

How to Prepare for the Precision Medicine Era

- A GENOMIC PERSPECTIVE ON HEALTH AND DISEASE

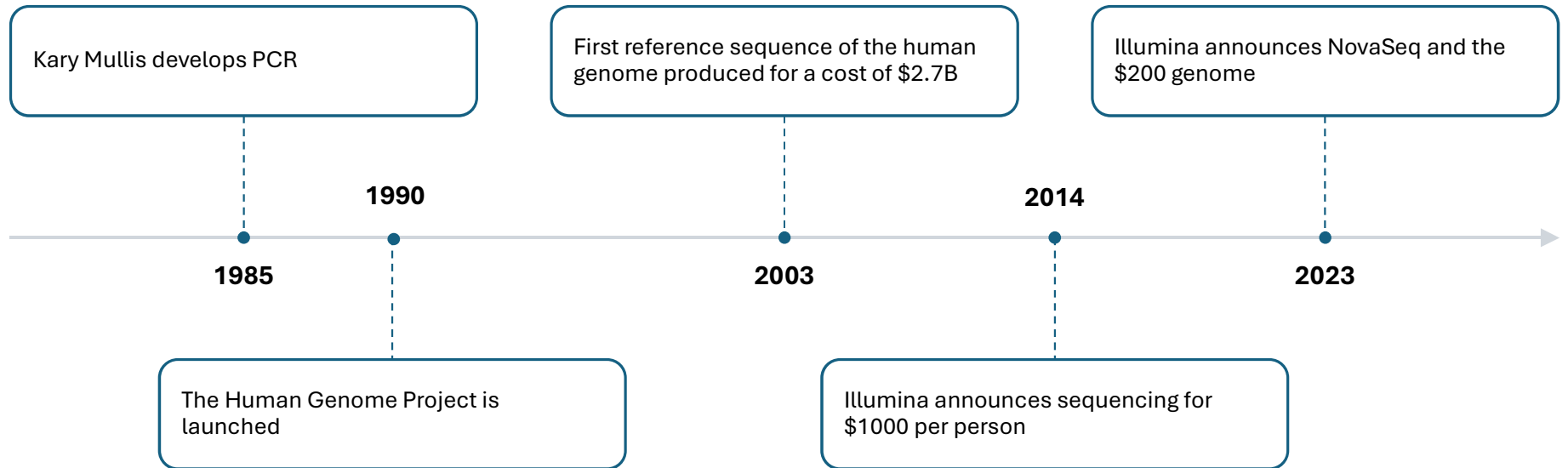
- Katheron Intson, PhD
- For: Parker University
- NeuroCon, Dallas, Texas
- July 27, 2024



“In each of us is written the story of our species.” –Jacques Monod

- Our DNA is composed of millions of base pairs that code for everything that makes us who we are
- Within DNA are “genes”, the units of heredity
- There are differences in the DNA sequences among individuals, called “genetic variants”
- These variants contribute to the uniqueness of each individual

The development of genome sequencing



Genetic variants influence:

Physical Traits

Behaviours

Disease Susceptibility

Variants shape the way we interact with our physical environment, and the way our environment interacts with us

PMID: 24045858

- White coat in *Panthera leo* is caused by a mutation to the Tyrosinase (TYR) gene
- Non-tawny lions camouflage poorly
 - Poorer hunters
 - Vulnerable to human hunting



Variants shape our personality and life

- **Dopamine Receptor D4 and Novelty-Seeking (*DRD4-7R*)**
 - Greater propensity for risk-taking, novelty-seeking, sensitivity to the environment, ADHD, addiction
- **Monoamine Oxidase A and Aggression (*MAOA-3L* and *MAOA-4R*)**
 - Higher levels of aggression and impulsivity; increased risk of antisocial behaviour if exposed to child abuse
- **Catechol-O-Methyltransferase and Stress Resilience (*COMT Val158Met*)**
 - Greater stress resilience following exposure to trauma, but at the cost of poorer learning + memory
- **Fat Mass & Obesity-Associated Protein and Energy Balance (*FTO* variants)**
 - Increased risk of obesity



It's becoming clear that common diseases are often constellations of **rare single-gene variants**

NEUROMUSCULAR DISEASE

PRACTICE GUIDELINES | VOLUME 31, ISSUE 12, P1313-1357, DECEMBER

[Download Full Issue](#)

The 2022 version of the gene table of neuromuscular disorders (nuclear genome)

EPILEPSY

ARTICLE

DOI: 10.1038/s41467-018-07524-z OPEN

Genome-wide mega-analysis identifies 16 loci and highlights diverse biological mechanisms in the common epilepsies

SCHIZOPHRENIA

Article

Rare coding variants in ten genes confer substantial risk for schizophrenia

MIGRAINE

ARTICLES

<https://doi.org/10.1038/s41588-021-00990-0>

nature genetics

OPEN

Genome-wide analysis of 102,084 migraine cases identifies 123 risk loci and subtype-specific risk alleles

PARKINSON'S

ARTICLE

The genetic spectrum of a cohort of patients clinically diagnosed as Parkinson's disease

AUTISM

Cell

Genomic architecture of autism from comprehensive whole-genome sequence annotation



Talk Outline

- How genetic variants are **linked** to disease
- How genetic diseases can be **treated** with precision and personalized medicine
- **Examples** of precision medicines
- How medicine is **changing**
- What **you** should be prepared for in your **practice**



The end of congenital disability?

- 6,800,000 Americans live with congenital disabilities
- Can be diagnosed prenatally, at birth, or in childhood
- 80% are due to genetic variants (mutations)



The end of neurodegeneration?

- 7,000,000 Americans live with neurodegenerative disease (Alzheimer's, Parkinson's, ALS)
- Can be early-onset, but usually diagnosed after age 65
- At least 5%, but likely more, are due to monogenetic (single-gene) variants

Emerging therapeutic modalities offer hope for reversal of genetic disease

Enzyme Replacement Therapies

Antisense Oligonucleotides

Gene therapy

Gene editing

A TALE OF MICE AND MEN: The discovery of GRI Disorder

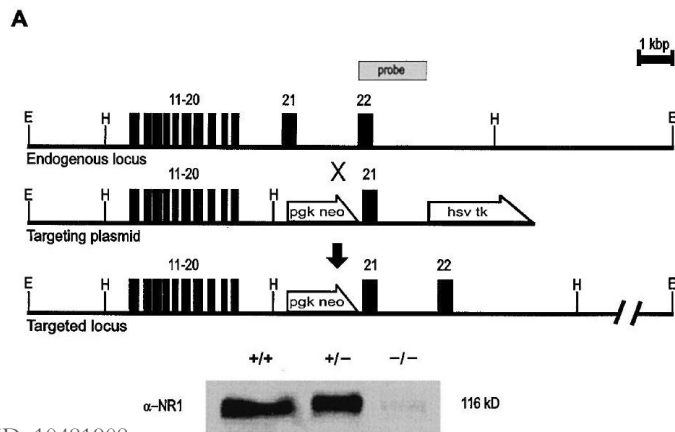
A non-syndromic epileptic
encephalopathy with gene
therapy in-development



Bryson MacArthur, first annual GRICon, 2018

An introduction to *Grin1*^{KD} mice

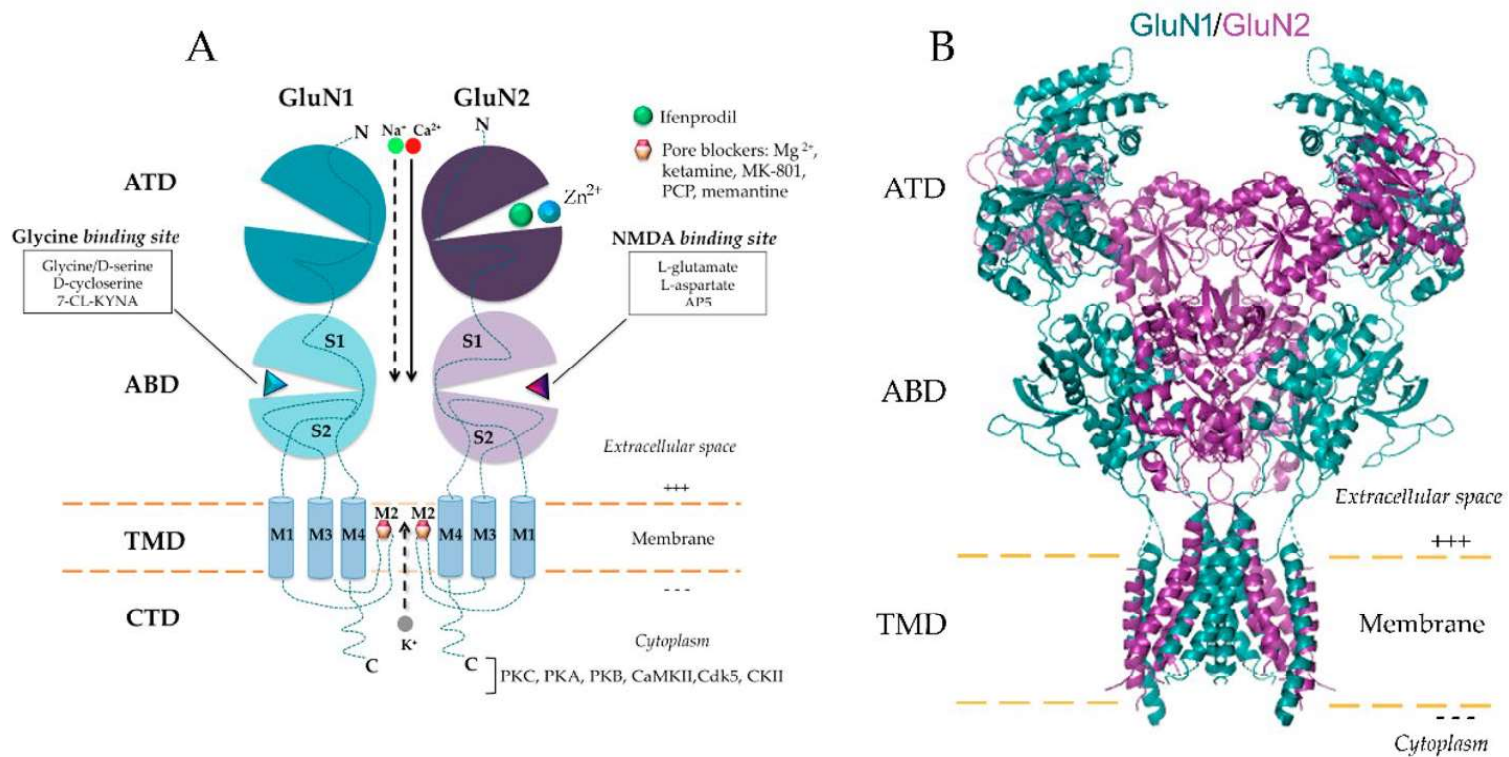
- Insertion of neomycin resistance gene into intron 20 of *Grin1* gene
- *Grin1*^{KD} mice have 7.3% Grin1 protein levels of WT mice; *neo* mutation produces hypomorphic allele
- *Grin1*^{KD} mice have altered motor, stereotypic, social, vocal, and sexual behaviours



Wild-type (typical) mouse

Grin1^{KD} mouse

Grin1 encodes the must-have subunit of the N-Methyl-D-Aspartate (NMDA) Receptor





It was all for the love of science until the *GRIN1* families found us...

These children had been diagnosed with rare or unique variants to the *GRIN1* gene through genome sequencing

Grin1^{KD} mice as a model of human *GRIN1* variants

MICE

Seizures

Progressive volume reductions in some brain structures + white matter

Impaired learning + executive function in tasks

Altered social communication measured via USV

Hyperlocomotion in OFT

Possible cortical blindness

PATIENTS

Epilepsy

Generalized cerebral atrophy

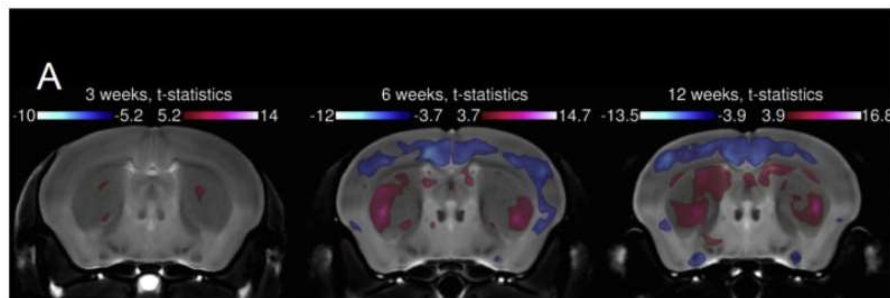
Intellectual disability

PMID: 27164704

Non-syndromic or severe speech delay

Hyperkinetic movement disorders

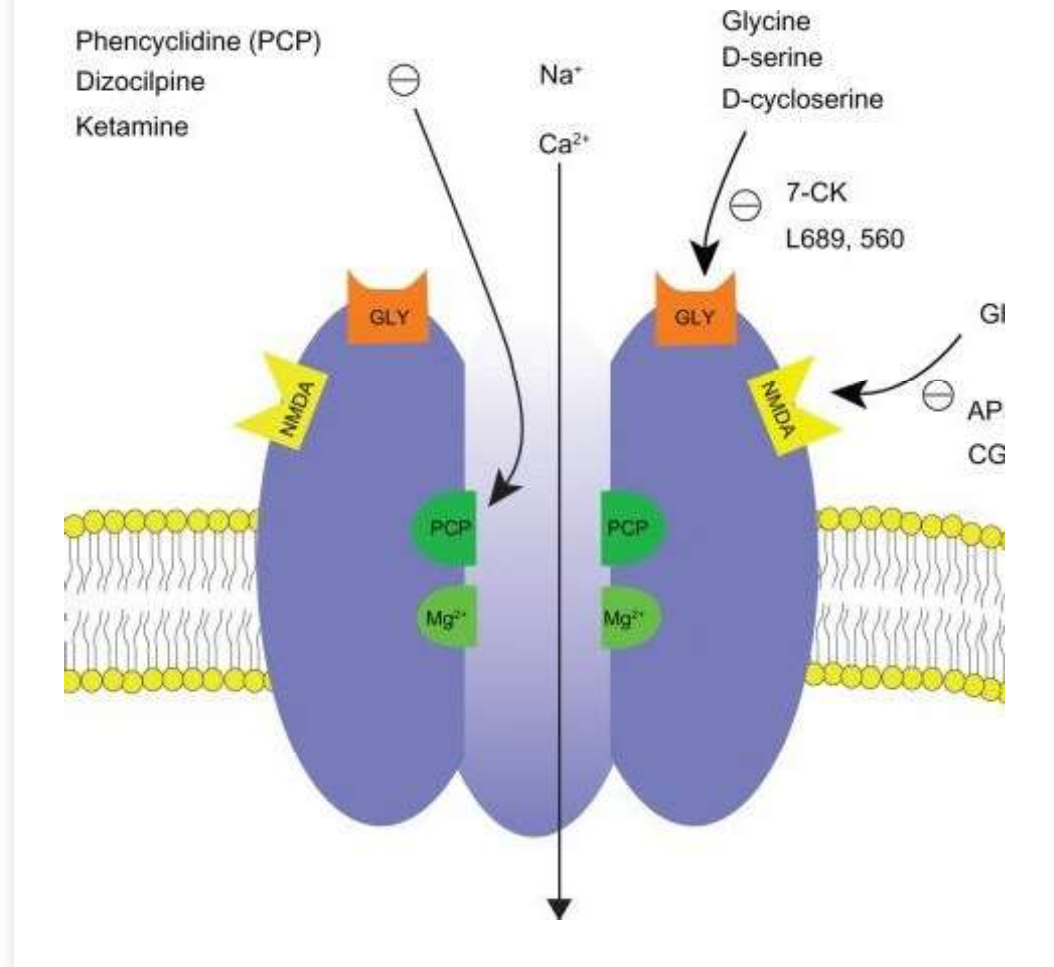
Cortical blindness



PMID: 31299220

What changes occur to the kids' NMDA receptors as a result of variants to the *GRIN1* gene?

- Total protein expression level
- Protein folding, degradation
- Subunit interaction and assembly
- Receptor trafficking
- Sensitivity to ligands
- Channel open time/probability, channel conductance, Mg⁺⁺ stickiness, leakiness
- Receptor desensitization or deactivation
- Binding of exogenous modulators
- Impaired excitation/inhibition balance
- Impaired synaptic plasticity



~~Small Molecule Drugs~~

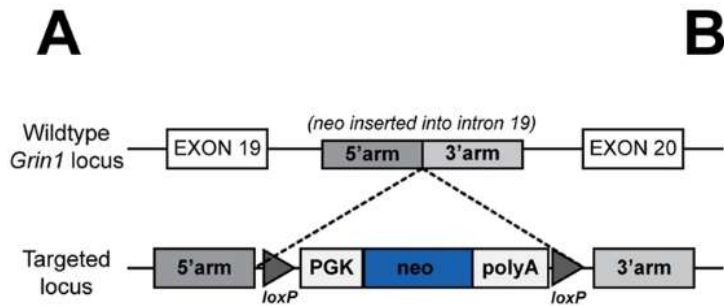
~~Enzyme Replacement Therapies~~

~~Antisense Oligonucleotides~~

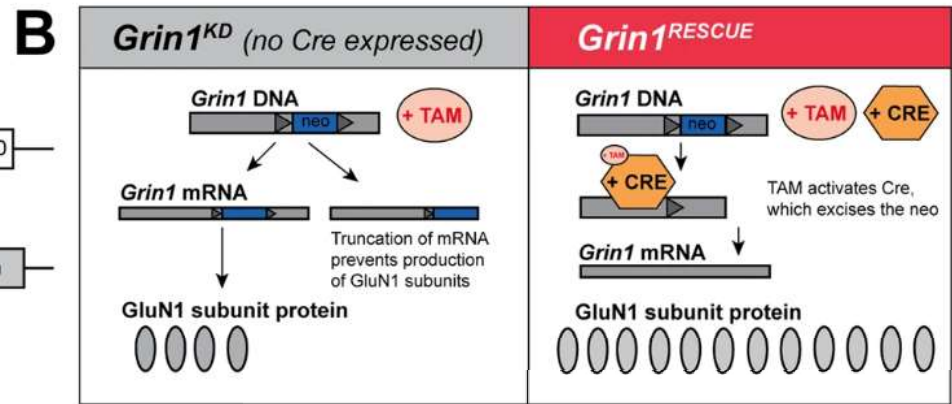
Gene therapy

Consequences of
lifelong Grin1
mutations in mice
can be rescued in
adulthood

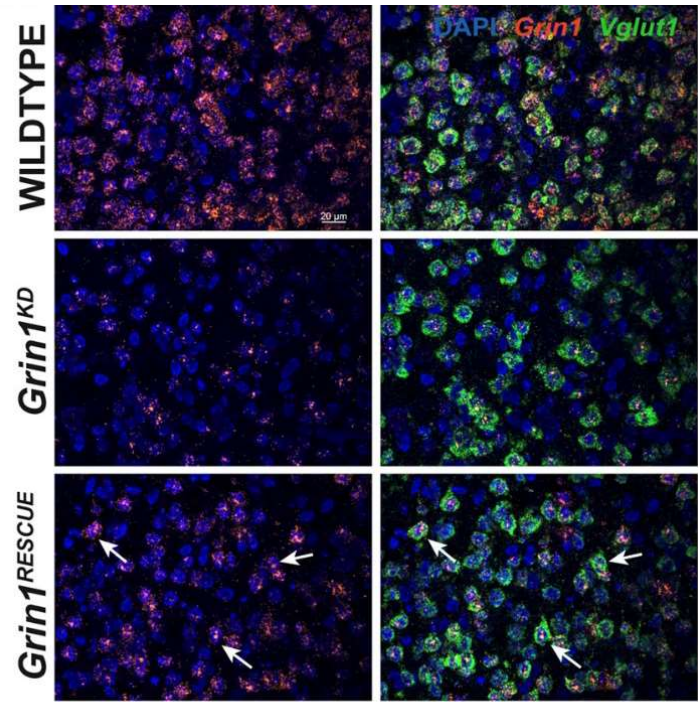




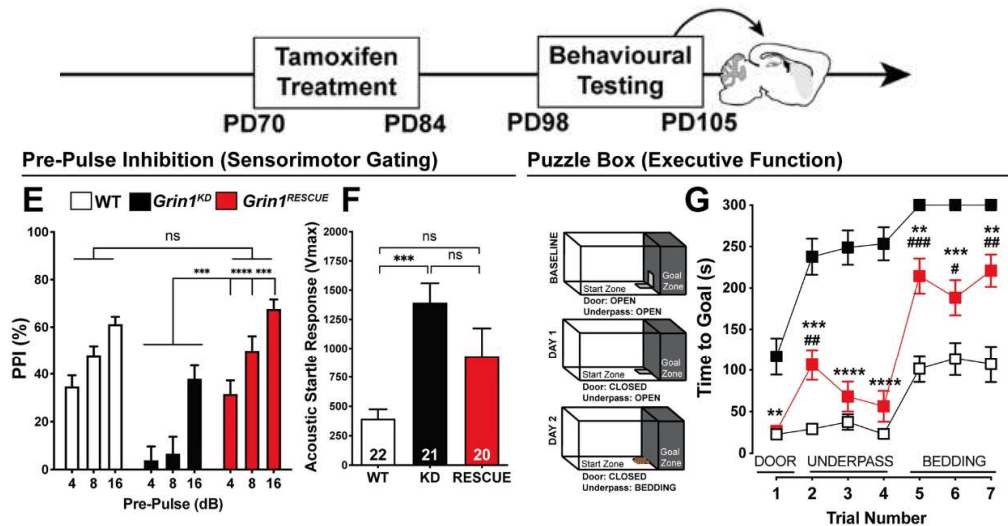
PMID: 10481908



Generation of *Grin1*^{RESCUE} mice

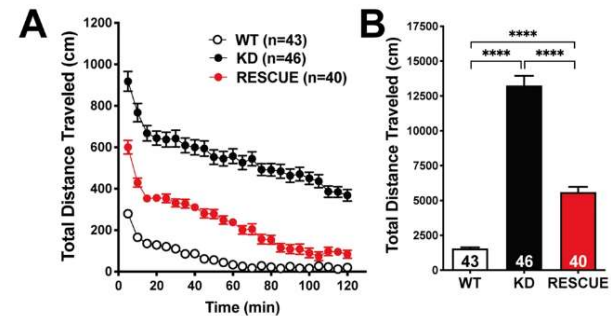


Cognitive impairments and seizures are improved by intervention in adult mice



*no seizures observed in treated mice

Novelty-Induced Locomotion





Takeaways

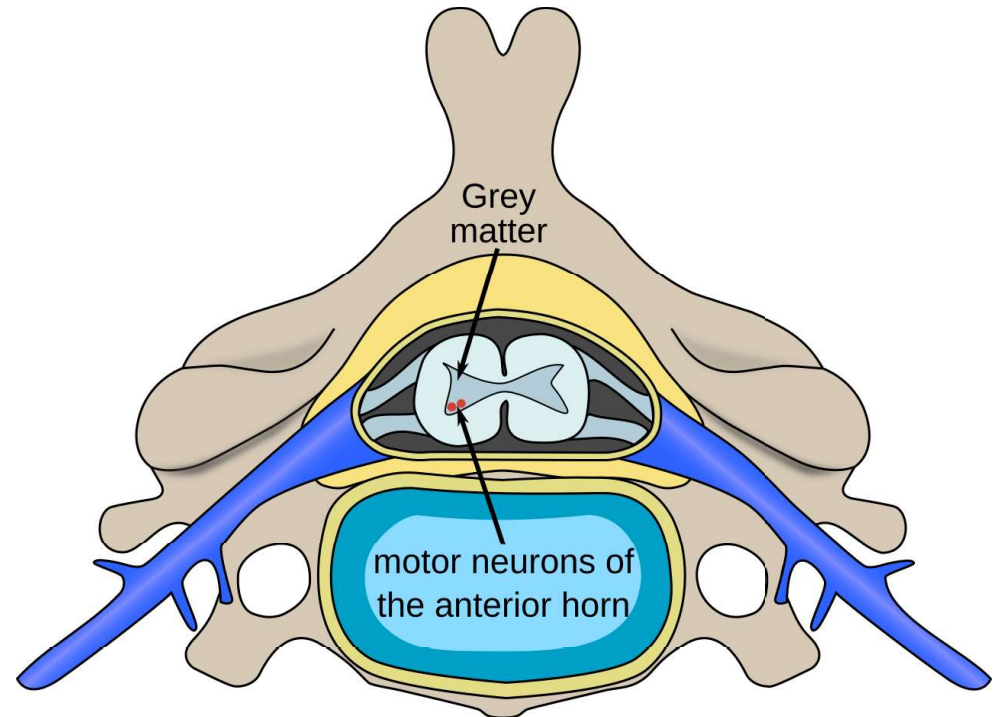
- Remarkable recovery from lifelong deficiencies in Grin1/NMDARs is possible in adult animals
 - Gene therapy likely holds promise for children with Grin1 variants
 - Gene therapy development now in-progress, preliminary results showing reversal of phenotype
 - If you are diagnosed with a genetic variant, try to get a sense of the gene's function in your body
-

Gene therapy
already in the
clinic



Example: Spinal Muscular Atrophy

- Progressive degeneration of motor neurons in the spinal cord beginning at age 6 months
- Usually autosomal recessive, caused by variants to Survival Motor Neuron 1 (SMN1) gene
- SMA Type 1, SMA Type 2, SMA Type 3, SMA Type 4 diagnoses based on onset/severity of symptoms



Historically, SMA Type 1 was a death sentence for children

Stage 1 (Birth to 6 months)

- Hypotonia, difficulty moving, difficulty feeding, shallow breathing

Stage 2 (6 months to 1 year)

- Regression of motor skills, loss of reflexes, respiratory issues, swallowing challenges

Stage 3 (1 year onward)

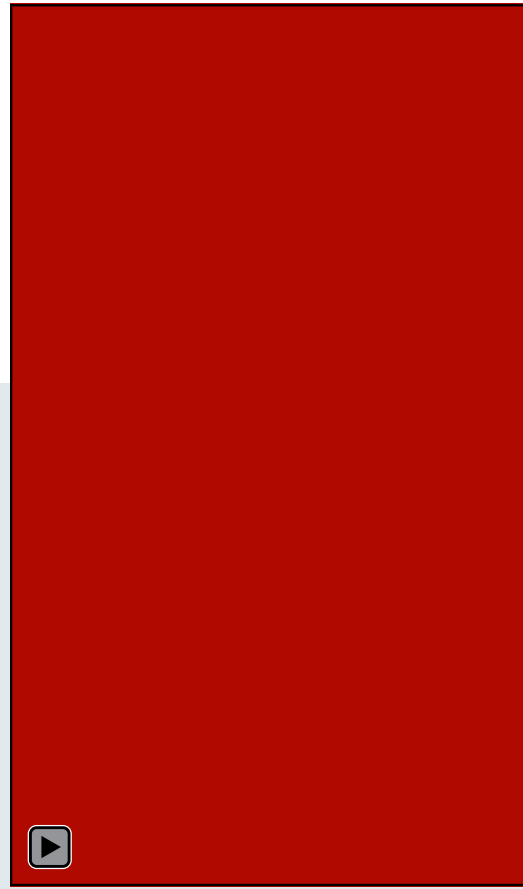
- Severe motor impairment, skeletal deformities, ventilation, food aspiration, infection

Death

Motor Milestones: 10.5 Months

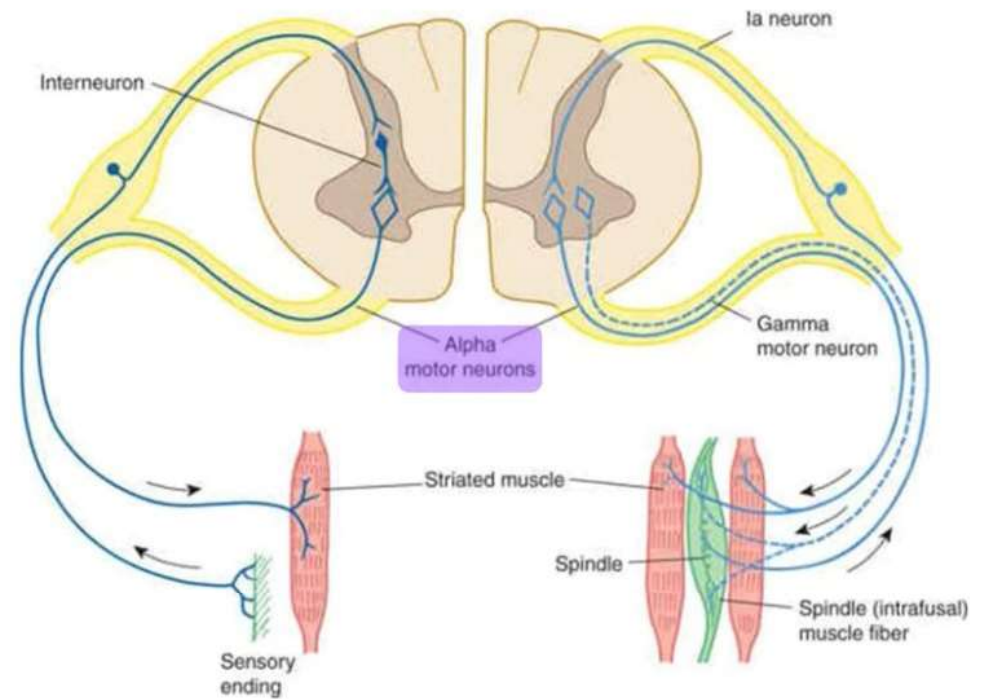


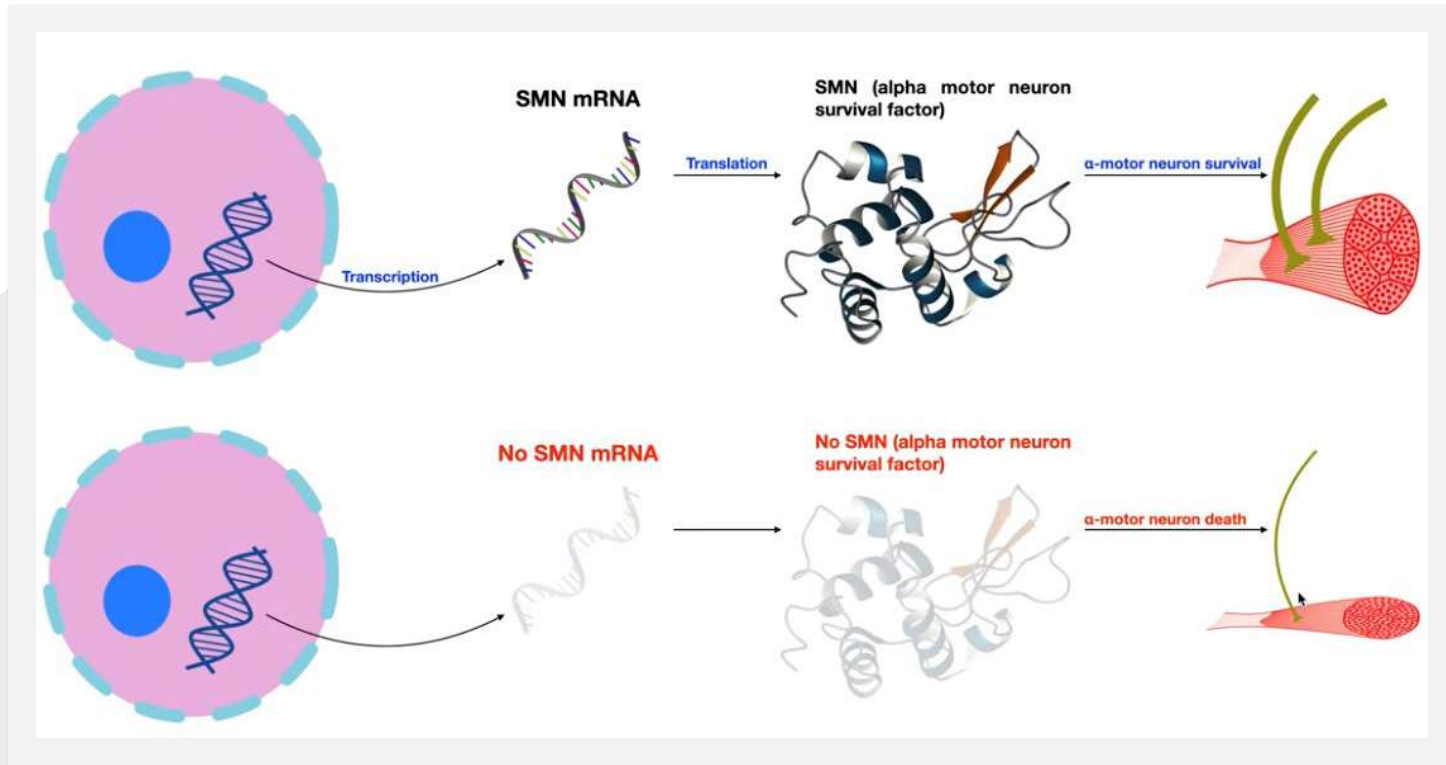
Typical Development



SMA Type 1/SMA+ Zolgensma

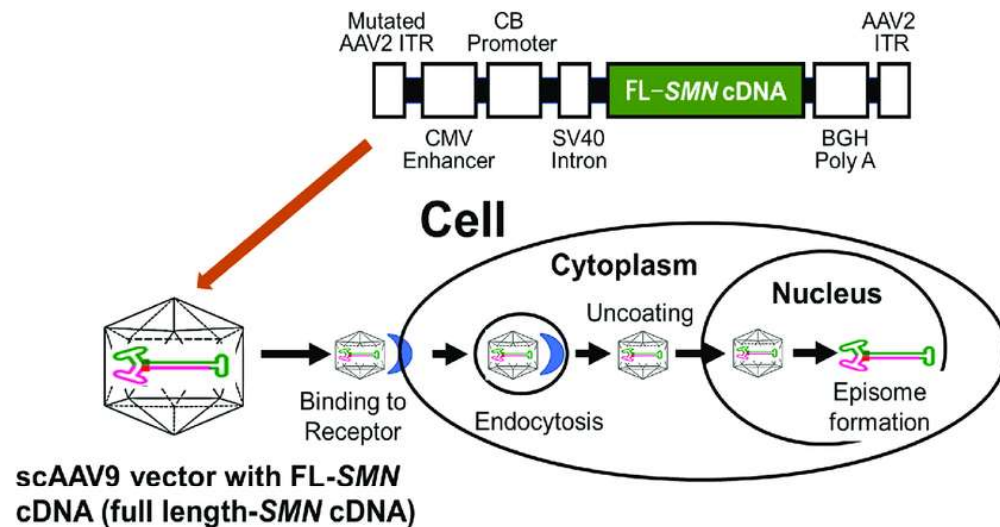
α -Motor Neurons



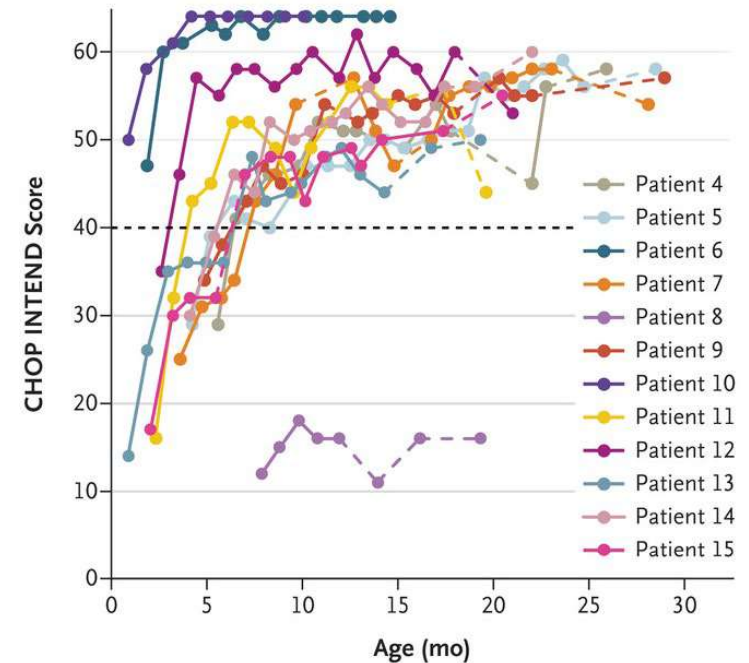


SMN1 protein promotes survival of α motor neurons

Gene therapy is capable of halting and reversing disability in SMA



PMID: 35606491



Historically, SMA Type 1 was a death sentence for children

Stage 1 (Birth to 6 months)

- Hypotonia, difficulty moving, difficulty feeding, shallow breathing

Stage 2 (6 months to 1 year)

- Regression of motor skills, loss of reflexes, respiratory issues, swallowing challenges

Stage 3 (1 year onward)

- Severe motor impairment, skeletal deformities, ventilation, food aspiration, infection

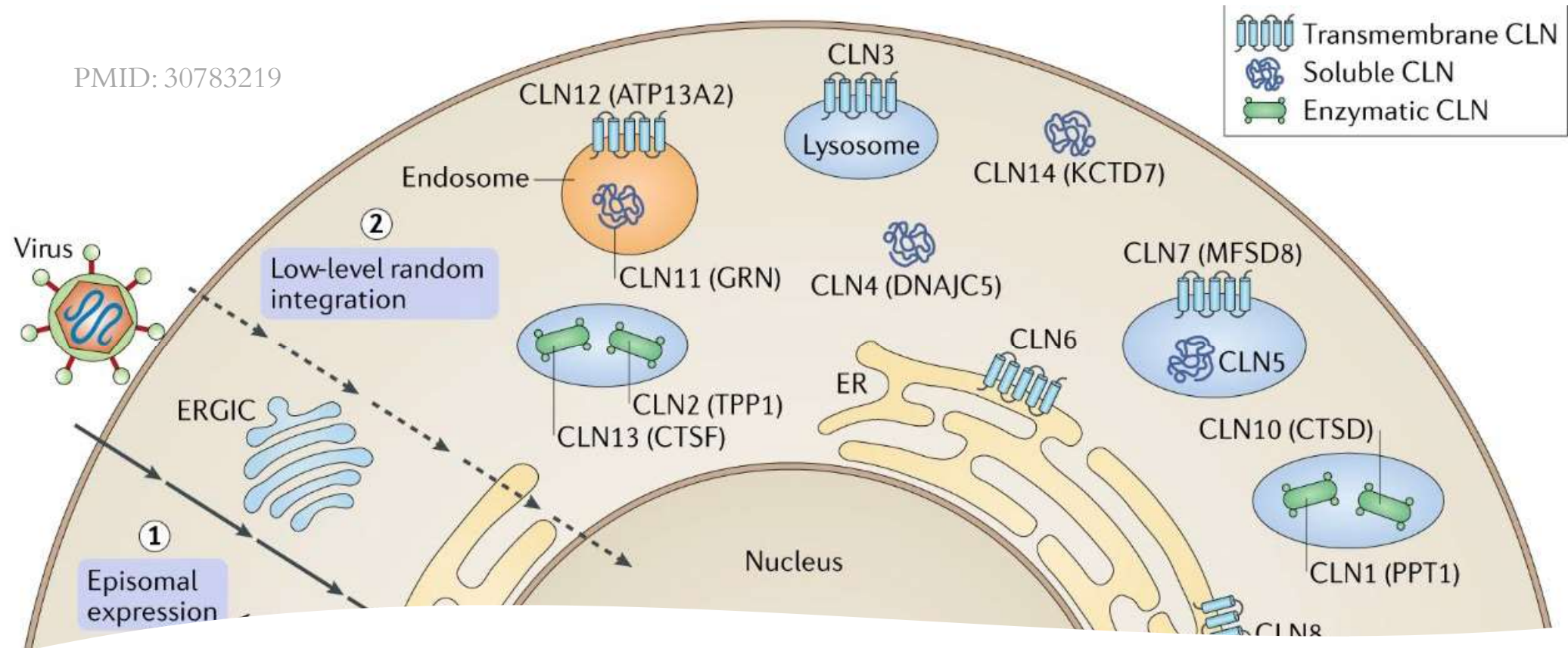
Death



Children with neuromuscular disorders who receive gene therapy may still struggle with:


- Scoliosis
 - Atypical curvature leading to discomfort/pain
- Joint contractures
 - Stiffness and reduced range of motion
- Hip dysplasia
 - Misalignment or instability of the hip joint
- Muscle weakness
- Fractures
 - Osteopenia, osteoporosis increasing risk of fractures

PMID: 30783219



Example: Batten's Disease

- Progressive accumulation of lipofuscins due to poor lysosomal clearance from neurons and other tissue
- Usually autosomal-recessive, caused by mutations to genes that encode proteins essential for lysosomal function (*PPT1*, *TPP1*, *CLN3*, *CLN5*, *CLN6*, *CLN7*, *CLN8*)



Personalized
medicine from
bench to
bedside

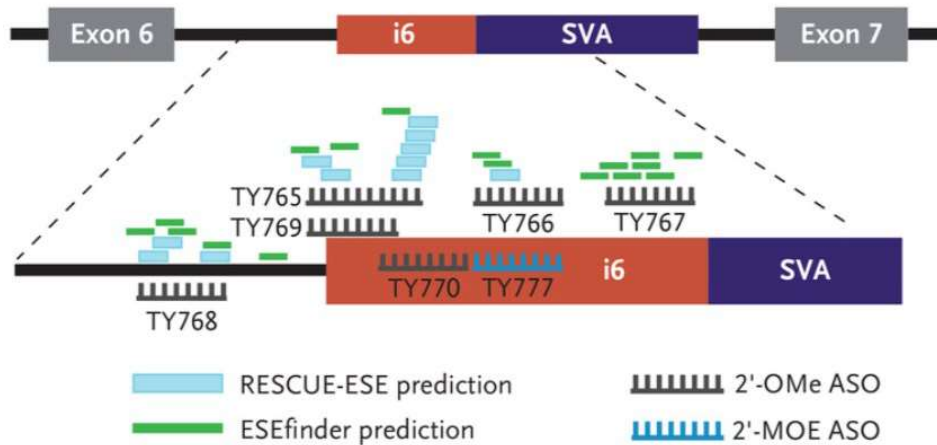




Mila, 2019

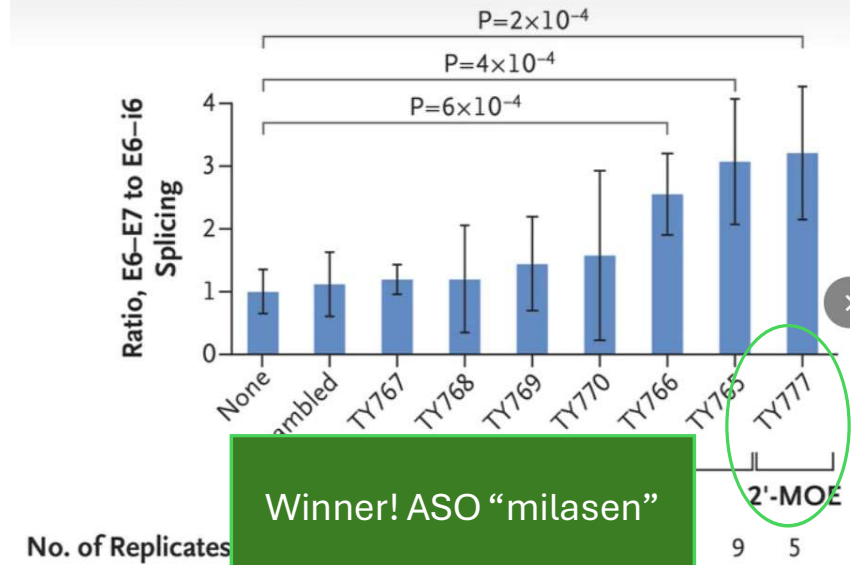
“N of 1” therapy in Batten’s Disease

- **Age 3:** Typical development
- **Age 4:** Vision difficulty
- **Age 5:** Language and social regression, clumsiness, stumbling
- **Age 6:** Rapid vision loss, falls, difficulty swallowing, motor speech disorder, epileptiform brain activity
- Discovery of loss-of-function *CLN7* variants



PMID: 31597037

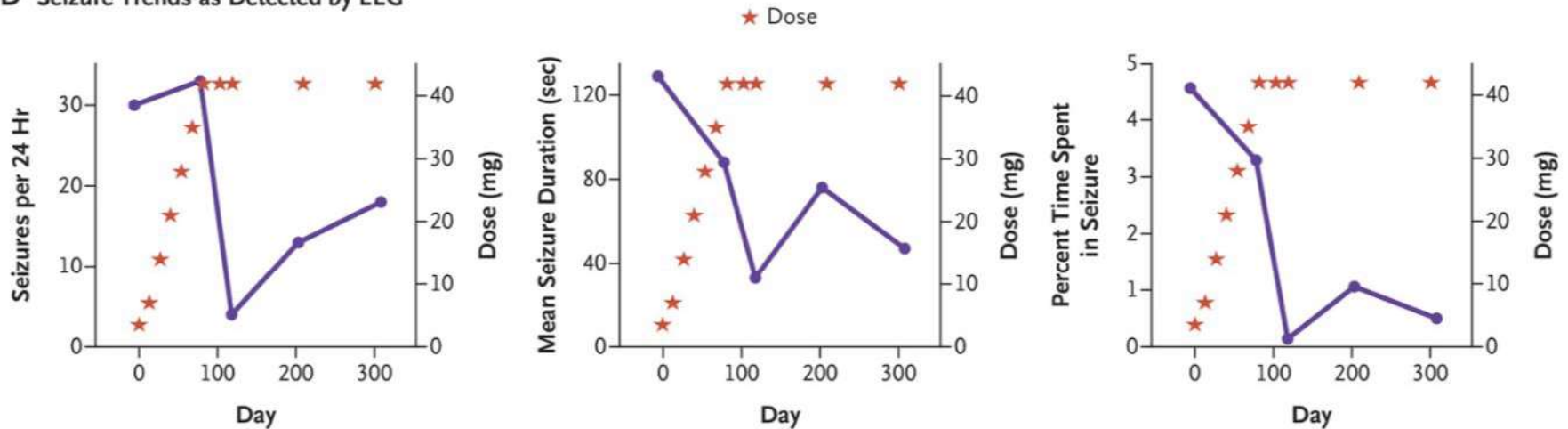
B Screen in Patient Fibroblasts



Bespoke antisense oligonucleotide therapy for CLN7 Batten Disease

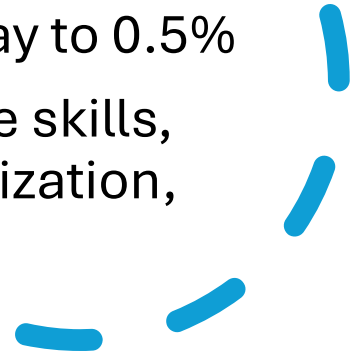
- The problem: therapeutics for Mila cannot be scaled to treat other CLN7 children due to the uniqueness of her variant

D Seizure Trends as Detected by EEG



Milasen reduced
Mila's seizures
over 200 days

- Mila's seizures were reduced from occurring for 5% of her entire day to 0.5%
- Some improvements in adaptive skills, such as communication, socialization, and daily living skills





Lessons learned from Mila

- Ultimately, focus on the brain lead to improvement of neurological symptoms, but not others
- Bespoke therapies are feasible, but costly
- ... at this present time



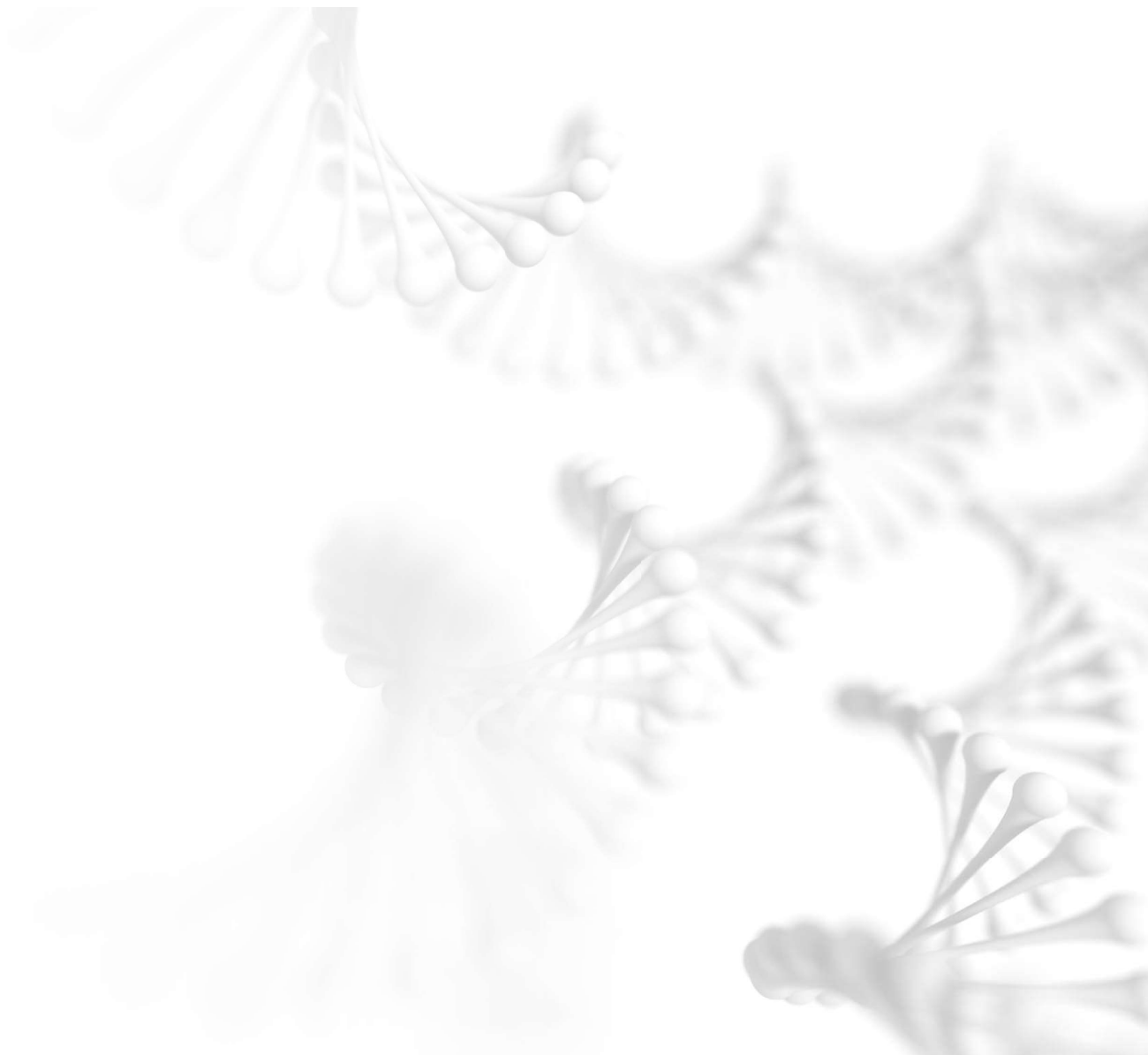


What's next? 5-year timeline

- Systematic prioritization of genetic subgroups in diseases of aging
 - E.g., LYS3884961 by Lilly/Prevail for Parkinson's patients with *GBA* variants
 - Gene editing in reprogrammed cell populations or localized to specific organs
 - EDIT-101 by Editas Medicine for blindness due to *CEP290* variant IVS26
 - CTX001 by CRISPR Therapeutics for gene-edited hematopoietic stem cell therapy due to *HBB* variants
 - Explosion of gene replacement therapeutic development for LoF variants
-



Gene editing in humans



The curious case of Lulu, Nana, and Amy

- Unpublished, shrouded in mystery
- Parents part of project involving HIV-positive fathers
- Twins conceived via IVF, additional child conceived via IVF
- Underwent CRISPR/Cas9-mediated germline deletion of 32 base pairs of the CCR5 gene
- Goal: confer resistance to HIV acquisition
- Questions regarding informed consent, scientific validity and priority, off-target effects, non-publication of results

Principal Investigator Dr. He Jiankui



The black box has been opened

- We are all living with medically-actionable, rare genetic variants
- Many of us will receive gene therapies within our lifetime
- It remains to be seen whether germline editing (heritable changes) will be ethically permitted

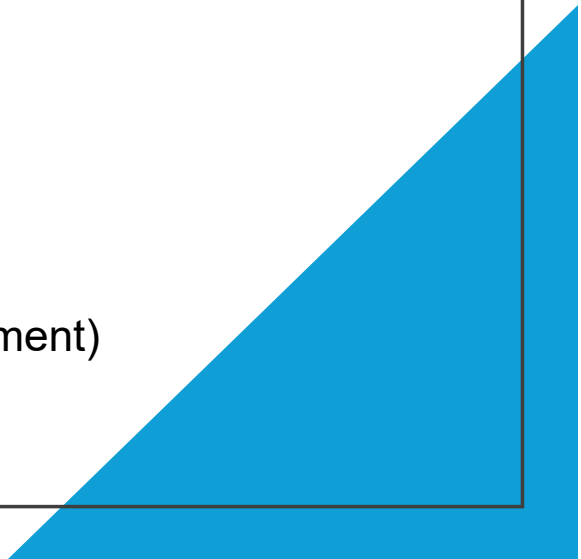




Impact of precision medicine on functional medicine

- Halted disease progression
 - Stability at advanced stages of disease requires ongoing care
 - Reversal of disability
 - Greater need for providers trained in rehabilitation
 - Expanded scope of practice + increased work in disciplinary teams
 - Greater need for geriatric and pediatric providers
 - Treatment of developmental disease and disorders of aging
-

Areas of expected change in functional medicine within your career

- Musculoskeletal issue focus in obesity recovery (Ozempic, *FTO* gene therapy)
 - Pediatric rehabilitation (neuromuscular disorders, neurodevelopmental disorders, epilepsies, cerebral palsy, mitochondrial disorders, lysosomal storage disorders)
 - Geriatric rehabilitation (neurodegenerative disorders)
 - Pain management demand in above populations
 - Fewer arthritis, migraine patients (gene therapies in development)
 - Greater development of subspecialties
- 

Thank you!



K.INTSON@UTORONTO.CA