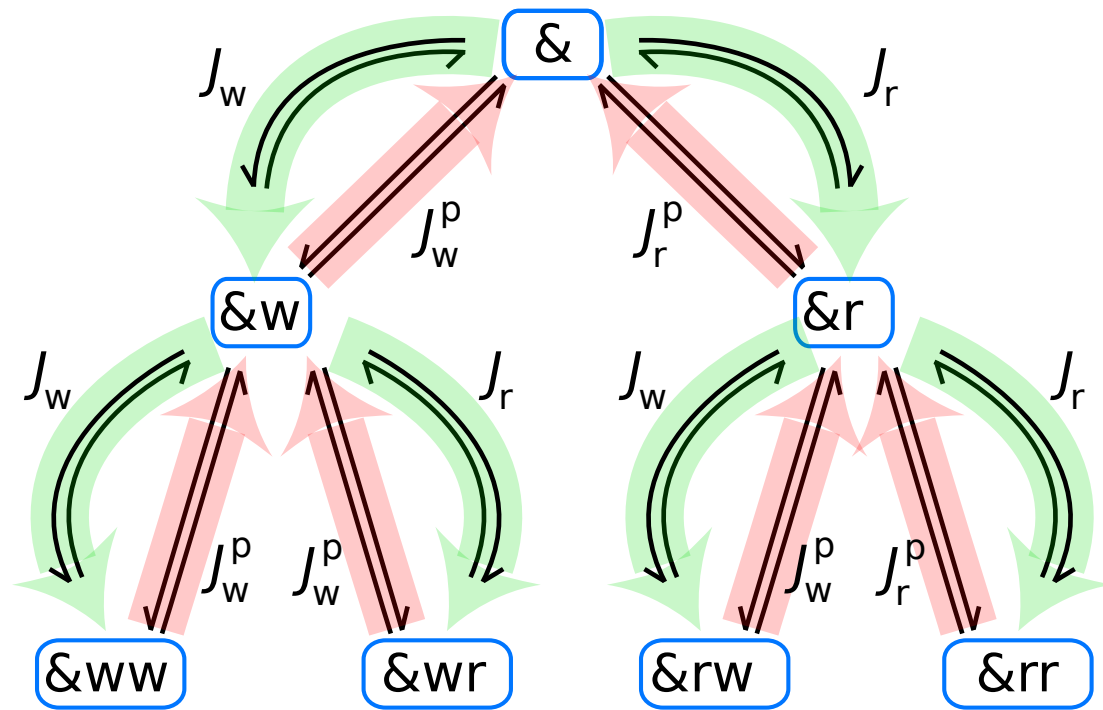
1) adiabatic: $\eta_{min} = e^{-\gamma-\bar{\gamma}}$

2) fast: $\eta_{min} = e^{-\delta}$

3) “forward proofreading”: $\eta_{min} = e^{-\gamma-\bar{\delta}}$

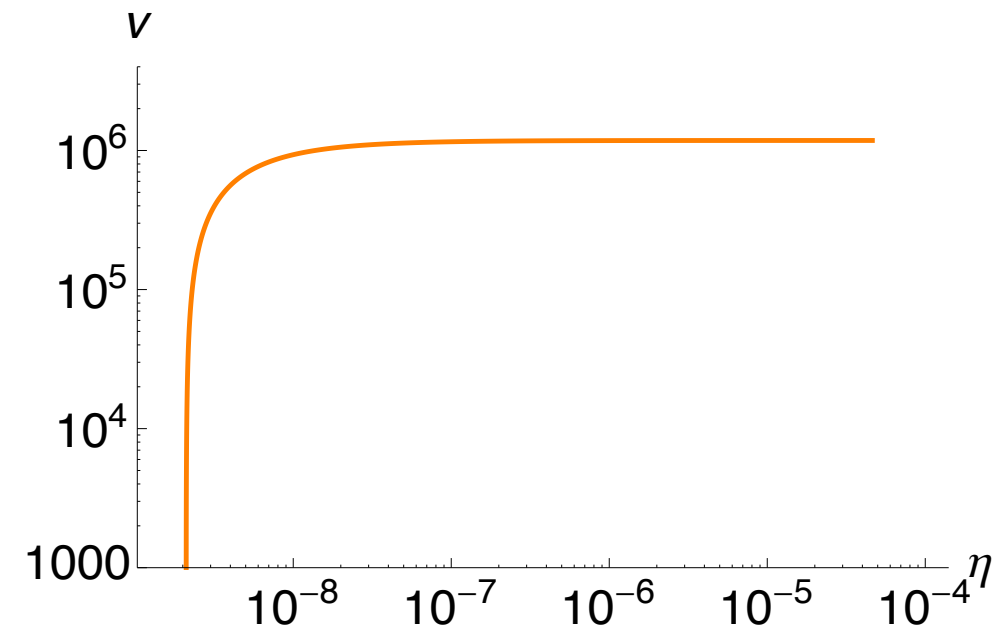
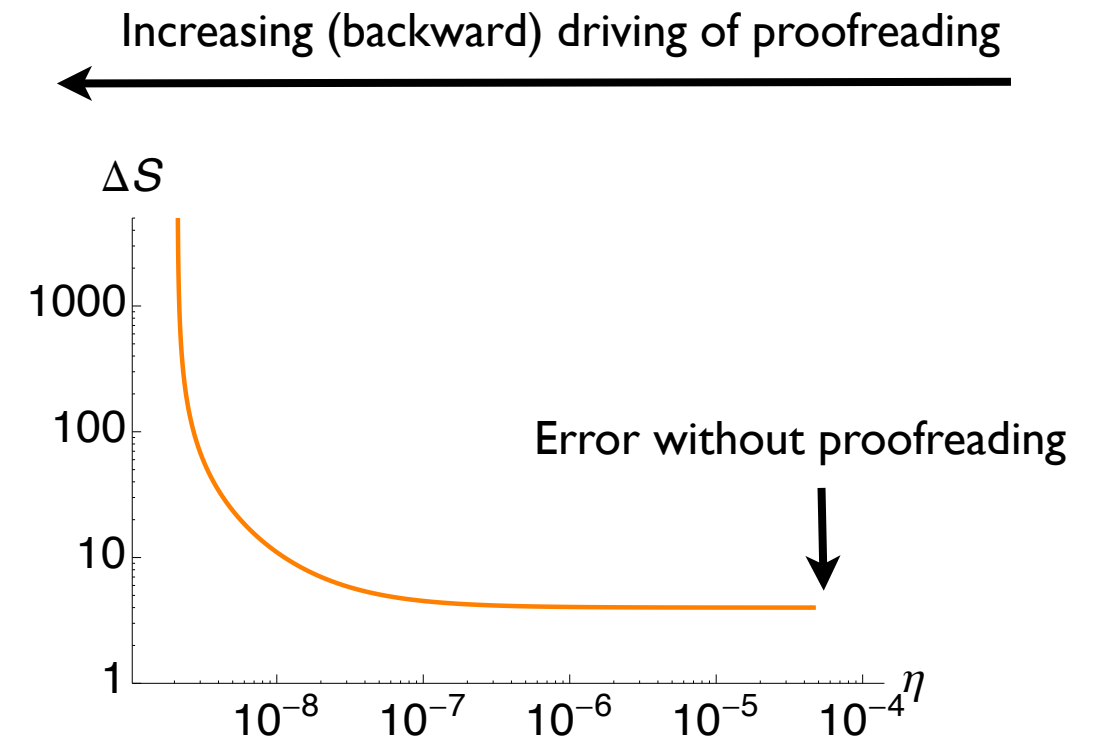


Proofreading



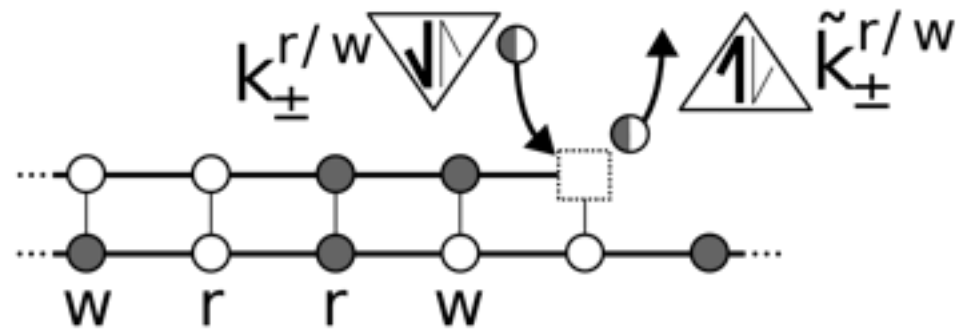
parallel reaction with a backward driving
(preferentially removing wrong monomers)

low error limit of proofreading:
high dissipation per step, vanishing velocity

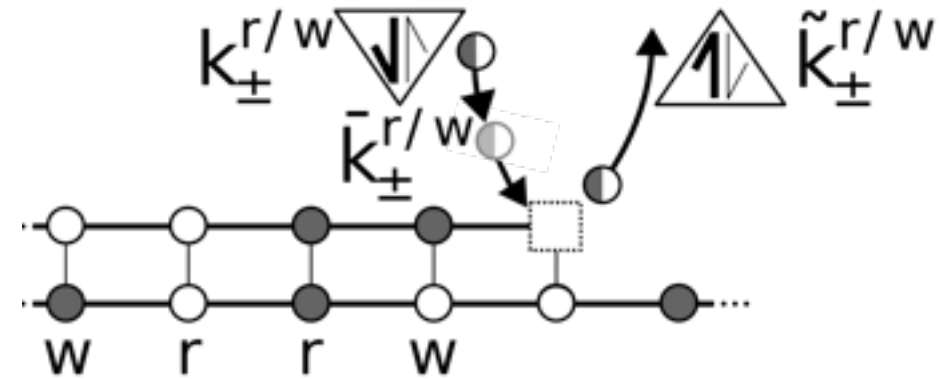


P. Sartori and SP, in preparation

Proofreading



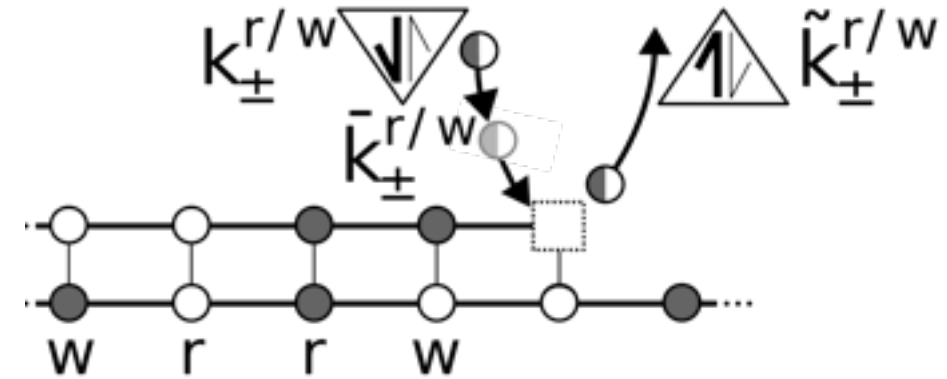
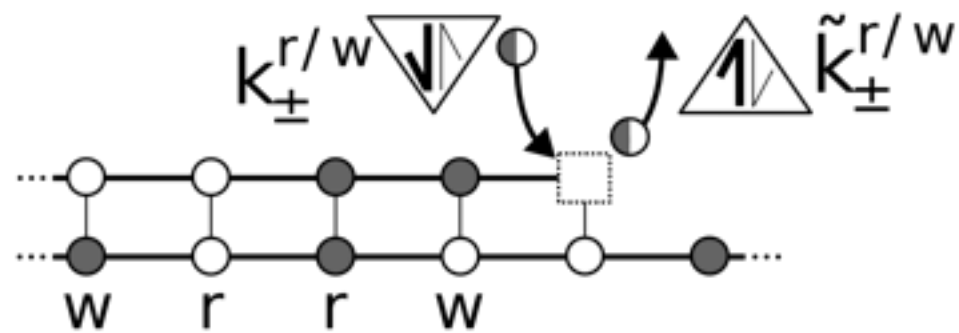
two one-step parallel reactions



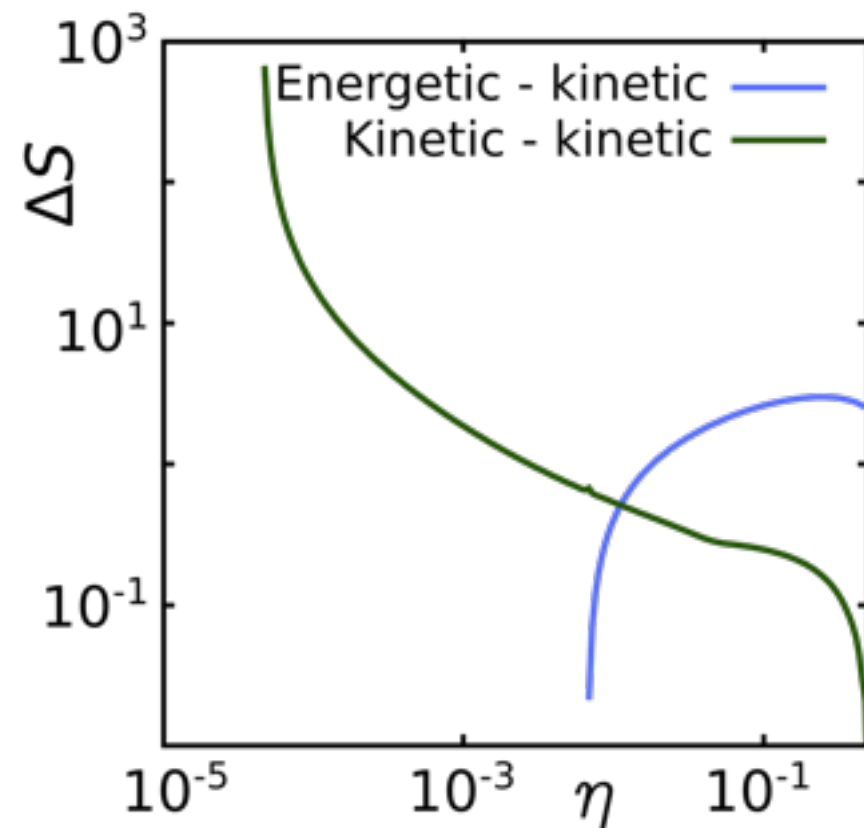
copying with an intermediate step +
proofreading

- the proofreading rates are characterized by δ_p and γ_p
- numerical minimization of the dissipation over the remaining parameters

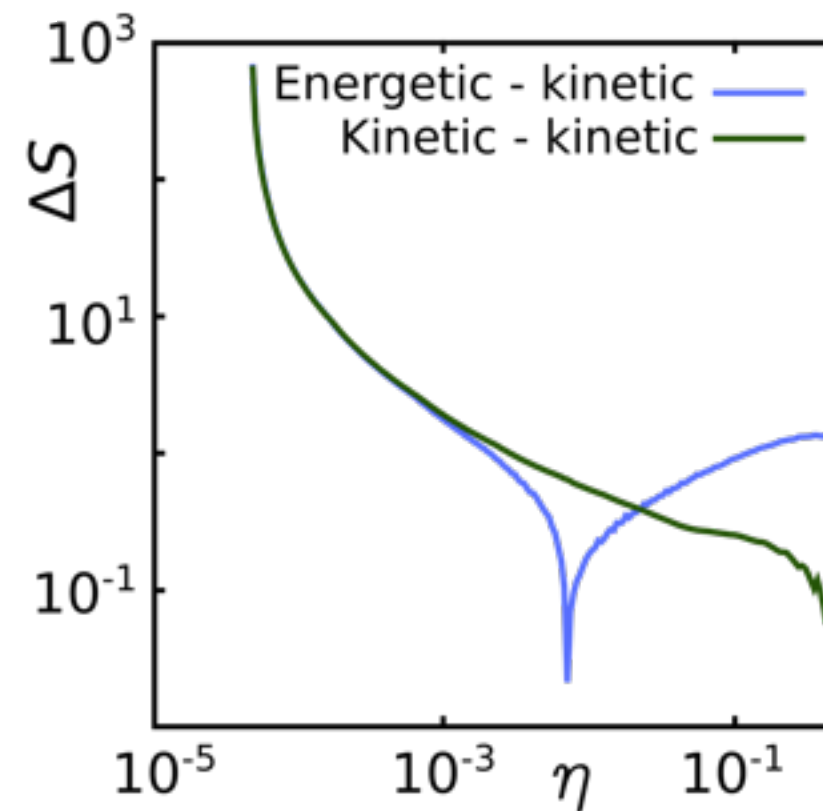
Proofreading models



KP without intermediate state

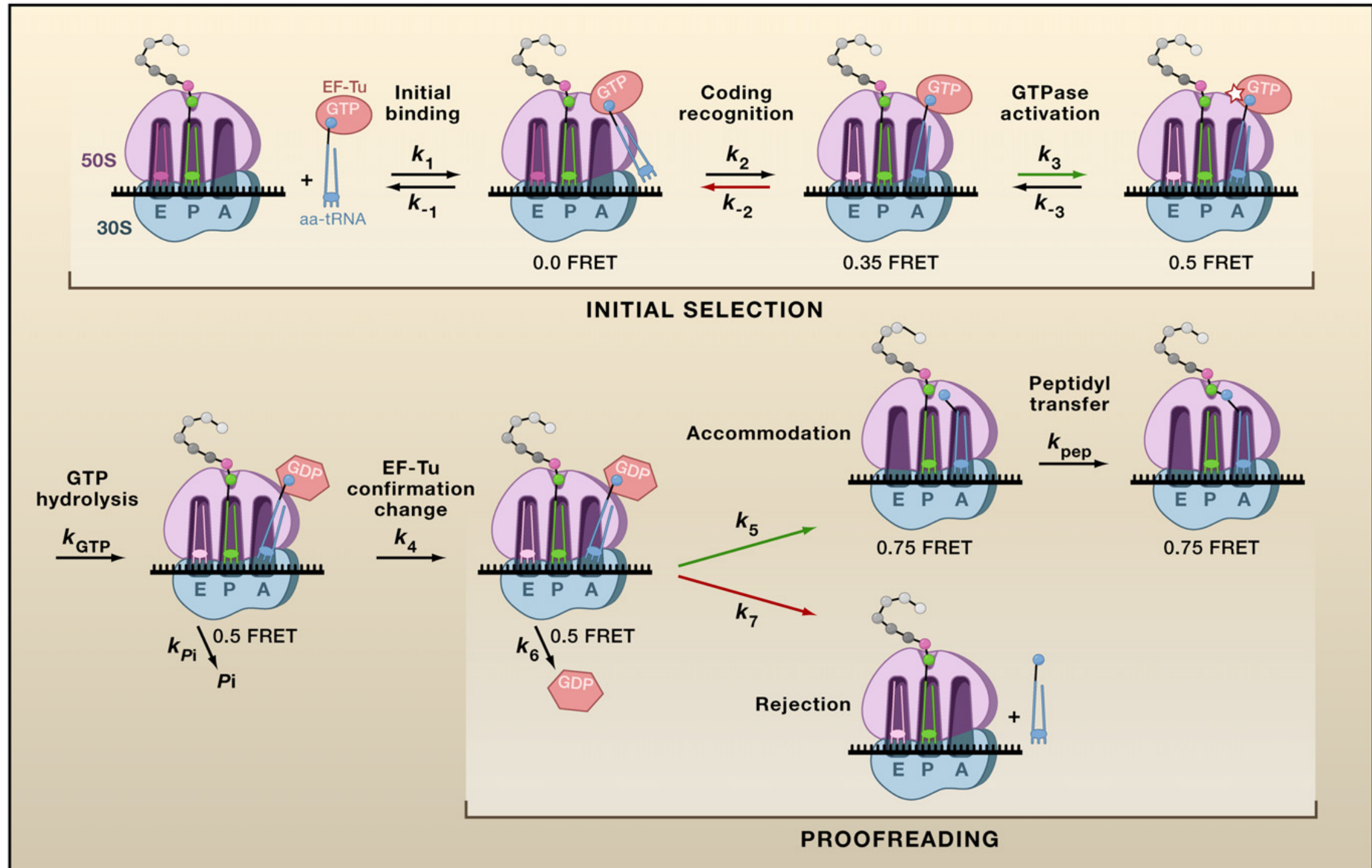


KP with intermediate state



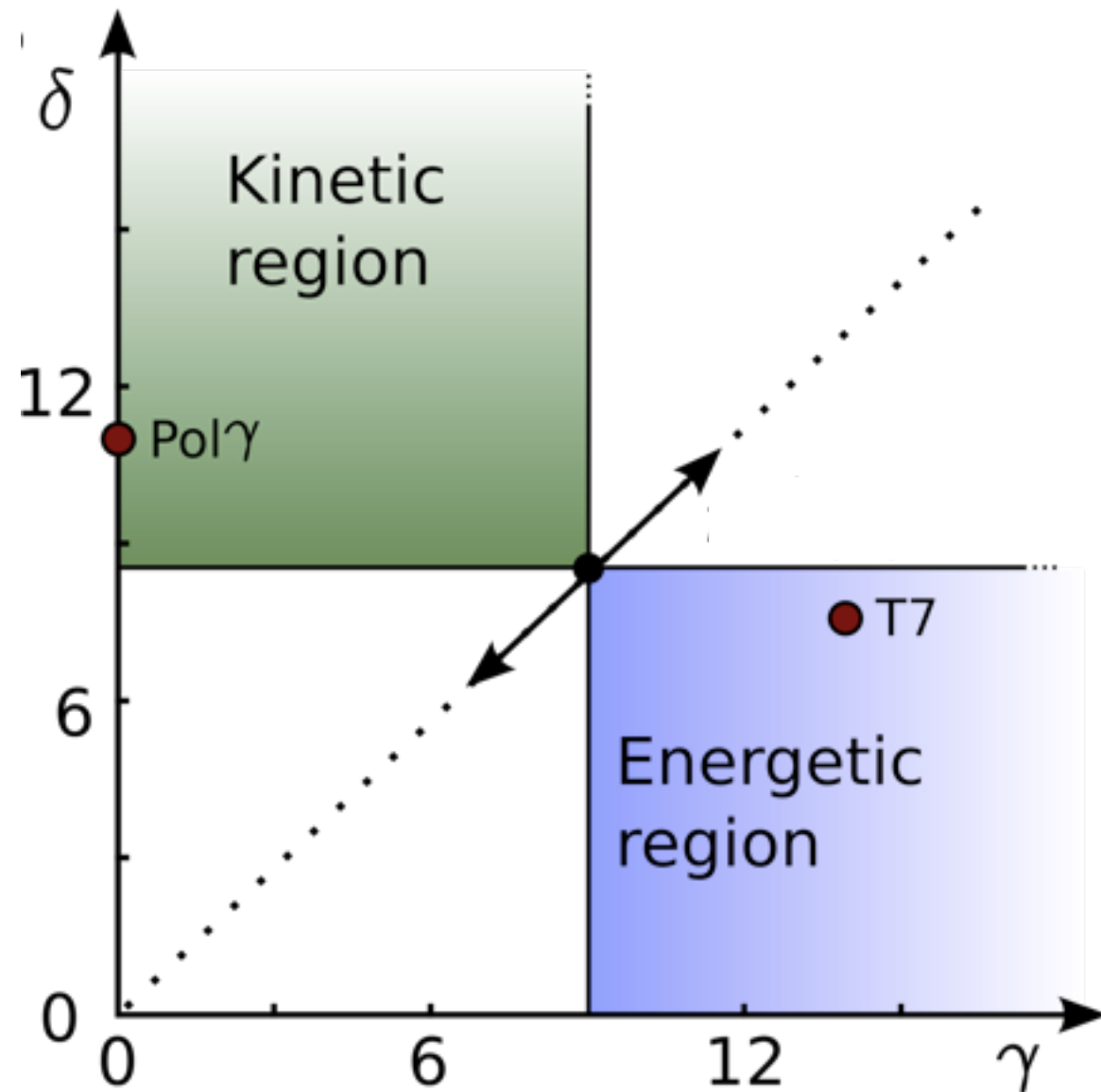
- 1) proofreading is always kinetic
- 2) energetic copy + kinetic proofreading requires intermediate state

tRNA selection pathway



Zaher and Green, Cell (2009)

Conclusions



- There are two different and *separated* regimes in stochastic copying: one based on energy barriers and one based on energy differences
- Reducing the error rate in the two regimes has different costs (more dissipation or less velocity)
- Experimental data suggest that both copying strategies are adopted in biology
- An analysis of more complex schemes shows how the strategies can be combined. Getting closer to realistic models