



Mieux comprendre la physiopathologie des amyloses pour mieux traiter

Christophe Sirac

Team BioPIC (Biology of Plasma cell, Immunopathology and Cancer)

CNRS UMR 7276 - INSERM U1262 - Control of the Immune Response B and Lymphoproliferations (CRIBL)

Université de Limoges



InsERM



Université
de Limoges





Liens d'intérêts

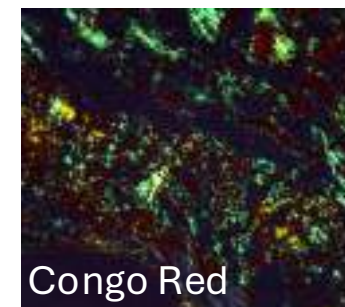
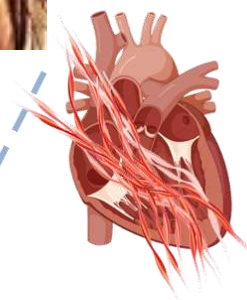
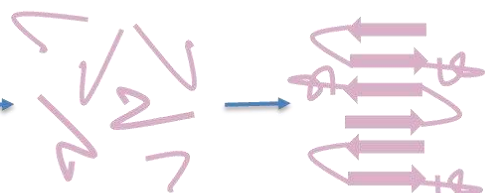
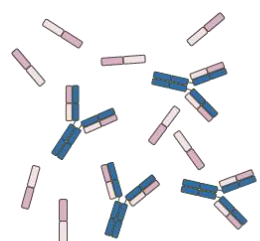
- Research grants from Attralus, Gate Bioscience, Sebia, AmylTherapeutics



Producing cells
The (not) good



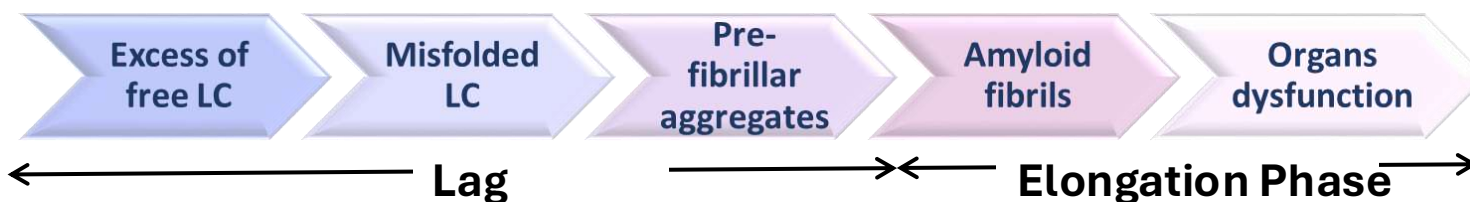
Unstable protein
The ugly



Deposits

The bad

**Organ
dysfunction**

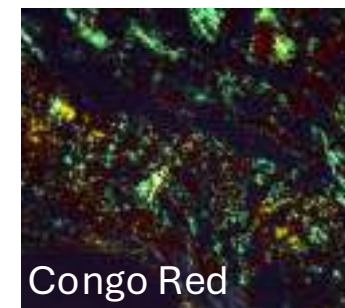
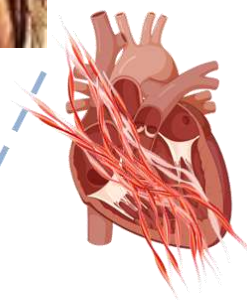
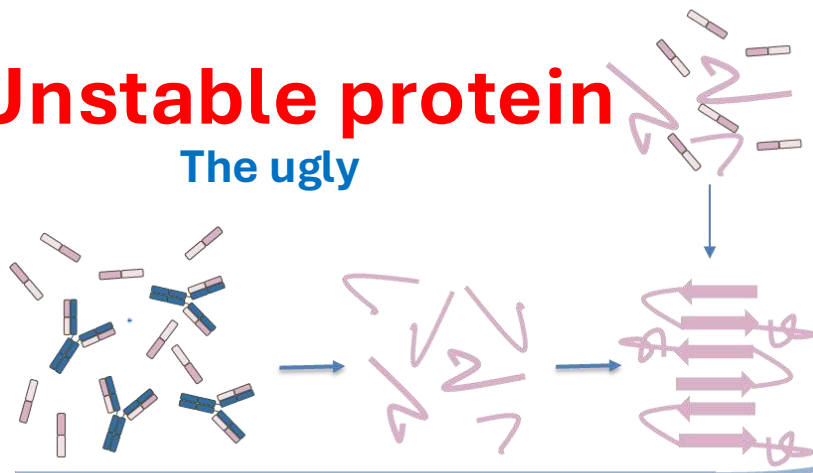




Producing cells
The (not) good



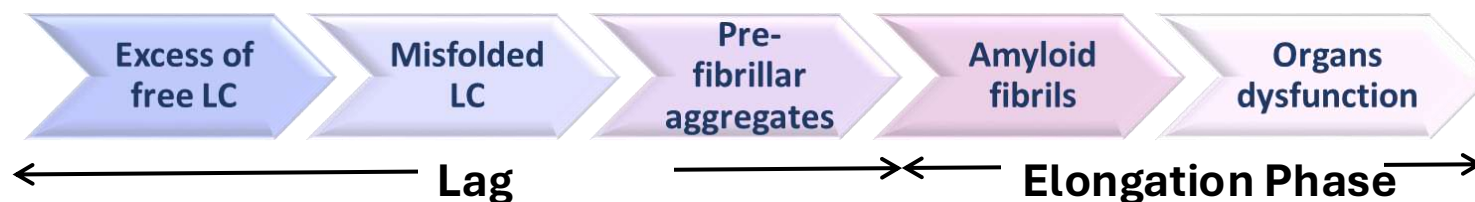
Unstable protein
The ugly



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**Organ
dysfunction**





Amyloses ATTR, AA...

But: Réduire la production de la protéine amyloïdogène **mais** ne pas tuer la cellule productrice
siRNA (patisiran), ASO (inotersen), Crispr-Cas9 (Nex-
Z)...

➤ **Double avantage:** Ciblage du foie et peu de variabilité de séquences dans les protéines

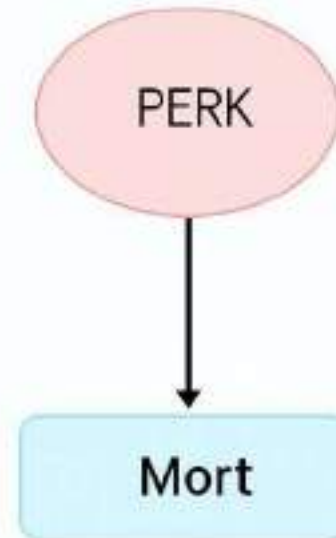
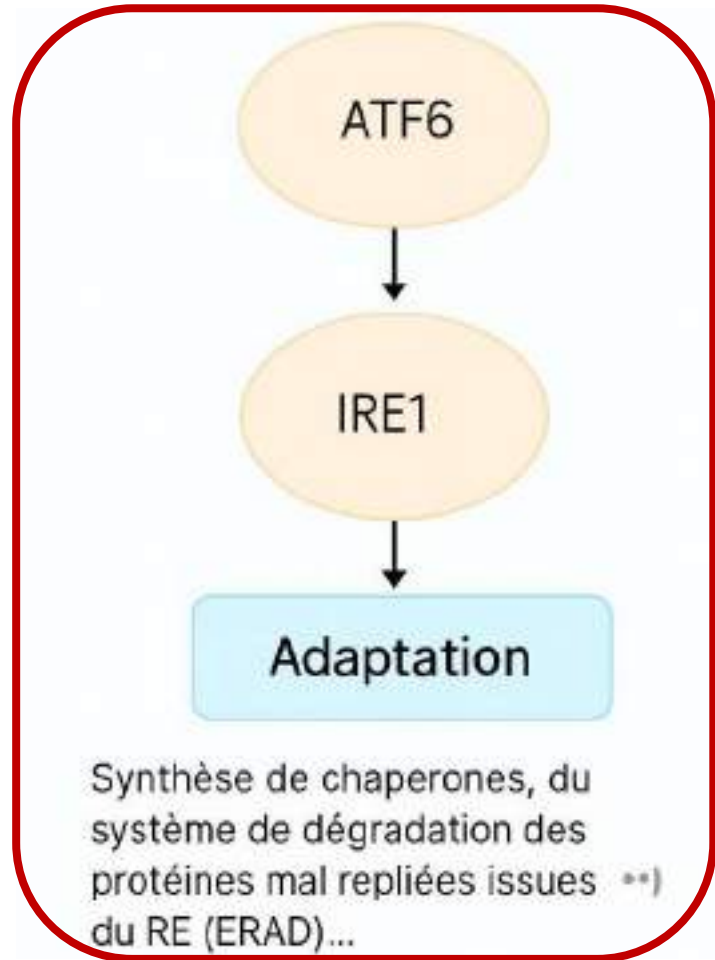


Amylose AL

Buts: Réduire la production de la LC amyloïdogène **et** tuer le clone plasmocytaire



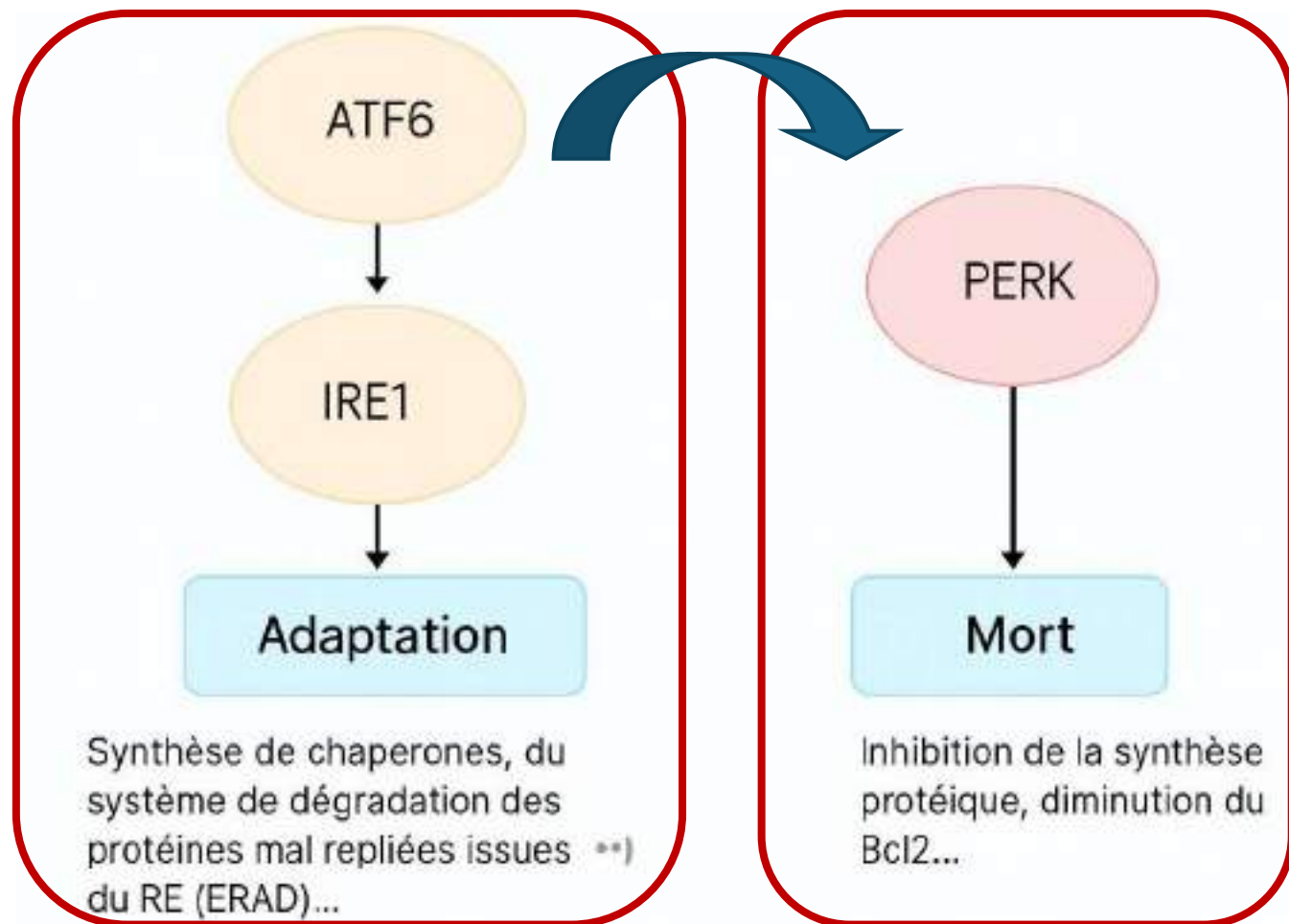
Comment supporter la production massive de protéines = système UPR (unfolded protein response)



Inhibition de la synthèse protéique, diminution du Bcl2...

**Talon d'Achille
des
plasmocytes =
Protéostase
intracellulaire**

Les cellules productrices privilégient les voies d'adaptation (ATF6, IRE1, XBP1)



Perturbation du système de protéostase

↓
Inhibiteurs du protéasome, autres...

↓
MORT

From www.bloodjournal.org by guest on February 4, 2015. For personal use only.

NEOPLASIA

Proteasome inhibitors induce a terminal unfolded protein response in multiple myeloma cells

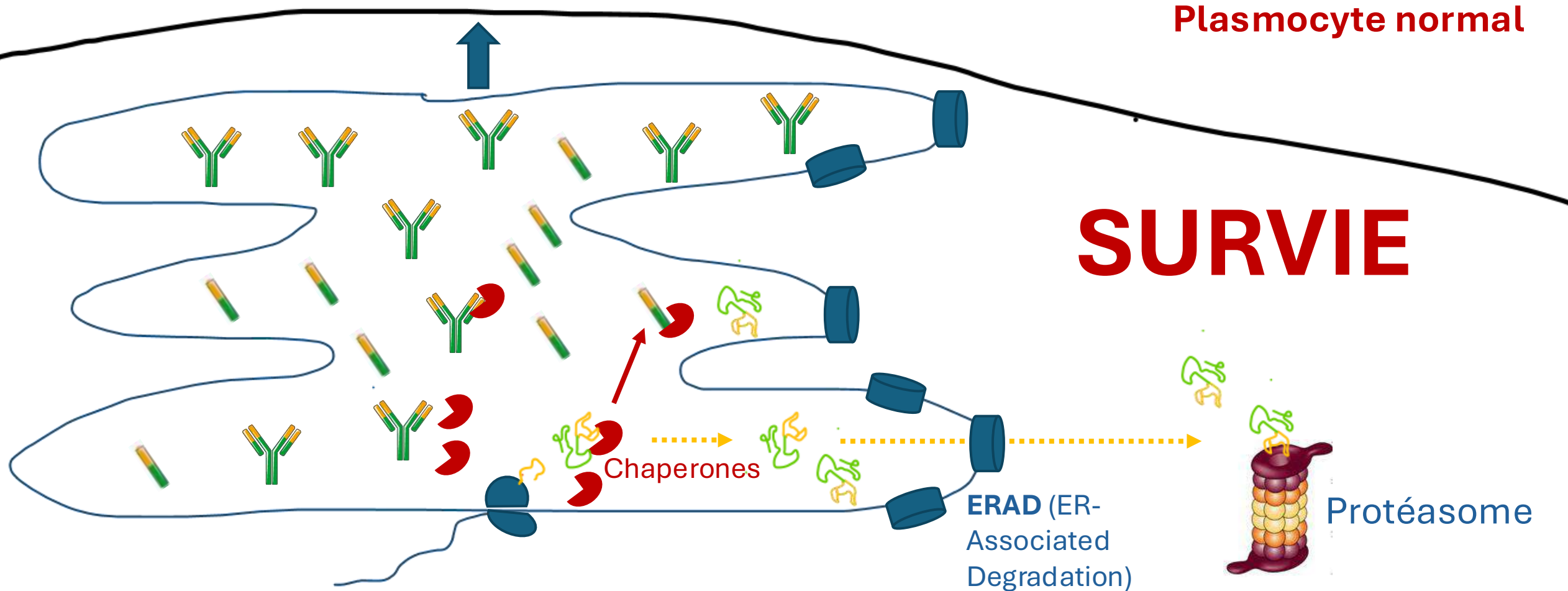
Esther A. Obeng, Louise M. Carlson, Delia M. Gutman, William J. Harrington Jr, Kelvin P. Lee, and Lawrence H. Boise



Sécrétion

Plasmocyte normal

SURVIE

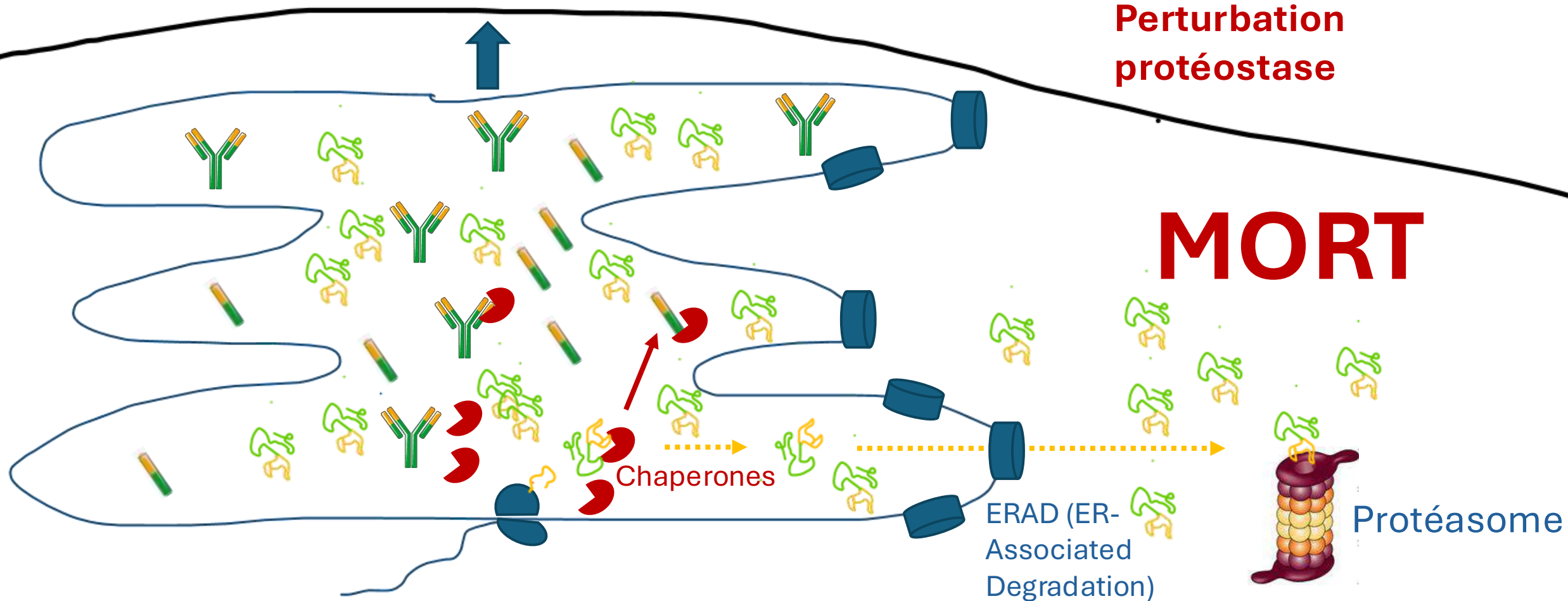




Sécrétion

Perturbation
protéostase

MORT





Blocage LC par siRNA: Accumulation chaînes lourdes dans ER → **terminal ER s**

LYMPHOID NEOPLASIA

One siRNA pool targeting the λ constant region stops λ light-chain production and causes terminal endoplasmic reticulum stress

Ping Zhou,¹ Xun Ma,¹ Lakshmanan Iyer,² Chakra Chaulagain,¹ and Raymond L. Comenzo^{1,2}

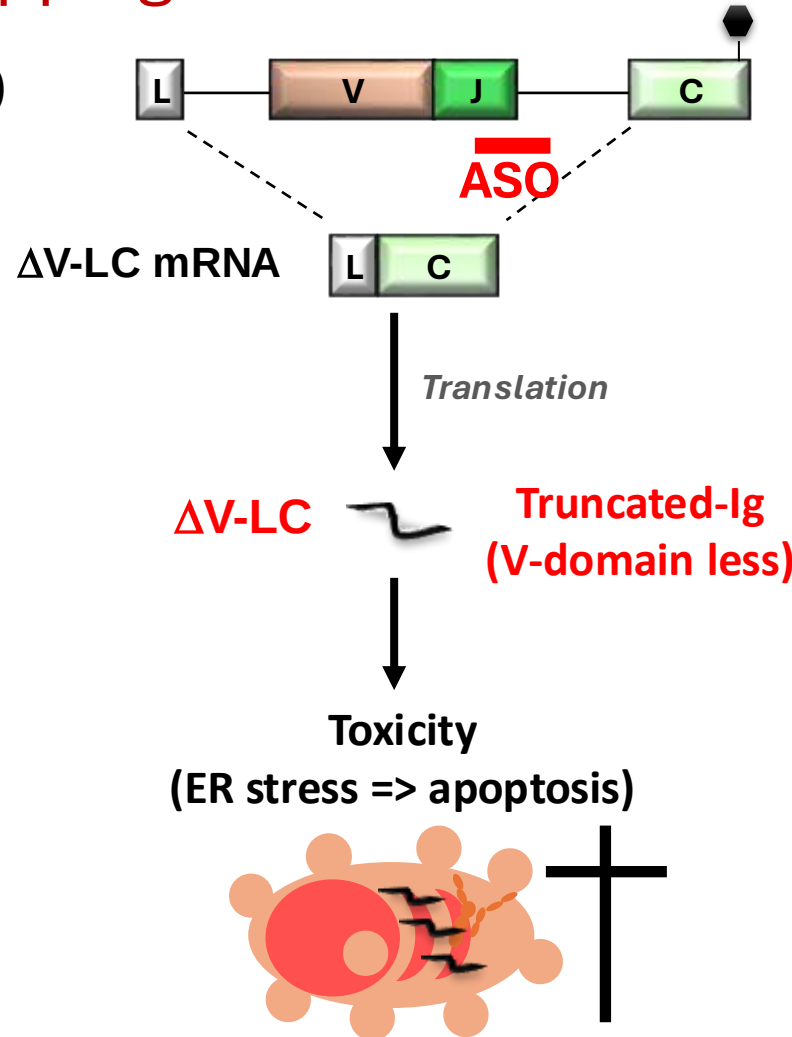
¹Tufts Medical Center, Boston, MA; and ²Tufts University School of Medicine, Boston, MA

- Peut être clone-spécifique (médecine personnalisée)



LC exon skipping: Accumulation CH dans ER → **terminal ER stress**

LC gene (clone specific)



Physiological and druggable skipping of immunoglobulin variable exons in plasma cells

[Mohamad Omar Ashi](#), [Nivine Srouf](#), [Jean-Marie Lambert](#), [Anne Marchalot](#), [Ophélie Martin](#), [Sandrine Le Noir](#), [Eric Pinaud](#), [Maria Victoria Ayala](#), [Christophe Sirac](#), [Jérôme Saulière](#), [Jérôme Moreaux](#), [Michel Cogné](#) & [Laurent Delpy](#)

Cellular & Molecular Immunology 16, 810–819 (2019) | [Cite this article](#)

Delpy et al. patent WO2017089359

- Peut être clone-spécifique (médecine personnalisée)



Amylose AL

But: Réduire la production de la LC amyloïdogène **et** tuer le clone plasmocytaire

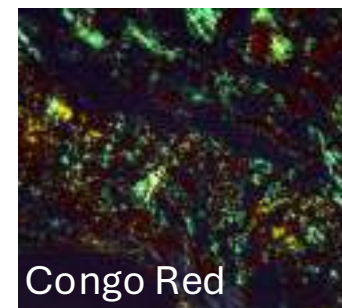
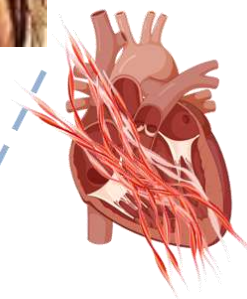
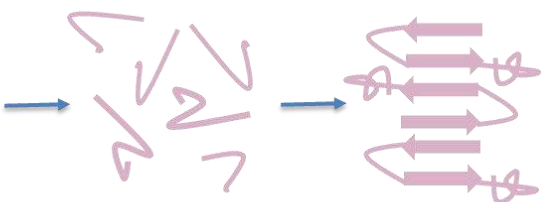
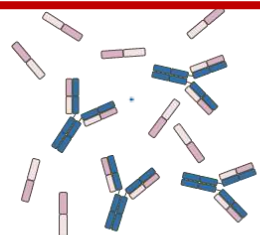
> Traitements anti-myélome font très bien le job (Dara, Bispé, Car-T...)



Producing cells
The (not) good



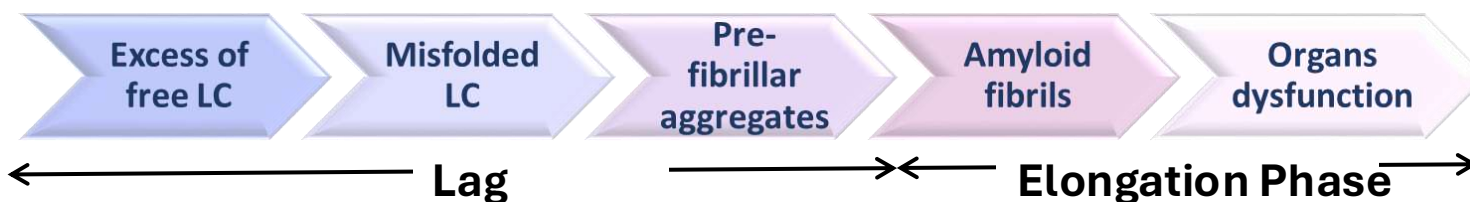
Unstable protein
The ugly



Deposits

The bad

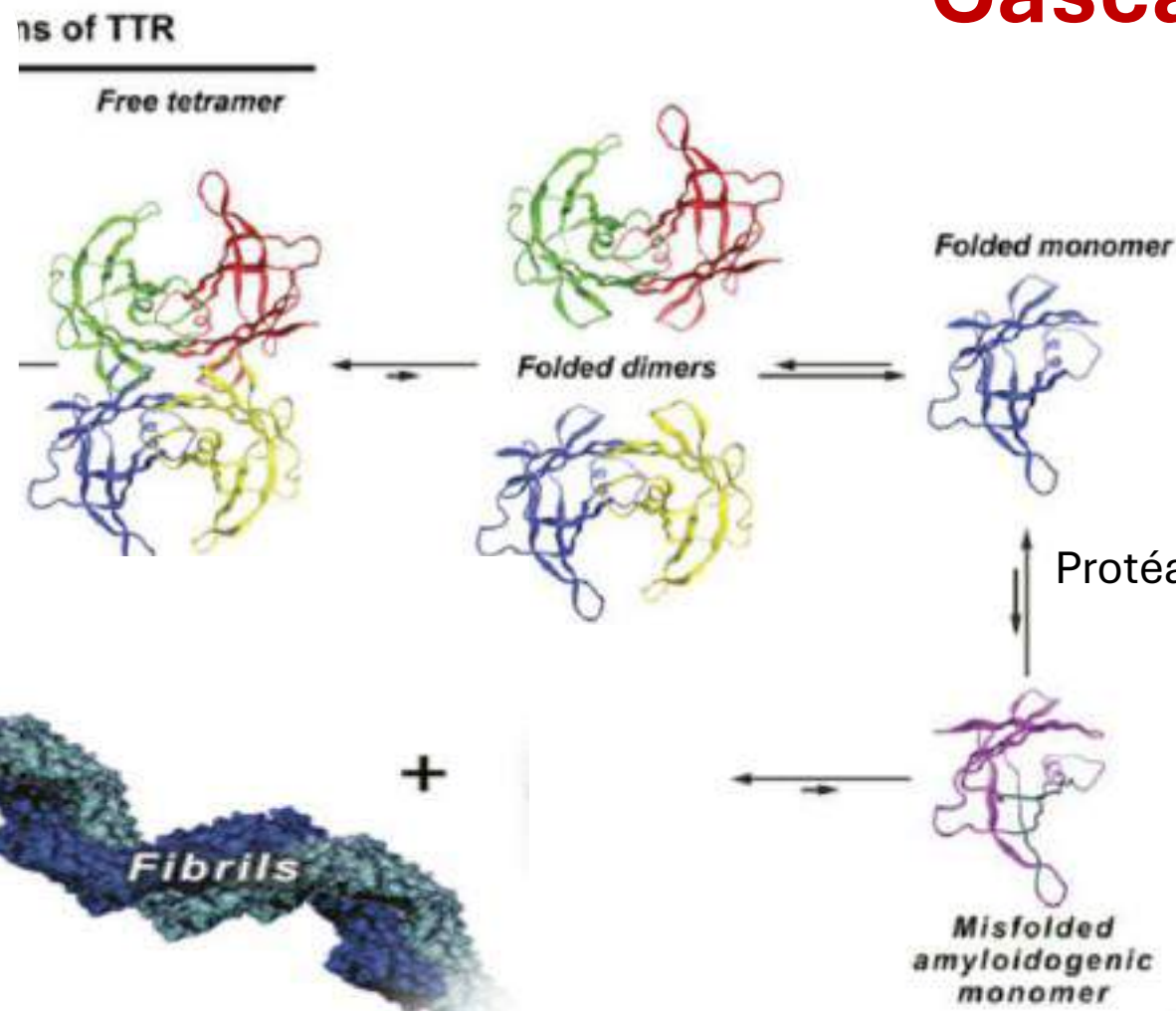
**Organ
dysfunction**





Cascade amyloïde de la TTR

Accélérée par certaines mutations
Travaux de Jeff Kelly (depuis 1992)



ARTICLE

Plasmin activity promotes amyloid deposition in a transgenic model of human transthyretin amyloidosis

Ivana Stanova¹, Rozita Adib^{1,2}, Stephan Elmerich¹, Michal R. Golos^{1,4}, Janet A. Gilbertson¹, Nicola Botcher¹, Diana Cavetti¹, Graham W. Taylor¹, Nigel Rendell¹, Glorisa A. Tennire¹, Guglielmo Verone¹, Riccardo Patarini¹, P. Patrizia Mangione^{1,2}, Julian D. Gilmore¹, Mark B. Pepys¹, Vittorio Bellotti^{1,2}, Philip M. Hawkins¹, Raya Al-Shaw¹ & J. Paul Sienkiewicz^{1,2}

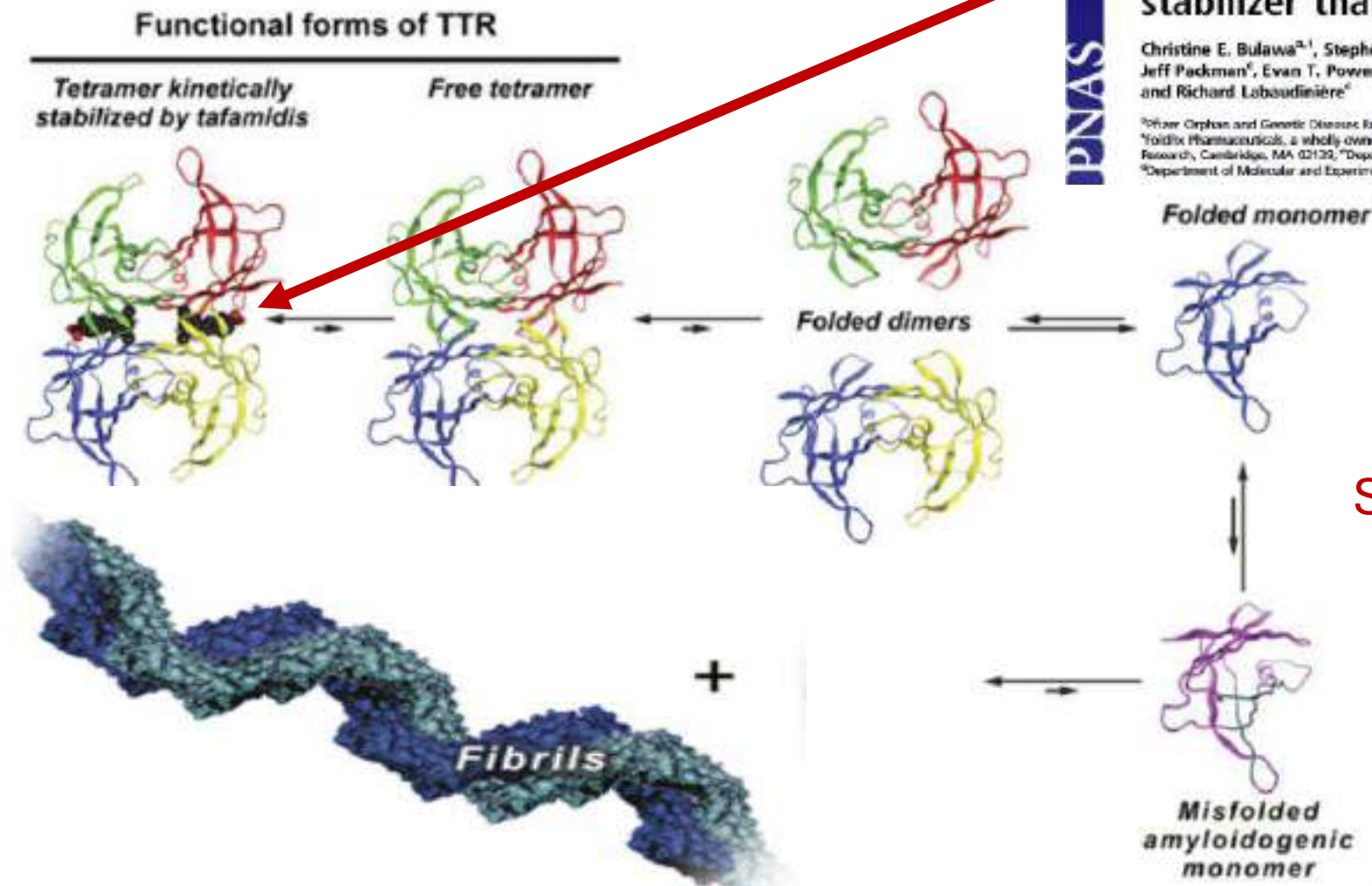


Tafamidis, a potent and selective transthyretin kinetic stabilizer that inhibits the amyloid cascade

Christine E. Bulawa^{a,1}, Stephen Connelly^b, Michael DeVit^c, Lan Wang^d, Charlotte Weigel^e, James A. Fleming^f, Jeff Peckman^g, Evan T. Powers^{h,i}, R. Luke Wiseman^g, Theodore R. Foss^h, Ian A. Wilson^{h,i}, Jeffery W. Kelly^{a,i,g}, and Richard Labaudinière^a

^aPfizer Orphan and Genetic Disease Research Unit, Cambridge, MA 02140; ^bDepartment of Molecular Biology, The Scripps Research Institute, La Jolla, CA 92037; ^cFoldix Pharmaceuticals, a wholly owned subsidiary of Pfizer, Cambridge, MA 02140; ^dCardiovascular and Metabolic Diseases, Novartis Institute for Biomedical Research, Cambridge, MA 02132; ^eDepartment of Chemistry, The Scripps Research Institute and ^fThe Skaggs Institute for Chemical Biology, La Jolla, CA 92037; ^gDepartment of Molecular and Experimental Medicine, The Scripps Research Institute, La Jolla, CA 92037; and ^hLife Technologies, Genentech Systems, Beverly, MA 01915

PNAS

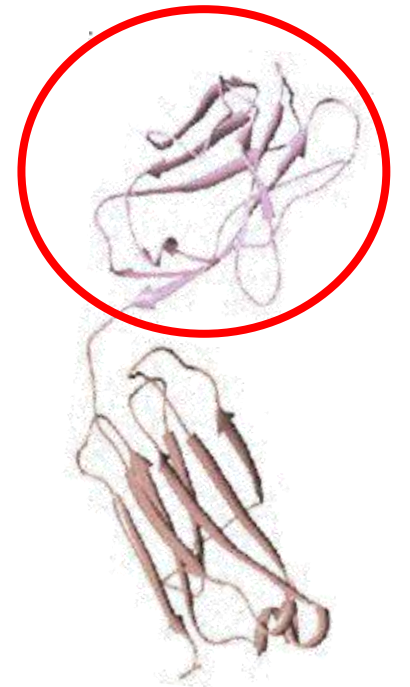


Stabilise tous les mutants et WT



Amylose AL: ça se complique...

- Role of the V domain: Not all LC are amyloidogenic
- 100 papers about specific mutations reducing stability of LC (*Ramirez-Alvarado, Wall, Kelly, Eisenberg, Merlini, Lavatelli, Buel, Buchner, Fandrich, Pozo-Yauner, Stevens, etc...*)
- ALbase with ~900 AL LC – Limoges ~350 AL LC
 - No common mutations
 - Multiple mutations in each VL





Emergence d'outils de prédiction d'amyloïdogénicité par machine-learning



Garofalo et al. 2021

83%



Rawat et al. 2021

79%

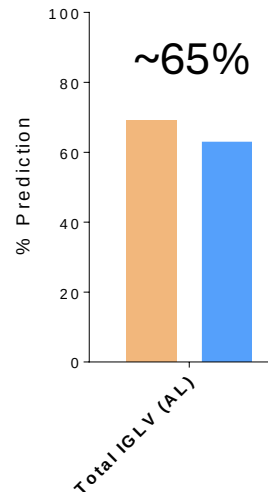


Zhou et al. 2023

93%



Limoges database, 350
IGLV AL, 580 IGLV
other monoclonal
diseases



Nombreux biais
(contrôles, données
cliniques)

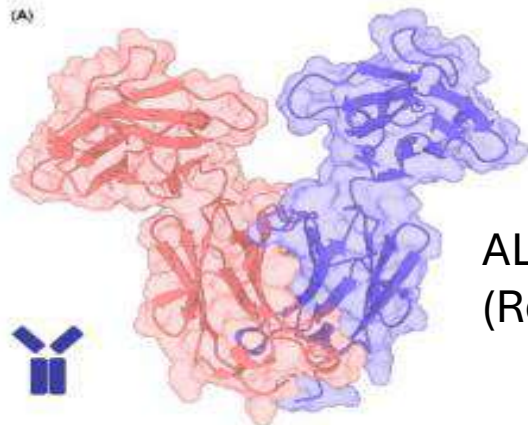
BUT:

- Prédire évolution dans MGUS ou MM
- Evaluer la dangerosité de la LC



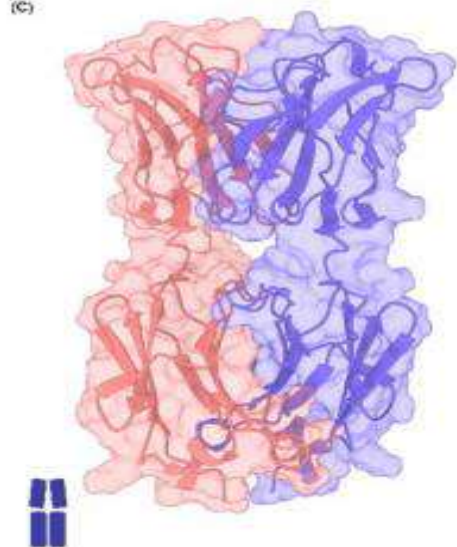
Dimère VL ouvert

(A)



AL55 structure
(Real crystal)

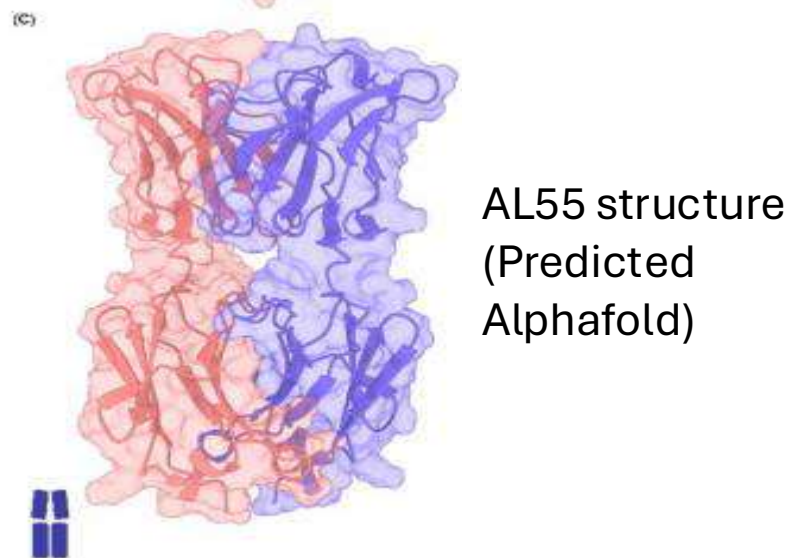
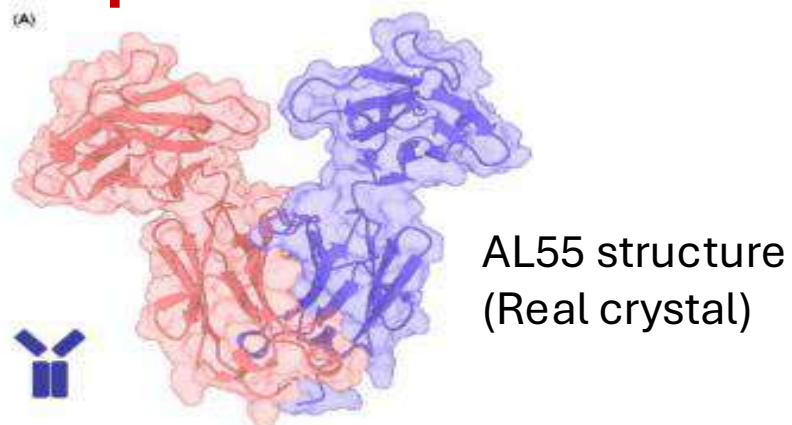
(C)



AL55 structure
(Predicted
AlphaFold)

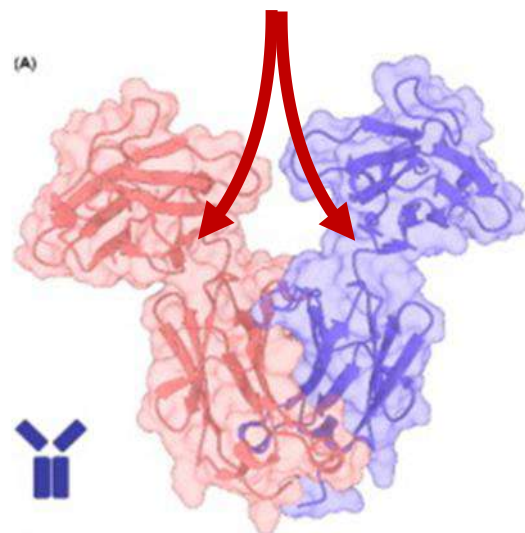


Open VL dimer



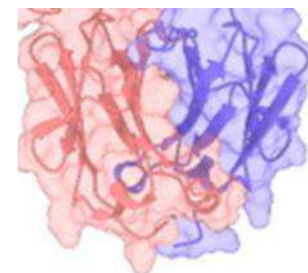
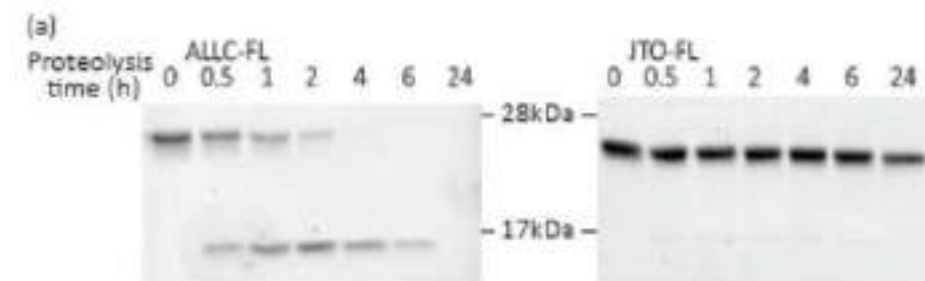
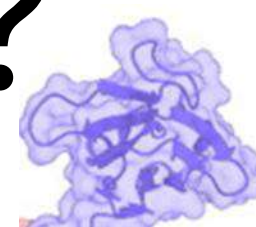
Puri et al. JMB 2025

Protéases



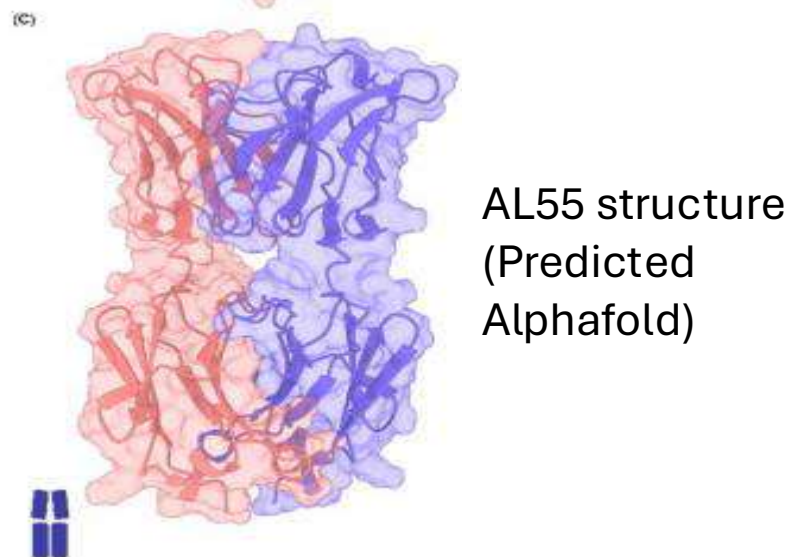
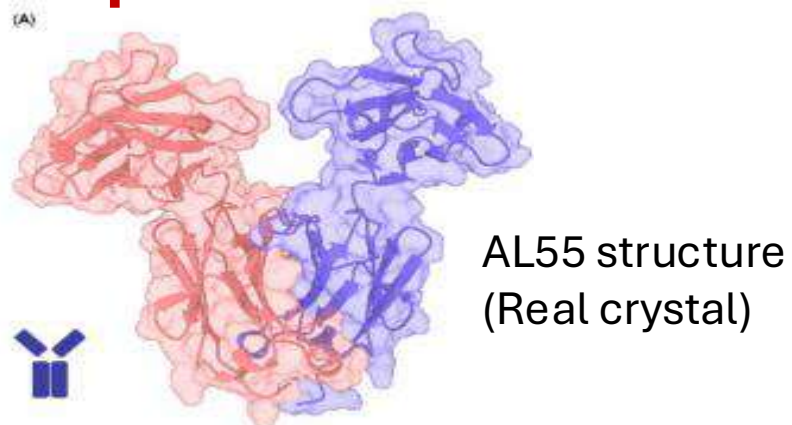
Morgan et al, JMB 2016

?



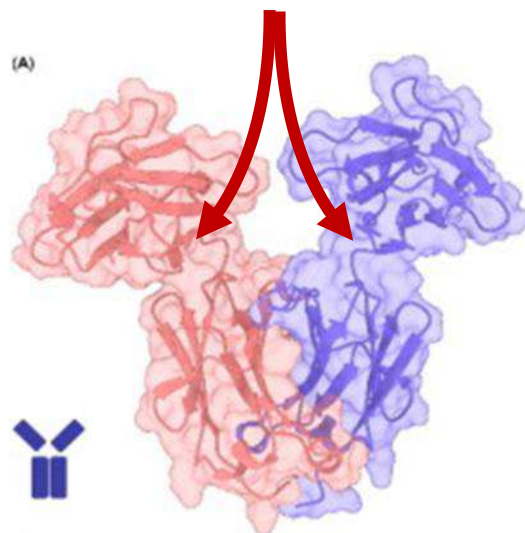


Open VL dimer

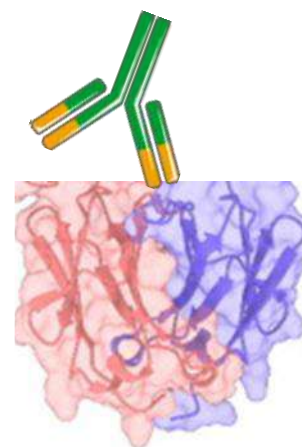
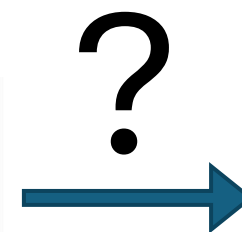
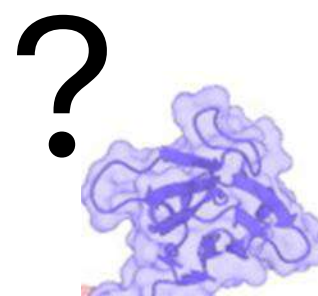


Puri et al. JMB 2025

Protéases



Morgan et al, JMB 2016



Diagnostic
Prédiction





536 Blood-Based Diagnostic Assay for λ Light Chain Amyloidosis through Quantification of an Amyloidogenicity-Indicating Neo-Epitope

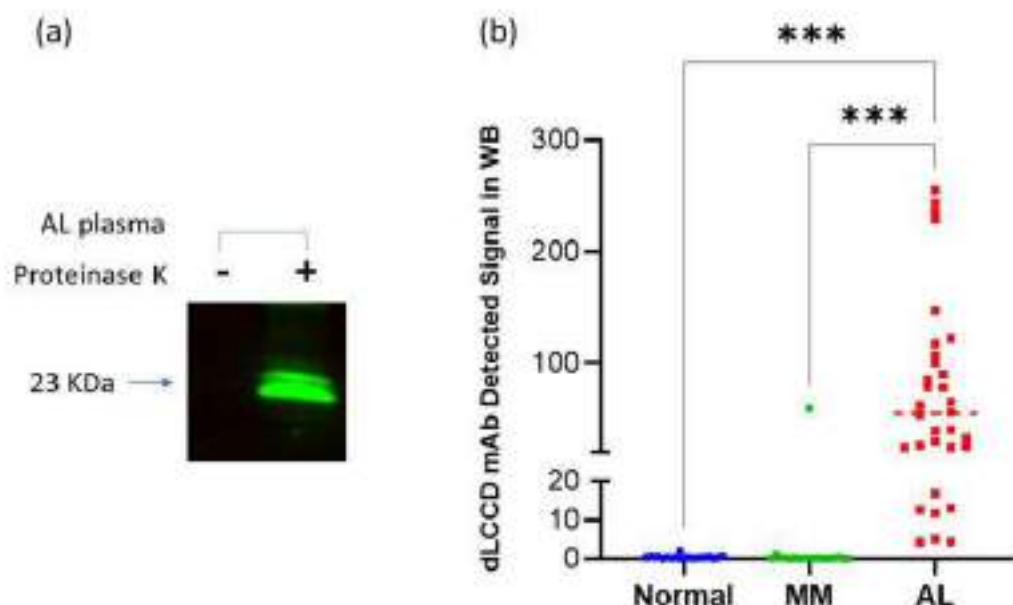


Fig. 1. dLCCD mAbs quantify dLCCD fragment specifically in monoclonal λ AL patient plasma samples after limited proteolysis with Proteinase K. (a) Example of an AL plasma sample before and after limited proteolysis, ran on a Western blot, and detected with a dLCCD mAb targeting the neo-epitope on dLCCD. A fluorophore-tagged secondary antibody was used for the detection and quantification. (b) dLCCD signal of protease-treated λ FLC in plasma samples from healthy normal donors (N=23), MM (N=19) and AL (N=32) patients. One-way ANOVA was used to compare normal or MM to AL samples, resulting in $p=0.0007$ and 0.0008 , respectively.

**Test sérique pour prédire et suivre
l'amyloïdogénicité d'une LC monoclonale
(uniquement lambda LC)**

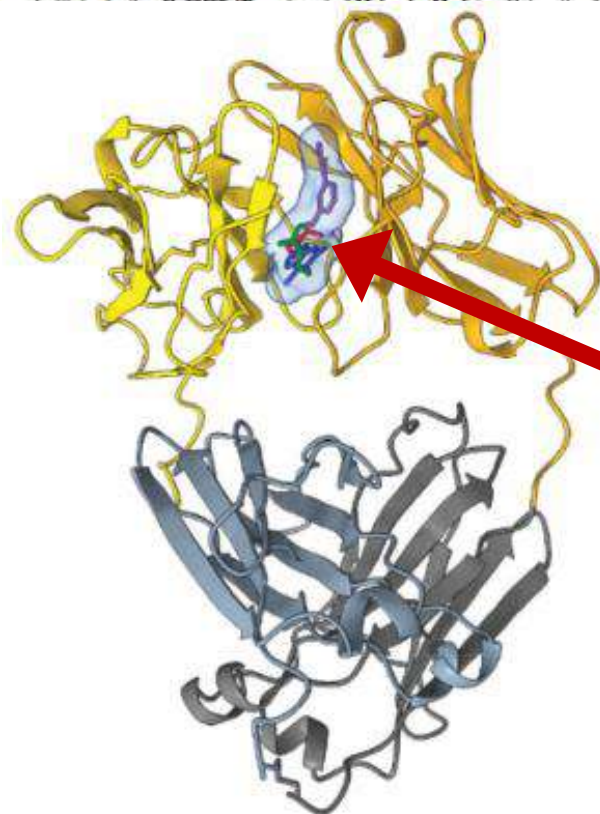


Stabilization of amyloidogenic immunoglobulin light chains by small molecules

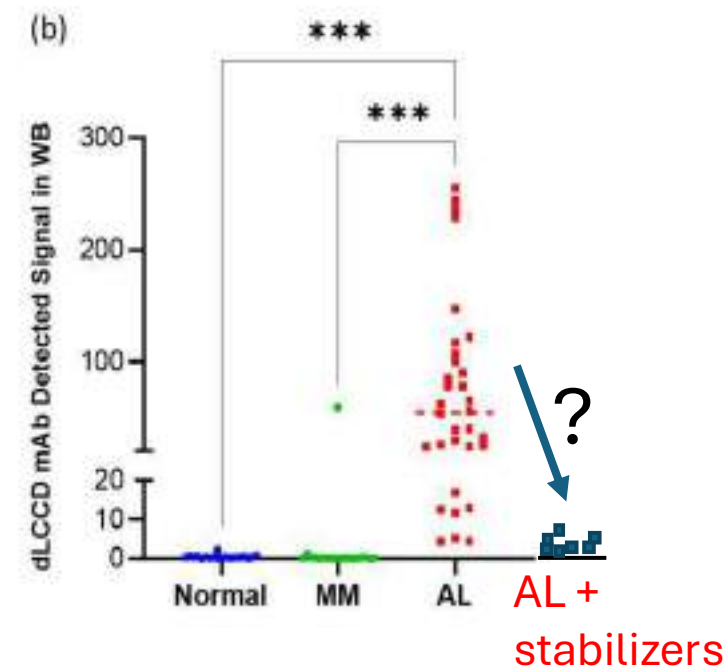
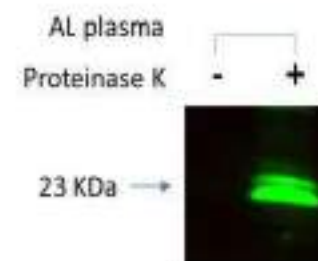
Gareth J. Morgan^{a,h,1,2,3}, Nicholas L. Yan^{a,h,1}, David E. Mortenson^{a,b}, Enrico Rennella^{c,d,e}, Joshua M. Blundon^{a,b}, Ryan M. Gwin^{a,b}, Chung-Yon Lin^{a,b}, Robyn L. Stanfield^f, Steven J. Brown^g, Hugh Rosen^g, Timothy P. Spicer^g, Virmeliz Fernandez-Vega^g, Giampaolo Merlini^{h,i}, Lewis E. Kay^{c,d,e,j}, Ian A. Wilson^k, and Jeffery W. Kelly^{a,h,k,2}

^aDepartment of Molecular Medicine, The Scripps Research Institute, La Jolla, CA 92037; ^bDepartment of Chemistry, The Scripps Research Institute, La Jolla, CA 92037; ^cDepartment of Molecular Genetics, The University of Toronto, Toronto, ON M5S1A8, Canada; ^dDepartment of Biochemistry, The University of Toronto, Toronto, ON M5S1A8, Canada; ^eDepartment of Chemistry, The University of Toronto, Toronto, ON M5S1A8, Canada; ^fDepartment of Integrative Structural and Computational Biology, The Scripps Research Institute, La Jolla, CA 92037; ^gDepartment of Molecular Medicine, The Scripps Research

Stabilizing amyloid LC (?)



**Stabilisateur
du dimère**





Stabilizing amyloid LC (?)

Utile?

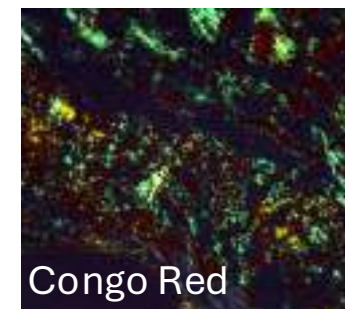
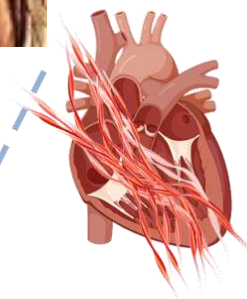
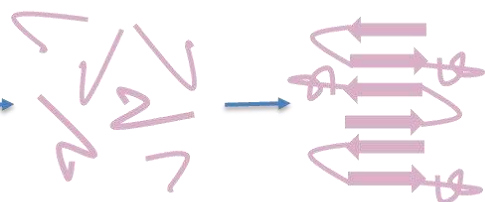
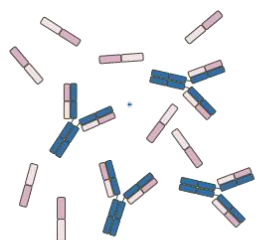
(à l'heure des bi-spécifiques, CAR-T...)



Producing cells
The (not) good



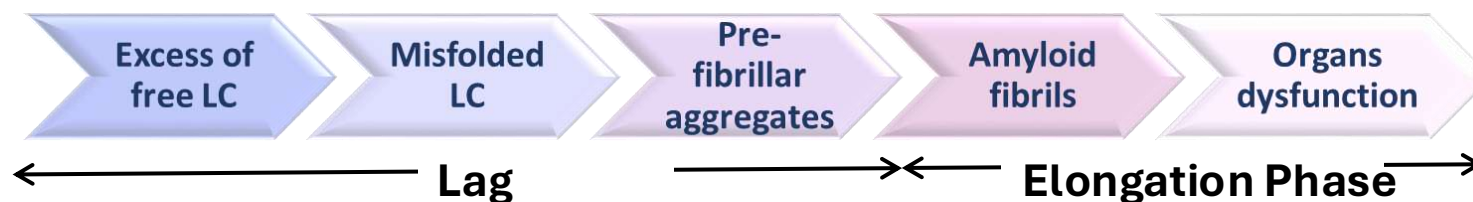
Unstable protein
The ugly



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**Organ
dysfunction**





Avant la Cryomicroscopie electronique (CryoEM)



Forme native
circulante

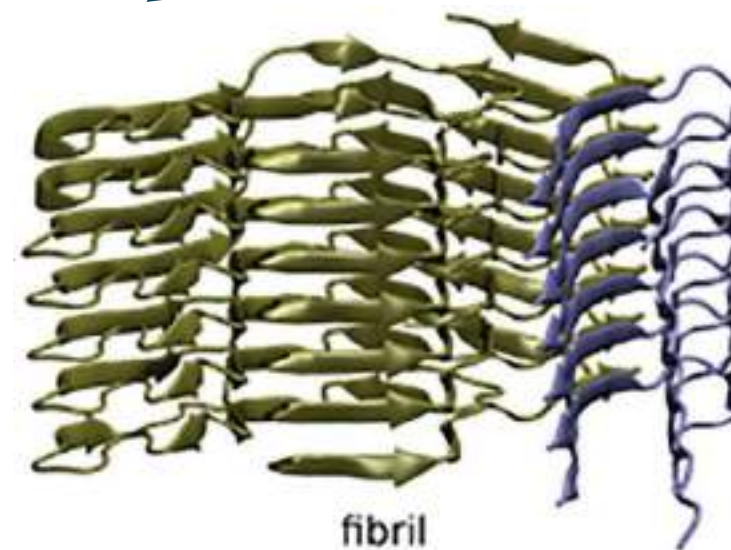
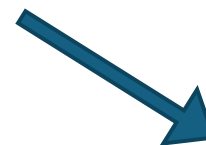
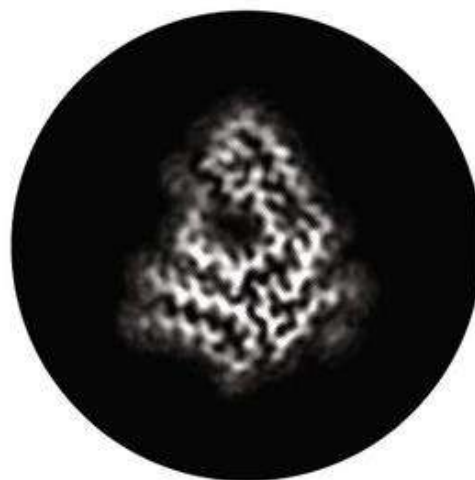
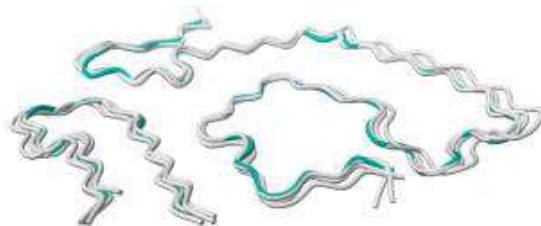




Avec la Cryomicroscopie electronique (CryoEM)



Forme native
circulante



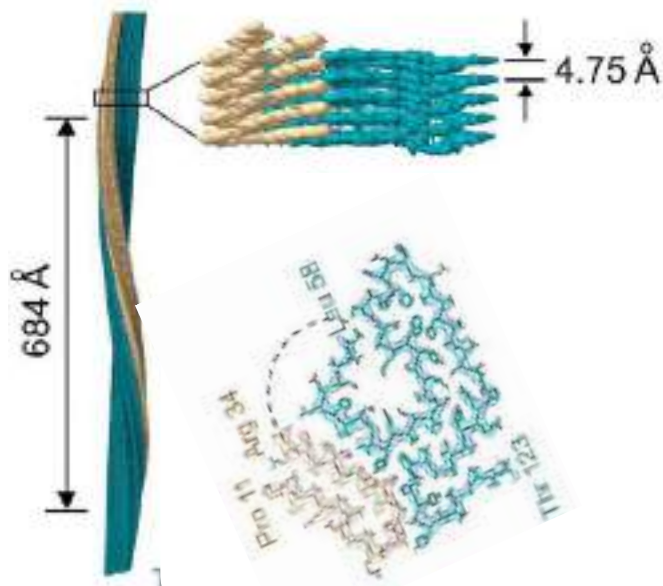
fibril



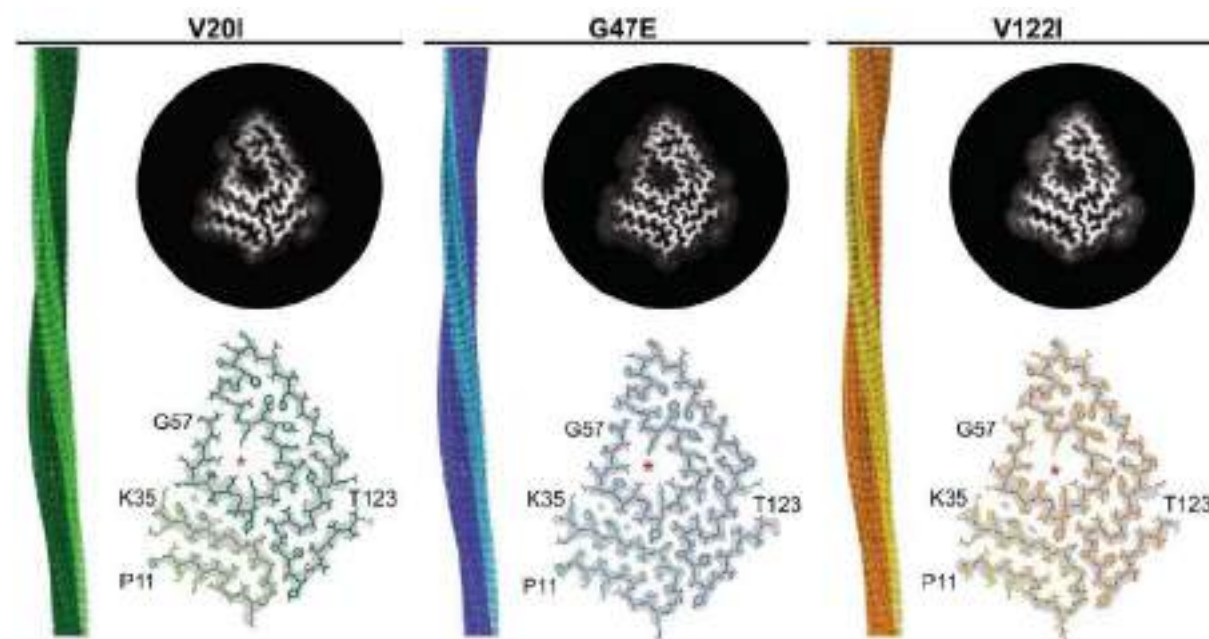
ATTR (mais aussi la plupart des amyloses systémiques)

- Homogénéité inter-organes
- Homogénéité inter-patients
- Homogénéité inter-mutants

a) ATTRwt-p1



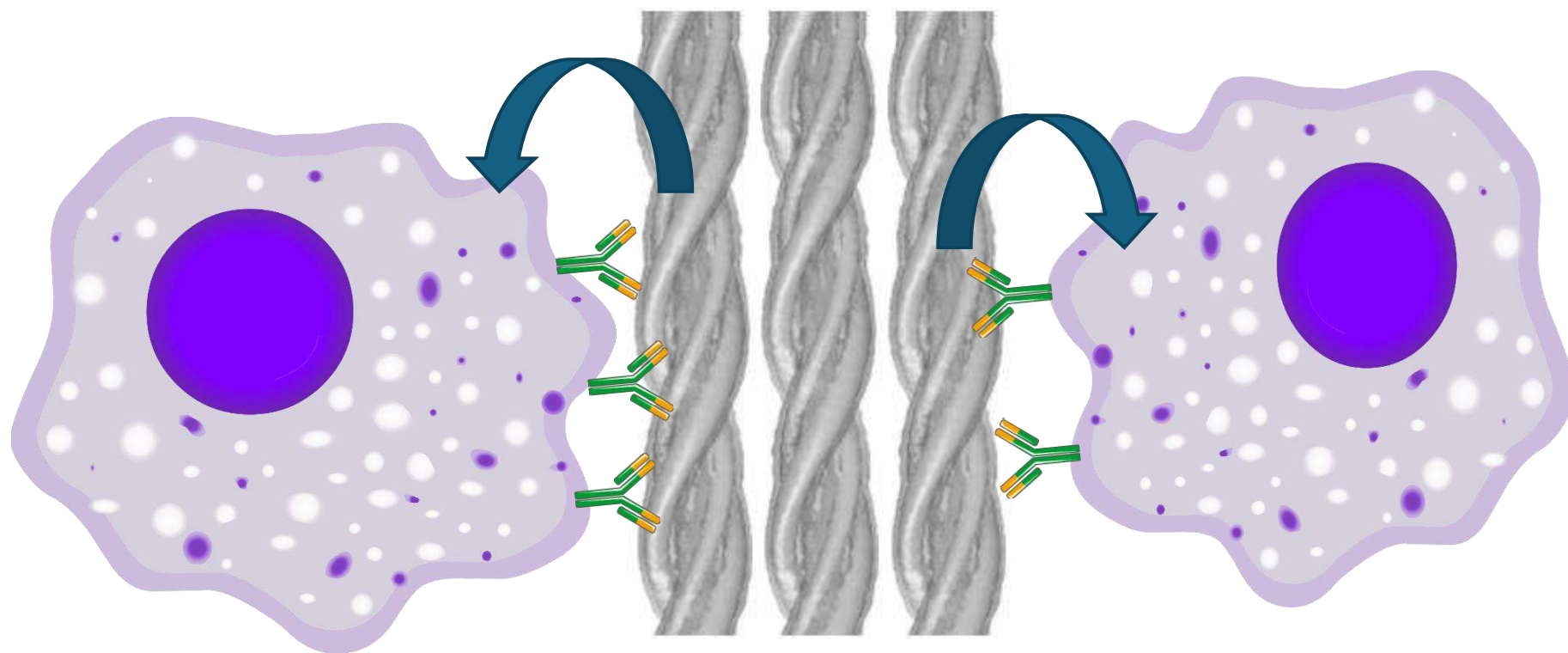
ATTR-mut





Rationnel pour les nouvelles générations d'anticorps anti-amyloïde

« Nettoyer » les organes



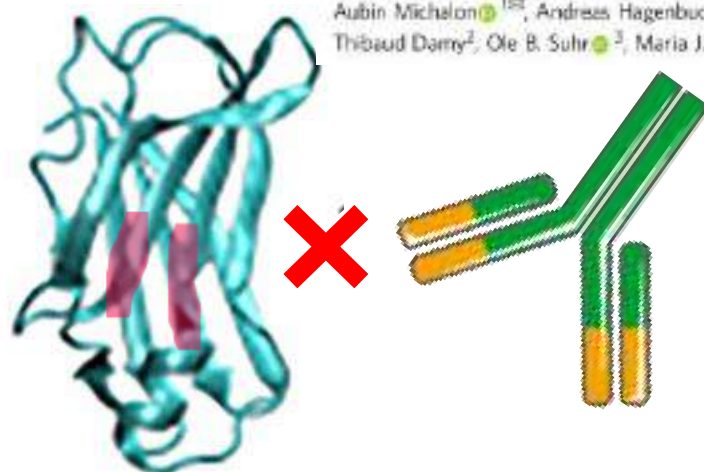
En complément des autres thérapies



Rationnel pour les nouvelles générations d'anticorps anti-amyloïde

A human antibody selective for transthyretin amyloid removes cardiac amyloid through phagocytic immune cells

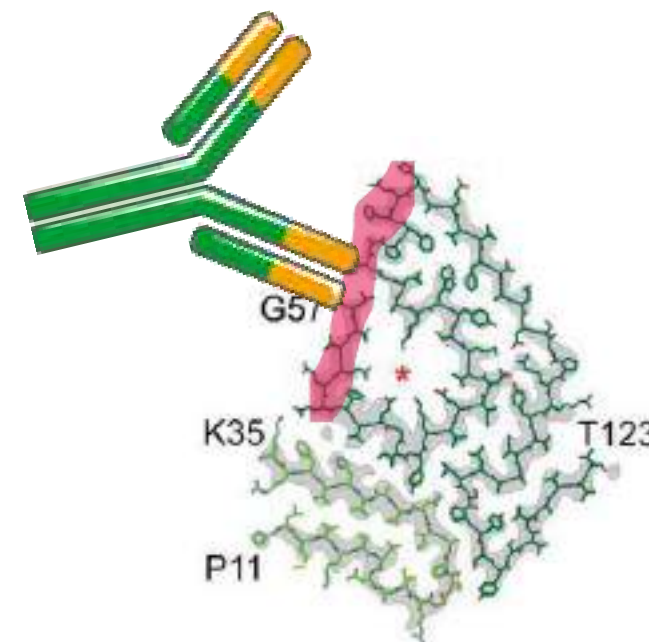
Aubin Michalon¹, Andreas Hagenbuch¹, Christian Huy², Evita Varela¹, Benoit Combaluzier¹, Thibaud Darny², Ole B. Suhr³, Maria J. Saraiva⁴, Christoph Hock^{1,2}, Roger M. Nitsch^{1,2} & Jan Grimm¹



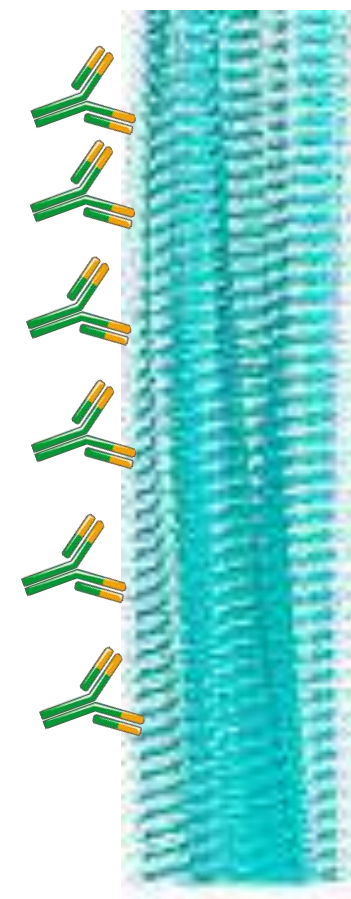
Forme native
circulante

Phase 1 Trial of Antibody NI006 for
Depletion of Cardiac Transthyretin Amyloid

Pablo Garcia-Pavia, M.D., Ph.D., Fabian aus dem Siepen, M.D.,
Erwan Donal, M.D., Ph.D., Olivier Lairez, M.D., Peter van der Meer, M.D., Ph.D.,
Arnt V. Kristen, M.D., Michele F. Mercuri, M.D., Ph.D., Aubin Michalon, Ph.D.,
Robert J.A. Frost, M.D., Ph.D., Jan Grimm, Ph.D., Roger M. Nitsch, M.D.,
Christoph Hock, M.D., Peter C. Kahr, M.D., and Thibaud Darny, M.D., Ph.D.

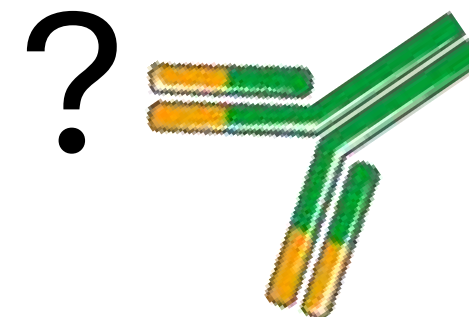
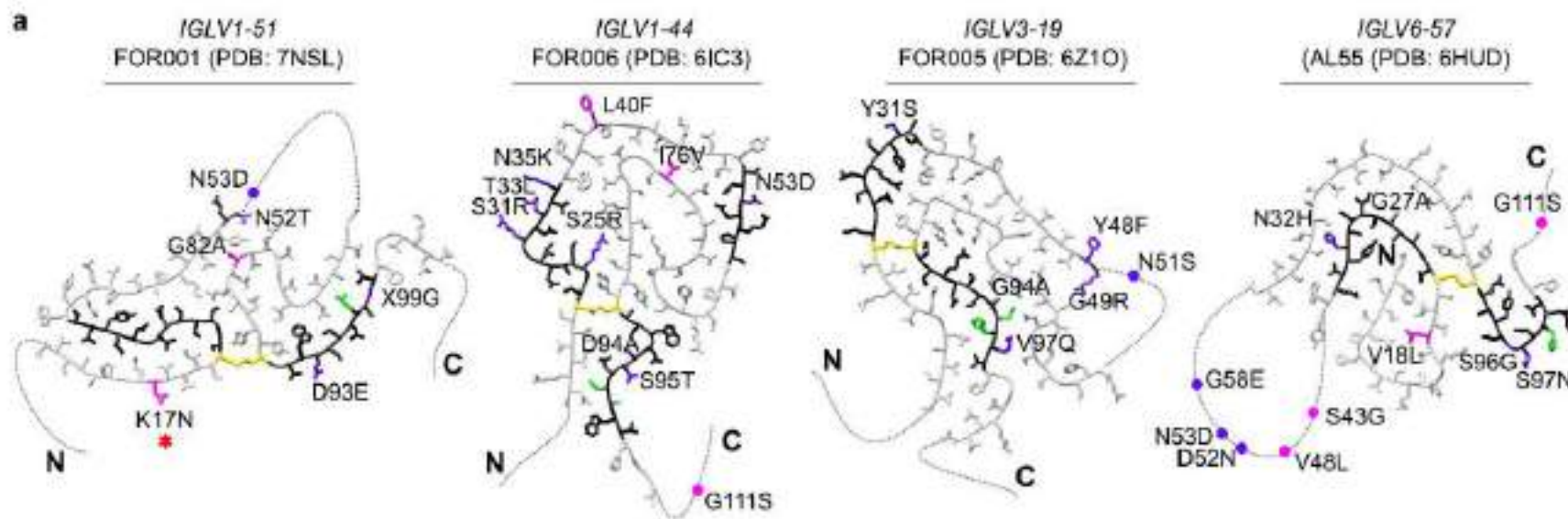


Forme
fibrillaire





Et dans l'amylose AL...

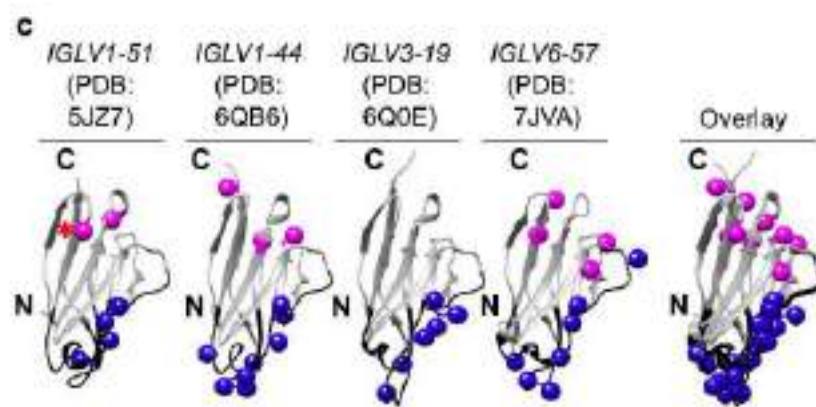


b

	CDR1	
FOR001	QSVLTQPPSVSAAPGQNVVTISCSGSSSNIGNNYVSWYQQL	40
FOR006	—VLTQPPSASGTPGQRTVITSCSGRSENIGRNLVKWKYQQF	40
FOR005	—SELTQDPVAVSVALGQTVRITCQGDG—LRYSASWYQOK	38
AL55	NFMLTQPHSVSESPGKTLTISCTGSSASIASHYVQWYQQR	40

	CDR2	
FOR001	PGTAPKLLIYEETDKRPSGIPDRFSGSKSGTSAT—LGITG	78
FOR006	PGTAPKLLIYSNDQRPSCVDPDRFSGSKSGTSAS—LAVSG	78
FOR005	PGQAPVLVIFRKSNRPSPGIPDRFSGSSSGNTAS—LTITG	76
AL55	PGGAPTTLLIYENDQRPSEVPDRFSGSIDSSSNSASLTISG	80

	CDR3	
FOR001	IQTAEADYYCQGWESSELAGVFGGSKTLTVLGQPKAAPS	118
FOR006	IQSEDEADYYCAWMDATLNAWVFGGSKTLTVLSQPKAAPS	118
FOR005	AQARDEADYYCNSRDSSANKQVFGGSKTLTVLGQPKAAPS	116
AL55	LKTEDEADYYCQSYDGN—NHWVFGGSKTLTVLSQPKAAPS	119*





Echec ou succès relatif des anticorps anti-AL



NEOD001 = 2A4 (Anti-AA fibrils)

Prothena Announces Phase 3 AFFIRM-AL Clinical Trial for Birtamimab in Patients with AL Amyloidosis Did Not Meet Primary Endpoint

2025-05-23



ISS: MGUS, Amyloidosis, and Other Non-Multiple Myeloma Plasma Cell Dyscrasias: Clinical and Epidemiological

Renal responses in a phase 1/2 study of at-02, a novel pan-amyloid depleter ig fusion protein for the treatment of patients with AL amyloidosis

Gregory Bell¹, Claire Sherman¹, Matt Meldorf¹, Mozen Hanna², Vasvi Singh³,
Graham Hillis⁴, Olga Motornova⁵, Brett Sperry⁶, Brian Drochman⁷, Ahmad Masri⁸



CAEL101 = 11-1F4 (Anti-Vk4 soluble)

Results did not achieve statistical significance for the primary endpoint in overall patient population

PUBLISHED

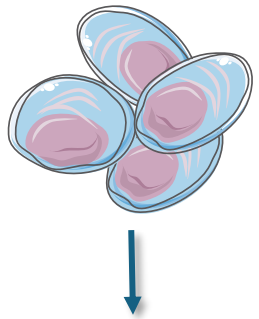
16 July 2025

CARES Phase III demonstrated anselamimab's potential to improve survival and cardiac outcomes in patients with advanced AL-kappa amyloidosis

Neurimmune expands collaboration with AstraZeneca to develop and commercialize fibril depleter NI009 for AL amyloidosis

Zurich, Switzerland - 04.12.2025

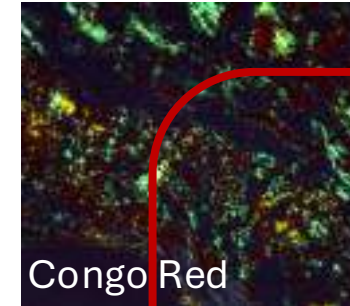
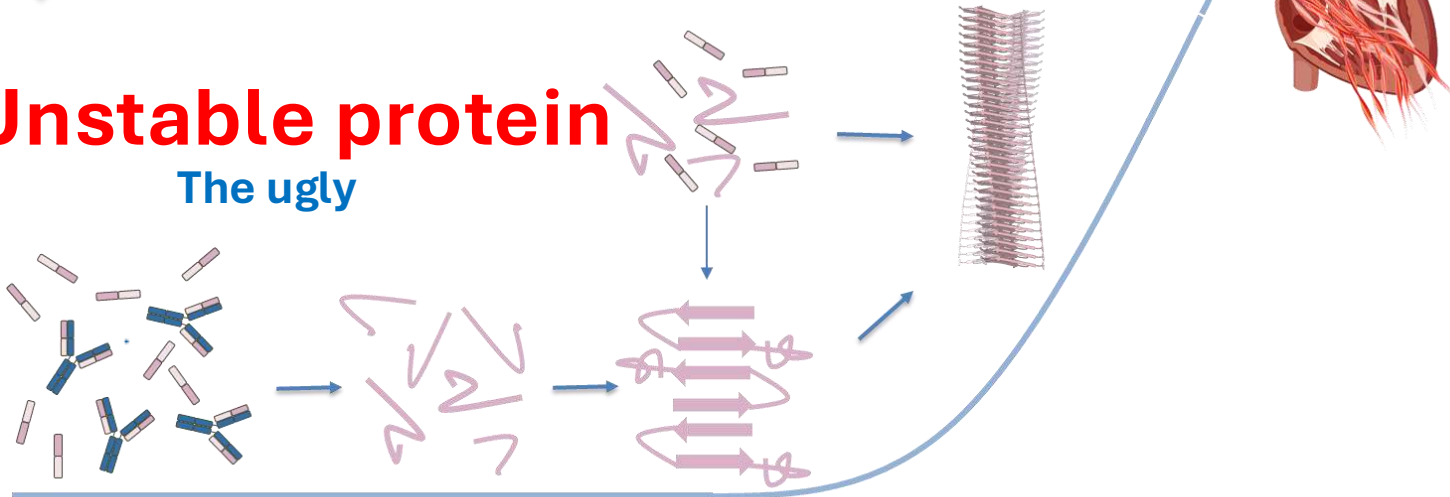
> Anti-Lambda : immunisation effectuées avec des fibrilles amyloïdes lambda de patients



Producing cells
The (not) good



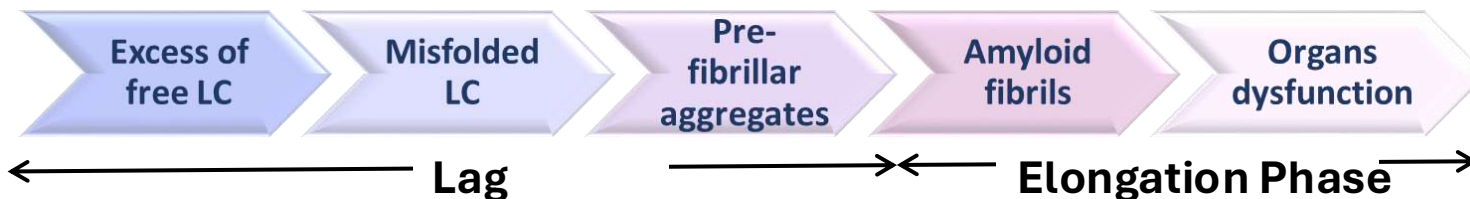
Unstable protein
The ugly



Deposits

The bad

**Organ
dysfunction**



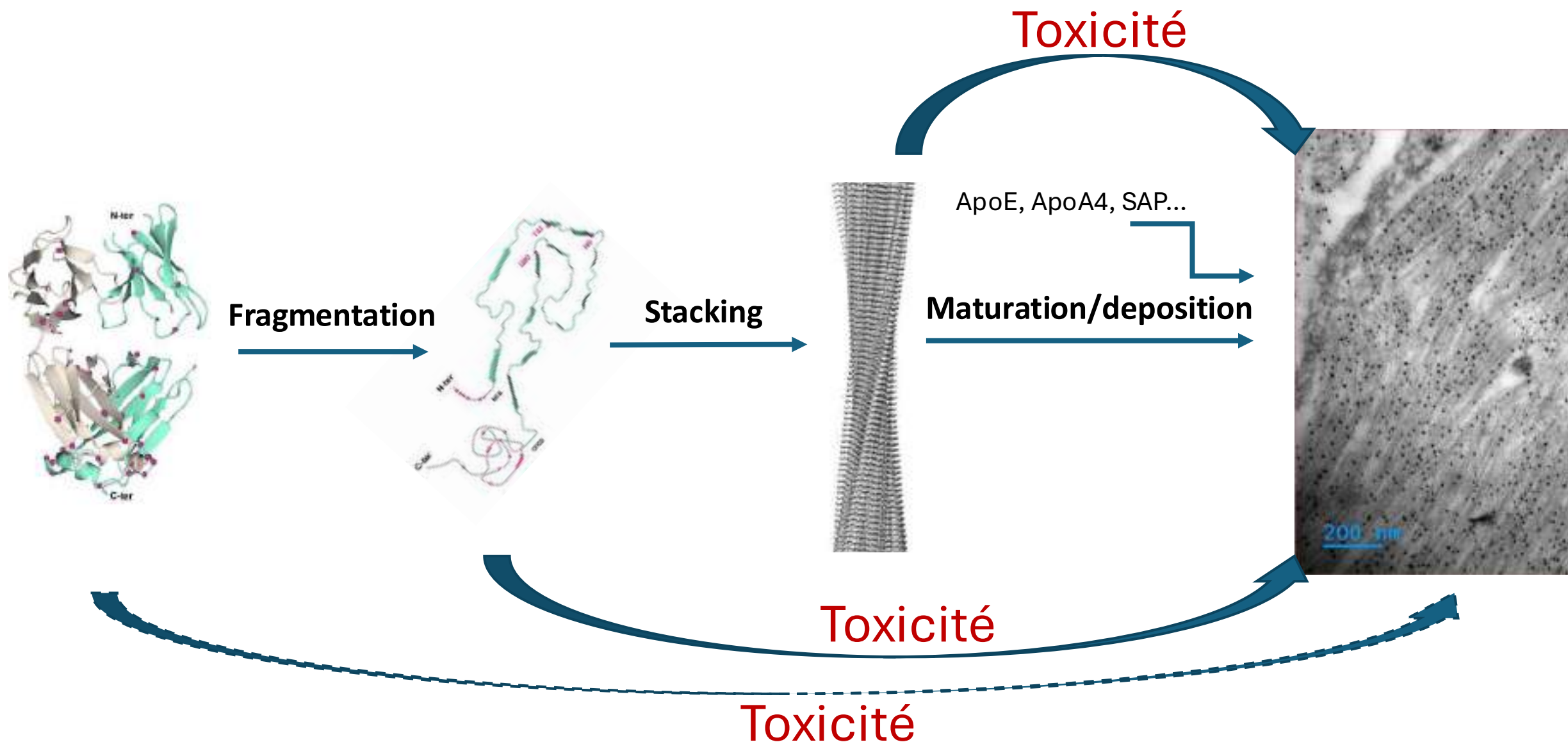


Toxicité

Tropisme

➤ Déclencheur





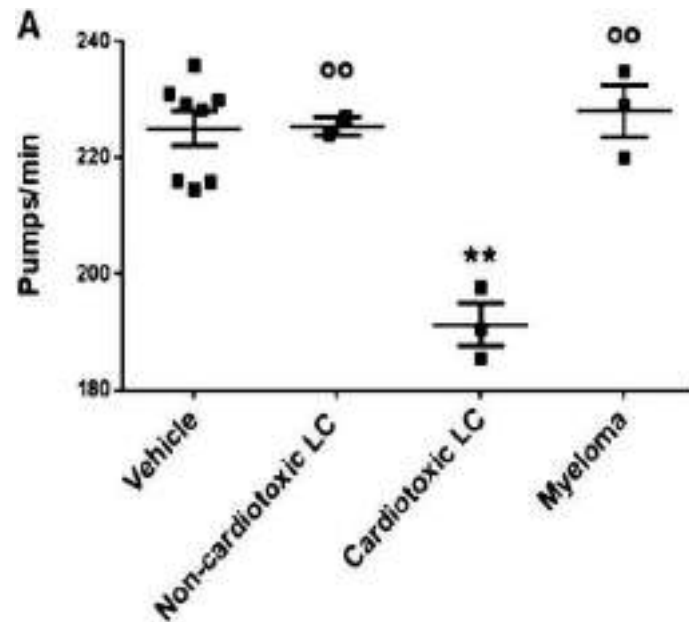


Difficulté à faire des modèles expérimentaux

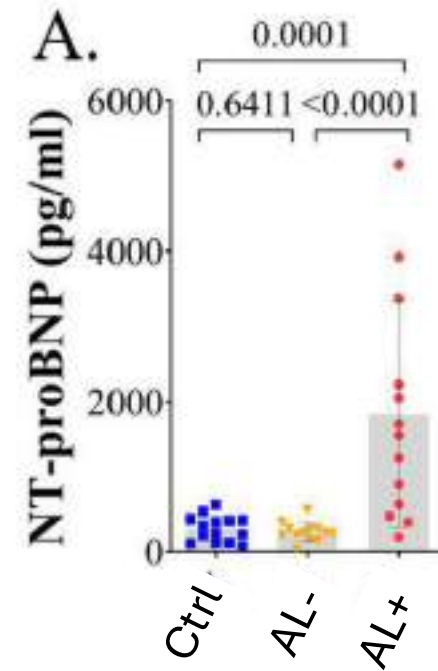
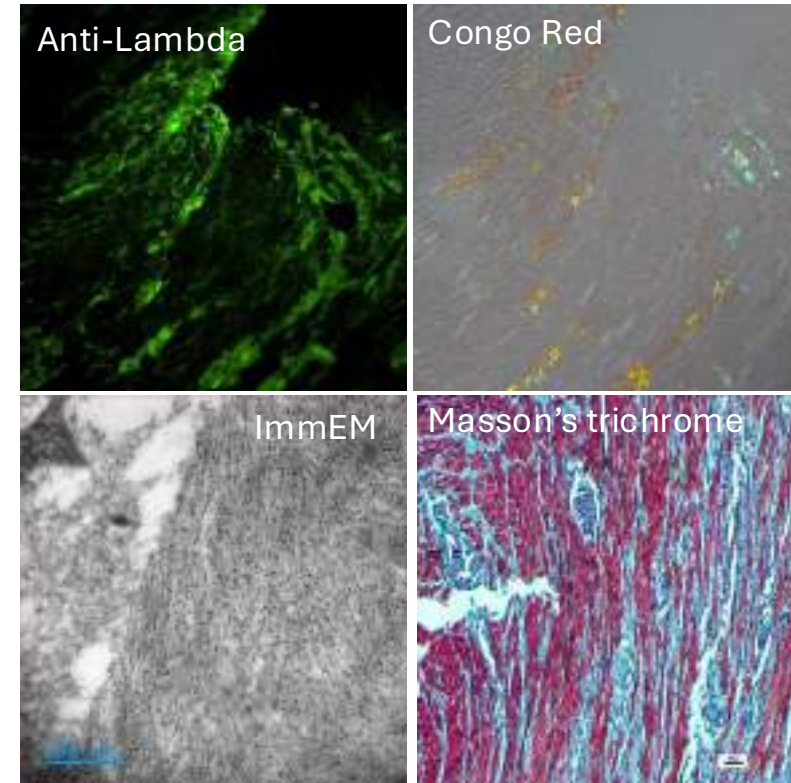
- Composition cellulaire complexe
- Matrice extracellulaire
- Flux/Force
- Système immunitaire



Modèles expérimentaux



Diomede et al, Blood, 2010



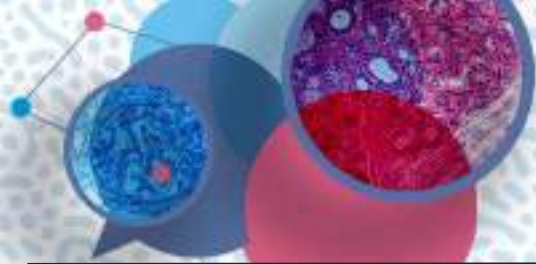
Martinez-Rivas et al, Nat Comm, 2024



Conclusion



Fais de la recherche fondamentale



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CRIBL Lab
Lab retreat
2023

F.Lavatelli and coll.
M.Ehrmann and coll.
P.Sicard
S.Ricagno and coll.

THANKS



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