



ProMIS Neurosciences: Selective targeting of pathogenic misfolded proteins, based on a proprietary discovery platform

**Toronto Stock Exchange (TSX) ticker: PMN.TO
OTCQB ticker: ARFXF.**

October 22, 2021

Forward looking statement: safe harbor

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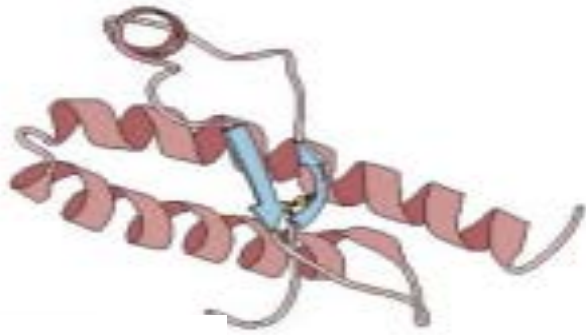
ProMIS Summary

- Differentiated ***technology platform*** – Computational approaches to rational design of selective antibodies, a unique ProMIS capability
- ***Growing portfolio*** of antibodies ***selective for mis-folded proteins*** implicated in neurodegenerative diseases
- High selectivity a ***ProMIS competitive advantage***. Lack of selectivity for mis-folded proteins likely the primary source of failures or limited success in prior competitor programs in neurodegenerative diseases
- ***Lead program PMN310 potential "best of the next generation" antibody therapy in Alzheimer's disease***: highly selective for toxic oligomer form of amyloid, differentiated from likely first generation products from Biogen, Eisai, and Lilly; PMN310 differentiated from first oligomer selective antibody from Acumen
- ***Growing portfolio*** contains differentiated, selective antibodies against toxic forms of TDP-43 (ALS, FTD); alpha synuclein (PD, MSA), tau, SOD1, and others
- Fluid-based biomarkers may enable ***rapid and capital efficient path to clinical readout*** and value inflection for all programs
- Multiple programs 21-36 months from clinical data – IND enabling work followed by SAD/MAD trial in patients

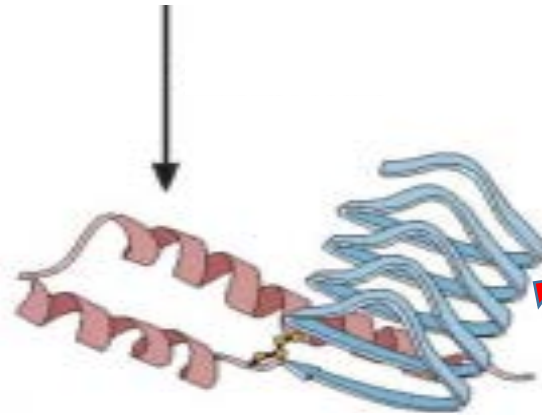
ProMIS Platform - Summary

- ProMIS is founded on a ***unique, revolutionary approach tailoring antibodies to defined targets***, using an interdisciplinary synthesis of physics and biology
- IBM's quantum computing group in April 2020 predicted that within five years [i.e. by 2025] quantum computing will enable ***"developing novel biologic products based on protein folding predictions"***
- ***ProMIS is 10+ years ahead*** of that prediction - by focusing on the problem of predicting conformational epitopes (amino acid sequence ***and shape***) exposed only on pathogenic, ***mis-folded proteins***
- Through sophisticated scaffolding, and other methods refined over several years, ProMIS creates ***peptide antigens*** that reproduce the amino acid sequence and conformation of the predicted epitopes to be used for immunization and generation of monoclonal antibodies.
- **ProMIS at AAIC 2021** – "*Conformational epitopes exposed on misfolded toxic forms of amyloid-beta, tau and alpha-synuclein directly contribute to their seeding activity*" - potential breakthrough in scientific understanding of disease, exposed epitopes play a direct role in disease progression, additional mechanism for ProMIS antibodies
- Continuous discussions with large pharma over the last five years have confirmed that as of now, to the best of our knowledge, ***no one has an antibody "rational design" capability like ProMIS'***

Mis-folded proteins have the same amino acid sequence as normal proteins...the only difference is the shape...ProMIS identifies conformational epitopes exposed only on mis-folded proteins



Normal protein – folds into a specific shape to perform its physiologic function



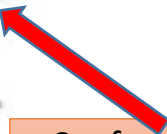
Toxic mis-folded form

Mis-folded protein...improper folding exposes toxic portions of the protein.....in a particular shape or conformation...



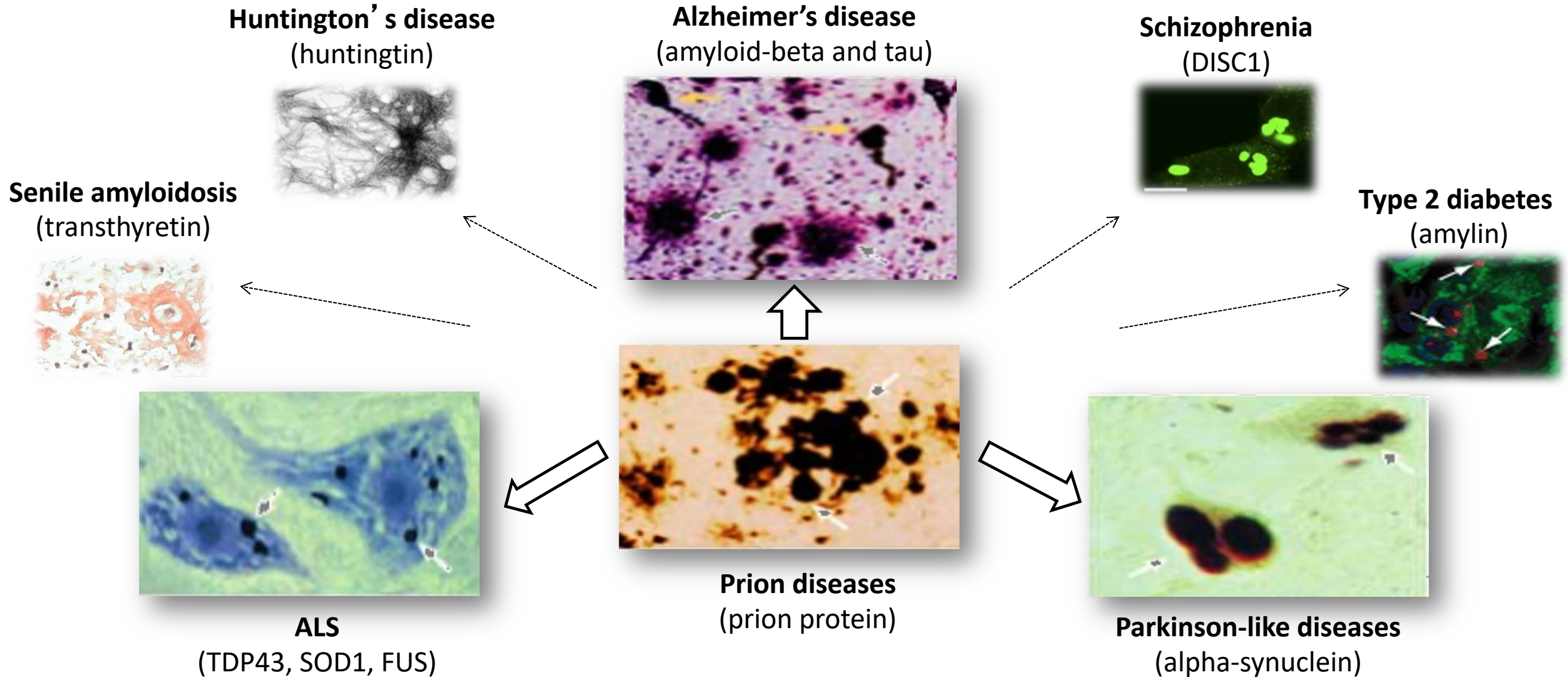
ProMIS platform predicts conformational epitopes – both amino acid sequence and shape, only exposed on toxic mis-folded proteins....

Conformational
Epitope predicted
by ProMIS
platform



Immunizations with those epitopes lead to selective antibodies

Alzheimer's, Parkinson's and ALS are protein misfolding diseases, where the toxic mis-folded proteins propagate in a prion-like manner



The ProMIS platform generates antibodies selective for the misfolded toxic forms of pathogenic proteins

Identification of epitopes selectively exposed on toxic misfolded form of the target protein using predictive computational algorithm



Immunization with disease-associated epitope and screening of monoclonal antibodies with desired binding profile and protective activity

Neutralize the toxic misfolded form

Don't interfere with the normal form, critical for brain health

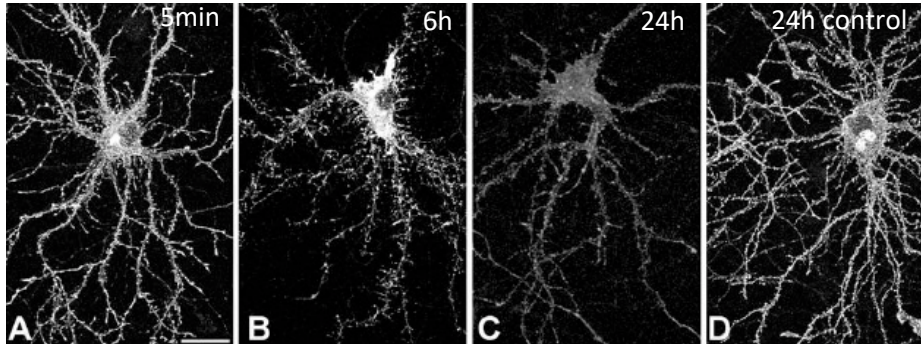


Successful track record thus far in generating antibodies selective for pathogenic forms of amyloid-beta, tau, alpha-synuclein, TDP-43, SOD1

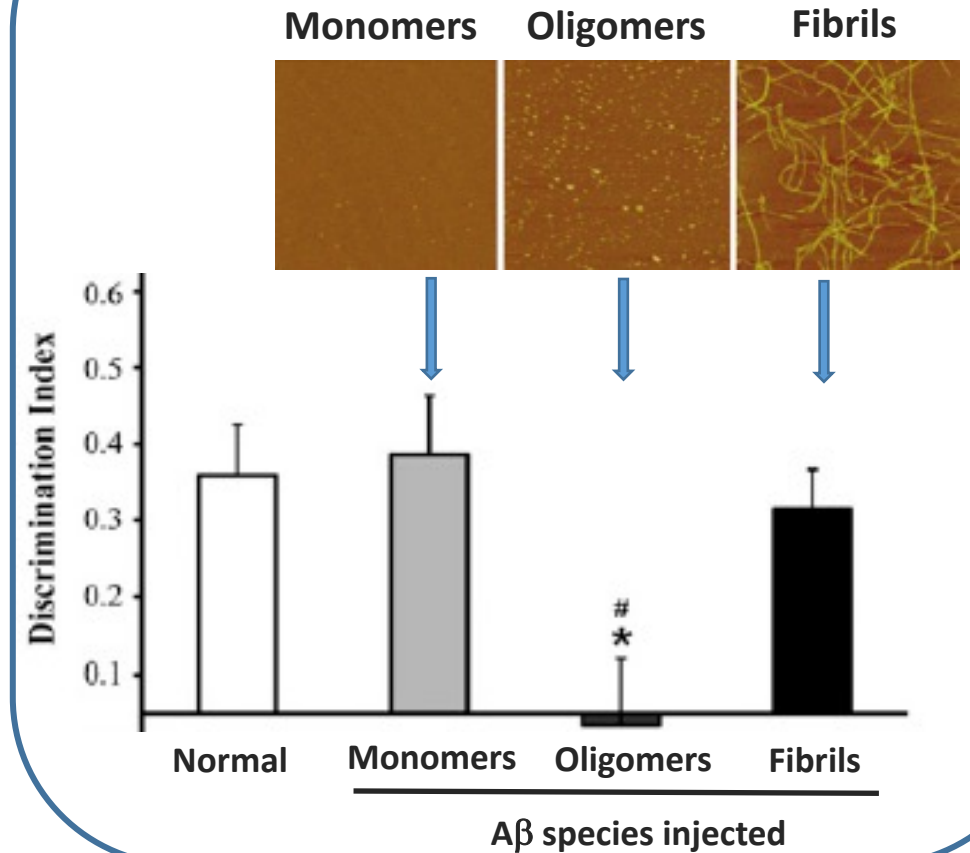
Alzheimer's disease: soluble toxic Amyloid-beta oligomers – not plaque or monomers – are the most neuropathogenic A β species

- Synapse abnormalities and memory impairment correlate poorly with plaque burden in human and mouse AD^{1,2}
- A β monomers and A β insoluble fibrils (plaque) have little or no demonstrable toxicity in vitro or in vivo³⁻⁵
- Soluble A β oligomers show the highest degree of neurotoxicity⁶
 - Toxicity in primary neuron cultures and brain slices^{3,5,7-9}
 - Induction of cognitive impairment in rodents^{3,4,10}

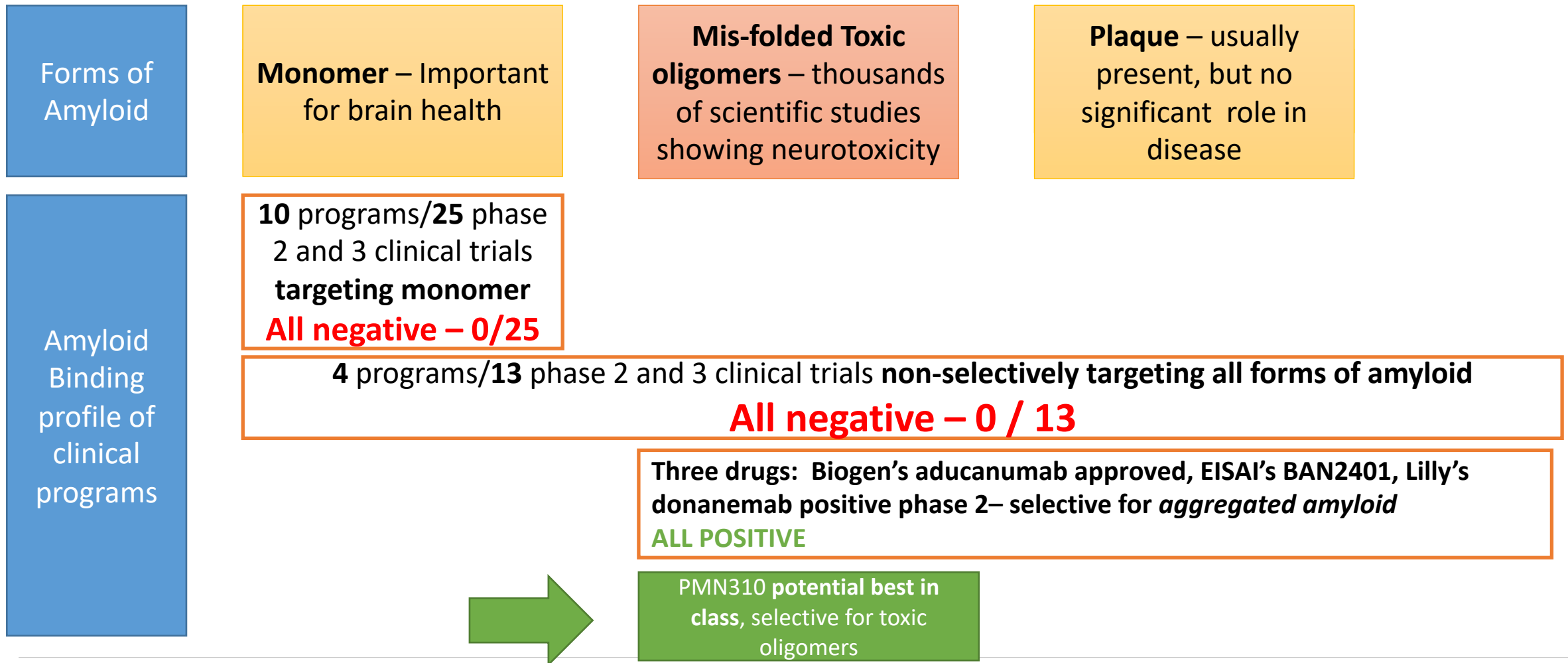
Synaptotoxicity of human A β oligomers on hippocampal neurons in vitro⁷



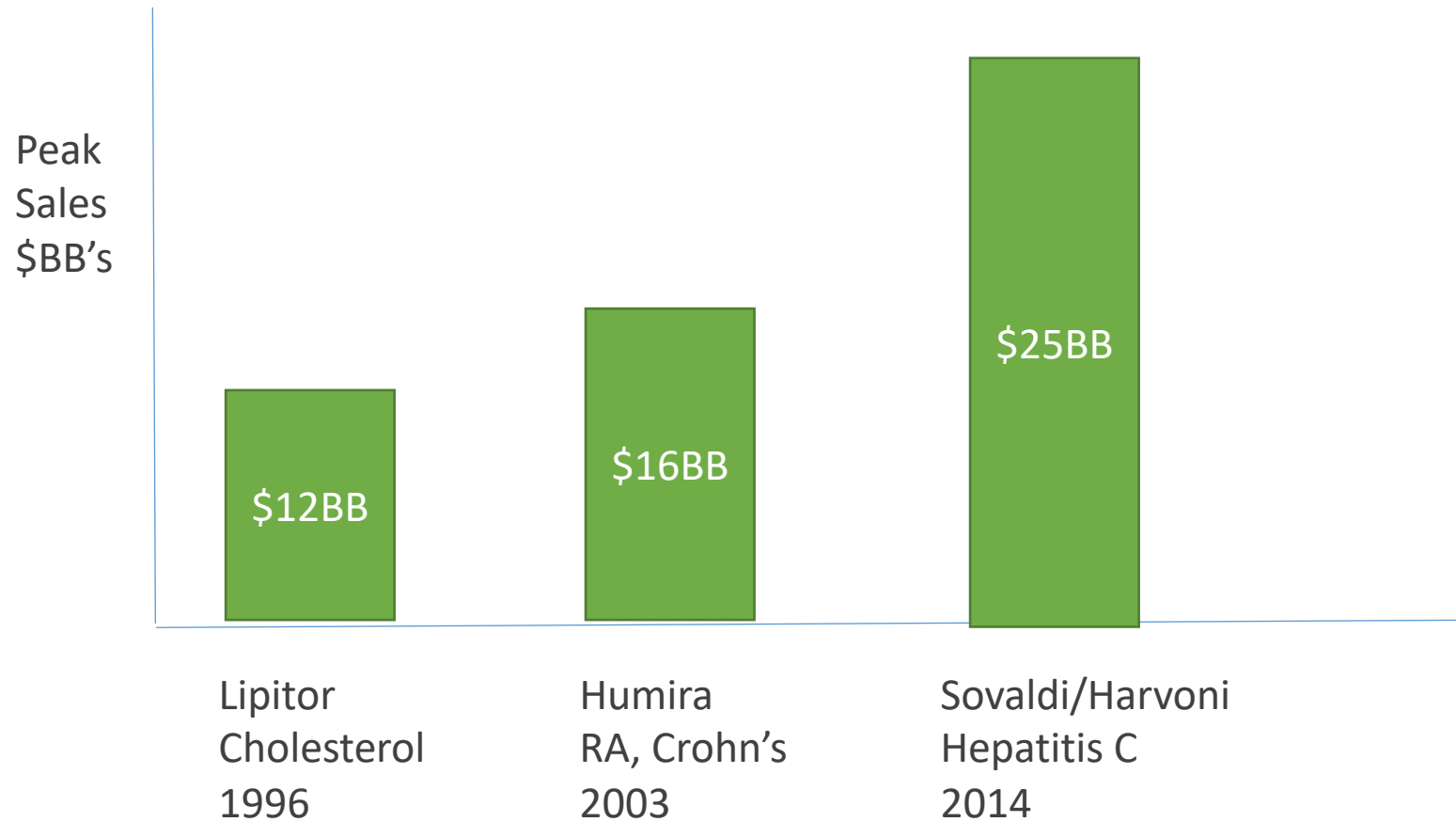
In vivo impairment of recognition memory by A β oligomers, not monomers and not fibrils¹⁰



Degree of selectivity for the correct (toxic) form of amyloid explains past clinical results



The three largest products in industry history were not first in class, but “best in class” – the inventors identified improvements to existing drugs



ProMIS following the “best in class” playbook:

- Took advantage of “proof of biology” developed by earlier products:
- Used ProMIS proprietary science platform to design an improved product, which may yield superior clinical results

PMN310: an anti-A β -oligomer antibody with strong potential to demonstrate best-in-class characteristics in Alzheimer's treatment

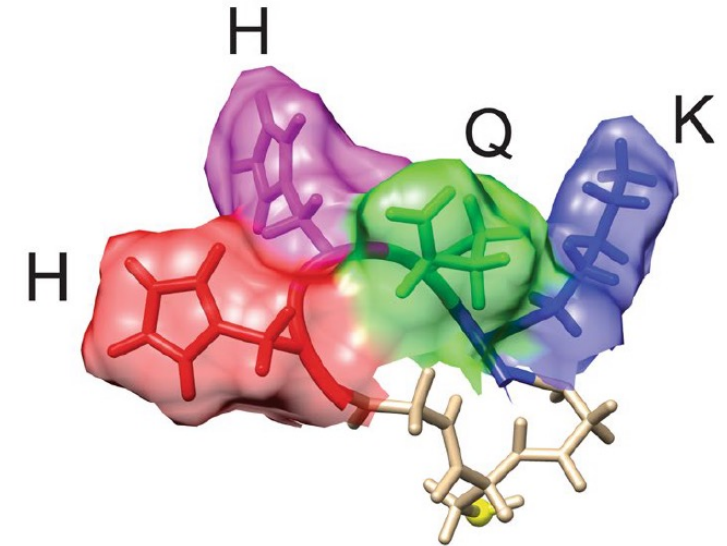
Aducanumab (Biogen), BAN2401 (Eisai), donanemab (Lilly) selective for all aggregated amyloid – plaque and oligomers

- Aducanumab (phase 3), BAN2401 (phase 2) and donanemab (phase 2) showed modest but meaningful efficacy in reducing cognitive worsening
- **Aducanumab approved by FDA – June 7, 2021**
- None bind monomer (the physiologic amyloid species)
- All bind amyloid plaque → ARIA-E (brain swelling)

PMN310 is a next-generation, best-in-class anti-amyloid therapy

- Highly selective for only toxic oligomers
- Dose expected not to be limited by off-target binding or side effects
 - Does not bind monomer
 - Does not bind plaque → **likely no ARIA-E side effect**
- All dosed PMN310 will be focused on neutralizing toxic oligomers → **potentially greater clinical efficacy**

PMN310 will utilize blood-based biomarkers in initial trials, enabling rapid timeframe and limited capital needs to value-creating clinical data



A β amino acids 13-16 (HHQK) form a unique, A β oligomer specific conformational epitope targeted by PMN310

There are three forms of amyloid, PMN310 is differentiated by selective binding of the toxic form (oligomers)

Bapineuzumab (Pfizer)

- Phase 2 failure
- Phase 3 failure
- ARIA-E side effect

Solanezumab (Eli Lilly)

- Phase 2 failure
- Phase 3 failure

Aducanumab (Biogen)

- Phase 2 & 3 success
- ARIA-E side effect

PMN310

- Selective binding to oligomers
- > Expected improvement in efficacy & safety

MONOMERS

- binding wastes therapeutic ammunition

FIBRILS (Plaque)

- binding wastes therapeutic ammunition
- contributes to ARIA-E side effect

OLIGOMERS*

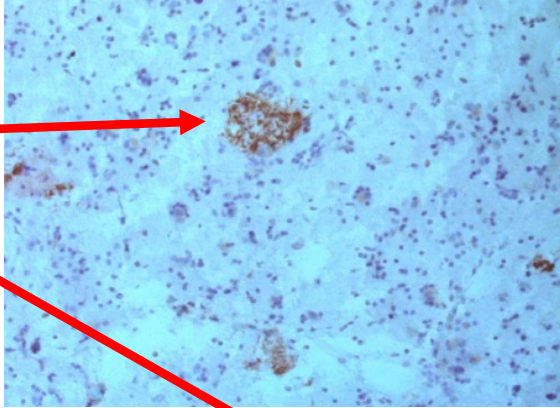
- the right target



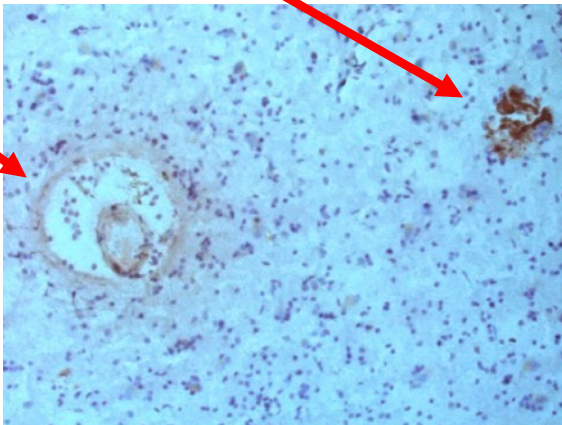
ARIA-E associated with aducanumab, BAN2401 & bapineuzumab; PMN310 lack of binding to A β plaque strongly suggests a *potential safety advantage - no ARIA-E*

Aducanumab

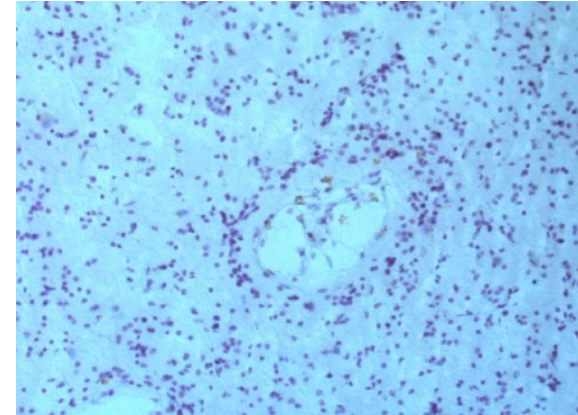
Plaque
binding



Vascular
deposit
binding

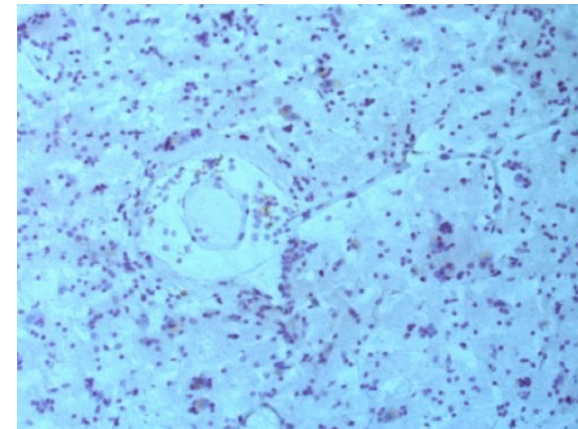


PMN310



No binding to plaque
or vascular deposits

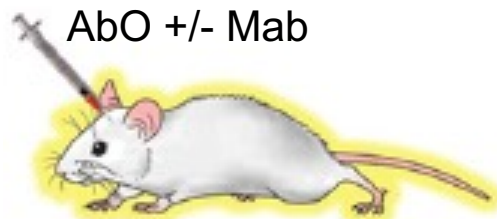
Most likely no ARIA-E*



Administration of PMN310 to mice prevents loss of short-term memory formation caused by toxic oligomers, by saving mouse neurons

THE EXPERIMENT

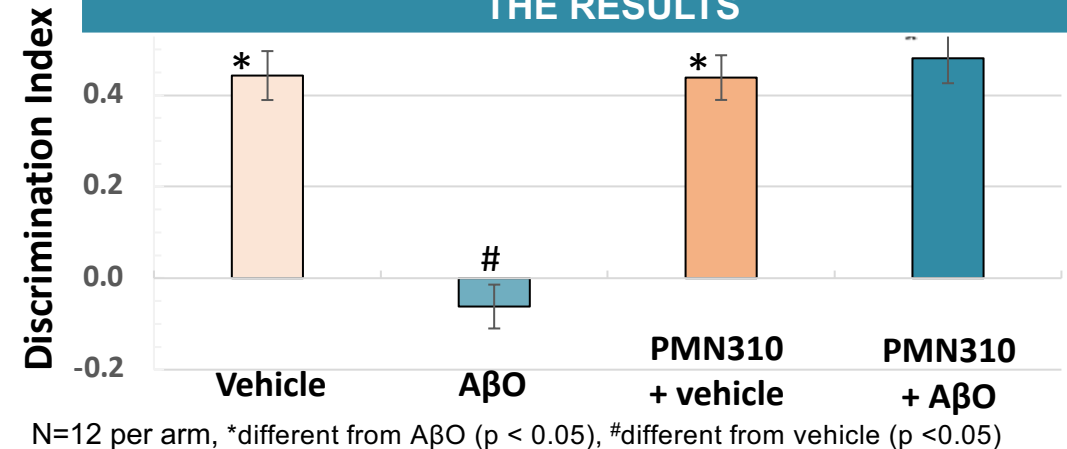
- Mice are tested for discriminating objects after brain injection of:
 - Buffer (vehicle) - normal response
 - Toxic A β oligomer
 - PMN310 and buffer (vehicle)
 - PMN310 and A β Oligomer



7 days



THE RESULTS

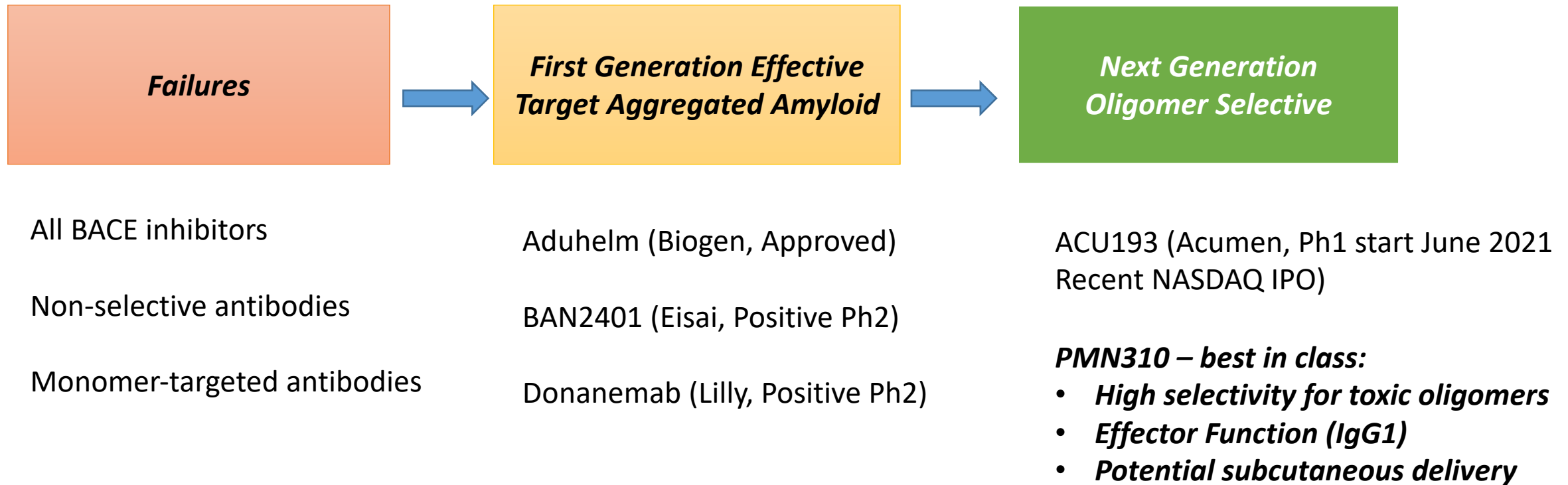


Novel Object Recognition Assay

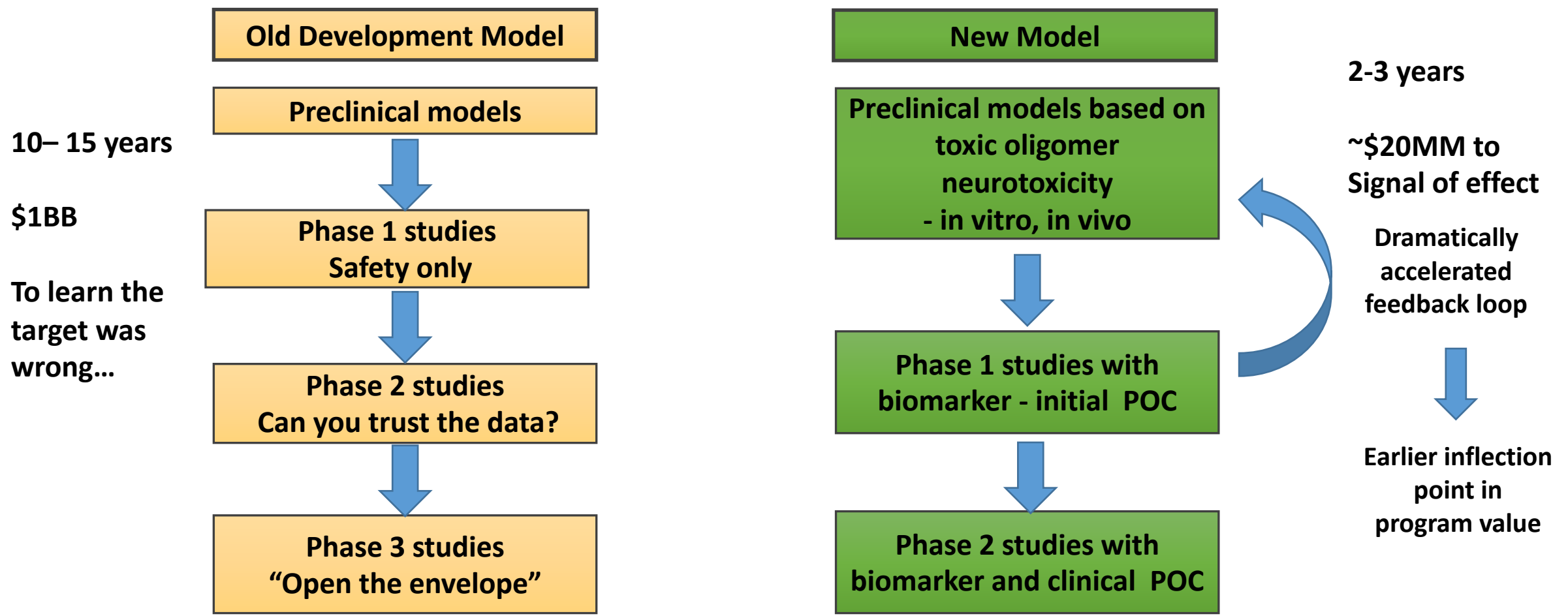
- Control mice remember a familiar object when re-exposed to it and spend more time exploring a new object
- Oligomer-injected mice lose the ability to discriminate between known and novel objects and spend equivalent amounts of time exploring both

PMN310 has the potential to be the “best of the next generation” therapy in Alzheimer’s

Amyloid beta-targeted therapies




A new development paradigm for Alzheimer’s and other neurodegenerative diseases using biomarkers will dramatically improve cost, risk....and time to success



Alzheimer's drug R&D: we're in the midst of a biomarker revolution!

Early Alzheimer's Disease: Developing Drugs

Guidance



HHS Public Access

Author manuscript
Alzheimers Dement. Author manuscript; available in PMC 2018 May 18.

Published in final edited form as:
Alzheimers Dement. 2018 April ; 14(4): 535–562. doi:10.1016/j.jalz.2017.12.001

NIA-AA Research Framework: Toward a New Definition of Alzheimer's disease

Clifford R. Jack Jr.^{a,*}, David A. Bennett^b, Kaj Blenn^c, Samantha Budd Haeberlein^f, David M. Holtzman^g, Karlawish^j, Enchi Liu^k, Jose Luis Molinuevo^l, Katherine P. Rankin^o, Christopher C. Rowe^p, Ronald M. van Dyck^q, Snyder^d, and Reisa Sperling^s

U.S. Department of Health & Human Services
Center for Biologics Evaluation and Research (CBER)

February 2018
Clinical/Medical



Almost there

The unexpected way we might one day diagnose Alzheimer's

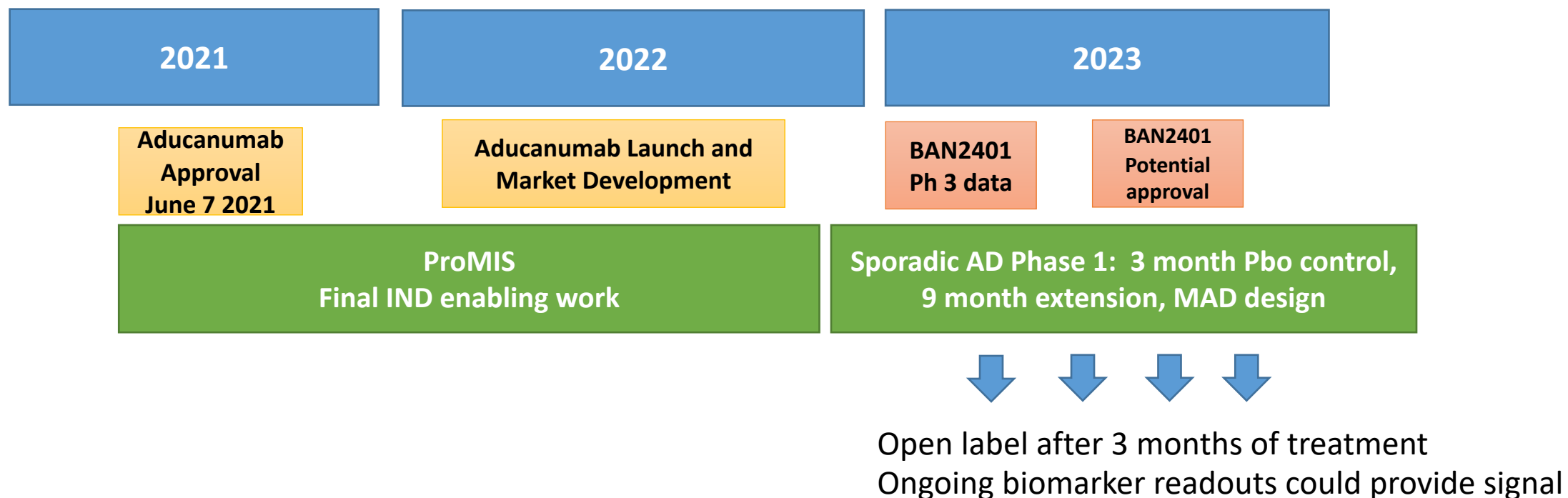
By Bill Gates | April 2, 2019

Jeff Bezos teams with Bill Gates to fund new and better ways to diagnose Alzheimer's

by [john carroll](#) — on April 4, 2019 10:10 AM EDT

PMN310: potential for value-creating clinical data in the near term

- likely positive market developments could amplify PMN value



- Recent advances in blood-based biomarkers may allow ProMIS to detect an objective treatment signal as early as Phase 1, potentially providing rapid & cost-effective proof-of-concept

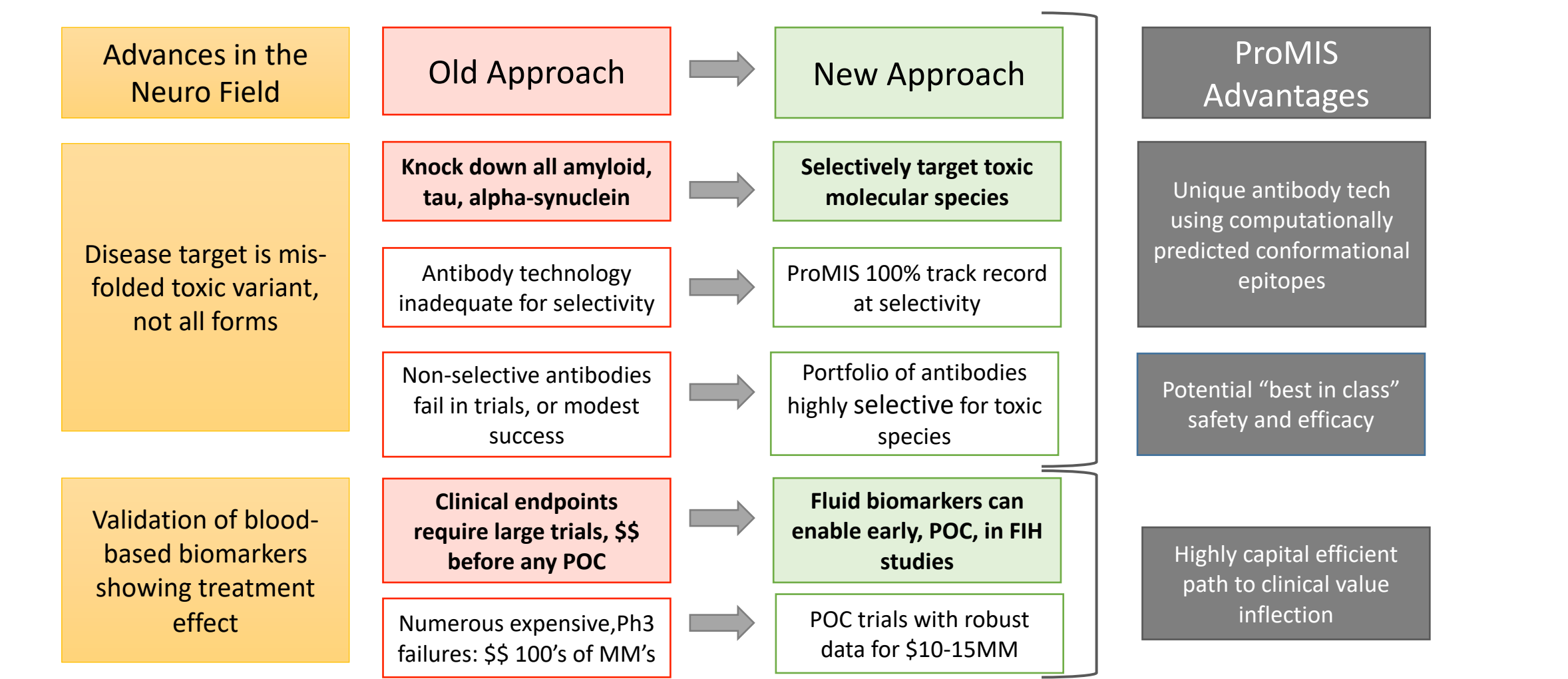
ProMIS: a broad differentiated portfolio; a unique technology platform

- Potential "best of the next generation" for all of neurodegenerative disease

Misfolded protein target	Lead indication	Other Indications	Status
Amyloid beta	Alzheimer's		IND enabling work ongoing
TDP-43	ALS	FTD, LATE	Lead antibodies
Alpha synuclein	Multiple System Atrophy	Parkinson's, LBD	Lead antibodies
tau	Alzheimer's	PSP, other tauopathies	Lead selection
SOD1	ALS		Lead antibodies
RACK1	ALS	HD, cancers	Immunizations
Ataxin2	ALS		Computational modeling
Disc1	ALS	Schizophrenia	Computational modeling
Amylin	T2Diabetes		Computational modeling

DLB: Dementia with Lewy bodies, FTD: Frontotemporal dementia, LATE: Limbic-predominant age-related TDP-43 encephalopathy, ALS: Amyotrophic lateral sclerosis, PSP: Progressive supranuclear palsy, AD: Alzheimer's disease, HD: Huntington's disease

Advances in the neurodegeneration field + ProMIS unique capabilities = value creation potential



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Thank You

Please feel free to contact us with any additional questions.

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Appendices

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- | | |
|---|---|
| 1) ProMIS unique antibody design platform | <ul style="list-style-type: none">- Computational modeling approaches combining Physics and Biology- 100% track record at achieving target molecular species selectivity |
| 2) PMN310 – lead program in Alzheimer’s | <ul style="list-style-type: none">- Comparator data support “best in class” potential- In vivo data: blocks cognitive deficit caused by amyloid-beta oligomers- Competitor clinical data support science – oligomer is target |
| 3) Alpha Synuclein program | <ul style="list-style-type: none">- Several antibody candidates with selectivity and functional benefit- Illustrates ProMIS’ competitive advantage in achieving selectivity |
| 4) TDP-43 program | <ul style="list-style-type: none">- Multiple antibodies with target selectivity- Functional benefit both as extracellular antibody and intracellular intrabody |

ProMIS Platform

The ProMIS platform generates antibodies selective for the misfolded toxic forms of pathogenic proteins

Identification of epitopes selectively exposed on toxic misfolded form of the target protein using predictive computational algorithm



Immunization with disease-associated epitope and screening of monoclonal antibodies with desired binding profile and protective activity

Neutralize the toxic misfolded form

Don't interfere with the normal form, critical for brain health



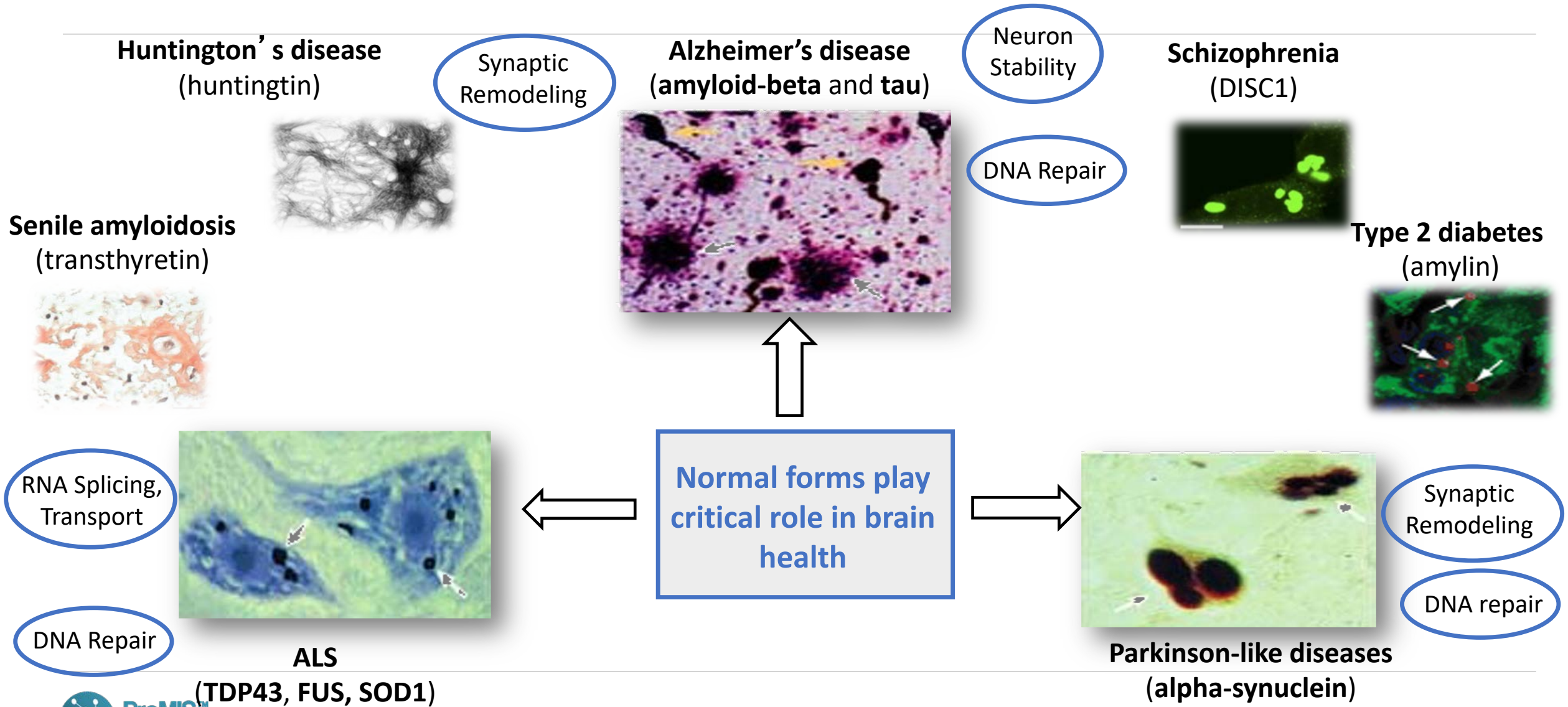
Successful track record thus far in generating antibodies selective for pathogenic forms of amyloid-beta, tau, alpha-synuclein, TDP-43, SOD1

Therapies must be selective for the mis-folded form – ProMIS has a unique advantage

Protein	Role of physiologic monomer	Disease Caused by Mis-Folded Form
Amyloid Beta	Synaptic remodeling	Alzheimer's
Alpha Synuclein	DNA repair	Parkinson's Lewy Body Dementia MSA
TDP43	RNA transcription	ALS FTD

- ProMIS has a portfolio of antibodies with high selectivity for the mis-folded form of each of these proteins
- Differentiated from first generation and competitor products
- ProMIS unique capability

The same proteins perform essential cellular functions when in their normal, physiological form



Collective Coordinates: Algorithm to predict misfolding-specific epitopes

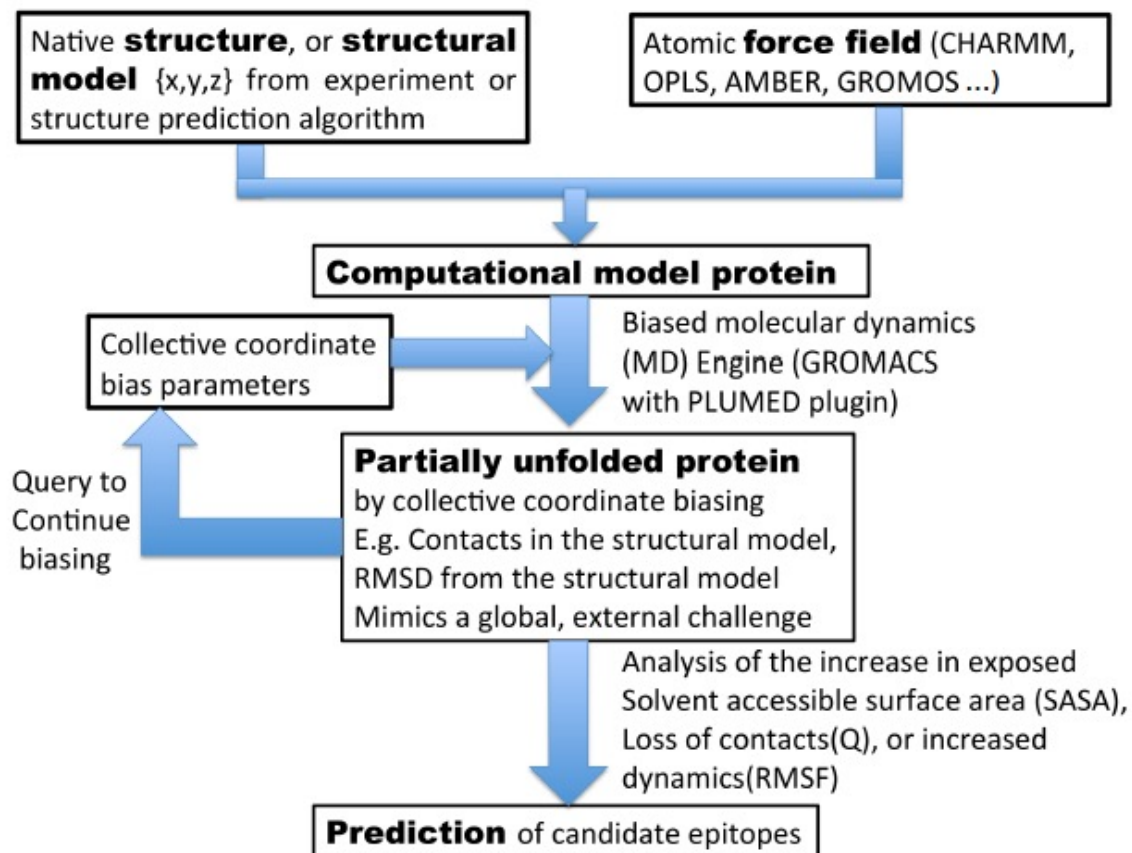
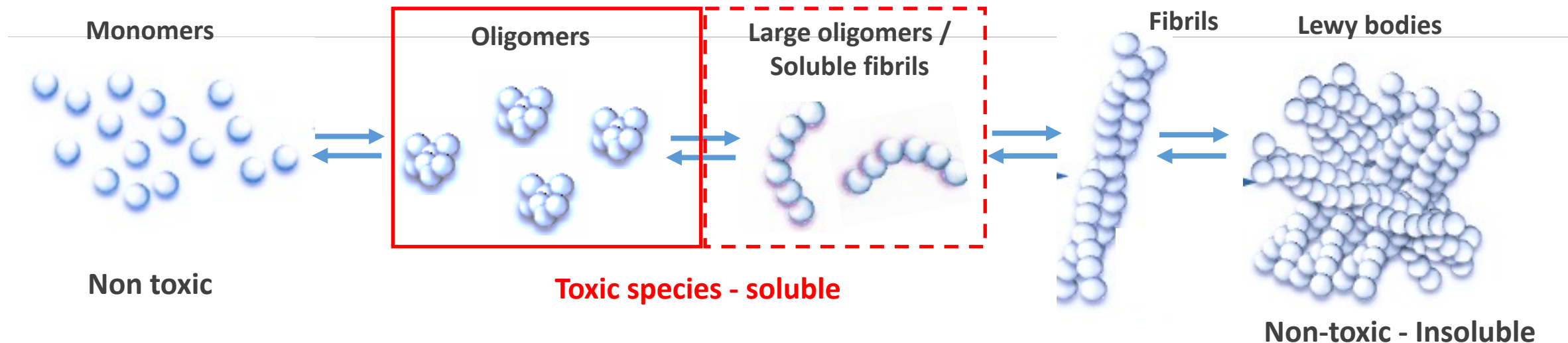


Figure 1: Flowchart of the steps in the prediction algorithm
J Phys Chem 2018

- General principles:
 - Weakly stable regions in the native fold may constitute target epitopes that are preferentially exposed in misfolded species.
 - These misfolding-specific epitopes can be distinguished from the same regions in the native fold by their conformational properties
 - By properly scaffolding the epitopes, antibodies may be raised to selectively bind to misfolded species
- Molecular dynamics simulations partially unfold a protein or fibril in order to predict misfolding-specific epitopes generated by regions that are most likely to locally unfold in response to global stress
- Several metrics used to measure local disorder: solvent exposed surface area, native contacts, root mean squared fluctuations
- Patent filing: Plotkin SS. Systems and methods for predicting misfolded protein epitopes by collective coordinate biasing. International application No: PCT/CA2016/051306. 2016
- Manuscript on Collective Coordinates applied to SOD1: Peng X, Cashman NR & Plotkin SS, J Phys Chem, 2018

ProMIS discovery platform applied to alpha-synuclein: Selectivity advantage over other antibodies

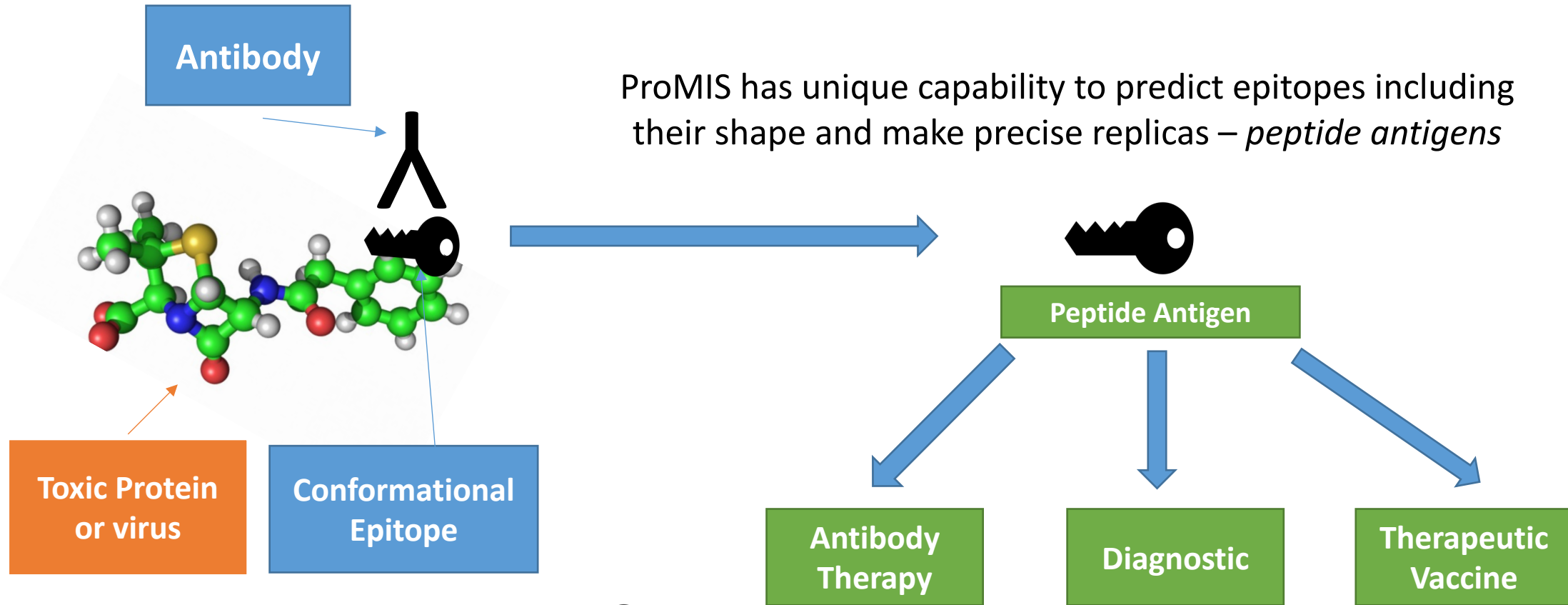


- Generation of monoclonal antibodies through immunization with peptide/monomer or synthetic oligomers gives rise to antibodies that cross-react to varying degrees with non-toxic monomers and/or insoluble fibrils
 - Binding to non-toxic species is likely to reduce the effective dose and diminish efficacy
 - Binding to monomers/physiological tetramers may interfere with normal alpha-synuclein function
 - Interaction with circulating alpha-synuclein may cause infusion reactions (observed with Prothena antibody at 60 mg/kg)
- The ProMIS platform allows for identification of conformational epitopes predicted to be exposed solely on toxic species -> Immunization with these epitopes allows for generation of antibodies selective for pathogenic α -syn
 - The selectivity and activity of the antibodies is validated empirically in vitro and in vivo
 - Reactivity with patient-derived material (native alpha-syn species) is an essential component of the validation process

FAQs

- What is unique about ProMIS technology platform?
 - ProMIS combines physics and biology to predict portions of mis-folded proteins exposed as epitopes; it requires physics to predict **shape**, and shape is the only difference between normal and toxic mis-folded proteins
- If the science is clear about toxic, mis-folded proteins, why are large companies not targeting selectivity?
 - Many large pharma programs failing/working modestly now were designed before the science was as clear
 - Without ProMIS unique approach to antibody design it is impossible to systematically create selective antibodies
- Will others eventually catch up, and compete?
 - Eventually, advances in computing power (like IBM quantum computing) will enable others to predict shape and create selective antibodies
 - ProMIS is building a large IP estate in the mis-folded protein area; there are only so many conformational epitopes exposed
 - ProMIS identifies a few for every target protein, creates antibodies, and files for IP

ProMIS unique technology platform: computation of conformational epitopes exposed only on mis-folded proteins, led to PMN310, and a therapeutic vaccine

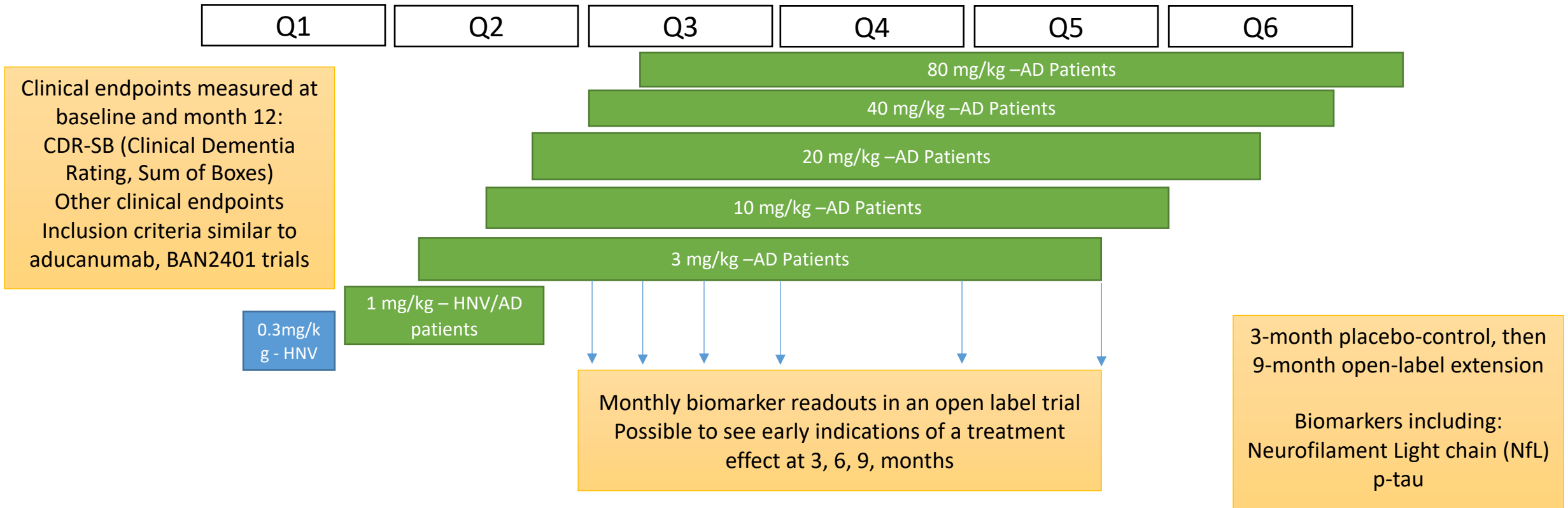


Antibodies bind to targets called *EPITOPES* (🔑) which have a specific conformation or shape

ProMIS Template Development Program- Summary

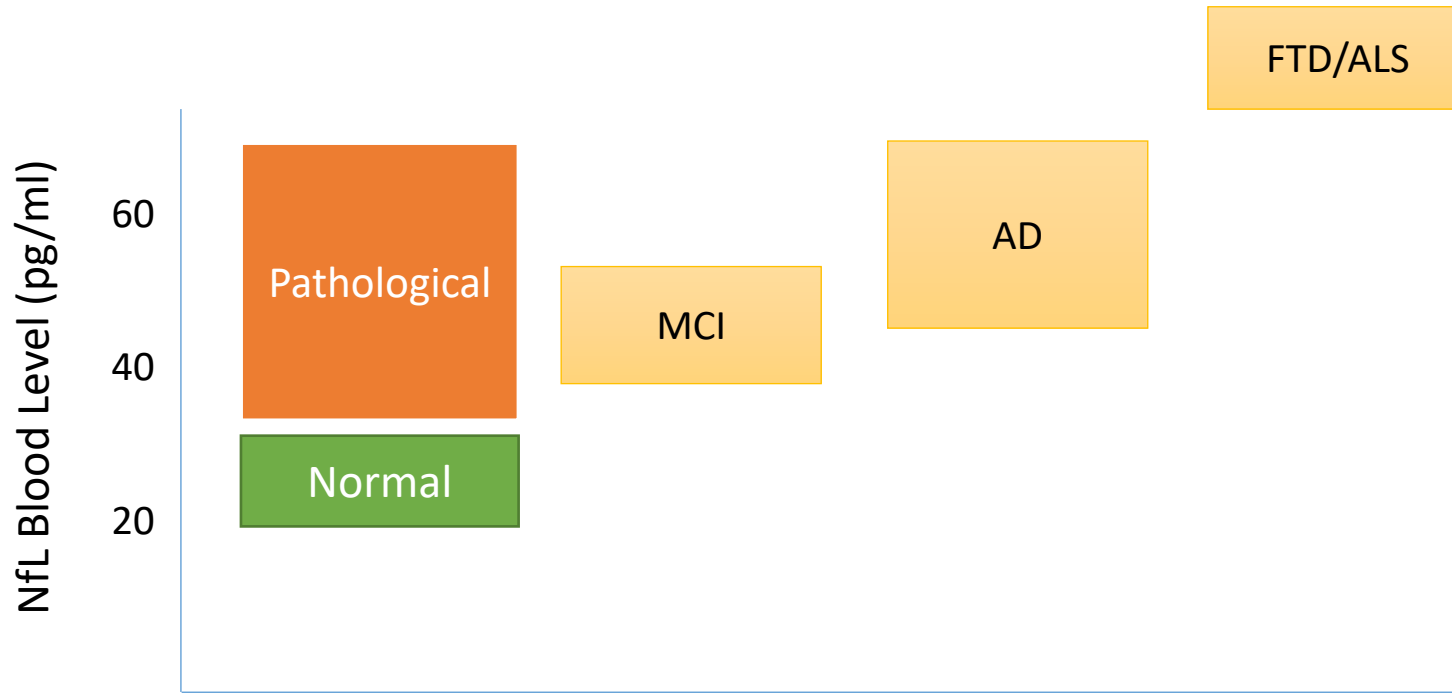
- First step is computational predictions of epitopes, creating peptide antigens, immunizing mice and/or rabbits to create initial antibody candidates
- Lead candidates selected through assays to screen for selectivity, including selective binding to patient bio-samples vs healthy controls, in vitro assays
- Lead candidates humanized (Vendor – Abzena), CHO cell line (Selexis), GMP manufacturing (KBI), and GLP tox conducted to enable IND
- All programs will treat patients in the first in human study (after healthy normal in low dose arms)
- All programs targeting disease populations with elevated biomarkers (like NfL – neurofilament light, an indicator of the rate of neuronal death; or p-tau, phosphorylated tau, a measure of the toxic tau created by toxic amyloid-beta oligomers), enabling a dose-dependent treatment response to be seen in monthly blood draws, quantitative assays, that could correlate with clinical endpoints
- Extremely capital efficient relative to past programs in neurodegenerative disease. ProMIS estimates:
 - Discovery - <6 months, <\$100k
 - IND enabling work: \$8MM - \$10MM
 - Clinical program to initial POC: \$10MM - \$15MM

Phase 1 trial design concept: biomarkers in addition to clinical endpoints can give a signal suggesting therapeutic benefit early in clinical development



HNV = healthy normal volunteers

Neurofilament light (NfL) is a measure of the rate of neuronal death.....



From JAMA Neurology, April 22 2019

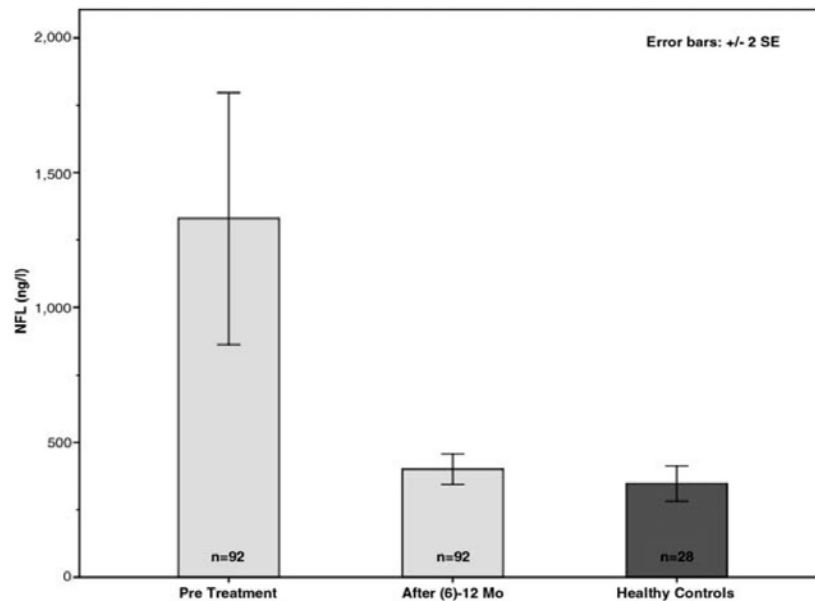
“ The findings suggest that plasma NfL can be used as a noninvasive biomarker associated with neurodegeneration in patients with AD and may be useful to monitor effects in trials of disease-modifying drugs”

Mattsson, et al, JAMA Neurology

Sources: from AAIC 2018 ABBVIE, Wash U, U Sorbonne;
Rohrer et al, 2016 AAN; Mattson et al, JAMA Neurology 2017

NfL has proven to be a valuable pharmacodynamic marker in multiple sclerosis and SMA (spinal muscular atrophy)

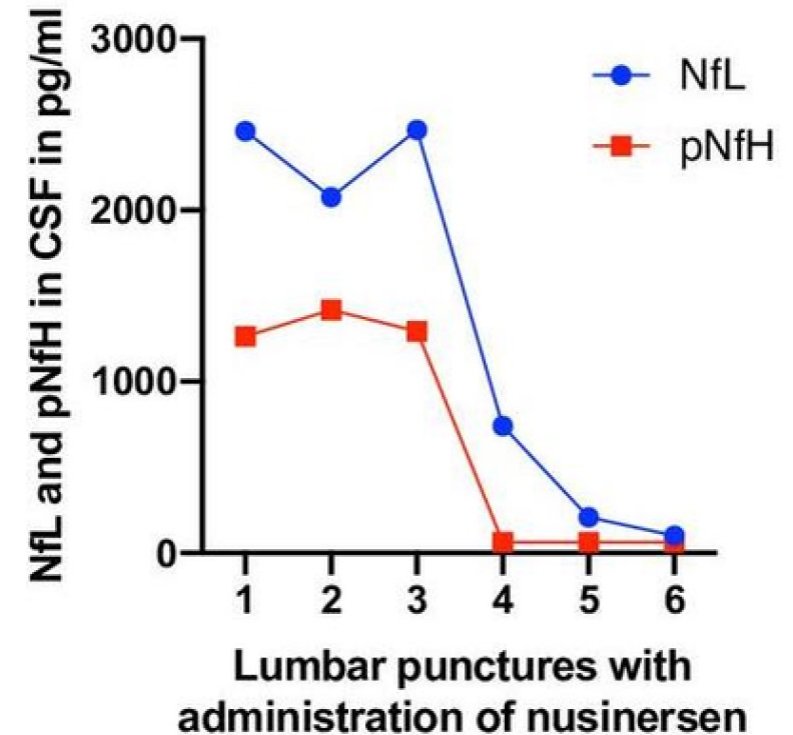
Tysabri in multiple sclerosis



Neurofilament light in cerebrospinal fluid following monthly administration of natalizumab (300 mg) in 92 patients with multiple sclerosis

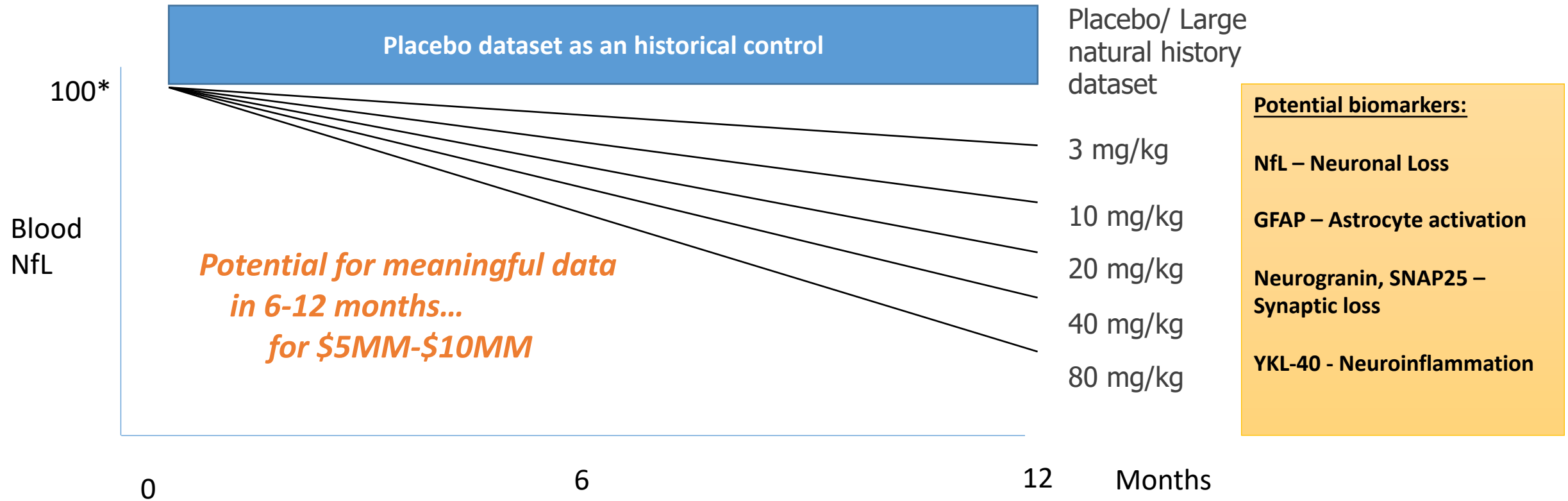
Gunnarsson, et al Annals of Neuro 2011

Spinraza in spinal muscular atrophy



Winter B. et al, J Neurol Neurosurg Psychiatry 2019

Hypothetical example – Phase 1 biomarker readout could show a disease-modifying treatment effect in a dose-dependent fashion *at a cost of ~\$10MM*



* 100 = patient baseline value

Dose escalation design: 3-month placebo-control, then 9-month open-label extension

ProMIS Alzheimer's Lead Program: PMN310

Alzheimer's Disease: the epidemic of this century

**MORE THAN
5 MILLION**
AMERICANS ARE LIVING WITH ALZHEIMER'S

“Will Bankrupt Medicare
if Therapy is not developed”

Direct and Indirect Costs
Today in the US \$500BB....
.... and the number
is tripling by 2030....

ALZHEIMER'S DISEASE IS THE
**6TH LEADING
CAUSE OF
DEATH**
IN THE UNITED STATES

**EVERY
66 SECONDS**

SOMEONE IN THE UNITED STATES
DEVELOPS THE DISEASE

IT
KILLS
MORE
THAN

**BREAST AND
PROSTATE CANCER
COMBINED**

Alzheimer's is at an inflection point like immuno-oncology ten years ago

"Neuroscience has the potential to be in the '20s what oncology has been in the last decade."

– Bill Anderson, CEO of Roche Pharmaceuticals (JPM 2020)

Immuno-Oncology

~2000's

- Numerous failures
- Skepticism, lack of capital

2010 -inflection

- First cancer vaccine approved (sipuleucel-T)
- First positive clinical data for PD-1 (nivolumab)

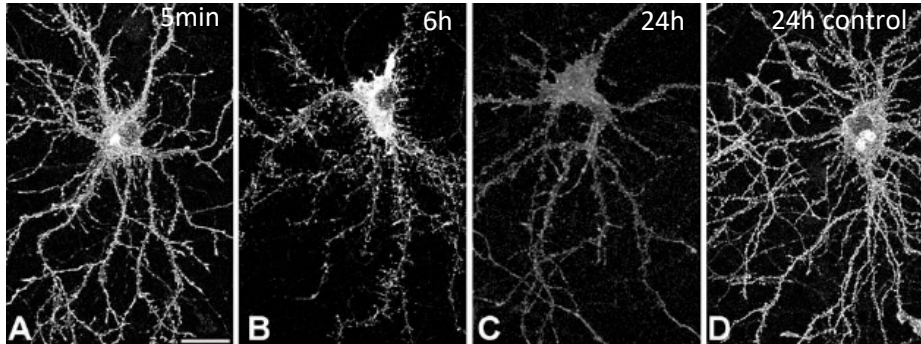
2017/2018 - validation

- Gilead acquires Kite (2009 start up) \$11.9BB
- Celgene acquires Juno (2013 start up) \$9BB

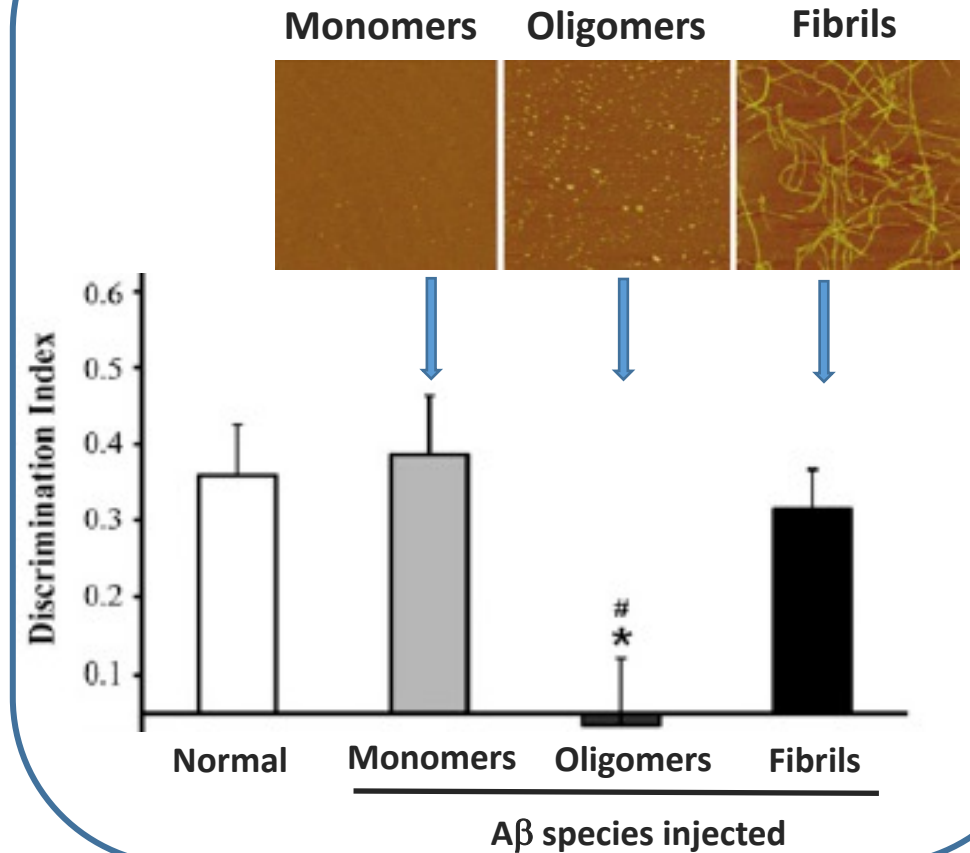
Alzheimer's disease: soluble toxic A β oligomers – not plaque or monomers – are the most neuropathogenic A β species

- Synapse abnormalities and memory impairment correlate poorly with plaque burden in human and mouse AD^{1,2}
- A β monomers and A β insoluble fibrils (plaque) have little or no demonstrable toxicity in vitro or in vivo³⁻⁵
- Soluble A β oligomers show the highest degree of neurotoxicity⁶
 - Toxicity in primary neuron cultures and brain slices^{3,5,7-9}
 - Induction of cognitive impairment in rodents^{3,4,10}

Synaptotoxicity of human A β oligomers on hippocampal neurons in vitro⁷



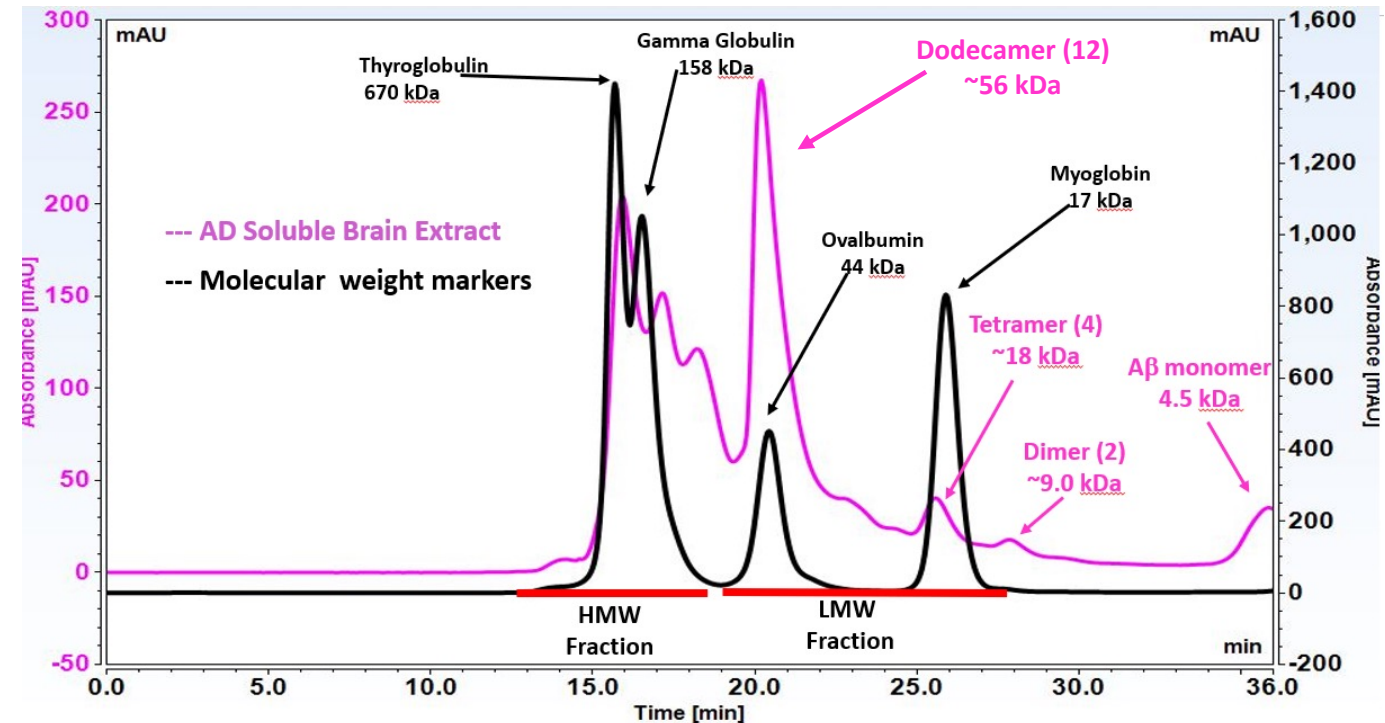
In vivo impairment of recognition memory by A β oligomers, not monomers and not fibrils¹⁰



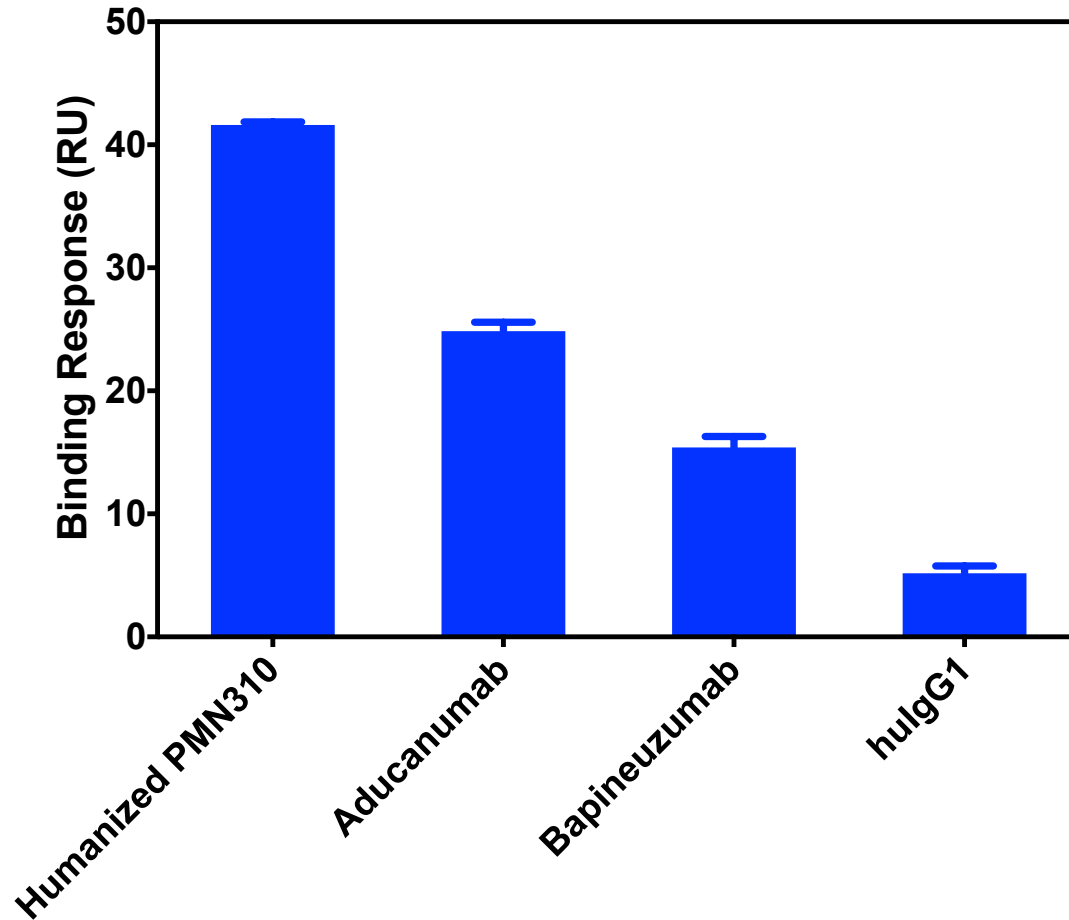
Not all oligomers are equivalent soluble low molecular weight toxic oligomers drive neurotoxicity and disease progression

- AD brain homogenates contain heterogeneous A β aggregates dominated by high molecular weight species including protofibrils (>70kDa, or ~15+ "monomers")^{1,2}
- Neurotoxic activity resides primarily in the **LMW (low molecular weight) fraction** (<70 kDa)¹
- A subset of the LMW oligomers are the most toxic oligomers (**dimer, tetramer, dodecamer**)^{1,3-7}

Size exclusion chromatography of AD brain extract



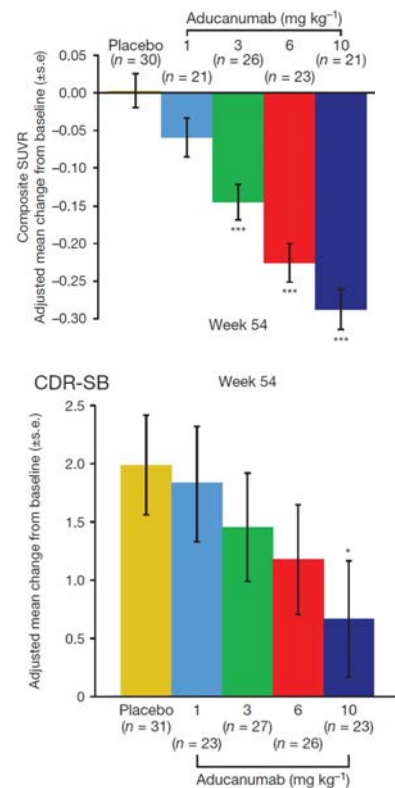
PMN310 shows superior binding to toxic oligomers from human AD brains vs other antibodies directed against amyloid-beta



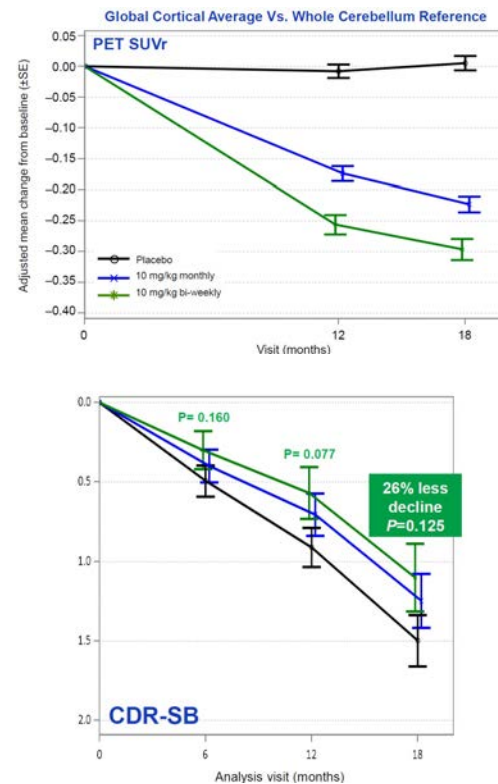
- Binding of antibodies to the toxic oligomer-enriched LMW fraction of soluble human AD brain extract was evaluated by surface plasmon resonance (SPR)
- Results representative of over 10 SPR runs with extracts from 11 different AD brains
- hulgG1 = Background control

Four positive trials...partially selective antibodies...higher dose over long duration leads to better outcomes, but all limited by brain swelling (ARIA-E)

Biogen Ph1b “PRIME” December 2014

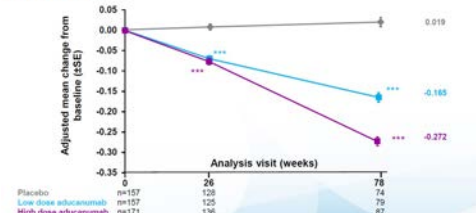


Eisai BAN2401 Phase 2 July 2018

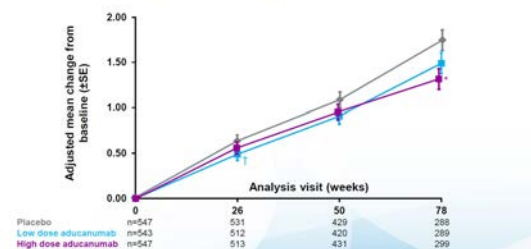


Biogen pivotal EMERGE Dec 2019*

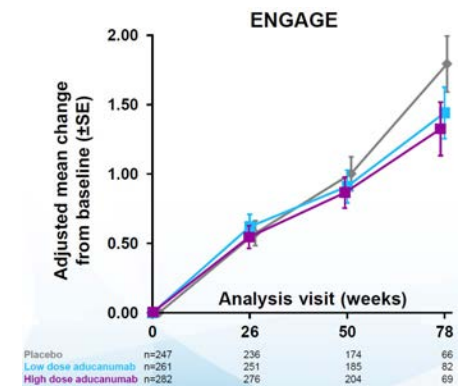
EMERGE: Longitudinal change from baseline in amyloid PET SUVR



EMERGE: Longitudinal change from baseline in CDR-SB



Biogen pivotal ENGAGE Dec 2019**



The goal in Alzheimer's is prevention – a large and growing market

Symptomatic
AD
~6MM

30%- 50% of
population over 60
with pathology, pre-
symptomatic
~30MM- 40MM

Population over 60 at risk –
86MM in US and Canada

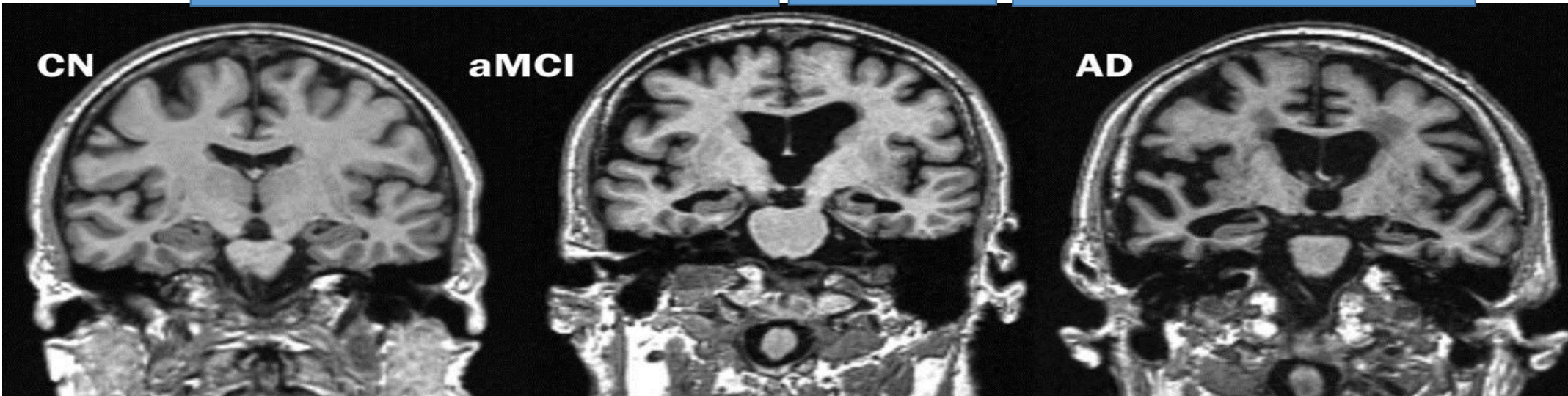
20+ years to develop symptoms, from onset of pathology (neuronal death)

Billions of neurons were killed leading to the Alzheimer's brain on the right.....toxic amyloid is a root cause

15 -20 years pre-symptomatic

5 years MCI

10 years Alzheimer's



Icelandic Genetic mutation:
No amyloid
No AD

Down Syndrome:
Excess amyloid
Early, High % AD

Healthy Brain

Brain with
Mild Cognitive Impairment

Brain with AD¹

Toxic, mis-folded amyloid plays a role throughout disease

Toxic, mis-folded amyloid hyper-phosphorylates tau

*Disease detection and progression measurable in blood – NfL (neuronal death),
p-tau – (phosphorylated tau)*

¹ Reviewed in Bloom 2014, JAMA Neurol

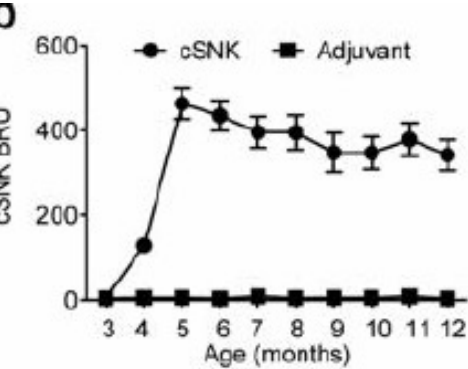
ProMIS therapeutic vaccine for Alzheimer's disease

- Large patient population with symptomatic AD in need of effective treatment ($n \geq 5M$, USA) → **Therapeutic vaccination**
- Ever greater number of individuals in the pre-symptomatic phase of the disease (potentially 40% of the population over 65, ~50M) –> **Prophylactic preventative vaccine**
- Vaccine approach will benefit from recent progress in the development of blood-based biomarkers of neurodegeneration to diagnose AD and identify individuals at risk of developing disease
- **ProMIS discovery platform being applied to devise a safe and effective vaccine to induce a specific immune response against toxic A β oligomers**
 - Identified a set of 6 conformational peptide epitopes selectively exposed on toxic A β oligomers
 - All 6 peptide epitopes shown to be capable of inducing selective and protective antibodies against toxic A β oligomers
 - Successful proof of concept vaccination study conducted with one of the peptides (cSNK) in a mouse model of AD (APP/PS1 mice): Neuronal protection and improvement in cognitive deficits²
 - Concept: multivalent vaccine with some or all 6 A β peptides. Tau peptides recently identified could also be included.
- **Immediate aims: Construct and test multivalent A β vaccine for ability to induce a protective antibody response**

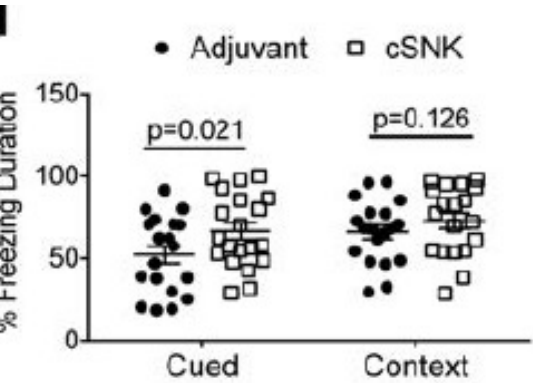
Benefit of vaccination with ProMIS A β oligomer (A β O) epitope

Mice vaccinated with conformational A β oligomer epitope (cSNK) coupled to KLH

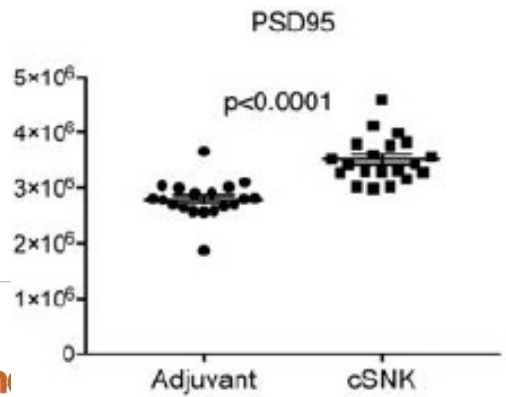
Robust, sustained antibody response (SPR)



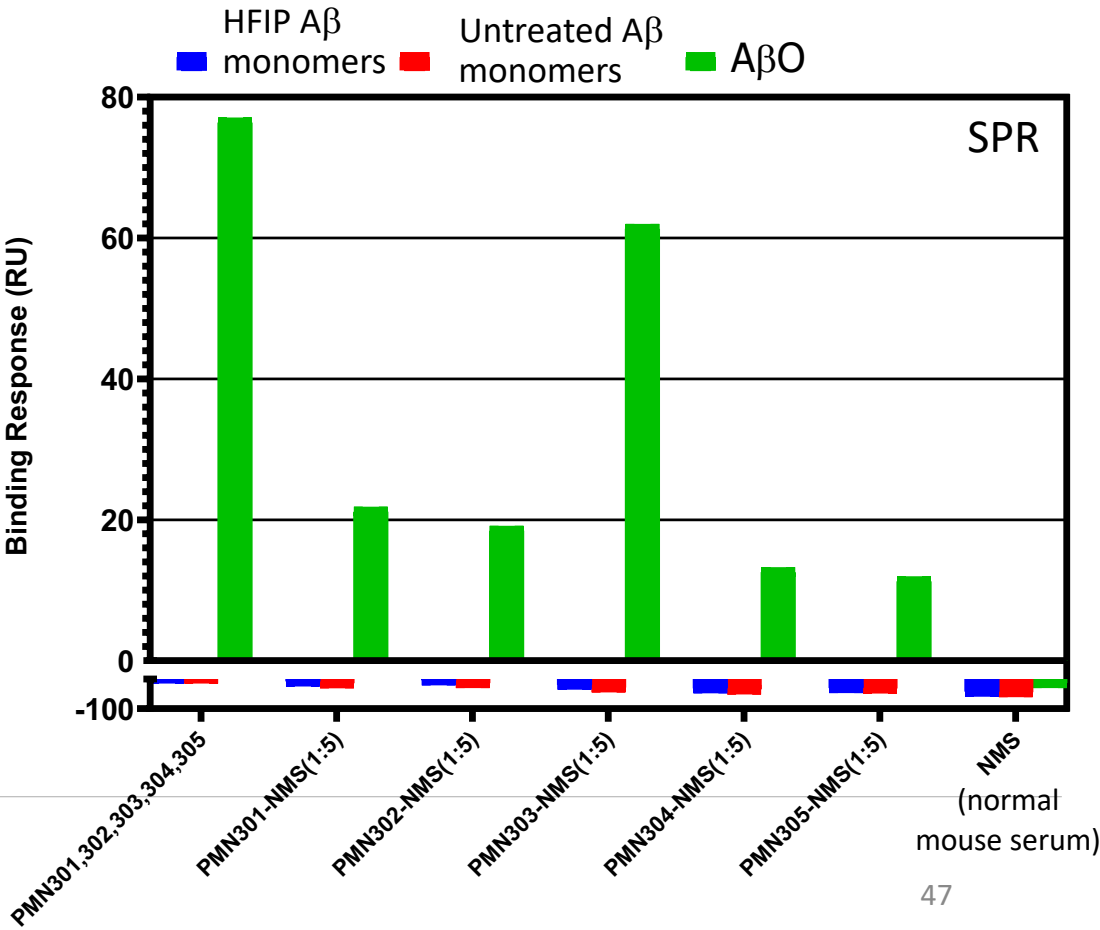
Improvement in behavioral deficits of APP/PS1 mice



Protection against synaptic damage



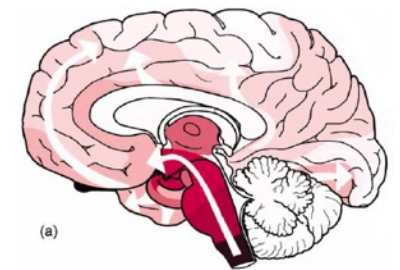
Biological support for multivalent vaccine approach: Greater binding with mixture of sera from mice immunized with different A β O epitopes vs immune serum against individual peptides



Alpha-Synuclein Program

Alpha-synuclein is the major driver of Parkinson's disease and other synucleinopathies

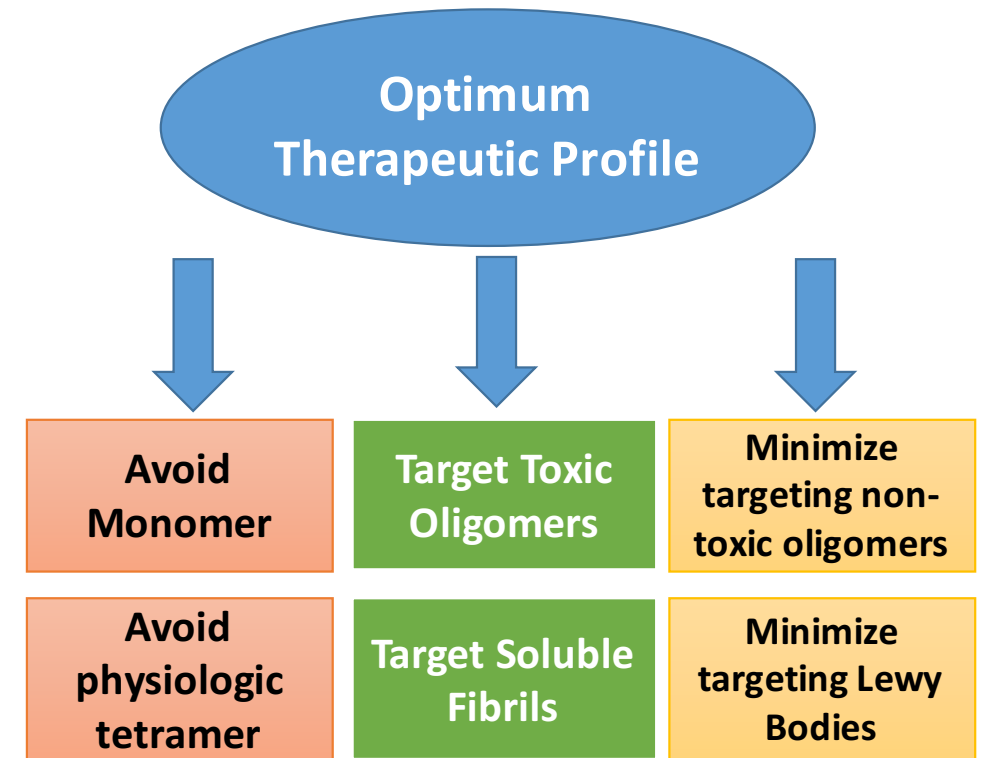
- Synucleinopathies: Parkinson's disease (PD) is a progressive neurodegenerative disorder characterized by loss of dopaminergic neurons located in the midbrain and the presence of intraneuronal inclusions (Lewy bodies/Lewy neurites) consisting mainly of aggregates of α -synuclein (α -syn). Accumulation of insoluble α -syn fibrils in the brain is also observed in dementia with Lewy bodies (LBD) and multiple system atrophy (MSA).
- Genetic mutations (A53T, A30P, E46K, H50Q, G51D) and duplications/triplications of the α -syn gene cause inherited familial PD¹
- Polymorphisms in the α -syn gene and its promoter are associated with an increased risk of sporadic PD¹
- Mice injected with aggregated α -syn from diseased patient brains develop synucleinopathy-like pathology and symptoms²
- Aggregated α -syn propagates from cell to cell in a prion-like manner in cell culture³ and in animals^{4,5} mimicking disease progression in patients



Braak, H et al, Neurobiol of Aging, 2003

Therapeutic imperative: selectively target only the toxic α -synuclein aggregates

- **Alpha-synuclein exists in different forms including normal, physiologically important conformations and toxic forms:**
 - Monomers play an important role in the regulation of synaptic vesicle trafficking and release as well as neuronal survival¹
 - Physiological α -syn tetramers inhibit aggregation and must be preserved for healthy α -syn homeostasis²⁻⁴
 - Lewy bodies consisting of insoluble α -syn are a marker of disease but unlikely to have direct neurotoxicity⁵
 - Recent evidence indicates that α -syn toxicity resides primarily with soluble oligomers^{6,7}
 - Progression of disease is likely mediated by oligomers and small soluble fibrils of α -syn that have been shown to propagate from cell to cell in a prion-like manner in vitro⁸ and in vivo⁹
- **Maximal efficacy and safety is expected to require selectivity for the toxic forms of α -syn, oligomers and/or small soluble fibrils, while avoiding physiologic forms of α -syn**



ProMIS' technology platform has created antibodies with greater selectivity than other α -synuclein-directed antibodies

Target Properties	PMN Antibodies	Prothena/ Roche	BioArctic/ ABBVIE	Neurimmune/ Biogen
No binding to monomers	Yes	X	+/-	X
No binding to physiological tetramers	Yes	X	+/-	X
Binding to oligomers/small soluble fibrils	Yes	Yes	Yes	Yes
Binding to native toxic α -syn in PD/LBD brain extract	Yes	Yes	Yes	Yes
Little or no binding to insoluble fibrils (Lewy bodies)	Yes	X	X	X

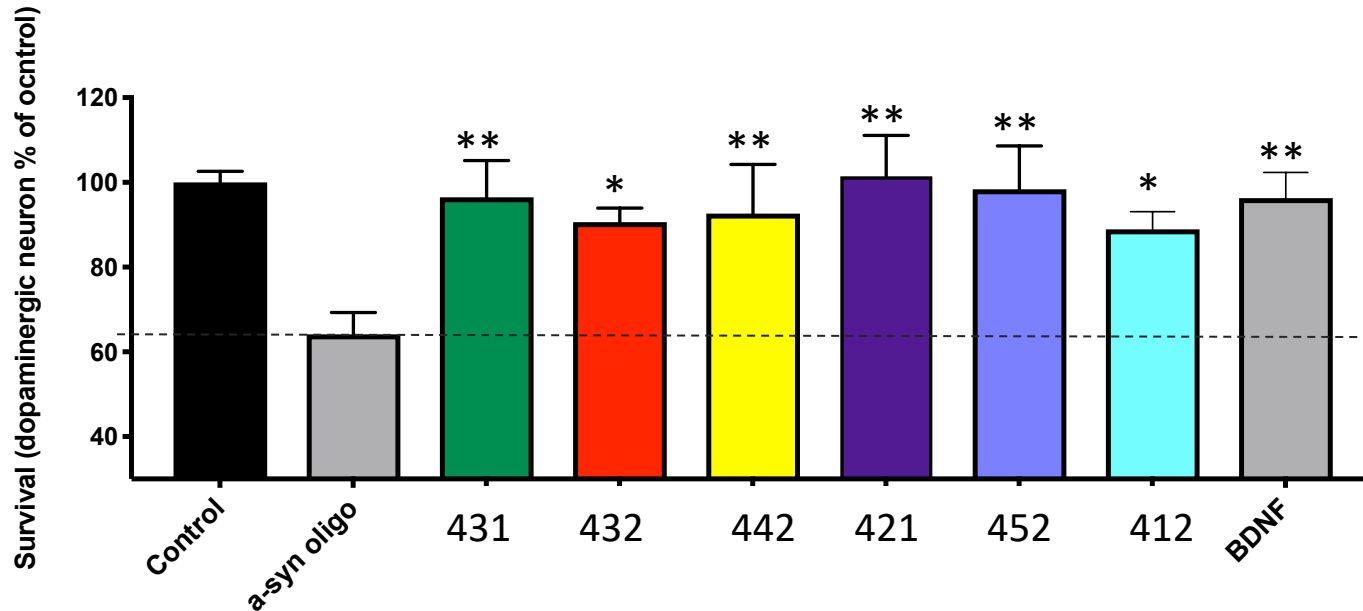


Limited benefit in PASADENA trial may be due to lack of selectivity

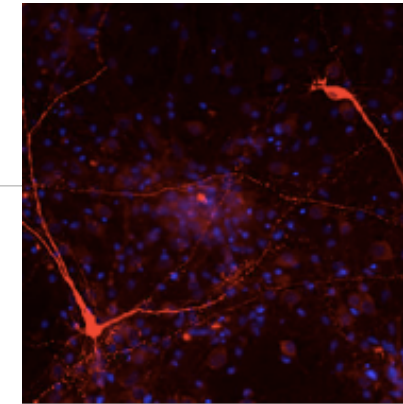


Discontinued: poor results may be due to lack of selectivity

ProMIS antibodies protect dopaminergic neurons against α -synuclein oligomer toxicity *in vitro*

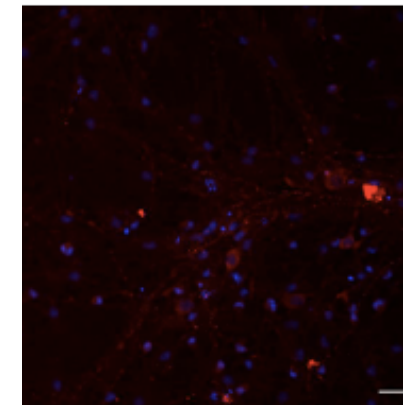


- Multiple antibodies provide neuroprotection in the same range as the brain-derived neurotrophic factor (BDNF) positive control
- As expected, antibodies alone had no effect on viability (not shown)



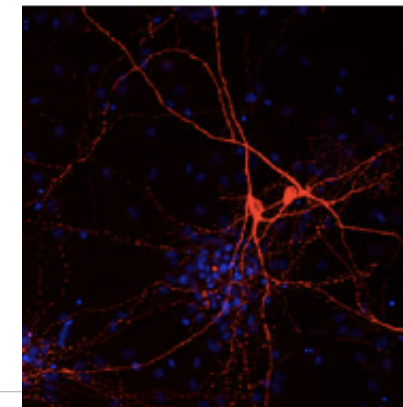
CONTROL

Normal neurons
in bright red



α -SYN OLIGOMERS

Neurons killed by
toxic oligomers



**PMN ANTIBODY +
 α -SYN OLIGOMERS**

Neuronal death
blocked by PMN
antibody

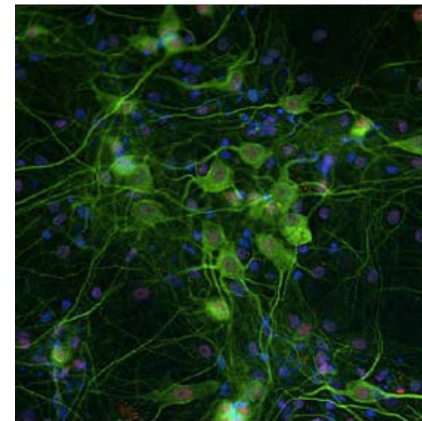
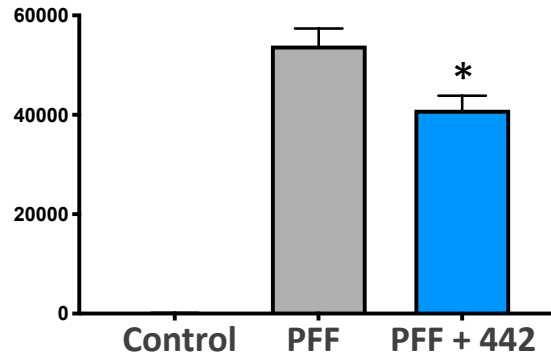
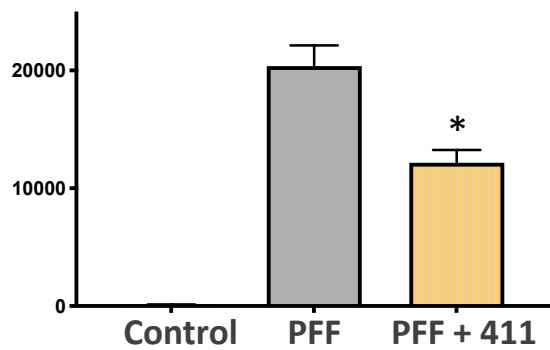
ProMIS antibodies inhibit α -syn propagation: Reduced PFF uptake and formation of aggregates

Human soluble α -syn fibrils +/- PMN antibody

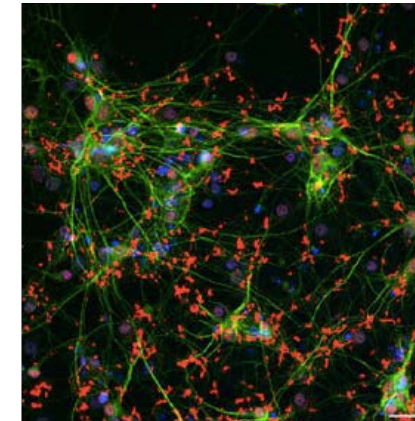
Staining for human α -syn aggregates



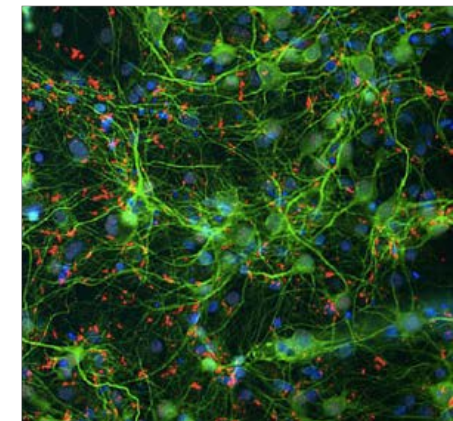
ProMIS antibodies significantly decrease formation of α -syn aggregates



CONTROL



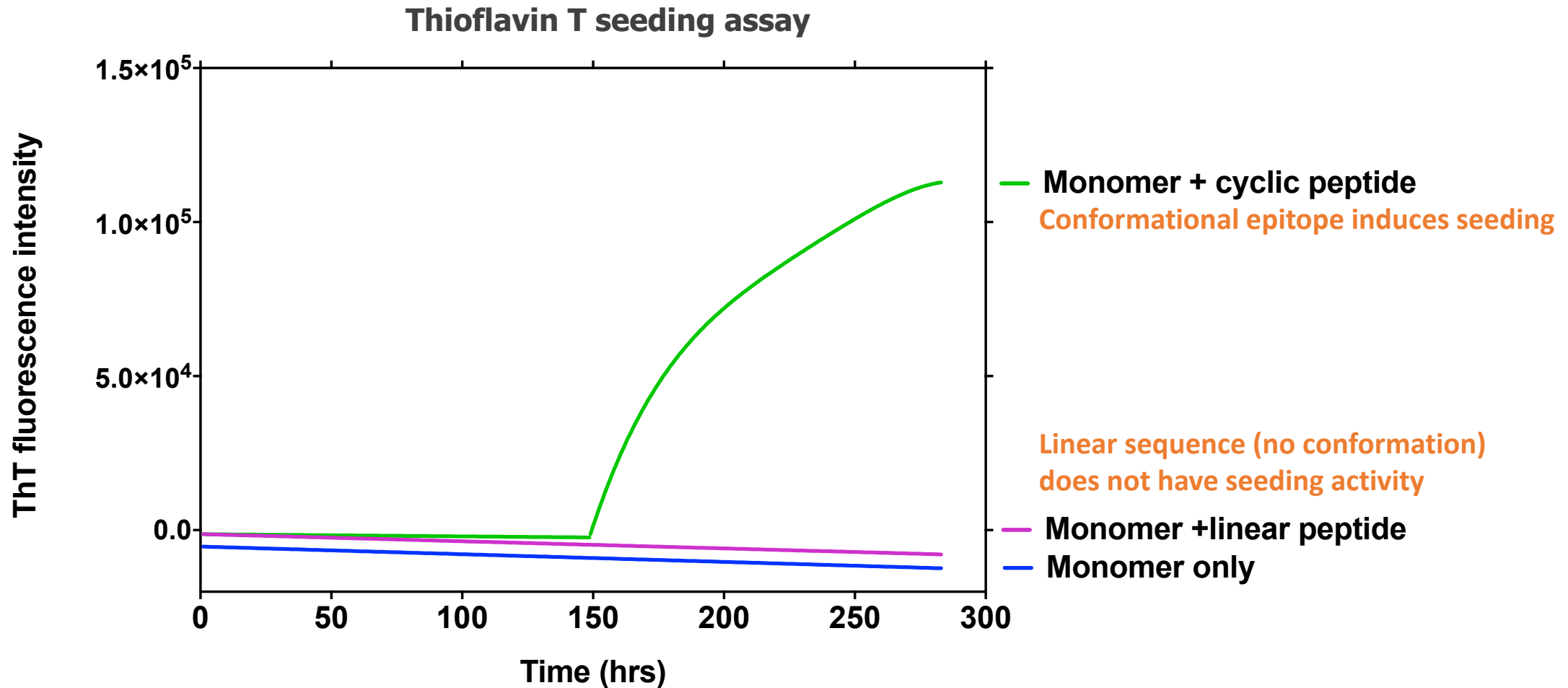
α -SYN SOLUBLE FIBRILS



α -SYN SOLUBLE FIBRILS +
PMN ANTIBODY 411

Human α -syn aggregates stained red

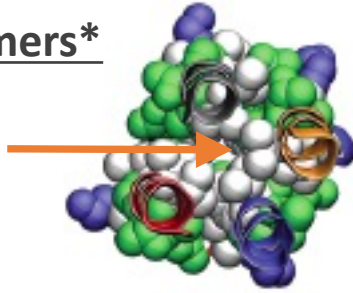
The misfolded portion of α -syn (conformational epitope) acts as a seed for toxic, prion-like propagation...confirming epitope prediction...explaining biology of disease



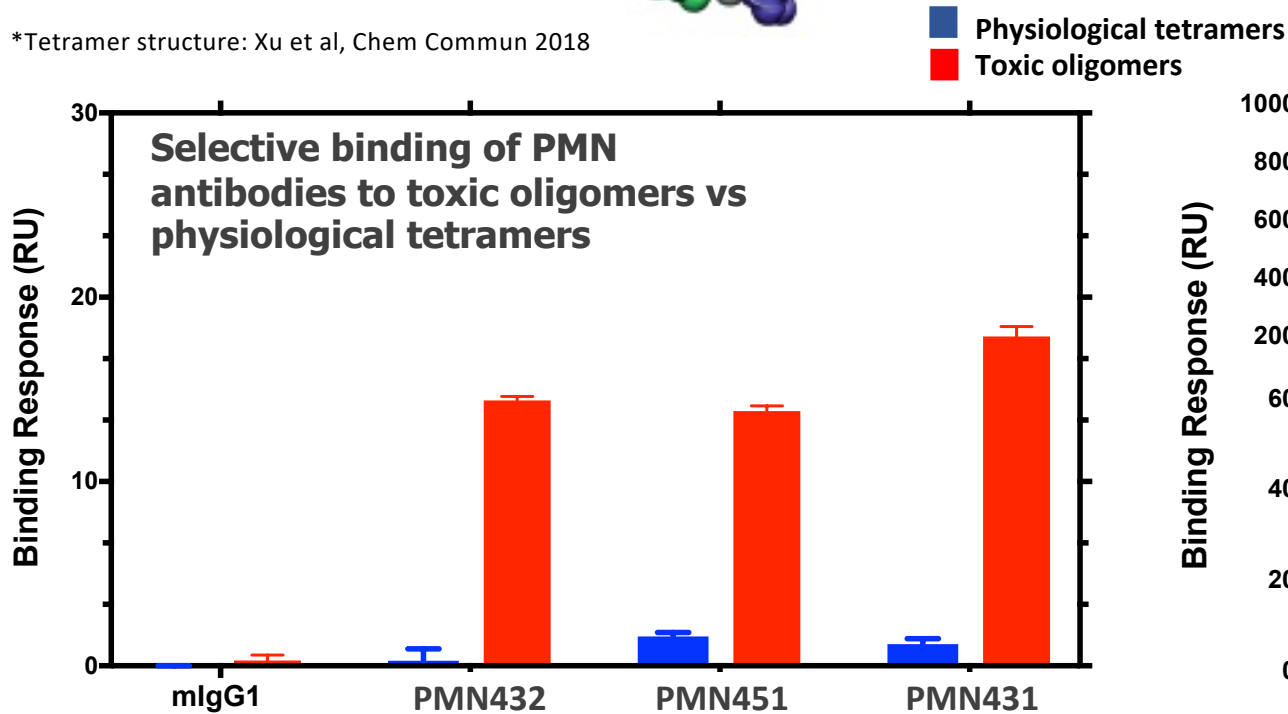
Only ProMIS antibodies bind toxic oligomers (tetramers) but not physiologic tetramers (SPR) – unrivalled ability to create multiple selective antibodies based on computational predictions of conformational epitopes

Physiologic tetramers*

Target epitope of PMN antibodies is buried in the hydrophobic core

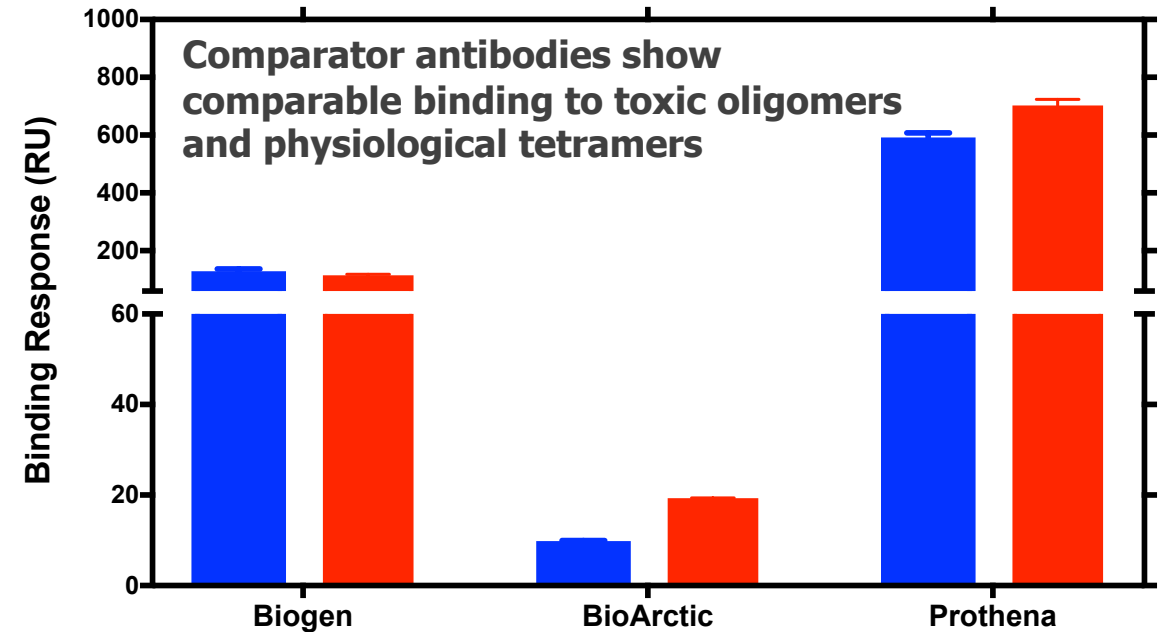
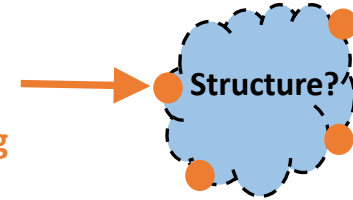


*Tetramer structure: Xu et al, Chem Commun 2018



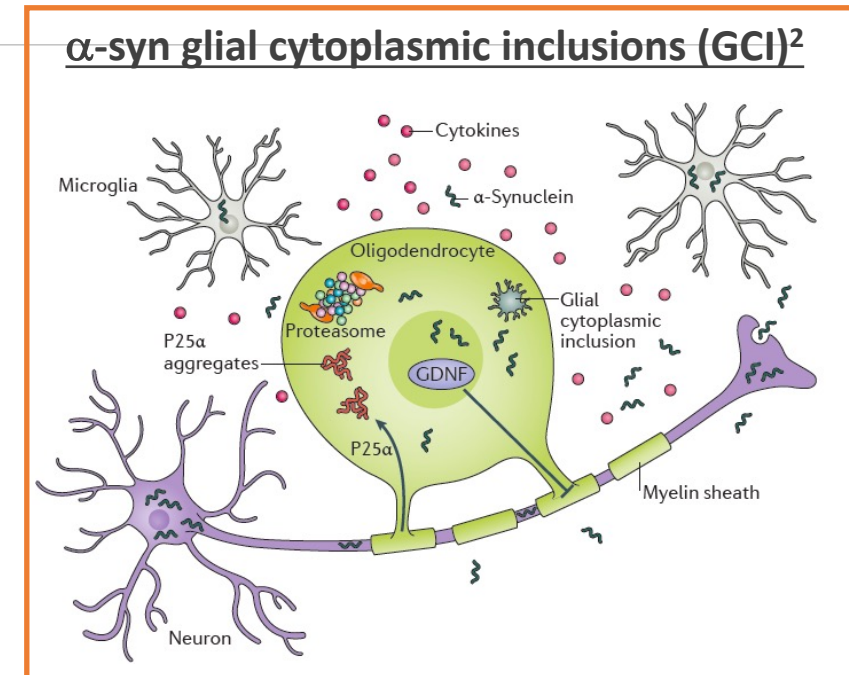
Misfolded Toxic oligomers

Target epitope of PMN antibodies is exposed and available for binding

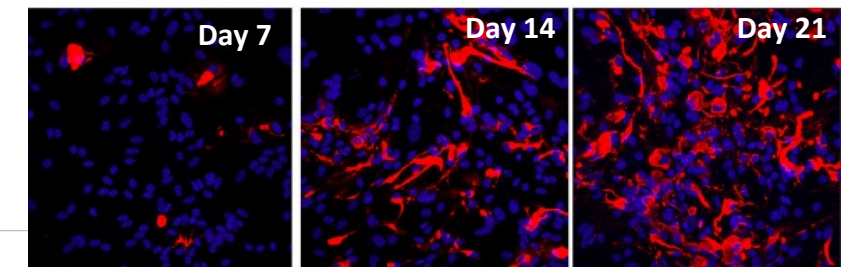


Multiple system atrophy (MSA) as a disease indication for therapeutic antibodies targeting toxic species of α -synuclein

- MSA is a rare neurodegenerative disease with an estimated prevalence of 3.4-4.9 cases per 100,000 population¹
- MSA is characterized by rapidly progressive autonomic failure and motor symptoms with predominant parkinsonian features (MSA-P) or dominant cerebellar features (MSA-C)^{1,2}
- There is no effective treatment and the mean survival from the onset of symptoms is 6-10 years^{1,2}
- Histologically characterized by α -synuclein aggregates in the cytoplasm of oligodendrocytes and to a lesser extent in neurons and other glial cells^{1,2}
- Alpha-synuclein aggregates from MSA brain homogenates propagate in a prion-like manner in vitro³ and in vivo⁴ and cause MSA-like neurodegeneration in mice⁵
- Suitability for clinical testing:
 - Rapid progression allows for earlier efficacy signal
 - High levels of NfL in serum represent a potential biomarker for effect on neuronal damage – early clinical PoC
 - No placebo effect observed in clinical trials to date
 - Rare disease but easy recruitment due to unmet need and existence of a global MSA Registry (GLOMAR) and supporting organizations^{1,2}



In vitro propagation of MSA α -syn aggregates³



ProMIS Alpha-Synuclein Program- Summary

- **Several antibody candidates** show desired selectivity and in vitro benefit, blocking neurotoxicity and propagation
- Alpha-synuclein protein a difficult challenge in molecular species selectivity – six molecular species of alpha-synuclein, two physiologically important that should be avoided, two toxic that should be targeted
- **ProMIS has better selectivity than competitor antibodies** in the clinic: Roche/Prothena antibody completely non-selective, mixed results in Phase 2 but proceeding; Biogen/Neurimmune antibody completely non-selective, discontinued.
- Only ProMIS antibodies can selectively target **toxic** tetramers of alpha synuclein, but not **physiologic** tetramers. ProMIS has achieved this with multiple antibodies, created with different conformational epitope predictions, illustrating the power of its unique platform
- Lead antibodies are undergoing humanization, next step is IND enabling work
- Target indications are MSA (multiple system atrophy, rare disease, elevated biomarkers) and Parkinson's disease

TDP-43 Program

Selective targeting of pathogenic TDP-43 for optimal safety and efficacy

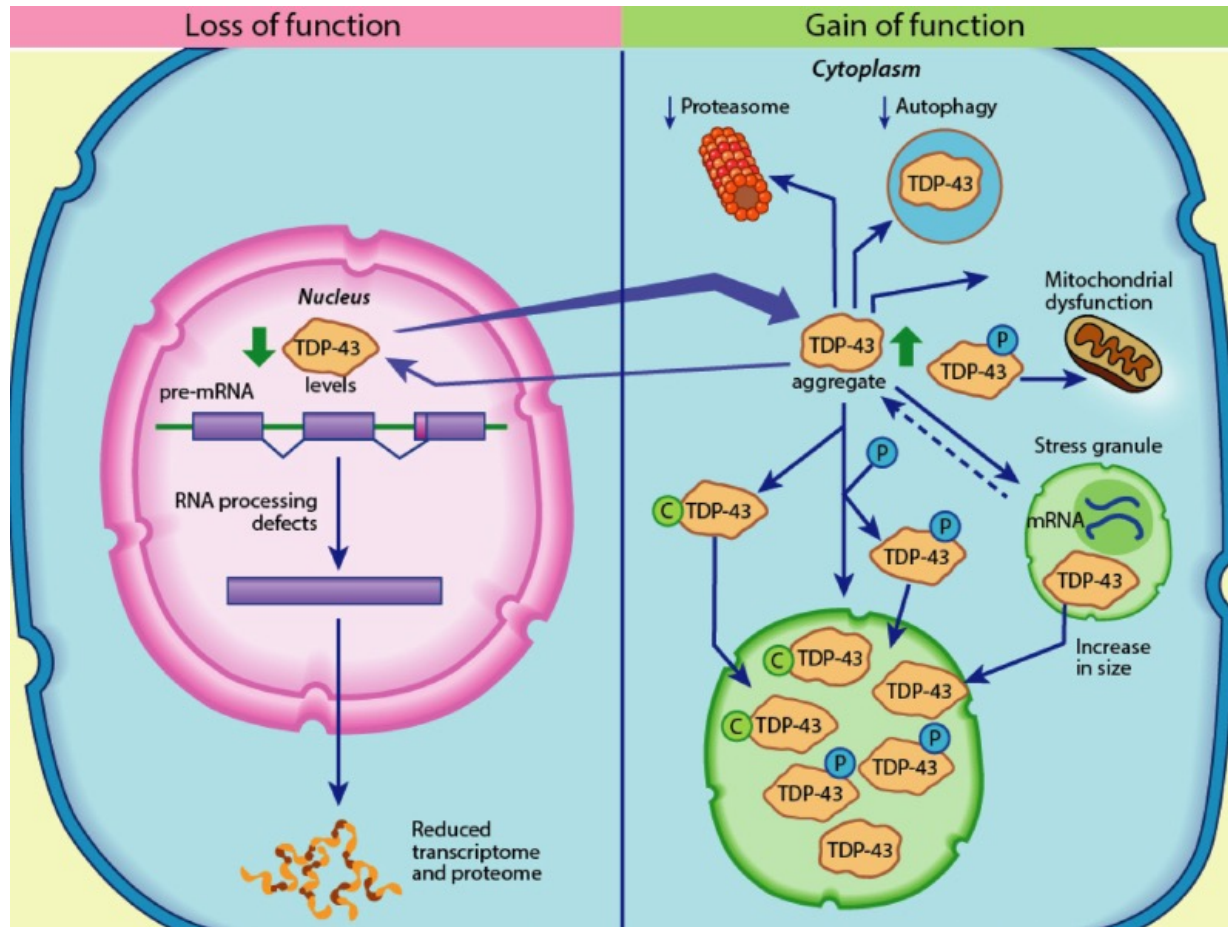


Figure from de Boer et al¹

TDP-43 is essential to neuronal cell survival¹

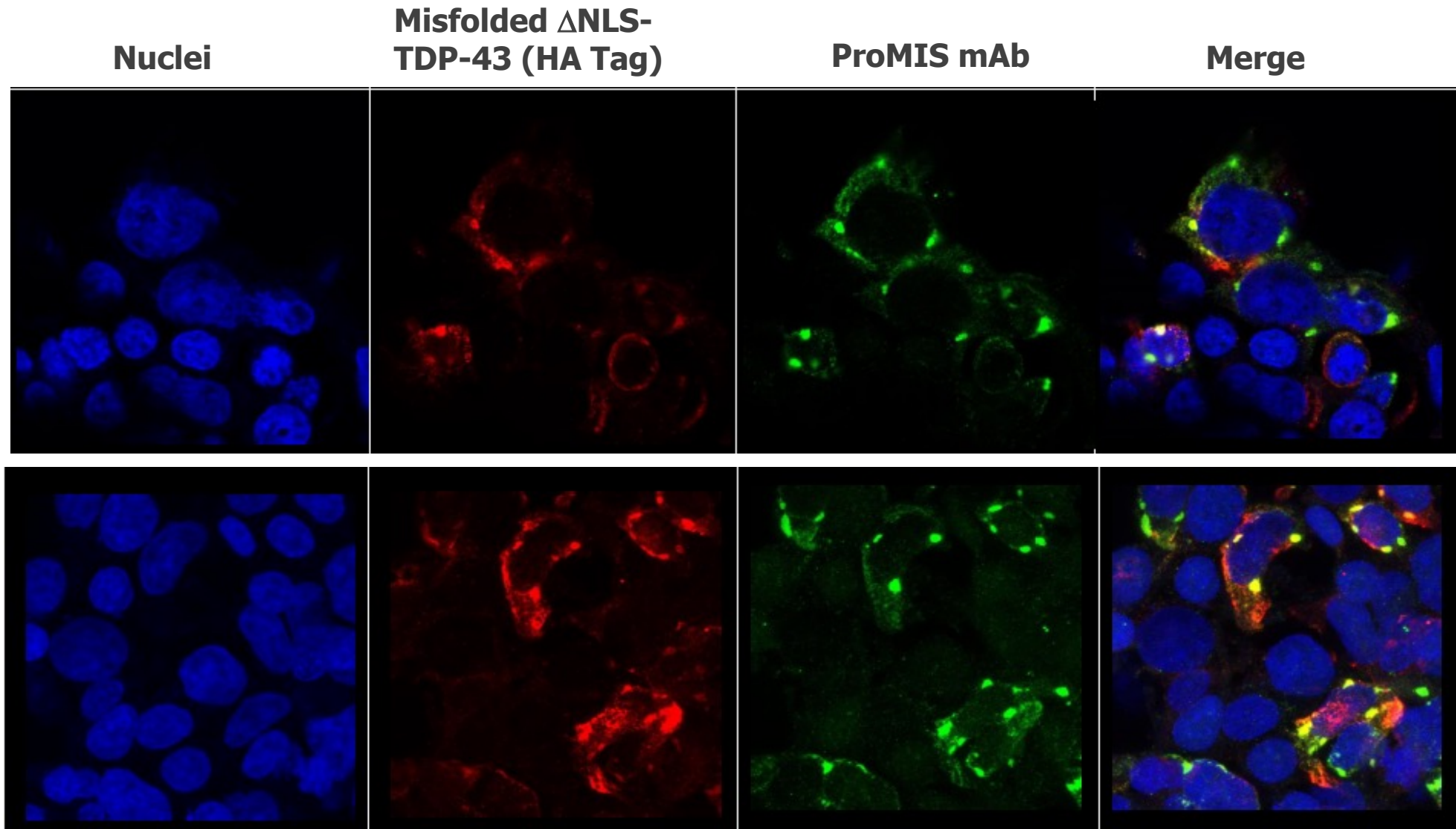
- TDP-43 is normally present in the nucleus of all cells and performs an essential role in RNA splicing, transport and stability
- Under stress conditions (e.g. oxidative stress) normal TDP-43 also forms protective stress granules in the cytoplasm

Misfolded TDP-43 gives rise to both loss of function and toxic gain of function¹

- Loss of function: Under disease conditions, misfolding of TDP-43 causes formation of mislocalized cytoplasmic aggregates. Nuclear depletion leads to defective splicing and transport of mRNA.
- Toxic gain of function: Cytoplasmic aggregates of misfolded TDP-43 are toxic and interfere with physiologic stress granule function. They also induce misfolding of other proteins into pathogenic aggregates²⁻⁴ - "TDP-43 Pathological Interactome"

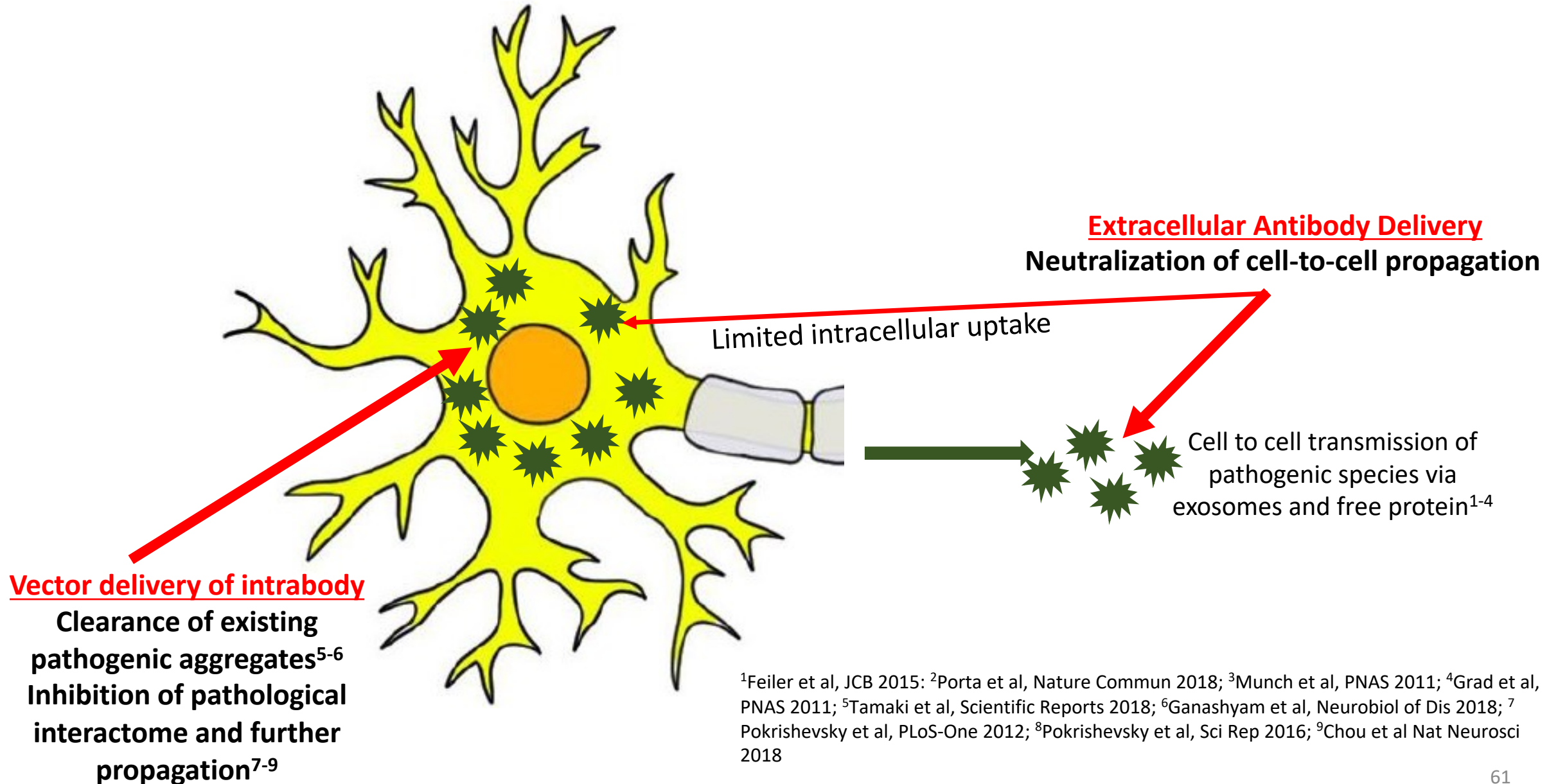
Targeting of pathogenic TDP-43 requires stringent selectivity for the misfolded form of the protein to avoid safety concerns

ProMIS mAbs to misfolded TDP-43 show excellent selectivity: react with mislocalized, aggregated Δ NLS-TDP-43 but not nuclear wild type TDP-43

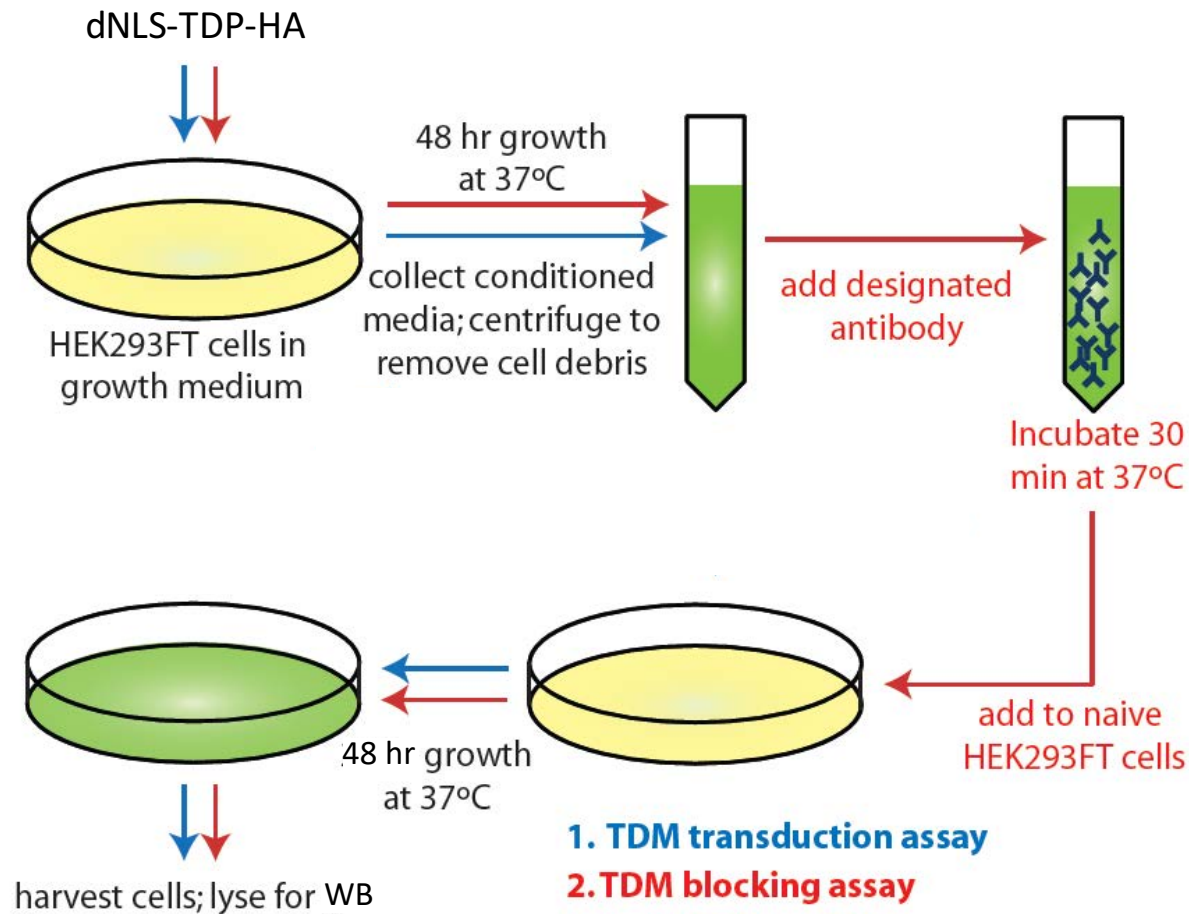


- HEK-293 cells transfected with Δ NLS TDP-43 lacking a functional nuclear localization signal
- Cells stained for HA tag (red) of overexpressed Δ NLS TDP-43 or with rabbit mAb to misfolded TDP-43 epitope at 2 μ g/ml (green).
- Nuclei stained with DAPI (blue)
- Images analyzed by confocal microscopy (Z-stacks)

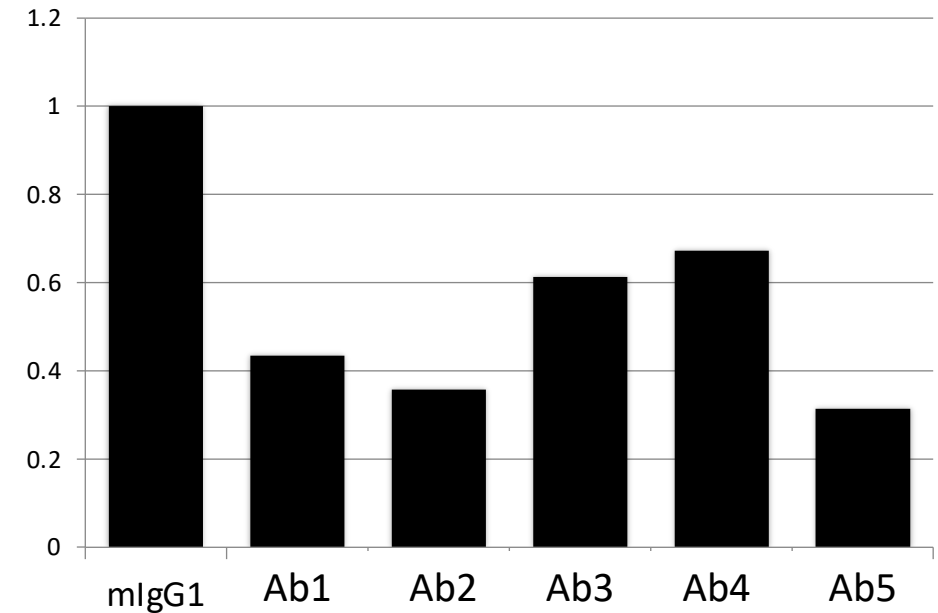
Antibody-Based Targeting of Pathogenic TDP-43 - Two approaches; extra cellular antibody, vectorized intrabody (gene therapy – with a partner)



Functional Assay: ProMIS *antibodies* inhibit cell-to-cell transmission of misfolding Δ NLS-TDP43 – suggests potential as extracellular antibody



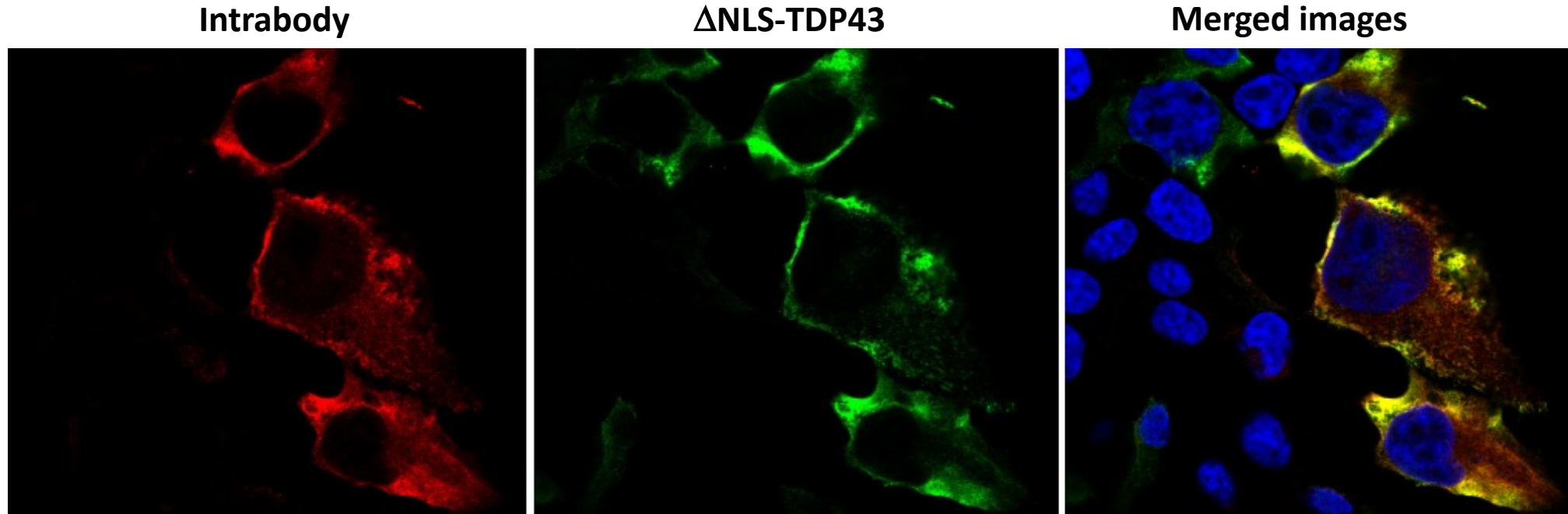
HA- Δ NLS-TDP-43 transmission relative to mIgG1 control



ProMIS mAbs inhibit transmission of misfolding TDP-43 from the conditioned medium of donor HEK293 cells transfected with Δ NLS-TDP-43 to naïve recipient cells

ProMIS *intrabodies* highly selective: interact only with cytoplasmic Δ NLS-TDP43 aggregates and not normal nuclear TDP43

HEK293 cells transfected with Δ NLS-TDP43 and single chain ProMIS intrabody



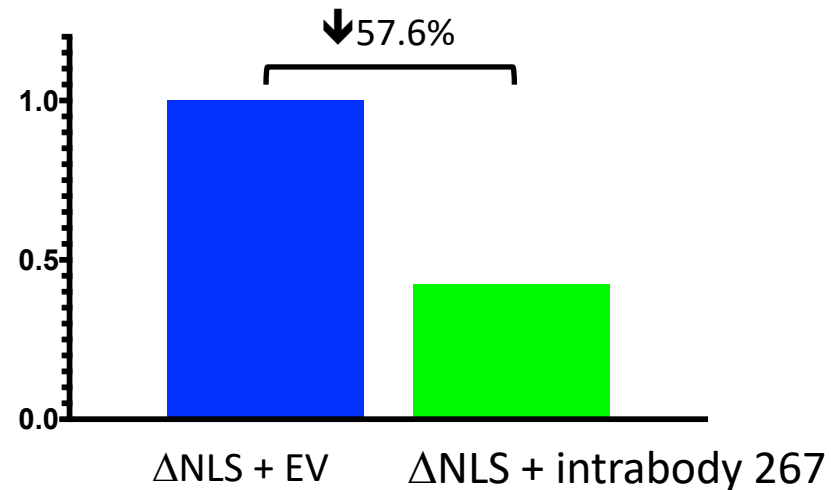
- Expression of ProMIS TDP43 intrabody is not toxic to cells
- Intrabody co-localizes with mislocalized, aggregated cytoplasmic Δ NLS-TDP43
- Intrabody does not interact with endogenous normal TDP43 in the nucleus

TDP-43 intrabody promotes clearance of misfolded TDP-43 aggregates inside the cell – functional benefit



Misfolded TDP43-selective intrabody with lysosomal targeting signal promotes degradation of TDP-43 aggregates without cellular toxicity

HA intensity relative to control “dNLS+EV”



ProMIS TDP-43 program: Summary

- Several antibody candidates show molecular species selectivity: binding to toxic mis-folded TDP-43, not normal physiologic TDP-43
- One candidate a lead priority for use as an extracellular antibody, shows functional benefit in blocking prion-like propagation in vitro
- Other candidates prioritized as “intrabodies”, targeting intracellular mis-folded TDP-43; high affinity binding, selectivity for toxic TDP-43, no detrimental impact on cell viability, clearance of pathogenic aggregates from inside cells
- ProMIS unlikely to enter gene therapy space directly, pursuing partnering discussions with large “peer” organizations with gene therapy/vector capability
- TDP-43 toxic “interactome” with other ProMIS targets – RACK1, SOD1, ataxin2 – potential synergistic portfolio for ALS disease-modifying treatment
- All candidates preclinical stage, initiating humanization, next step is IND enabling work

Experienced Leadership Team

Name	Title	Years of Experience	Prior Experience
Gene Williams	Chairman & CEO	25+	<ul style="list-style-type: none"> Former SVP at Genzyme, with senior roles integrating commercialization, drug development, and deal making Recently the CEO of Dart Therapeutics, an Orphan Disease drug development company Founder and director of Adheris, which became the largest company in the patient adherence/compliance area
Elliot Goldstein	President	25+	<ul style="list-style-type: none"> Held positions as SVP of Strategic Product Development at SmithKline Beecham (now GSK) Chief Operating Officer and Chief Medical Officer of Maxygen Chief Operating Officer at DART Therapeutics
Neil Cashman	Chief Science Officer	25+	<ul style="list-style-type: none"> Holds the Canada Research Chair in Neurodegeneration and Protein Misfolding Diseases, Serves as the Director of the University of British Columbia ALS Centre, Awarded the Jonas Salk Prize for biomedical research in 2000
David Wishart	Chief Physics Officer	25+	<ul style="list-style-type: none"> Distinguished University Professor in the Departments of Biological Sciences and Computing Science at the University of Alberta Co-Director of The Metabolomics Innovation Centre Bristol-Myers Squibb Research Chair in Pharmaceutical Sciences 1995-2005 Fellow of the Royal Society of Canada
Dan Geffken	CFO	25+	<ul style="list-style-type: none"> Founding Managing Director of Danforth Advisors Served as the Chief financial officer of Homology, Inc, GenePeeks, Inc., Transkaryotic Therapies, Inc., Cidara, Inc., Apellis, Inc. and Stealth BioTherapeutics, Inc.
Johanne Kaplan	Chief Development Officer	25+	<ul style="list-style-type: none"> Former VP of Research at Genzyme Associate Immunopathologist at SmithKline Beecham where she established an Immunotoxicology program Her work has resulted in over 60 scientific publications and multiple patents
Gavin Malenfant	Chief Operating Officer	25+	<ul style="list-style-type: none"> Head of Operations at Sarepta, in support of approval/ launch of EXONDYS 51 At Genzyme, head of operations for research and development At Genzyme, ead rare disease program management organization



Scientific Advisory Board

Name	Years of Experience	Prior Experience	Affiliations
Sharon Cohen, MD	20+	<ul style="list-style-type: none"> Medical Director & Principal Investigator of Toronto Memory Program FRCPC in neurology from Royal College of Physicians of Canada and a fellowship in Behavioural Neurology from the University of Toronto 	 Toronto Memory Program
Rudy Tanzi, PhD (Chairman)	20+	<ul style="list-style-type: none"> Professor of Neurology at Harvard University, Vice Chair of Neurology, Director of Genetics & Aging Research Unit, Co-Director McCance Center for Brain Health at Mass General Hospital 	 HARVARD UNIVERSITY MASSACHUSETTS GENERAL HOSPITAL
Bill Mobley, MD, PhD	25+	<ul style="list-style-type: none"> Dean for Neurosciences Initiatives, Distinguished Professor of Neurosciences, and Florence Riford Chair for Alzheimer Disease at the University of California, San Diego 	 UC San Diego SCHOOL OF MEDICINE
James Kupiec, MD	20+	<ul style="list-style-type: none"> Former VP, Global Clinical Leader for Parkinson's disease, and Clinical Head of the Neuroscience Research Unit for Pfizer, Inc. Clinical focus on development of therapies for neurodegenerative disorders 	 Ciba sanofi~synthelabo Pfizer
C. Warren Olanow, MD	25+	<ul style="list-style-type: none"> Previous Henry P & Georgette Goldschmidt Professor & Chairman, Department of Neurology at Mount Sinai School of Medicine, presently Professor Emeritus Department of Neurology & Department of Neuroscience, CEO of CLINTREX 	 MOUNT SINAI SCHOOL OF MEDICINE
Andre Strydom, MD, PhD	25+	<ul style="list-style-type: none"> Professor Institute of Psychiatry, Psychology and Neuroscience at King's College London Honorary Consultant psychiatrist, South London and the Maudsley NHS Foundation Trust 	 KING'S COLLEGE LONDON NHS South London and Maudsley NHS Foundation Trust
Michelle Hastings, PhD	20+	<ul style="list-style-type: none"> Professor and Director, Center for Genetic Diseases, Rosalind Franklin University of Medicine and Science Faculty Member at the Chicago Medical School 	 ROSALIND FRANKLIN UNIVERSITY OF MEDICINE AND SCIENCE 