



Control and design of photochemistry

photoisomerization and excitation energy transfer

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FOKKE & SUKKE

WETEN WAAR HET IN DE WETENSCHAP OM DRAAIT

know what science is all about

... ZEER INDRUKWEKKEND, COLLEGA ...

... very impressive, colleague ...

MAAR WERKT
HET OOK IN
THEORIE?

but does it also work in theory?

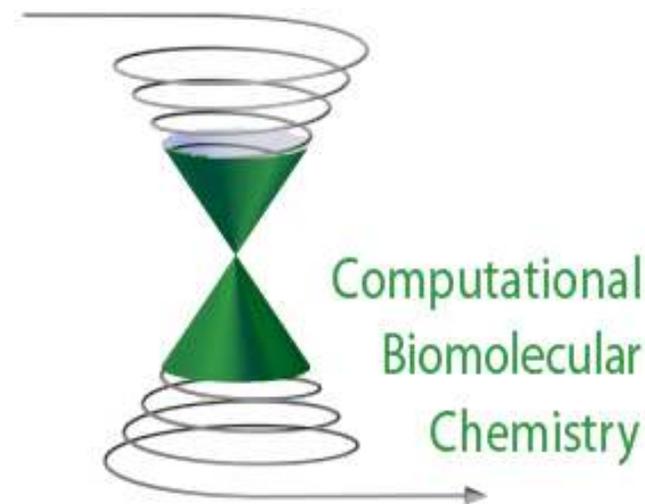


Photo-isomerization

rotation of double bond after photon absorption

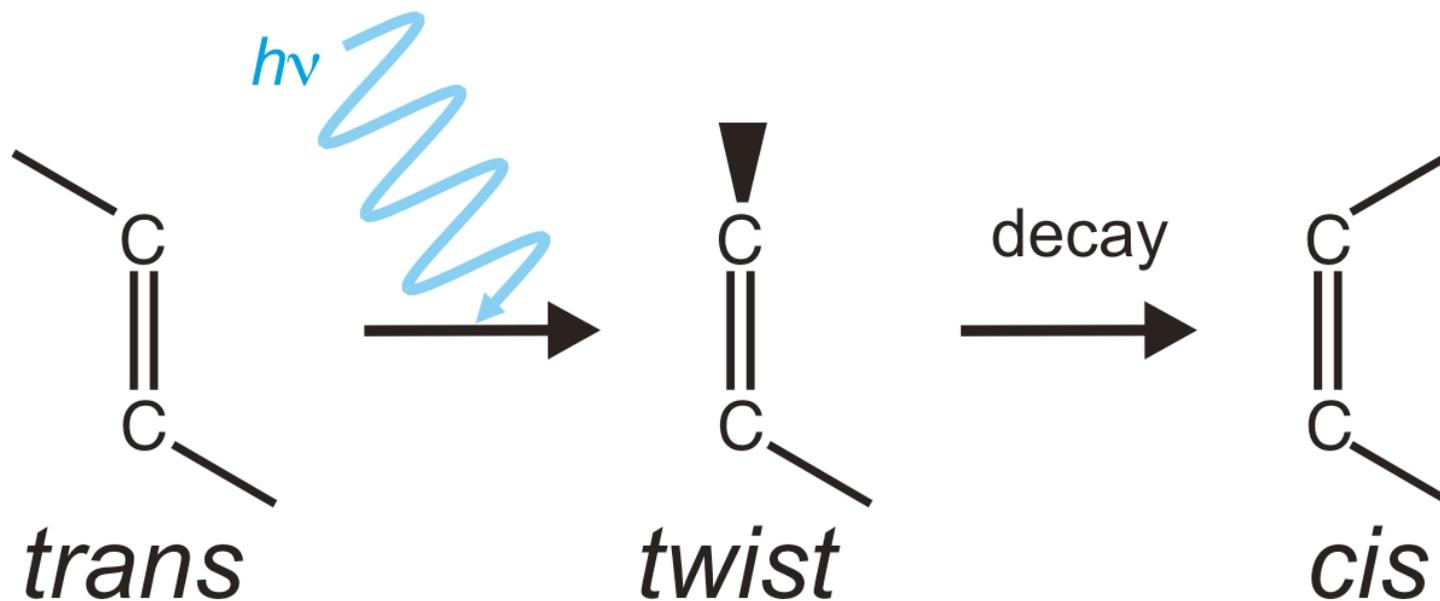
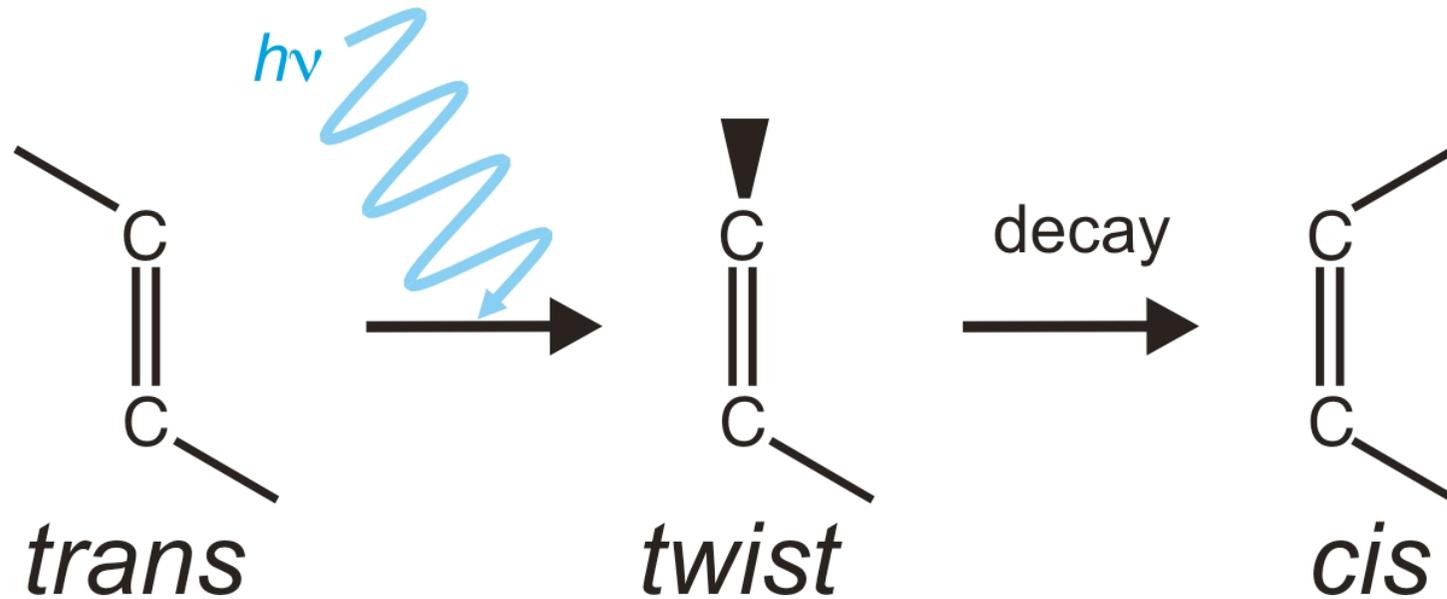


Photo-isomerization

rotation of double bond after photon absorption



ubiquitous in photo-biology, e.g. :

photosynthesis

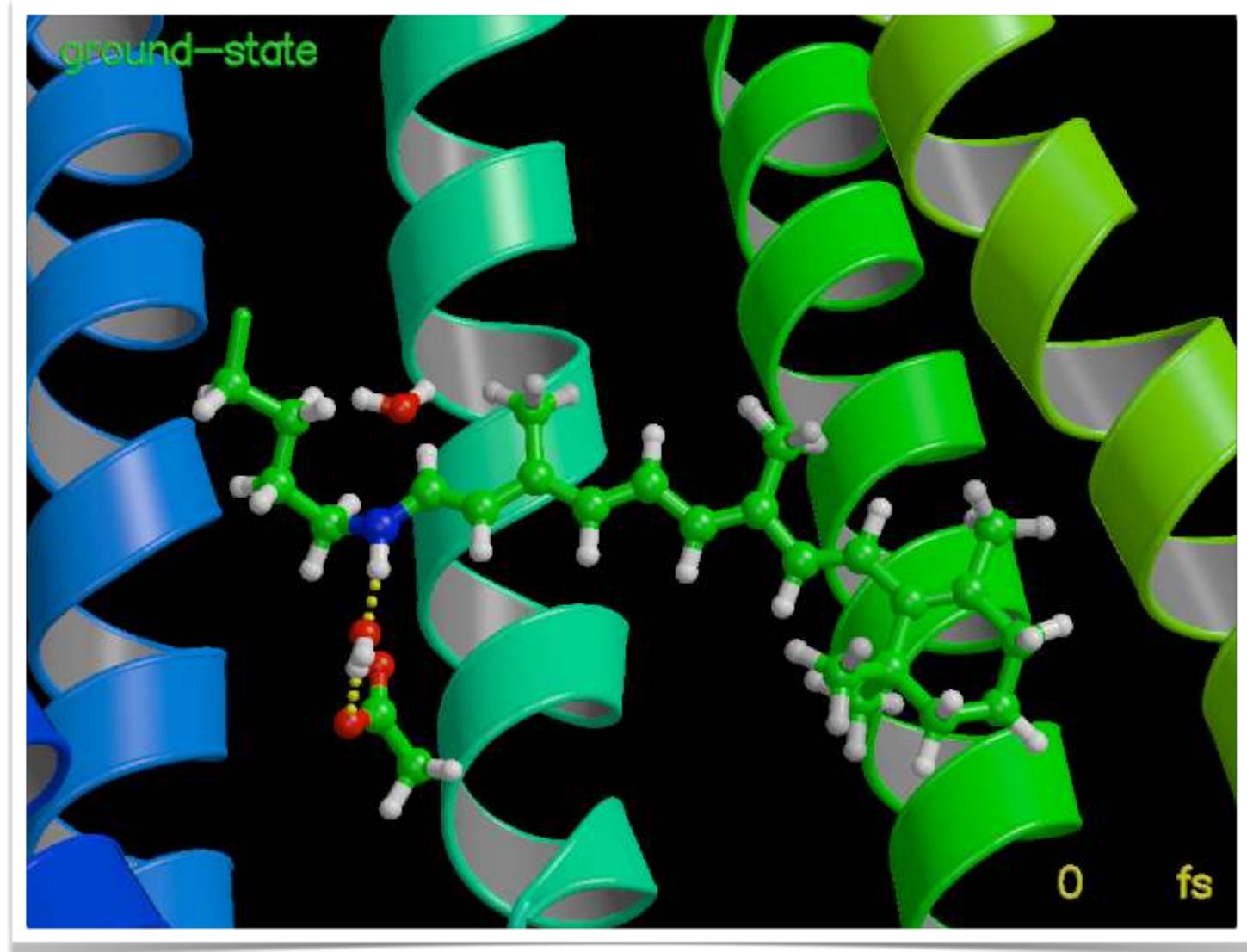
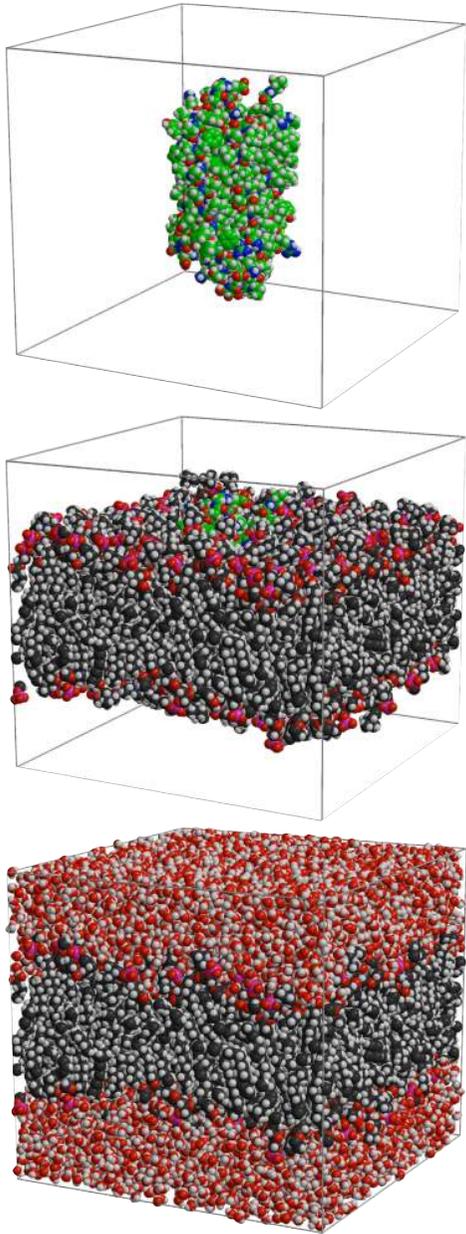
bacteriorhodopsin

light sensing

rhodopsin, phytochromo, photoactive yellow protein, ...

photoisomerization in bacteriorhodopsin

observe while it happens in MD simulations

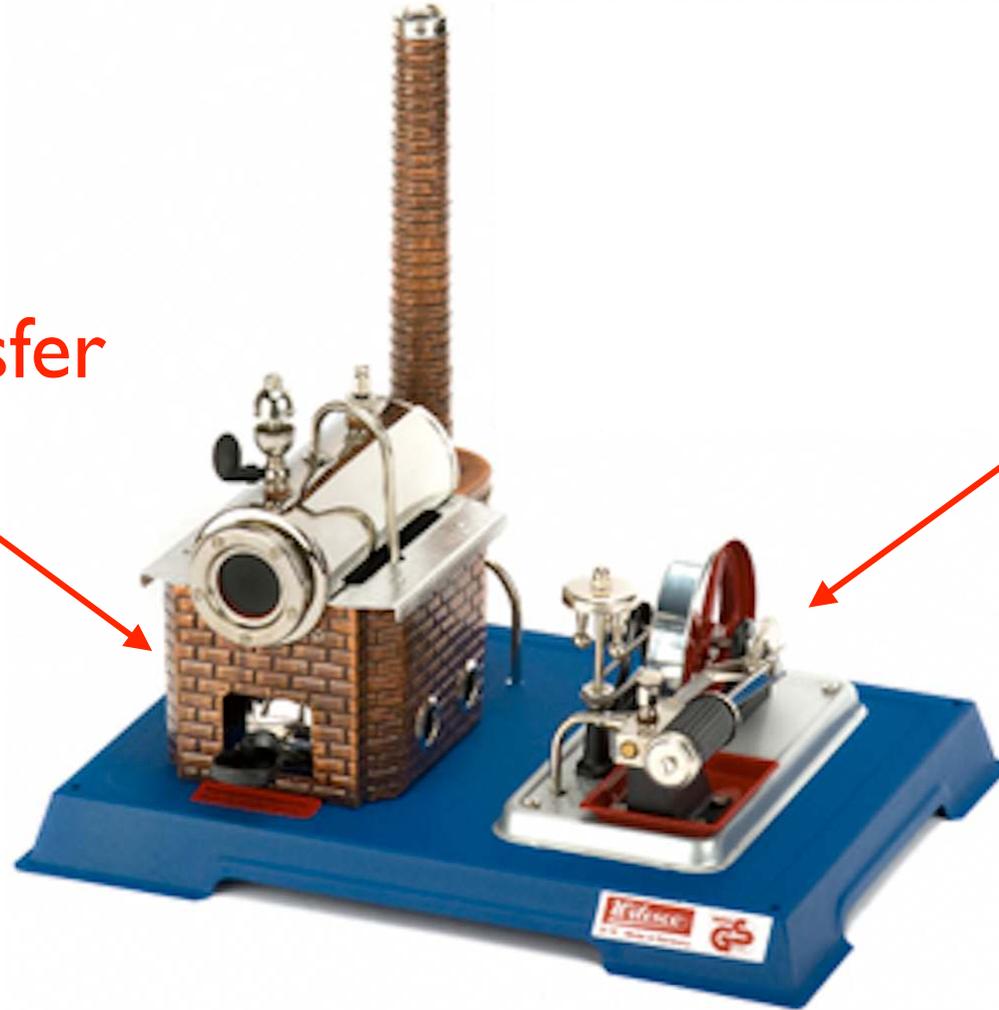


Our ultimate goal

arteficial molecular machines

energy transfer

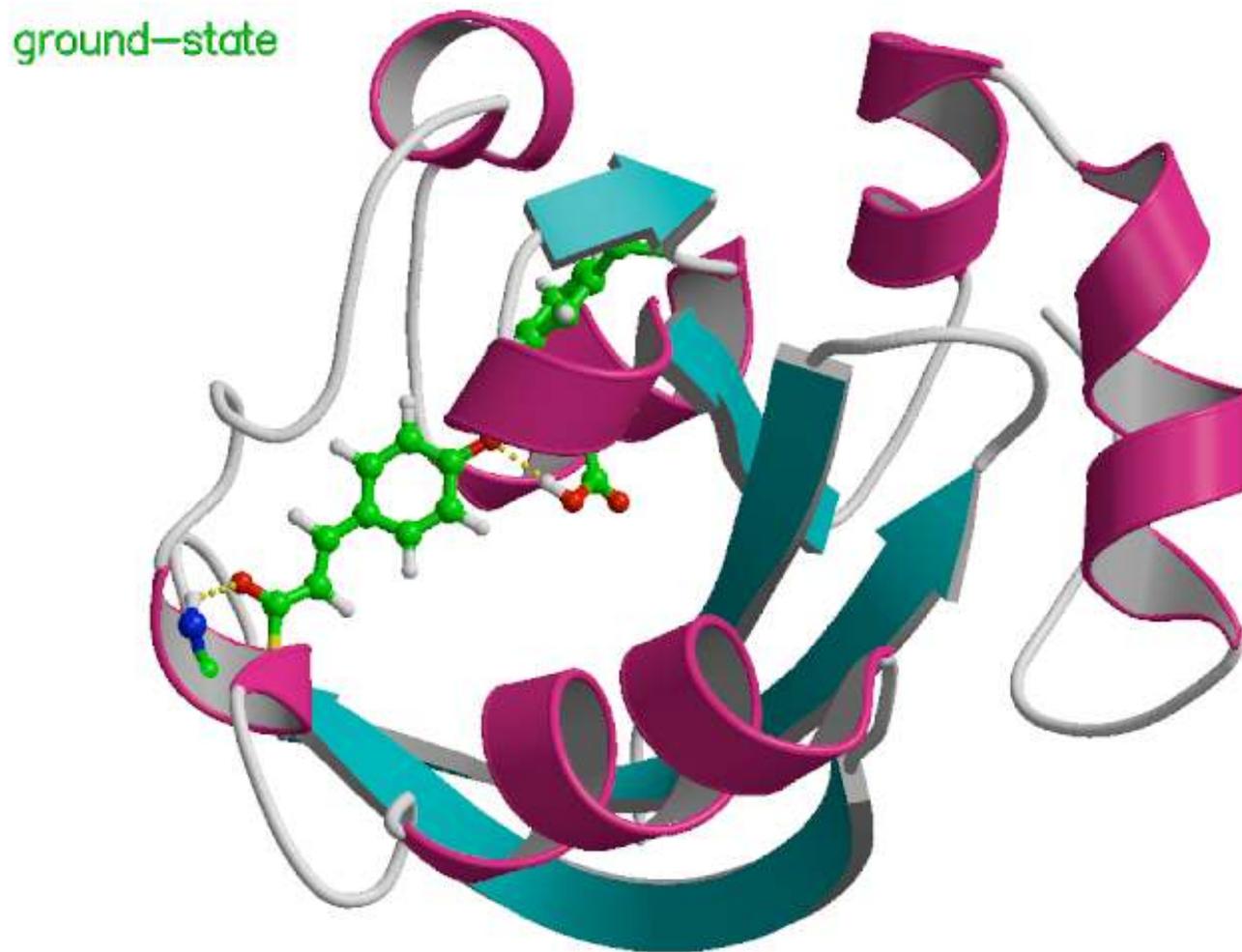
isomerization



Get inspired by nature

e.g. photo-isomerization in photoactive yellow protein

learn & mimic the effect of the protein environment

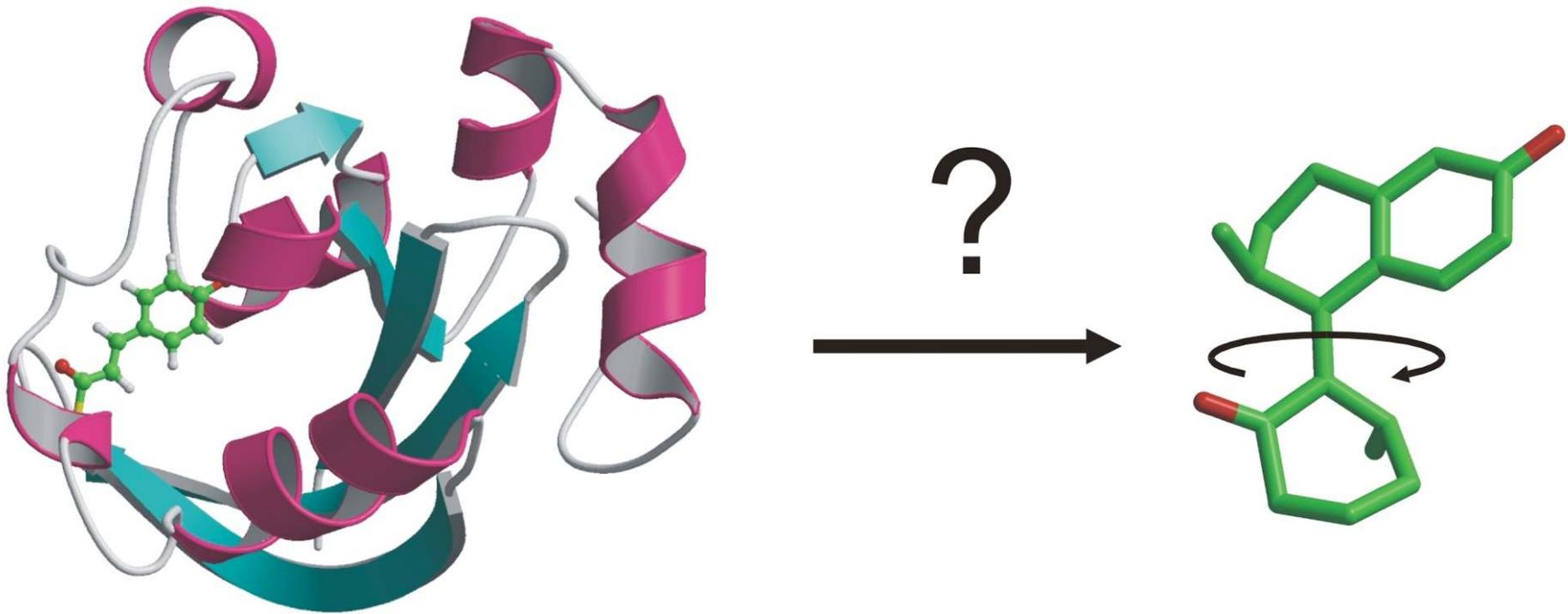


0 fs

Get inspired by nature

photo-isomerization in photoactive yellow protein

learn & mimic the effect of the protein environment



however....

still too complex, even in our simulations

Reducing complexity in MD simulations

maximally correlated motion in trajectory ($\mathbf{x}(t)$)

find vector $\mathbf{a} \in R^{3N}$ that correlates with observable $f(t)$

$$p_a(t) = [\mathbf{x}(t) - \langle \mathbf{x} \rangle] \cdot \mathbf{a}$$

observable

quantum yield, energy gap, lifetime, ...

maximize Pearson coefficient

$$R = \frac{\text{cov}(f, p_a)}{\sigma_f \sigma_a}$$

reducing dimensionality: basis

normal modes: eigenvectors of Hess matrix

principal components: eigenvectors of covariance matrix

$$C_{ij} = \langle (x_i - \langle x_i \rangle)(x_j - \langle x_j \rangle) \rangle$$

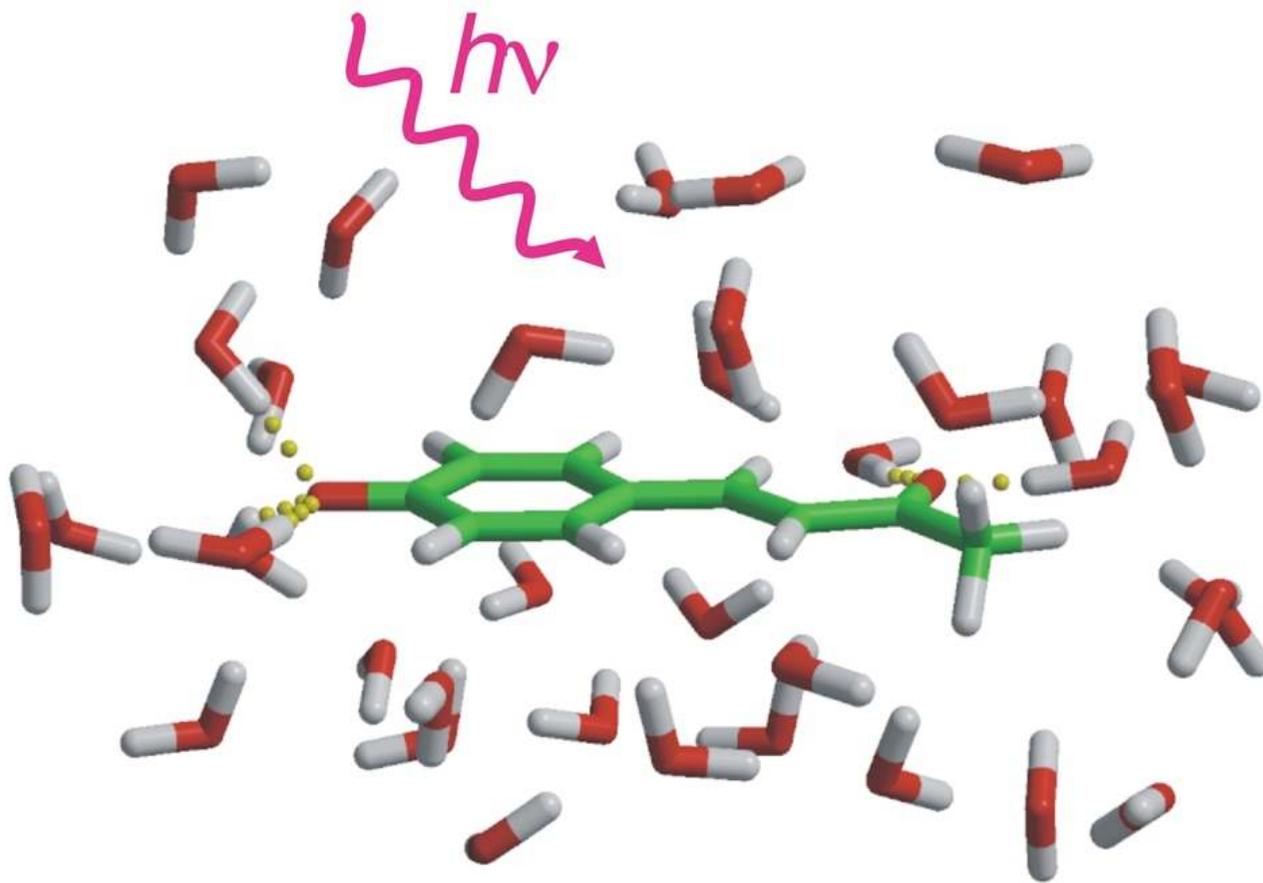
Simpler model systems

photo-isomerization in isolation and solution

lower complexity

systematic improvement of theory

high quality experimental data



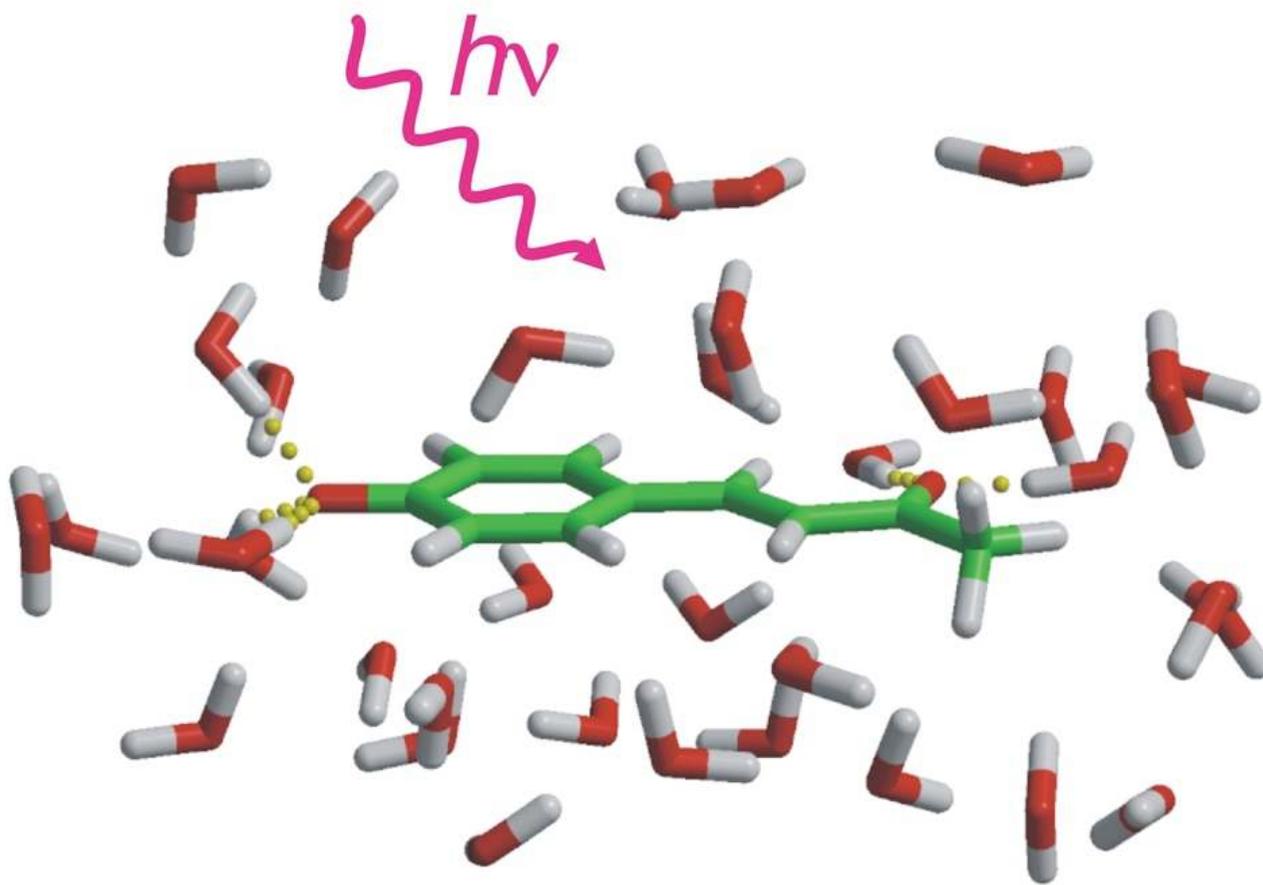
Simpler model systems

photo-isomerization in isolation and solution

lower complexity

find correlation between conformation & quantum yield

control quantum yield

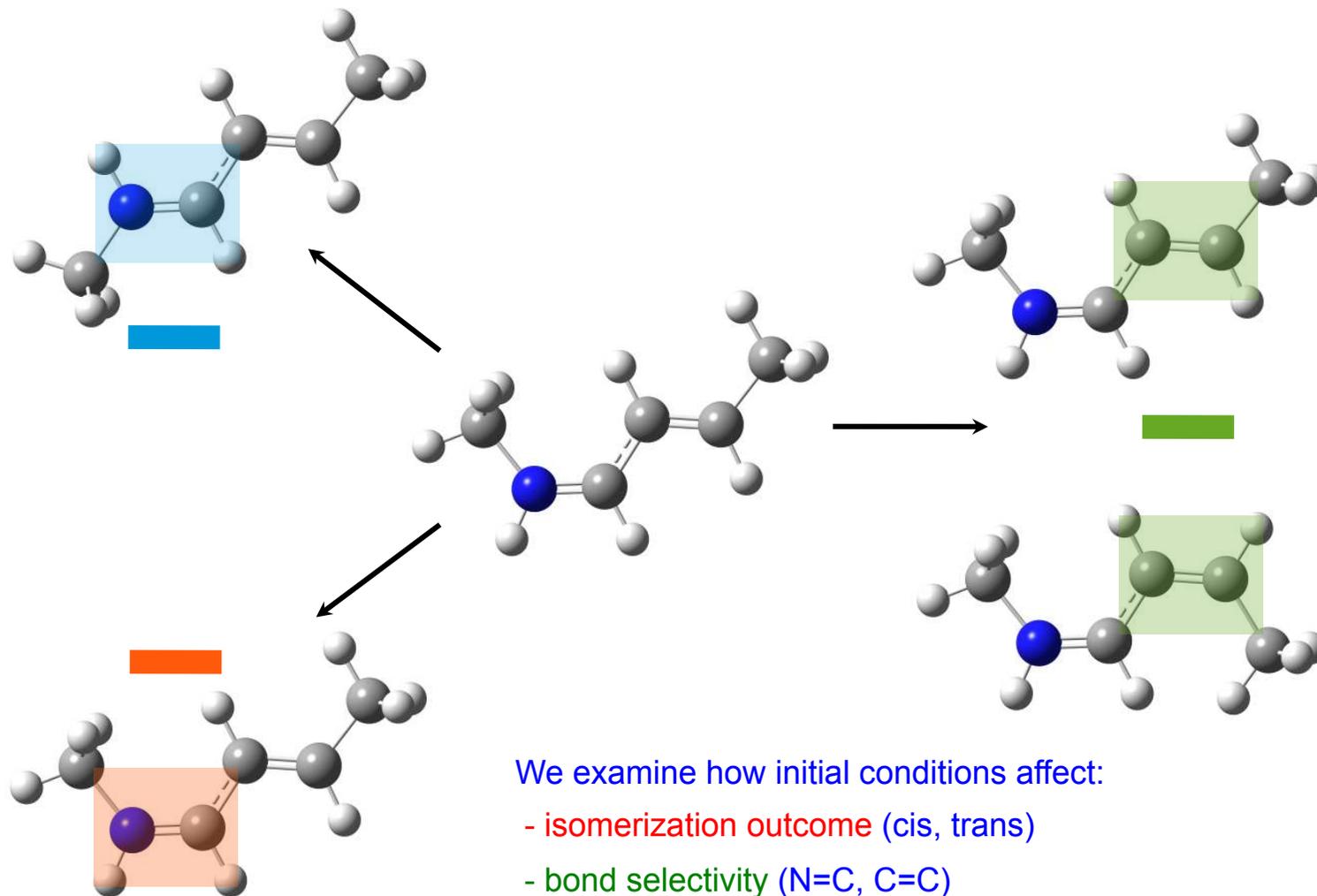


Simpler model systems

safety in numbers: many simulations

statistical analysis conformation-outcome

protonated schiff base (retinal model)



Non-adiabatic molecular dynamics

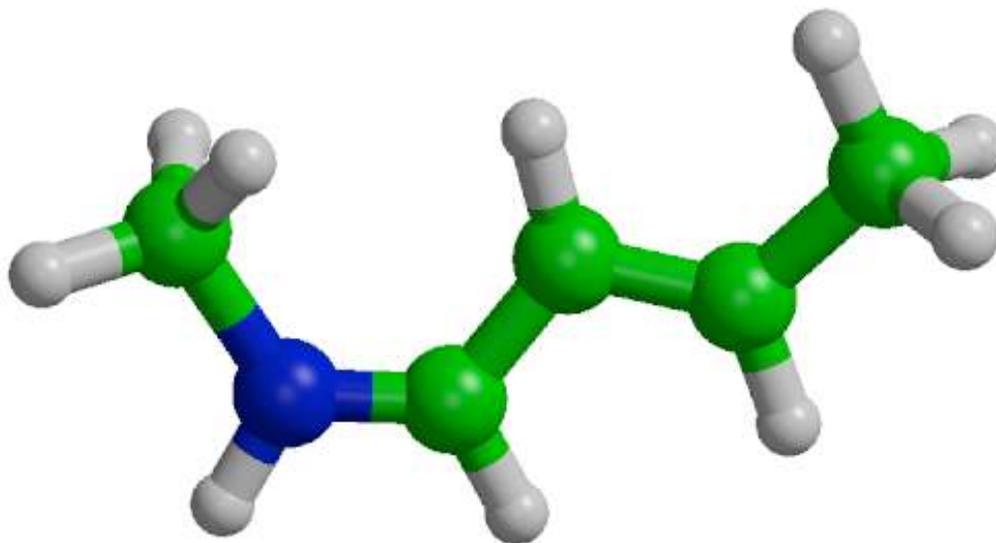
comparing diabatic hopping with fewest switches

photoisomerization of protonated Schiff base

aim a: find out if initial conditions determine outcome

aim b: control outcome

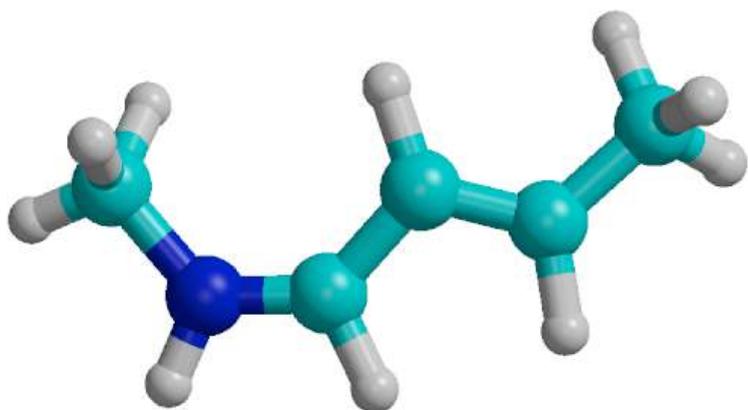
aim c: compare hopping algorithms



simulations

CASSCF(4,4)/6-31G*, diabatic & fewest switches surface hopping

excited-state

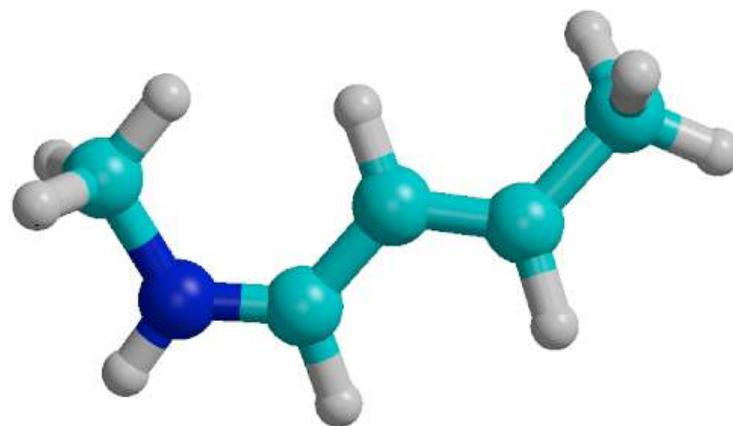


QY: 44.6%/42.4%

average lifetime: 115.8 fs/75.2 fs

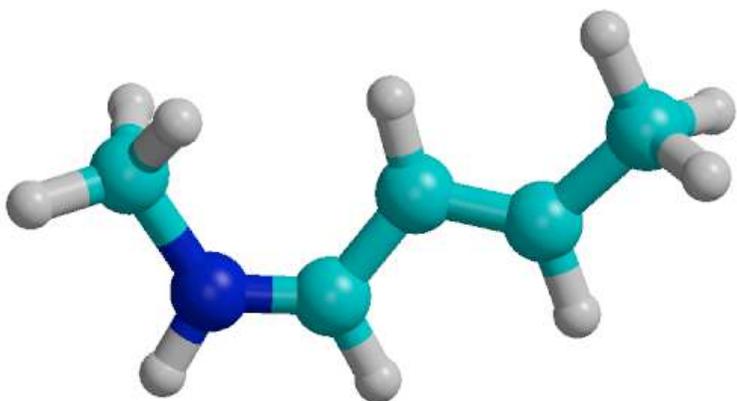
0 fs

excited-state



0 fs

excited-state

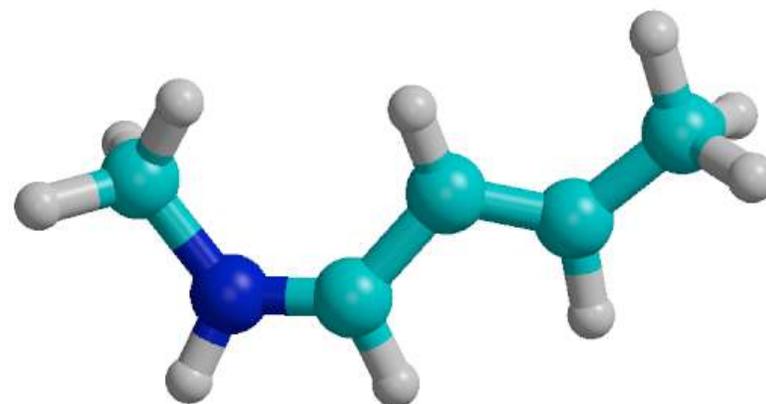


QY: 35.5%/34.8%

average lifetime: 139.5 fs/83.7 fs

0 fs

excited-state



QY (both): 19.9 %/22.8%

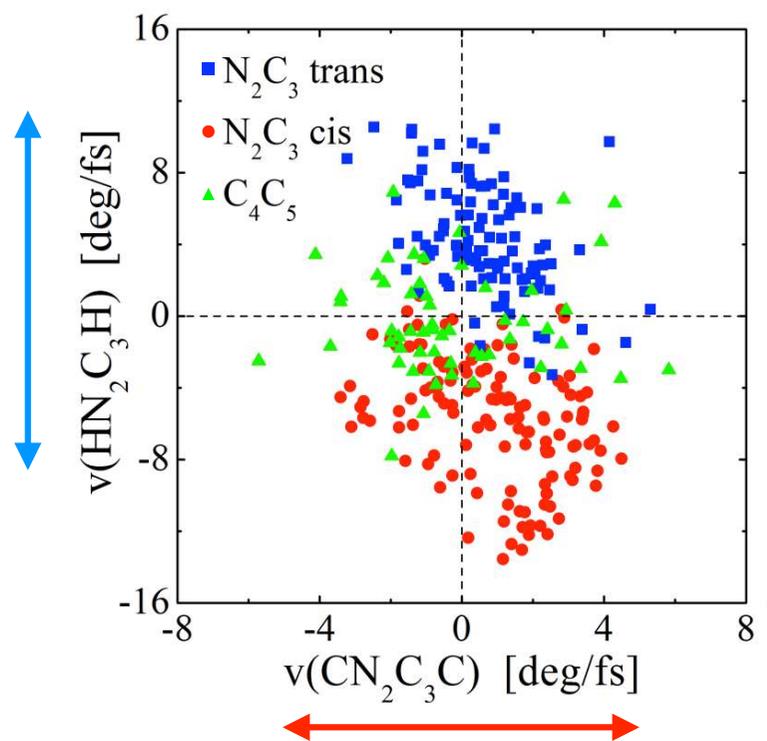
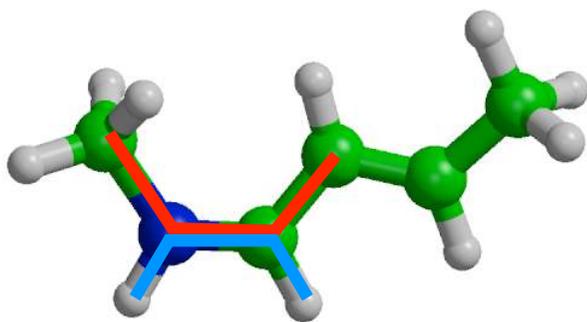
average lifetime: 60.2 fs/54.6 fs

0 fs

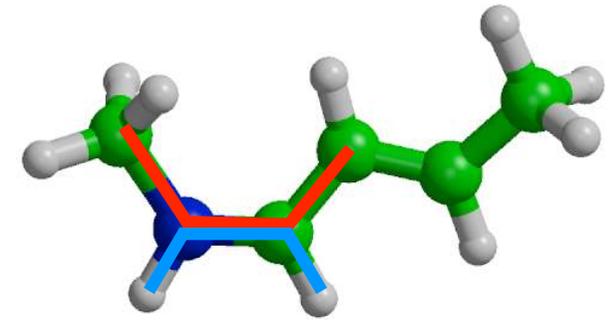
Simpler model systems

free unbiased simulations

what determines outcome: hydrogen-out-of-plane motion

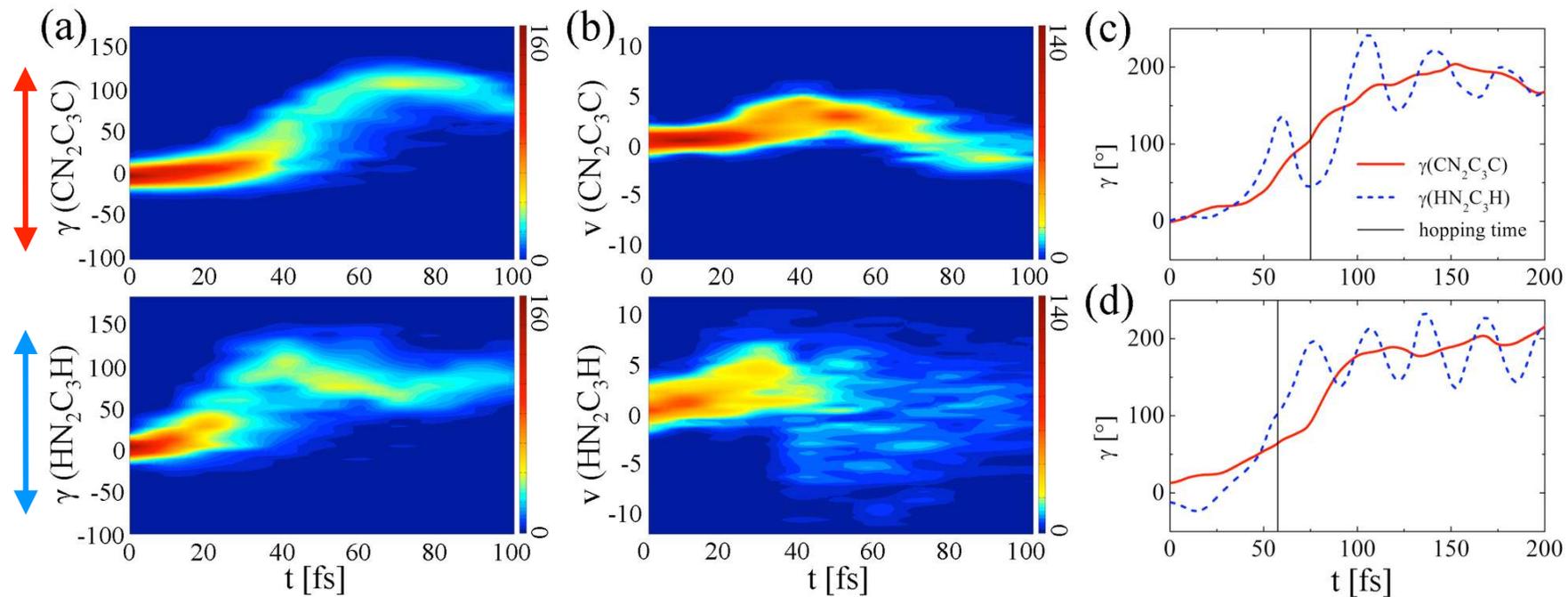


Simpler model systems



free unbiased simulations

phase between $\text{HN}=\text{CH}$ and $\text{CN}=\text{CC}$



evolutionary approach: optimize for synchronicity

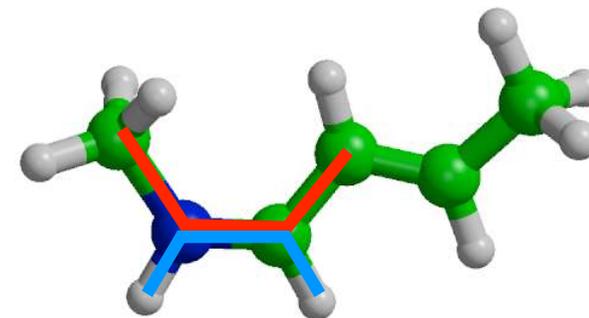
constrain dihedral angles from synchronous simulations

generate new ensemble

Simpler model systems

free unbiased simulations

thermal ensemble



Outcome	N ₂ C ₃ cis	N ₂ C ₃ trans	C ₄ C ₅
τ_{DSH} [fs]	96 ± 1	132 ± 2	51 ± 1
N _{i,DSH}	132	105	59
P _{i,DSH} [%]	44.6	35.5	19.9
τ_{FSH} [fs]	65 ± 1	74 ± 1	46 ± 1
N _{i,FSH}	208	171	112
P _{i,FSH} [%]	42.4	34.8	22.8

optimizing synchronicity

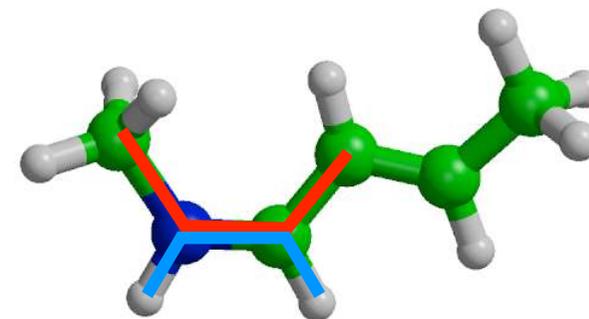
thermal ensemble

Therm. Ens.	γ_1 [°]	γ_2	γ_3	γ_4	P_{sync}	N_{traj}
unconstr. FSH	-	-	-	-	7.63 %	491
unconstr. DSH	-	-	-	-	1.0%	296

Simpler model systems

free unbiased simulations

thermal ensemble



Outcome	N ₂ C ₃ cis	N ₂ C ₃ trans	C ₄ C ₅
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optimizing synchronicity

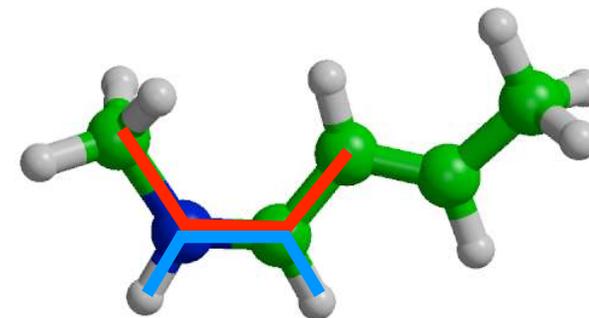
new ensemble with fixed dihedrals

Therm. Ens.	γ_1 [°]	γ_2	γ_3	γ_4	P_{sync}	N_{traj}
unconstr. FSH	-	-	-	-	7.63 %	491
unconstr. DSH	-	-	-	-	1.0%	296
1 FSH, γ_i - [14]	-17.5	168.6	174.1	-	6.06 %	99
2 FSH	13.0	150.0	-172.4	-	18.09 %	187
2 DSH	13.0	150.0	-172.4	-	16.24 %	193
3 FSH	-20.63	-154.4	172.9	-	14.57 %	199
3b FSH	-20.63	-154.4	172.9	11.1	14.21 %	197

Simpler model systems

free unbiased simulations

thermal ensemble



Outcome	N ₂ C ₃ cis	N ₂ C ₃ trans	C ₄ C ₅
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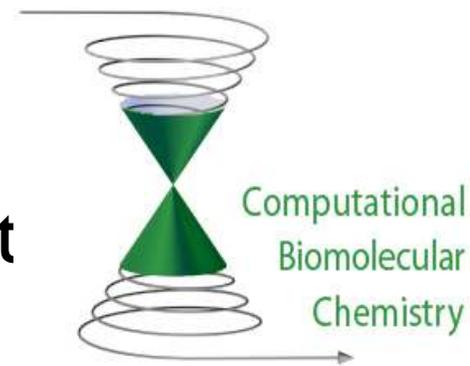
optimizing synchronicity

second generation

Outcome	N ₂ C ₃ cis	N ₂ C ₃ trans	C ₄ C ₅
P _{i,DSH} [%]	39.9	46.1	14.0
P _{i,FSH} [%]	47.8	39.9	12.4

challenge: fixing dihedrals by chemical modification?

Acknowledgements



members from the Grubmüller department



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IC London

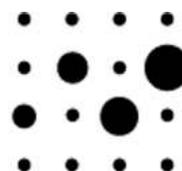


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(Toulouse, Fr.)

funding



DAAD



VolkswagenStiftung