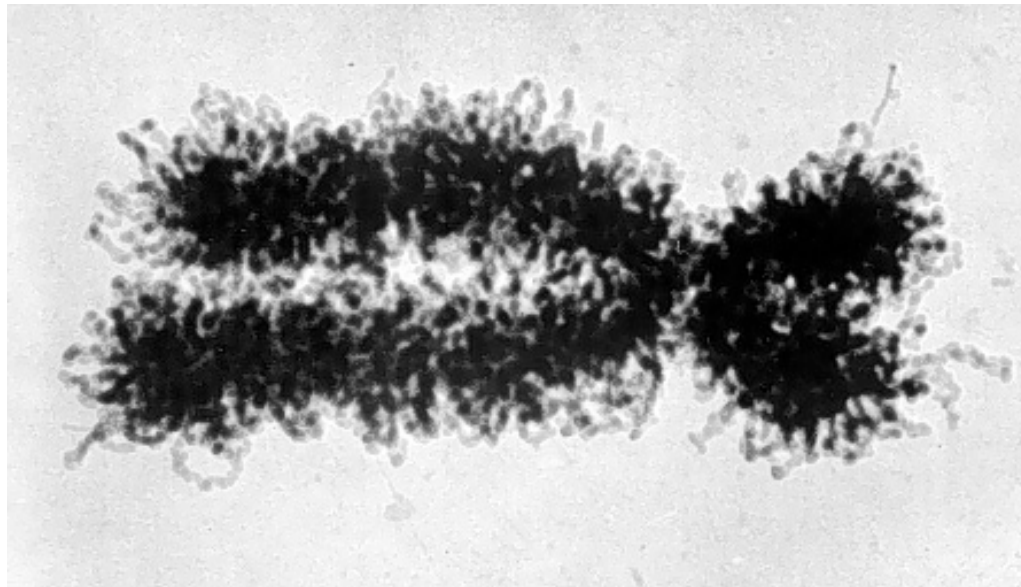


DNA Flexibility

Electron micrograph of a metaphase chromosome

•cm's of DNA

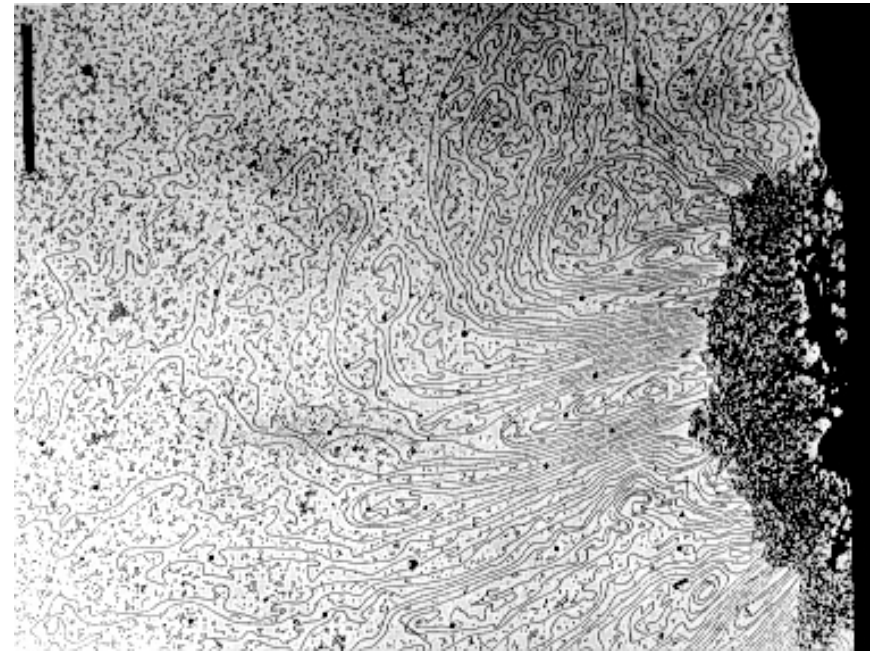
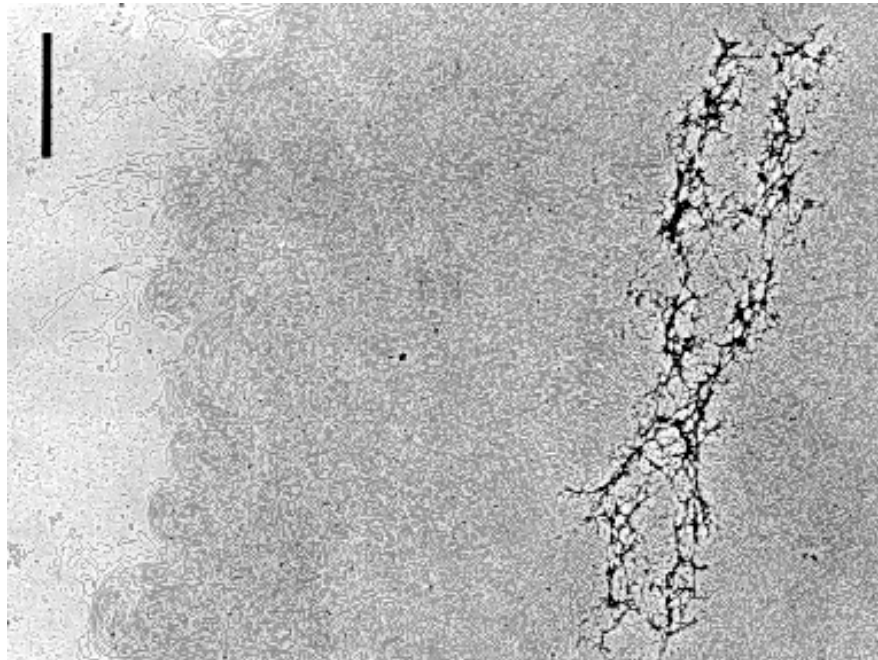


] Few hundred
nm's

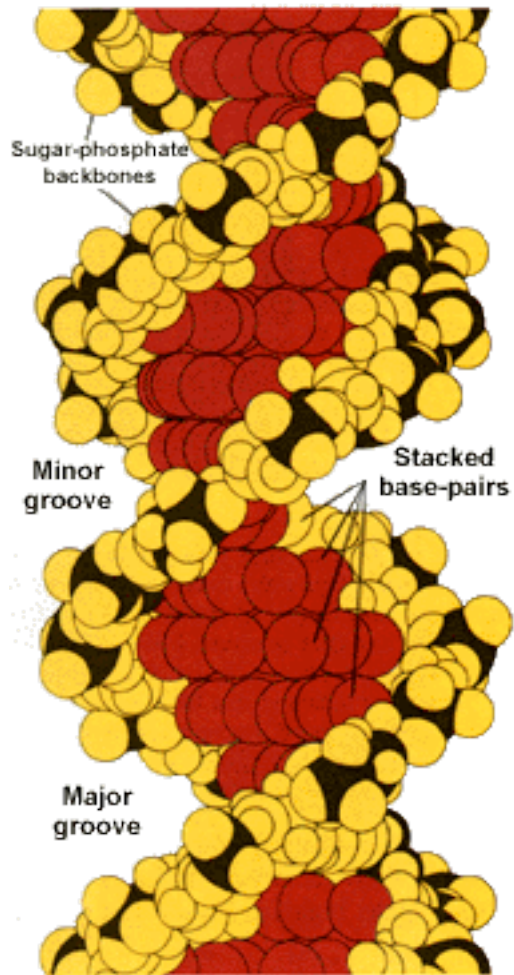


few μ 's

Metaphase chromosome, after removal of histones

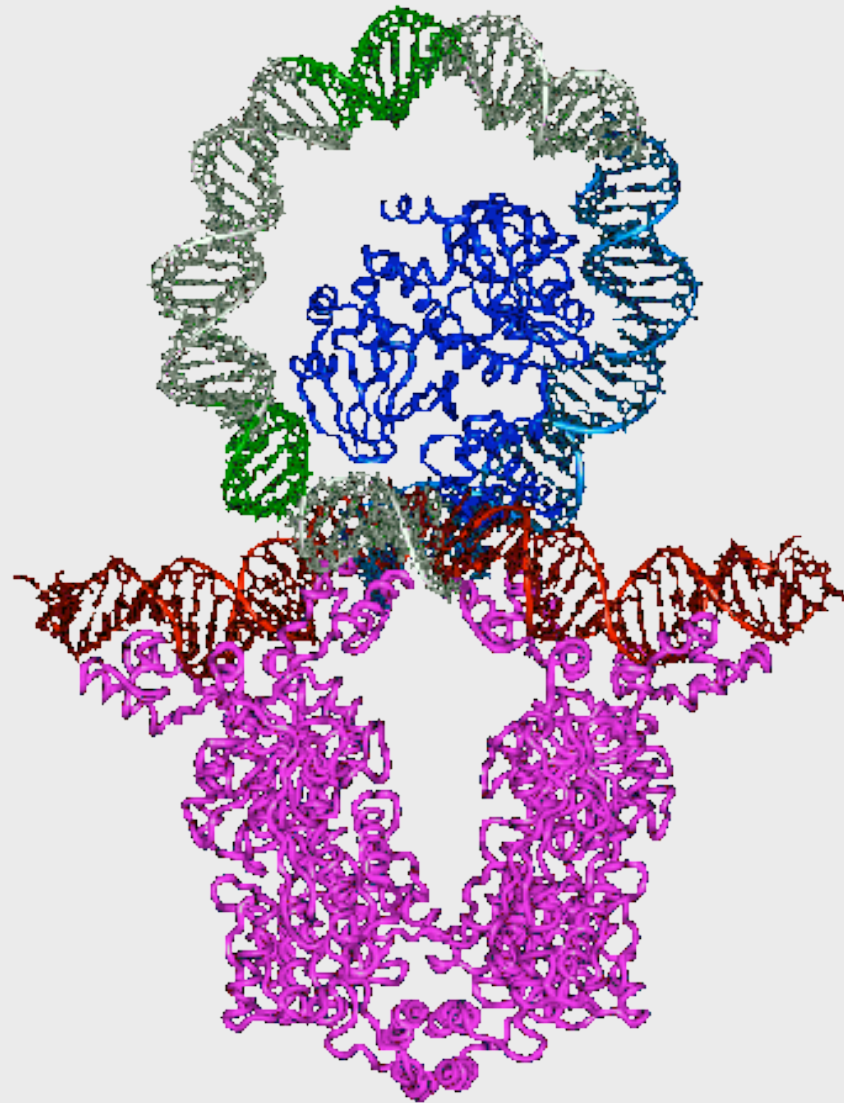


DNA is a stiff polymer

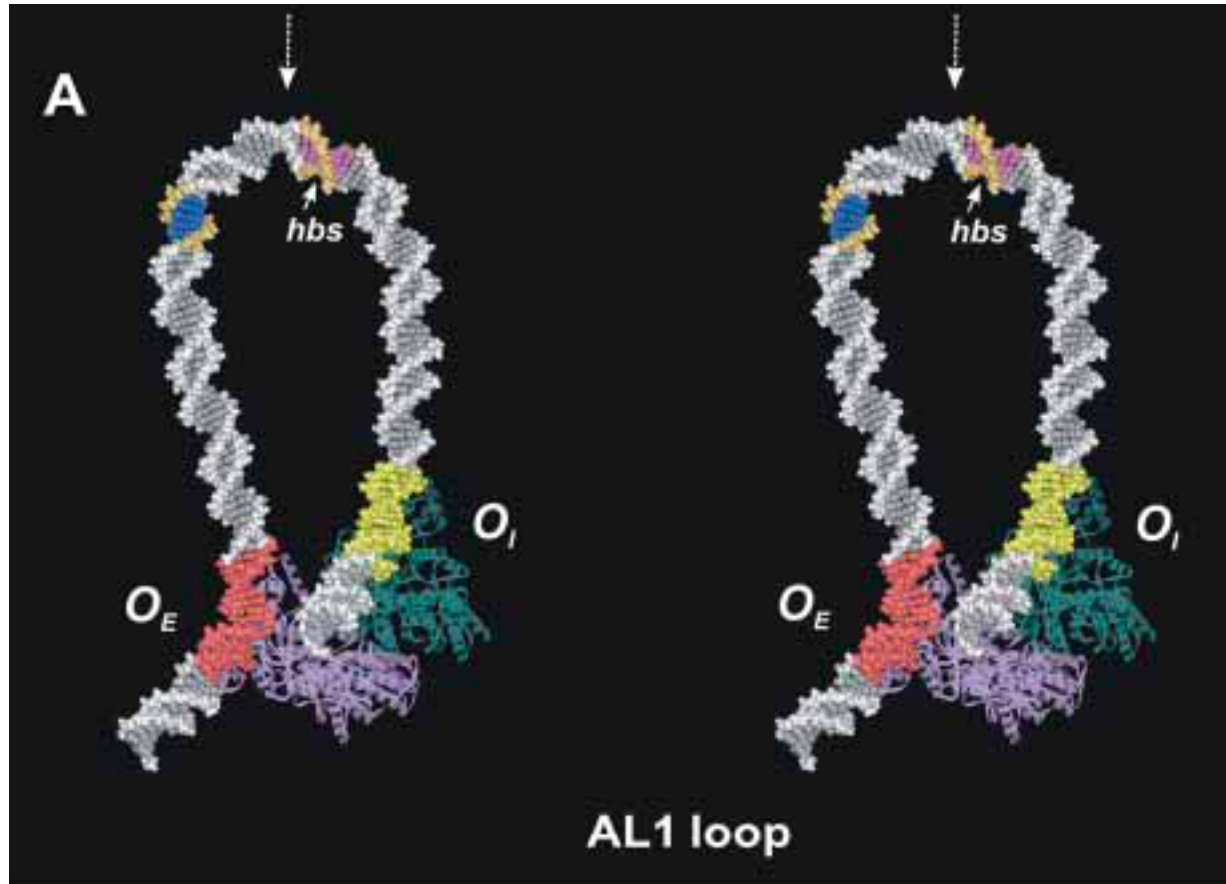


- Phosphate-phosphate repulsion
- Hard sphere repulsions of bases

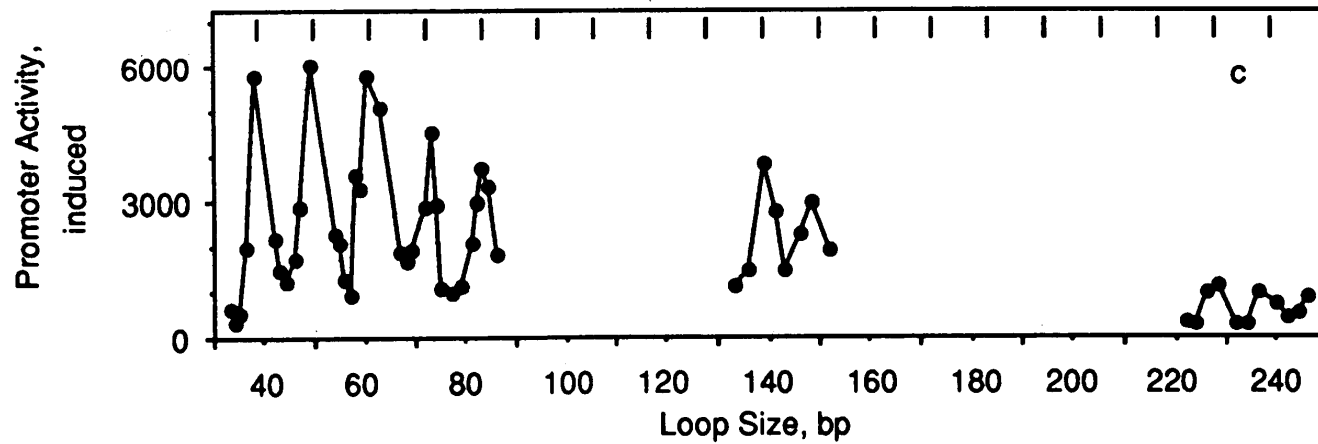
Sharply looped DNA in the *lac* operon



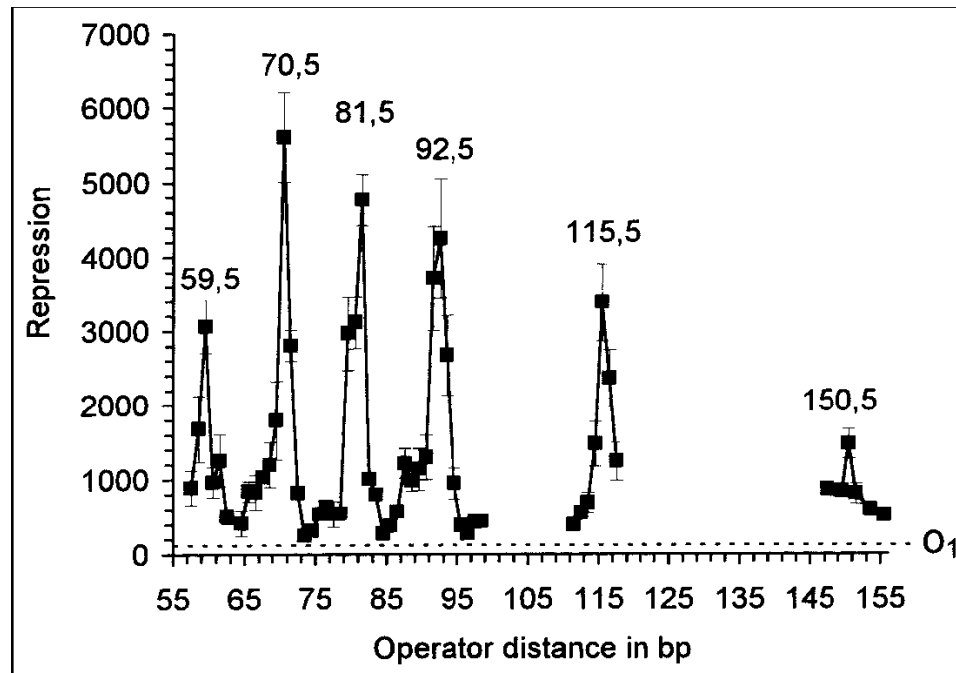
Sharply looped DNA in the Gal repressosome



Sharply looped and *twisted* DNA *in vivo*



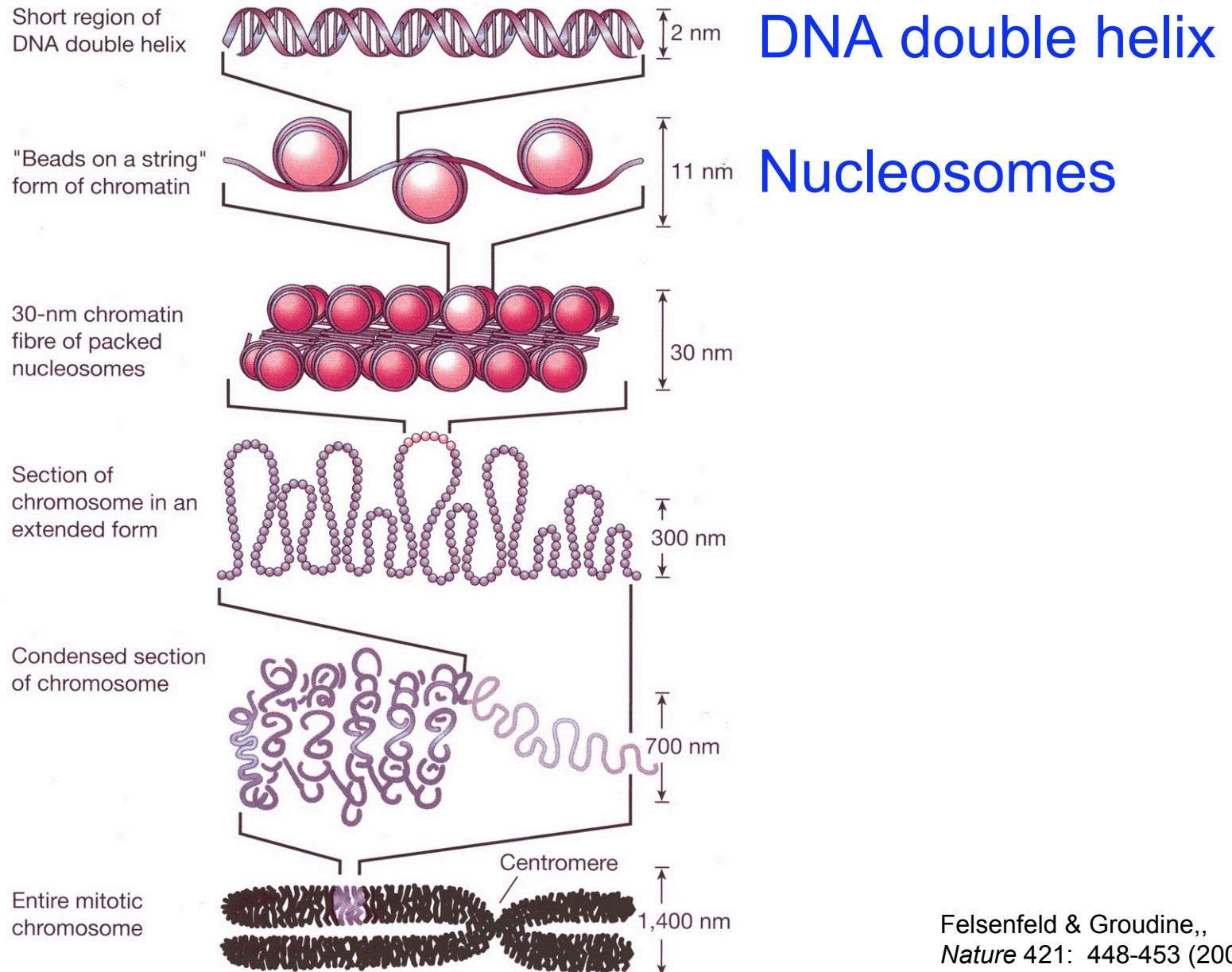
araCBAD



lac

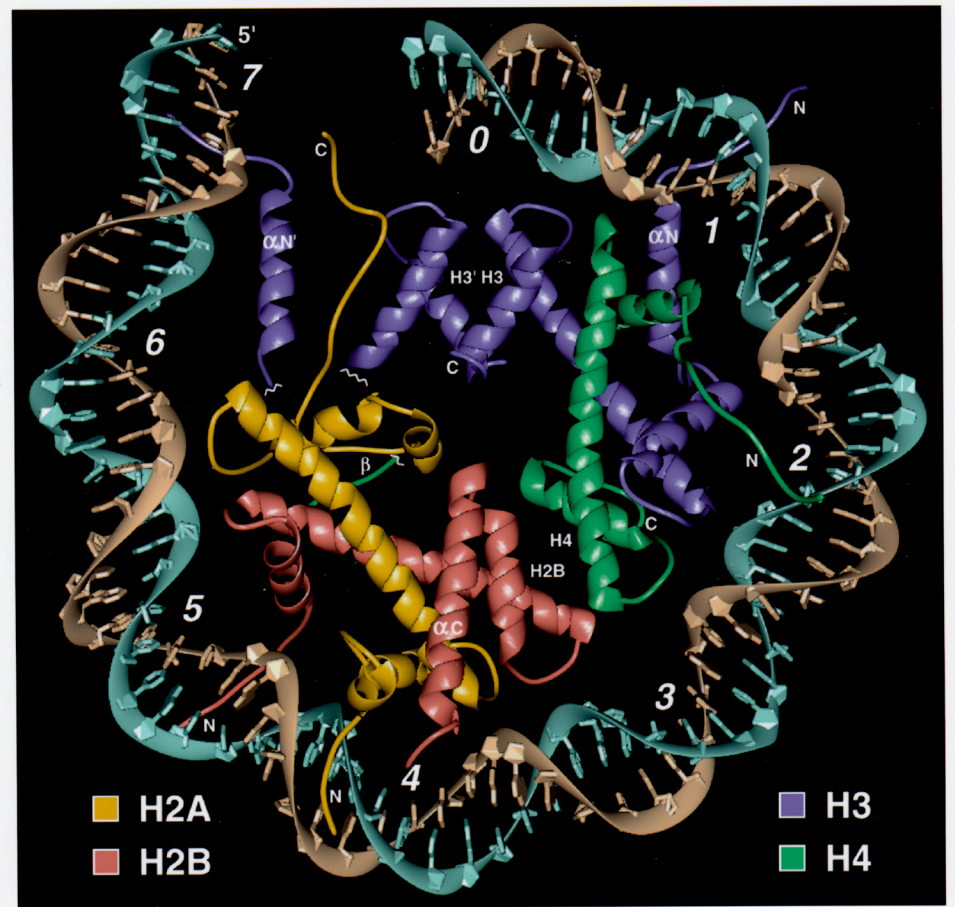
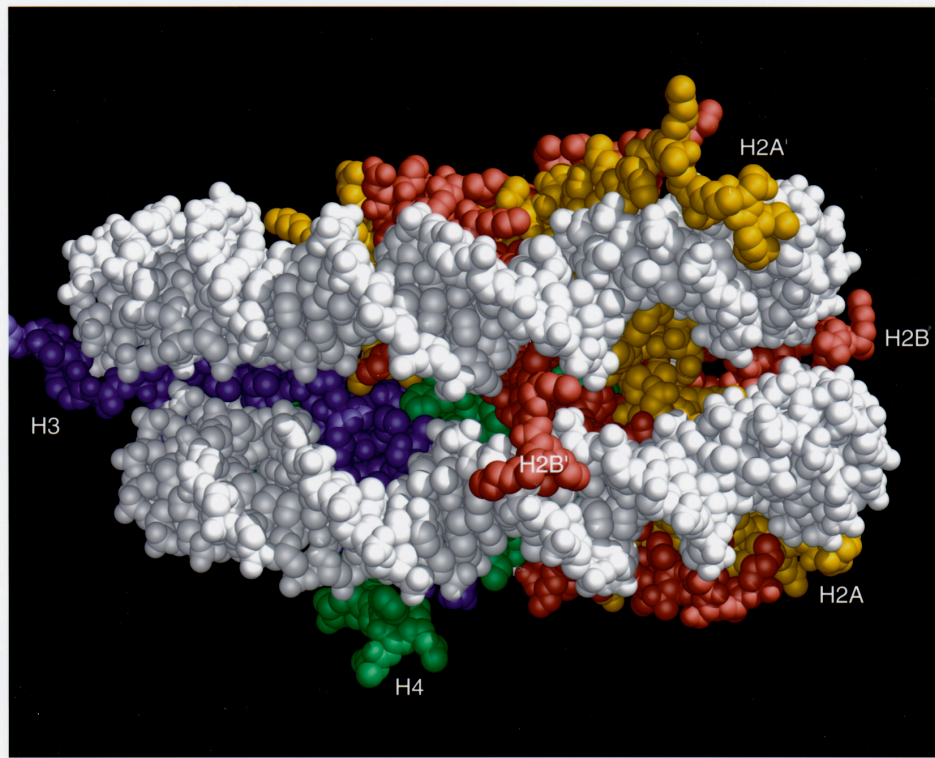
Lee & Schleif, 1989;
Müller et al., 1996

Hierarchical DNA folding in eukaryotic chromosomes



Felsenfeld & Groudine,,
Nature 421: 448-453 (2003)

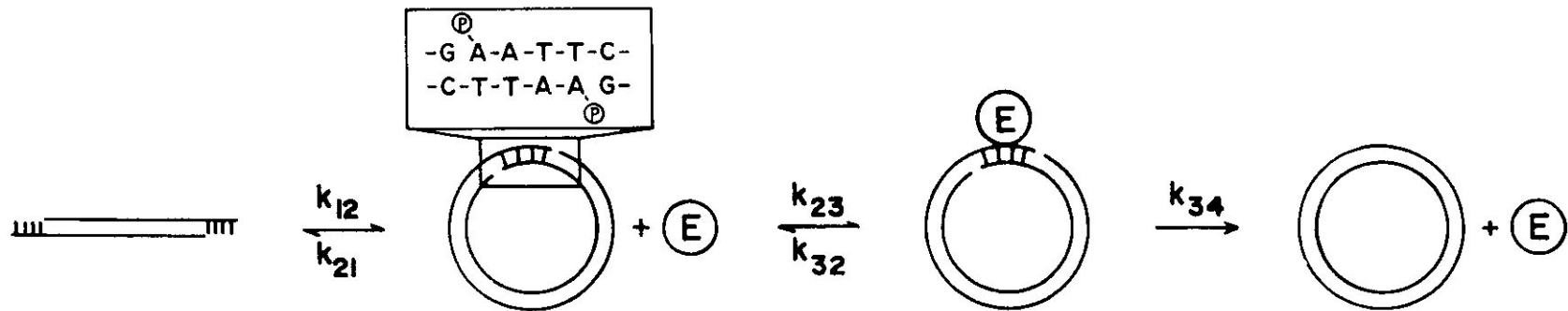
Most eukaryotic DNA is sharply looped



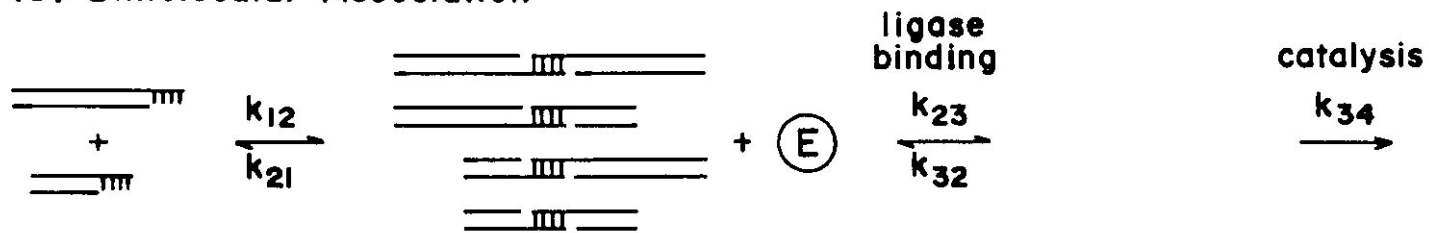
~80 bp per superhelical turn

Cyclization assay for DNA flexibility

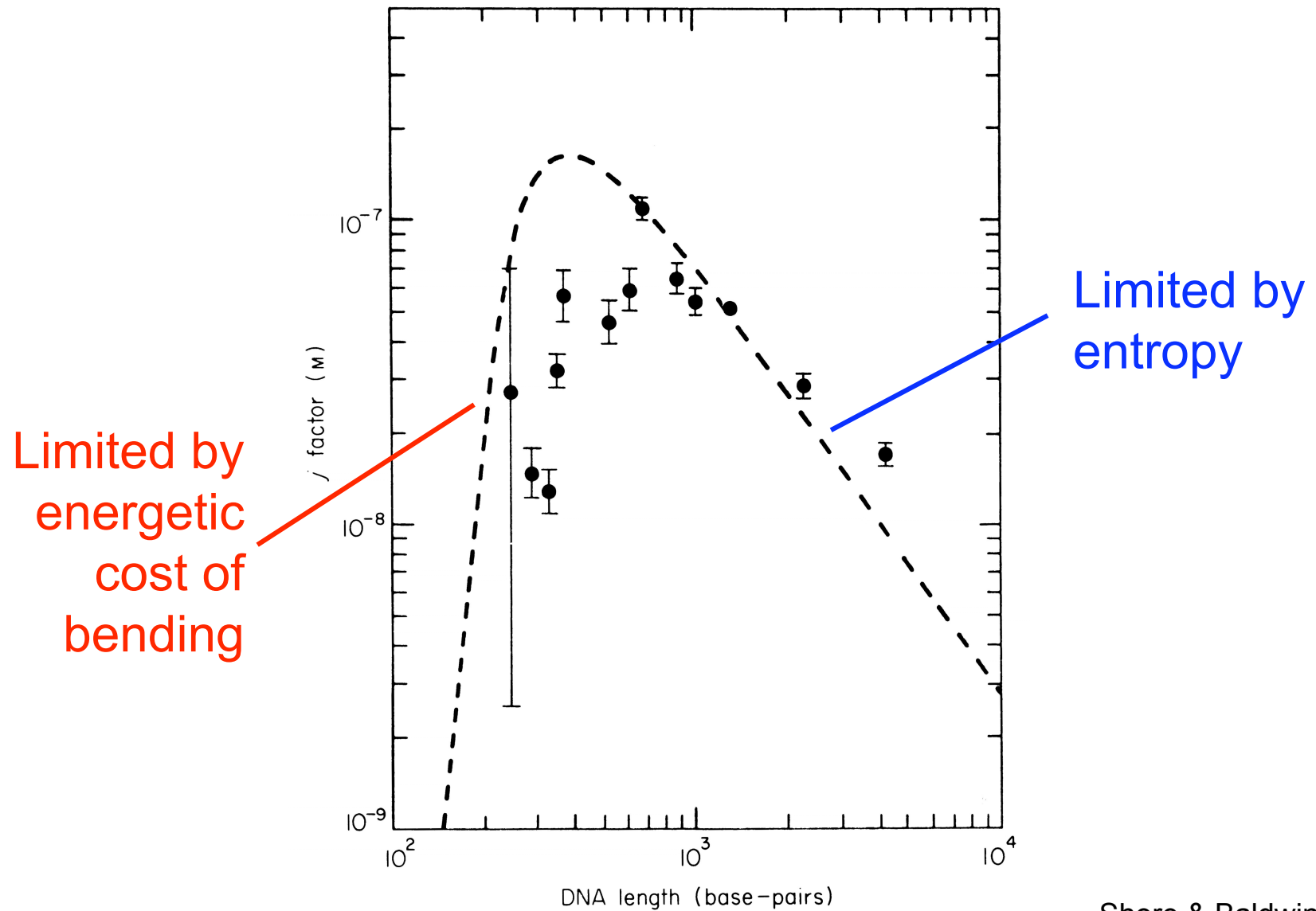
(a) Cyclization



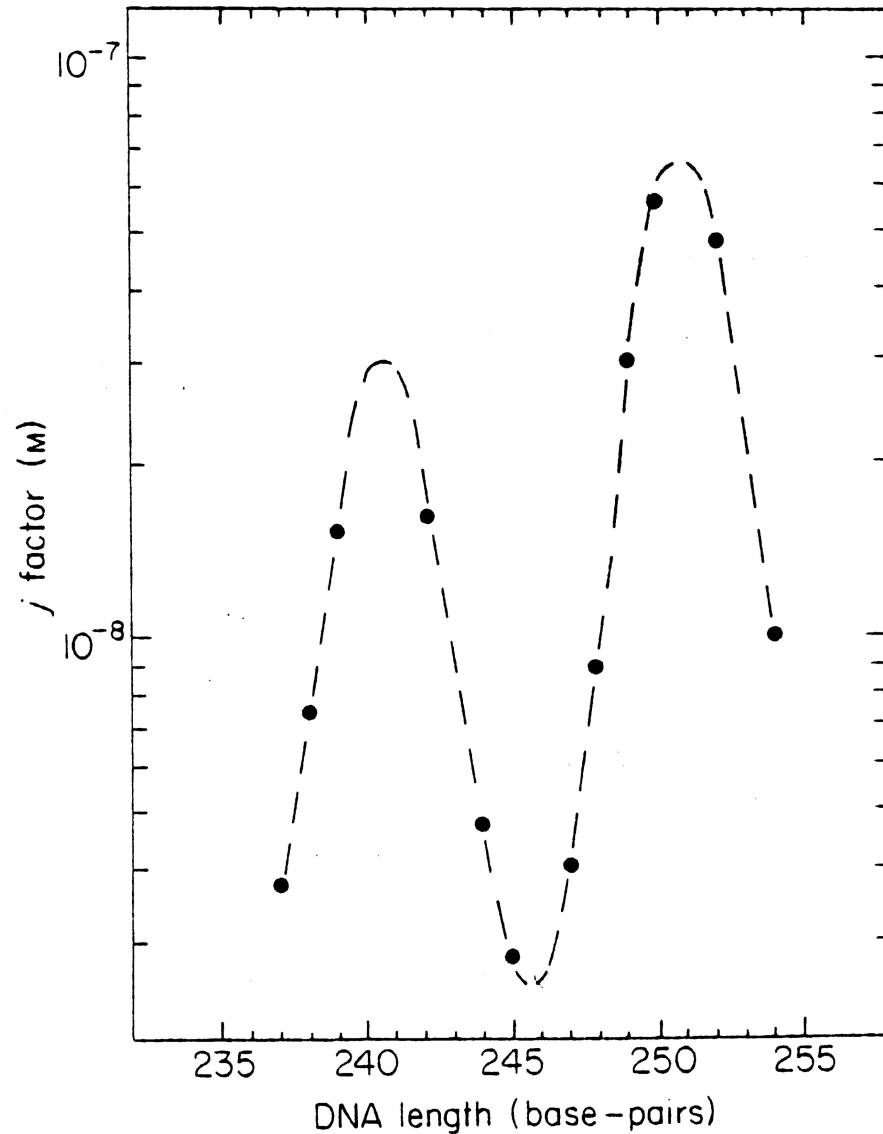
(b) Bimolecular Association



J depends on DNA length



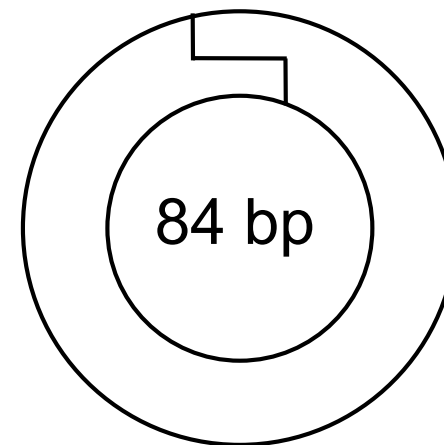
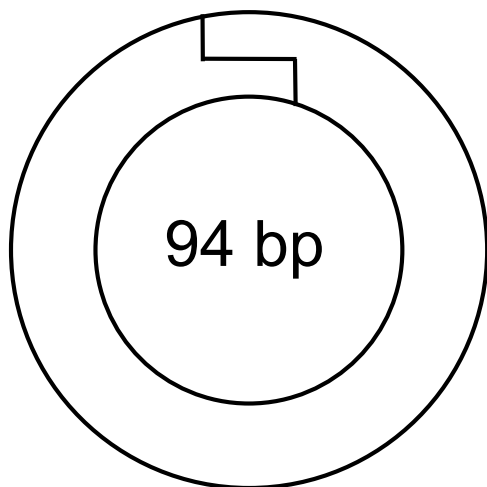
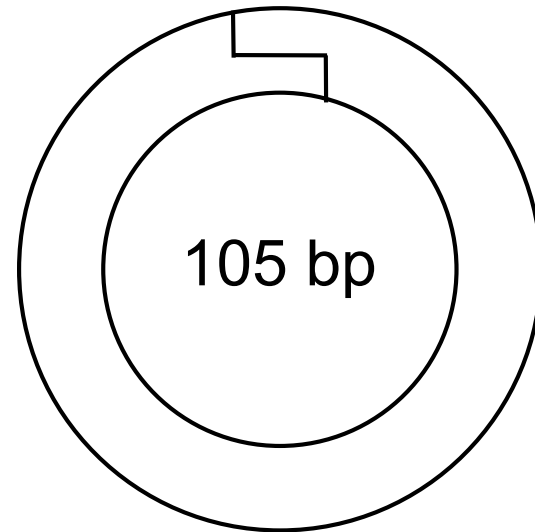
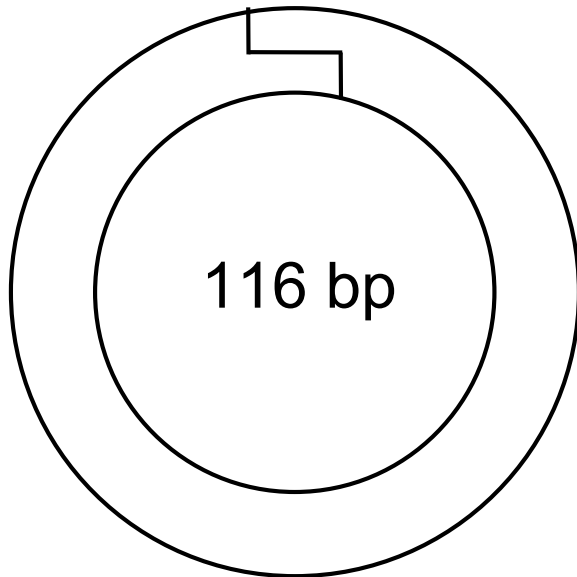
J depends on total DNA *twist*



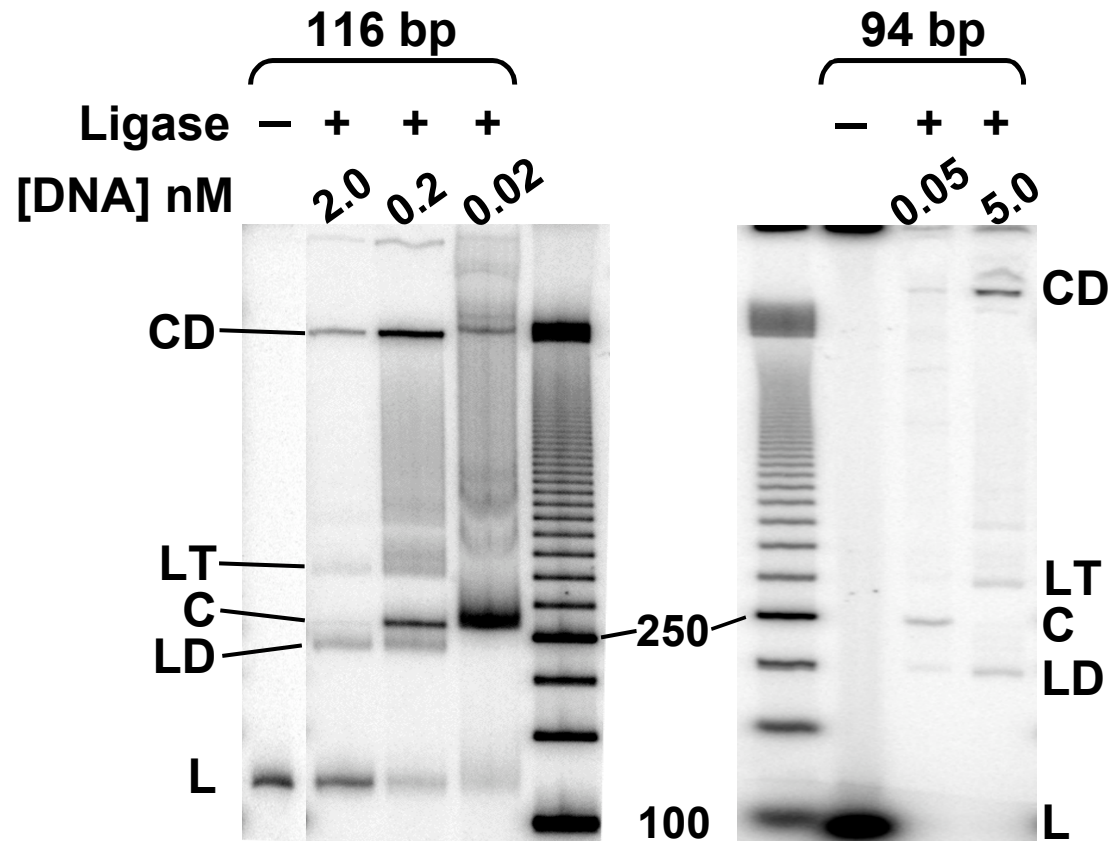
- Period equals DNA helical repeat

- Amplitude reflects the DNA torsional stiffness

Very small DNA circles!



116 bp and 94 bp circles are easy to make

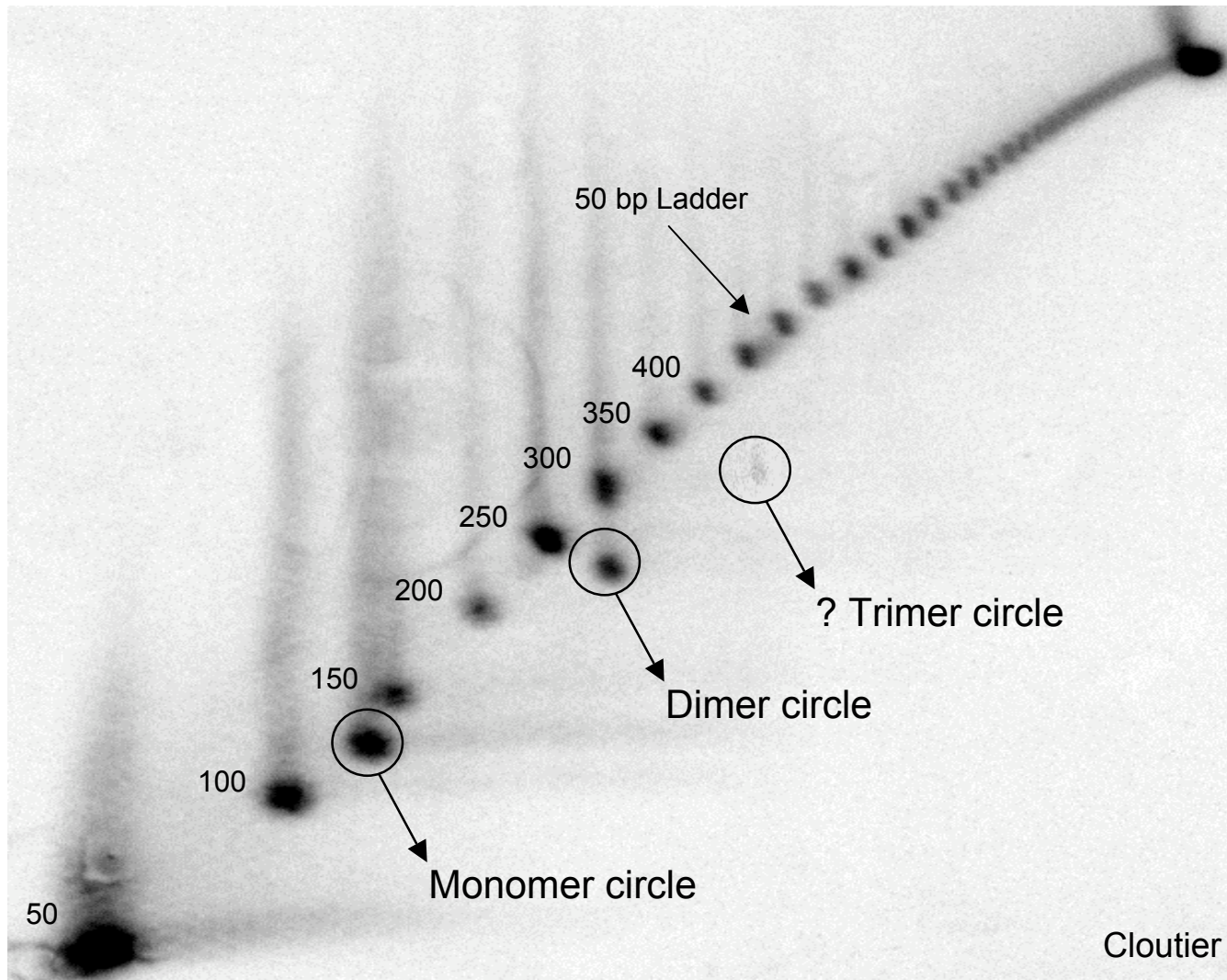


How do we know they are circles?

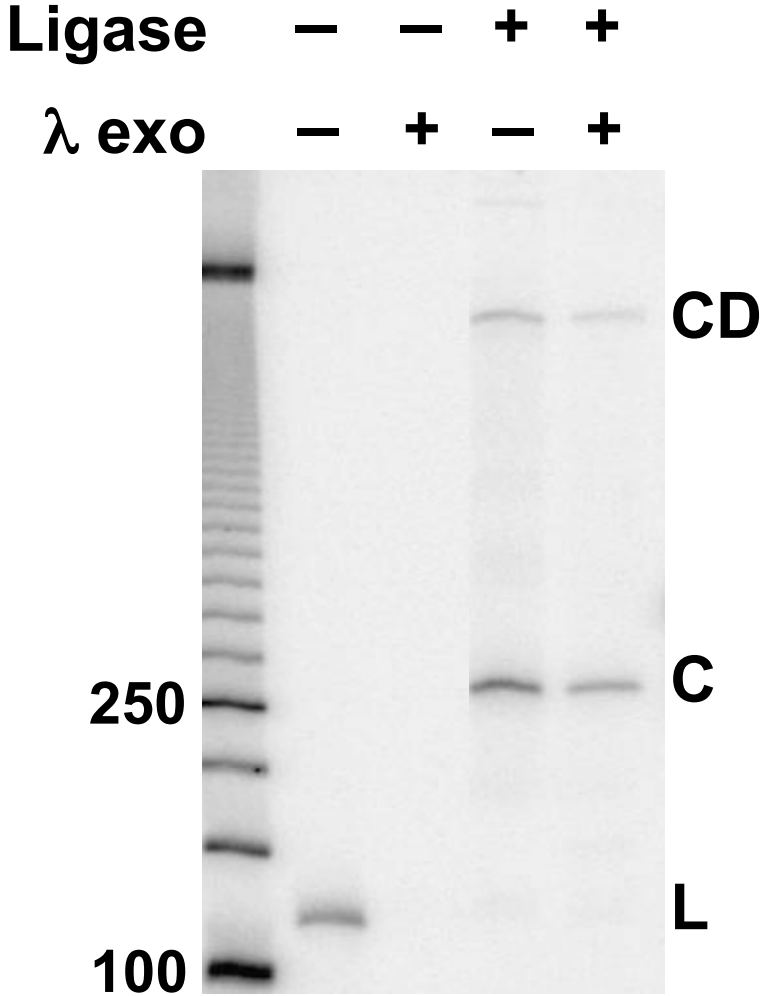
- (Monomeric) circles are favored at low concentration
- Circles run off the ladder of linear oligomers
- Circles run off diagonal in a topology-sensitive 2-D gel assay
- Circles resist digestion by exonuclease

Circles run off the diagonal in 2-D gel

Ligation of 116 bp DNA at 100 pM



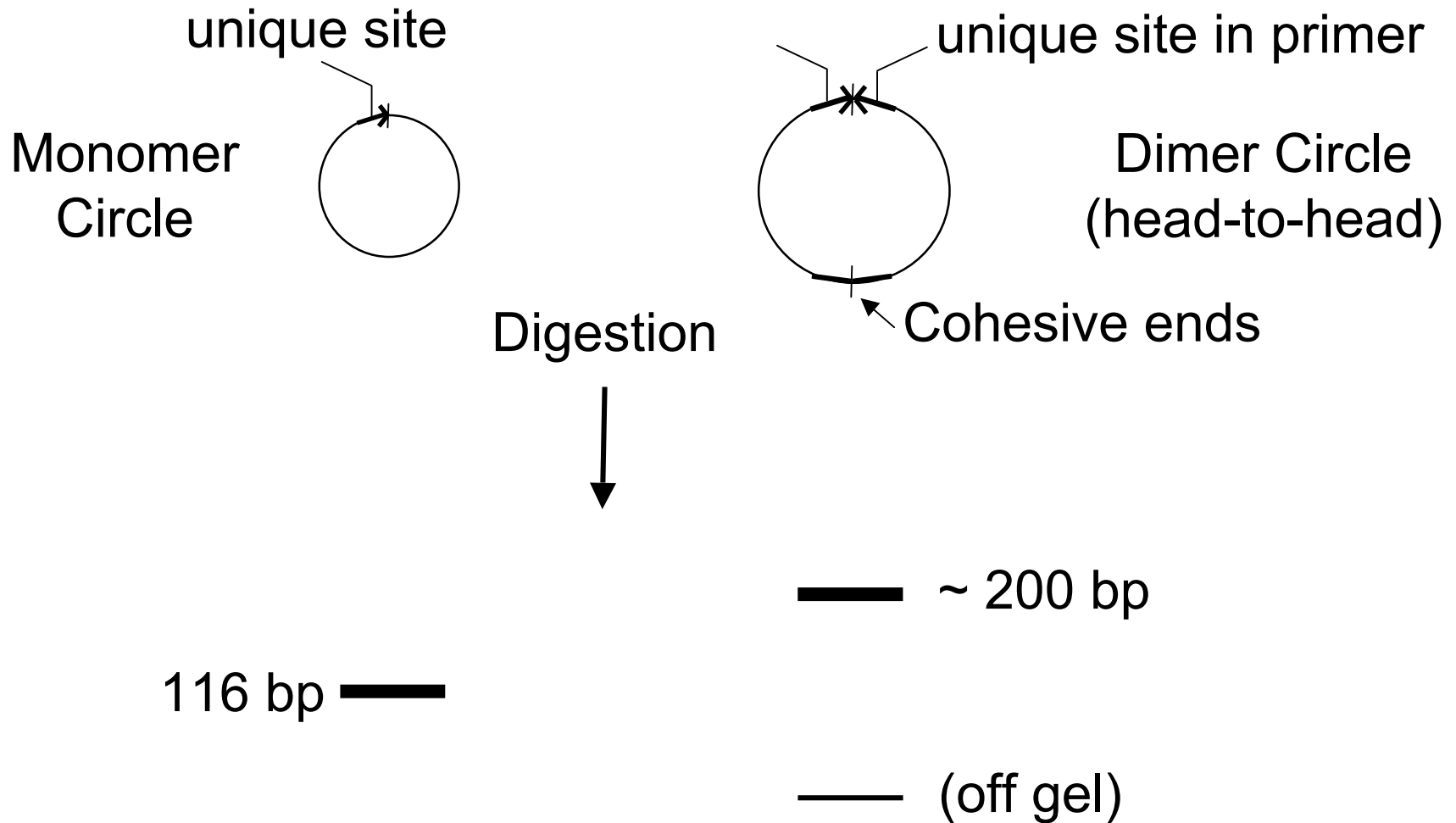
Circles resist digestion by exonuclease



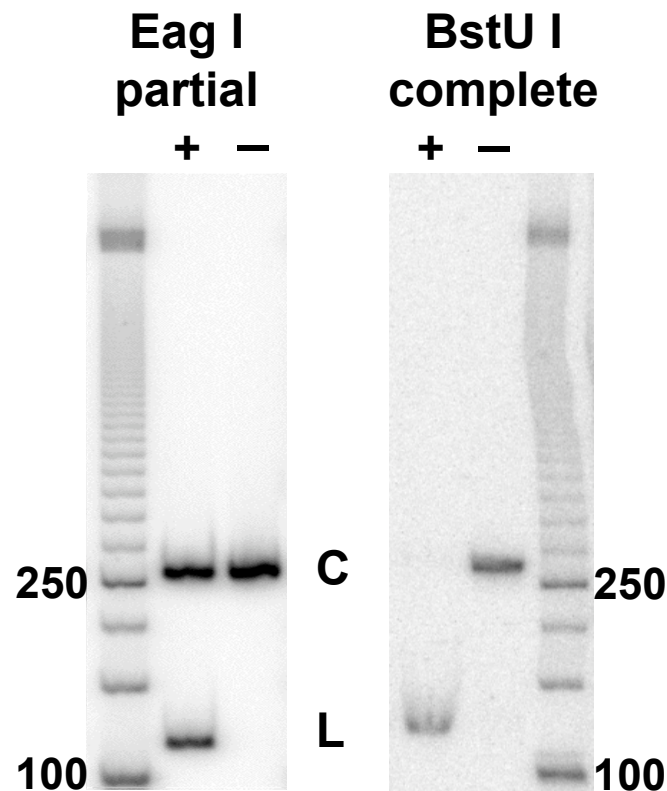
How do we know circles are monomeric?

- Monomeric circles are favored at low concentration
- Monomer circles run near monomer linears in agarose gels
- Partial restriction digestion yields only linear monomer
- Complete restriction digestion nearby cohesive site yields only linear monomer

Restriction enzyme digestion distinguishes monomers from oligomers

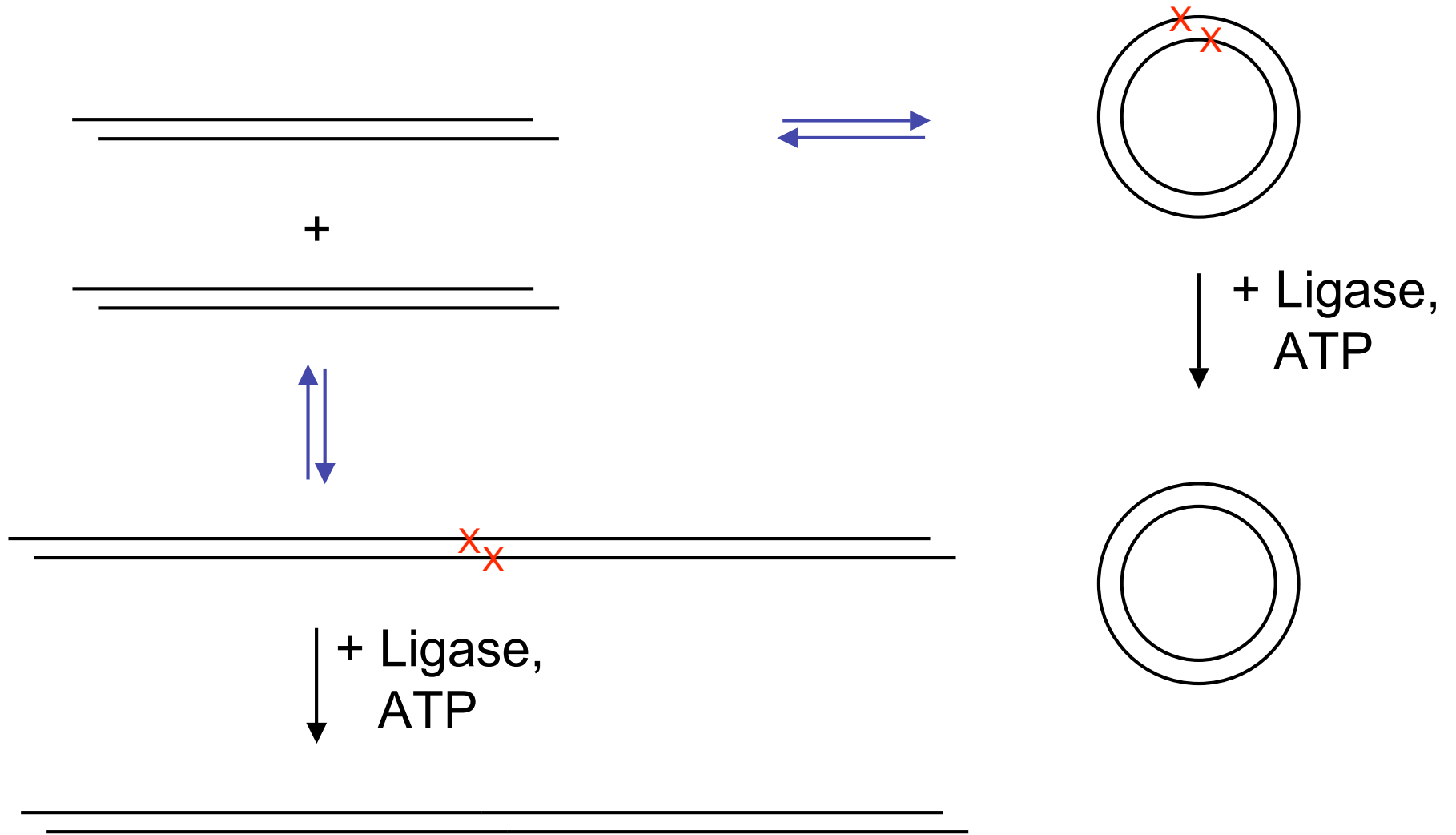


Restriction enzyme digestion distinguishes monomers from oligomers

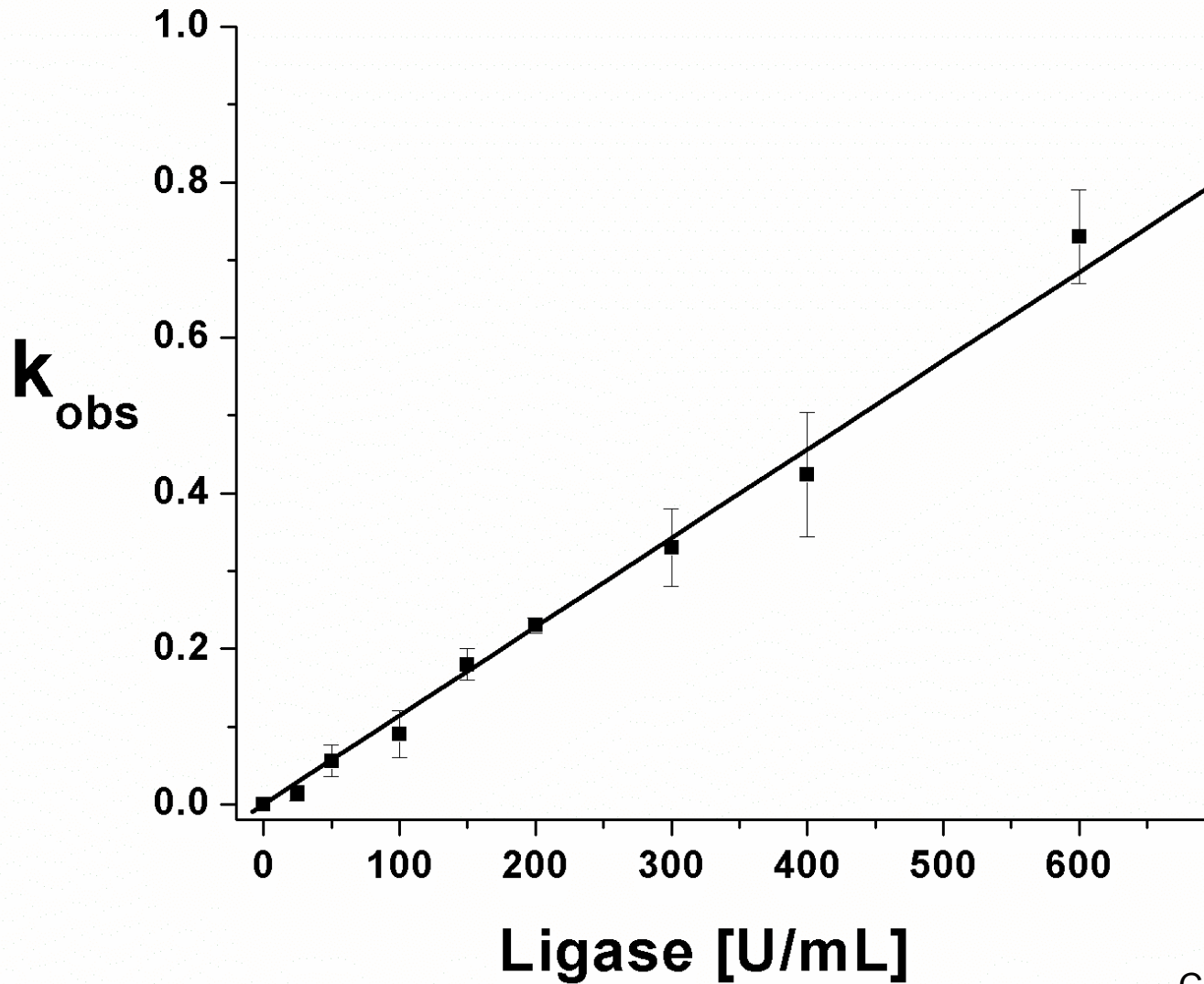


- Partial digestion by Eag I yields only linear monomer
- Complete restriction digestion by BstU I yields only linear monomer

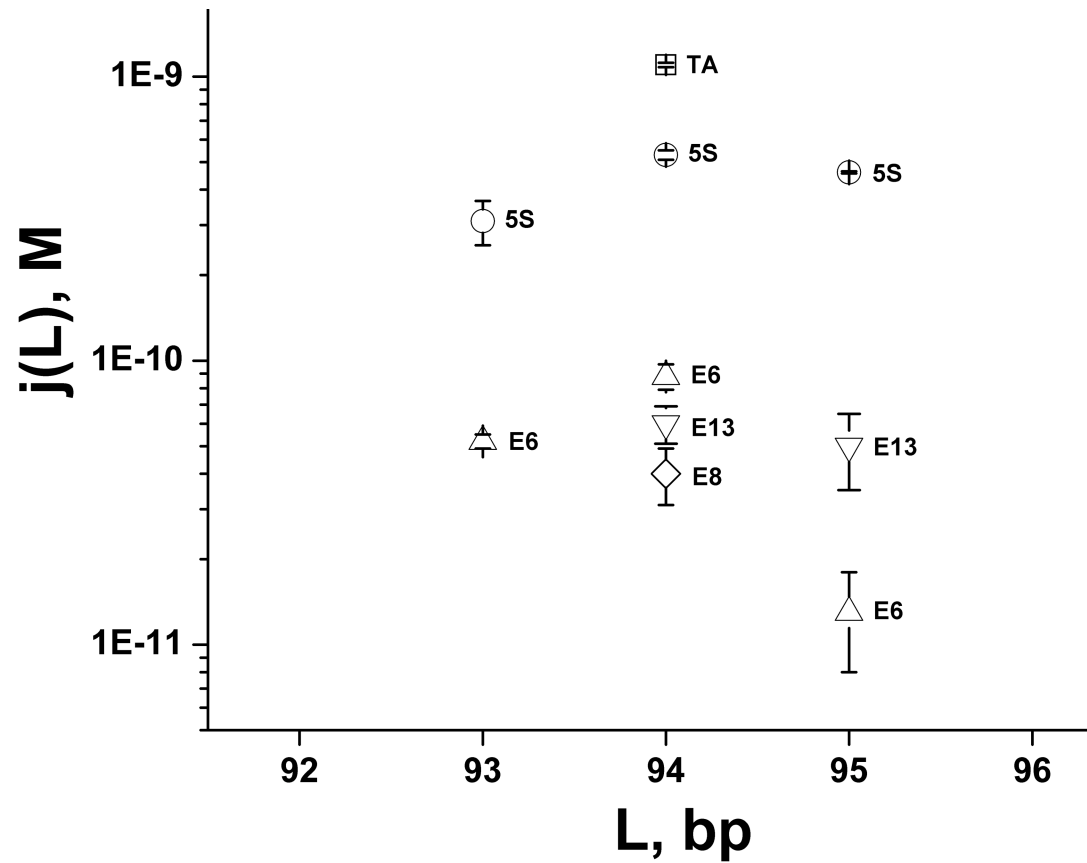
Quantitative measurement of J factor



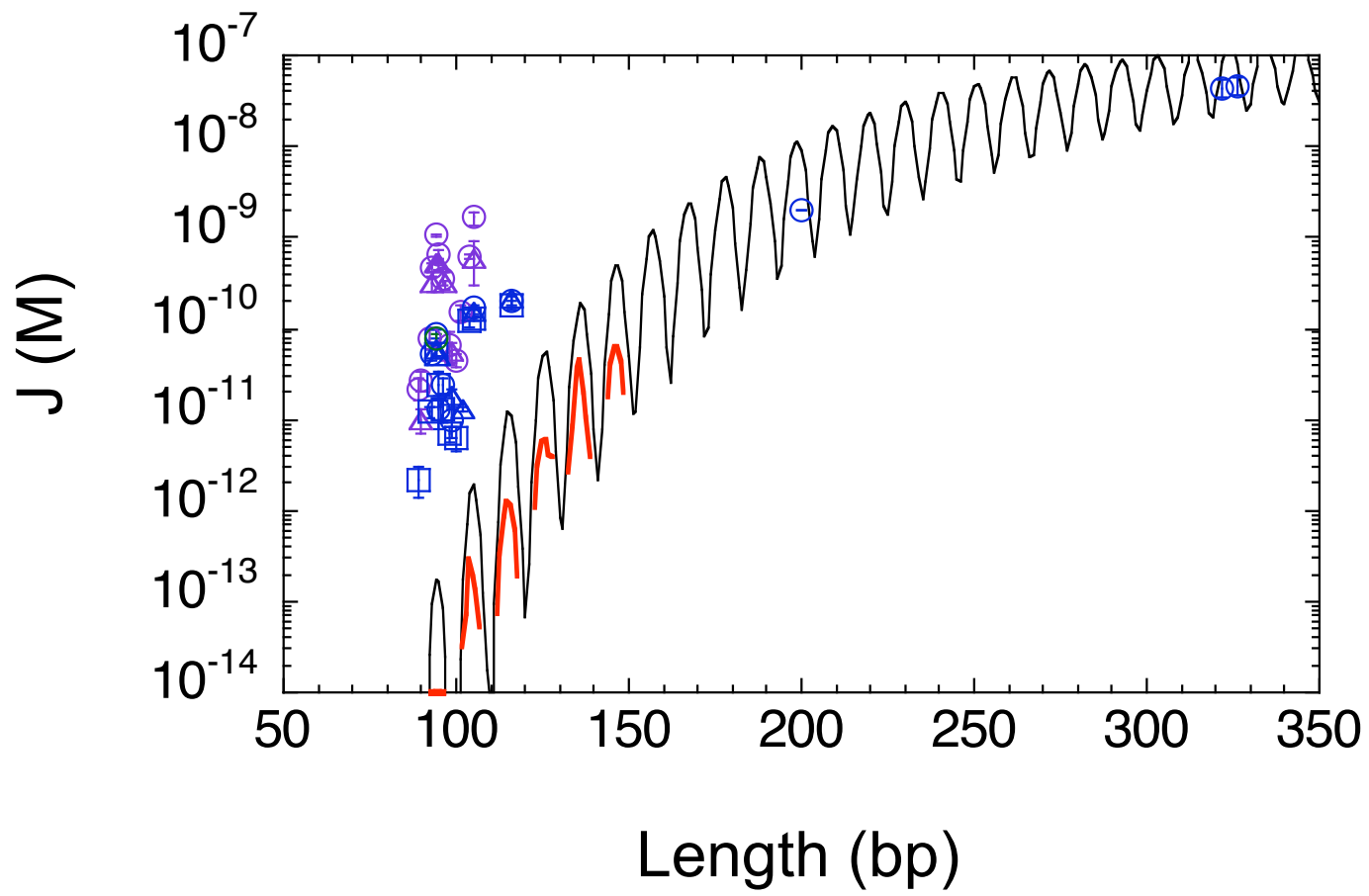
Cyclization reactions with 94bp DNAs are first order in ligase concentration



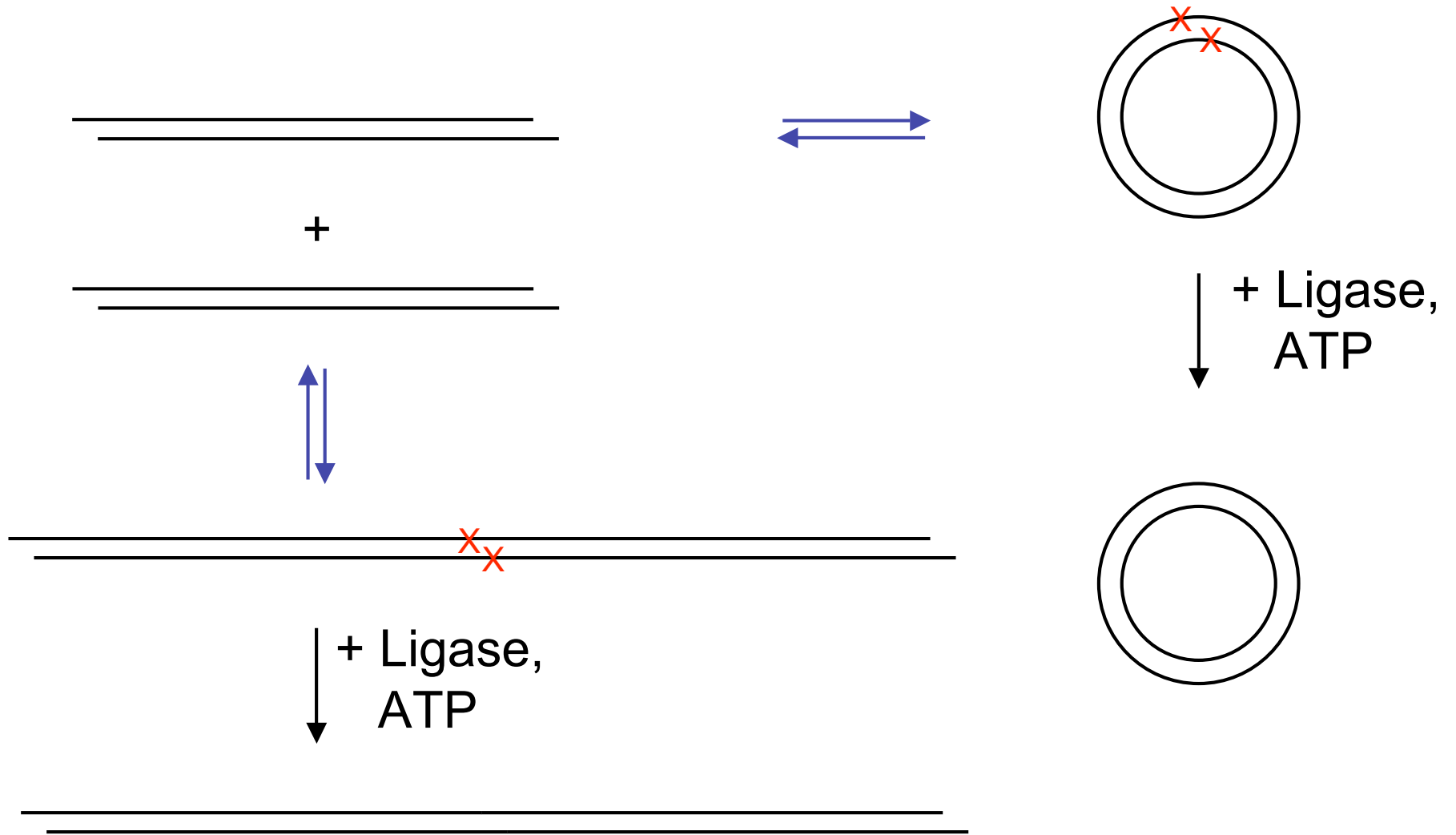
Differing DNA sequences differ in inherent cyclizability



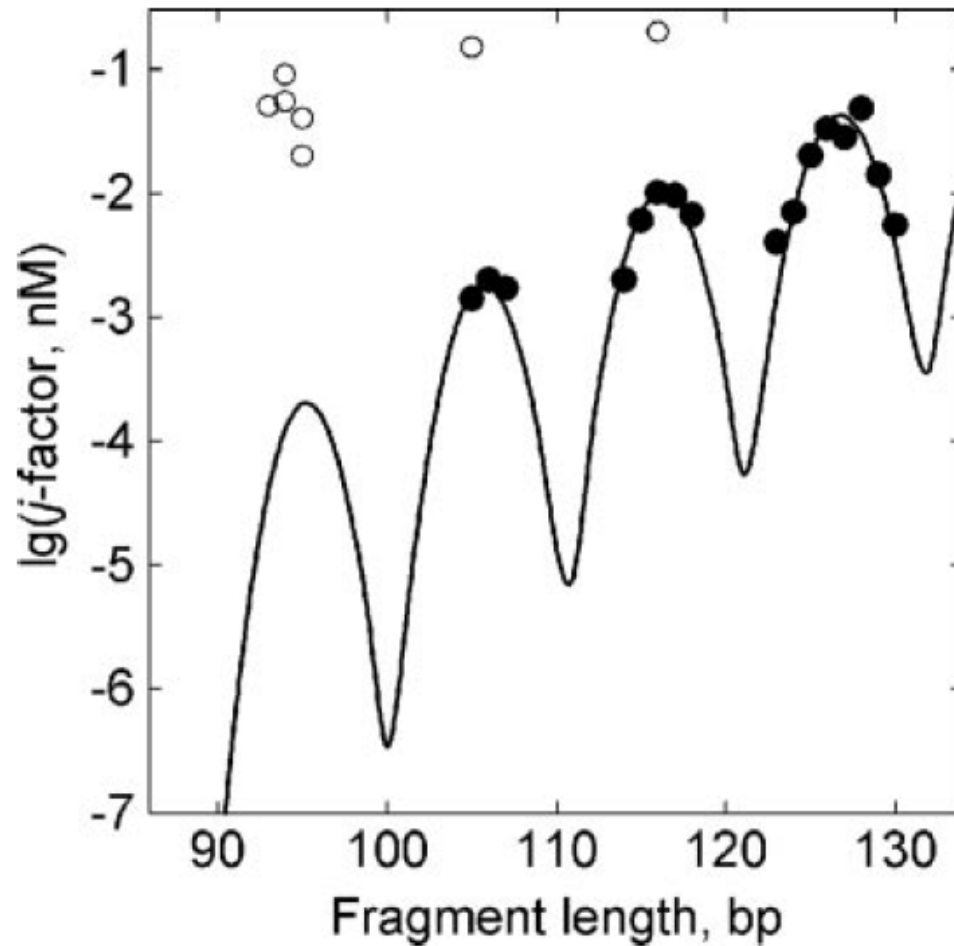
Sharply bent DNA appeared to be much softer for sharp looping than predicted



Quantitative measurement of J factor

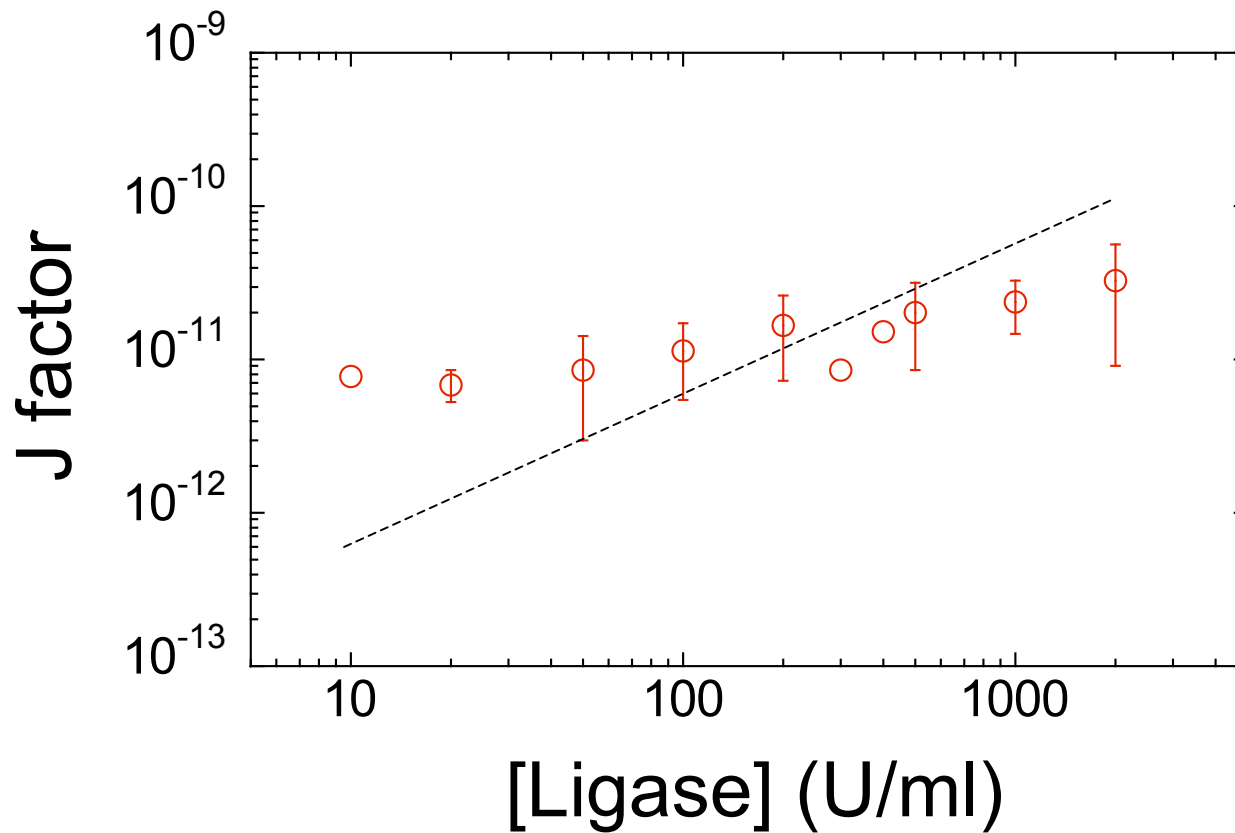


DNA may *not* be softer for sharp looping than predicted

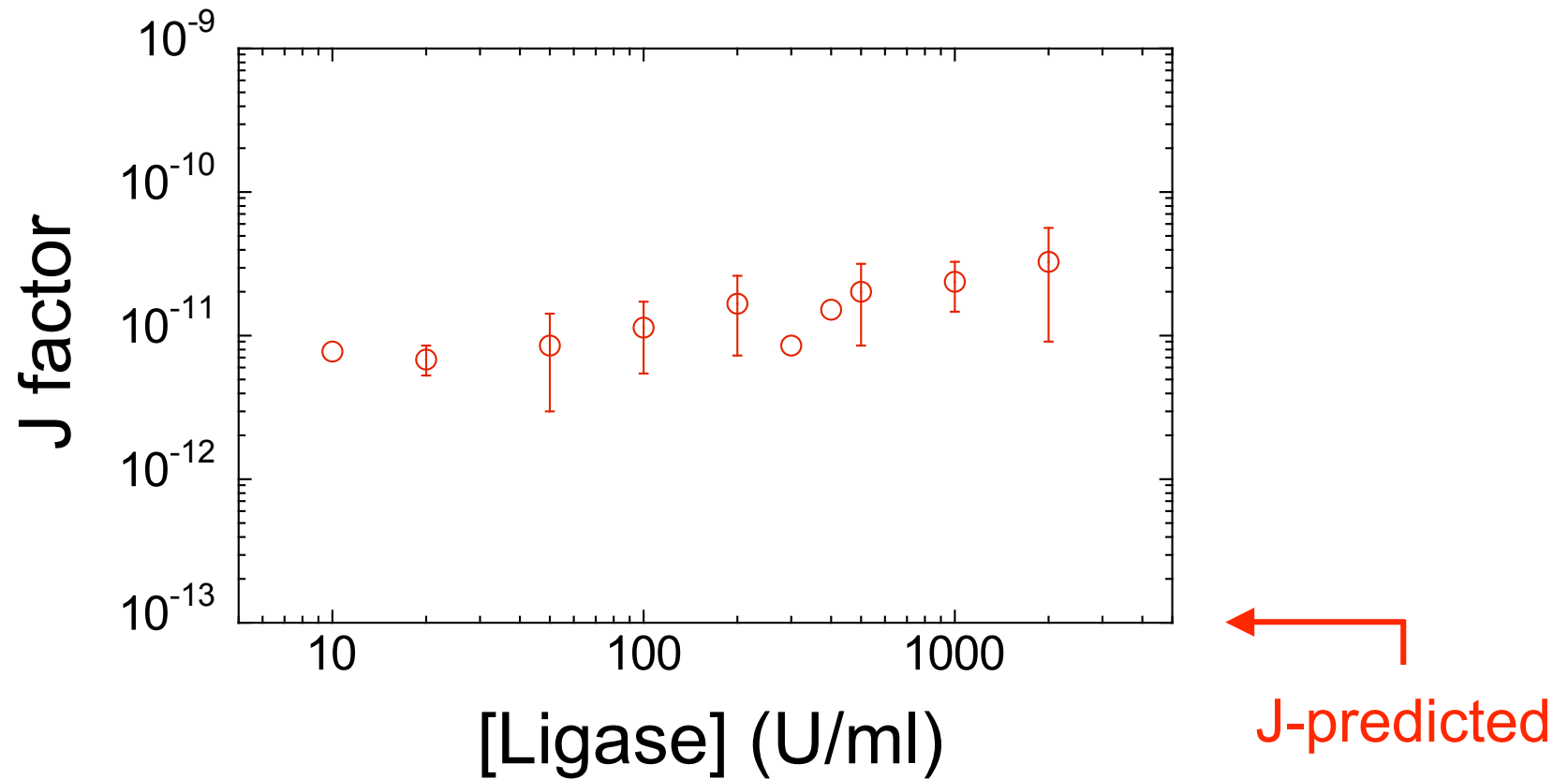


Measured J factors are independent of [ligase]

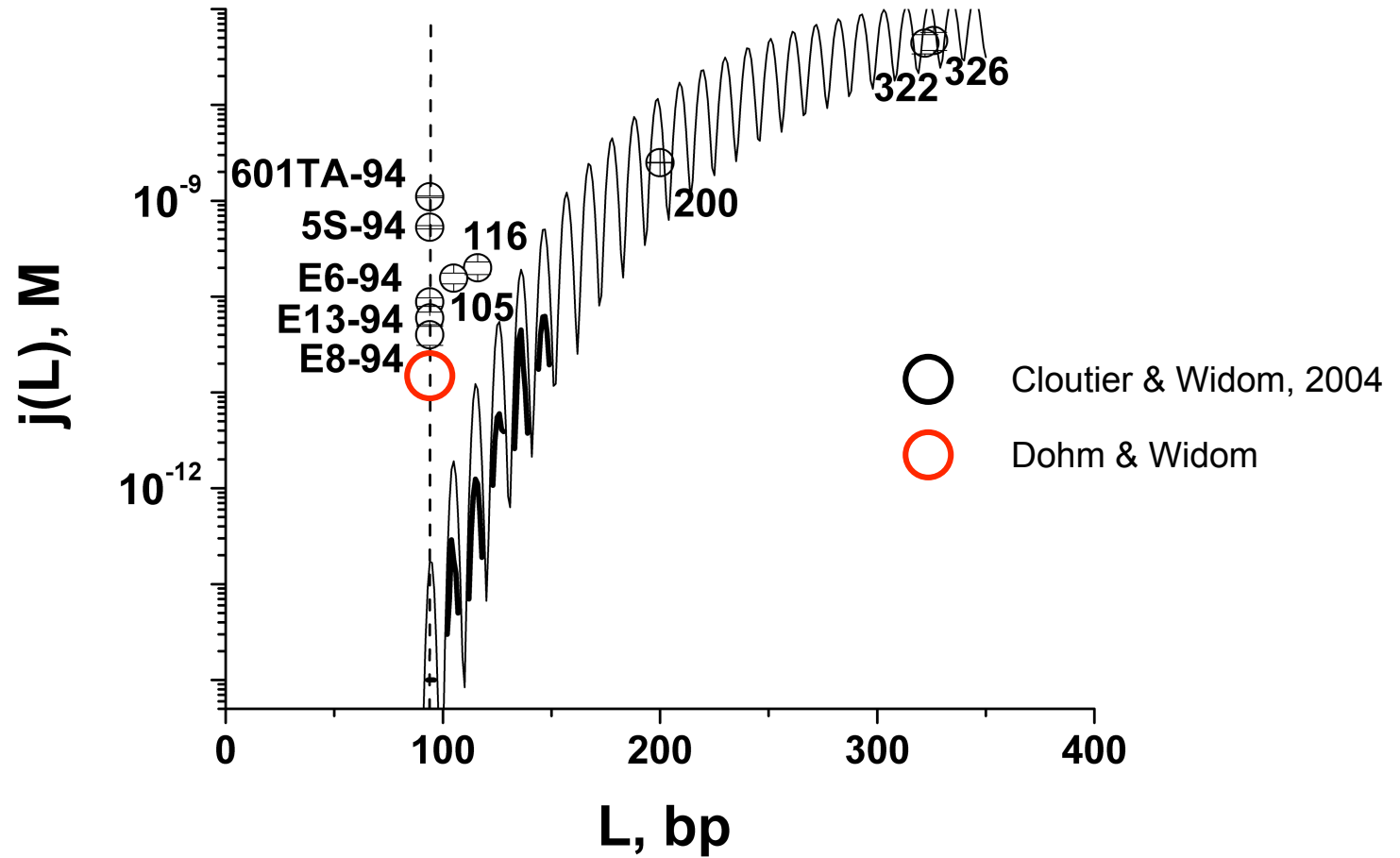
- bimolecular joining reactions, too, are first order in [ligase]



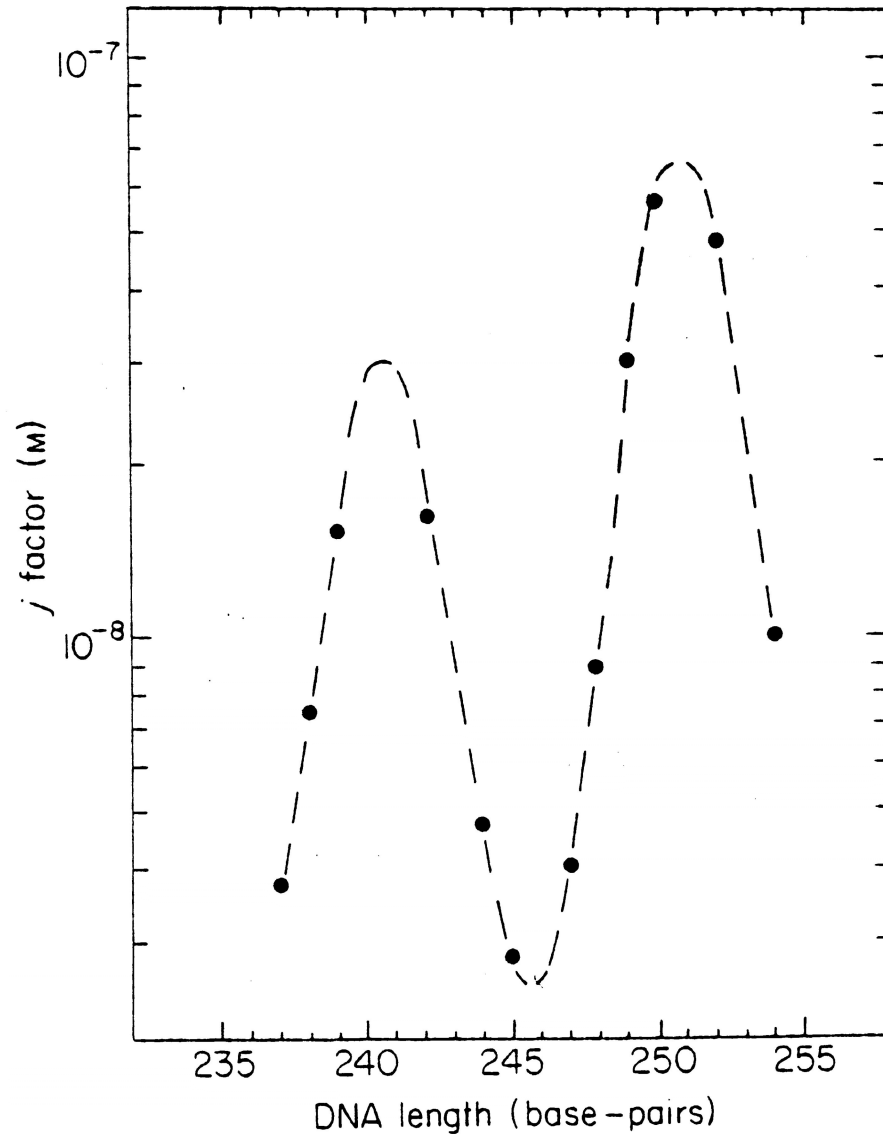
Measured J factors greatly exceed prediction



Measured J factors greatly exceed prediction



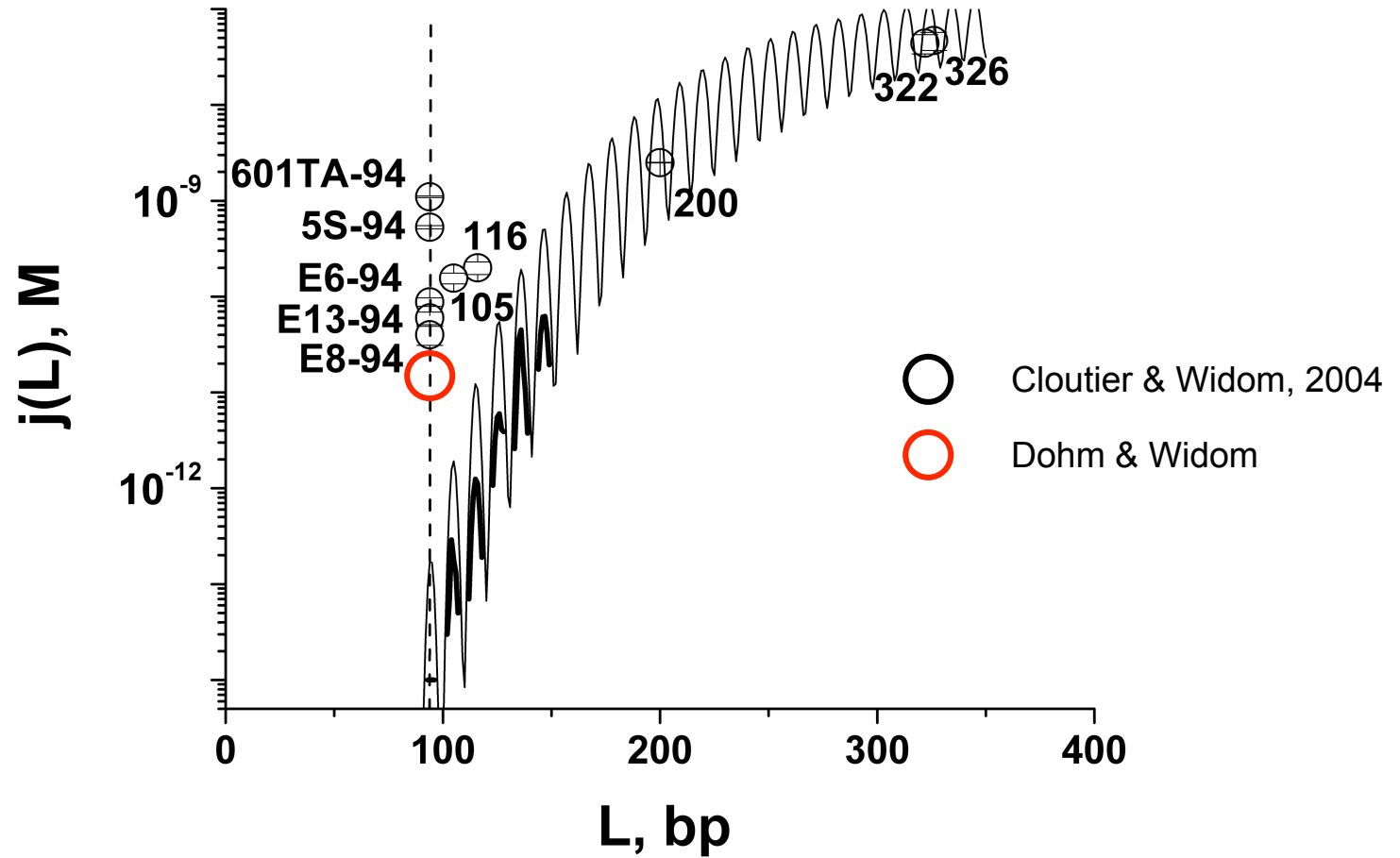
J depends on total DNA *twist*



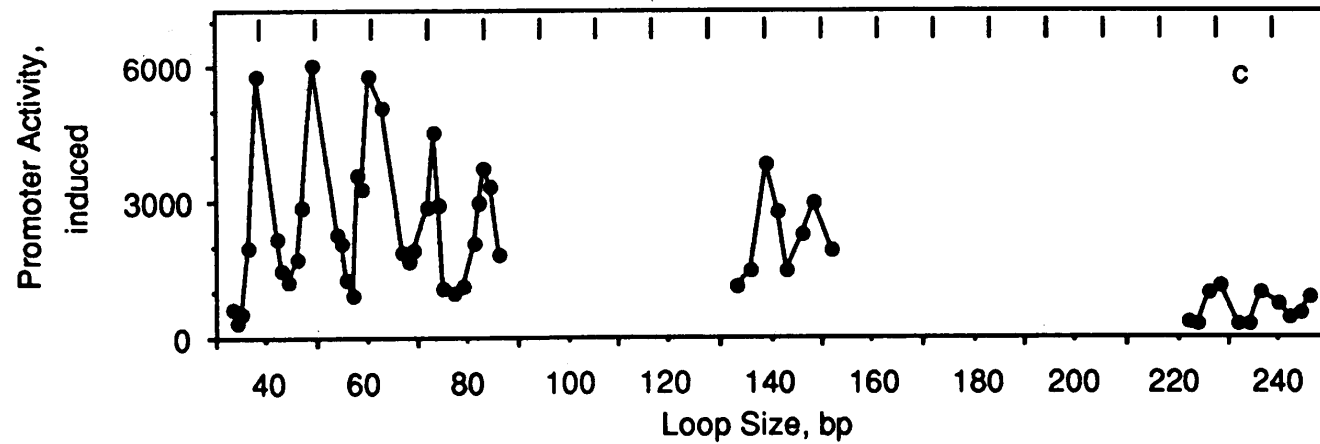
- Period equals DNA helical repeat

- Amplitude reflects the DNA torsional stiffness

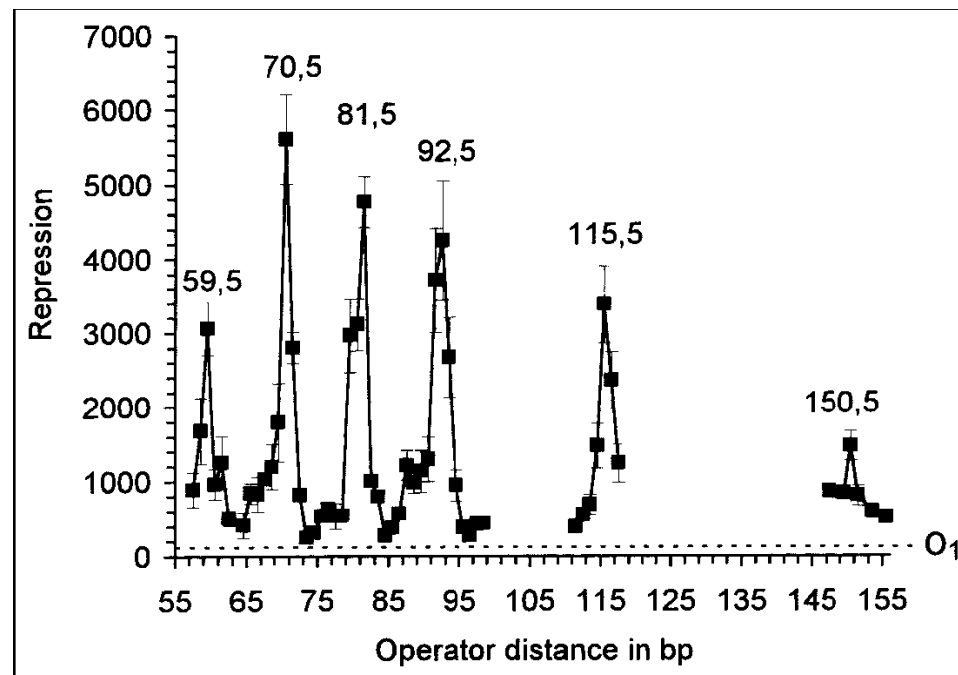
Measured J factors greatly exceed prediction



Sharply looped and *twisted* DNA *in vivo*



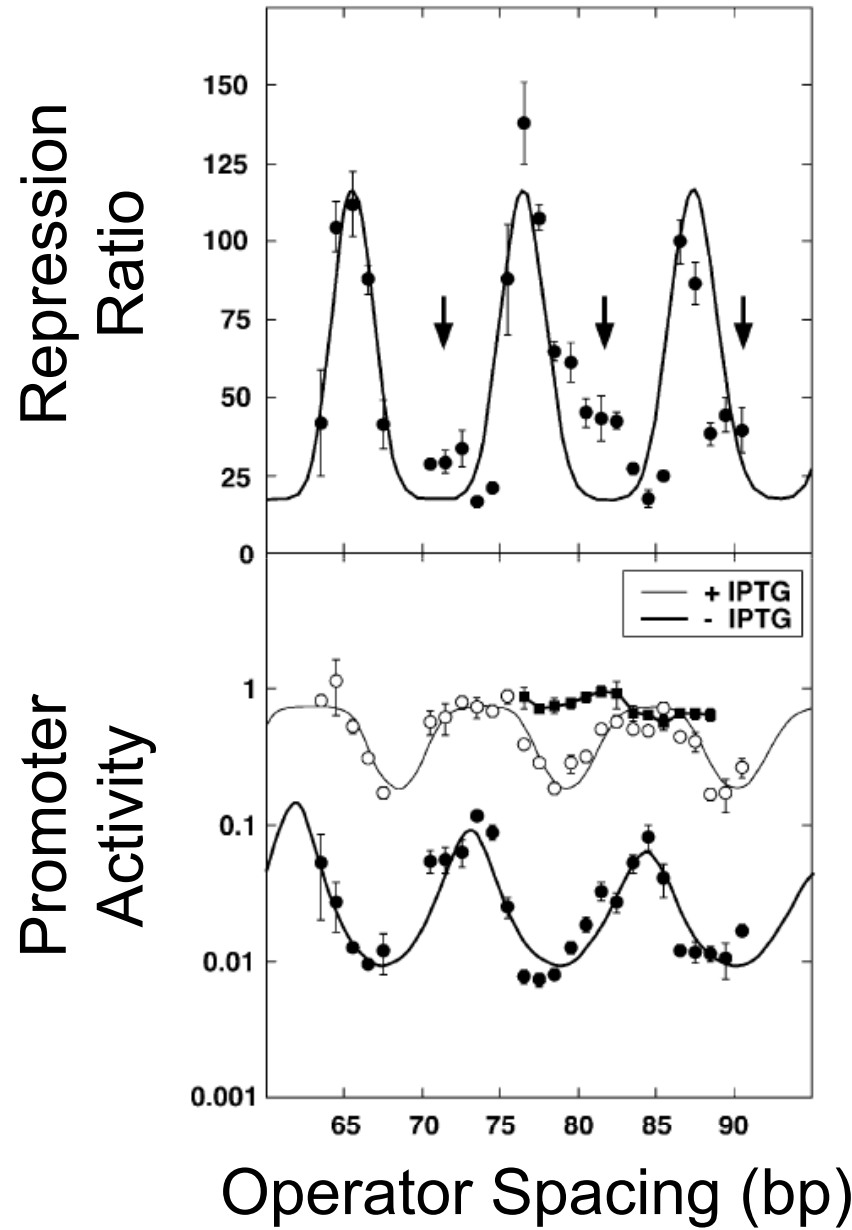
araCBAD



lac

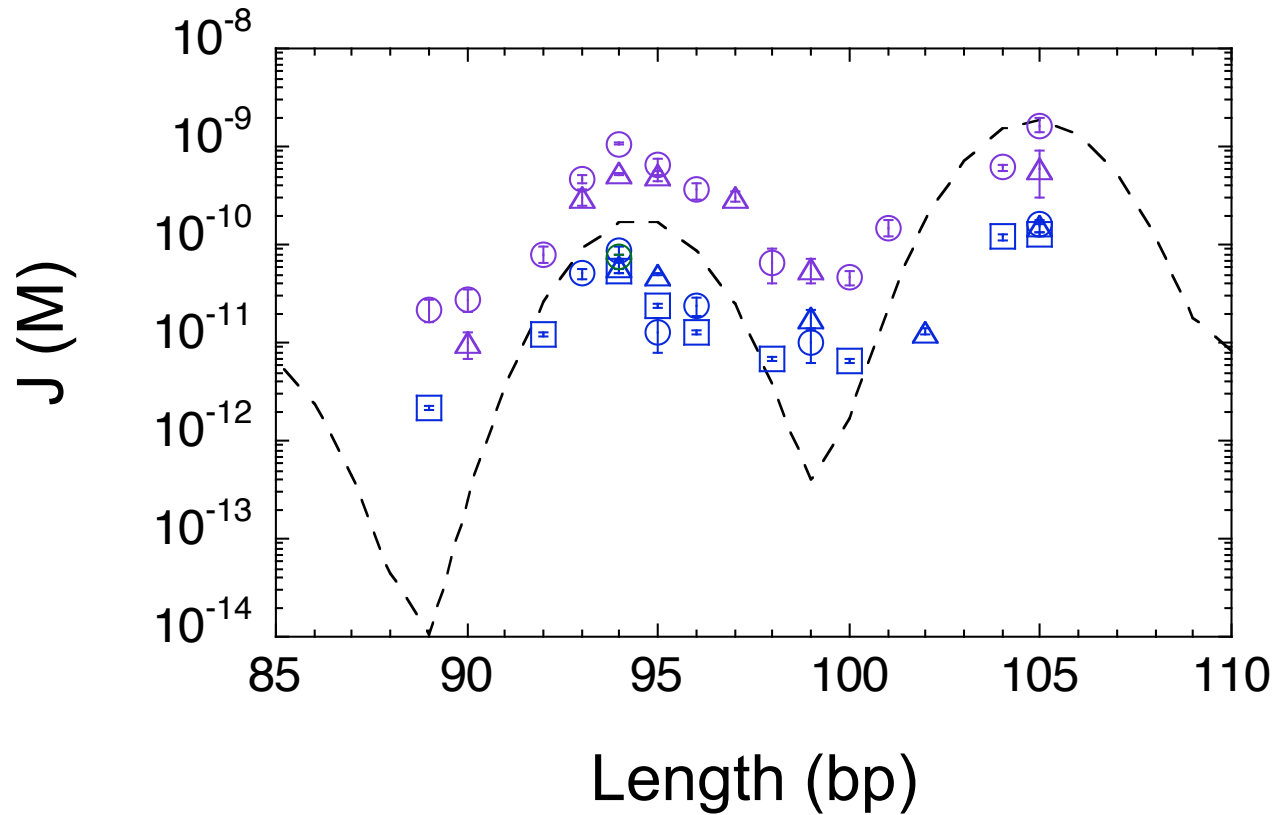
Lee & Schleif, 1989;
Müller et al., 1996

Sharply looped and *twisted* DNA, *in vivo*

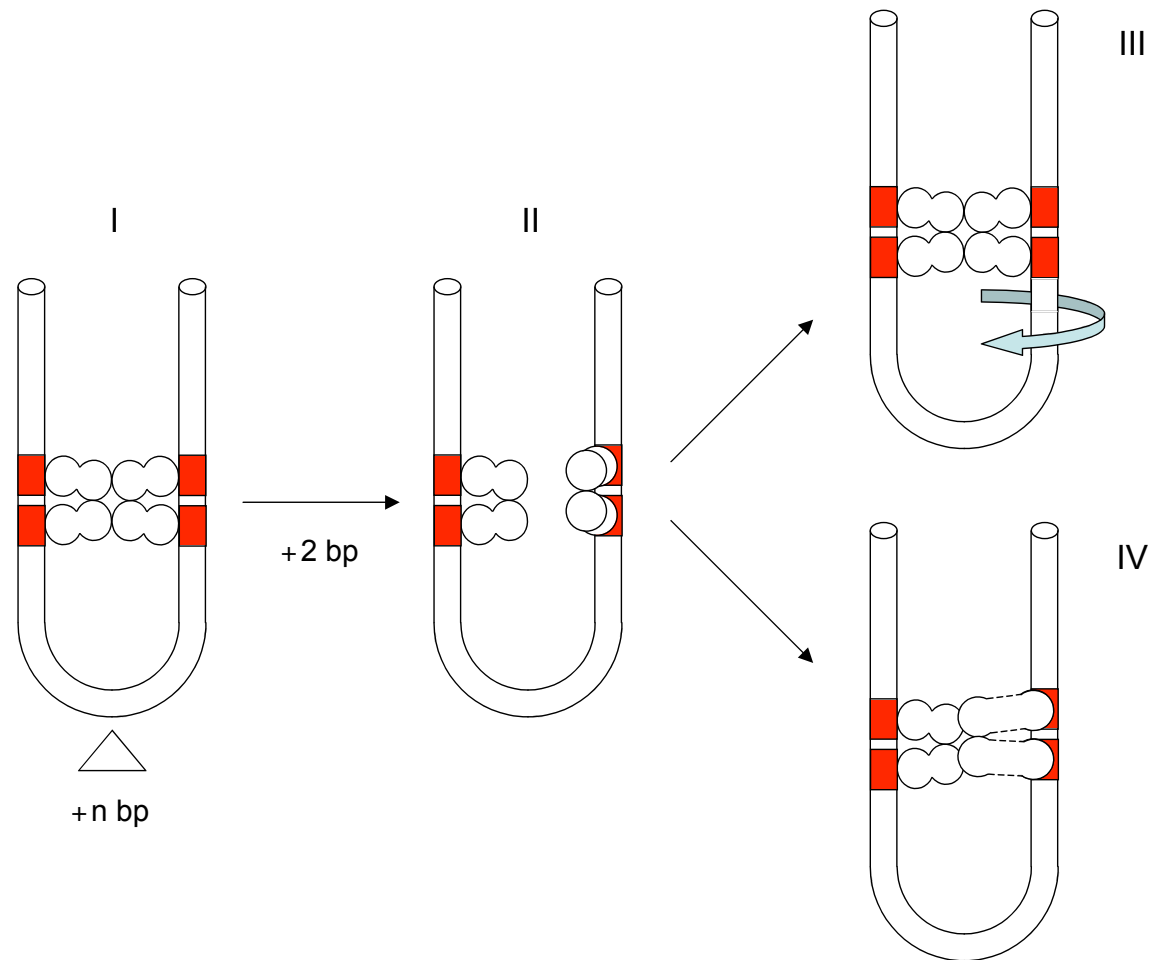


• *lac* promoter

Sharply bent DNA also appears to be much softer for twisting than predicted

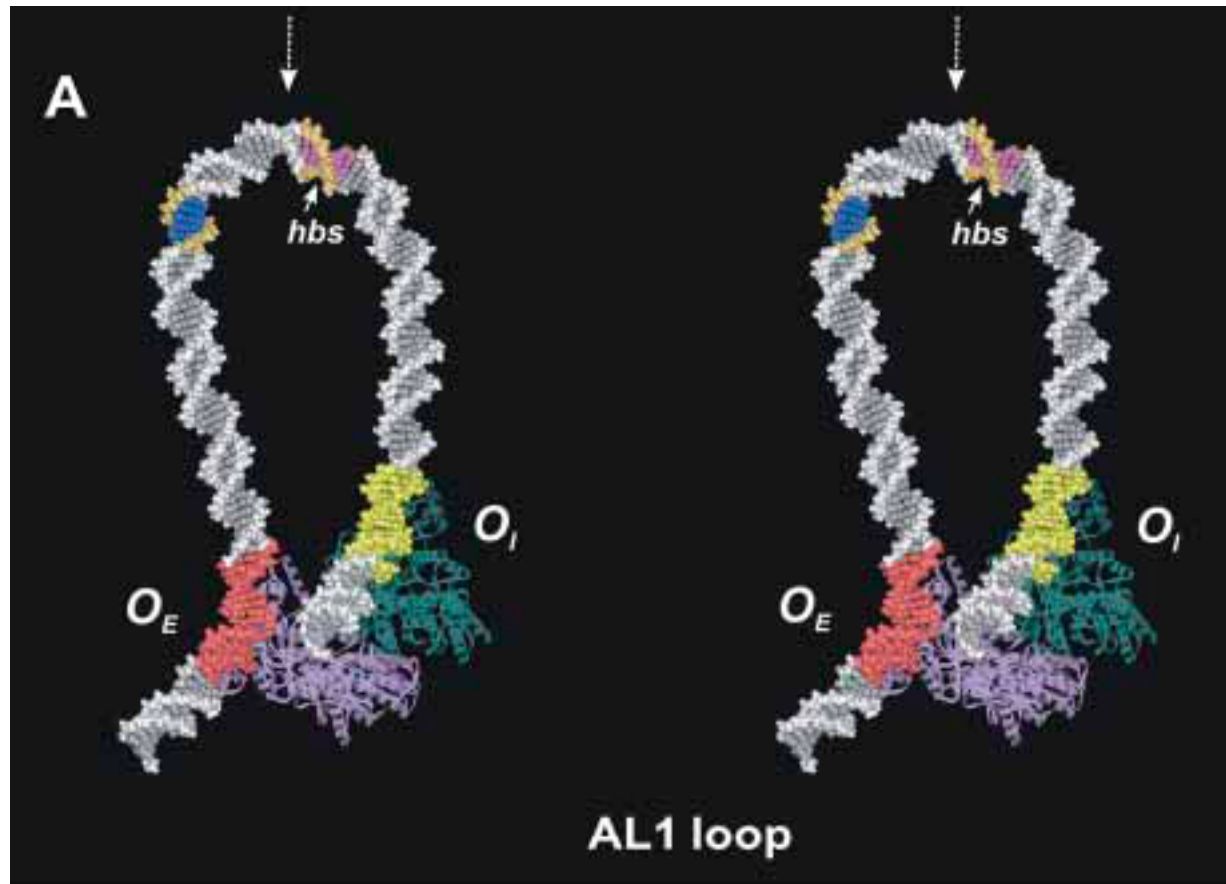


Formation and stability of sharply looped protein–DNA complexes

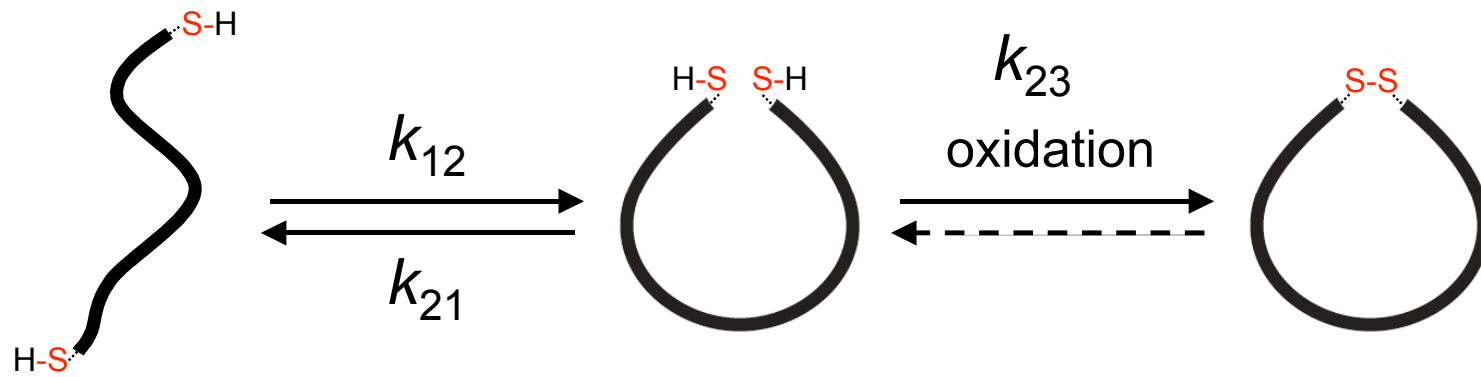


Looping vs cyclization

- Sharply looped DNA in the Gal repressosome

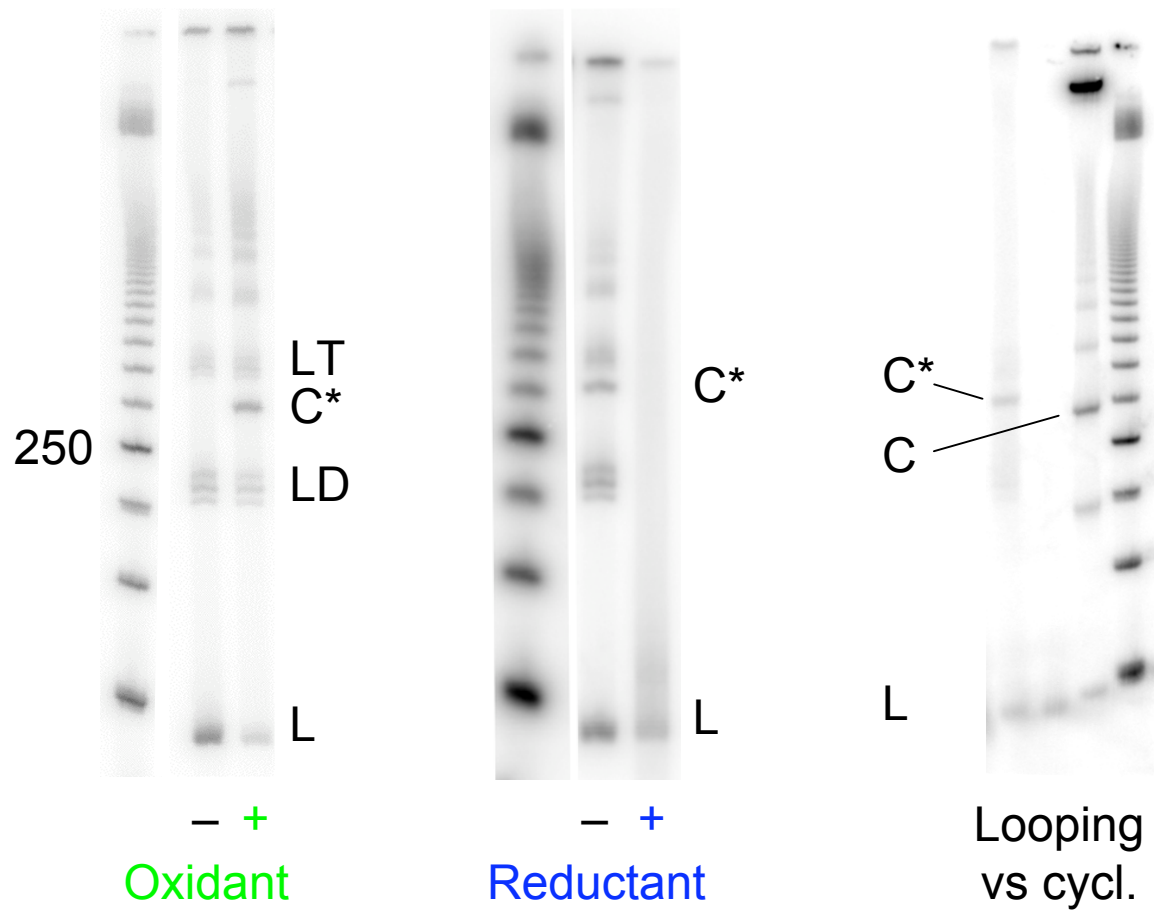


Chemical trapping assay for looping equilibria and kinetics

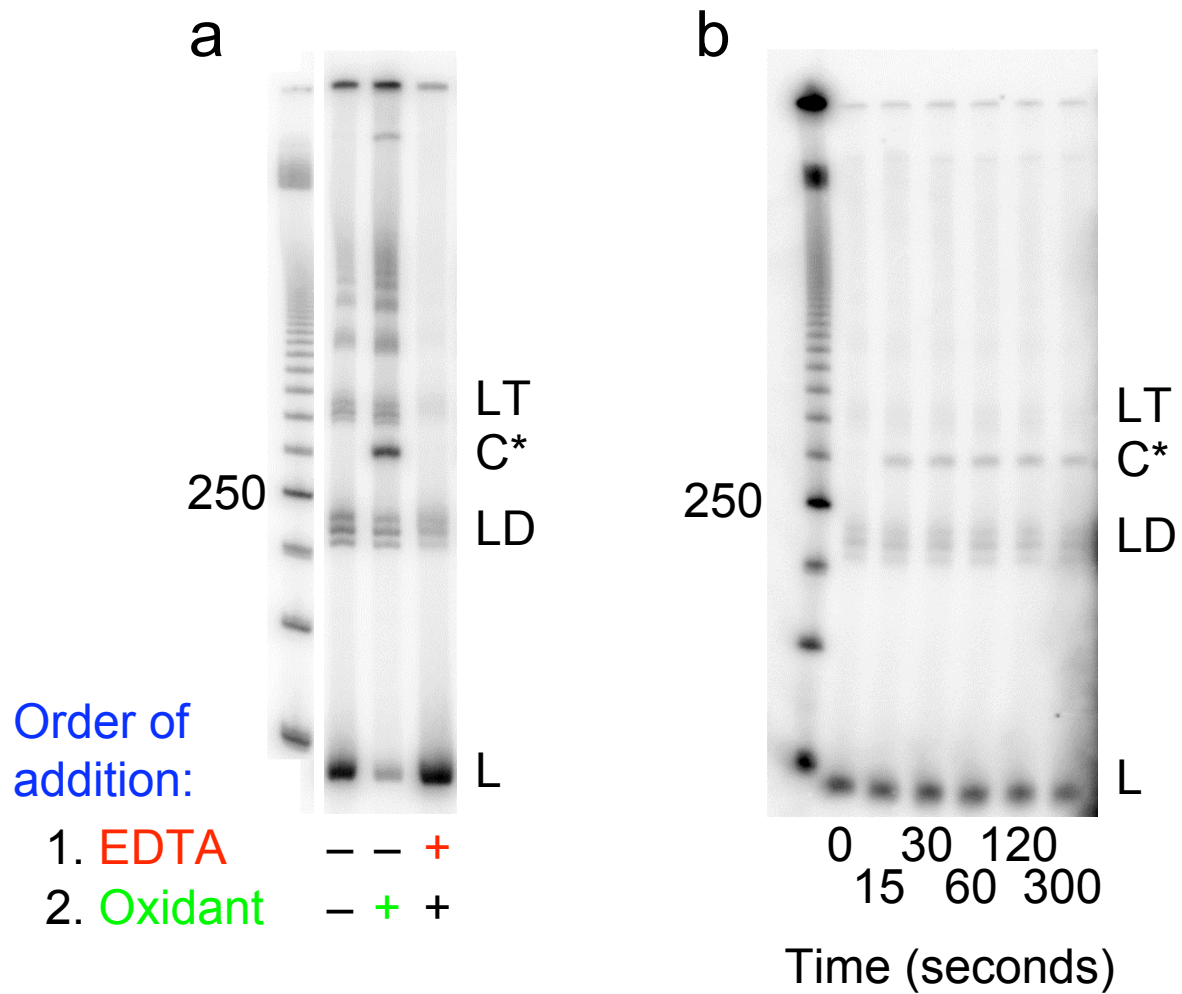


Chemical trapping assay for looping equilibria and kinetics

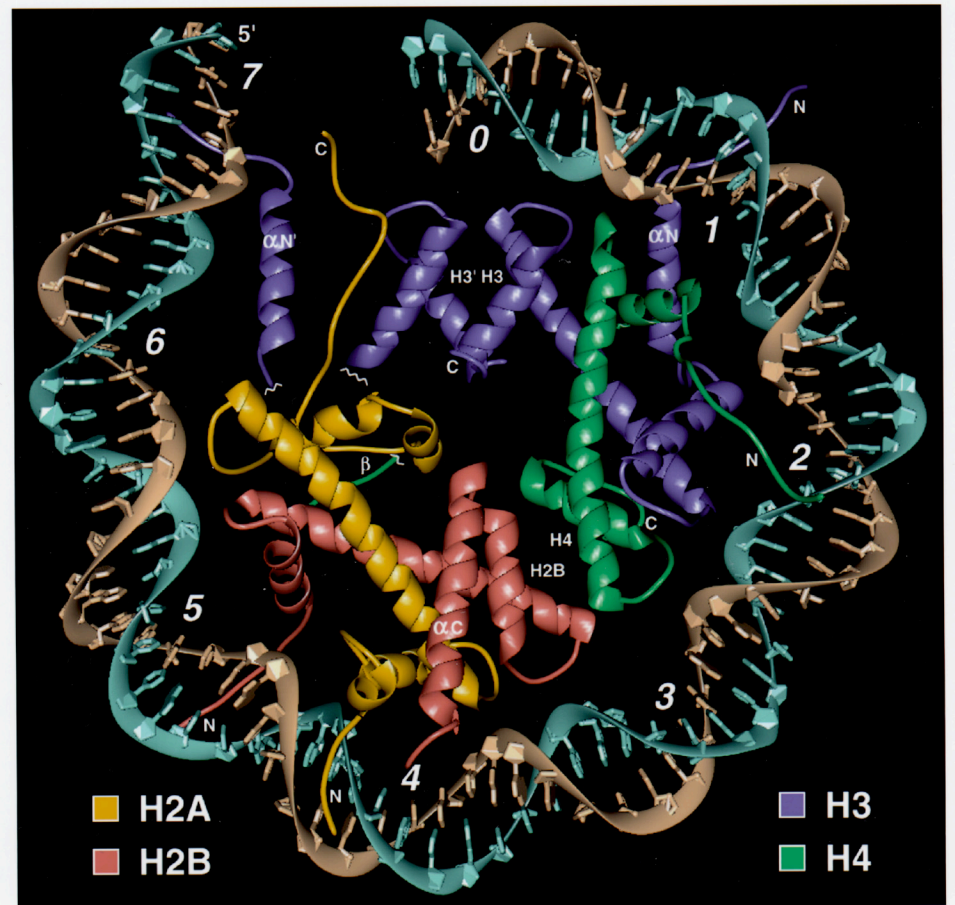
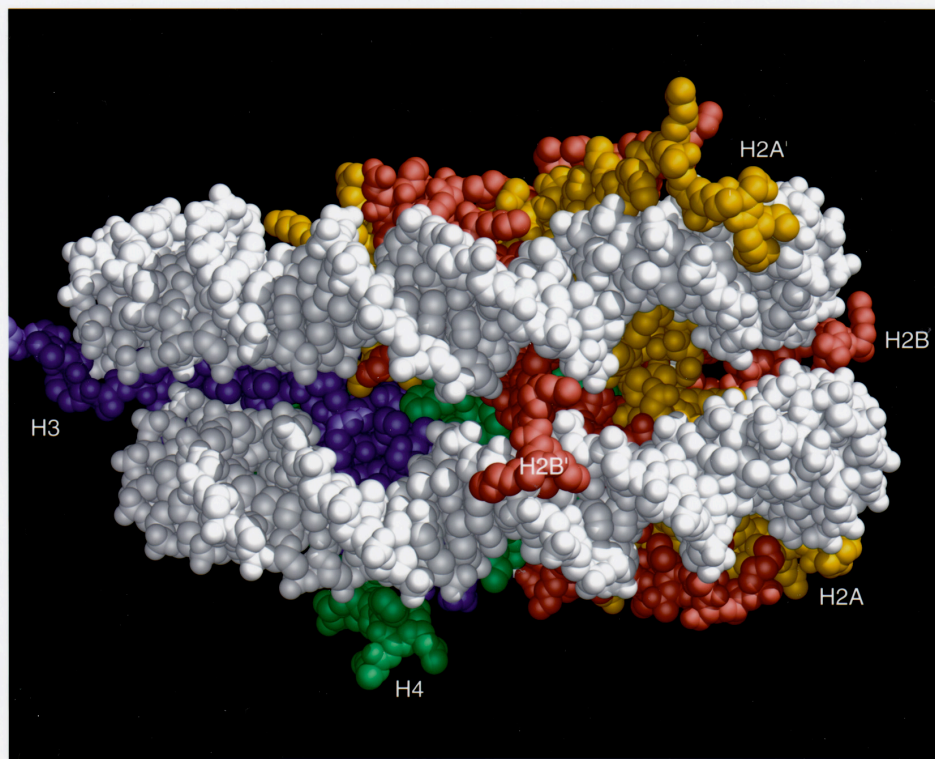
•94 bp loops



Rapid spontaneous looping of 94 bp DNAs

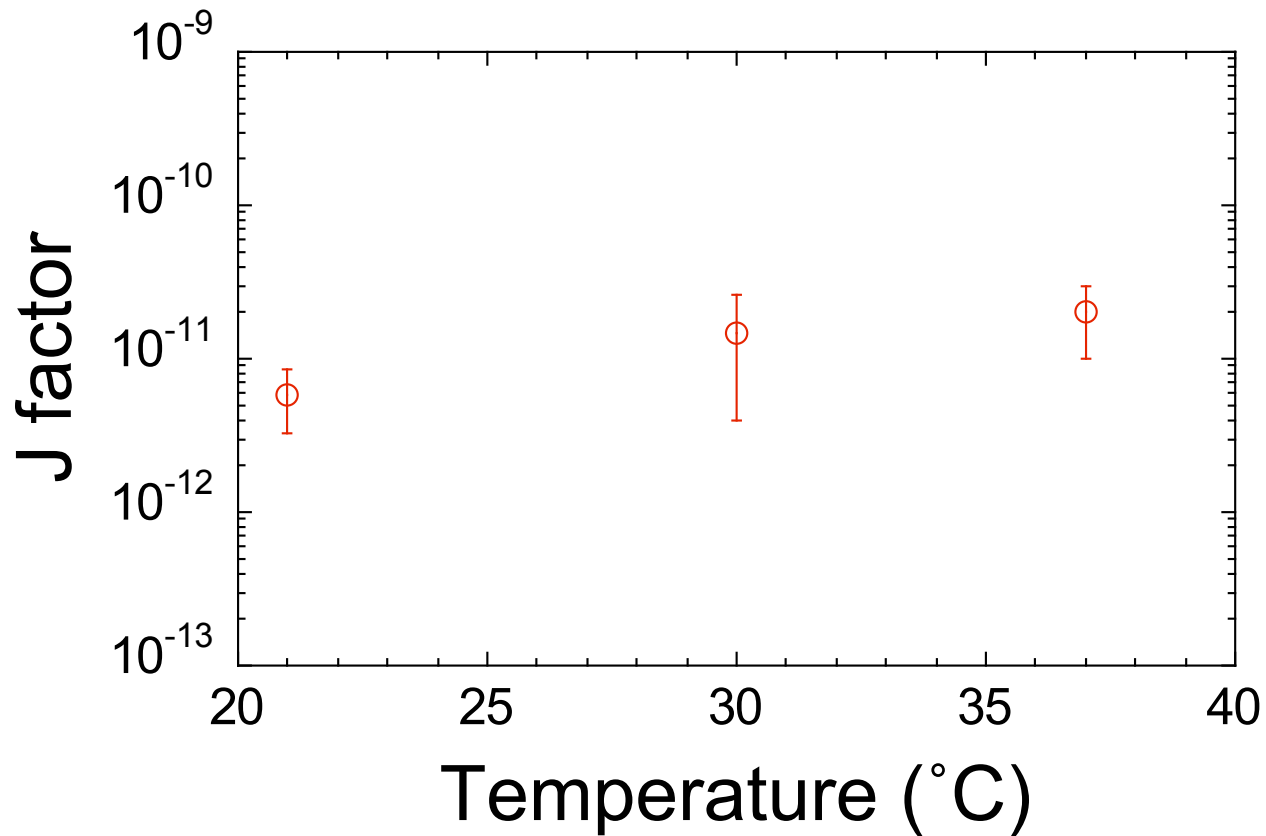


Structural basis of sharply looped protein–DNA complexes

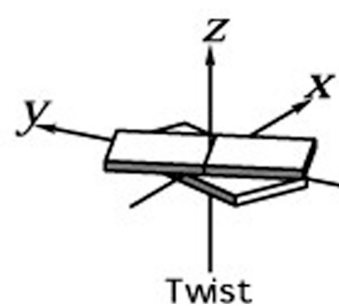
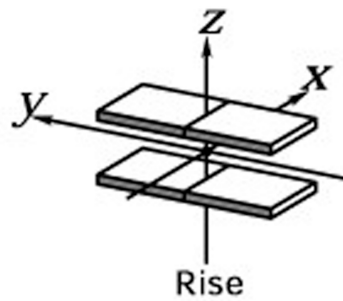
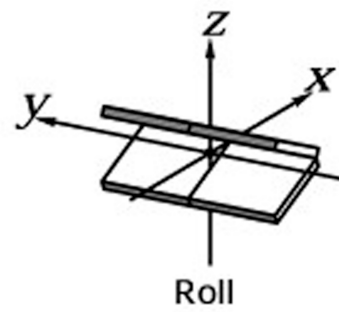
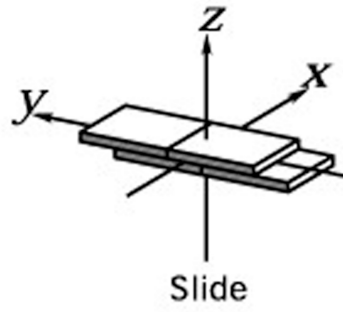
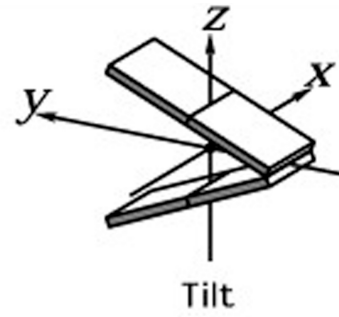
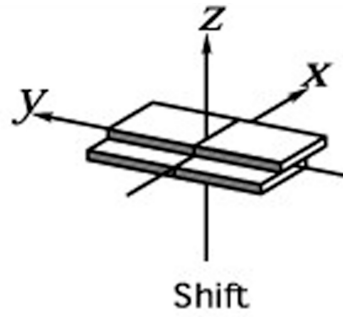


Luger et al., 1997
Richmond & Davey, 2003

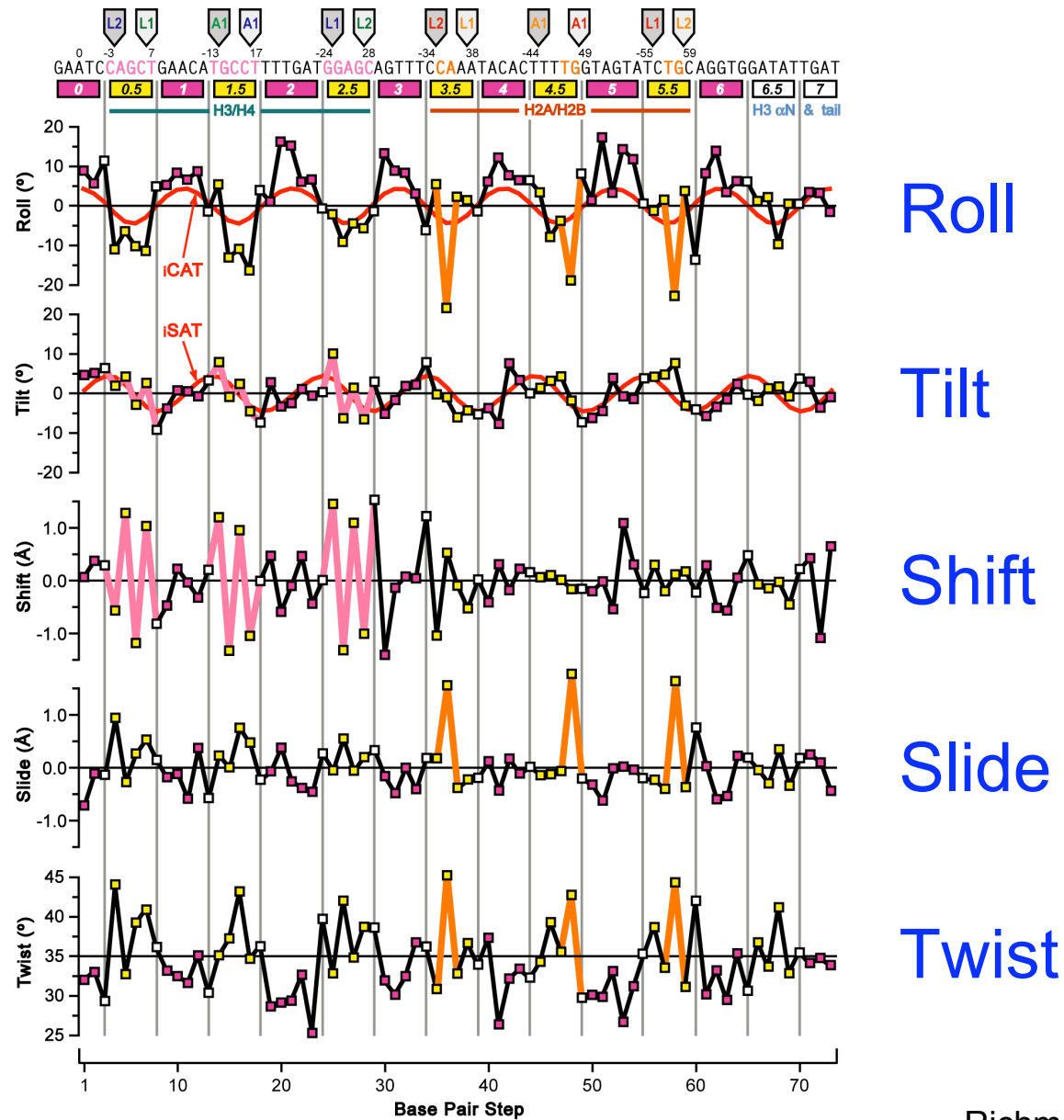
J factors are weakly dependent on temperature



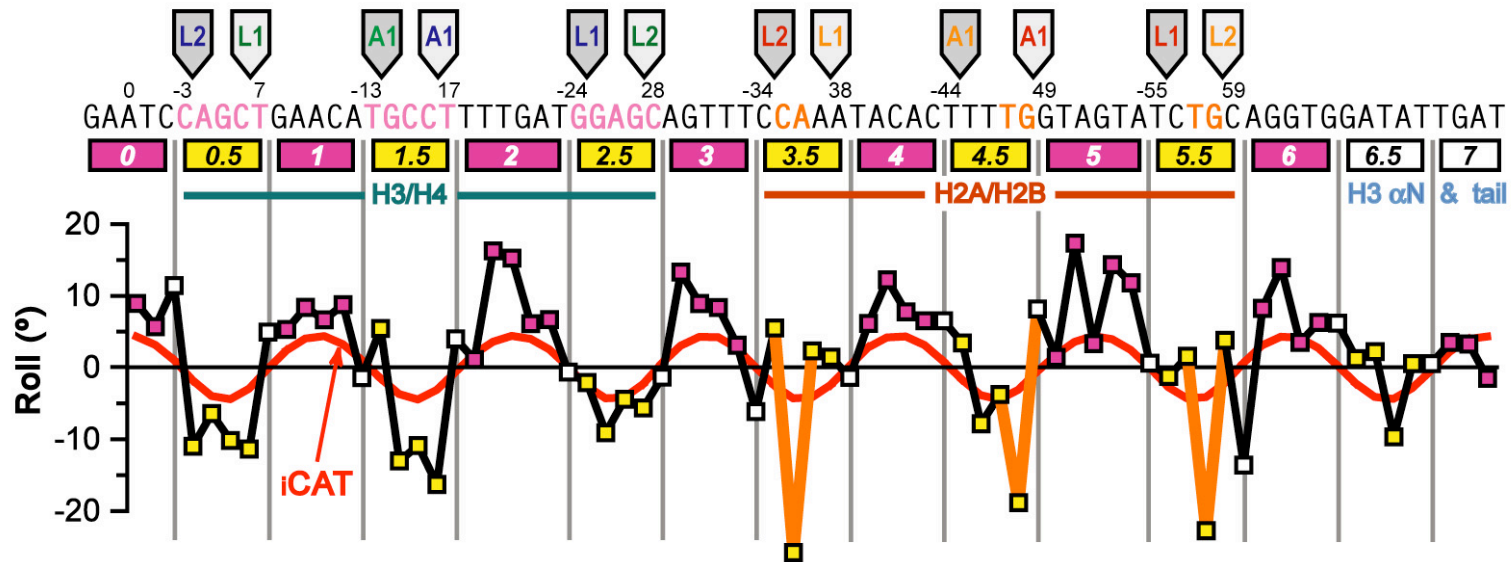
Basepair steps as fundamental units of DNA mechanics



Correlated deformations for sharp DNA wrapping



Structural basis of sharply looped protein–DNA complexes



- Small distortions, and localized larger distortions along the full wrapped DNA length

Acknowledgements

Tim Cloutier

Julie Dohm

Karissa Fortney

Dan Grilley