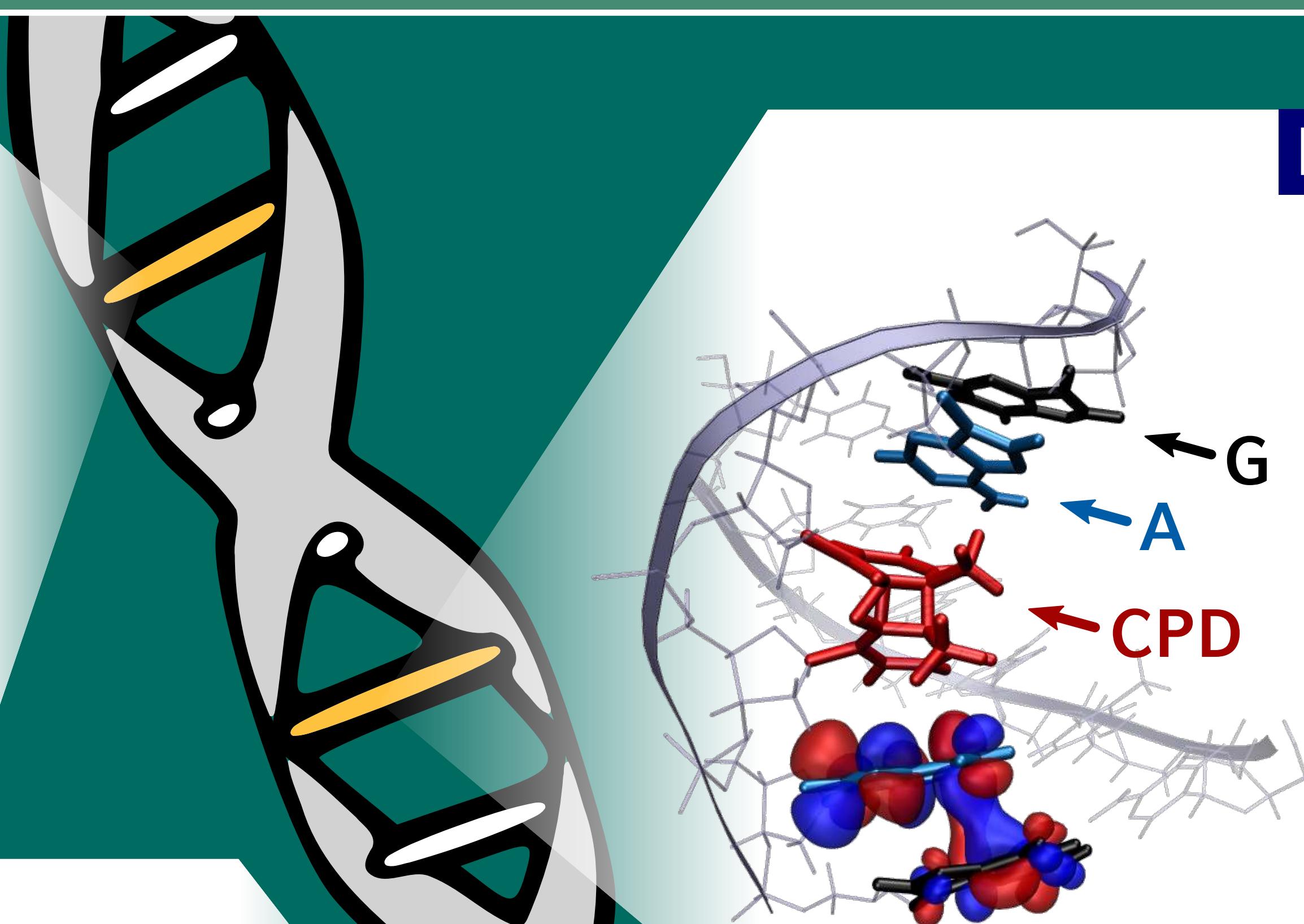


# UV-INDUCED DNA SELF-REPAIR MECHANISM TRACED WITH QUANTUM CHEMISTRY

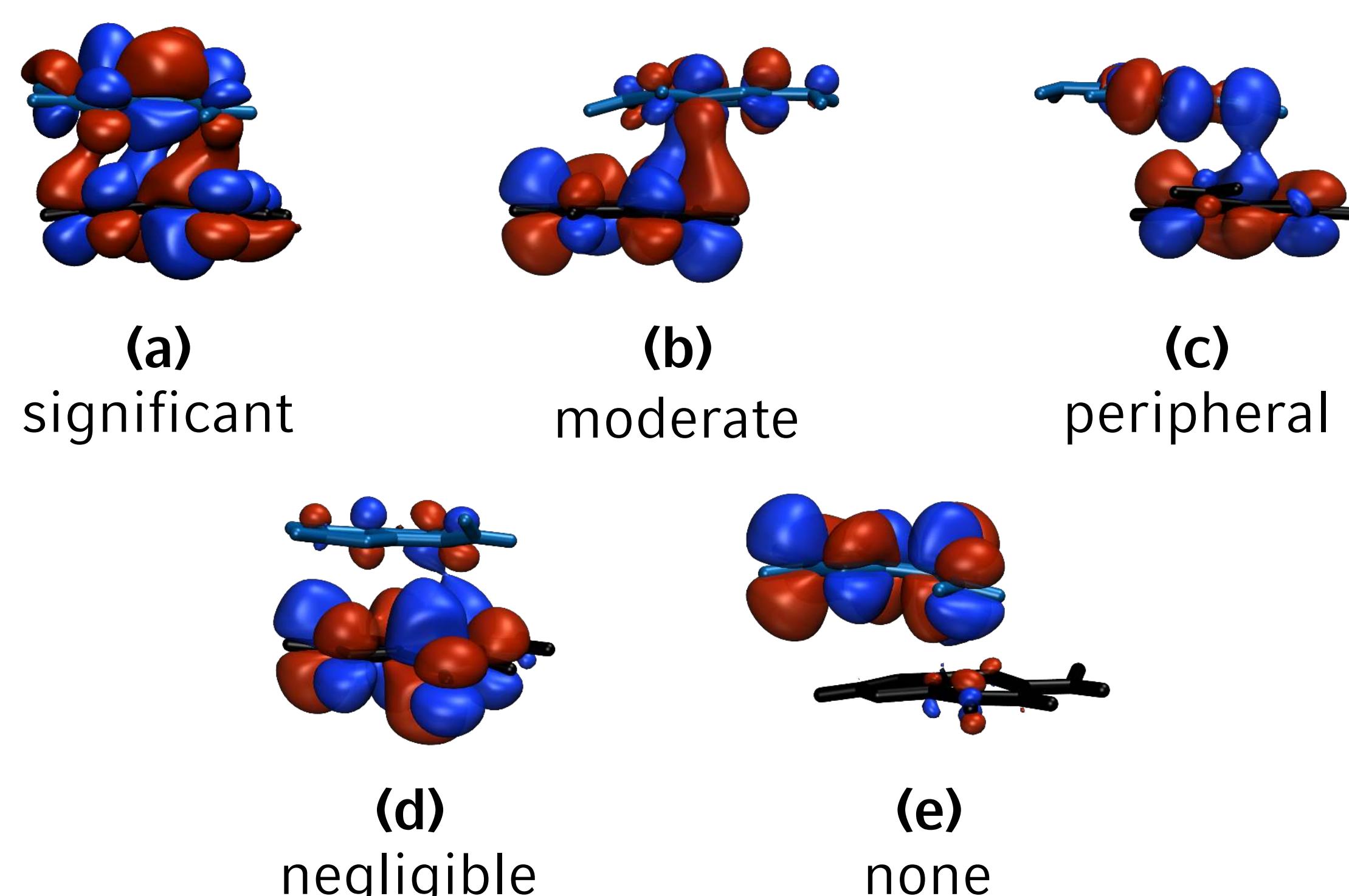
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**ABSTRACT**

- Specific DNA sequences promote self-repair of the cyclobutane pyrimidine dimer (CPD) photolesion<sup>[1]</sup> after UVB excitation (290nm)
- Theoretical investigations support photolyase-like repair mechanism

**EXPERIMENTAL BACKGROUND**

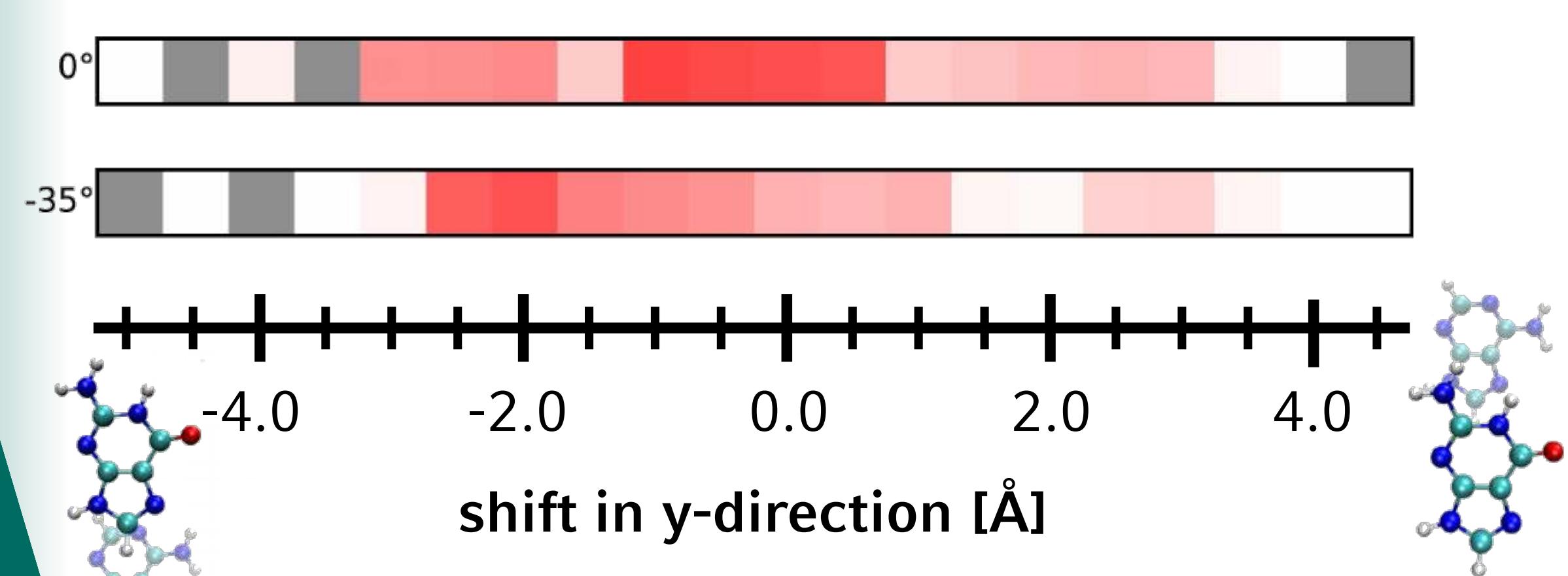
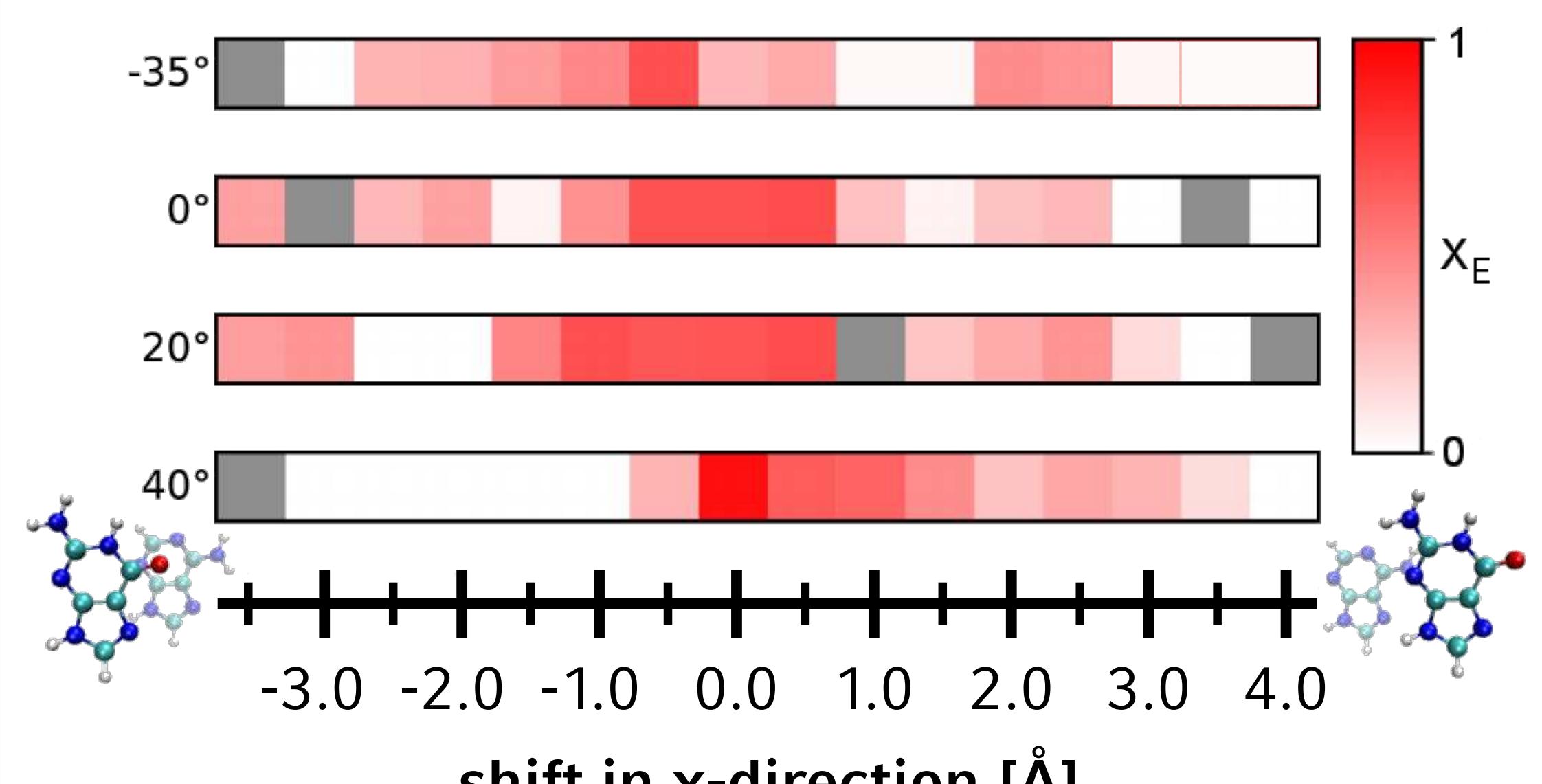
- DNA sequence: 5'-GA-T=T-AG-3'
- Selective excitation at G, not at A, induces CPD self-repair
- Proposed mechanism<sup>[1]</sup>
  1. (GA)\* exciplex formation
  2. G→A electron transfer
  3. A<sup>-</sup>→T=T electron injection
  4. CPD ring opening reaction

**CLASSIFYING ORBITAL OVERLAP****Definition of Exciplexic Strength  $X_E$** 

- Exciplex → excitation into shared orbitals
- Excitation coefficients  $c$  from TD-DFT calculation
- State-specific:  $X_{E,n} = \frac{\sum_{ab} c_{a,b}}{\sum_{abcde} c_{a,b,c,d,e}}$
- Averaged over  $N$  states:  $X_E = \frac{1}{N} \sum_{n=1}^N X_{E,n}$   
(here:  $N = 5$ )
- Takes values between 0 and 1

**STEP 1: (GA)\* EXCIPLEX FORMATION****TD-DFT/ $\omega$ B97X-D/cc-pVDZ**

- Conformational scans of isolated G-A system
- Heat map for  $X_E$  of different structures

**STEP 2: G→A CHARGE-TRANSFER****CASSCF/cc-pVDZ**

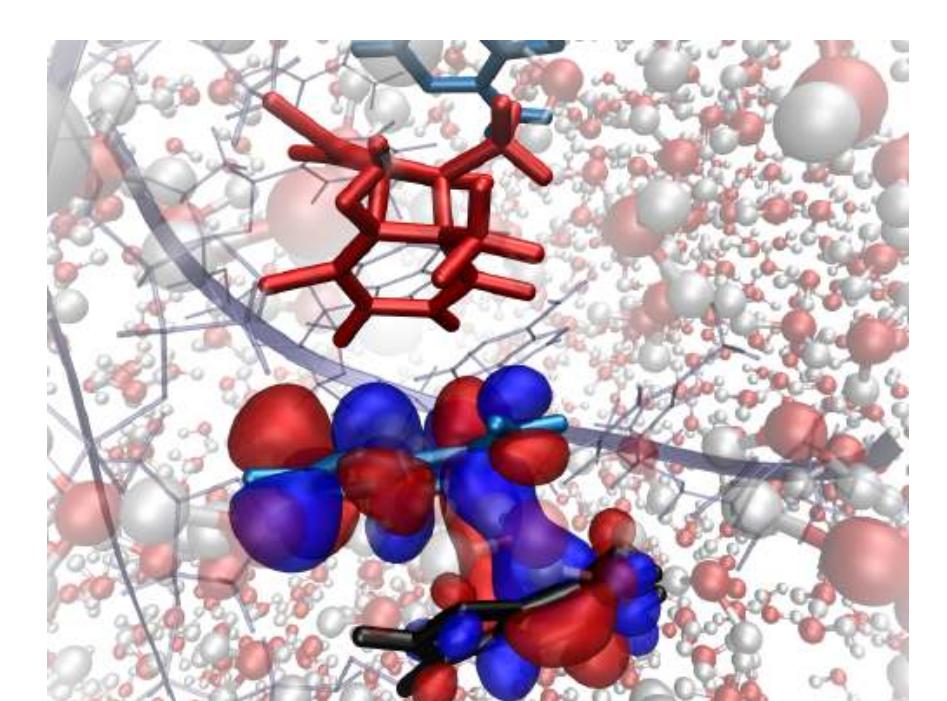
- Symmetric  $\pi\pi^*$  active spaces
- Orbital localization for easier interpretation
- Only G→A charge transfer in agreement with redox potentials<sup>[2]</sup> and experiment
- Excitation energies overestimated by 2–3eV  
(experiment: 4.28eV)

**Excitation energies in eV and characters of excitations**

CAS	4,4	6,6	8,8	10,10	12,12	14,14	16,16
$S_0 \rightarrow S_1$	6.91	6.57	5.74	6.46	6.32	5.66	5.64
	$\pi_G \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$
$S_0 \rightarrow S_2$	7.27	6.62	6.60	7.09	7.08	7.11	6.19
	$\pi_A \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_G \rightarrow \pi_G^*$
$S_0 \rightarrow S_3$	7.42	7.20	7.18	7.21	7.18	7.21	7.00
	$\pi_G \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_{G/A}^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_G^*$
$S_0 \rightarrow S_4$	7.63	7.41	7.63	7.38	7.37	7.37	7.30
	$\pi_A \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$

**INCLUDING THE ENVIRONMENT****QM(CASPT2)/MM//cc-pVDZ**

- 5'-GA-T=T-AG-3' sequence in water
- Two representative conformations
- G and A in QM region
- Excitation energies in experimental range
- Steps 1 and 2 of the proposed mechanism theoretically validated

**Excitation energies in eV and characters of excitations**

CAS	geometry 1		geometry 2	
	12,11	14,13	12,11	14,13
$S_0 \rightarrow S_1$	4.40	4.73	4.47	4.16
	$\pi_G \rightarrow \pi_G^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_G \rightarrow \pi_G^*$
$S_0 \rightarrow S_2$	4.93	5.00	5.49	5.03
	$\pi_A \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_G^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$
$S_0 \rightarrow S_3$	5.22	5.59	6.05	5.87
	$\pi_G \rightarrow Ry_A^*$	$lp_G \rightarrow \pi_G^*$	$\pi_A \rightarrow \pi_A^*$	$\pi_A \rightarrow \pi_A^*$
$S_0 \rightarrow S_4$	5.67	5.70	6.41	6.02
	$\pi_G \rightarrow Ry_A^*$	$\pi_G \rightarrow \pi_A^*$	$\pi_G \rightarrow \pi_A^*$	$lp_G \rightarrow \pi_G^*$

**REFERENCES**

- [1] D. B. Bucher, C. L. Kufner, A. Schlueter, T. Carell, W. Zinth, *J. Am. Chem. Soc.* **138**, 186 (2016).  
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[3] R. Szabla, H. Kruse, P. Stadlbauer, J. Šponer, A. L. Sobolewski, *Chem. Sci.* **9**, 3131 (2018).