## **Deep Genomics** Decode the genome, create life changing therapies

### **BRENDAN J FREY, PHD, FRSC**

CEO AND FOUNDER, DEEP GENOMICS COFOUNDER, VECTOR INSTITUTE FOR ARTIFICIAL INTELLIGENCE PROFESSOR, ENGINEERING & MEDICINE, UNIVERSITY OF TORONTO

### Deep Genomics team, Bocce Ball, October 2021

Multidisciplinary: Machine learning, computational biology, software engineering, experimental biology, chemistry, drug development, business development.





TORONTO moderna Somerck facebook Stanford Shire Google Min genzyme Spirer V Mettore Pharmacia COLOR

## Team in place to execute our mission









Ferd Massari MD Chief Medical Officer PHARMACIA 2 Pfizer



Matt Cahill MBA JD PhD Head of Finance, Bus. Ops







Sen Dir Predictive Systems Google TORONTO VECTOR INSTITUTE



Jeffrey Brown PhD Head of Preclinical Res.

Voyager WAVE Bristol Myers Squibb Strategic Advisors



**Steve Jurvetson** SPACEX

#### Scientific Advisors



Steven Dowdy



Stephen Scherer



#### **Board Members**



Adam D'Augelli true Ventures



Eric

Marcussen

Alex Morgan khosla ventures

**Jennifer Cook** 

Roche

GRAIL

Yann LeCun

facebook MYU 🥞 AT&T

Genentech

bridgebio

**Brendan Frey** Matt Cahill









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## RNA therapies: Medicines are digital information

CCCAAATGCACTCCTGG

## Digital RNA Biology Platform

10<sup>17</sup><sub>Bytes</sub> X Data

Artificial Intelligence

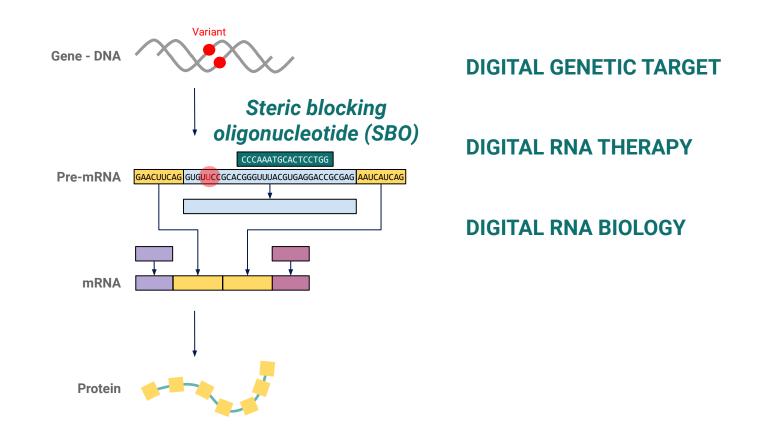


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Programming the best RNA therapies for almost any gene in any genetic condition

## A digital framework for untangling complexity





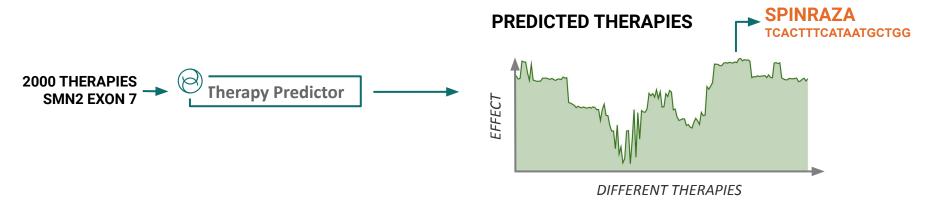
Discovering Spinraza for Spinal Muscular Atrophy - in one afternoon on a computer





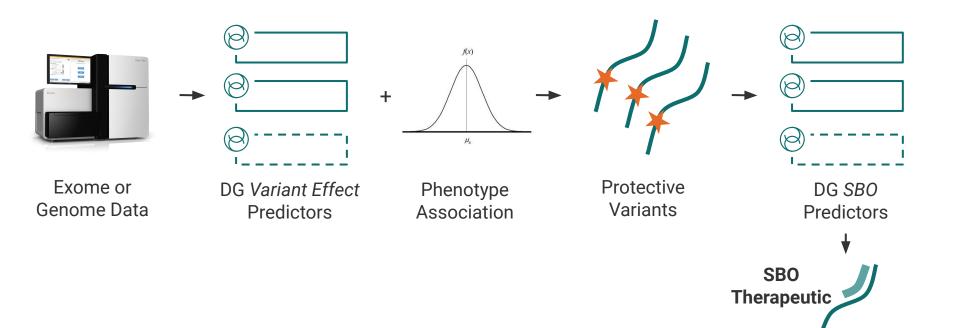
### PREDICTED EXON TARGETS

**Top 10% - SMN2 EXON 7** 10% to 20% 20% to 30% 30% to 40% 40% to 50%



## Discovering novel complex disease targets using exome or genome sequencing data

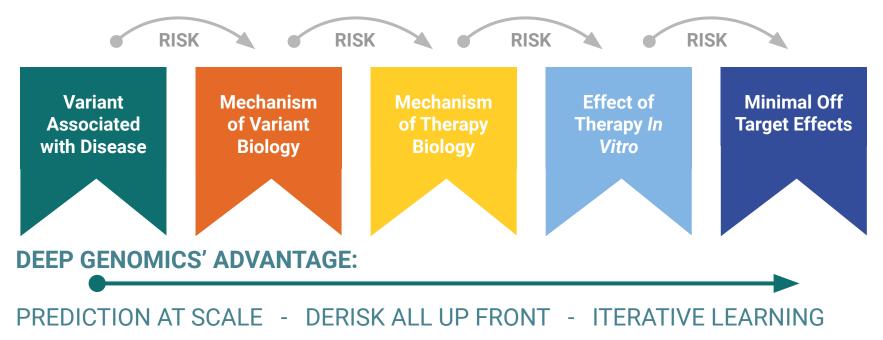




# Our digital AI Workbench predicts drug discovery outcomes up front

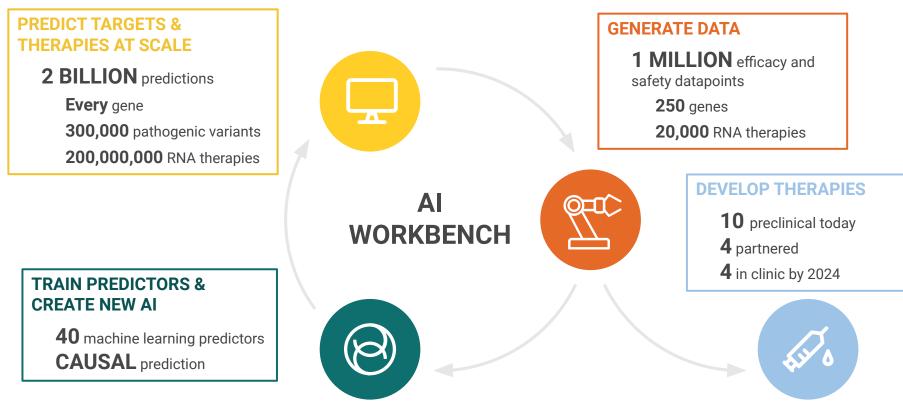


TRADITIONAL APPROACH - SEQUENTIALLY DERISK - EXPERIMENTAL TRIAL & ERROR - BESPOKE

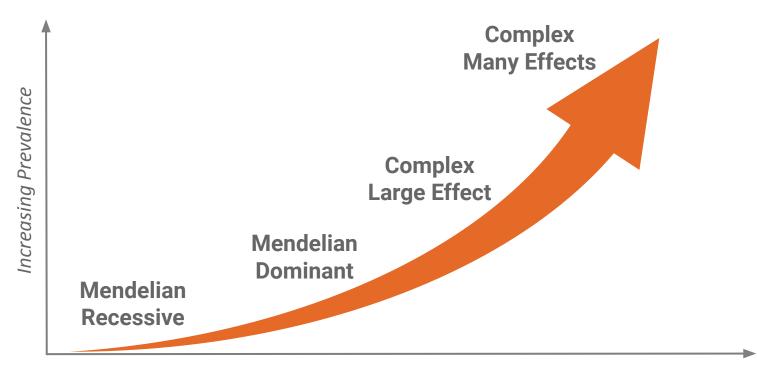


# Data driven prediction, positive feedback loops, and exponential growth





Tackling complexity - Our digital AI Workbench unlocks targets across the spectrum of genetic types



Increasing Biological Complexity

# Predictors drive portfolio productivity, and we have 40 of them

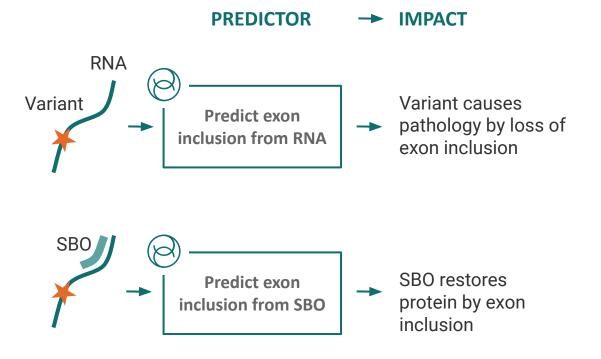


### **EXAMPLE APPLICATIONS**

- Identify novel targets
- Design therapies
- Predict safety and toxicity
- Identify new patient populations
- Predict drug properties

### **SBO PREDICTORS**

- Protein restoration
- Protein expression Increase
- Protein knockdown



# AI Workbench and "plug-and-play" SBO technology powers high-value, low risk therapies



### THERAPEUTIC AREA

#### CNS

Frontotemporal Dementia Niemann-Pick Disease Type C Pediatric Epilepsy Parkinson's Disease

#### **Metabolic**

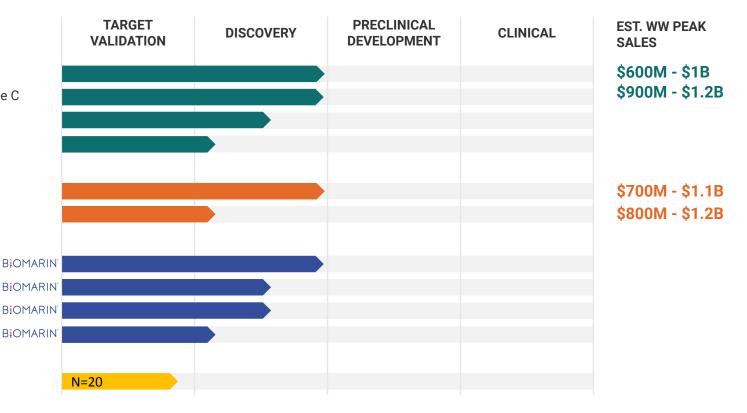
Wilson Disease Refractory Gout

#### Partnered

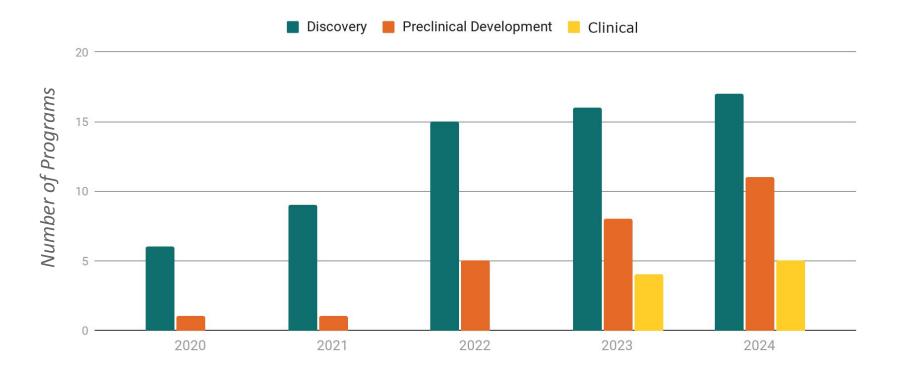
Undisclosed Undisclosed Undisclosed Undisclosed

#### **Target Identification**

Under Assessment



## DG Portfolio: At tipping point to significant clinical expansion



## Company goals to mid-2024



### PREDICTORS

Expand AI for complex disease, SBO effect and SBO safety

### **GENES SCREENED**

100

60 complex, 40 Mendelian Enabled by AI and robotics

28

## 80

### TARGETS WITH PATENTED LEADS

**Complex and Mendelian** 



### EXPAND PARTNERSHIPS

Broaden pipeline, access non-dilutive funding PRECLINICAL PROGRAMS Internal, partners, CRO

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PROGRAMS IN THE CLINIC



### Key question for Canadian genomics community Some big datasets are exponentially better than others - which datasets should public money be used to generate?

- 1. Good: Genome sequencing produces high quality data
- 2. Proof points from industry: Biotechs request sequencing money, demonstrate their capacity to use it, then GC signs a cheque with the understanding that the data is public.
- 3. Input from Industry: Council of Canadian Biotech Entrepreneurs a committee to help identify big genomics datasets of strategic value to Canada



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Thank You