

AI to redefine Antibody discovery workflow

Who are we?

French TechBio company

Spin-off from CNRS/INRAe

2017



Artificial Intelligence for antibody drug

Unique technology and expertise in Europe for applying AI/ML for drug discovery

Early adopters purchased our products

Based in EU, US, South Korea and Israel, biotech & pharma companies

40



Antibodies characterized

Applied wihtin 9 therapeutic areas, can be used for other biodrug

1st Artificial Intelligence based vaccine

Al used to design peptide for CoVepiT, OSE Immunotherapeutics



Our DNA is to combine Computational and Biology

A team of 15 people dedicated to your success

- 10+ years of research in AI/ML in drug discovery
- International experience working on US (DARPA) and European programs
- 100+ peer reviewed scientific publications and 4 patents
- KOLs in immuno-oncology (French hospital, Medtech, Biotech and pharma companies)
- Make antibody faster, better and safer using Al

POWERED BY:







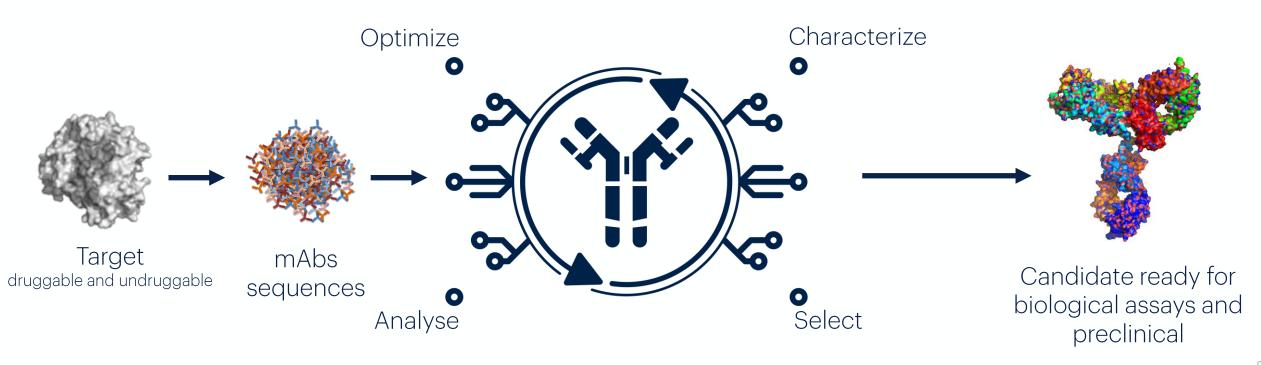




We transform years into 1 month with digitalization

MAbSilico solution - Computational antibody drug design

MAbFactory - AI & ML



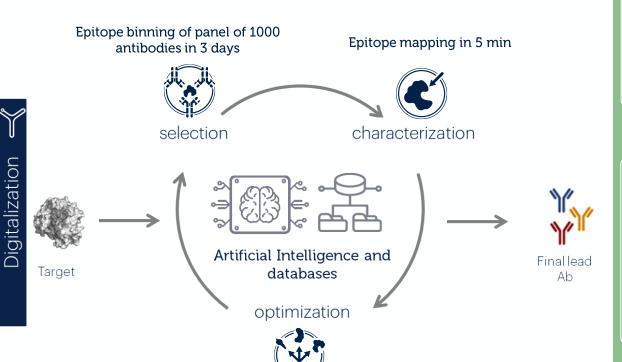
- Decrease the risk, Accelerate drug discovery by x500, all therapeutic areas
- Pick and design candidates ready for biological functional assays
- Y Parallelization of billion of simulations to design the best candidate to the target



Digitalization of drug discovery

Only antibody sequences and target name required

No new data training/ML required



Forecast off-targets in minutes

- Accelerate 500x drug discovery
- Increase 5x the success rate of market approval
- **Reduce 10x drug discovery costs**
- **Expand IP protection**
- Validated by industry (pharma & biotech)

MAbSilico assets & achievements

Databases proprietary and public

- ¥ 4000 structures
- 70M sequences
- ¥ 400k Abs, 3k targets
- ¥ 1300 Abs with kd



- Y Patents, publications
- Y Next generation sequencing (NGS)
- * Public databases

3 algorithms

Docking Protein-protein docking



Similarity CDR similarity measurement



Affinity KD prediction





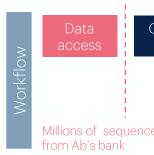
Antibodies drugs in preclinic and clinic

2 Abs in preclinic, 2 Abs in clinic (phase 2 or 3) and 1 peptide-based vaccine in phase 1 from early adopters



MAbSilico workflow - Computational antibody drug design

Parallelization of million of simulations in days



Competitive analysis

Ab identification

Epitope mapping

Affinity prediction & ranking

Ab formatting PTM + Hydrophobic + Immunogenecity

Millions of sequences

Target selection for data extraction & modeling

Off-target prediction

> Delivery of up to 50 Ab sequences with complete characterization





Competitive

landscape

analysis of target of interest



Abs sequences (scFv, VH, VL)



MAbSubstitute FTO antibody proposal



MAbCross Off-targets identification



MAbTope Epitope mapping



MAbAffinity Affinity prediction



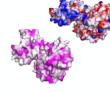
MAbHuman Humanization



MAbProductivity PTM, hydrophobicity and immunogenicity



Target explorer MAbFactory



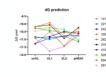
Encodina into one sinale alphabet



Screening against original patented sequences



Give comprehensive insights about the original target and how the selected antibody works



Compute in silico AG values (Regression and Classification tasks) & rank lead candidates



Screen MAbSilico database of human frameworks to select appropriate scaffold



Immunogenic peptides' regions Hydrophobic patches and immunogenic peptides sequences



Functional assays Manufacturability Kd/ΔG measurements Epitope Validation

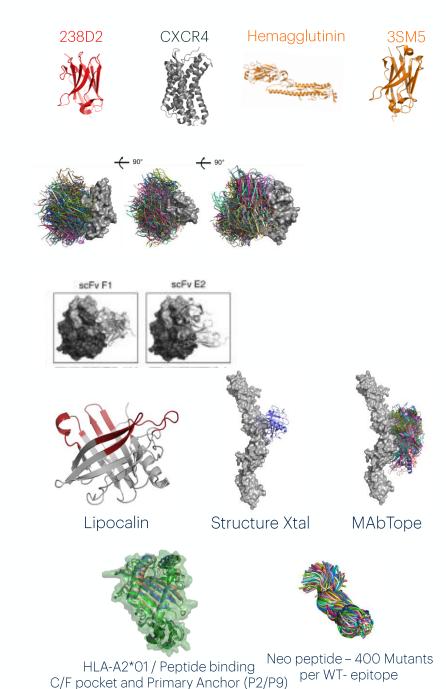
AI used for biologics drug discovery

- **VHH:** method of CDR similarity developed on VHH against CXCR4 (off-target prediction and binder replacement)
- **Bispecific Ab:** BiCKI® platform of OSE Immuno with anti-PD-1 as central skeleton (OSE-279)
- ** CAR-T: ScFv can be used for CAR-T, work done on anti-cath-D

 A. Yahya et al., BMJ 2019

Novel format: proof of concept for epitope mapping on lipocalin (4 hyper-variables loop ~ 4 CDRs)

Peptide: in silico affinity maturation for CoVepiT (covid19 vaccine in clinical trial) <u>V. Gauttier et al., BioRxiv 2020</u>



05

Use cases of drug design & discovery programs

Computational peptide design for covid-19 vaccine - CoVepiT clinical phase I

Design new lead candidates against RBD from SARS-CoV-2

End-to-end framework that can serve as anti-pandemics platform

Design of an anti-A2A design using MAbSubstitute

Design new lead candidates to replace a breast cancer triple negative drug

Use of MAbFactory to select and design of anti-TIGIT

MAbFactory: SaaS platform for antibody characterization

Examples of MAbTope, fast and high-throughput unbiased computational epitope mapping

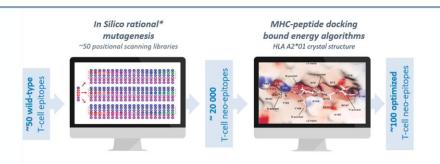
	Dupilumab / IL4Rα	OSE-127 / IL-7Rα	Eculizumab / C5	OSE-230 / ChemR23	Lipocalins	Time
⊕ + 0 → (⊕) + ⊘ + ⊗	 The Dupilumab (DUPIXENT®) is targeting the IL4Rα and is FDA-approved for Eczema, Atopic dermatitis, Several forms of severe asthma. Epitope was not known until 2020 	• New Ab targeting the IL- 7Rα to prevent the expansion of pathogenic memory immune cells and chronic autoimmune attacks.	 Eculizumab targets the complement C5 and is indicated in nocturnal paroxystic hemoglobinuria (NPH) and in atypical hemolytic and uremic syndromes (aHUS). The epitope was only partially solved and did not explain the biological observations (absence of interspecies crossreactivity) 	New Ab targeting ChemR23 (GPCR) which is overexpressed at the inflamed site of IBD, Arthritis, CLE, ANCA vasculitis, Asthma, Sepsis, COPD, ARDS, CHP, COVID-19 New concept in the management of severe & chronic inflammation	Lipocalins are a novel class of binding molecules with 4 hyper-variables loops (~ 4 CDRs) which transport small hydro- phobic molecules	• Al vs. biology, from minutes to months
MAbTope	Very highly probable Highly probable Probable Possible	Very highly probable Highly probable Probable Prossible	90° Highly probable Probable Possible	Very highly probable Highly probable Probable Prossible	transthyretin fibronectin CTLA-4	Minutes
Biological assays	Experimentally Crystallography validated (PDB: 6wgl) residues (cytometry)	Peptide array HDX	Experimentally validated residues (peptide array) C5-pep1 C5-pep2 Page 10	Experimentally validated peptide (ELISA)	Crystallography PDB:2WQA PDB:3BX7	Months
Publications	Tahir et al., 2021	nature communications Lyssia et al., 2018	MOLECULAR IMMUNOLOGY Brachet et al., 2017	Science Advances Trilleaud et al., 2021		

Computational peptide design for covid-19 vaccine

Clinical phase 1

Design a multi-target CD8 T cell peptide COVID-19 vaccine targeting for design targeting several structural (S, M, N) and non-structural (NSPs) SARS-CoV-2 proteins.

- Find potential universal vaccine against future coronaviruses.
- Adapt an immunologically optimized multineoepitopes-based peptide vaccine platform for SARS-CoV-2.
- Identify and optimize peptides' immunogenicity using computational approach.



^{*} Random mutagenesis -> 28 000 billion peptides possibilities Rational design based on OSE Know-How -> 20 x 103 peptides

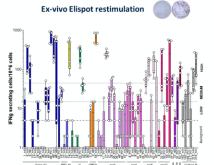




- 55WT epitope out of 35.000 identify as best HLA-A2 binder with strong immunogenicity.
- Design of 55 neoepitopes validated experimentally (IFN_γ measurement) which induce strong proportion of virus specific CD8 T-cells.
- First human data: Immunodominant epitopes tested in COVID-19 convalescent positively versus unexposed subjects.
- In clinical development phase 1



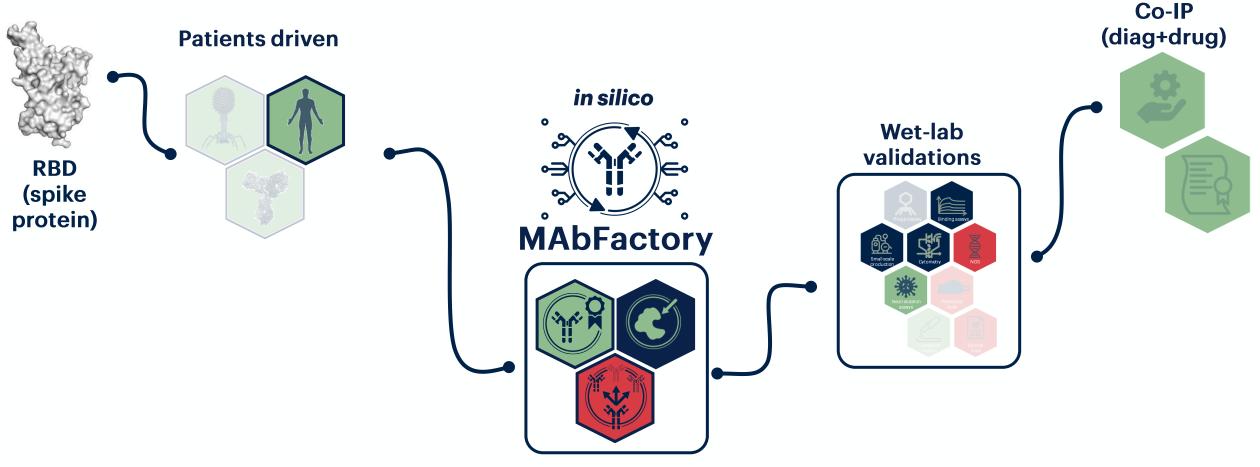
C/F pocket and Primary Anchor (P2/P9)





End-to-end framework that can serve as anti-pandemics platform, from patients to lead candidates

Humabdiag <u>project</u> - ongoing



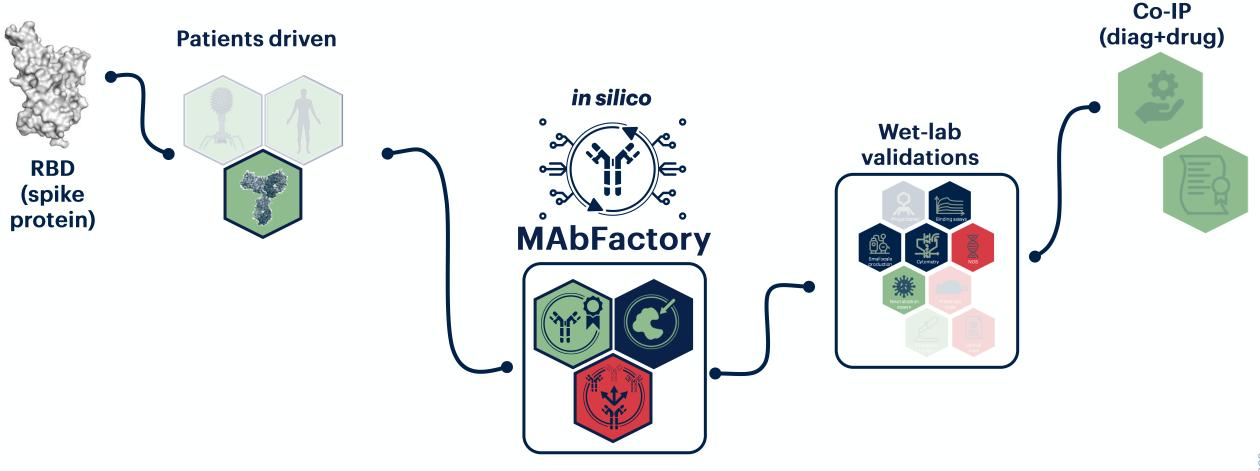








End-to-end framework that can serve as anti-pandemics platform, from patients to lead candidates MabCOVID- ongoing











Design new lead candidates against RBD from SARS-CoV-2 Ongoing project

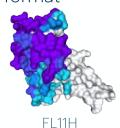


RBD

- Wuhan-neutralizing IgG isolated from patients as starting Ab
- Design of FL11H and FL24H using NGS sequences

candidates

- Two distinct predicted epitopes
- Production in diabody (VH-VH) format



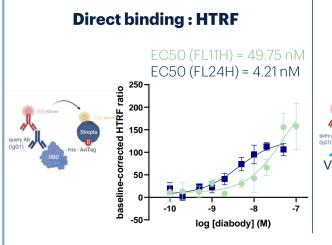


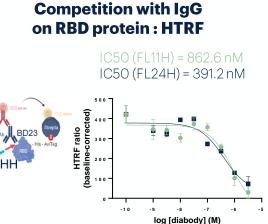


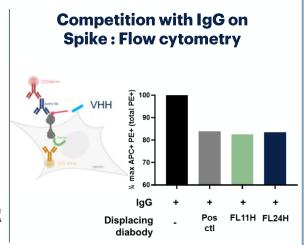
Conclusions

- Potential therapeutic mAb design by computer in days
- Format to optimized
- Effect on Omicron variant
- Ongoing: affinity maturation

In vitro validation of binding and neutralization effect



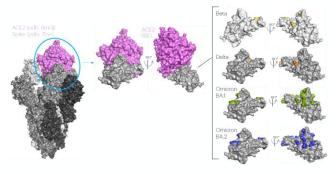




Neutralization assays (IC50 concentration in µg/ml)

Antibody	Wuhan	Delta	Omicron
FL11H (diabody)	4,3	1,4	Ongoing
FL24H (diabody)	28,6	9	Ongoing
Imdevimab (IgG)	0,006	0,006	Ongoing
Hyb. (IgG)	0,044	0,39	Ongoing

SARS-CoV-2 variants

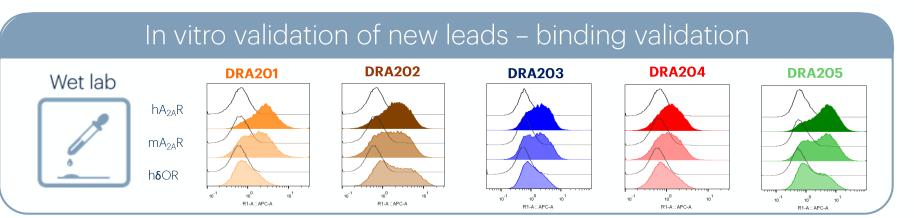


MAbSubstitute

Use cases

$\overline{\mathsf{A}_{2\mathsf{A}}\mathsf{R}}$ $(\mathsf{A}_{1},\mathsf{A}_{2\mathsf{A}},\mathsf{A}_{2\mathsf{B}},\mathsf{A}_{3})$

- GPCR (Gi & β-arr)
- P1 class purinergic receptor, Autism Spectrum Disorder, Metastatic spread, Dementia, Sleep disorders



Leads

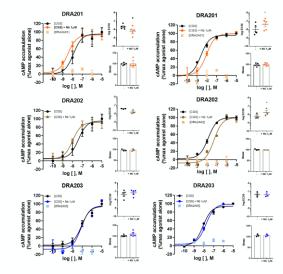
- Original: ScFv, Antagonist
- New: 5 sequences VHH

Conclusions

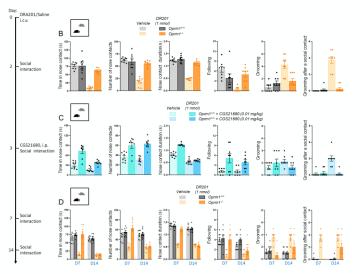
- Multiple alternative binders
- Allosteric Modulation & Cross specific
- In vivo validation

In vivo validation of new leads and biological activities

Pharmacological profile of candidate antibodies on human and murine A2a receptors



Effects of DR-2A-O1 administered ICV on social interaction in the *Oprm1* knockout mouse model of autism



Design new lead candidates to replace a breast cancer triple negative drug Ongoing project



TROP2

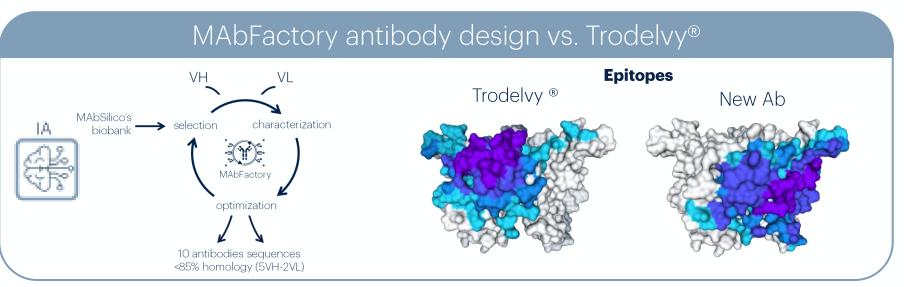
- Trophoblast cell-surface antigen 2 –TROP2
- Type I transmembrane glycoprotein, Triple Negative Breast Cancer, Valuable Biomarker in metastatic spread

Leads

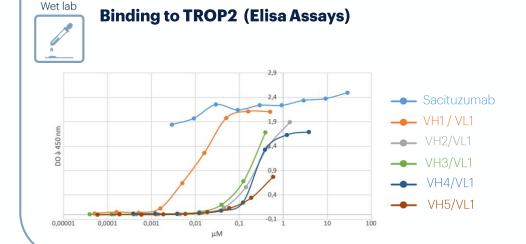
- Sacituzumab; therapeutic ADC antagonist Trodelvy®
- New: 10 sequences (5 VH and 2 VL, in scFv)

Conclusions

- Multiple alternative binders
- Allosteric Modulation & Cross specific
- In vivo validation





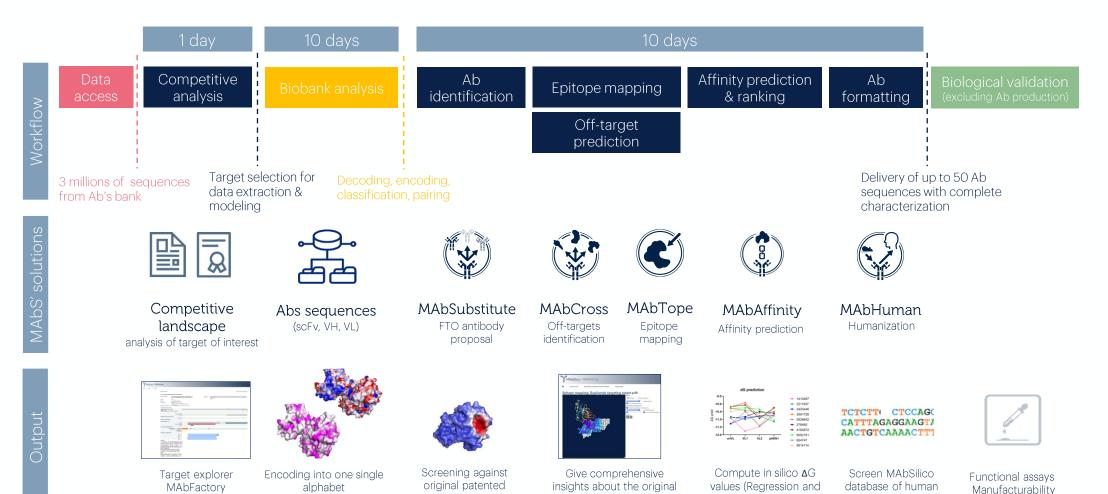


Binding to TROP2 (Monolith) Sacituzumab expressed as ScFv for comparison

Antibody (scFv)	Affinity	
sacituzumab	285 nM	
VH A / VL 1	287 nM	

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New leads against TIGIT selected & designed from a naïve bank of 3M antibodies



sequences

target and how the

selected antibody works

Classification tasks) &

rank lead candidates

frameworks to select

appropriate scaffold

Kd/∆G measurements

Epitope Validation

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Use of MAbFactory to select and design of anti-TIGIT (part 2)

Ongoing project

TIGIT

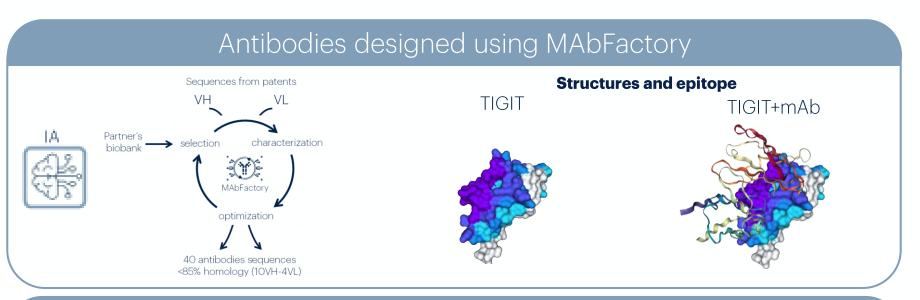
- T-cell immunoreceptor with Ig and ITIM domains
- Immune check-point

Leads

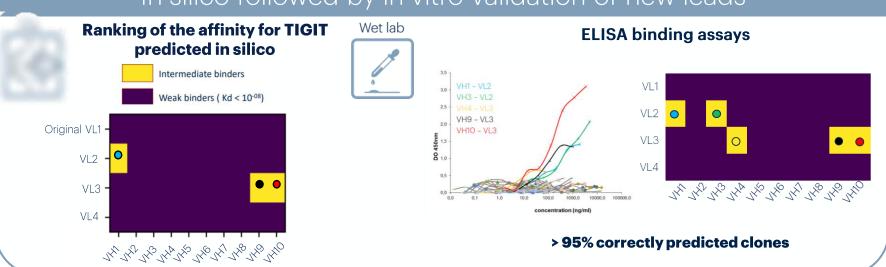
- 10 anti-TIGIT disclosed in patents
- New: 40 sequences (10 VH & 4
 VL, in scFv) from 3M sequences

Conclusions

- Selection of binders from biobank in days
- in silico humanization of murine lead candidates
- Affinity predicted validated by biological assays

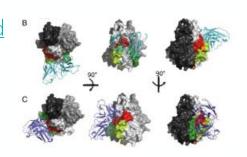


In silico followed by in vitro validation of new leads

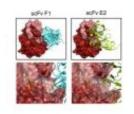




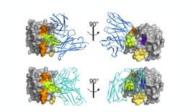
*MAbTope: A Method for Improved Epitope Mapping, J Immunol October 15, 2018



*Immunotherapy of triple-negative breast cancer with cathepsin Dtargeting antibodies, Journal for **ImmunoTherapy of Cancer** - Feb 2019, 05th



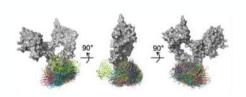
¥4C3 Human Monoclonal Antibody [...], **Front. Immunol.,** 25 September 2020



*A recycling anti-transferrin receptor-1 [...], **mAbs** - 15 Dec 2018



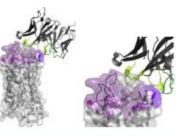
* Biasing human epidermal growth factor receptor 4 [...], Cancer Sciences, 16 May 2020



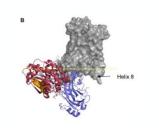
Y Agonist anti-ChemR23 mAb [...], Science Advances, 02 Apr 2021

YIL-7 receptor blockade blunts antigen-

specific memory T cell [...], NATURE



Y G protein-dependent signaling triggers a β-arrestin-scaffolded[...]



FASEB J. 32,1154-1169 (2018)



COMMUNICATIONS (2018)

¥1 EU patent (<u>WO2018087494A1</u>)



Ready to use the AI-driven solutions for antibody discovery faster, safer and better



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