



COLLABORATIVE  
**TRAJECTORY  
ANALYSIS  
PROJECT**

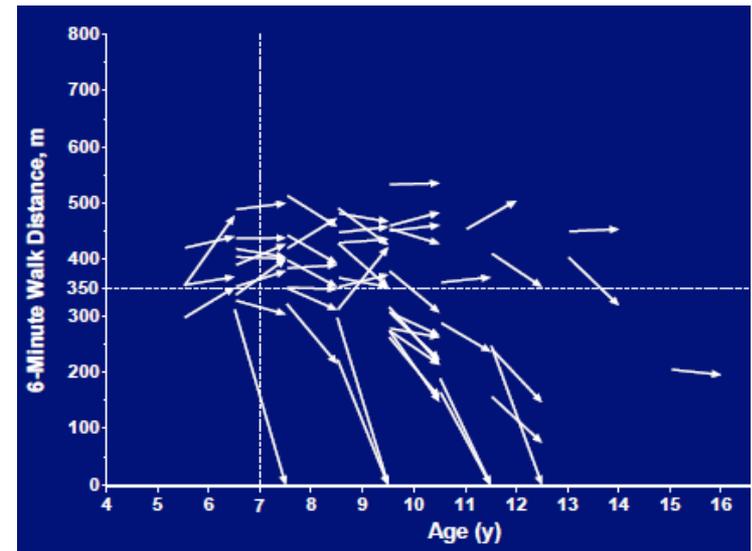
**Enabling the right trial design, the first time**

*Supporting new therapies to patients sooner*

# Impetus

***Pivotal trials in Duchenne have failed to meet primary endpoints – largely due to high patient heterogeneity***

<u>TRIAL</u>	<u>PHASE</u>	<u>PATIENTS**</u>	<u>Met end point?</u>
DEMAND III	Ph 3*	186	no
PTC 007	Ph 2*	174	no
DMD-ACT	Ph 3*	228	no
<u>Tadalafil</u>	Ph 3*	331	no



- Burden and disappointment for patients and families
- Uncertainty and wasted resources for drug development
- Disincentive for investors

# Natural History Landscape 2013/14

- Trial design based on thought-leader input
- No direct access to natural history data
  - Arduous learning curve for who has what data
  - Lengthy cycle-time for requested analytics
  - Significant duplication of effort to access the basics
- The first publication to show individual patient trajectories did not appear until late 2013

# **cTAP mission**

**Collaborate to learn from patient data**

**Adopt successful approaches from other fields**

**Create tools and insights for drug development**

**Share broadly**

**Deliver near-term impact**

# cTAP Members and Collaborators

<b>Clinical experts and registries</b>	<ul style="list-style-type: none"><li>• Eugenio Mercuri; Catholic University, Rome</li><li>• Nathalie Goemans; UZ Leuven</li><li>• Francesco Muntoni; University College London</li><li>• Brenda Wong; Cincinnati Children's Hospital</li><li>• Fondazione Telethon</li><li>• Northstar UK</li></ul>
<b>Sponsors</b>	<ul style="list-style-type: none"><li>• BioMarin</li><li>• Pfizer</li><li>• PTC Therapeutics</li><li>• Sarepta Therapeutics</li><li>• Shire PLC</li><li>• Solid Biosciences</li><li>• Catabasis</li><li>• BMS</li></ul>
<b>Patient advocates</b>	<ul style="list-style-type: none"><li>• Cure Duchenne</li><li>• Parent Project Muscular Dystrophy</li></ul>
<b>Analytic and Collaboration leads</b>	<ul style="list-style-type: none"><li>• Analysis Group</li><li>• Susan J. Ward</li></ul>

# What we do

- Access longitudinal data from de-identified patients in natural history and placebo arm studies
- Apply statistical methods to explain phenotypic variation and reduce variance
- Publish findings and share broadly (including with regulators)
- Embed in tools to optimize trial design and analysis
- Share costs: data access, analytics, publication, collaboration platform

# Progress: Data Access

**1,261** boys

**33** care centers

**10,508** clinic visits

## **Functional Assessments**

Ambulation > **4000**

Pulmonary, Cardiac > **350**

boys  $\geq$  3 yrs follow-up > **400**

## **Patient Characteristics**

steroid status, history

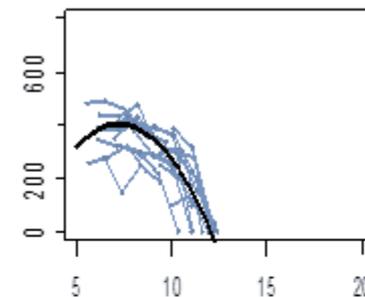
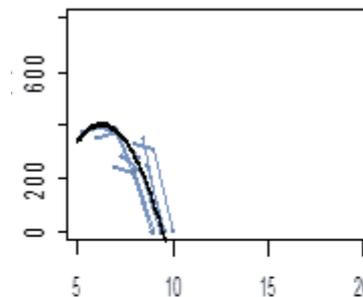
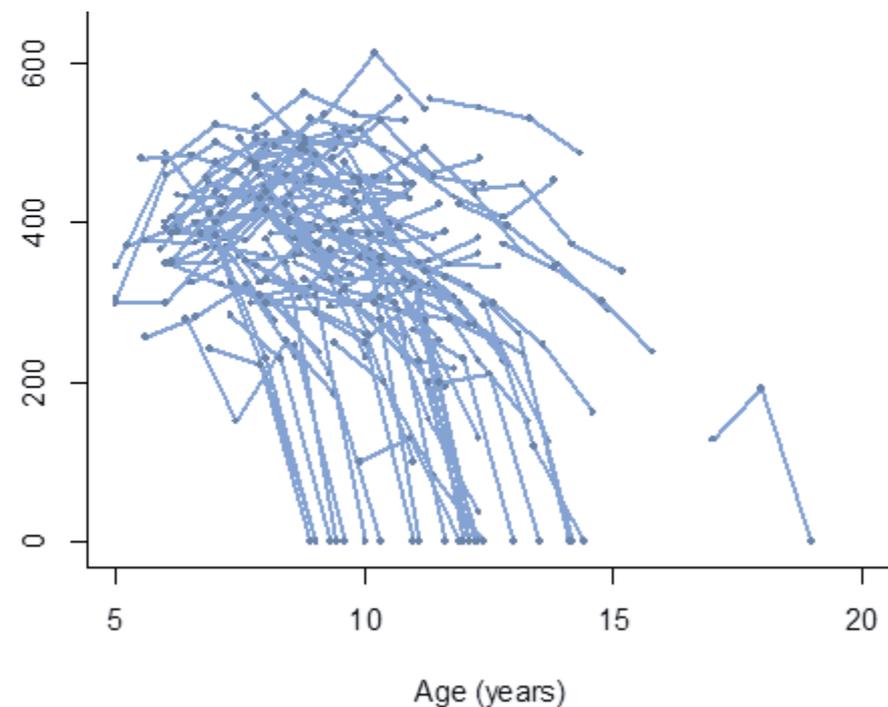
dystrophin genotype

body composition (incl. fat)

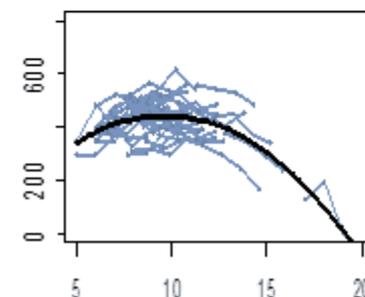
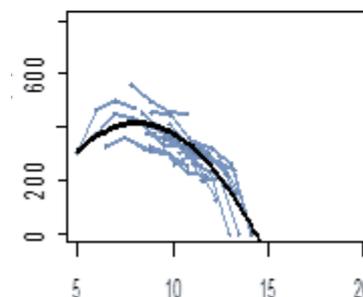
- **Largest clinical database in Duchenne**
- **Majority of data previously never shared**
- **Semi-annual updates**
- **Attractive Data Use/Business structure**
- **New Data, in progress**
  - New data fields: PUL, NSAA sub-scores
  - Large EU cohort
  - Placebo Arms from pivotal trials

# Progress: Analytics (1) Structure

## Walking distances in 6 minutes (m)

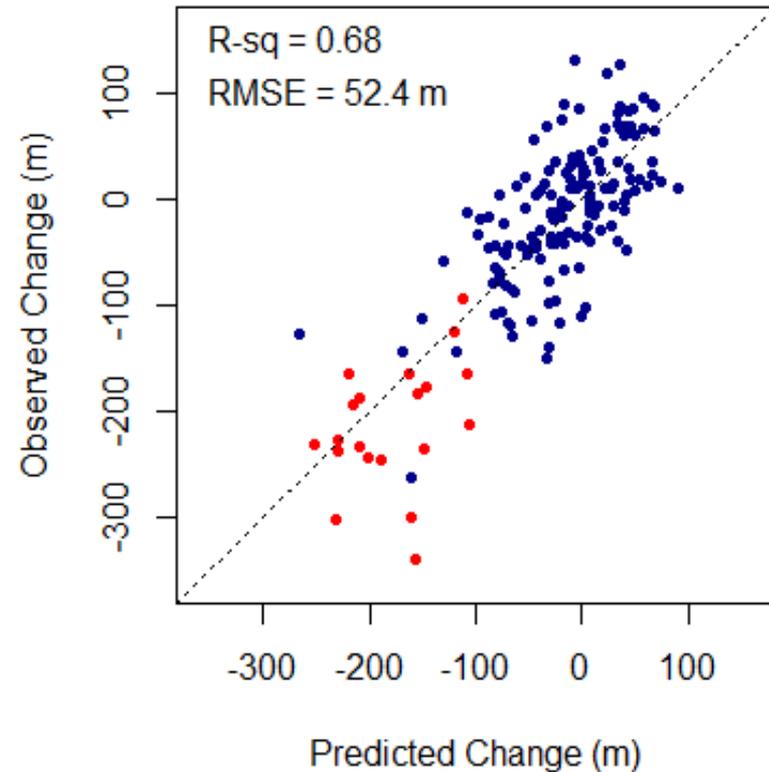
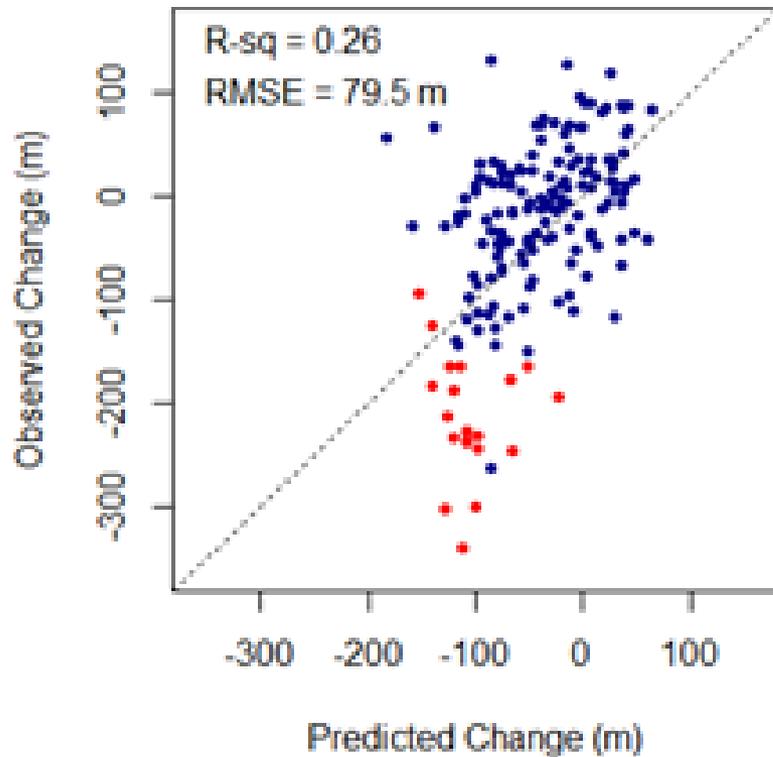


*~ 50% decrease in variance*



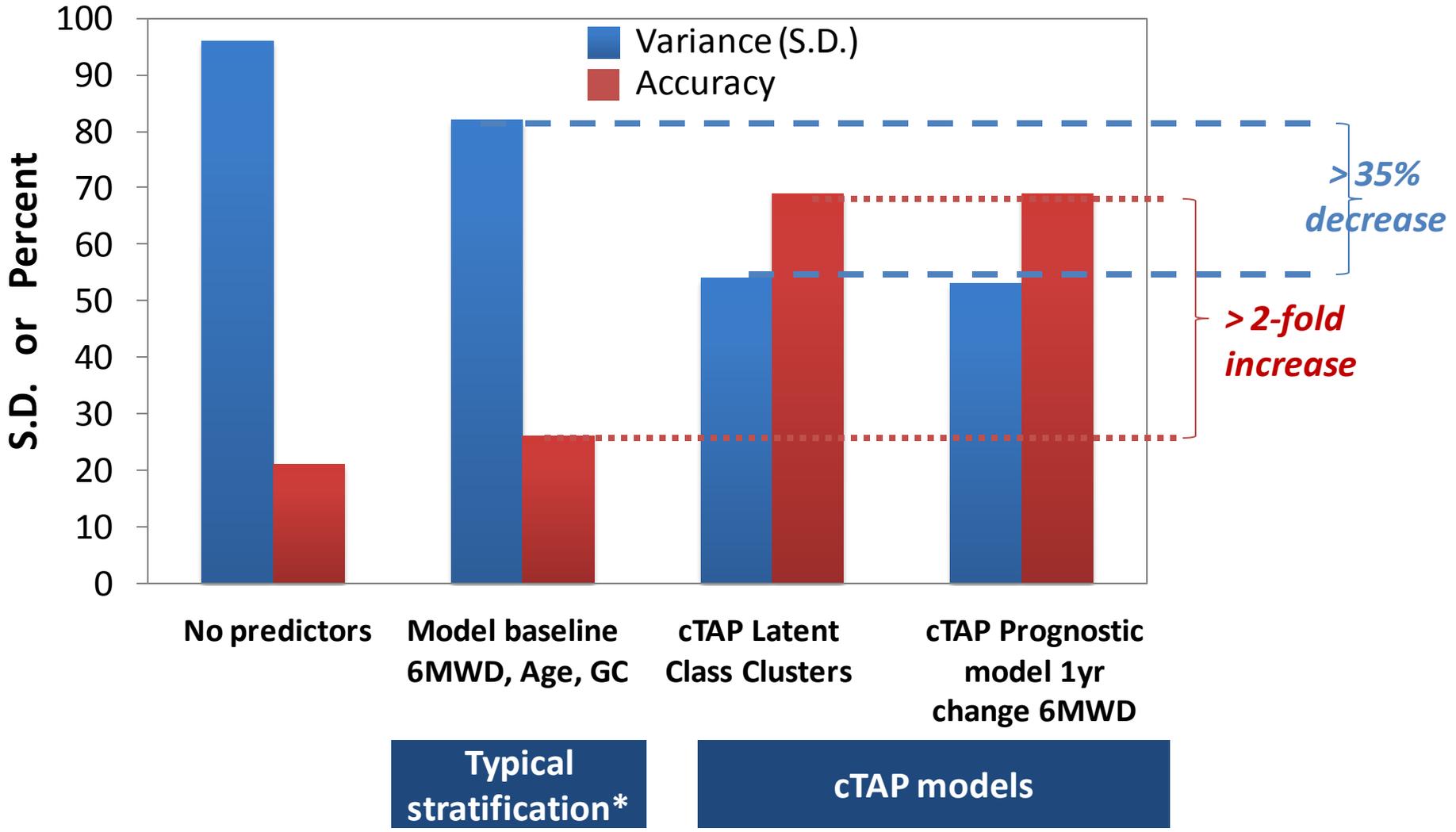
Eugenio Mercuri, et al., 2016. Categorizing natural history trajectories of ambulatory function measured by the 6-minute walk distance in patients with Duchenne muscular dystrophy. *Neuromuscular Disorders* 26 (2016) 576–583 doi: 10.1016/j.nmd.2016.05.016

# Progress: Analytics (2) Prognostic Model



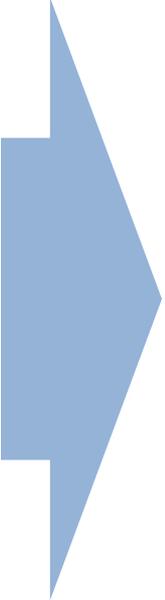
Nathalie Goemans et al., 2016. Individualized prediction of changes in 6- minute walk distance for patients with Duchenne muscular dystrophy. *PLoS One*. 0164684 . doi.org/10.1371

# Summary: cTAP models markedly reduce variance of change in 6MWD above and beyond typical stratification strategies

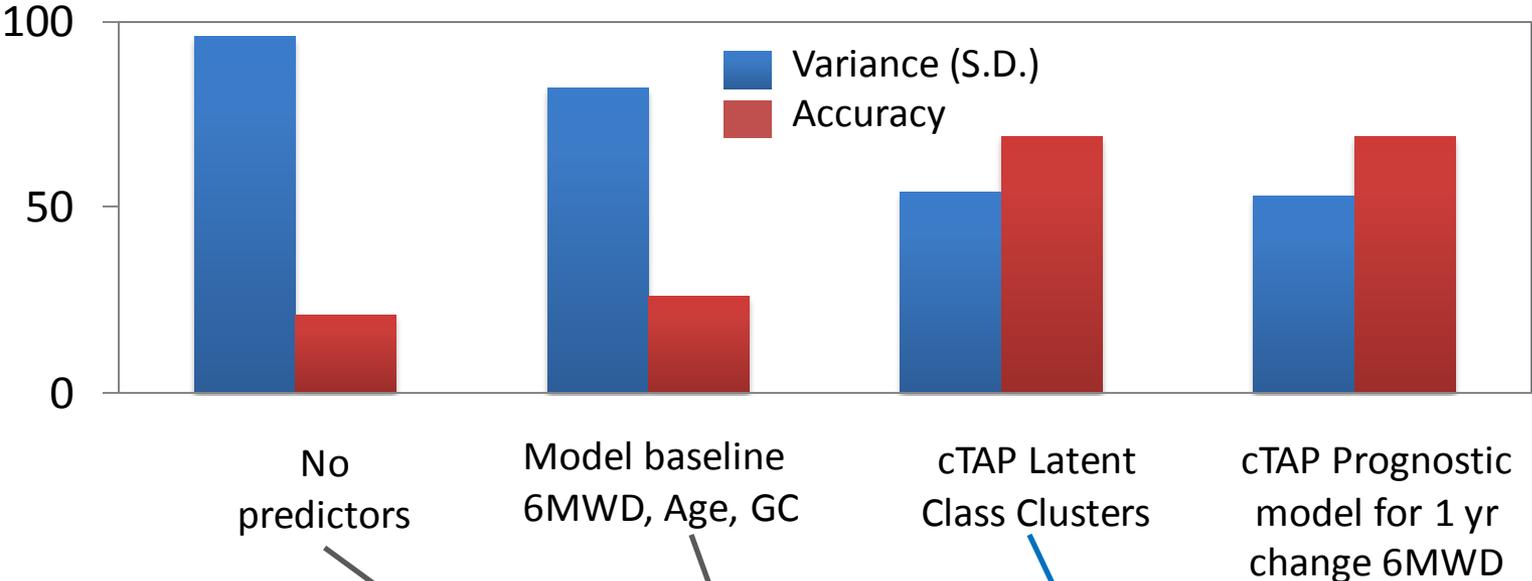


# Translation to Drug Development

## *Describe, Predict, Simulate*

- 
- Inform trial design and analysis
  - Enable natural history controls
  - Inform biomarker evaluation
  - Establish value of endpoints for regulators and payers

# Impact on Confidence in Trial results



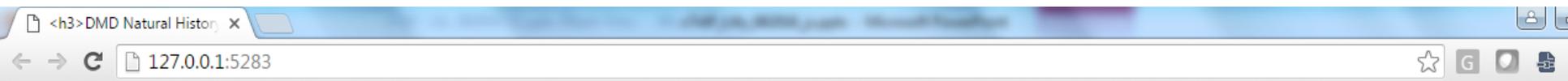
unexplained variation (m)	100	90	80	70	60	50	40
approx power (%)	60%	69%	79%	89%	96%	99%	100%

*Marked impact on power, over and above 'standard' adjustments*

# Tools: Duchenne Discovery portal

- Real-time, flexible, dynamic analytics
- User defined
  - outcome measure
  - segmentation parameters
- Computes
  - Patient characteristics at baseline
  - 1 year change in outcome measure
- Modularized additions
  - Correlations between outcome measures
  - Interactive per patient trajectory

# Illustration: Summarize patient characteristics



## DMD Natural History Explorer

### Outcome measure

NSAA total score

### Inclusion criteria

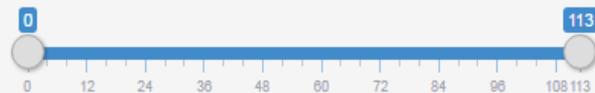
#### NSAA total score



#### Age (years)



#### Duration of steroid use (years)



Sample selection

Baseline characteristics

Follow-up time

Outcomes

Definitions

View

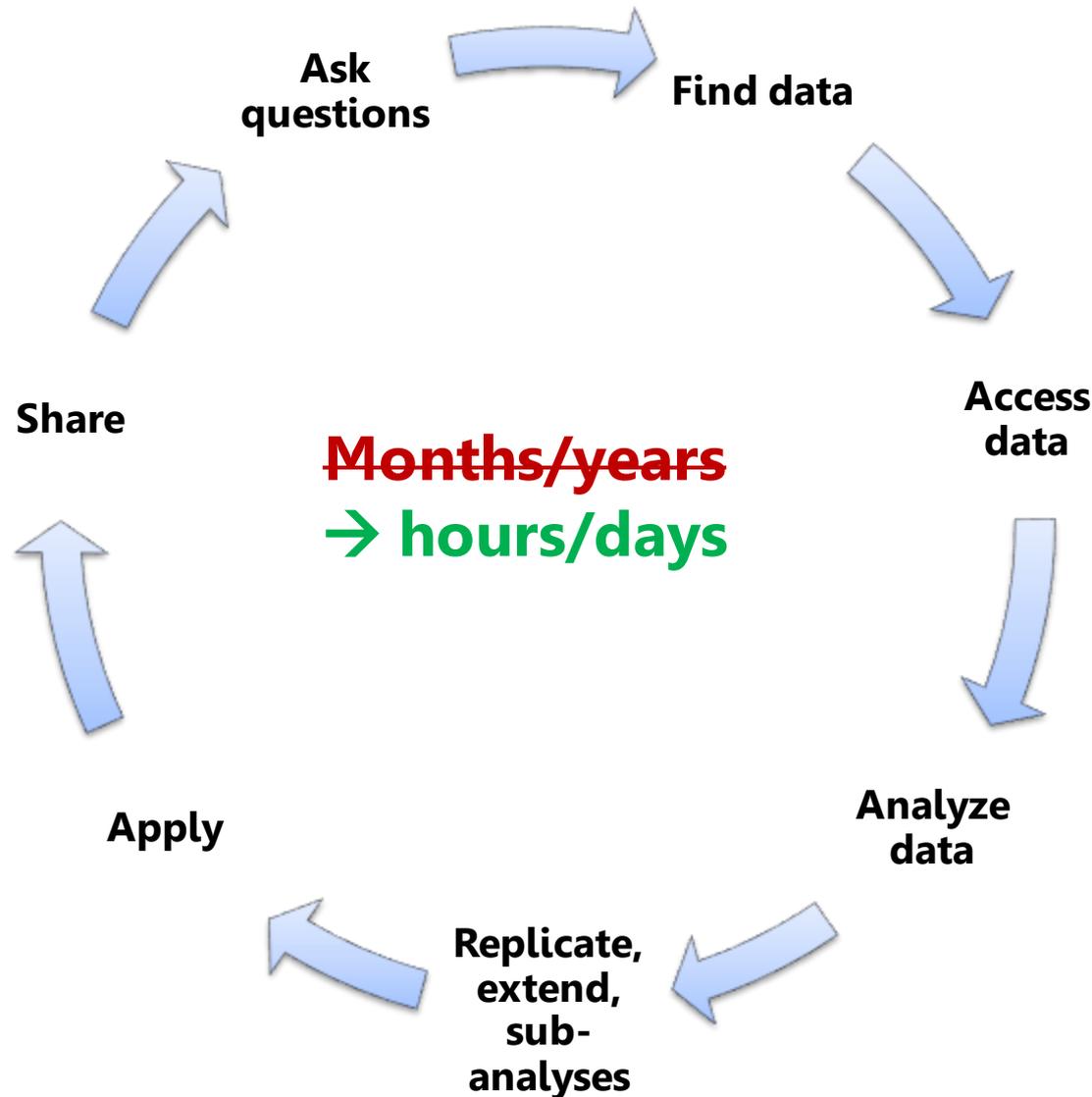
Configure

More details

Variable	Included patients (n=369)
Age, years	8.3 ± 1.5
Duration of steroid use, years	2.1 ± 1.5
NSAA total score	23 ± 7.5
Linearized NSAA score	63 ± 17.7
10MWR, seconds	6.7 ± 2.7
Rise from supine, seconds	6.5 ± 5.4
Dystrophin mutation type	
Deletion	223 (60.4)
Duplication	33 (8.9)
Other	16 (4.3)
Point mutation	46 (12.5)
Unknown/missing	51 (13.8)

Means and standard deviations are shown for continuous characteristics; counts and percentages are shown for categorical characteristics, unless otherwise noted

# Portal renders laborious data/analytic access redundant



- Who has data?
- What study measures, sample sizes are available?
- Is the data quality adequate?
- What is the process for data access (if one exists)?
- Contract & data use agreement negotiation
- Scientific plan / SAP
- Risks of delay or failure due to: limited responsiveness, hidden agendas, politics, etc.

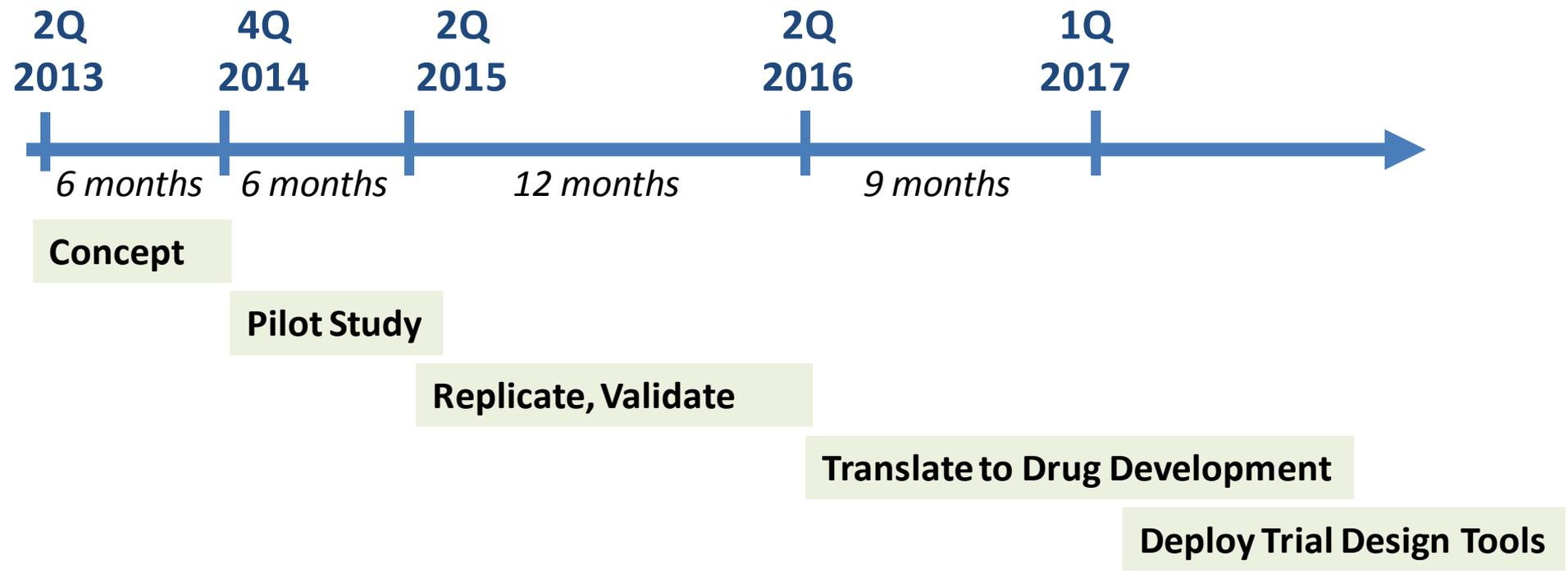
# cTAP Benefits to drug developers

- Access analytics on previously inaccessible data
- Get timely answers to systematic questions that drive trial design
- Tangible, practical support to accelerate academic understanding of heterogeneity in Duchenne natural history
- Collaborative access to best in class analysis solutions
- Insights and tools for deployment in drug development
- Shared benefits, shared Costs
- Modular research approach – pay only for your company's priorities
- Credibility of consortium structure with regulators
- Publish and communicate broadly to benefit entire community
- **New members** enjoy immediate access to all unpublished findings, collaborative insights and new tools

# Opportunities for Collaborating Clinical Experts & Registries

- Contribute to a data collaboration that benefits patients ... while retaining full control of data use and authorship
- Collaborate with peers to quickly share & replicate across multiple databases
- Collaborate with Pharma with efficiency and balance
- Unprecedented access analytics to accelerate research
- Author more publications, strengthen grant applications
- Receive funding support data collection
- Access tools to aid data cleaning

# Rapid Progress



# Objectives 2016-2017

- Extend clustering and prognostic modeling beyond 6MWD to other end-points
- Demonstrate consistency (or lack thereof) between natural history databases, clinical sites, and placebo arms
- Tie near term functional changes to long-term outcomes
- Establish matching criteria to support historical controls
- Determine prognostic value of glucocorticoid history
- Assess cost of clinical events and milestones
- Conduct Open Workshop(s); include regulators
- Continue to share findings promptly



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## Contact Us

**Susan J. Ward, Ph.D.**

Co-Founder and Collaboration Lead

Executive Director

The TAP Collaboration

[info@ctap-duchenne.org](mailto:info@ctap-duchenne.org)

**James Signorovitch, Ph.D.**

Co-Founder and Analytic Lead

Vice President

The Analysis Group Inc.

[James.signorovitch@analysisgroup.com](mailto:James.signorovitch@analysisgroup.com)