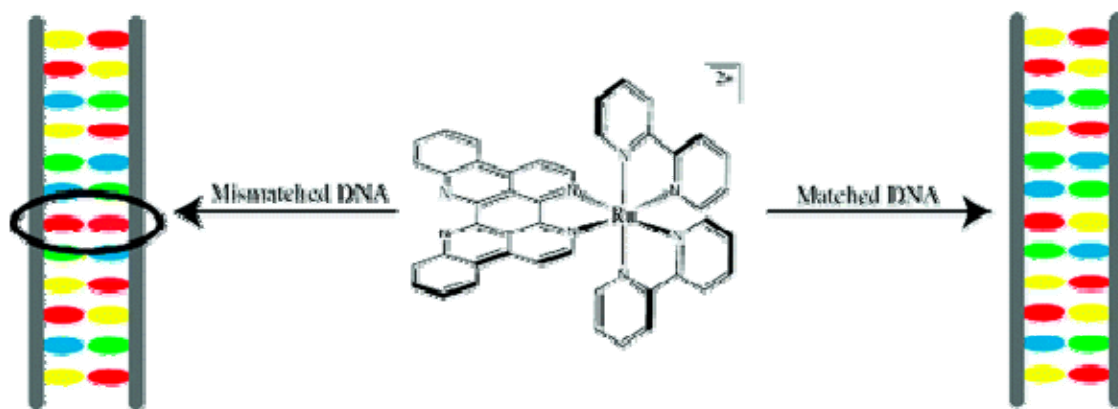


Binding of $\text{Ru}(\text{bpy})_2(\text{eilatin})^{2+}$ to Matched and Mismatched DNA

Brian M. Zeglis and Jacqueline K. Barton*

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

Received April 11, 2008



Abstract:

The DNA-binding properties of $\text{Ru}(\text{bpy})_2(\text{eilatin})^{2+}$ have been investigated to determine if the sterically expansive eilatin ligand confers specificity for destabilized single-base mismatches in DNA. Competitive DNA photocleavage experiments employing a sequence-neutral metallointercalator, $\text{Rh}(\text{bpy})_2(\text{phi})^{3+}$ (phi = 9,10-phenanthrenequinonediimine), and a mismatch-specific metalloinsertor, $\text{Rh}(\text{bpy})_2(\text{chrysi})^{3+}$ (chrysi = chrysene-5,6-quinonediimine), reveal that the eilatin complex binds to a CC mismatched site with an apparent binding constant of $2.2(2) \times 10^6 \text{ M}^{-1}$. Nonetheless, the selectivity in binding mismatched DNA is not high: competitive titrations with $\text{Rh}(\text{bpy})_2(\text{phi})^{3+}$ show that the complex binds also to well-matched B-form sites. Thus, $\text{Ru}(\text{bpy})_2(\text{eilatin})^{2+}$, despite containing the extremely expansive eilatin ligand, displays lower selectivity for the mismatch than does $\text{Rh}(\text{bpy})_2(\text{chrysi})^{3+}$, a metalloinsertor containing the smaller, though still bulky, chrysene-5,6-quinonediimine ligand. In summary, the size and shape of the eilatin ligand allow stacking with both well-matched and mismatched DNA.