

Développement d'une Méthode de Chimie Quantique Mêlant Plusieurs Niveaux de Théorie : Applications à l'Etude des Etats Electroniques de Macromolécules

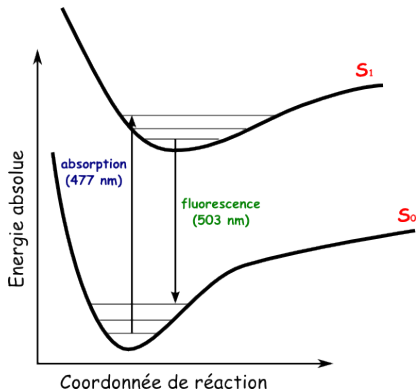
Pierre-François Loos

Equipe de Chimie et Biochimie Théoriques,
UMR 7565 CNRS-UHP, Institut Jean Barriol (FR CNRS 2843),
Faculté des Sciences et Techniques, Nancy-Université, B.P. 239,
France

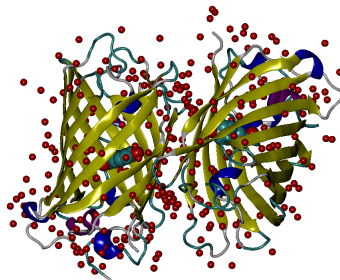
13 Juin 2008

Green Fluorescent Protein (GFP)¹

Excitation et émission



PDB ID : 1GFL² — *Aequorea victoria*



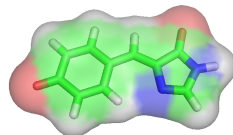
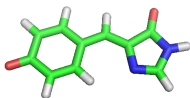
¹Zimmer, *Chem. Rev.* **2002**, *102*, 759.

²Yang et al. *Nat. Biotechnol.* **1996**, *14*, 1246.

Système modèle & effets de solvant

Chromophore de la GFP¹ : état **B**

Modèle SCRF : solvation implicite



Méthodes	Etat B
SAC-CI ³	558
TD-B3LYP ⁴	403
SAC-CI(Onsager) ³	549
TD-B3LYP(PCM) ⁴	417
Exp	477

³Das et al., *J. Comput. Chem.* **2003**, *24*, 1421

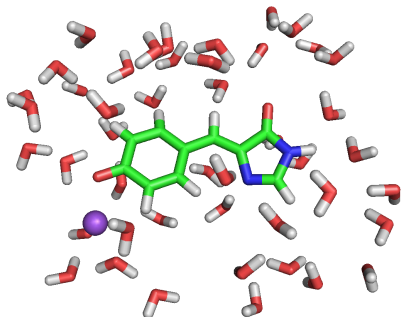
Méthodes hybrides

QM:MM : solvation explicite

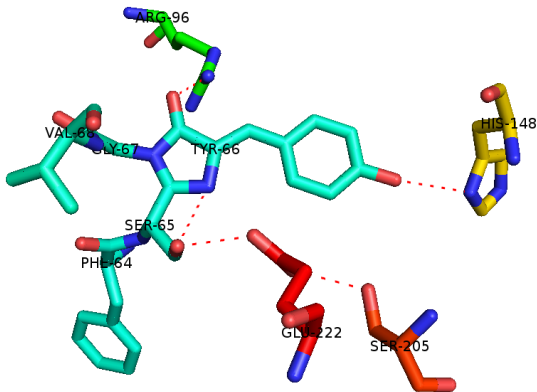
CASPT2/TIP3P⁵ : **B** (434 nm)

QM/MM : environnement biologique

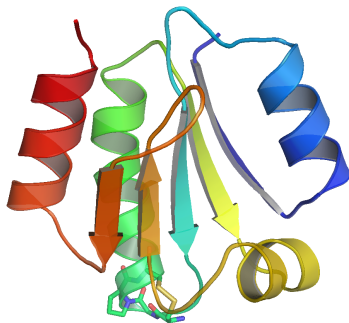
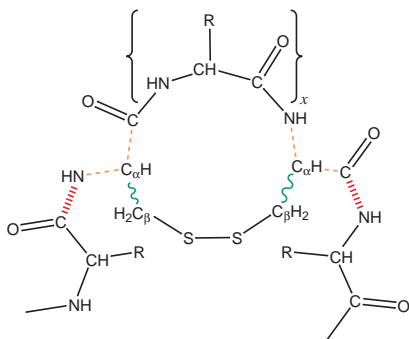
CASPT2/CHARMM⁵ : **B** (442 nm)



Site actif de la GFP : liaisons covalentes protéine-chromophore



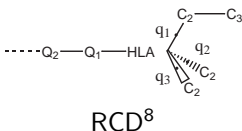
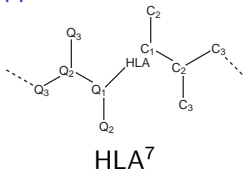
QM/MM, ou l'art de la partition



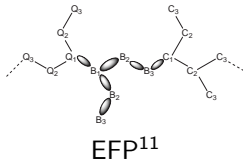
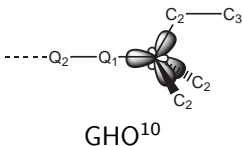
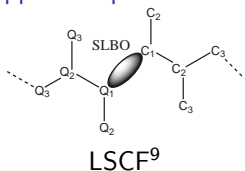
Thioredoxin⁶ (1TOF)

⁶Dumont et al. *Chem. Phys. Lett.*, in press; Dumont et al. *J. Phys. Chem. B*, en révision; Dumont et al. *J. Chem. Theor. Comput.*, soumis.

Approches Link-Atom



Approches par orbitales localisées



⁷Maseras et al. *J. Comput. Chem.* **1995**, *16*, 1170.

⁸Lin et al. *J. Phys. Chem. A* **2005**, *109*, 3991.

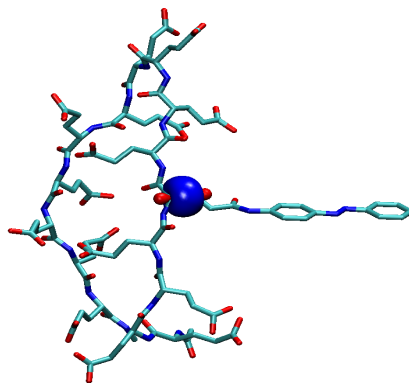
⁹Assfeld et al. *Chem. Phys. Lett.* **1996**, *263*, 100.

¹⁰Pu et al. *J. Phys. Chem. A* **2004**, *108*, 632.

¹¹Kairys et al. *J. Phys. Chem. A* **2001**, *104*, 6656.

Calcul LSCF/MM

- Local Self-Consistent Field :
partie quantique



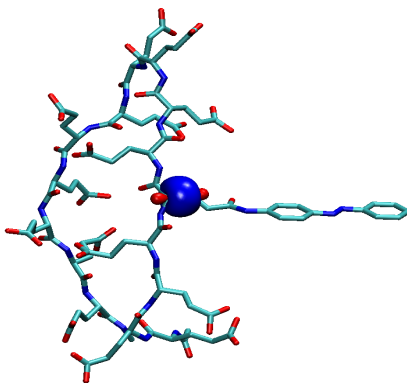
Calcul LSCF/MM

- Local Self-Consistent Field :
partie quantique
 - Optimisation sous contrainte
de la fonction d'onde :

$$F \cdot C = \underbrace{S \cdot C \cdot E}_{\text{variationnelle}} + \underbrace{S \cdot L \cdot A}_{\text{gelée}}$$

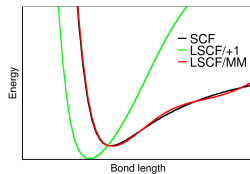
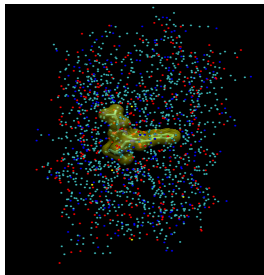
- Frontière QM/MM
*Strictly Localized Bond
Orbital (SLBO)* \iff

$$|I\rangle = \sum_{\mu \in \{X, Y\}} I_{\mu I} |\mu\rangle$$



Calcul LSCF/MM

- Local Self-Consistent Field : interface QM \rightleftharpoons MM



¹²Ferré et al. *J. Comput. Chem.*, **2002**, *23*, 610.

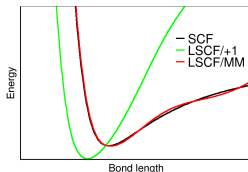
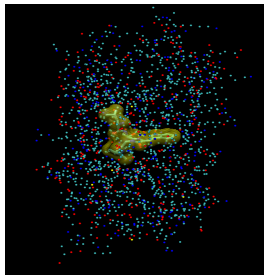
Calcul LSCF/MM

- Local Self-Consistent Field : interface QM \rightleftharpoons MM
 - Electronic Embedding : polarisation explicite de la fonction d'onde

$$\sum_{A \in \text{MM}} \sum_{\mu\nu} P_{\mu\nu}^T \left\langle \mu \left| \frac{q_A}{|\mathbf{r} - \mathbf{R}_A|} \right| \nu \right\rangle$$

- Atome Y quanto-classique :
 - quanto : base gaussienne QM + charge nucléaire (+1)
 - classique : paramètres non-liés (charge, vdw) + certains liés
- Potentiel de liaison frontière¹² :

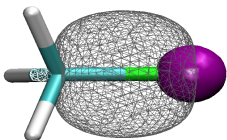
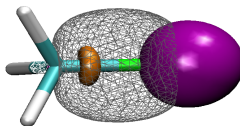
$$E_{X,Y,I} = (A + Br + Cr^2) e^{Dr} + \frac{E}{r}$$



¹²Ferré et al. *J. Comput. Chem.*, **2002**, *23*, 610.

Comment s'affranchir du potentiel frontière ?

- Prise en compte des e- de cœur de l'atome frontière (1 e- SLBO + 2 e- de cœur)
- Augmentation de la charge nucléaire (+3)

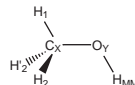
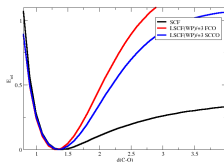
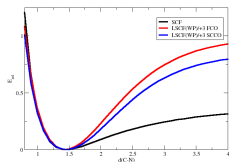
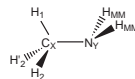
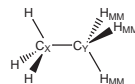
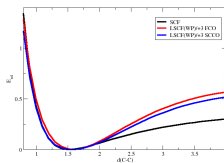
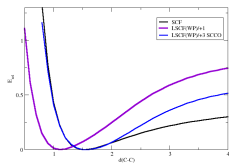
FCO¹³SCCO¹⁴

¹³Fornili et al. *Chem. Phys. Lett.* **2006**, 427, 236.

¹⁴Loos et al. *Comput. Lett.* **2007**, 4, 473.

Profils de dissociation des différents types de liaisons

Ethane (C-C)



Méthylamine (C-N)

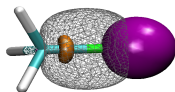
Méthanol (C-O)

Orbitale de cœur gelée (FCO) ou auto-cohérente (SCCO) ?

- SCCO plus diffuse que FCO



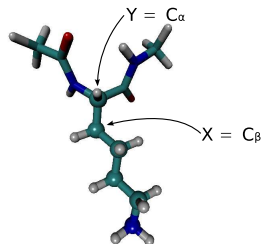
- Délocalisation partielle de la SCCO vers l'atome X



- SCCO évite la détermination de la FCO !
- Performances géométriques et énergétique quasi-identique

Tripeptide Ace-Lys-NMe

- QM : HF/6-311G**
- MM : Amber *ff99* + RESP
- SLBO et FCO : ELMO¹⁵



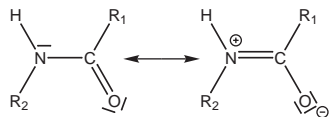
	$d(C_\alpha-C_\beta)$	ΔE
FBP	1,519(-0,017)	239,0(3,6)
FCO	1,573(0,037)	237,6(2,2)
SCCO	1,589(0,053)	238,0(2,6)



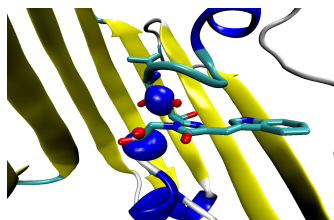
¹⁵Stoll et al. *Theor. Chem. Acc.* **1980**, 57, 169; Fornili et al. *J. Mol. Struct. (THEOCHEM)* **2003**, 632, 157.

Traitement de la liaison peptidique¹⁶

- Prise en compte de 2 e⁻ de valence pour l'atome d'azote
- Permet de représenter le doublet de l'azote
 ⇒ augmentation de la charge nucléaire (+5) de l'atome d'azote frontière



Formes mésomères

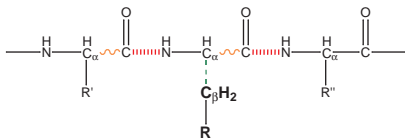


Enhanced Cyan Fluorescent Protein (10XD)

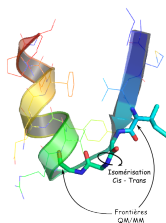
¹⁶Loos et al. *AIP Conf. Proc.* **2007**, 963, 308.

Une alternative intéressante aux partitions communes

- Partition naturelle pour les polypeptides et protéines
- Utilisable quelque soit le champ de force (traitement des charges classiques)
- Description symétrique de l'acide aminé
- Densité convenable à la jonction



Différents localisations

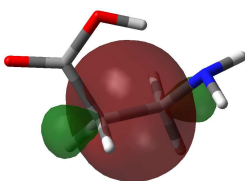
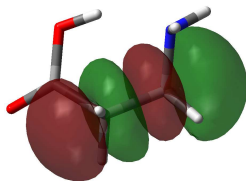


DHFR (1DRE) : C_α-C(=O)¹⁷

¹⁷Loos et al. *Theor. Chem. Acc.* **2007**, *118*, 165.

Vers une SLBO aut-cohérente : SCSLBO¹⁸

$$F' = \begin{pmatrix} F'_{11} & F'_{12} \\ F'_{12} & F'_{22} \end{pmatrix} = \begin{pmatrix} \langle l_l | \hat{F} | l_l \rangle & \langle l_l^* | \hat{F} | l_l \rangle \\ \langle l_l | \hat{F} | l_l^* \rangle & \langle l_l^* | \hat{F} | l_l^* \rangle \end{pmatrix}$$

SLBO $|l_l\rangle$ SLABO $|l_l^*\rangle$

$$|l_l\rangle_{\text{new}} = C_1^l \cdot |l_l\rangle_{\text{old}} + C_2^l \cdot |l_l^*\rangle_{\text{old}}$$

$$|l_l^*\rangle_{\text{new}} = -C_2^l \cdot |l_l\rangle_{\text{old}} + C_1^l \cdot |l_l^*\rangle_{\text{old}}$$

¹⁸Loos et al. *J. Chem. Theor. Comput.* **2007**, 3, 1047.

Comment obtenir une SLABO ?

- idem que SLBO : rotation dans l'espace des MOs virtuelles
- Projection PB → GB :


$$|I_I^*\rangle = \sum_{\mu\nu}^{\text{GB}} |\mu\rangle (\mathbf{S}^{-1})_{\mu\nu} \langle\nu|I_I^*\rangle_{\text{PB}}$$

- "Sens chimique" :

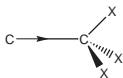
$$\text{SLBO} : |I_I\rangle = I_{X,I}|h_I^{X_I}\rangle + I_{Y,I}|h_I^{Y_I}\rangle$$

$$\text{SLABO} : |I_I^*\rangle = N_I^* \left(I_{X,I}|h_I^{X_I}\rangle - I_{Y,I}|h_I^{Y_I}\rangle \right) = I_{X,I}^*|h_I^{X_I}\rangle + I_{Y,I}^*|h_I^{Y_I}\rangle$$

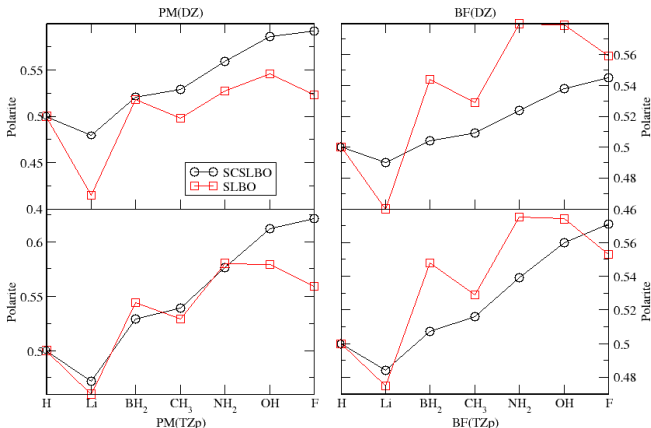
- Méthode de Head-Gordon et collaborateurs¹⁹

¹⁹Subotnik et al. *J. Chem. Phys.* **2005**, *123*, 114108. 

Effets de substituants : HF/6-31G* et HF/6-311G**



$$P_I = \frac{q_{C_X}}{2} = \sum_{\mu \in C_X} \sum_{\nu} |l_{\mu I}| |l_{\nu I}| S_{\mu\nu}$$

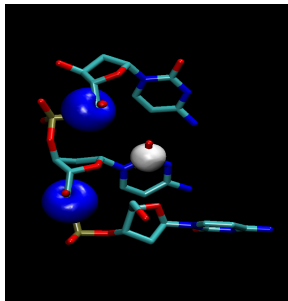


Energie d'ionisation de cœur

- Permet d'obtenir des informations sur la structure et la nature des molécules¹
- Etudes expérimentales sur des macromolécules²

Etudes théoriques

- Etats excités \implies effondrement variationnel et contrainte d'orthogonalité³
- Systèmes macromoléculaires \implies calculs LSCF/MM

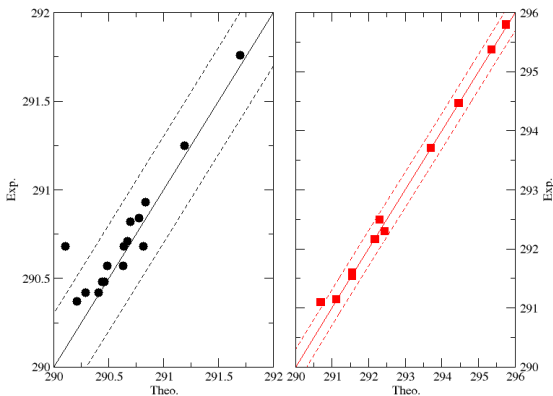


¹Schwarz et al. *Angew. Chem. Int. Ed.* **1974**, *13*, 454.

²voir par exemple : Gordon et al. *J. Phys. Chem. A* **2003**, *107*, 8512.

³Ferré et al. *J. Chem. Phys.* **2002**, *117*, 4119.

Molécules organiques : PBE0(BF)/6-311++G**//B3LYP/6-311++G**
(RMSD = 0,16 eV)



atomes C, H
RMSD = 0,09 eV

atomes C, H, N, O
RMSD = 0,19 eV

Etats de cœur ionisés de la liaison peptidique :

PBE0/6-311++G**//B3LYP/6-311++G**

Déviation maximale

Th. vs Th. & Exp. vs Th.

- Ref.¹-This work: $\simeq 0.3$ eV
- Exp.²-This work: $\simeq 0.2$ eV
- Exp.-Ref.: $\simeq 0.2$ eV

⇒ Détermination précise des énergies d'ionisation de cœur (C 1s)



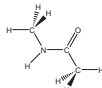
Formamide

*trans* N-méthylformamide*cis* N-méthylformamide

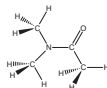
N,N-diméthylformamide



Acétamide



N-méthylacétamide



N,N-diméthylacétamide

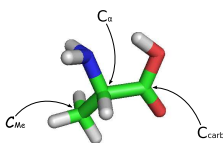


Numérotation des atomes

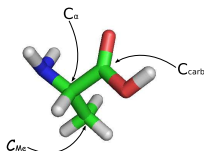
¹Chong et al. *J. Phys. Chem. A* **2002**, *106*, 356.

²Joly et al. *J. Atomic Data and Nuclear Data Tables*, **1984**, *31*, 433.

Conformations: $\Delta E < 1$ kcal/mol



Conf. 1

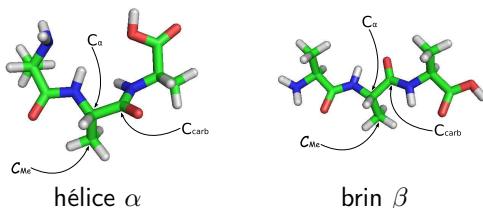


Conf. 2

Energie d'ionisation C 1s (eV) : Alanine

	Conf. 1	Conf. 2	Exp. ²
C _α	292.45(-0.15)	292.16(+0.14)	292.30
C _{carb}	294.60(+0.70)	294.92(+0.38)	295.30
C _{Me}	291.11(+0.09)	290.89(+0.31)	291.20

²Powis et al. *J. Phys. Chem. A* **2003**, 107, 25.

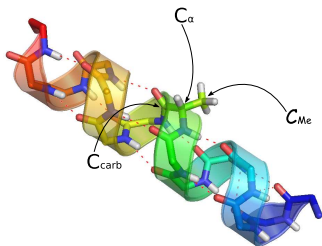
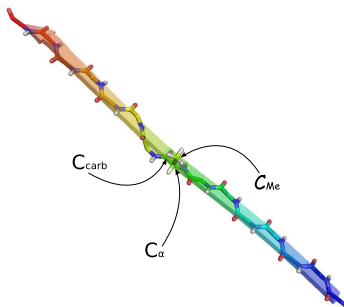
Ala-Triptide: conformation en hélice α et en brin β 

Energies d'ionisation C 1s (eV) : Ala-Triptide

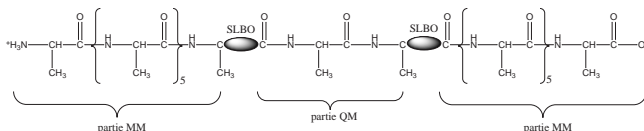
	hélice α	brin β	
C_{α}	291.87	292.00	$\simeq 0.3\text{-}0.4$ eV
C_{carb}	293.69	293.87 ³	$\simeq 2.0$ eV
C_{Me}	290.50	290.62	$\simeq 0.3\text{-}0.4$ eV

³ C_{carb} N-methylacetamide 293.37 eV

Poly-Ala-pentadecapeptide

hélice α brin β

Partition QM/MM du poly-Ala-pentadecapeptide

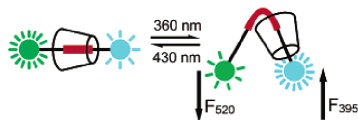


Energies d'ionisation C 1s (eV) : Pentadecapeptide

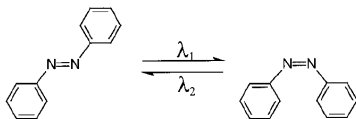
	PBE0/Amber		PBE0/Amber*	
	Hélice α	Brin β	Hélice α	Brin β
C_{α}	-0.26	-0.62	+0.05	+0.08
C_{carb}	-0.09	-0.76	+0.63	+0.23
C_{Me}	-0.71	-0.32	+0.47	+0.11

Applications industrielles et technologiques des dérivés de l'azobenzène

- Colorant en 'absorption' : 60-70% de la production mondiale¹



Isomérisation photoréversible : TAB \rightleftharpoons CAB



⁷Loos et al. *J. Chem. Theor. Comput.* **2008**, 4, 637.

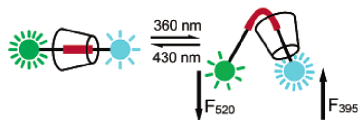
¹Zollinger, H. In *Color Chemistry, Syntheses, Properties and Applications of Organic Dyes and Pigments*; Wiley-VCH, Weinheim, 3 ed., 2003.

²Natansohn et al. *Chem. Rev.*, **2002**, 102, 4139.

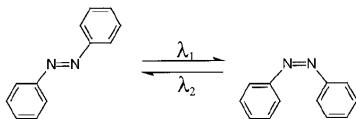
³Qu et al. *Org. Lett.*, **2004**, 6, 2085.

Applications industrielles et technologiques des dérivés de l'azobenzène

- Colorant en 'absorption' : 60-70% de la production mondiale¹
- Périphériques de stockage²
- Moteurs moléculaires³



Isomérisation photoréversible : TAB \rightleftharpoons CAB



¹Loos et al. *J. Chem. Theor. Comput.* **2008**, 4, 637.

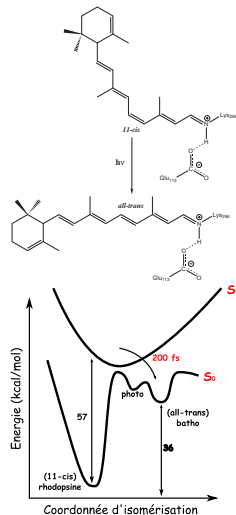
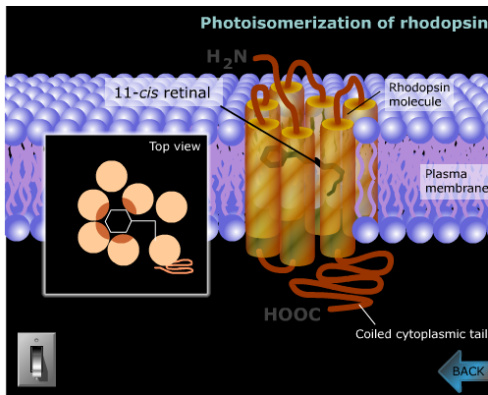
¹Zollinger, H. In *Color Chemistry, Syntheses, Properties and Applications of Organic Dyes and Pigments*; Wiley-VCH, Weinheim, 3 ed., 2003.

²Natansohn et al. *Chem. Rev.*, **2002**, 102, 4139.

³Qu et al. *Org. Lett.*, **2004**, 6, 2085.

<http://www.blackwellpublishing.com/matthews/rhodopsin.html>

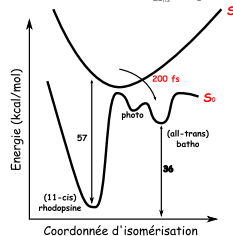
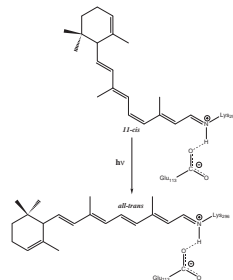
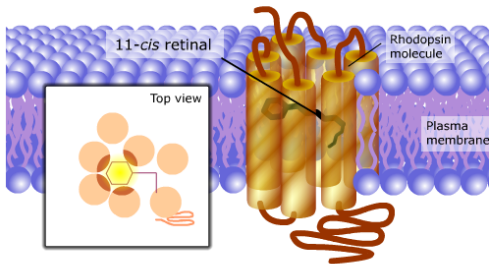
Energie lumineuse \rightleftharpoons Energie mécanique



<http://www.blackwellpublishing.com/matthews/rhodopsin.html>

Energie lumineuse \rightleftharpoons Energie mécanique

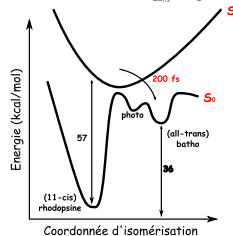
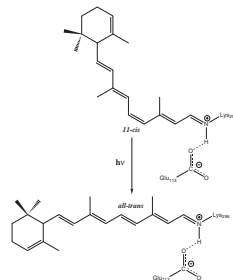
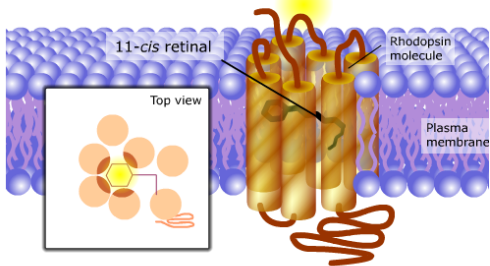
Photoisomerization of rhodopsin



<http://www.blackwellpublishing.com/matthews/rhodopsin.html>

Energie lumineuse \rightleftharpoons Energie mécanique

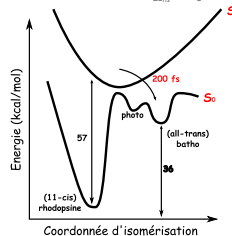
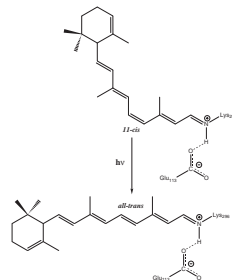
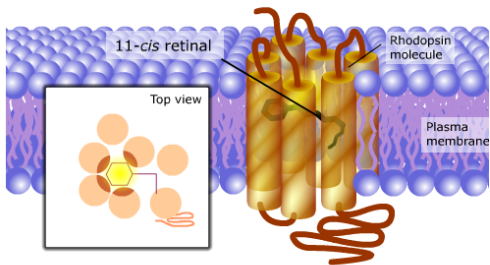
Photoisomerization of rhodopsin



<http://www.blackwellpublishing.com/matthews/rhodopsin.html>

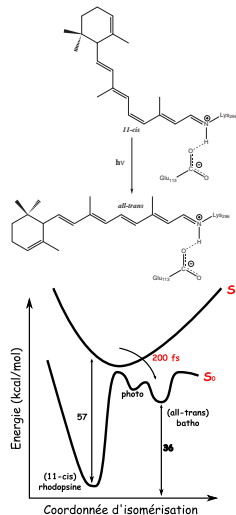
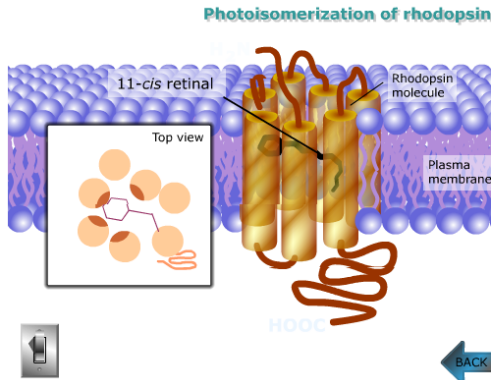
Energie lumineuse \rightleftharpoons Energie mécanique

Photoisomerization of rhodopsin



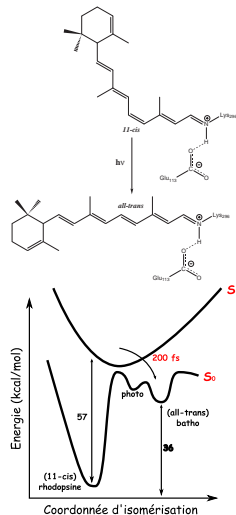
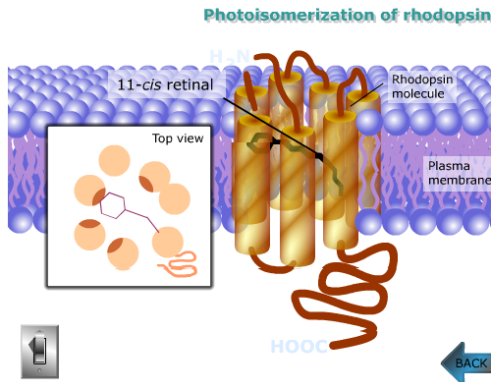
<http://www.blackwellpublishing.com/matthews/rhodopsin.html>

Energie lumineuse \rightleftharpoons Energie mécanique



<http://www.blackwellpublishing.com/matthews/rhodopsin.html>

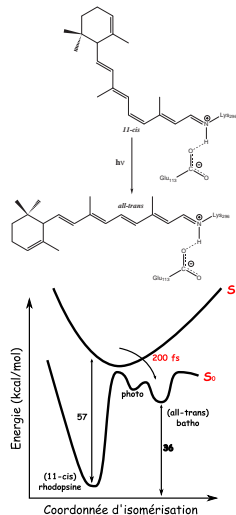
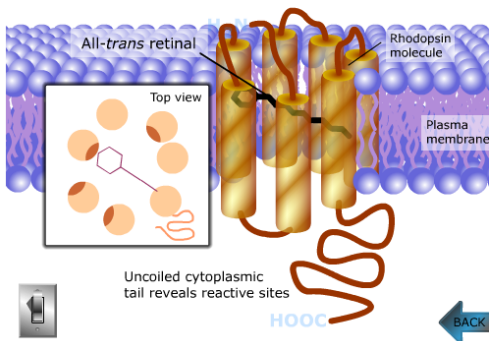
Energie lumineuse \rightleftharpoons Energie mécanique



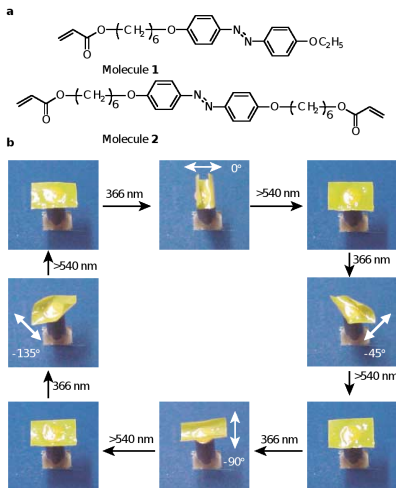
<http://www.blackwellpublishing.com/matthews/rhodopsin.html>

Energie lumineuse \rightleftharpoons Energie mécanique

Photoisomerization of rhodopsin

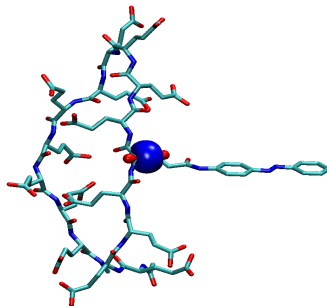
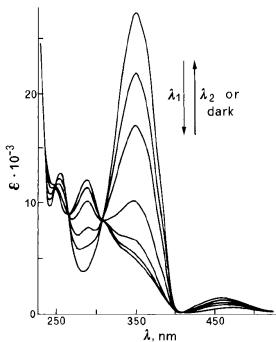


Yu, Y.; M. Makoto, M.; T. Ikeda, T. *Nature*, **2003**, *425*, 145.



Spectre UV-vis : TD-DFT

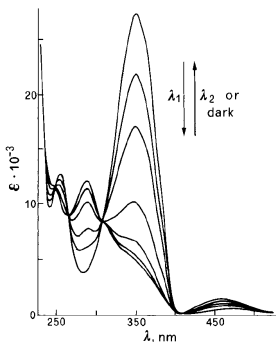
Système macromoléculaire : calculs MM



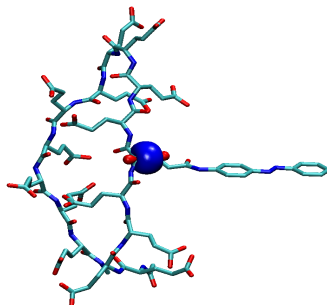
⁶Briquet et al. *Chem. Phys. Lett.*, **2006**, 417, 190.

Spectre UV-vis : TD-DFT

- Géometries: B3LYP/6-311G(d)
- Spectre UV-vis : TD-DFT
PBE0/6-311+G(d)⁶



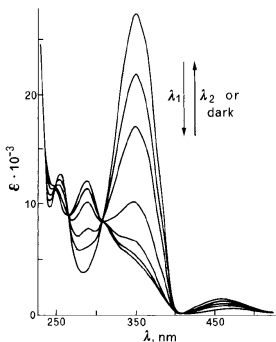
Système macromoléculaire : calculs MM



⁶Briquet et al. *Chem. Phys. Lett.*, **2006**, *417*, 190.

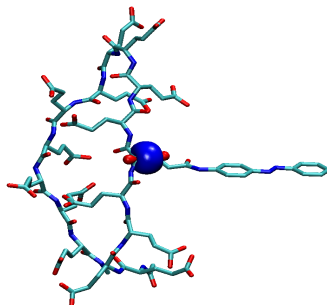
Spectre UV-vis : TD-DFT

- Géometries: B3LYP/6-311G(d)
- Spectre UV-vis : TD-DFT
PBE0/6-311+G(d)⁶



Système macromoléculaire : calculs MM

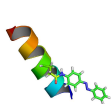
- Champ de force : Amber *ff99*
- GLU protoné
- N et C-terminus: groupement NME et ACE



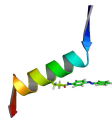
⁶Briquet et al. *Chem. Phys. Lett.*, **2006**, 417, 190.

Géométries QM/MM des Poly(acide L-glutamique) + TAB

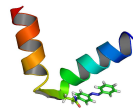
■ Structure α



Hélice α

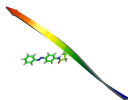


β - α - β

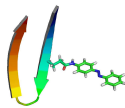


α - β - α

■ Structure β



Brin β

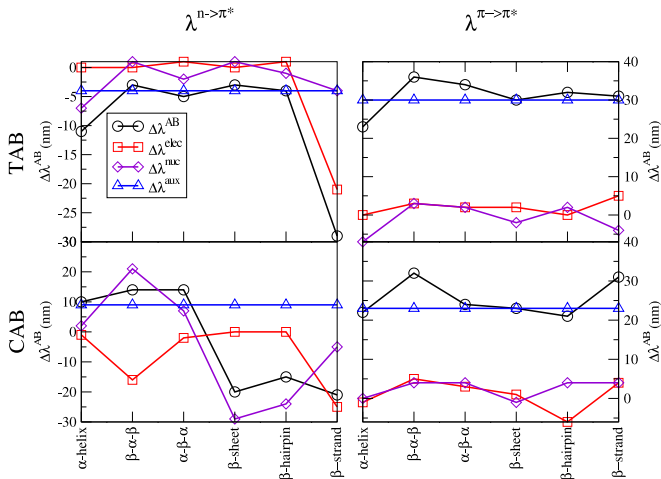


Épingle β



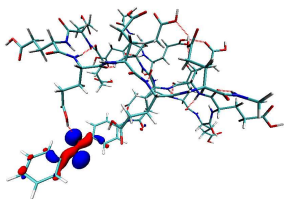
Feuillet β

$$\Delta\lambda^{AB} = \Delta\lambda^{elec} + \Delta\lambda^{nuc} + \Delta\lambda^{aux}$$

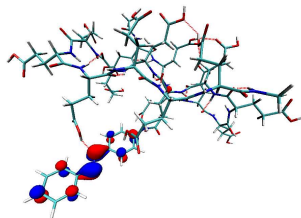


Orbitales frontières de la transition $n \longrightarrow \pi^*$

Orbitale non-liante (n)

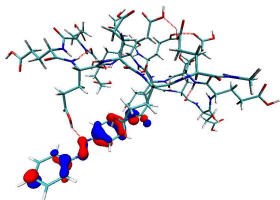


Orbitale anti-liante (π^*)

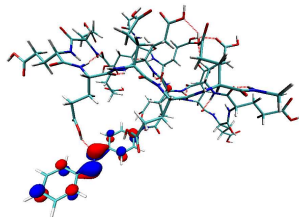


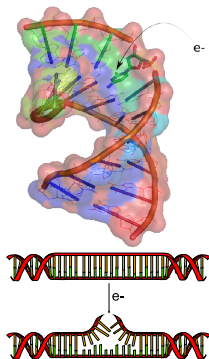
Orbitales frontières de la transition $\pi \longrightarrow \pi^*$

Orbitale liante (π)

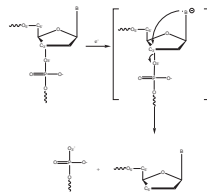


Orbitale anti-liante (π^*)



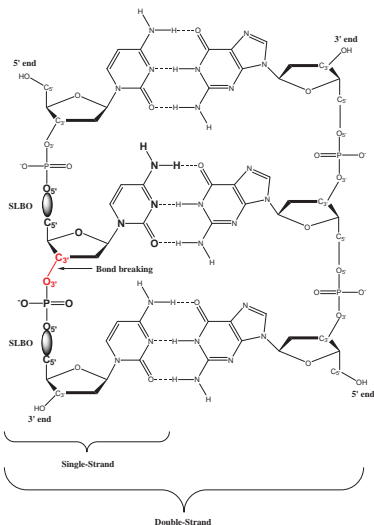
Lésion simple-brin de l'ADN¹

Mécanisme proposé



¹Boudaiffa et al, *Science*, **2000**,287, 1658.

Partition QM/MM



Conclusions

Perspectives

