## **SPARK**

Accelerating Research and Improving Outcomes in Autism Spectrum Disorder

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Igniting autism research Improving lives

## Significant Challenges in Developing Effective Treatments for ASD

- Heterogeneity in symptoms and etiology (roughly 40% genetic/60% environmental)
  - Some progress in genetics, but much more to learn
  - Some progress in environmental factors, but largely unknown
- How are symptoms linked to etiology?
- Objective measures required for translational and clinical research
- Only treatments are behavioral and resource-intensive



## How are symptoms linked to etiology?





## **Objective measures needed**





### More efficient treatments needed

- Only available treatments are behavioral and resource-intensive
- 10 years of intensive intervention = 20,000 hours of 1:1 therapy = \$1m
- How much impact do such interventions have on long-term outcome?



## Research efforts so far, but much more data needed

- Etiology
  - Genetics on 18K individuals with autism so far → 60 genes, but hundreds left to discover
  - Some progress in environmental factors, but largely unknown, perhaps up to 60% of cases
- Heterogeneity
  - Genetics-first approaches to study by genetic subtype
- What behaviors are proximal to etiology?
  - Genetic animal models of autism: what happens at a cellular and behavioral level?
- Objective measures required for translational and clinical research
  - EU-AIMS
  - Asdbiomarkers.org
  - Targeted grants in this area by SFARI

SPARK

## SPARK must scale





## What SPARK will do for individuals and families affected with autism

- Participants are partners, not subjects
- Commitment to return individual genetic results related to autism to medical professional that participant designates
- Return individual results on standardized behavioral questionnaires
- Inform community of aggregate results
- Opportunities to interact with autism experts online



# What SPARK will do for the research community

- Full understanding of the genetic architecture of autism
- Enable more efficient recruitment (genotype-based if desired) to entire research community at no cost
- All behavioral and genetic data will be made accessible to any qualified researcher
- Some embargoes on entire genetic dataset but data will be released quickly after it is generated



## **Genetic analysis in SPARK**

- Exome sequencing of trios to accelerate gene discovery
- Sequencing of unaffected siblings when possible
- State of the art genome-wide genotyping for GWAS
- SPARK genomics consortium will analyze first 10,000 families



## What's next for SPARKforAutism.org

- Wrapping up pilot phase (recruitment of 500 trios)
- National launch on April 21
- Release data in scheduled releases beginning in Q3, 2016
- Open cohort to recruitment by research community in 2016
- Recruit and collect specimens from at least 10,000 trios through April 2017
- Report on pilot and genetic analysis of first 10,000 families



#### The SPARK Team: From Exomes to Twitter



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