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#### **RESEARCH INTERESTS**

- Biochemistry of metalloenzymes
- Cytochromes P450, Flavins
- Enzyme catalysis and mechanism
- Coenzyme Q biosynthesis
- Fe-S hydroxylating non O<sub>2</sub>-dependent enzymatic system

#### **SUMMARY**

Coenzyme Q (CoQ) is an endogenous lipid soluble molecule found in bacterial plasmatic or mitochondrial inner membranes, where it works as an essential electron carrier in the respiratory chain. Today, several inactivating mutations in genes affecting the biosynthesis of CoQ have been identified in severe pathologies as myopathies, nephropathies, or cerebellar ataxia. CoQ biosynthesis is complex and requires no less than a dozen of proteins, but a clear assignment of catalytic or biological function for many participating proteins is still missing. We aim at getting a better knowledge at the fundamental level of the biosynthesis of CoQ in *E.coli*, using a judicious combination of molecular genetics, site-directed mutagenesis, biochemistry, molecular biophysics (rapid-kinetics) and structural biology (X-Ray crystallography). The core of this project will be to provide the first *in vitro* biochemical characterization of the proteins involved in the biosynthesis of CoQ in *E.coli*, to elucidate their exact function, to solve their X-Ray structures and to study their molecular mechanism.

I am also interested in RNA modifying enzymes, notably TrhP and RlhA, two Fe-S proteins that share biochemical and structural identity with UbiU and UbiV involved in anaerobic CoQ biosynthesis in *E.coli*.

#### **PUBLICATIONS**

##### **2021**

- An enzymatic activation of formaldehyde for nucleotide methylation.  
Bou-Nader, C, Stull, FW, Pecqueur, L, Simon, P, Royant, F, Fontecave, M, **LOMBARD, M**, Palfey, BA, Hamdane, D.  
*Nature Comm.* 2021, 12(1):4542, doi: 10.1038/s41467-021-24756-8.
- Dihydrouridine synthesis in tRNAs under reductive evolution  
Faivre, B, **LOMBARD, M**, Fakroun, S, Vo, CDT, Goyenvalle, C, Guérineau, V, Pecqueur, L, Fontecave, M, de Crécy-Lagard, V, Brégeon, D, Hamdane, D.  
*RNA biol.* 2021, doi: 10.1080/15476286.2021.1899653

## 2020

- The O<sub>2</sub>-independent pathway of ubiquinone biosynthesis is essential for denitrification in *Pseudomonas aeruginosa*  
Vo, CDT, Michaud, J, Elsen, S, Faivre, B, Bouveret, E, Barras, F, Fontecave, M, Pierrel, F, **LOMBARD, M\***, Pelosi, L.  
*J Biol Chem.* 2020, doi: 10.1074/jbc.RA120.013748
- Structural and functional characterization of 4-Hydroxyphenylacetate 3-hydroxylase from *Escherichia coli*.  
Deng, Y, Faivre, B, Back, O, **LOMBARD, M**, Pecqueur, L, Fontecave, M.  
*ChemBioChem.* 2020, doi: 10.1002/cbic.201900277

## 2019

- Ubiquinone Biosynthesis over the Entire O<sub>2</sub> Range: Characterization of a Conserved O<sub>2</sub>-Independent Pathway.  
Pelosi, L, Vo, C-D-T, Abby, S, Loiseau, L, Rascalou, B, Hajj Chehade, M, Faivre, B, Goussé, M, Chenal, C, Touati, N, Binet, L, Cornu, D, Fyfe, C, Fontecave, M, Barras, F, **LOMBARD, M\***, Pierrel, F.  
*mBio.* 2019, 10, e01319-19, doi: 10.1128/mBio.01319-19.
- A soluble metabolon synthesizes the isoprenoid lipid Ubiquinone.  
Hajj Chehade, M, Pelosi, L, Fyfe, CD, Loiseau, L, Laussel, L, Rascalou, B, Brugiére, S, Vo, CDT, Kazemzadeh, K, Couté, Y, Ciccone, L, Fontecave, M, Barras, F, **LOMBARD, M**, Pierrel F.  
*Cell Chem Biol.* 2019, 26, 482-492. doi: 10.1016/j.chembiol.2018.12.001.

## 2017

- Power of protein/tRNA functional assembly against aberrant aggregation.  
Bou-Nader, C, Pecqueur, L, Cornu, D, **LOMBARD, M**, Dezi, M, Nicaise, M, Velours, C, Fontecave, M, Hamdane, D.  
*Phys Chem Chem Phys.* 2017, 19(41):28014-28027. doi: 10.1039/c7cp05599d.
- Flavin-dependent epitranscriptomic world.  
**LOMBARD, M**, Hamdane D.  
*Arch Biochem Biophys.* 2017, 632:28-40. doi: 10.1016/j.abb.2017.06.011.
- The UbiK protein is an accessory factor necessary for bacterial ubiquinone (UQ) biosynthesis and forms a complex with the UQ biogenesis factor UbiJ.  
Loiseau, L, Fyfe, C, Aussel, L, Hajj Chehade, M, Hernández, SB, Faivre, B, Hamdane, D, Mellot-Draznieks, C, Rascalou, B, Pelosi, L, Velours, C, Cornu, D, **LOMBARD, M**, Casadesús, J, Pierrel, F, Fontecave, M, Barras, F.  
*J Biol Chem.* 2017, 292(28):11937-11950. doi: 10.1074/jbc.M117.789164.

## 2016

- A chemical chaperone induces inhomogeneous conformational changes in flexible proteins.  
Hamdane, D, Velours, C, Cornu, D, Nicaise, M, **LOMBARD, M**, Fontecave, M.  
*Phys Chem Chem Phys.* 2016, 18(30):20410-21. doi: 10.1039/c6cp03635j.

- Coenzyme Q Biosynthesis: Evidence for a Substrate Access Channel in the FAD-Dependent Monooxygenase Coq6.  
Ismail, A, Leroux, V, Smadja, M, Gonzalez, L, **LOMBARD, M**, Pierrel, F, Mellot-Draznieks, C, Fontecave, M.  
*PLoS Comput Biol.* 2016, 12(1): e1004690. doi: 10.1371/journal.pcbi.1004690.

## 2015

- Expression in yeast, new substrates, and construction of a first 3D model of human orphan cytochrome P450 2U1: Interpretation of substrate hydroxylation regioselectivity from docking studies.  
Ducassou, L, Jonasson, G, Dhers, L, Pietrancosta, N, Ramassamy, B, Xu-Li, Y, Loriot, MA, Beaune, P, Bertho, G, **LOMBARD, M**, Mansuy, D, André, F, Boucher, JL.  
*Biochim Biophys Acta.* 2015, 1850(7):1426-37. doi: 10.1016/j.bbagen.2015.03.014.

## 2014

- Biosynthesis and physiology of coenzyme Q in bacteria.  
Aussel, L, Pierrel, F, Loiseau, L, **LOMBARD, M**, Fontecave, M, Barras, F.  
*Biochim Biophys Acta.* 2014, 1837(7):1004-11. doi: 10.1016/j.bbabi.2014.01.015.

## 2013

- Ubil, a new gene in *Escherichia coli* coenzyme Q biosynthesis, is involved in aerobic C5-hydroxylation.  
Hajj Chehade, M, Loiseau, L, **LOMBARD, M**, Pecqueur, L, Ismail, A, Smadja, M, Golinelli-Pimpaneau, B, Mellot-Draznieks, C, Hamelin, O, Aussel, L, Kieffer-Jaquinod, S, Labessan, N, Barras, F, Fontecave, M, Pierrel, F.  
*J Biol Chem.* 2013, 288(27):20085-92. doi: 10.1074/jbc.M113.480368.

## 2011

- Intermolecular electron transfer in two-iron superoxide reductase: a putative role for the desulforedoxin center as an electron donor to the iron active site.  
Bonnot, F, Duval, S, **LOMBARD, M**, Valton, J, Houée-Levin, C, Nivière, V.  
*J Biol Inorg Chem.* 2011, 16(6):889-98. doi: 10.1007/s00775-011-0788-5.
- A new cytochrome P450 belonging to the 107L subfamily is responsible for the efficient hydroxylation of the drug terfenadine by *Streptomyces platensis*.  
**LOMBARD, M\***, Salard, I, Sari, MA, Mansuy, D, Buisson, D.  
*Arch Biochem Biophys.* 2011, 508(1):54-63. doi: 10.1016/j.abb.2011.01.008.

## 2010

- Biooxidation of methyl group: Part 2. Evidences for the involvement of cytochromes P450 in microbial multistep oxidation of terfenadine.  
El Ouarradi, A, **LOMBARD, M**, Buisson, D.  
*J Mol Catal B.* 2010, 67(3-4), 172-178. doi: 10.1016/j.molcatb.2010.07.017.