

## Lecture 3 Applications of EPR spectroscopy to the study of protein dynamics

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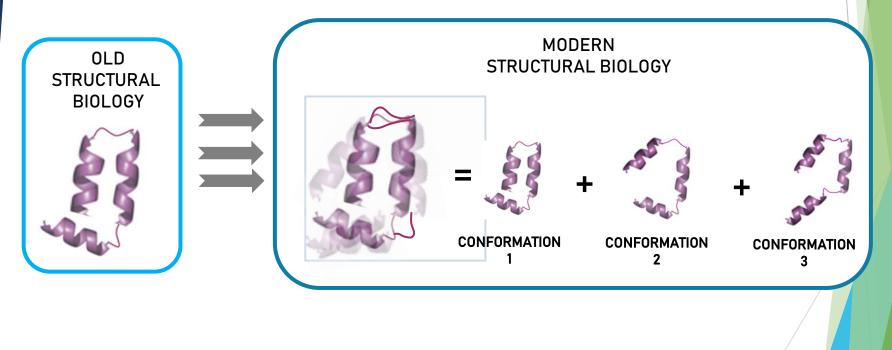


This project has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 101004806

#### INTRODUCTION



#### **MODERN : STRUCTURE = ENSAMBLE OF CONFORMATIONS**

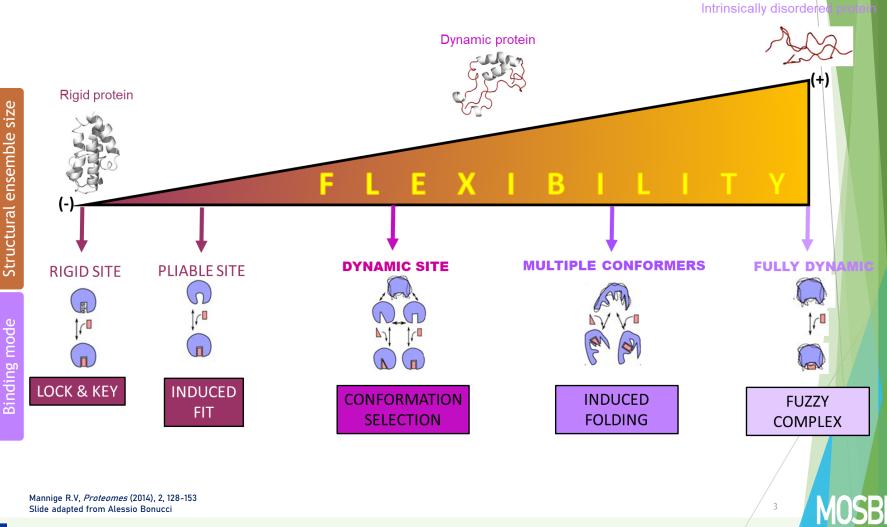


Mannige R.V, *Proteomes* (2014), 2, 128-153 Slide adapted from Alessio Bonucci



#### **INTRODUCTION**

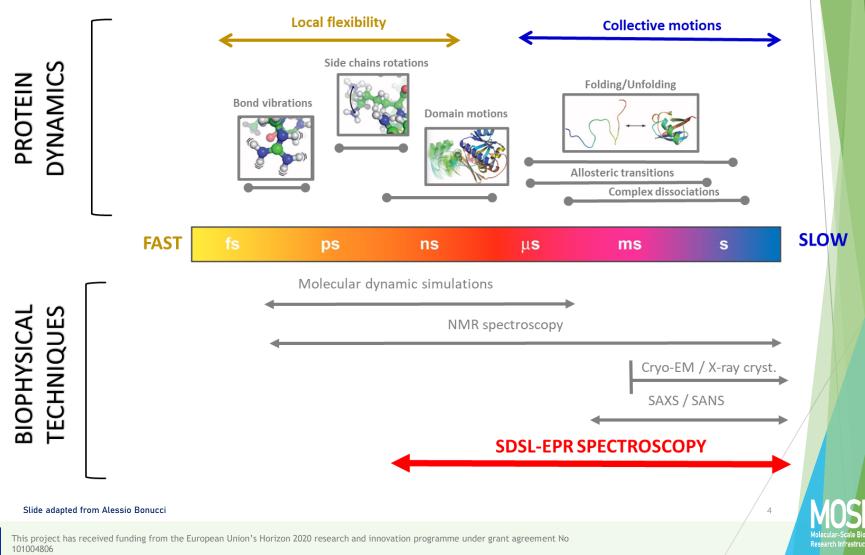
#### **INVESTIGATIONS ON PROTEIN DYNAMICS ARE FUNDAMENTAL** TO DECIPHER THE FUNCTIONS OF A DYNAMIC PROTEINS



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#### INTRODUCTION

#### Different approaches can be exploited to probe protein dynamics

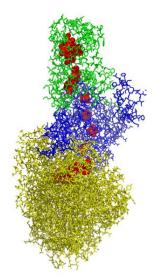


Paramagnetic species in biological systems

EPR = Electron Paramagnetic Resonance (or ESR = Electron Spin Resonance)

Paramagnetic species

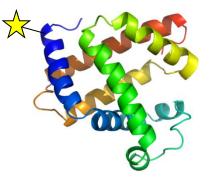
# Intrinsic paramagnetic centers



Metal centers or clusters: Fe, Cu, Ni, Mo ... Radicals: Semiquinone, Tyr °, S °, ...

Catalytic mechanisms, ET mechanisms, ...

Extrinsic paramagnetic label



Site-Directed Spin Labeling (SDSL)

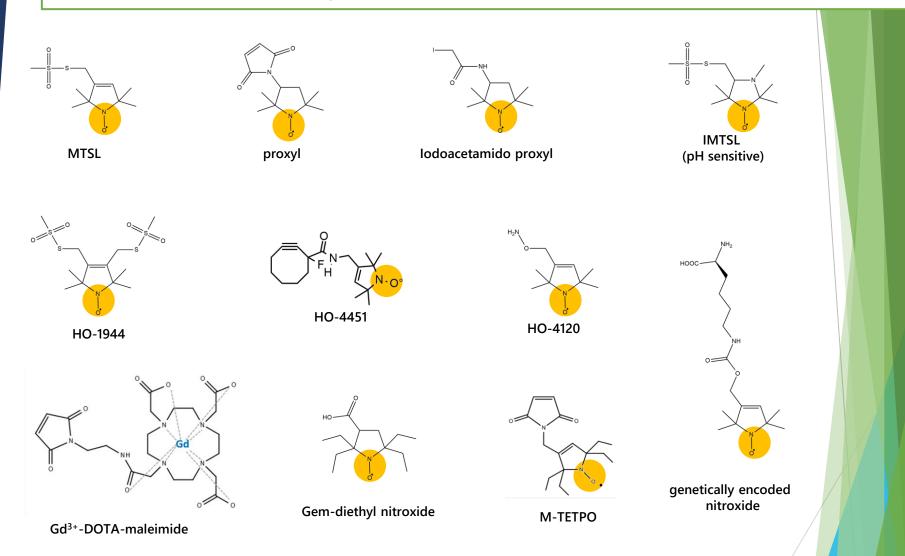
Structural transitions in proteins, protein-protein interactions, ...

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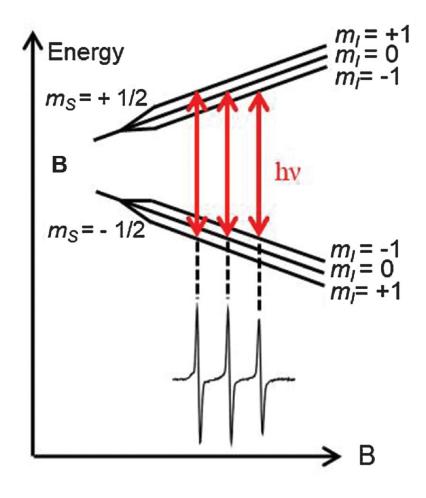
#### Spin labels for biomolecules

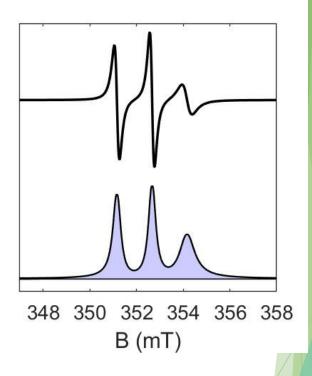


Le Breton, N. *et al. Front. Mol. Biosc.* 2015 ; M. Martinho *et al., Book of the RSC, Electron Paramagnetic Resonance* 2018 *vol. 26* ; Lorenzi, M. *et al. Angew. Chem. Int. Ed.* 2011 ; Mileo, E. *et al. Bioconjugate Chem.* 2013 ; Fleissner, M.R. *et al. PNAS* 2009 ; Kalai, T. *et al. Tetrahedron Lett.* 2011 ; Kucher, S. *et al. JMR* 2017 ; Schmidt *et al. JACS* 2014 ; Paletta, *et al. Org. Lett.* 2012 ; Fleissner, M.R. *et al PNAS* 2011 ; Smirnov, *et al. JACS* 2004 6



### EPR spectroscopy of nitroxide spin labels



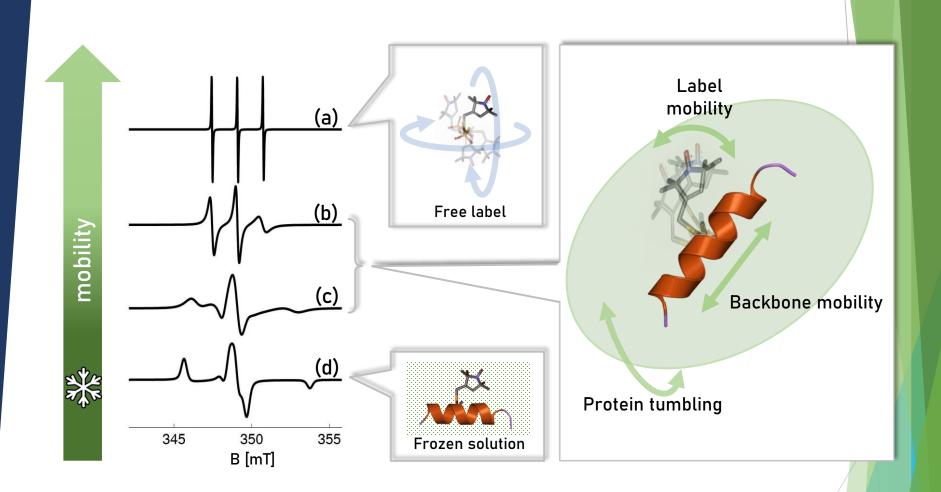


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M. Martinho et al., Book of the RSC, Electron Paramagnetic Resonance vol. 26 2018



#### EPR spectroscopy of nitroxide spin labels

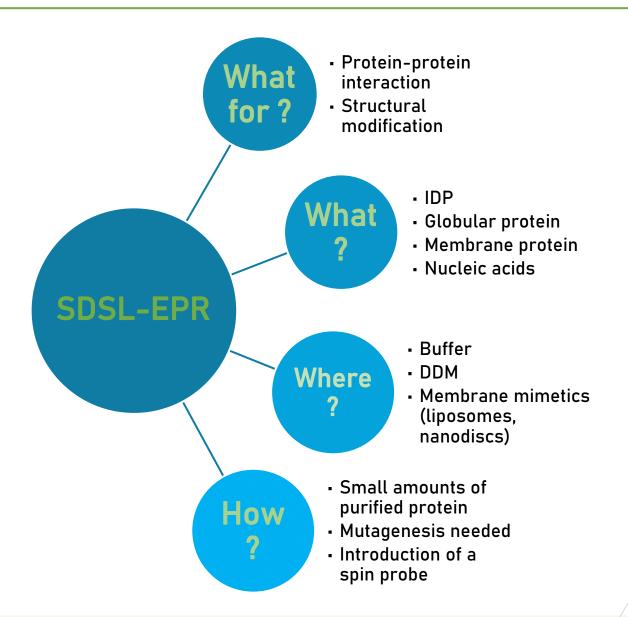


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#### Fournier, E, et al. Actualité Chimique 2019 443, 13-16



#### SDSL-EPR : A powerful technique





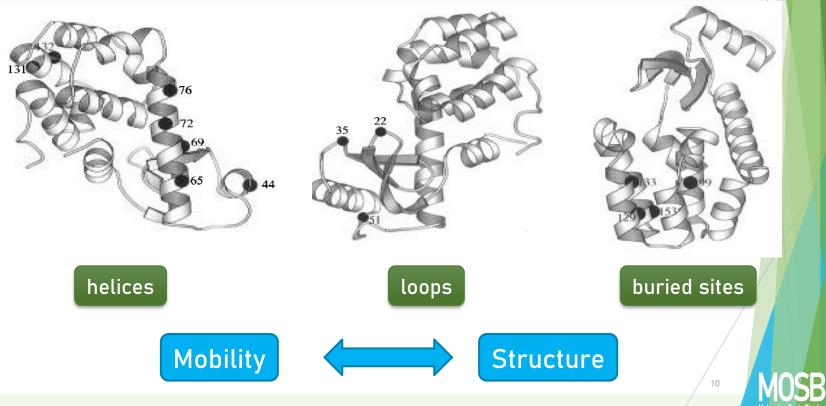
#### The pioneer work

Biochemistry 1996, 35, 7692-7704

## Motion of Spin-Labeled Side Chains in T4 Lysozyme. Correlation with Protein Structure and Dynamics<sup>†</sup>

Hassane S. Mchaourab,<sup>‡</sup> Michael A. Lietzow,<sup>‡</sup> Kalman Hideg,<sup>§</sup> and Wayne L. Hubbell<sup>\*,‡</sup>

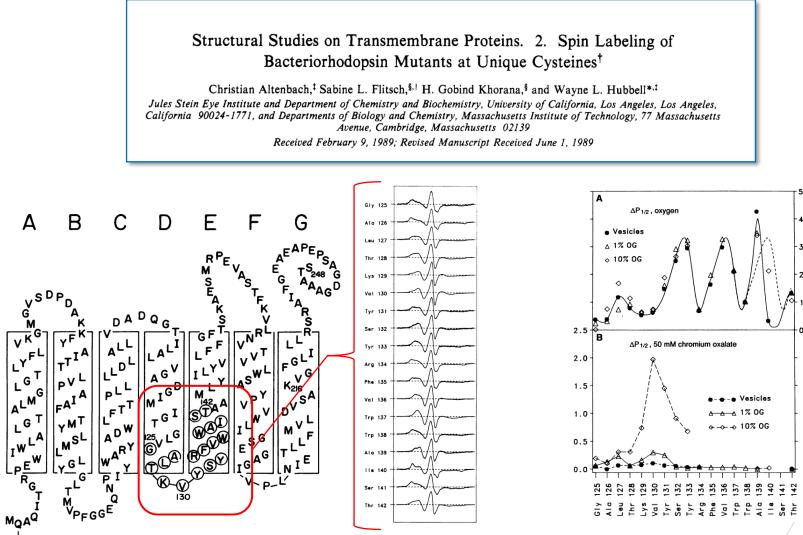
Jules Stein Eye Institute and Department of Chemistry and Biochemistry, University of California, Los Angeles, California 90095-7008, and Central Research Laboratory, Chemistry, University of Pecs, P.O. Box 99, H-7643 Pecs, Hungary



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#### The pioneer work for membrane proteins



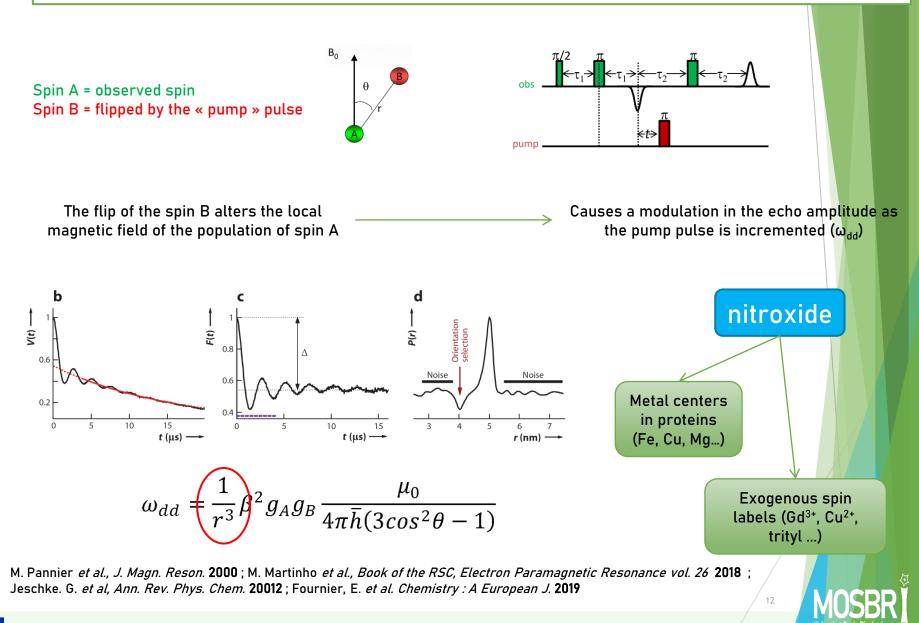
EPR spectra of spin-labeled bacteriorhodopsin mutants reconstituted in soybean lipid vesicles

The accessibility parameter ΔP<sub>1/2</sub> for oxygen (Å) or chromium oxalate (B) versus position of the spin label in the bacteriorhodopsin sequence.

Altenbach, C. *et al. Biochem.* **1989** *28*, 7806-7812 ; Altenbach, C. *et al. Science* **1990** *248*, 1088-1092



#### **DEER** : distance measurements

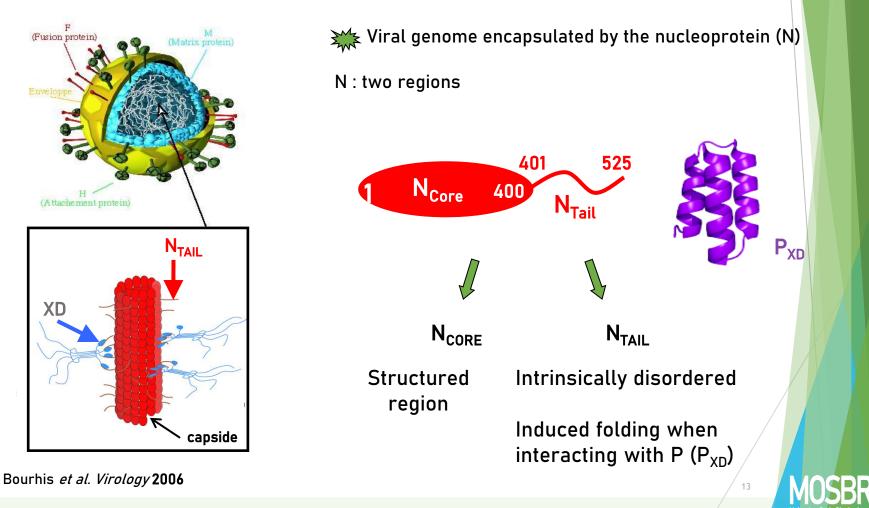




### Application 1 : Induced folding in nucleoprotein virus

Main cause of childhood mortality in developing countries (164 000 deaths, OMS 2008). To date, no antiviral treatment exists.

 $\rightarrow$  Understanding the mechanism of action of proteins involved in the replicative complex

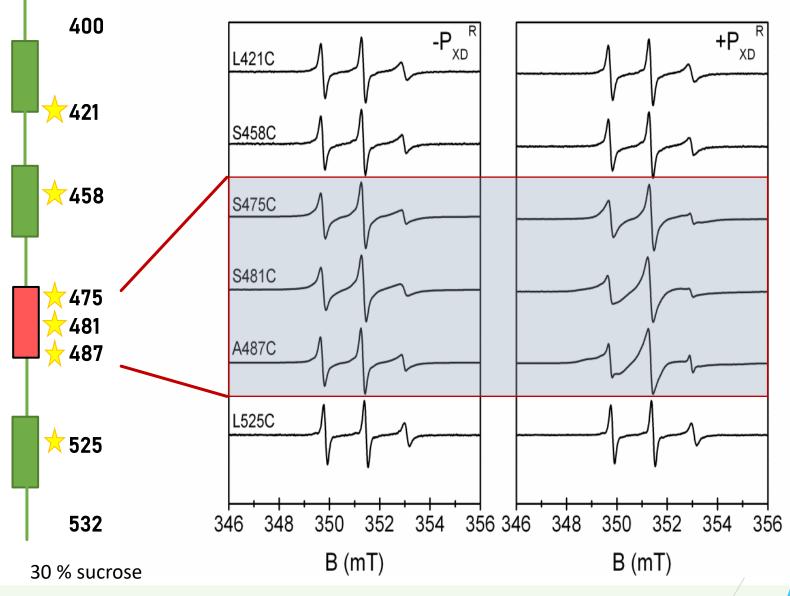


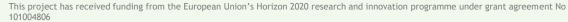


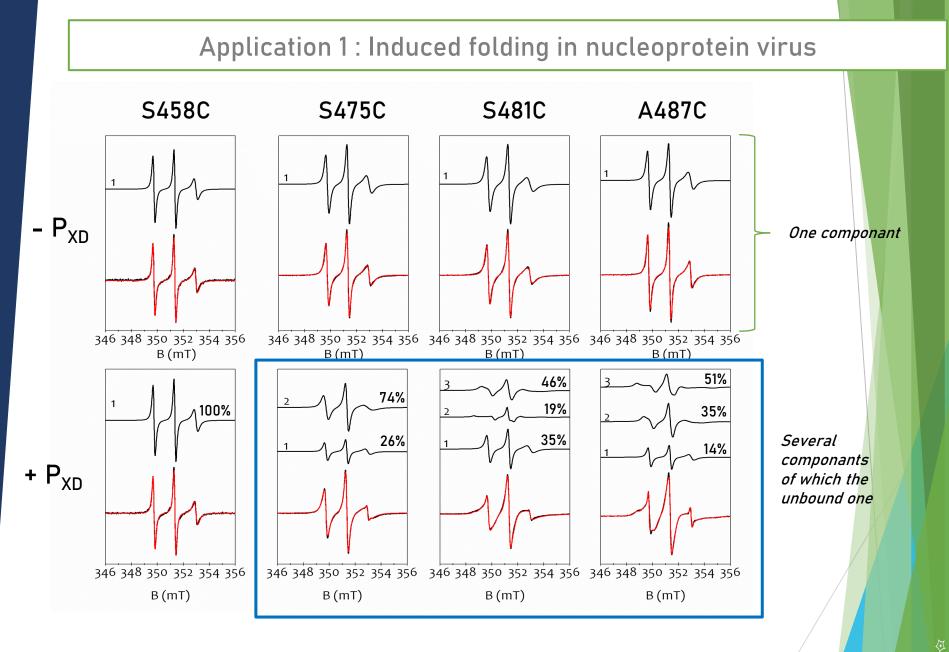
### Application 1 : Induced folding in nucleoprotein virus

Molecul

Research Infrastructur

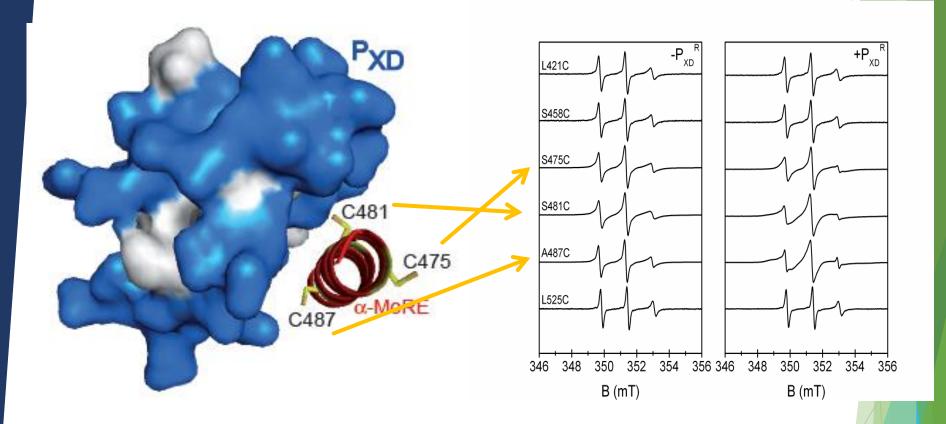








### Application 1: Induced folding in nucleoprotein virus



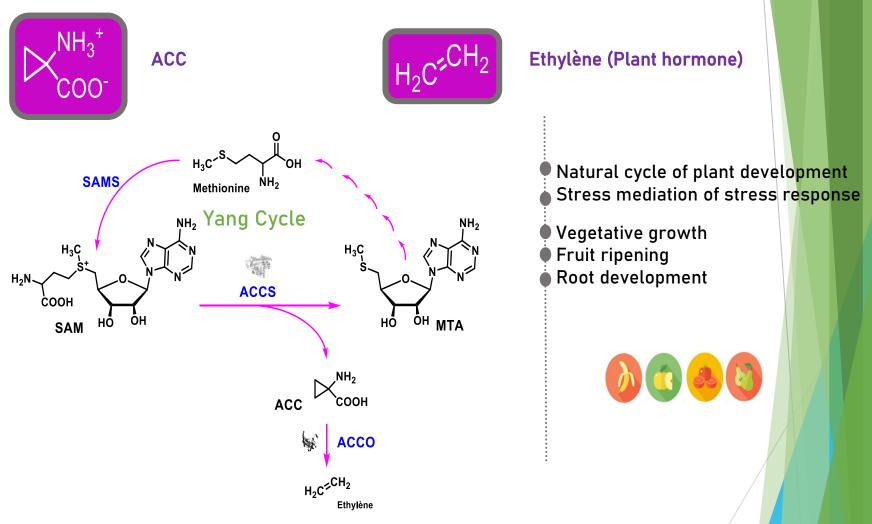
Validation of structural model of Hev N<sub>TAIL</sub>-P<sub>XD</sub> complex

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#### Habchi *et al,* JBC 2011

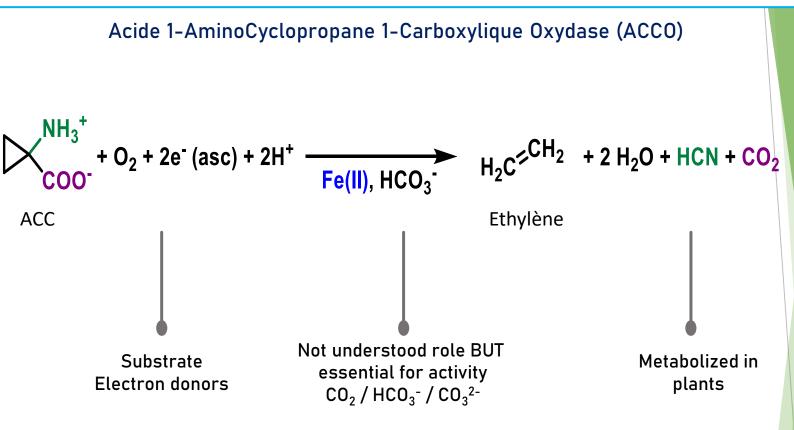


#### Acide 1-AminoCyclopropane 1-Carboxylique Oxydase (ACCO)



D.P. Murr et al., Plant Physiology. 1975 ; A. Bleecker et al., Annu. Rev. Cell. Dev. Biol. 2000









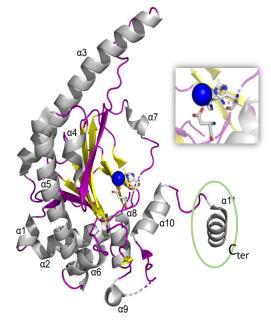
## 2

Crystallographic Structure from *Petunia hybrida* Z. Zhang *et al., Chem. Biol.* **2004** 

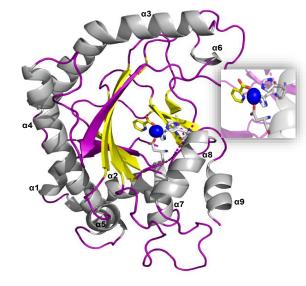
#### Crystallographic Structure from *Arabidopsis thaliana* X. Sun *et al., Nat. Commun.* **2017**

## 3

Structural models from *Malus domestica* and *Lycopersicum esculentum* Z. Zhang *et al., Chem. Biol.* 2004 ; L. Brisson et *al., JBIC.* 2012



2 formes : apo and holo : ACCO·Fe(II)
Tetramere : C<sub>term</sub> extremity from each monomere interacts with C<sub>term</sub> extremity from 2<sup>nd</sup> monomere C<sub>term</sub>



- 2 forms : ACCO·Zn(II) with inhibitors
- 17 truncated residues in C<sub>term</sub> part

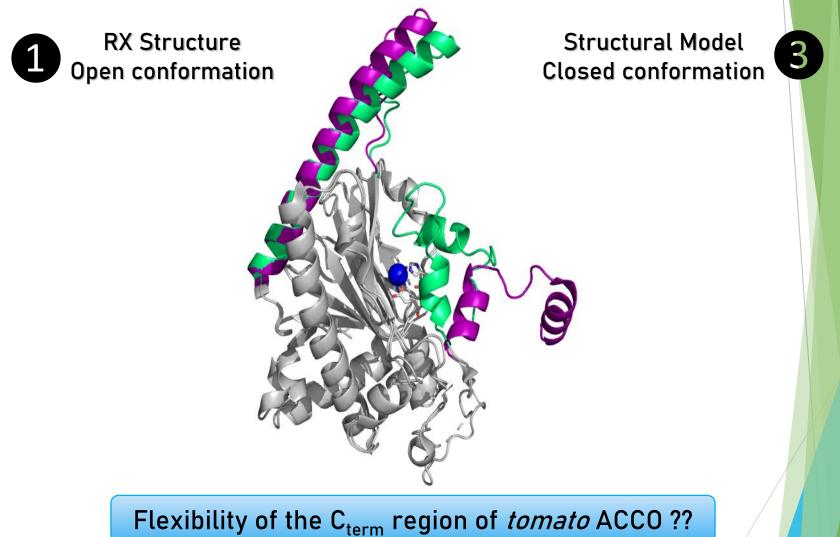
#### Not active conformation

Inhibited form

# Proposed active conformation

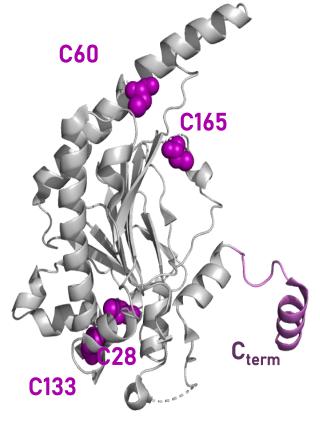
Other conformation of C<sub>term</sub> part



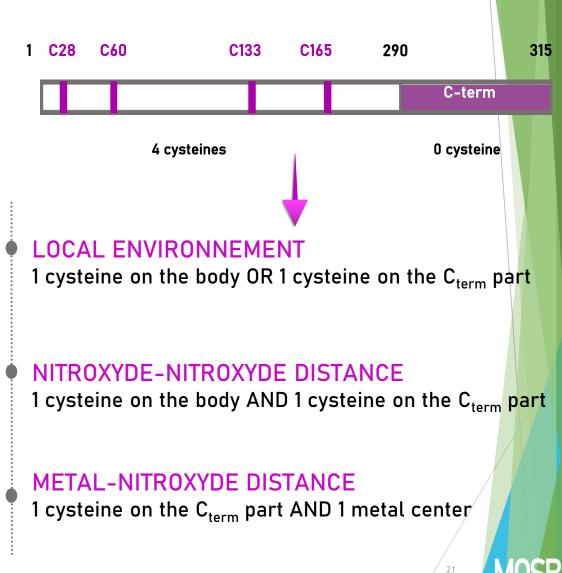








Tomato ACCO

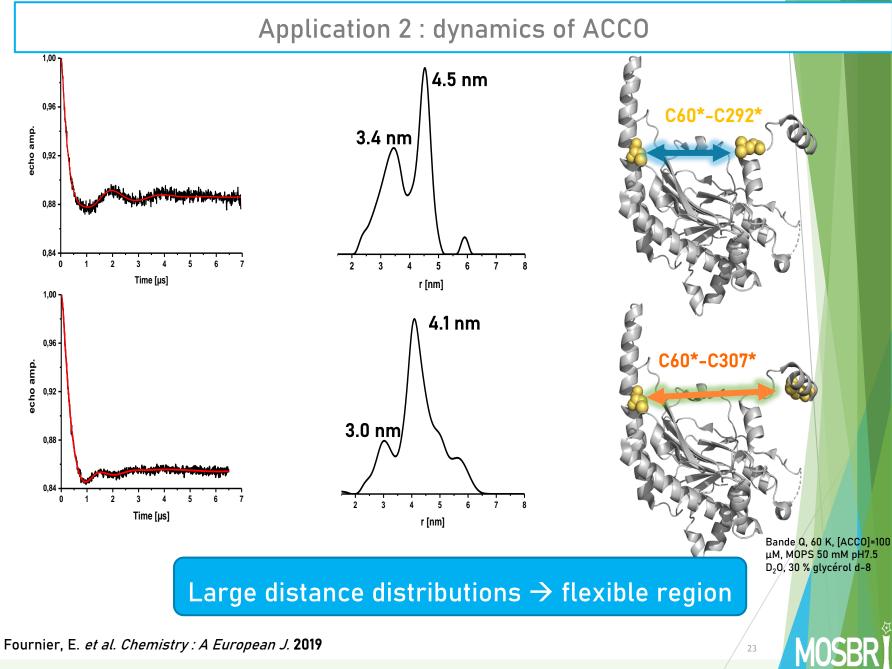




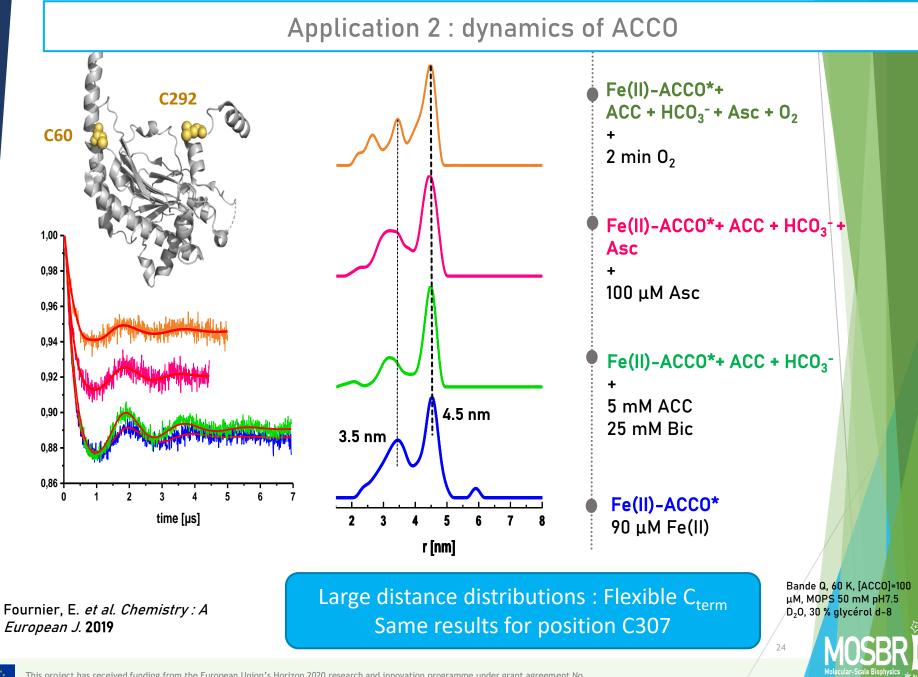


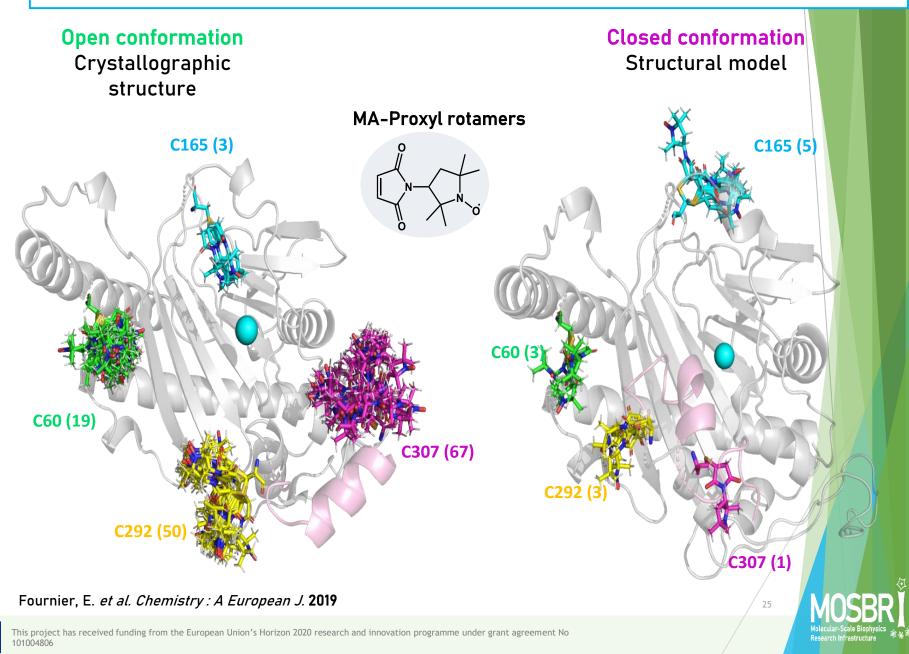


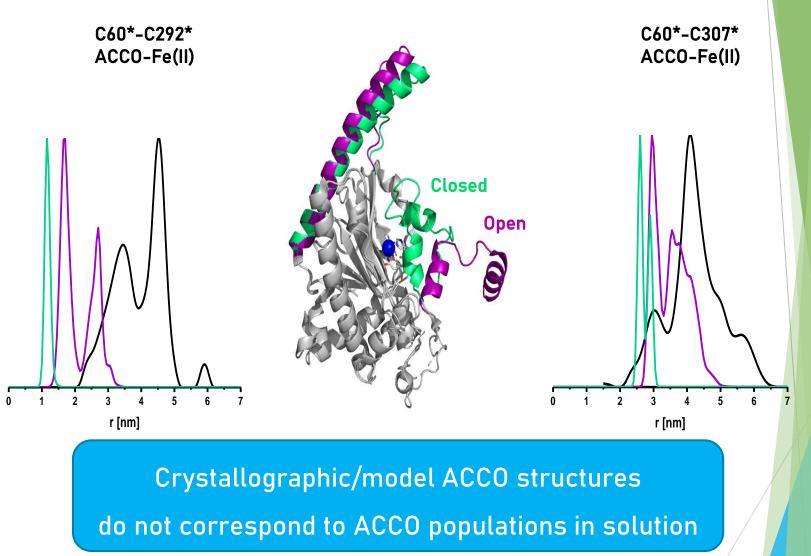
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Fournier, E. et al. Chemistry : A European J. 2019



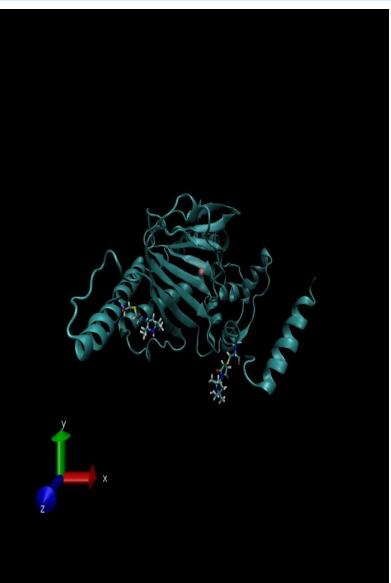




The University of Manchester

Sam P. de Visser Nick Fowler

C60\*-C292\* ACCO-Fe(II)

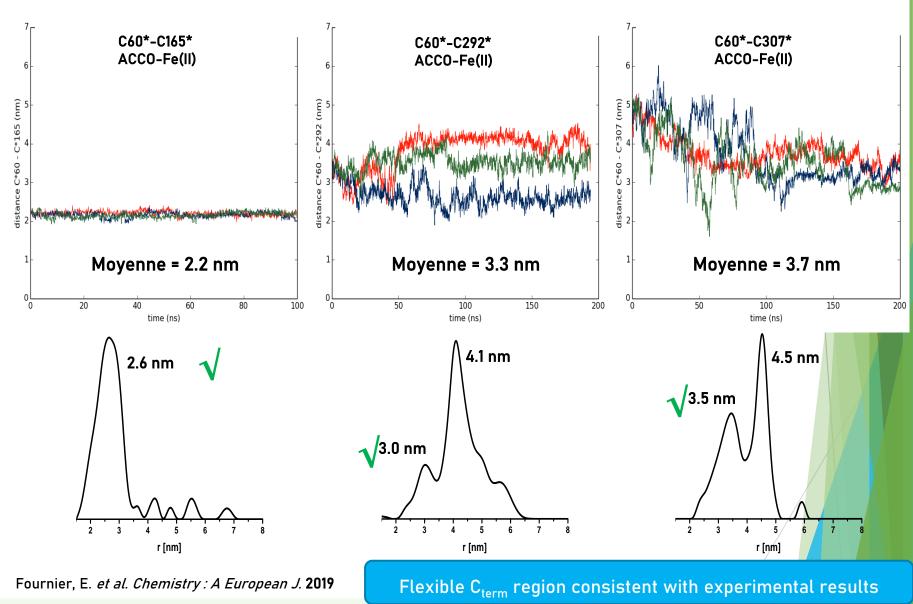


Fournier, E. et al. Chemistry : A European J. 2019



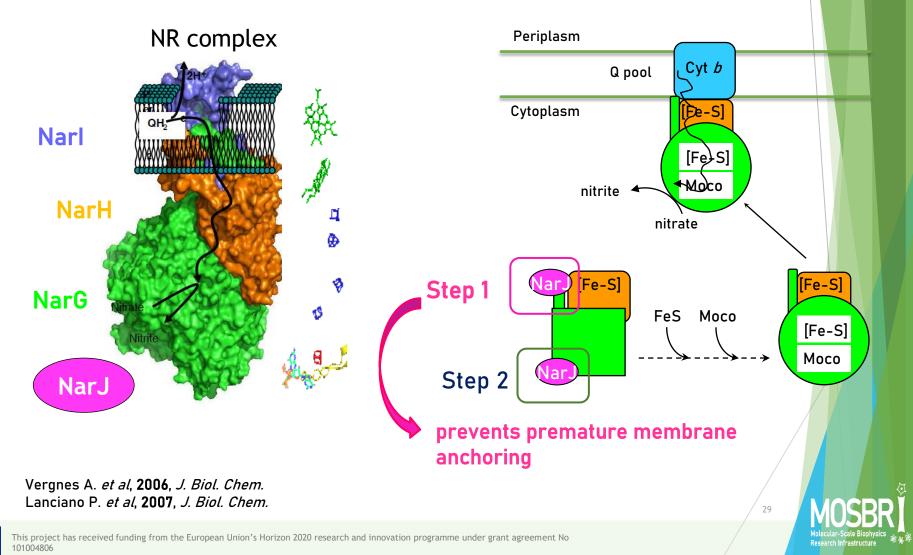
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Molecular-Scale Biophysics Research Infrastructure Application 3 : chaperone protein NarJ : looking for an interaction site

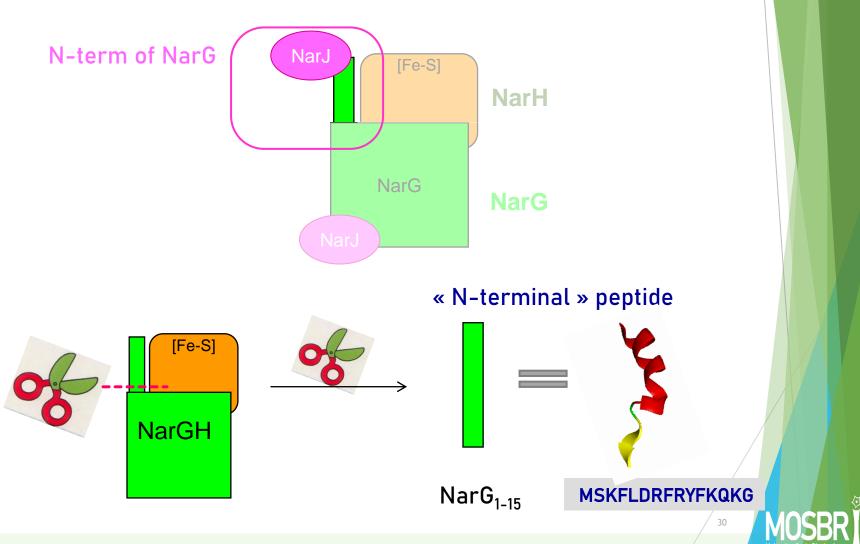
**NarJ** is involved in the **biogenesis** of the respiratory nitrate reductase (NR) complex *in E. coli* 



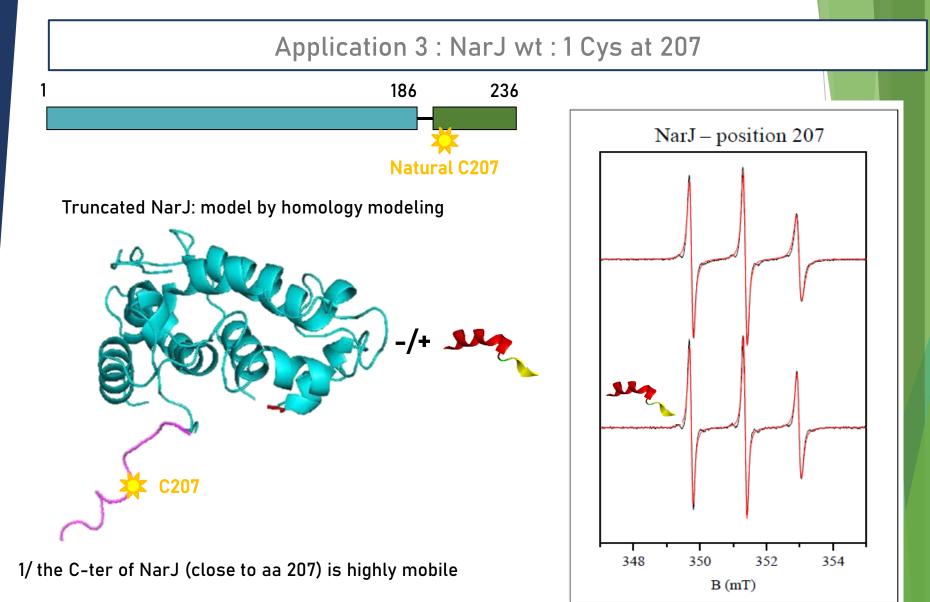


Application 3 : chaperone protein NarJ : looking for an interaction site

#### initial step : the recognition between NarJ and NR



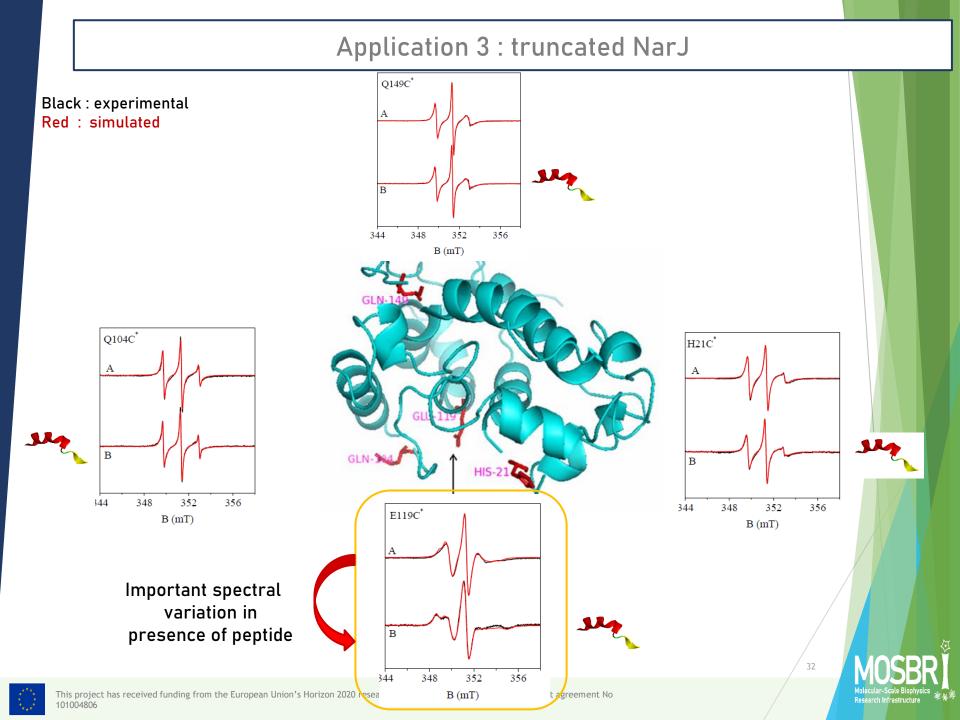


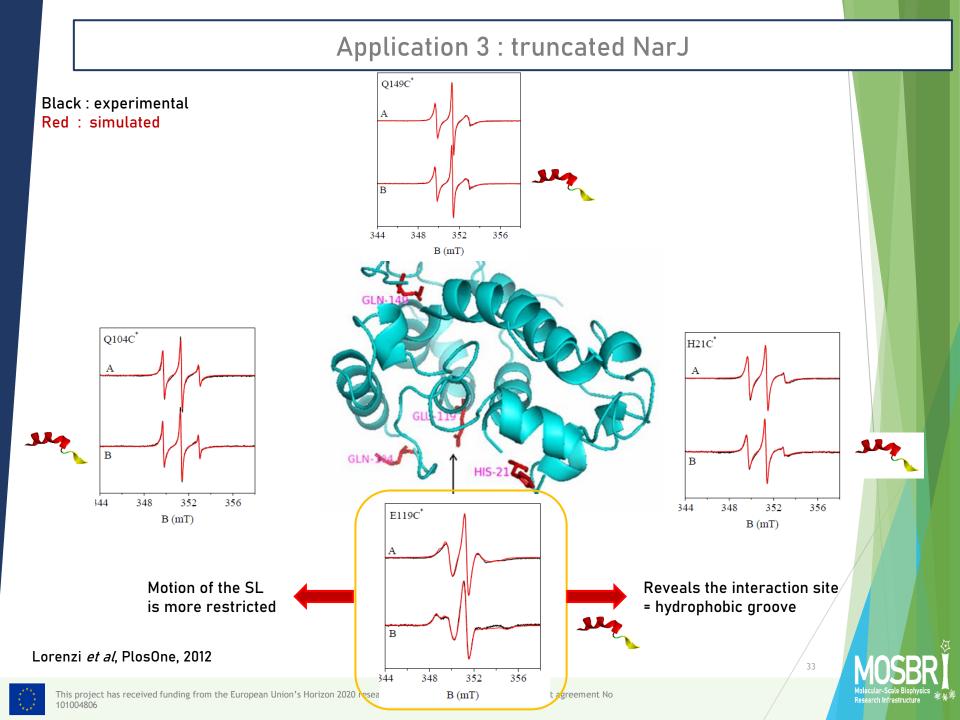


2/ Remains disordered in the presence of the peptide partner

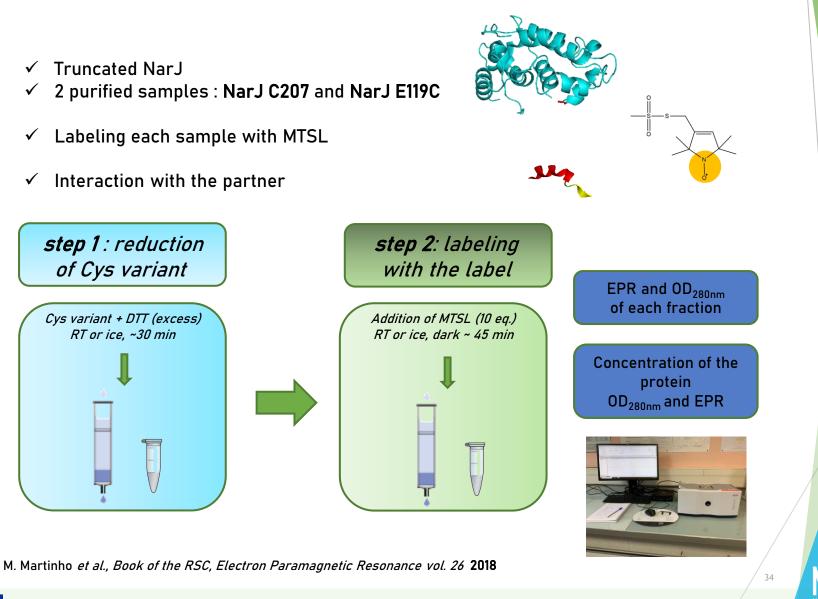




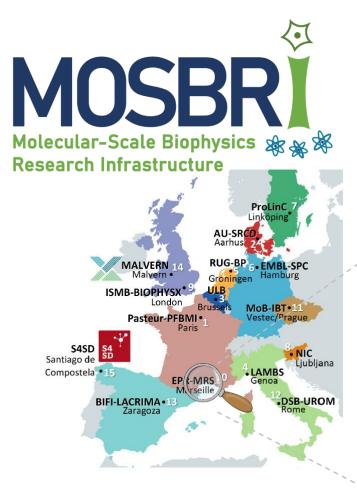




#### Application 3 : In the practicals



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# *Thank you for your attention !*

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