

The poly dA helix: a new structural motif for high performance DNA-based molecular switches

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ABSTRACT

We report a pH-dependent conformational transition in short, defined homopolymeric deoxyadenosines (dA₁₅) from a single helical structure with stacked nucleobases at neutral pH to a double-helical, parallel-stranded duplex held together by AH...H+A base pairs at acidic pH. Using native PAGE, 2D NMR, circular dichroism (CD) and fluorescence spectroscopy, we have characterized the two different pH dependent forms of dA₁₅. The pH-triggered transition between the two defined helical forms of dA₁₅ is characterized by CD and fluorescence. The kinetics of this conformational switch is found to occur on a millisecond time scale. This robust, highly reversible, pH-induced transition between the two well-defined structured states of dA₁₅ represents a new molecular building block for the construction of quick-response, pH-switchable architectures in structural DNA nanotechnology.

INTRODUCTION

Structural DNA nanotechnology is an emerging field that uses DNA to create either rigid architectures or dynamic switches (1–4). Dynamic, DNA-based nanodevices may also be described as molecular switches. They are based on structural transitions between two well-defined conformations of DNA upon the application of a stimulus. Several devices have been developed based on B-DNA assemblies employing differential hybridization of complementary strands, metal ions and indeed protons (5–11). Here we describe the poly dA helix as a new structural motif that functions as a molecular switch, which at low pH forms a parallel-stranded double helix and at neutral pH exists as a structured, single helix.

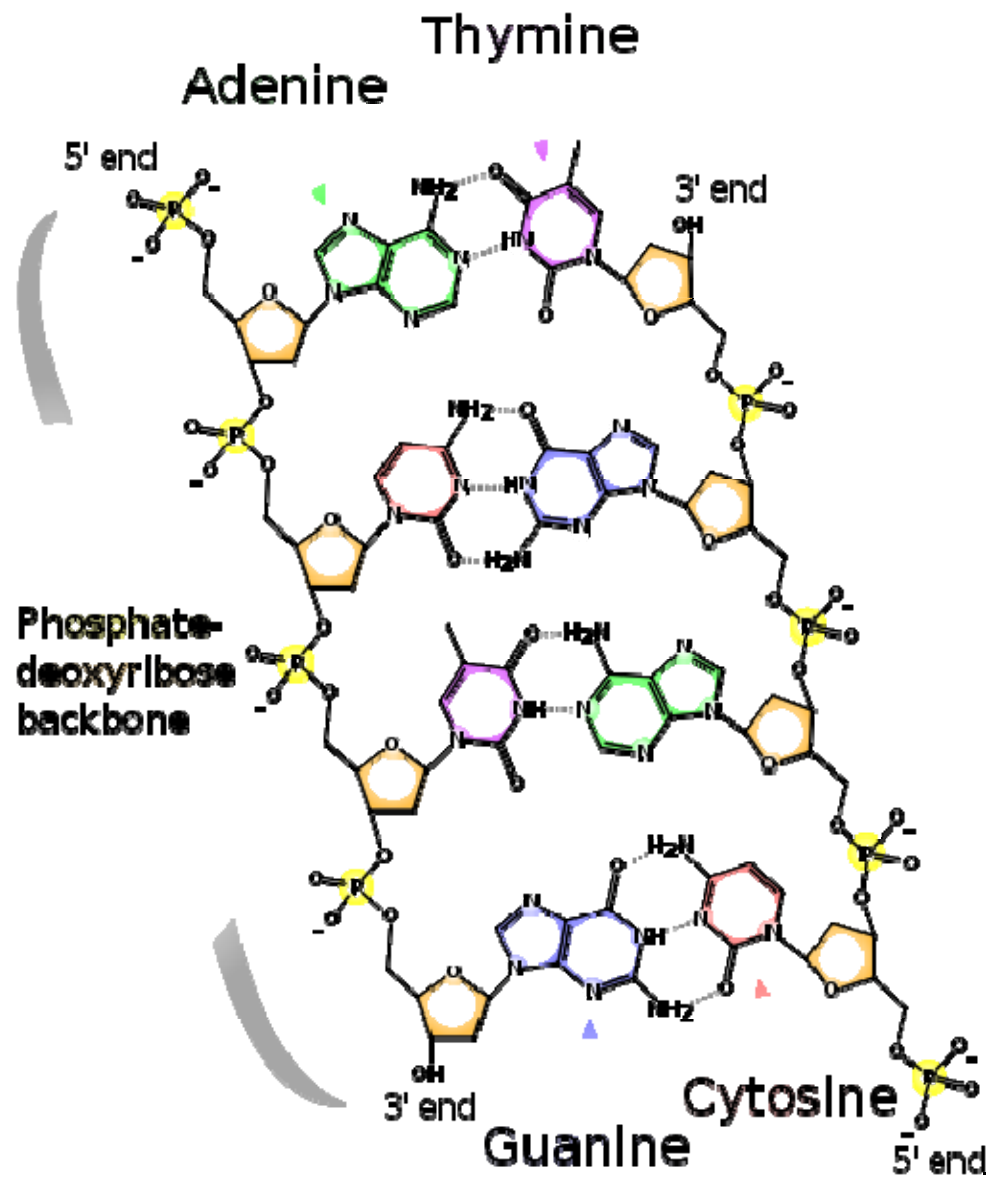
Early studies on understanding the structure, base-pairing scheme and base stacking properties of DNA and RNA duplexes used synthetic homopolymeric DNA and RNA as they were considered simplified model systems. Eventually it was found that these

synthetic homopolymers actually formed different unusual conformations involving non-Watson-Crick base pairing. Poly rC and poly dC formed i-tetraplexes (12,13), while poly rG and poly dG formed G-quadruplexes (14–16). Interestingly, poly rA formed a parallel-stranded double helix, called pi-helix at acidic pH due to N1 protonation of the adenines at pH < 5 (17–19). At neutral pH poly rA was found to exist as a single, right-handed helix with nine nucleotides per pitch of 25.4 Å (20,21). In fact, characteristic of the distinct nature of this helix, there are proteins called poly rA binding proteins (PABPs) that specifically bind poly rA over any random ssRNA (22,23). At neutral pH poly dA is known to exist as a structured single helix, similar to poly rA (24–27), except that the nucleobases in poly dA are more strongly stacked than in poly rA and are in the C2'-endo configuration. However, the behavior of poly dA at acidic pH is still unknown. We were encouraged by the fact that poly rA could form these structures, with no indication of any special role for the 2'OH. Further we also found a sprinkling of short DNA sequences that had an over-representation of adenines that formed parallel duplexes at acidic pH (28–30), all of which contained A–A base pairs. But, these sequences would be expected to exist as unstructured single strands at neutral pH.

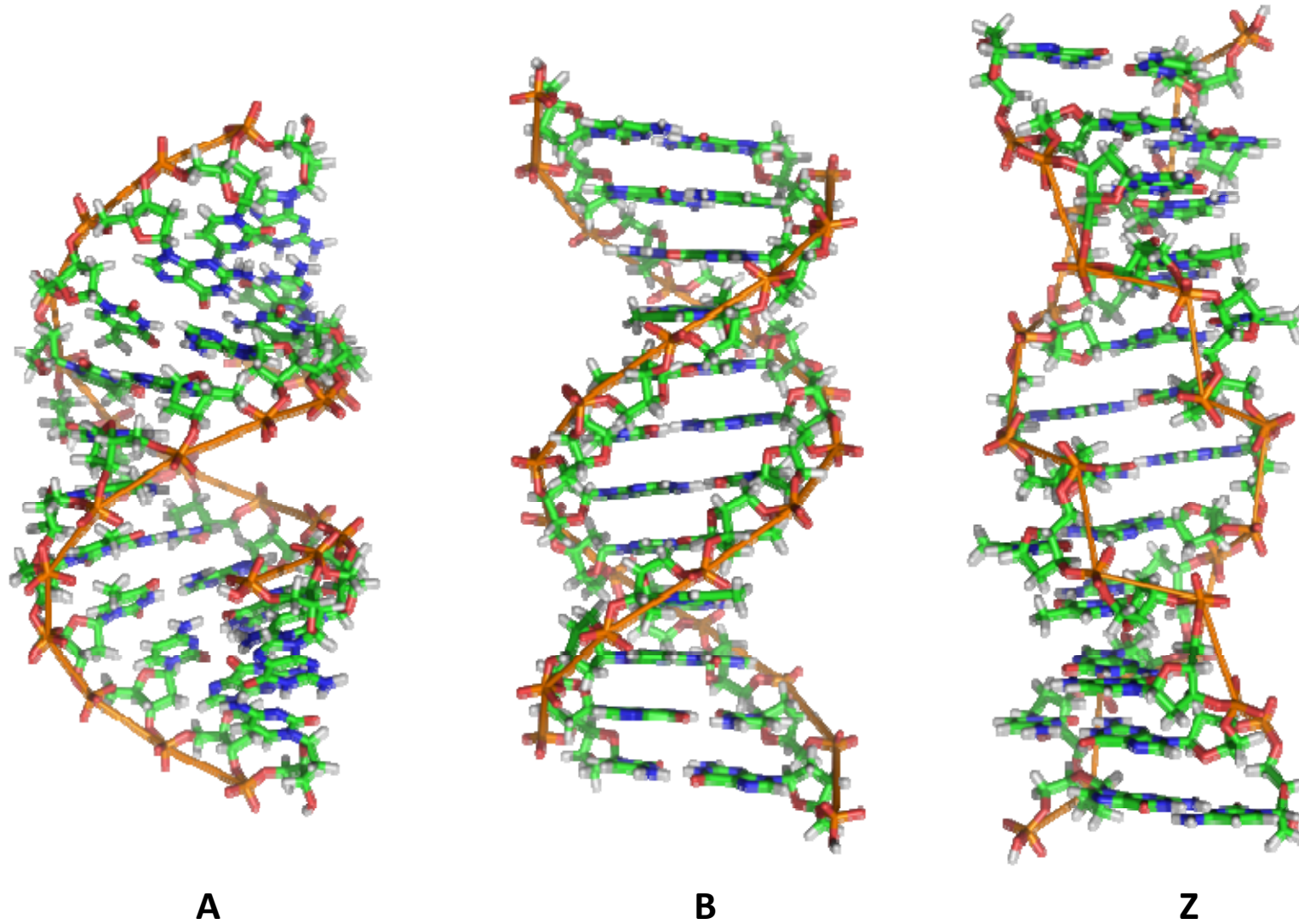
We have been interested in developing alternative, non-B-DNA building blocks that rely on non-Watson-Crick base pairing, for applications in structural DNA nanotechnology (31–35). Given that poly rA exists as a right-handed, parallel-stranded, double helix at acidic pH (17) and a structured right-handed single helix at neutral pH we reasoned that poly dA may have potential as a new building block for DNA based pH-switches if it is able to recapitulate poly rA behavior. In order to see whether poly dA alone could form a duplex at acidic pH and if so, could it switch reversibly between its structured single helical state at pH 7 to a structured duplex at acidic pH, we investigated a segment of poly dA. We chose a segment of poly dA 15 nucleotides long, because this is within the limits of the observed persistence length of the poly dA single helix (36). Using gel electrophoresis, circular dichroism (CD) spectroscopy and concentration dependent

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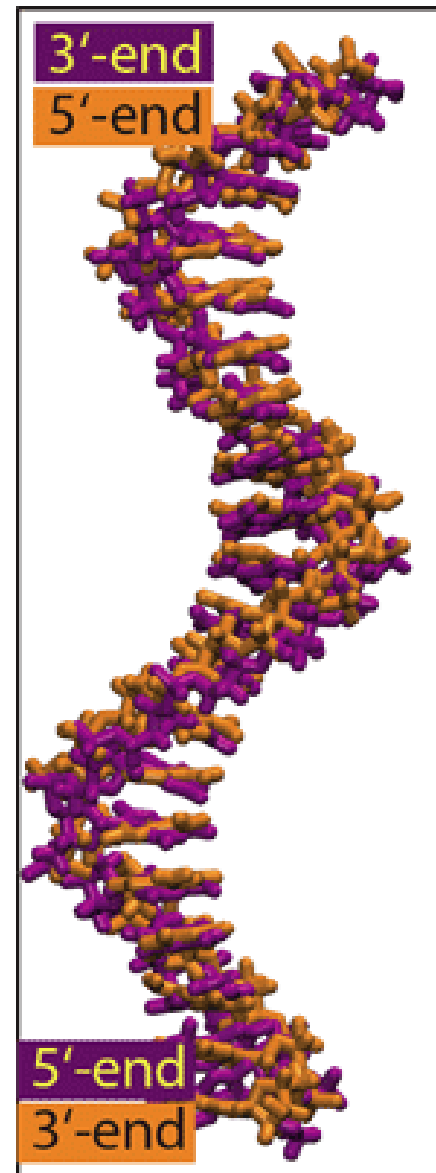
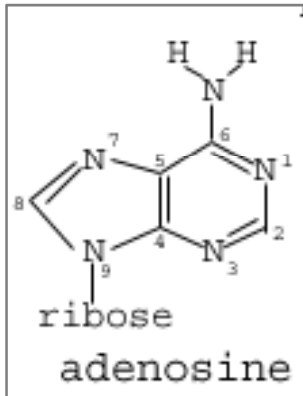
DNA



DNA helical structure



PolydA



http://images.google.com/imgres?imgurl=http://www.ks.uiuc.edu/Research/hemolysin/3versus5/PICTURES/comparison.gif&imgrefurl=http://www.ks.uiuc.edu/Research/hemolysin/3versus5/&usg=__q8FdF7xyFo6lqxUhHzRRNY2Rqg=&h=404&w=150&sz=22&hl=en&start=16&um=1&tbnid=SEAv21to sf2MDM:&tbnh=124&tbnw=46&prev=/images%3Fq%3DpolydA%2Bstructure%26hl%3Den%26rls%3Dcom.microsoft:en-US%26sa%3DG%26um%3D1

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We report a pH-dependent conformational transition in short, defined homopolymeric deoxyadenosines (dA₁₅) from a single helical structure with stacked nucleobases at neutral pH to a double-helical, parallel-stranded duplex held together by AH⁺-H⁺A base pairs at acidic pH. Using native PAGE, 2D NMR, circular dichroism (CD) and fluorescence spectroscopy, we have characterized the two different pH dependent forms of dA₁₅. The pH-triggered transition between the two defined helical forms of dA₁₅ is characterized by CD and fluorescence. The kinetics of this conformational switch is found to occur on a millisecond time scale. This robust, highly reversible, pH-induced transition between the two well-defined structured states of dA₁₅ represents a new molecular building block for the construction of quick-response, pH-switchable architectures in structural DNA nanotechnology.

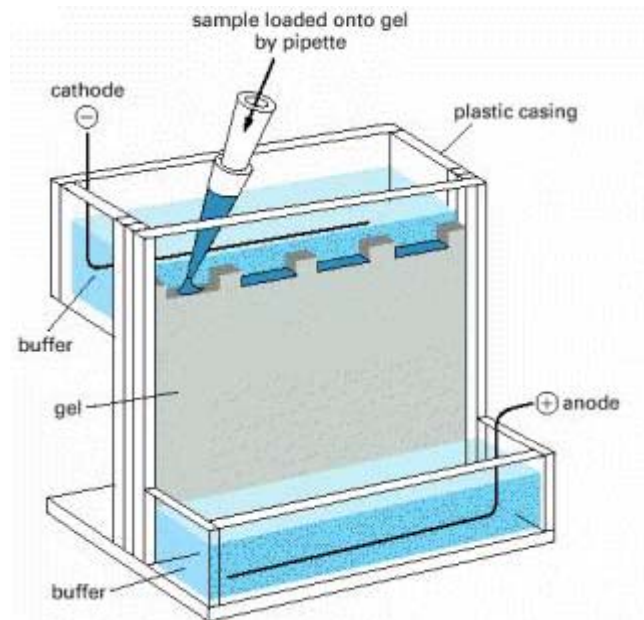
synthetic homopolymers actually formed different conformations involving non-Watson-Crick base pairs. Poly rC and poly dC formed i-tetraplexes (12,13), poly rG and poly dG formed G-quadruplexes (14,15). Interestingly, poly rA formed a parallel-stranded helix, called pi-helix at acidic pH due to N1 protonation of the adenines at pH <5 (17–19). At neutral pH poly rA was found to exist as a single, right-handed helix with 11 nucleotides per pitch of 25.4 Å (20,21). In fact, characteristic of the distinct nature of this helix, there are proteins called poly rA binding proteins (PABPs) that specifically bind poly rA over any random ssRNA (22,23). At neutral pH poly dA is known to exist as a structured single helix similar to poly rA (24–27), except that the nucleobases in poly dA are more strongly stacked than in poly rA and are in the C2'-endo configuration. However, the behavior of poly dA at acidic pH is still unknown. We were intrigued by the fact that poly rA could form these structures with no indication of any special role for the adenine. Further we also found a sprinkling of short sequences that had an over-representation of ad

DNA

Table 1. Poly dA sequences used in this study

Name	Sequence
Poly dA ₁₅	5'-d(AAAAAAAAAAAAAAAAAA)-3'
dTA6	5'-d(TAAAAAA)-3'
3'-Dabcyl-dA ₁₅	5'-d(AAAAAAAAAAAAAAAAAA)-Dabcyl-3'
3'-TMR-dA ₁₅	5'-d(AAAAAAAAAAAAAAAAAA)-TMR-3'
5'-TAMRA-dA ₁₅	5'-TAMRA-d(AAAAAAAAAAAAAAAAAA)-3'

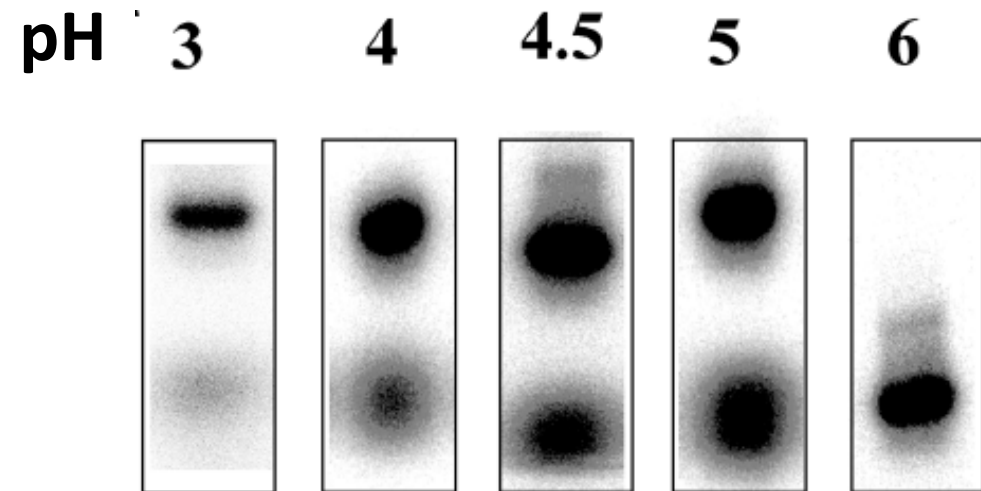
Gel-electrophoresis



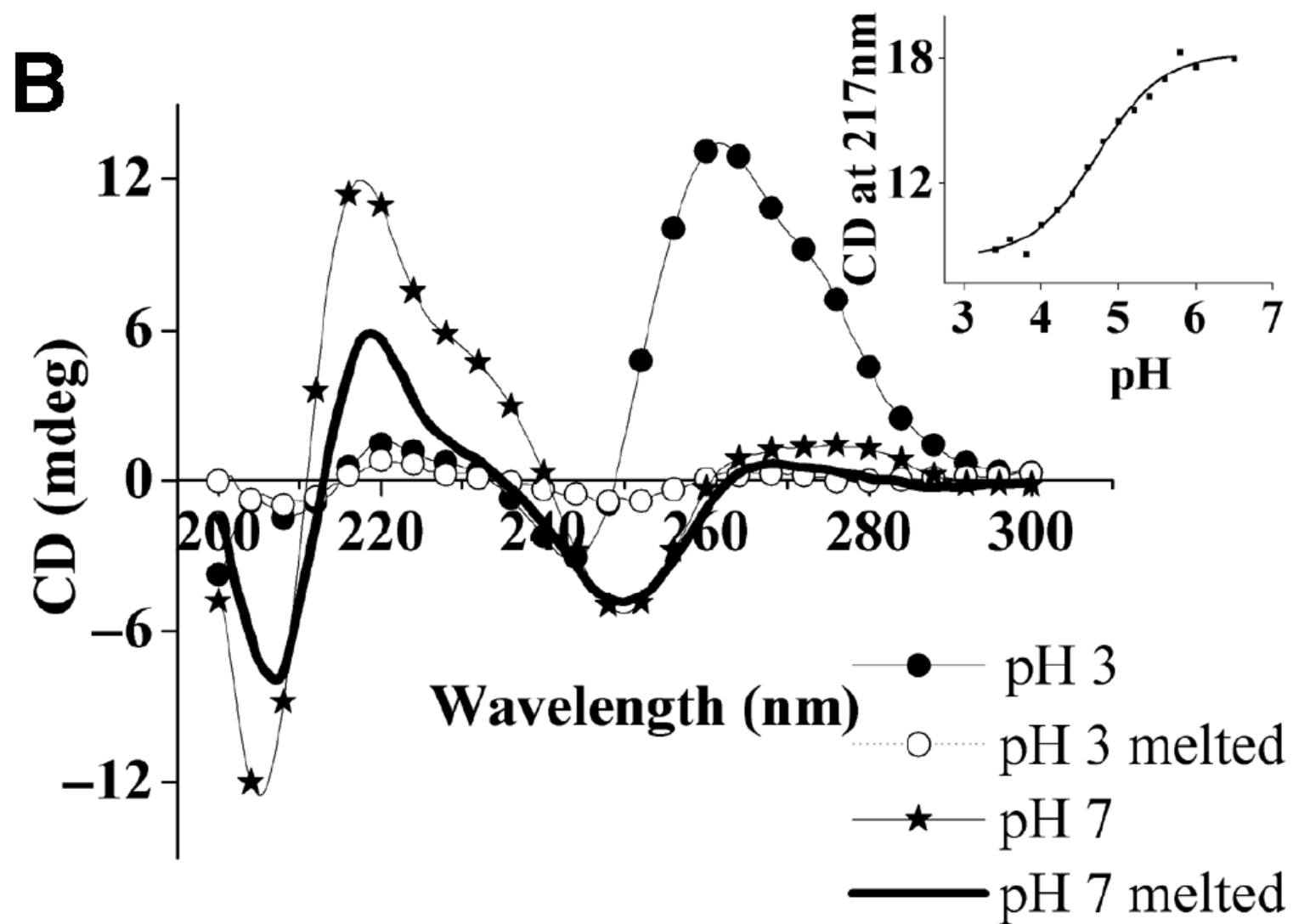
An Electrophoresis Apparatus.

A solution of proteins is applied to indentations made in a thin gel that is held between two clear plates of plastic. An electric current is applied, and over time the proteins separate.

Poly dA₁₅

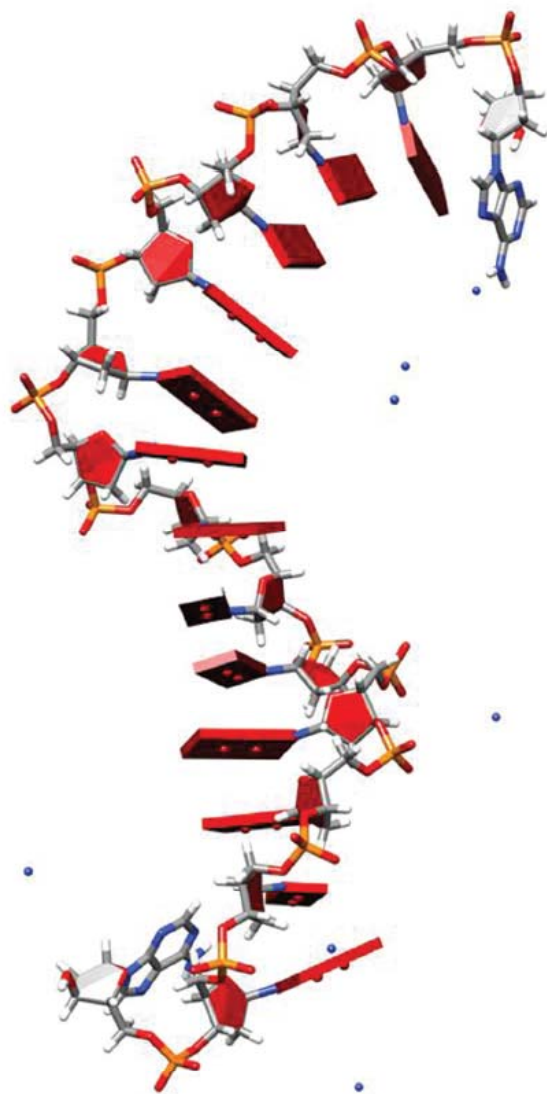


CD spectrum

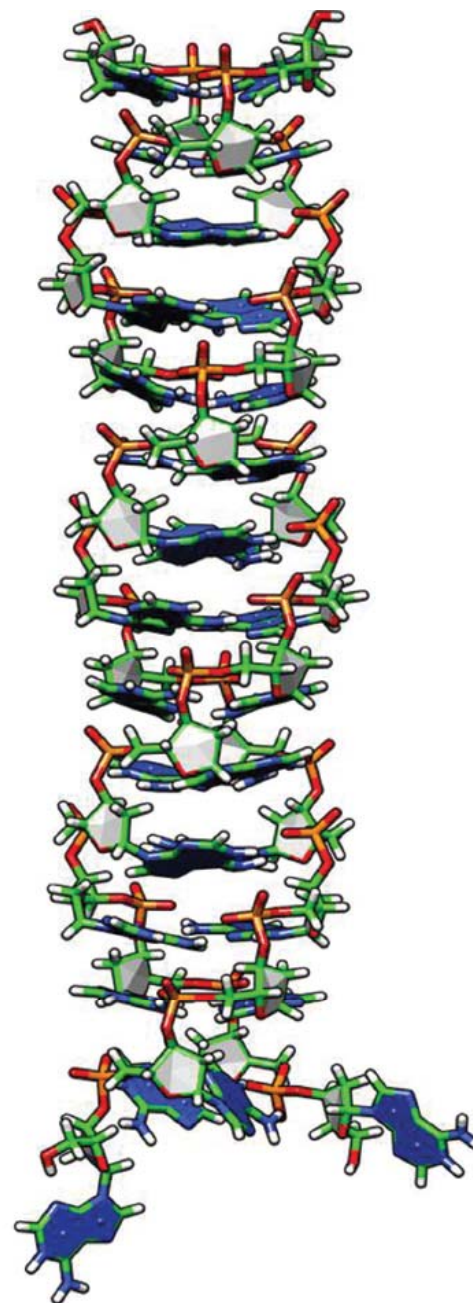


MD

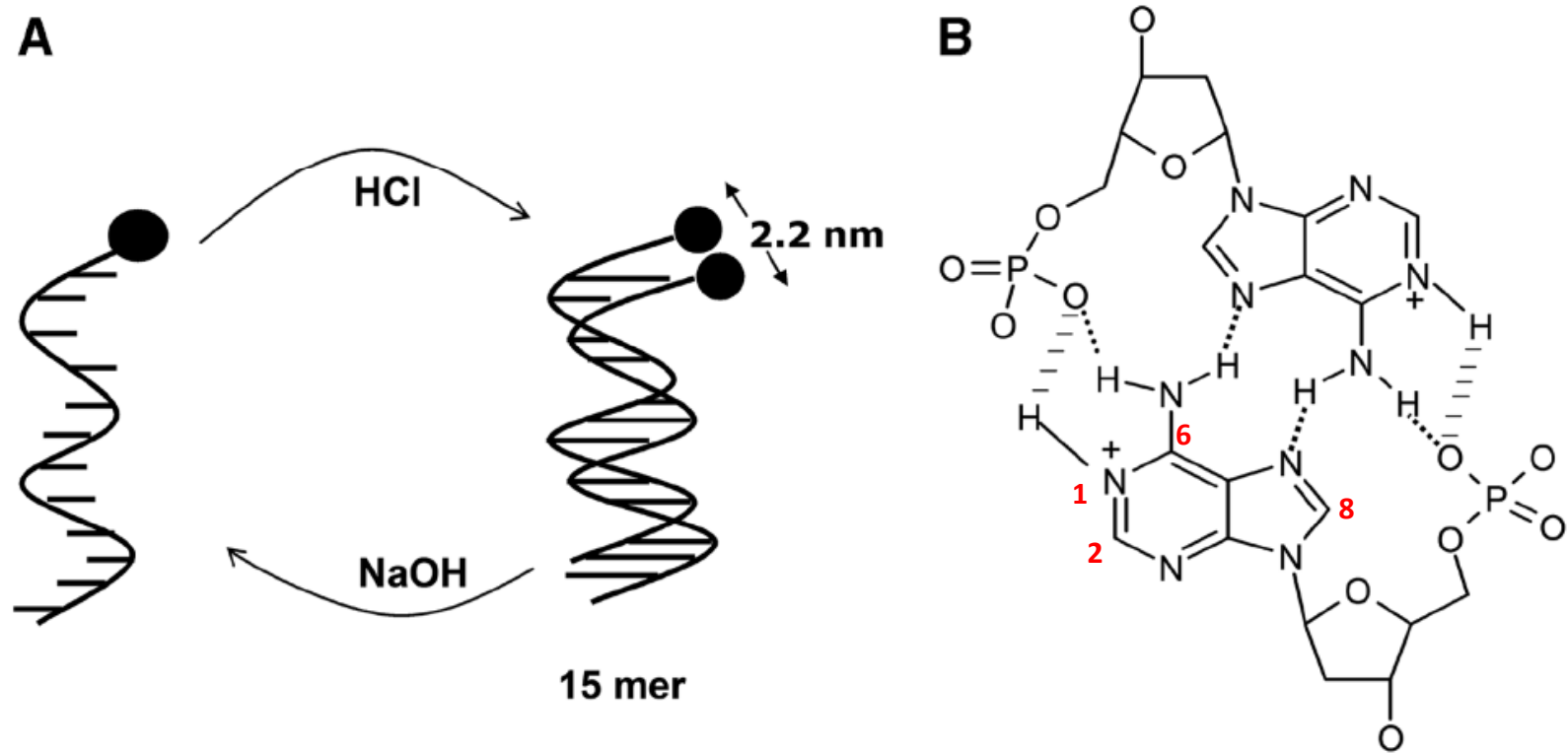
A



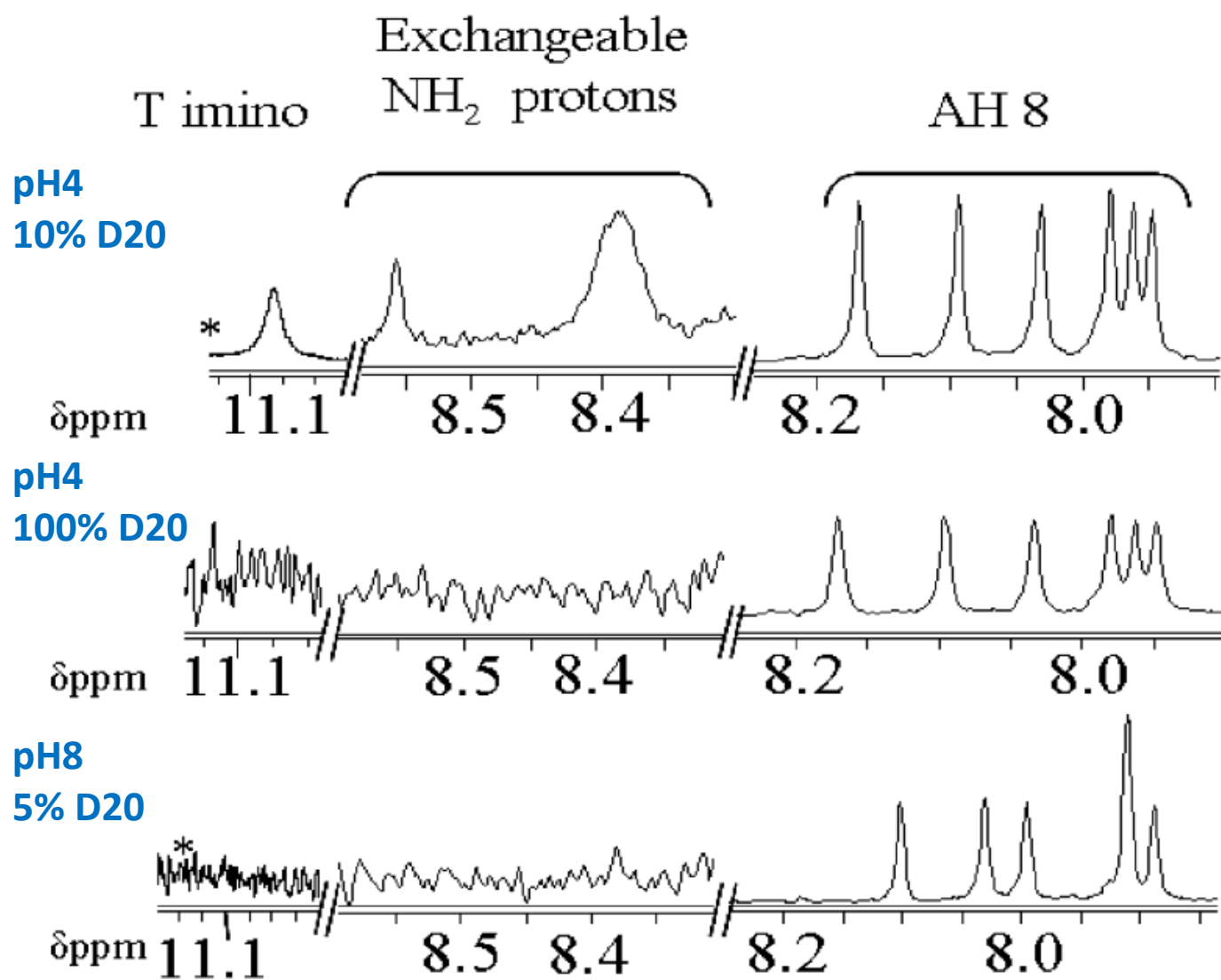
B



The scheme

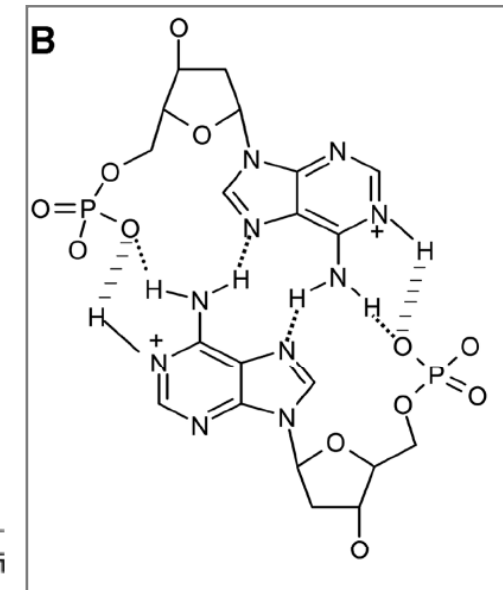
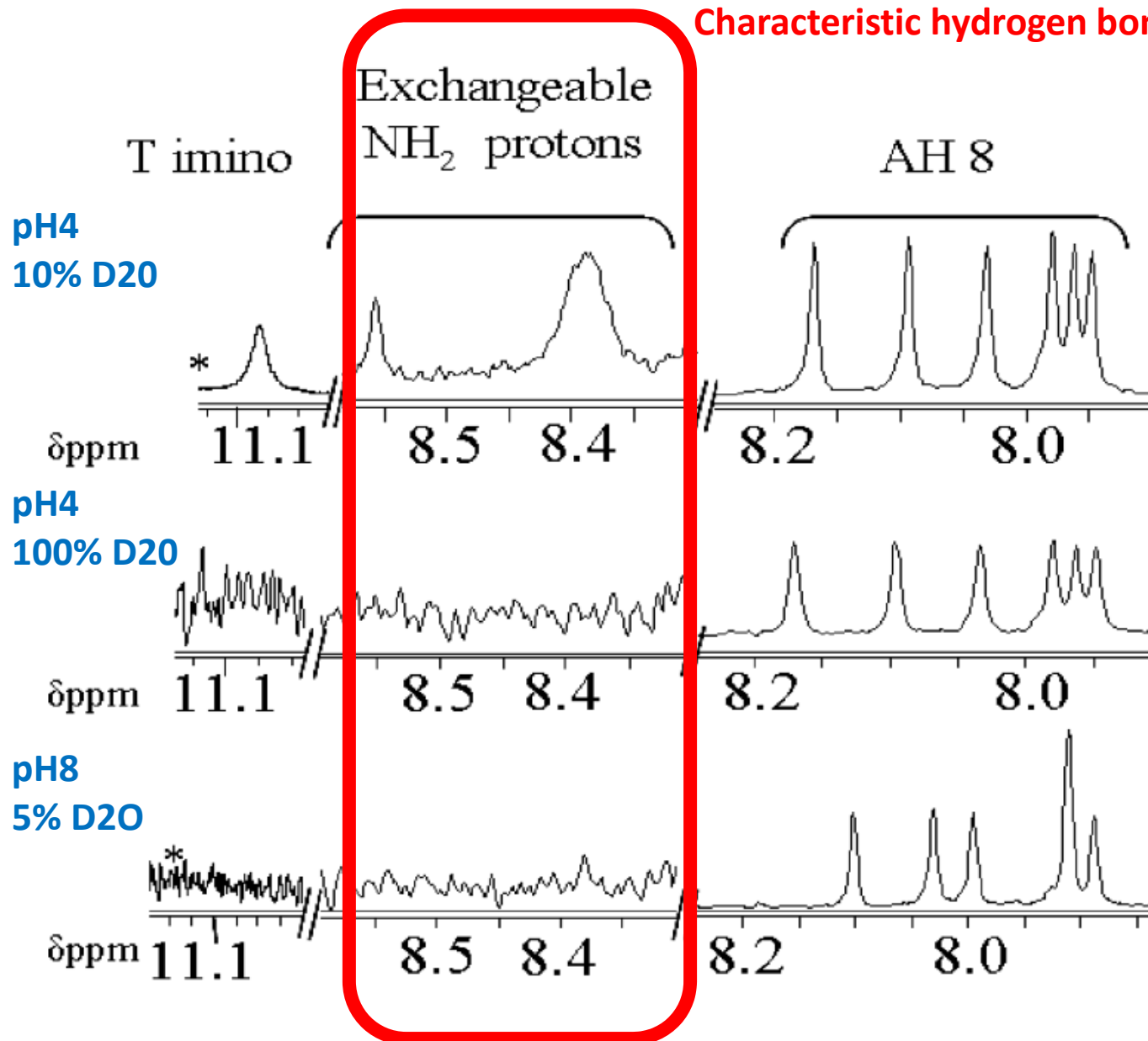


1D-NMR



1D-NMR

This NH₂ protons usually showed up at 6-7 δ ppm.
Characteristic hydrogen bonding in A-A base pairing.



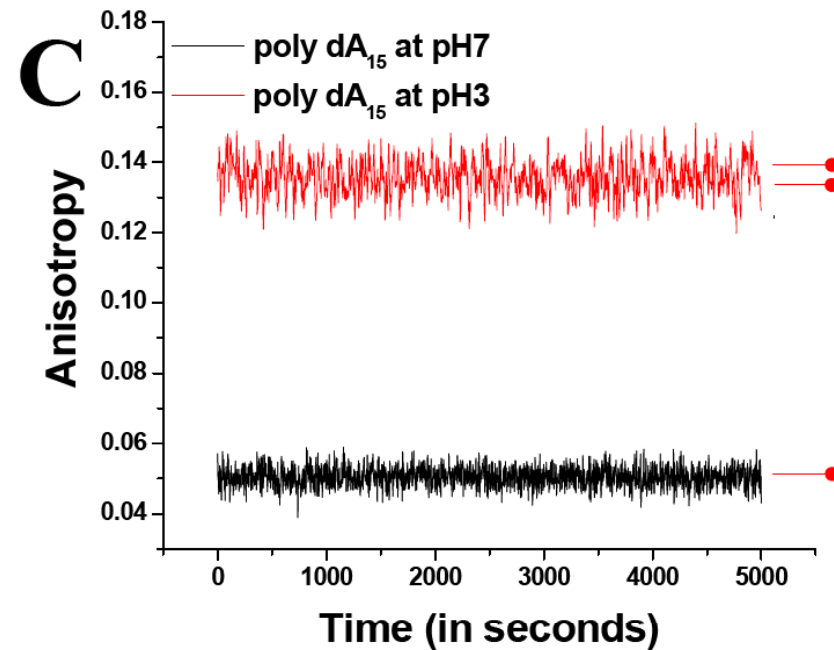
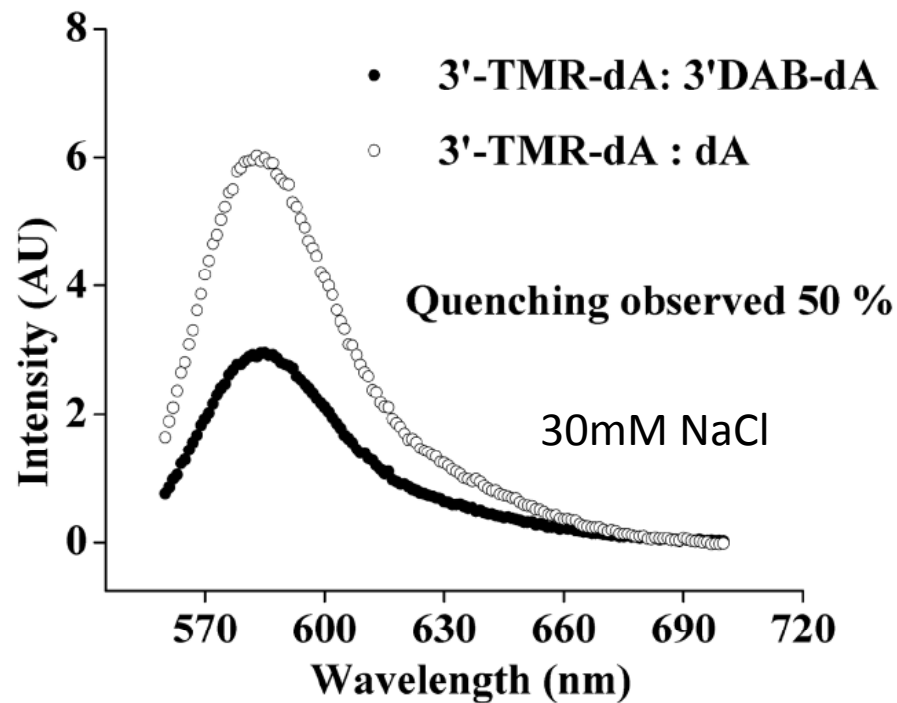
Fluorescence Quenching

3'-Dabcyl-dA₁₅

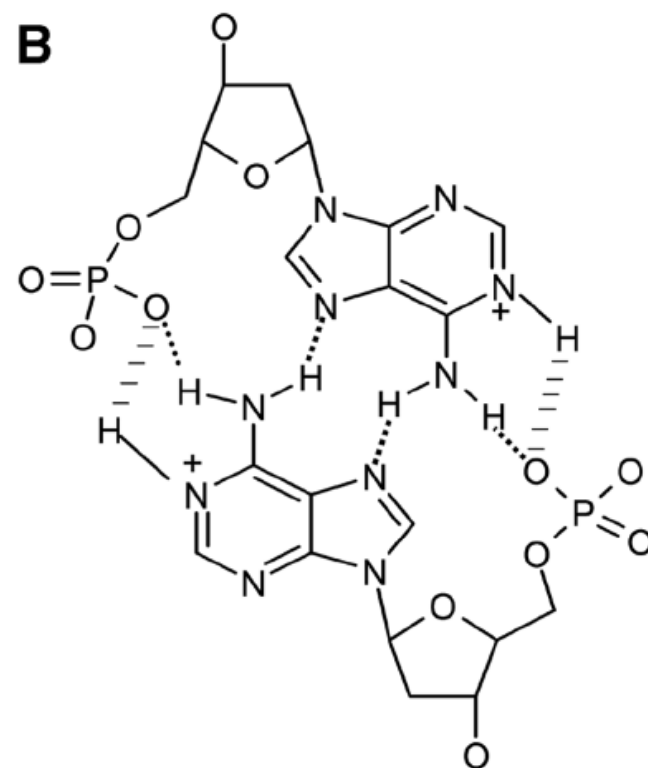
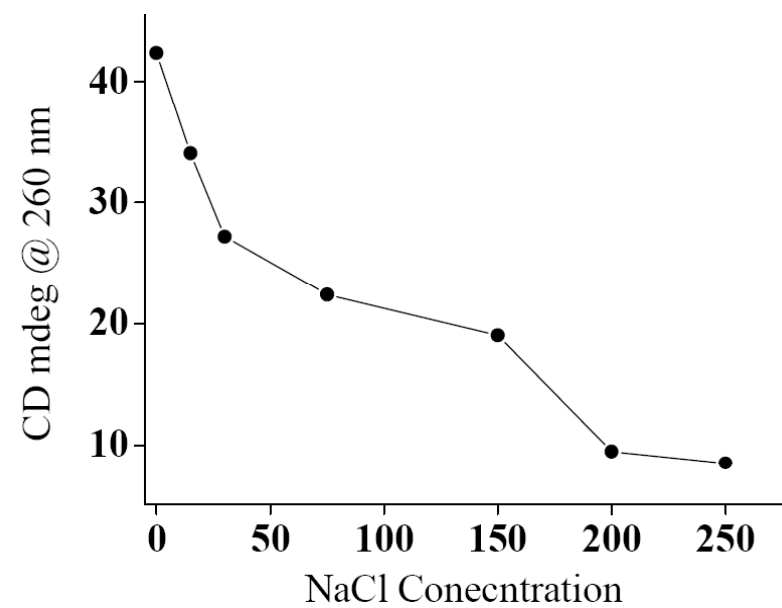
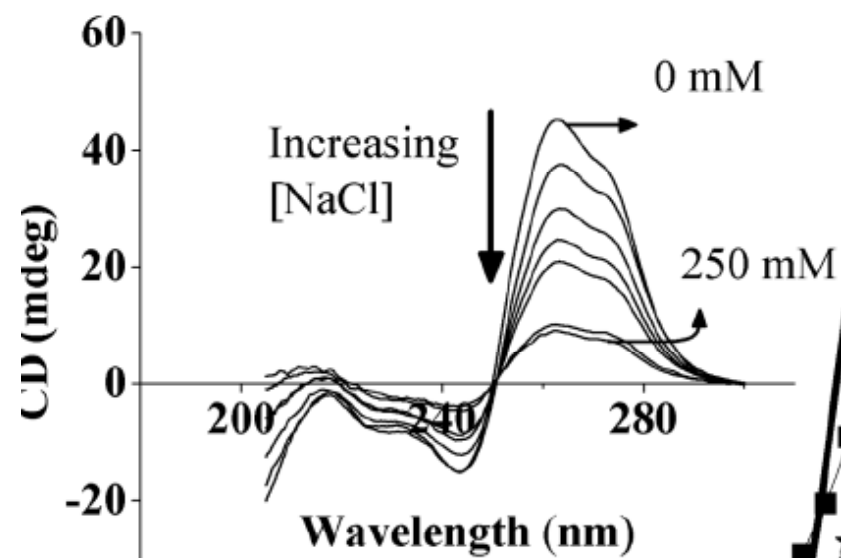
3'-TMR-dA₁₅

5'-d(AAAAAAAAAAAAAAAAAA)-Dabcyl-3'

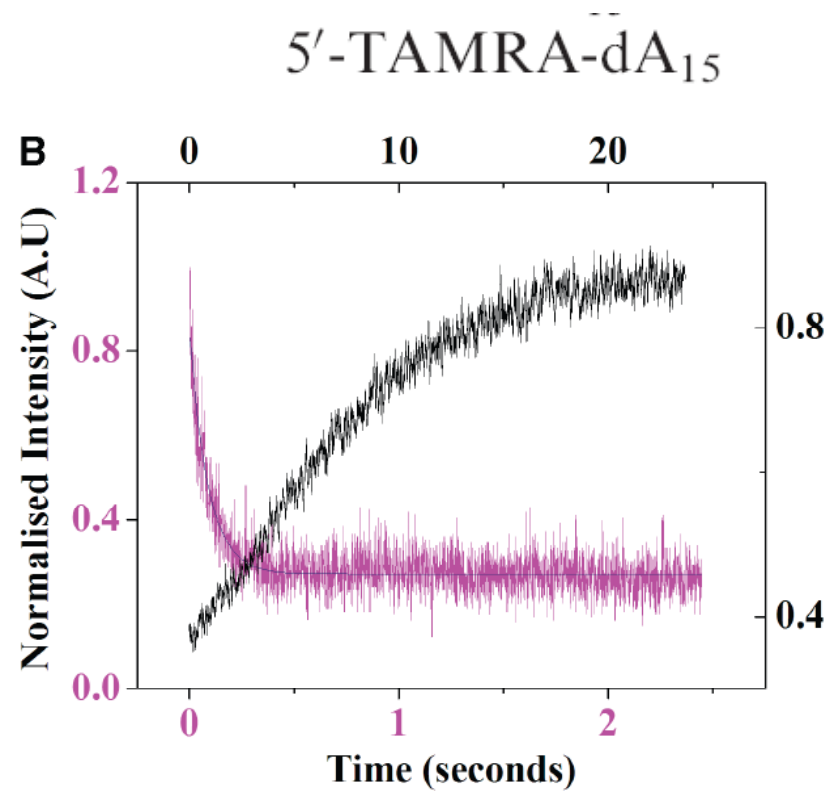
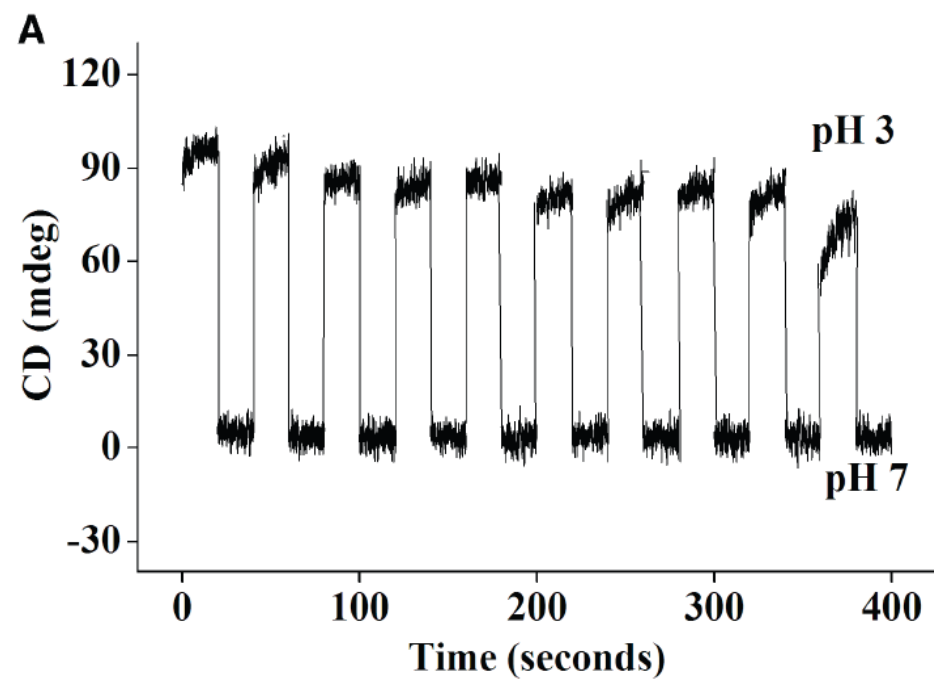
5'-d(AAAAAAAAAAAAAAAAAA)-TMR-3'



Salt dependence - CD



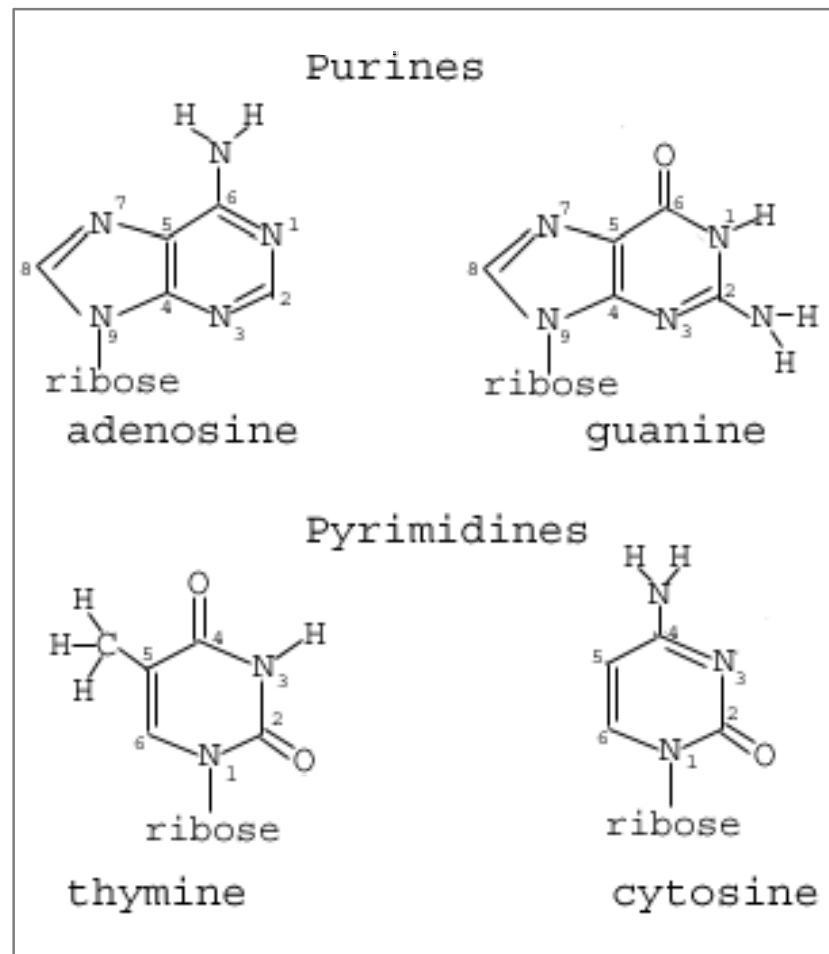
Structural transition is reversible



CONCLUSIONS

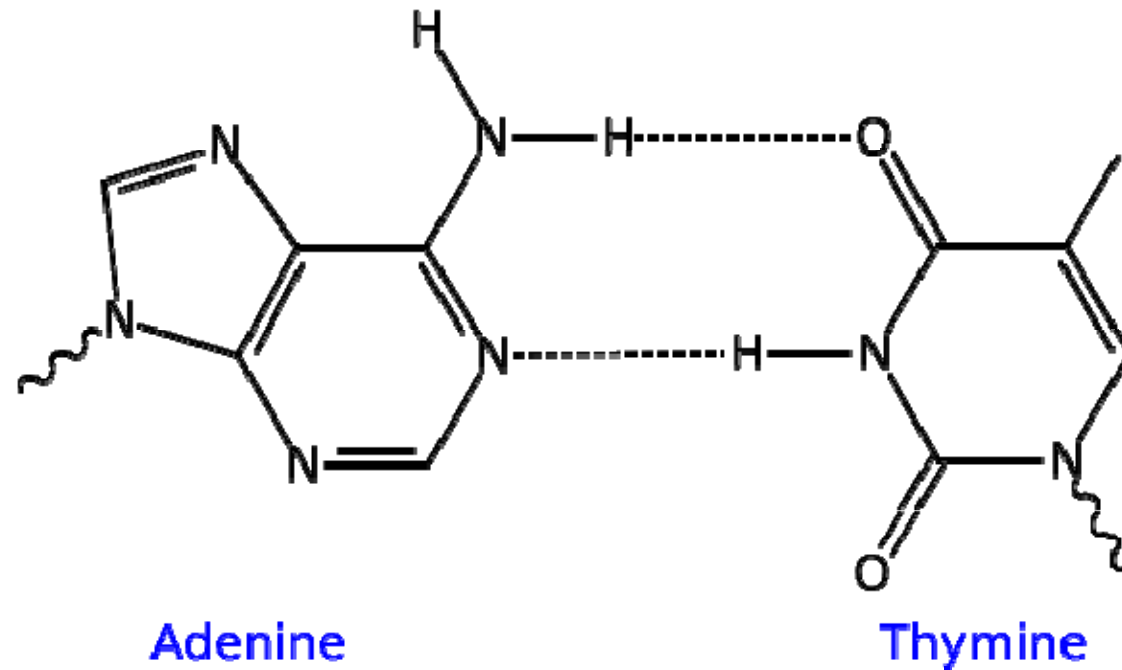
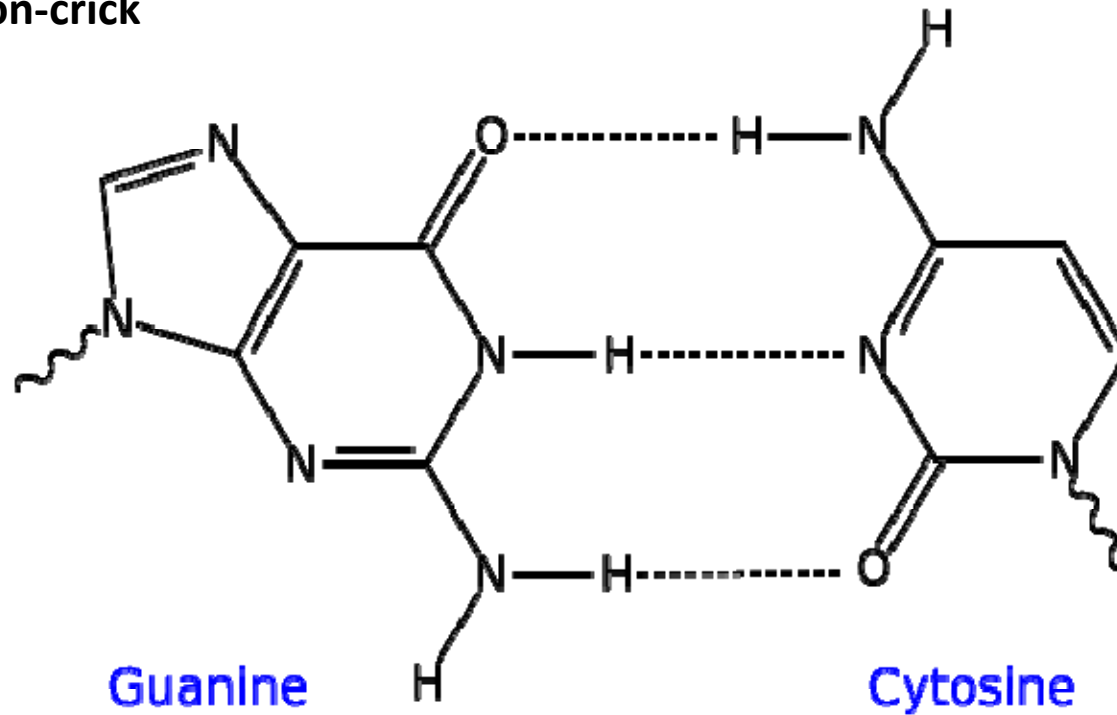
- 1) Poly dA exist as a structured single helix at neutral pH**
- 2) Poly dA forms a right-handed parallel-stranded double helix at acidic pH**
- 3) the duplex at low pH are held by Hoogsteen pairing bet protonated dA**
- 4) This can be used as a proton driven molecular switch**

Supp.



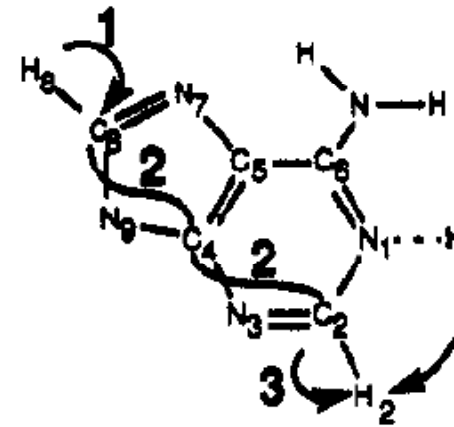
http://img.sparknotes.com/figures/7/749a4182b7527e44d289a612e420f40c/dna_bases.gif

Watson-crick

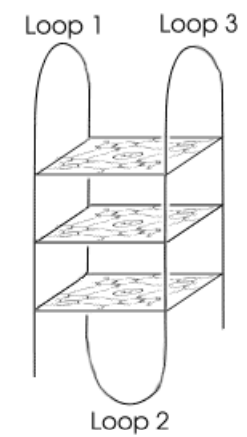
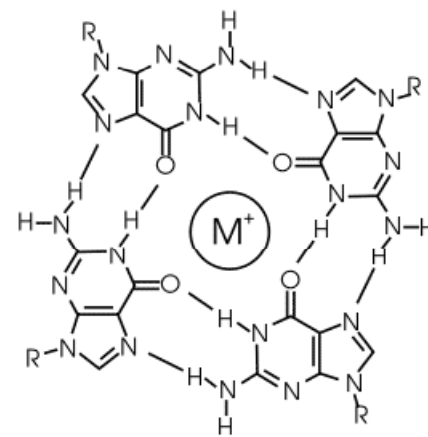
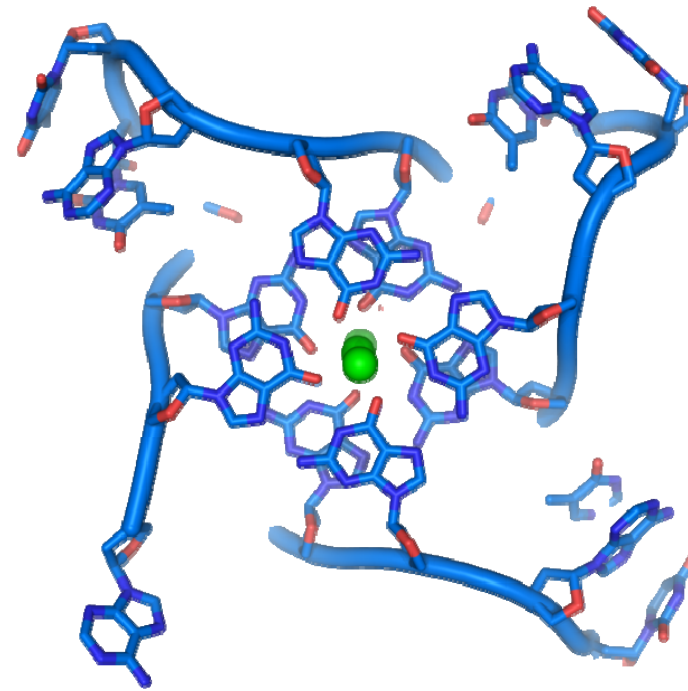
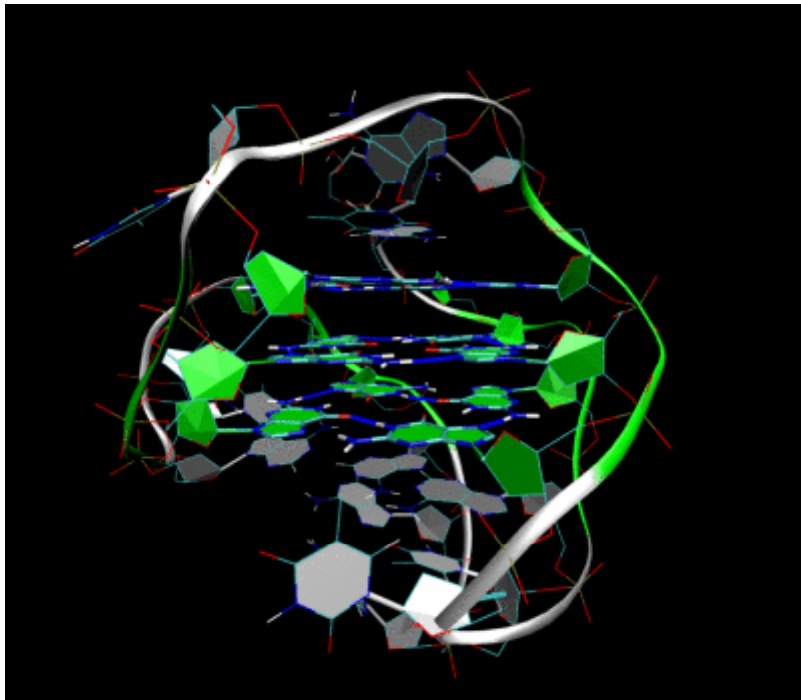


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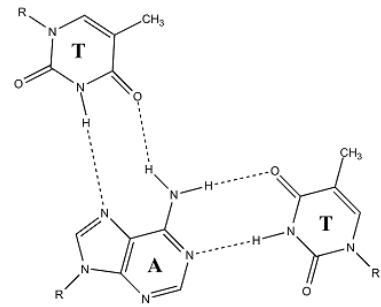
ADENINE



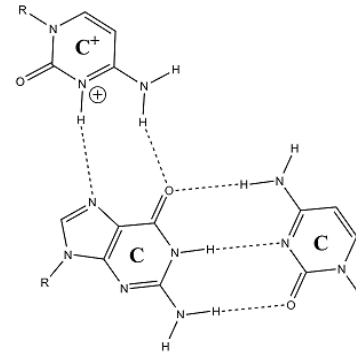
G-quadruplex



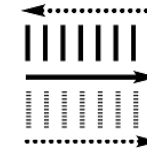
Hoogsteen base pair



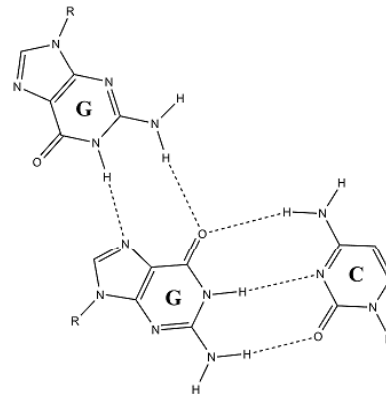
TA*T



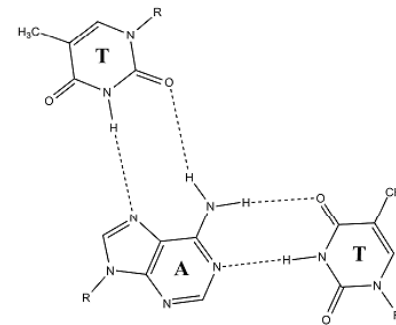
CG*C⁺



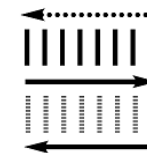
YR*Y



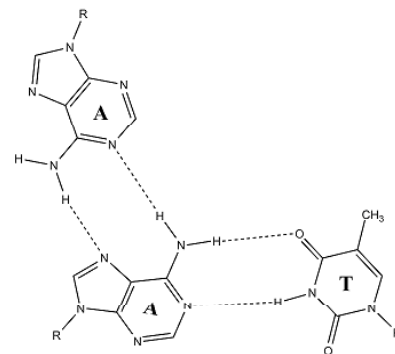
CG*G



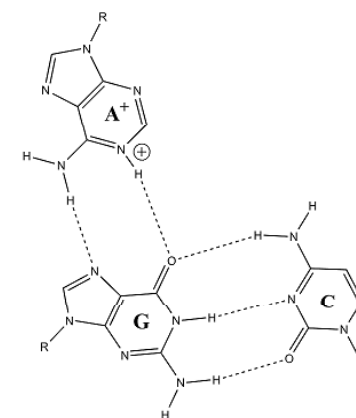
TA*A



YR*R

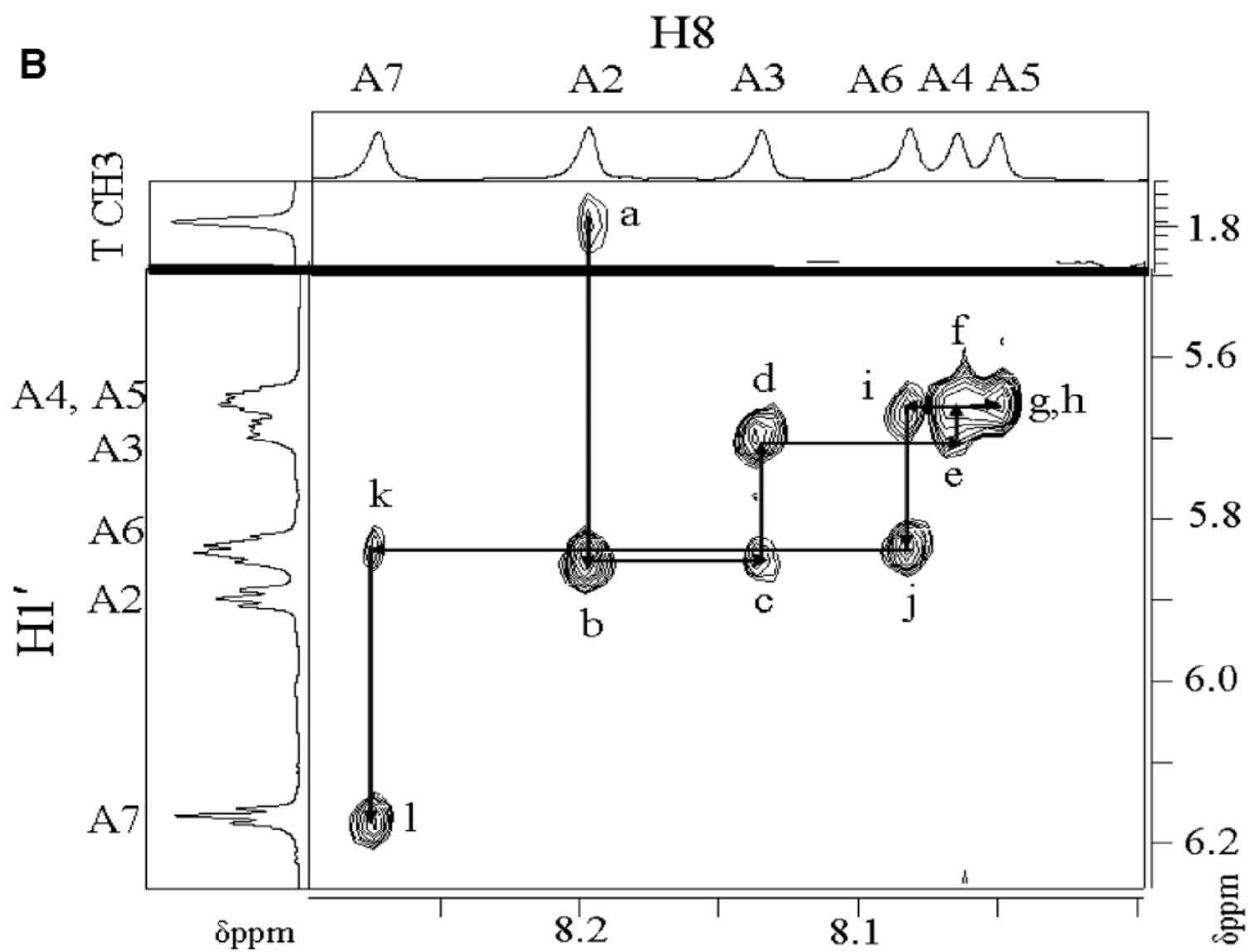


TA*T



CG*A⁺

2D-NMR



Thermal stability

