Insertion of a Bulky Rhodium Complex into a DNA Cytosine-Cytosine Mismatch: An NMR Solution Study

Christine Cordier, Valérie C. Pierre, I and Jacqueline K. Barton*

Contribution from the Division of Chemistry and Chemical Engineering, California Institute of Technology, Pasadena, California 91125

jkbarton@caltech.edu

Received May 31, 2007

Abstract:

The bulky octahedral complex Rh(bpy)₂chrysi³⁺ (chrysi = 5,6-chrysenequinonediimine) binds single-base mismatches in a DNA duplex with micromolar binding affinities and high selectivity. Here we present an NMR solution study to characterize the binding mode of this bulky metal complex with its target CC mismatch in the oligonucleotide duplex (5'-CGGA*C*TCCG-3')₂. Both NOESY and COSY studies indicate that Rh(bpy)₂chrysi³⁺ inserts deeply in the DNA at the mismatch site via the minor groove and with ejection of both destabilized cytosines into the opposite major groove. The insertion only minimally distorts the conformation of the oligonucleotide local to the binding site. Both flanking, well-matched base pairs remain tightly hydrogen-bonded to each other, and 2D DQF-COSY experiments indicate that all sugars maintain their original C₂-*endo* conformation. Remarkably, ³¹P NMR reveals that opening of the phosphate angles from a B_I to a B_{II} conformation is sufficient for insertion of the bulky metal complex. These results corroborate those obtained crystallographically and, importantly, provide structural evidence for this specific insertion mode in solution.

[Full text in html] [Full text in pdf]