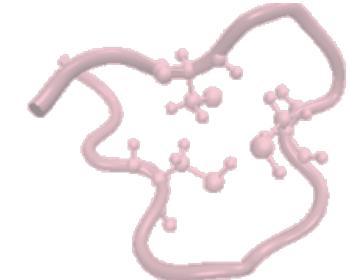


Développement de peptides antitumoraux ciblant l'interaction pro-cathepsine D/LRP-1



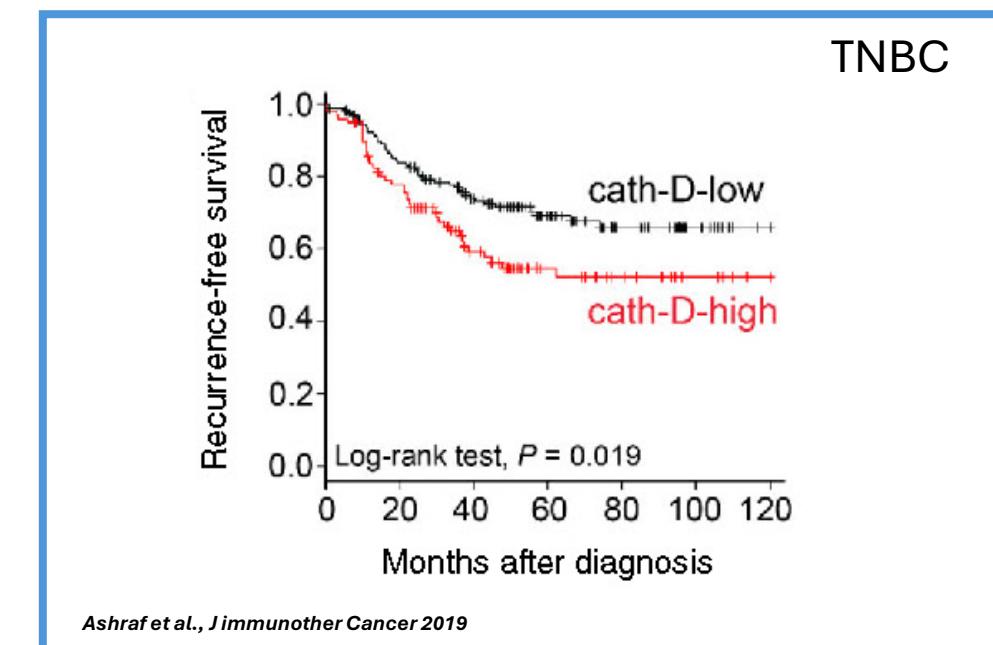
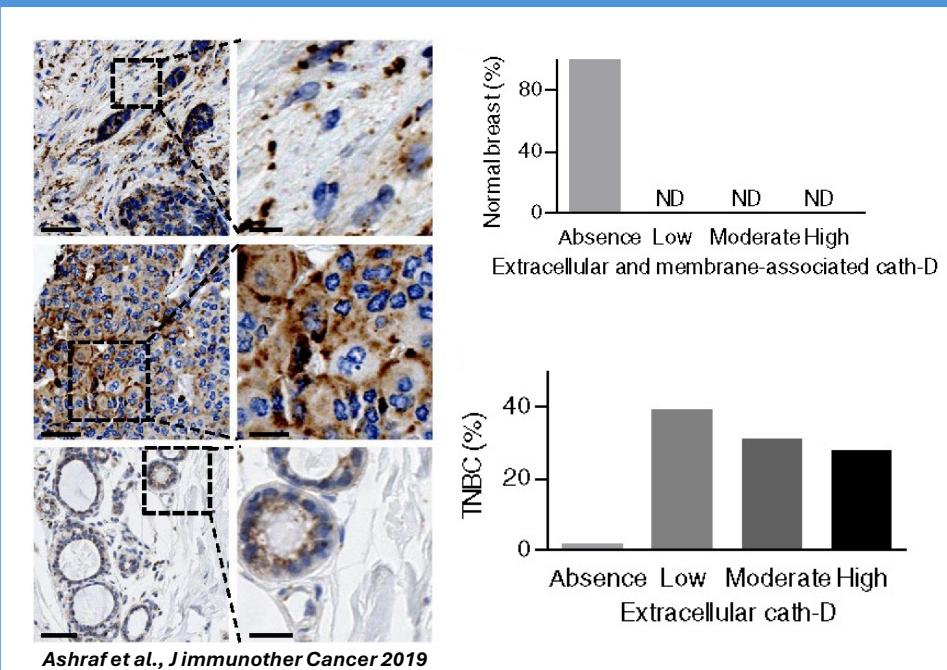
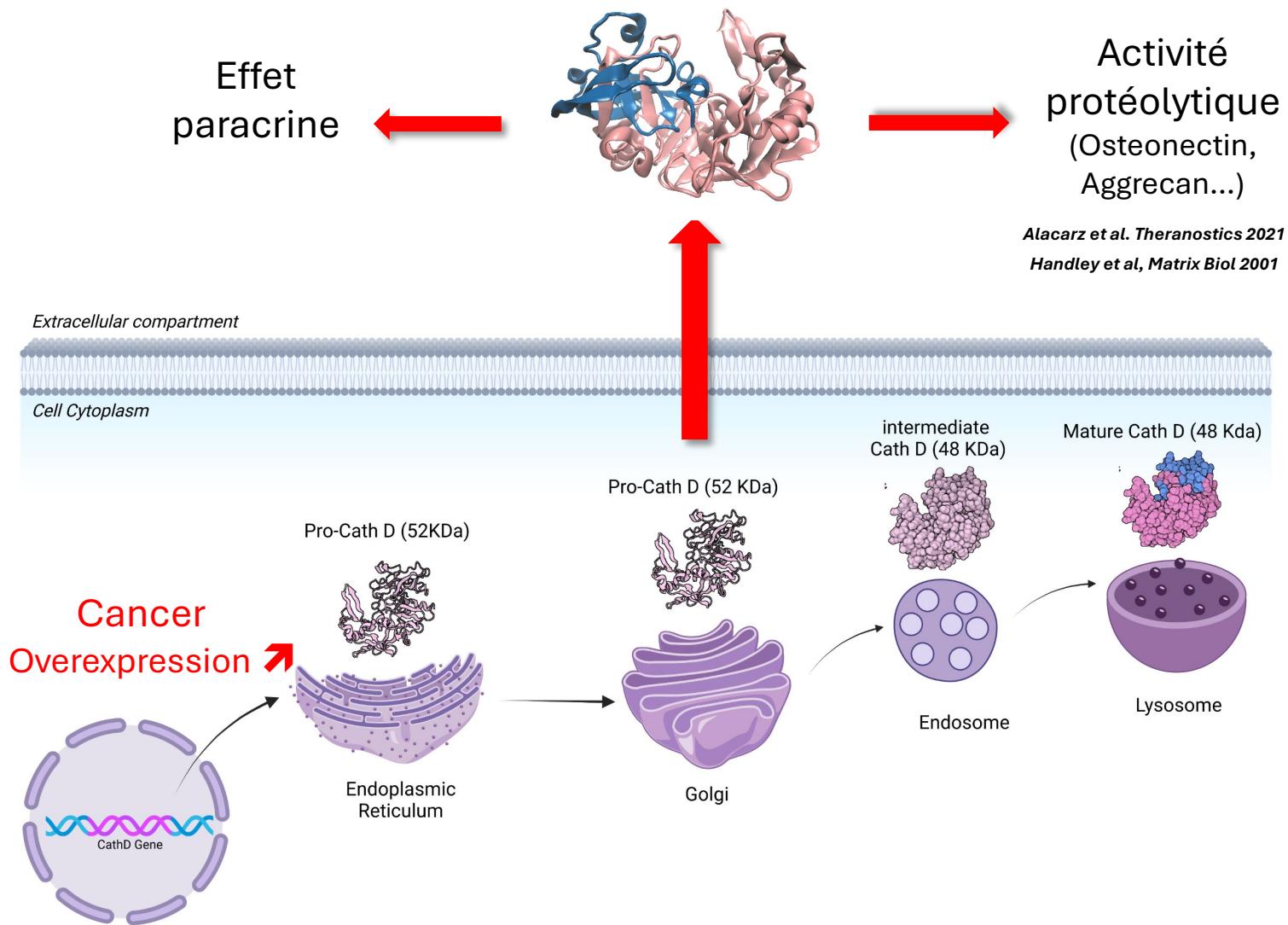
Tesnim Al-Khalifa, Elodie Lelièvre, Stéphanie Baud, Lise Chazée, Adeline Porcherie, Bruno Kieffer, Albin Jeanne, Stéphane Dedieu et Nicolas Etique

**UMR CNRS 7369 Matrice Extracellulaire et Dynamique Cellulaire
(MEDyC) Reims, France**

Equipe « Matrice extracellulaire, Cancer et cibles thérapeutiques »



La (pro-)Cathepsine D



(Pro-)Cathepsine D et stroma tumoral

JCB: ARTICLE

Catalytically inactive human cathepsin D triggers fibroblast invasive growth

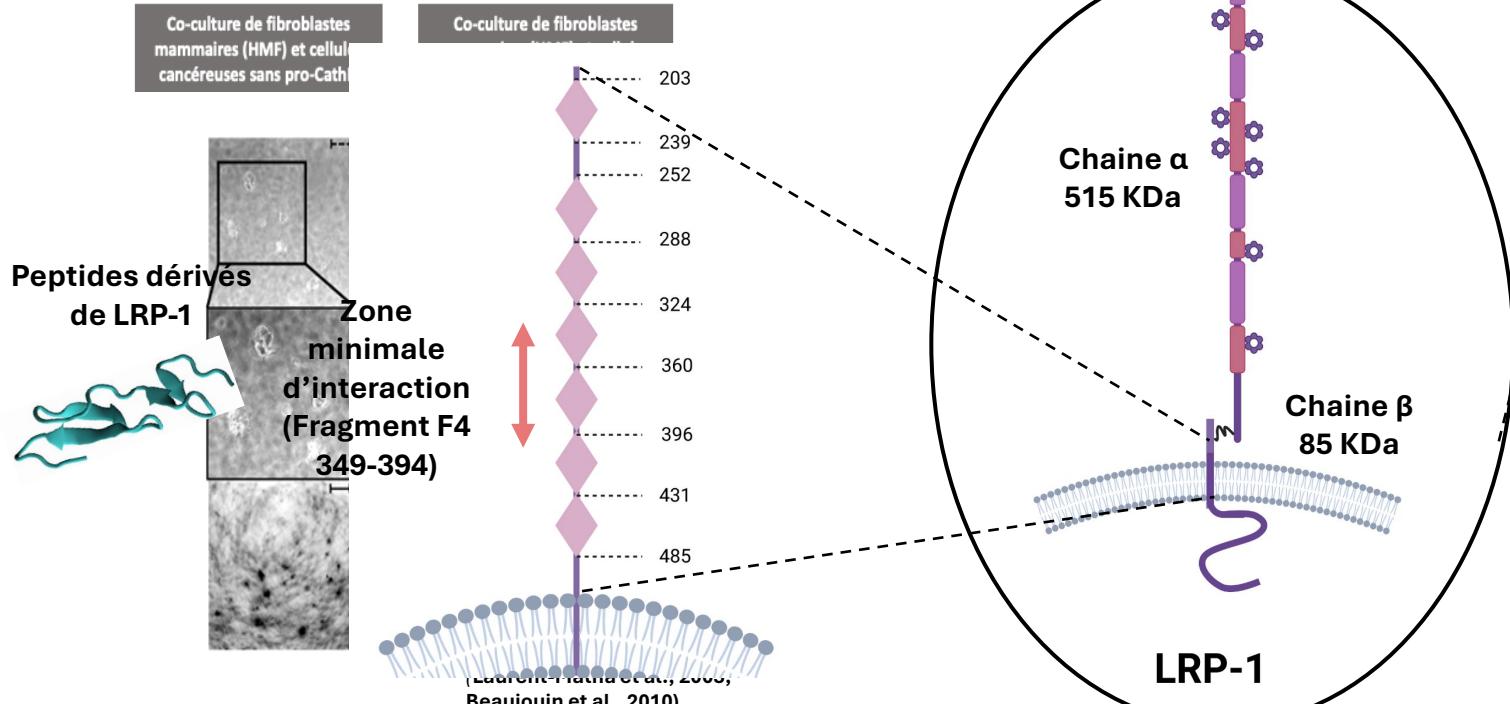
Valérie Laurent-Matha,¹ Sharon Maruani-Hermann,¹ Christine Prébois,¹ Mélanie Beaujouin,¹ Murielle Glondu,¹ Agnès Noël,² Marie Luz Alvarez-Gonzalez,² Sylvia Blacher,² Peter Coopman,³ Stephen Baghdiguian,⁴ Christine Gilles,² Jadranka Loncarek,⁵ Gilles Freiss,¹ Françoise Vignon,¹ and Emmanuelle Liaudet-Coopman¹

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Research Article

Pro-cathepsin D interacts with the extracellular domain of the β chain of LRP1 and promotes LRP1-dependent fibroblast outgrowth

Mélanie Beaujouin¹⁻⁴, Christine Prébois¹⁻⁴, Danielle Derocq¹⁻⁴, Valérie Laurent-Matha¹⁻⁴, Olivier Masson¹⁻⁴, Sophie Pattingre¹⁻⁴, Peter Coopman⁵, Nadir Bettache⁵, Jami Grossfeld⁶, Robert E. Hollingsworth⁷, Hongyu Zhang⁸, Zemin Yao⁸, Bradley T. Hyman⁸, Peter van der Geer⁹, Gary K. Smith¹⁰ and Emmanuelle Liaudet-Coopman^{1-4,*}

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⁶Genetics Research, GlaxoSmithKline, Five Moore Drive, Research Triangle Park, NC 27709, USA

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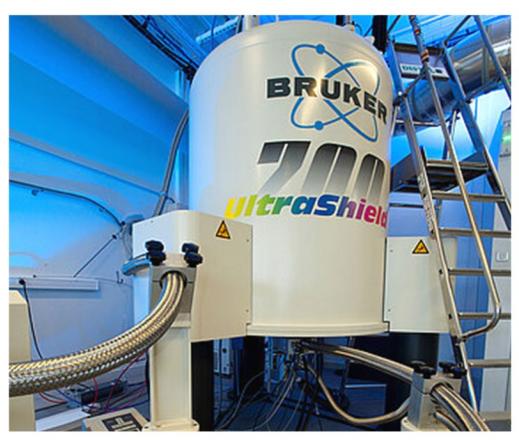
⁸Alzheimer Disease Research Laboratory, Massachusetts General Hospital, Harvard Medical School, Charlestown, MA 02129, USA

⁹Department of Chemistry and Biochemistry, San Diego State University, 5500 Campanile Drive, MC 1030, San Diego, CA 92182-1030, USA

¹⁰Screening and Compound Profiling, GlaxoSmithKline, Five Moore Drive, Research Triangle Park, NC 27709, USA

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Etude de l'interaction Cathepsine D/LRP-1

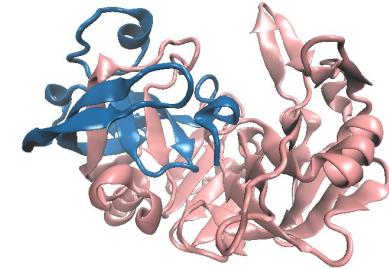


Collaboration Pr Bruno Kieffer
IGBMC

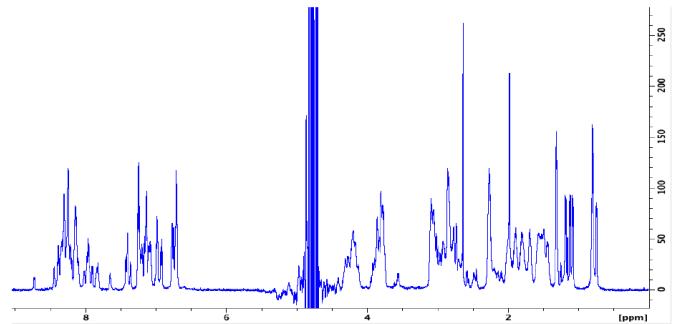


Collaboration Pr Stéphanie Baud
MEDyC

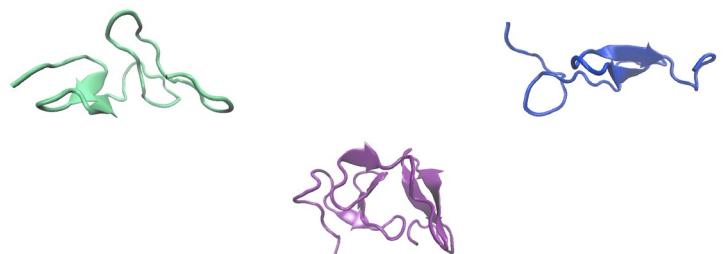
Choix des structures



Structure de la Cathepsine D (pCD)



Spectre RMN de la zone minimale d'interaction de LRP-1 (AA 349-394)



Simulations de dynamique moléculaire des fragments de LRP-1

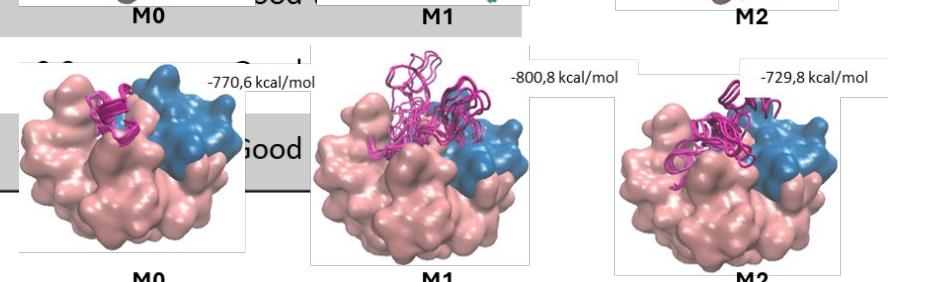
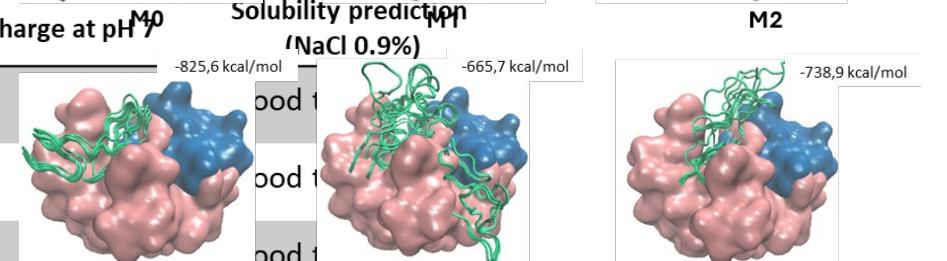
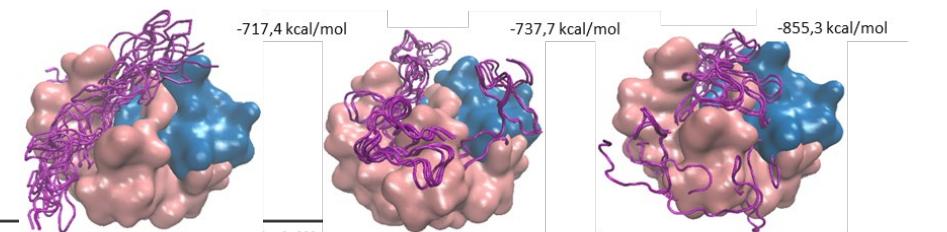
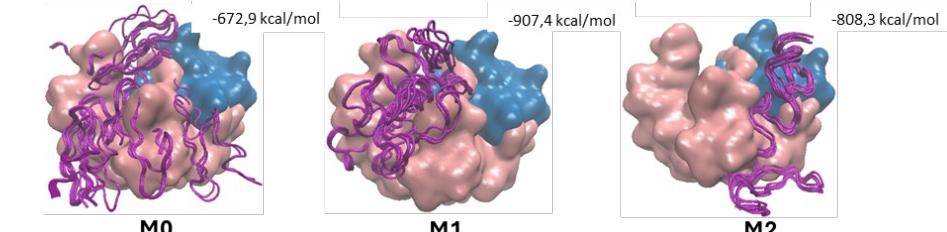
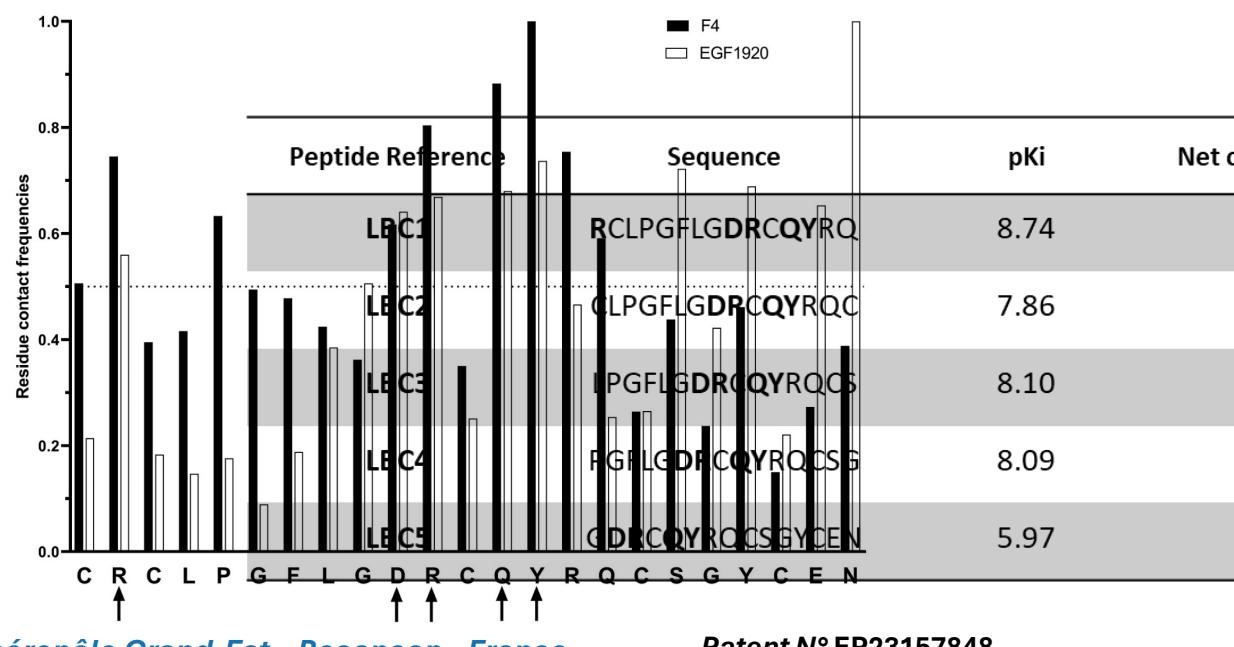
Identification des résidus clés de LRP-1

Chaine β LRP1

LRP1 45 AA (Fragment AA 349-394)

---STCTVNQGNQPQC**R**CLPGFLG**DRCQY**RQCSGYCENFGTCQMAADGSRQCRC**TAYFEGSRC**---

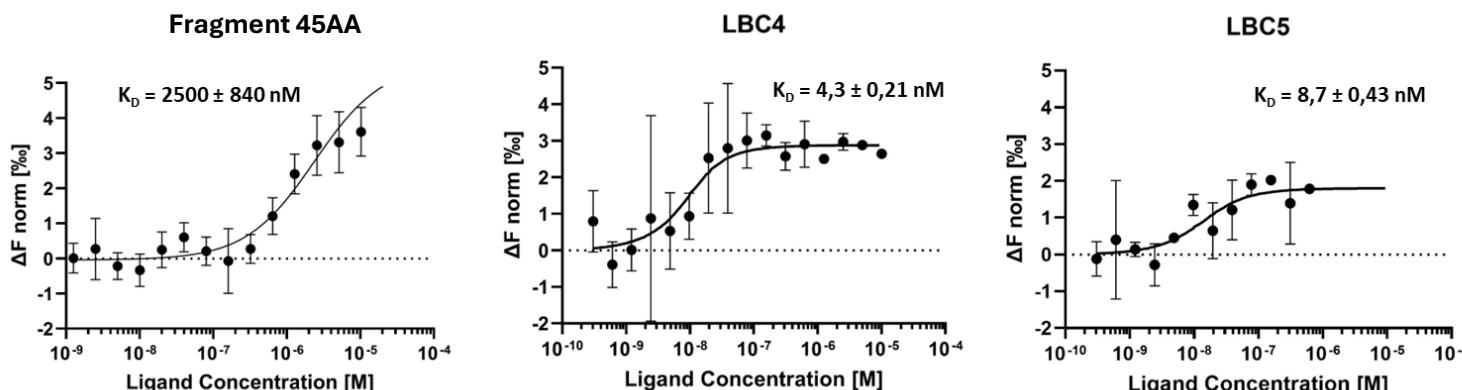
Design de 5 peptides
de 15 AA...



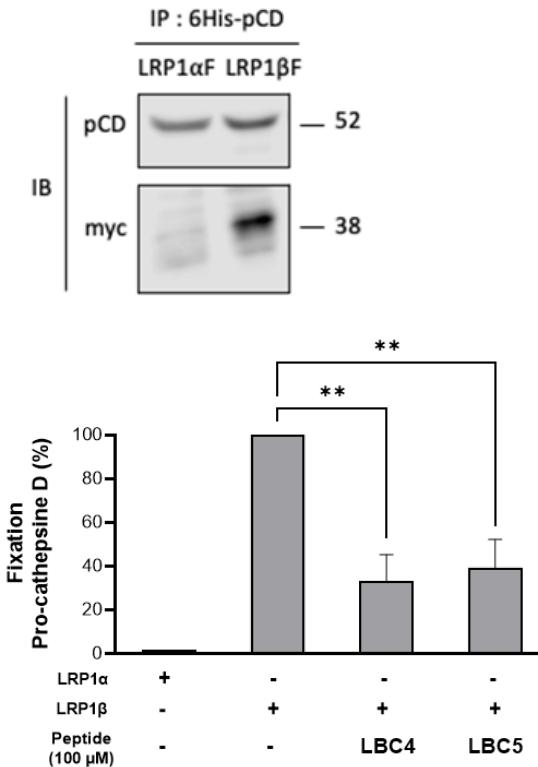
Test et sélection des peptides

- Etude de l'affinité avec la pCD par MST

Peptide	Sequence	Net Charge at pH 7	Water Solubility	K_D (nM)
Fragment 45 AA	CRCLPGFLGDRCQYRQCSGYCENFG TCQMAADGSRQCRCTAYFEGS	ND	Yes	2500 ± 840
LBC1	RCLPGFLGDRCQYRQ	1,9	Yes	52.5 ± 2.62
LBC2	CLPGFLGDRCQYRQC	0,8	Yes	$> 10\,000$
LBC3	LPGFLGDRCQYRQCS	0,9	Yes	340 ± 34
LBC4	PGFLGDRCQYRQCSG	0,9	Yes	4.3 ± 0.21
LBC5	GDRCQYRQCSGYCEN	-0,2	Yes	8.7 ± 0.43



- Etude du blocage de l'interaction par CoIP



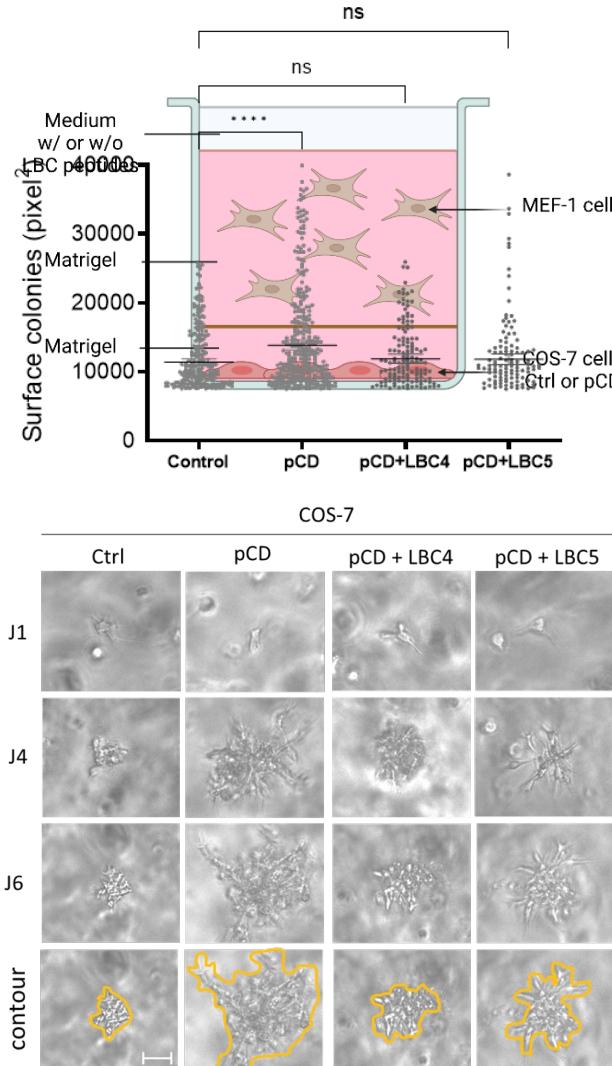
Les peptides LBC4 et LBC5 présentent chacun :

- une bonne affinité pour la pCD
- une capacité à bloquer l'interaction pCD/LRP-1

Effets *in cellulo* des peptides LBC4 et 5

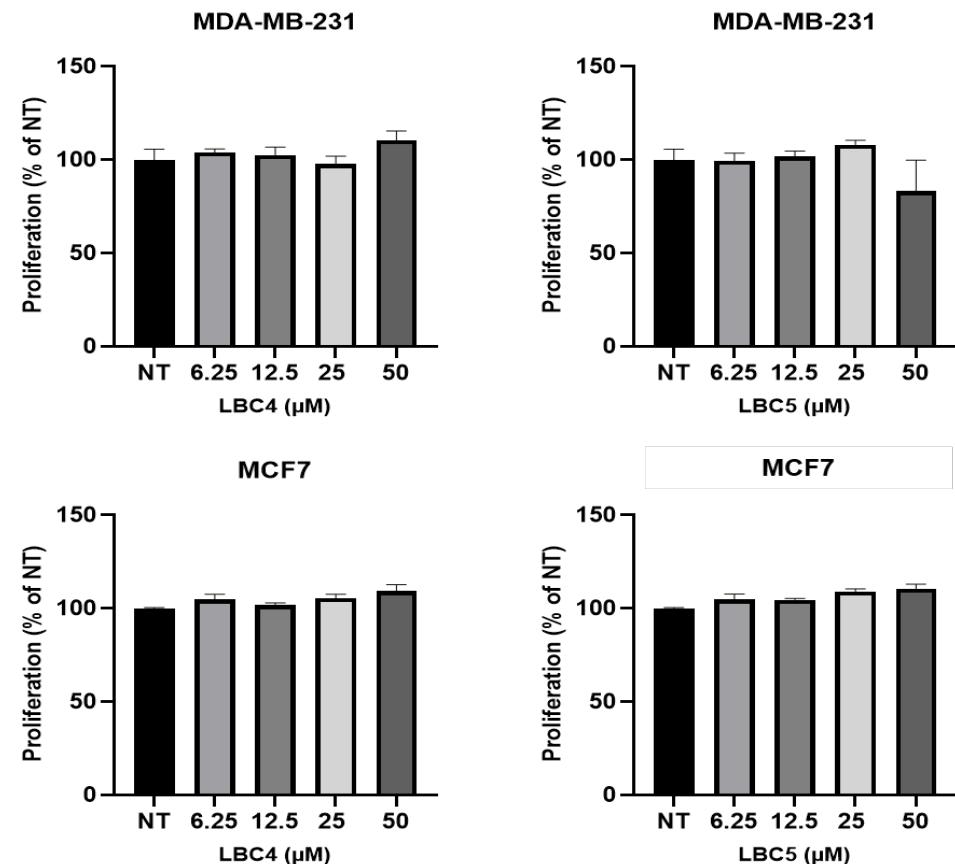
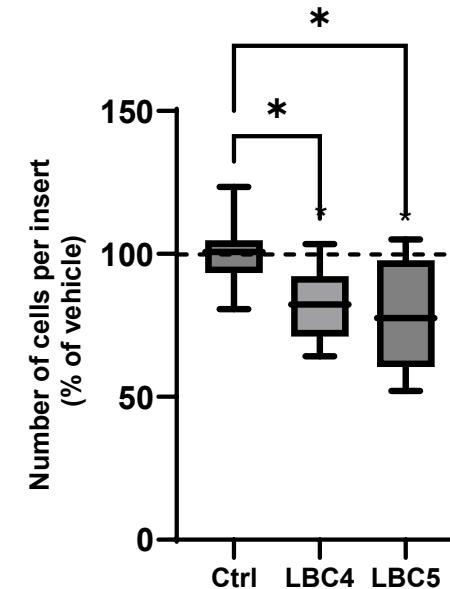
Cell Proliferation Assay

Sur les fibroblastes :



Sur les cellules tumorales :

Cell Invasion Assay

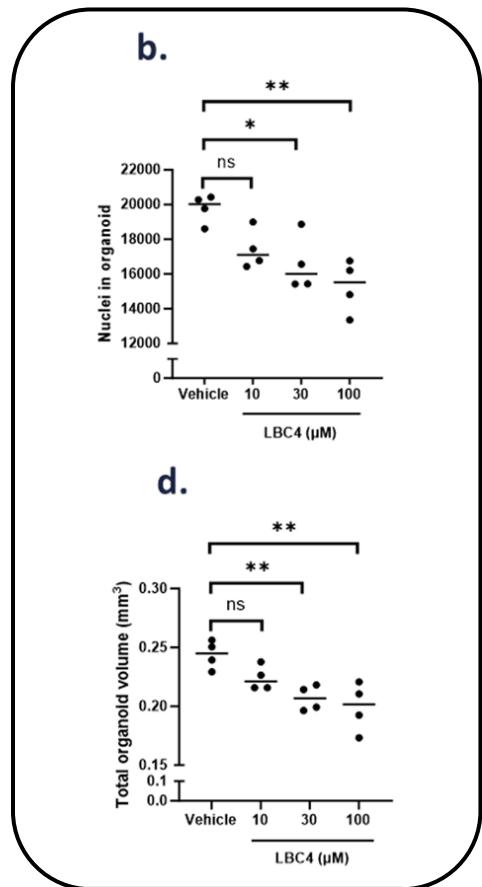
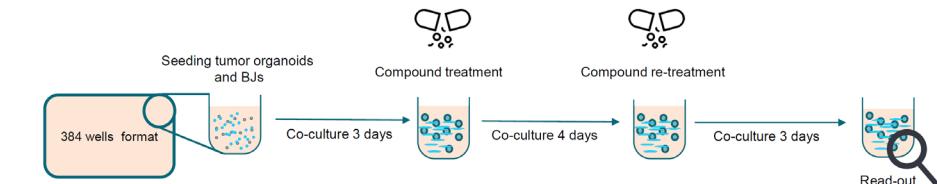


Les peptides LBC4 et LBC5 :

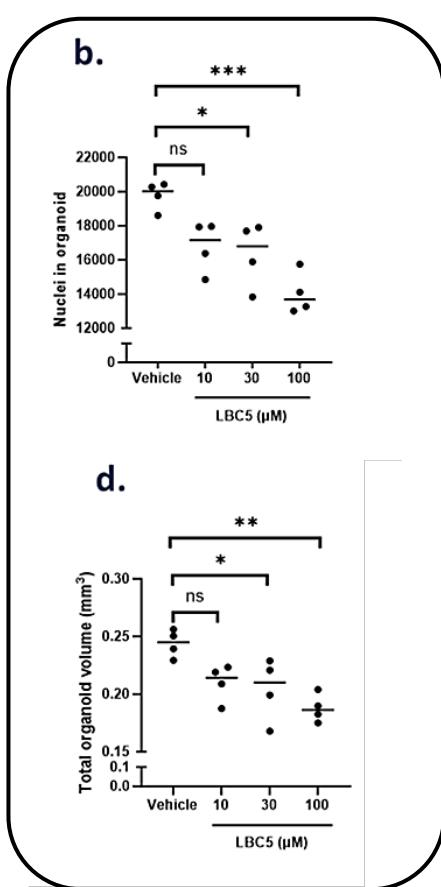
- Bloquent l'effet de la pCD sur la croissance des fibroblastes
- diminuent l'invasion des cellules tumorales
- Ne présentent pas d'effet cytotoxiques

Effets anti-tumoral des peptides LBC4 et 5

- Dans des organoïdes (PDO) issus de patientes TNBC...



LBC4



LBC5

- Dans un modèle immunocompétent (EO771/C57BL6)

J0: injection dans le Fat

Pad mammaire de cellules E0771



LBC4
(10 mg/kg)



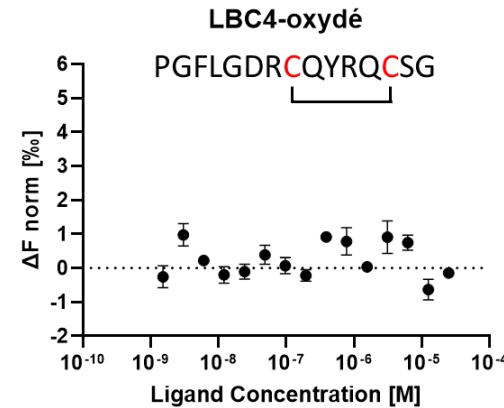
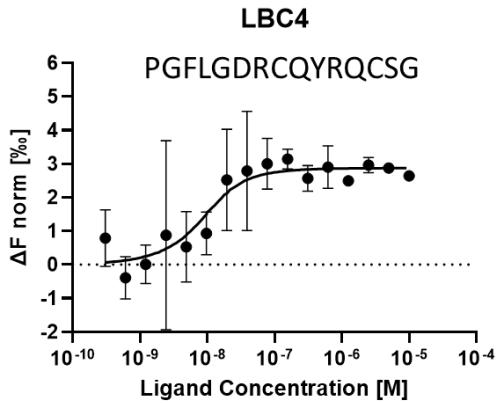
Traitement tous les 3 jours jusqu'à VT = 1500 mm^3

\approx J11 :
Randomisation et début traitement
(VT > 50 mm^3)

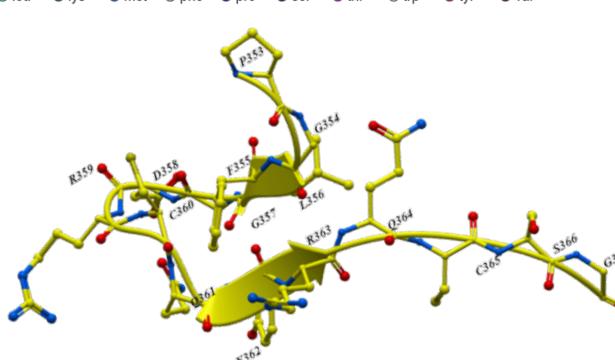
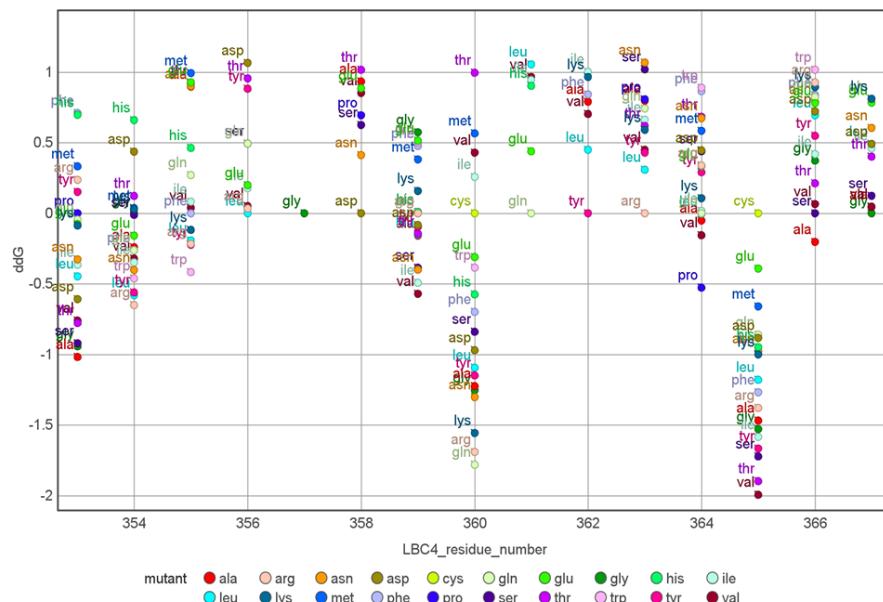
CONFIDENTIEL

Optimisation du peptide LBC4

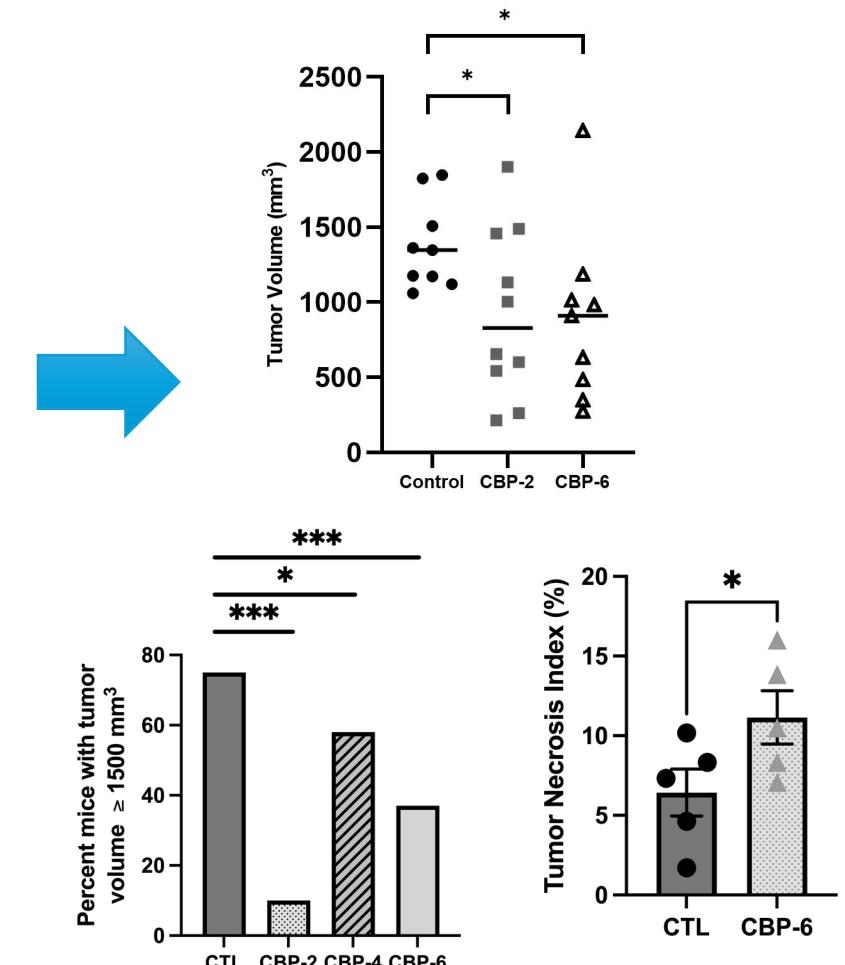
Perte d'affinité en milieu oxydant



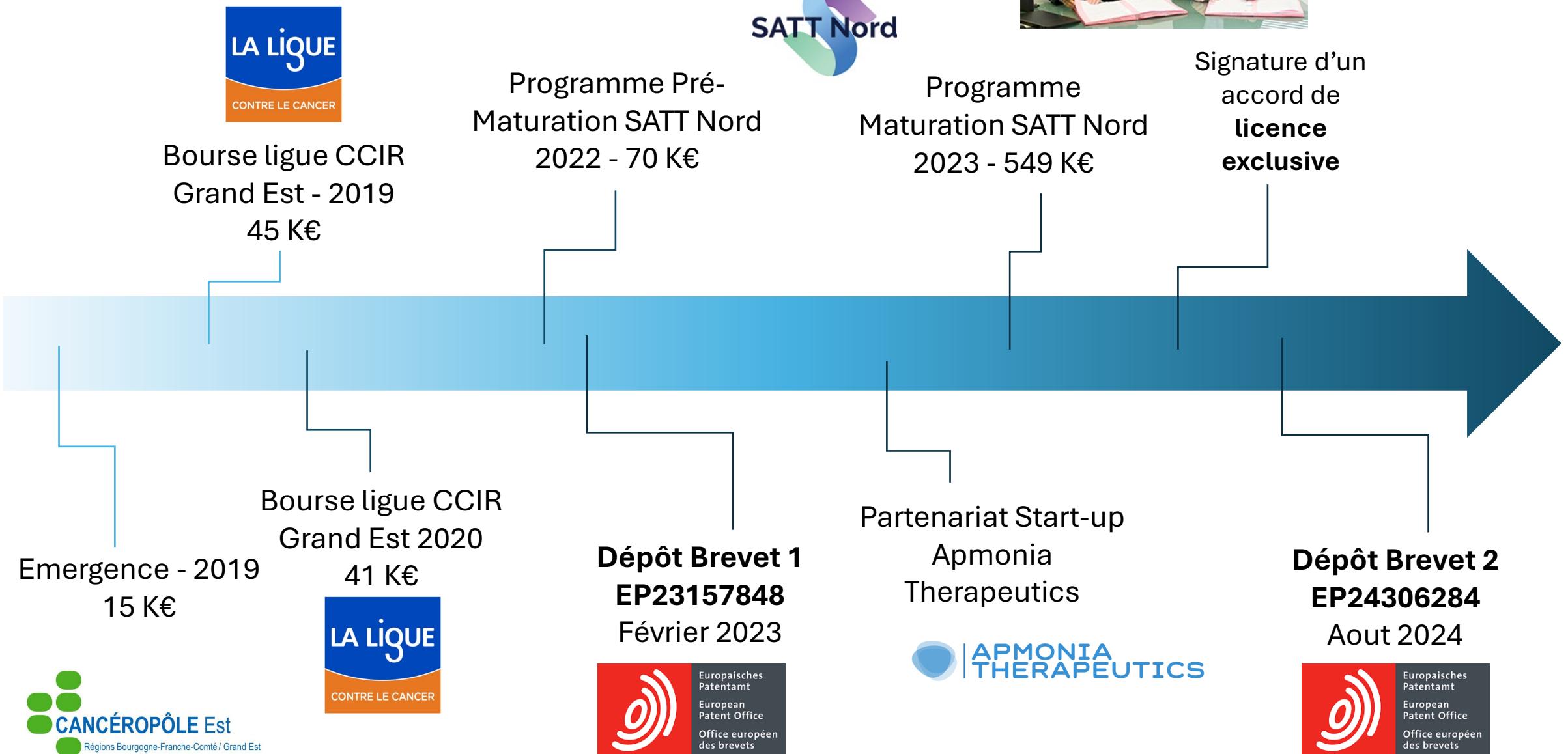
Optimisation du peptide LBC4 par mutagenèse in silico



Tests in vivo d'un nouveau lot de peptides optimisés dérivés de LBC4 (peptides CBP)



Le programme « Emergence », un levier



Remerciements

UMR CNRS/URCA 7369 MEDyC, Reims

Dir. Pr. DUCA Laurent

Co-Dir. Dr BREZILLON Stéphane

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GRIMBERT Frederic

Dr BRAURE Manon

DELBECQUE Axelle



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Dr PORCHERIE Adeline

ETIENNOT Marion



FINANCEURS

