

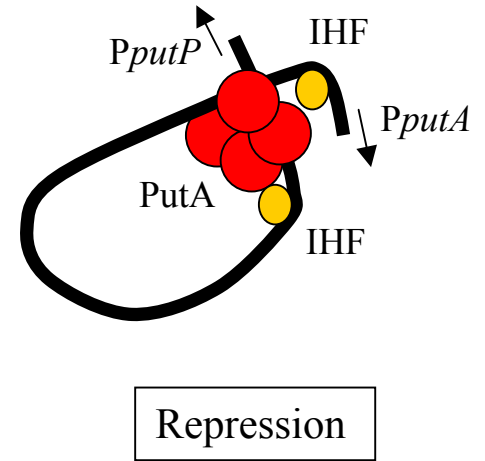
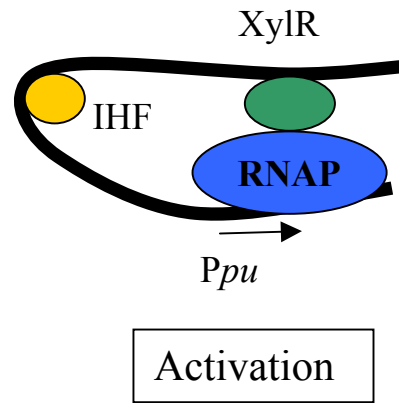
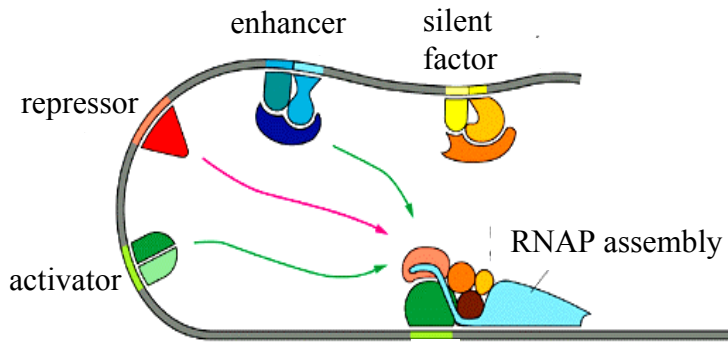
# **Theory of protein induced DNA looping and its role in the control of transcription**

David Swigon

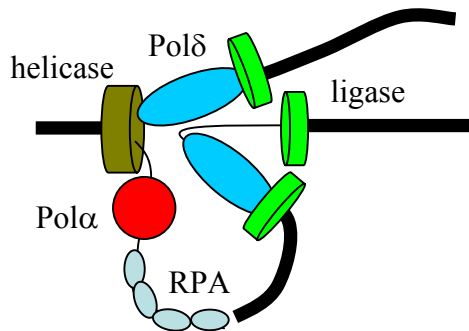
Burroughs Wellcome Fund  
Career Awards at the Scientific Interface  
Final round  
October 2002

The functionality of molecular biological processes is determined the mechanics and dynamics of complex protein-DNA assemblies.

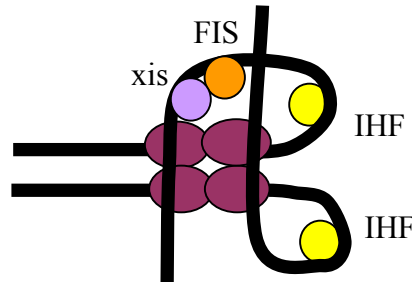
### Transcription



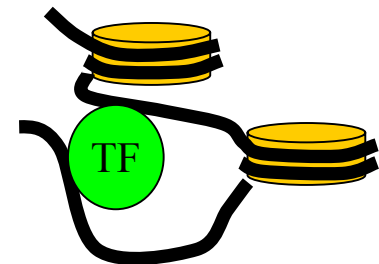
### Replication



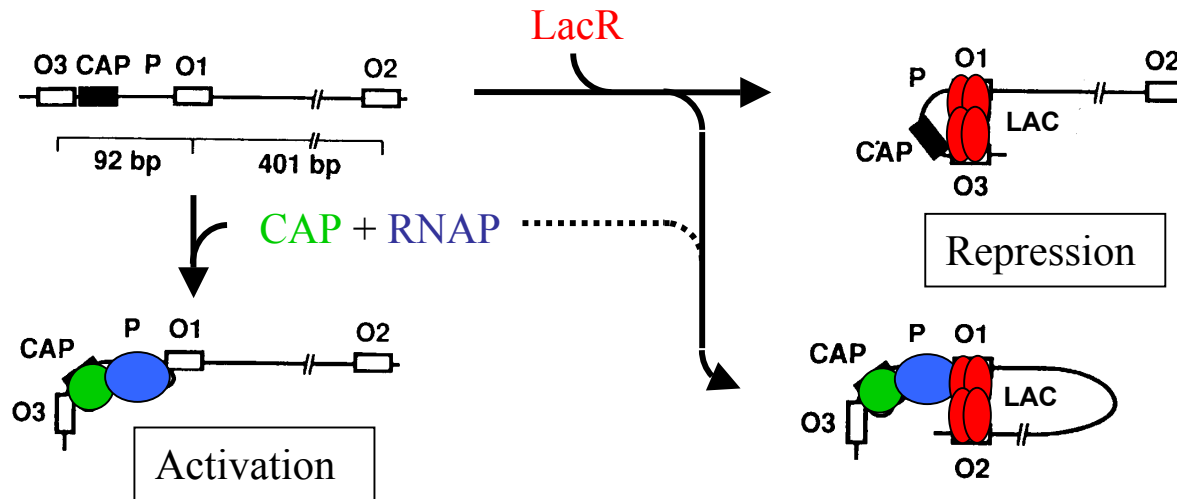
### Recombination



### Chromatin remodeling



# Example: Regulation of the *Lac* operon



Lac repressor protein (LacR) binds to O1 and one of O2 and O3.

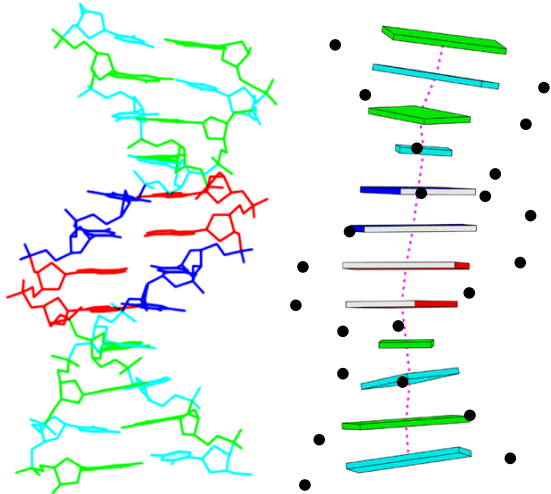
Absence of O2 and O3 decreases repression by 40 fold. [Oehler et al., *EMBO J.*, 1994]

Distance between O3/O2 and O1 affects repression 1-30 fold. [Müller et al., *J. Mol. Biol.*, 1996]

**The presence of multiple LacR binding sites combined with mechanical deformation of DNA and LacR gives rise to a highly efficient and controllable repressive mechanism.**

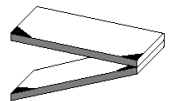
# Modeling of the *Lac* operon

DNA



Atoms

Base pairs



Tilt  $\theta_1$



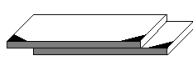
Roll  $\theta_2$



Twist  $\theta_3$



Shift  $\rho_1$



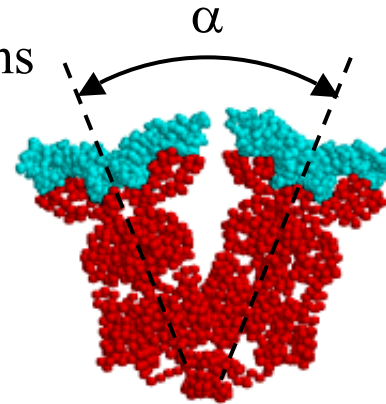
Slide  $\rho_2$



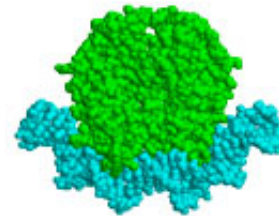
Rise  $\rho_3$

$$\Psi = \sum_{n=1}^N \psi^n(\theta_i^n, \rho_j^n) + \sum_{i < j} \frac{\delta_i \delta_j e^{-\kappa r_{ij}}}{4\pi\epsilon r_{ij}}$$

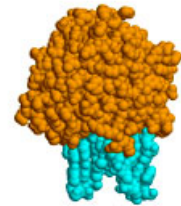
Proteins



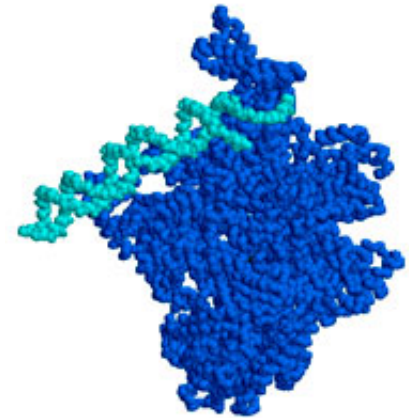
LacR (pdr026)



CAP (pdr023)



DNaseI (pde005)



RNAP (pd0305)

## Computational techniques developed to date:

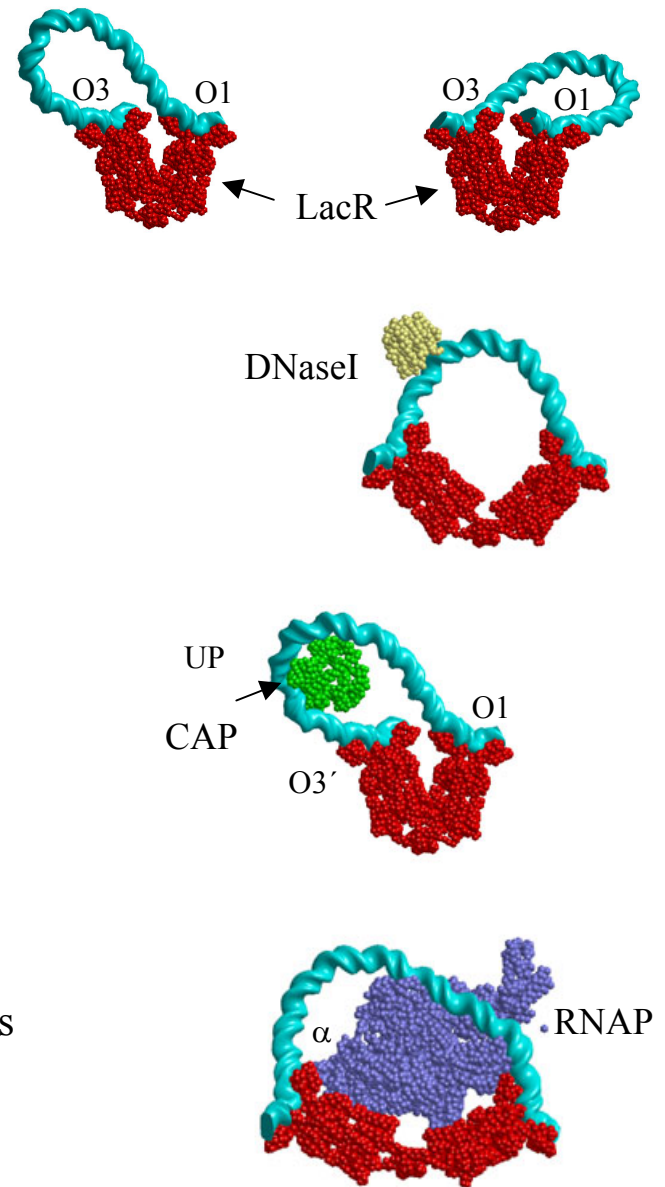
- calculation of DNA equilibrium configurations  
[Coleman, Olson, & Swigon, *J. Chem. Phys.*, to appear April 2003]
- MC algorithm for closure probability & free energy
- large-scale protein deformability, normal modes

## New findings about the structure and dynamics of LacR-CAP-DNA assembly:

- DNase I footprinting reveals LacR extended in short loops
- Loop enhances repression by inhibiting CAP binding
- CAP binding allowed if O3 site is shifted upstream to O3'
- Cooperative effect between RNAP and LacR binding

## Work in progress:

- Dynamics of loop formation
- Kinetic model of Lac repression and CAP activation
- Modeling of prokaryotic operons Ara, NtrC, Put
- Experimental verification of structural results and mechanisms
- Design of transcription control elements



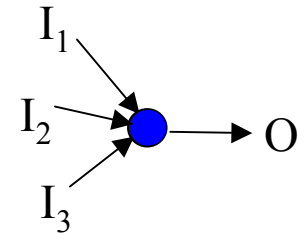
# Future Directions

Use mechanical principles to bridge the gap between structural biology, macromolecular assembly, and biological function.

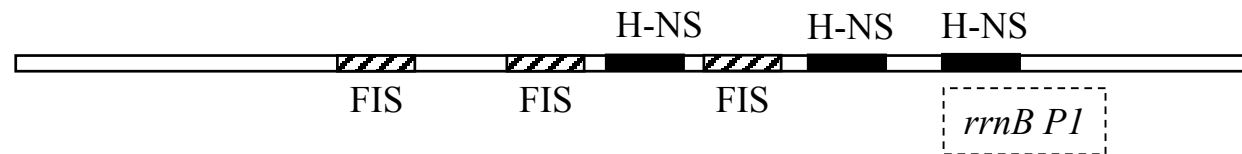
**Molecular biomechanics:** investigate the mechanics of transcription, replication, recombination; study the action of enzymes and molecular motors

**Genetic networks:** incorporate transcription-assembly mechanics in modeling the dynamics of genetic networks

$$[O] = f([I_1], [I_2], [I_3])$$



**Bioinformatics:** apply conformational principles to the identification of promoters and transcription factor binding sites



**Pharmaceutics:** design drugs that alter DNA mechanics; apply to transcription regulation in gene therapy and development of antimicrobial agents