



Genomic sequencing of newborns?

Presented to the Advisory Committee on Heritable Disorders in
Newborns and Children

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The Promise

“I think it won’t be too many years before parents will be able to go home from the hospital with their newborn babies with a genetic map in their hands that will tell them, here’s what your child’s future will be like.”

Bill Clinton, *Tallahassee Democrat*, 10/11/1996, A12

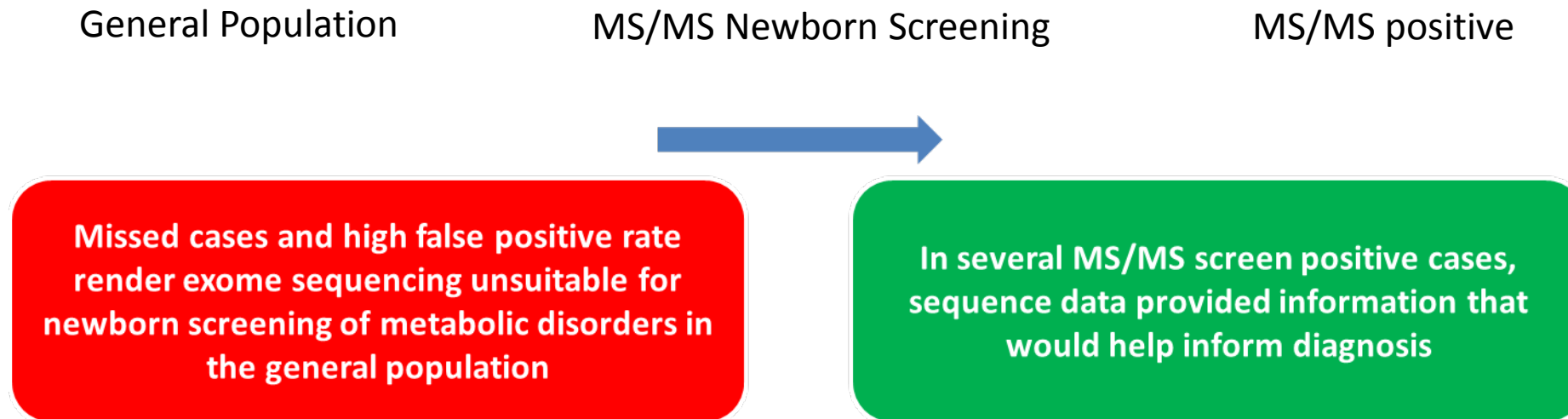
The hope

- Sequencing will yield
 - A correct diagnosis
 - That hadn't been already made
 - For a treatable disease
 - And treatment will lead to a better outcome.
- And will do this consistently and cost-effectively
- In broader and broader populations

Preliminary Conclusions from
UCSF study of newborn
bloodspots from the California
Biobank.

Presented at ASHG 2017

WES not recommended as a STAND ALONE tool for primary public health NBS for IEMs



....But DNA may play key role as second-tier test
depending on the disorder, the gene(s), the variant(s)...

A Spectrum of Sequencing in NBS

Aaron Goldenberg



Unique Features of NBS

- Blood spot collection is “unconsented”
- Legal justification transformed
- Although this varies from state to state, samples are stored in the state of California “biobank” available for research

Public Health Context

- Sequencing would identify numerous conditions that do not meet the legal and ethical justification for state mandated screening, and for which states cannot provide follow-up care
 - Preservation of screening programs is critical for public health and equality

Public Health Context

- However, sequencing could be used as a secondary or adjunct tool for detecting conditions that meet traditional newborn screening criteria.
 - Additional consideration needed regarding:
 - Return of secondary results
 - Storage of data

Concerns raised by state NBS programs

- Workforce and costs
- Education and communication
- Incidental findings
- Impact of private companies, potential to drive implementation
- Burden on parents
- Impact on original intent of public health NBS

Aaron Goldenberg, 2017 ELSI Congress

General Conclusions

- Thus far, WES has only been shown to be useful in clinical populations where diagnostic uncertainty is a barrier to good care.
- There is not (yet?) evidence that sequencing every newborn would be sufficiently beneficial to children or families, to justify using it in routine care, public health, or as a DTC service.

General Conclusions: Remaining Questions

- Are our recommendations simply a result of lack of “data,” and thus time bound?
- Will accumulation of evidence “solve” the dilemmas of sequencing newborns?



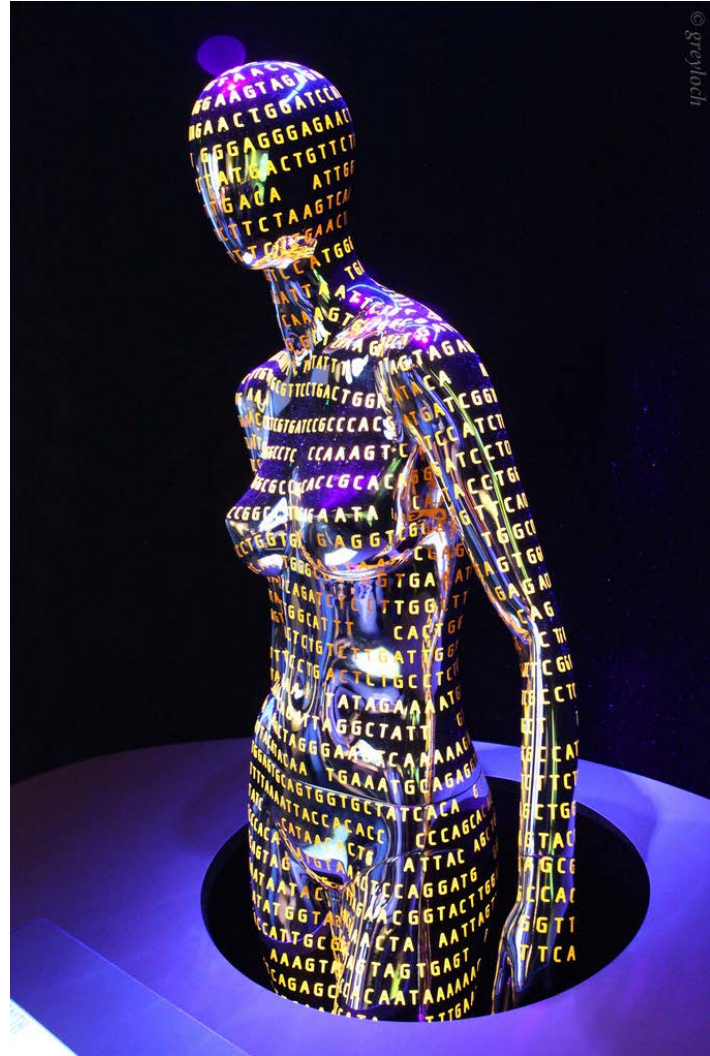
***How is this test different
from other tests?***

Why is this test different?

- Uncertainty of findings
- Interpretation requires broad data sharing
- Return of secondary (unexpected) findings

Why is this test different?

Data, Data, Everywhere



Data, Data, Everywhere

- What should be returned?
- What is “actionable?”
- Managing “variants of uncertain significance”



CORRIGENDUM: Recommendations for reporting of secondary findings in clinical exome and genome sequencing, 2016 update (ACMG SF v2.0): a policy statement of the American College of Medical Genetics and Genomics

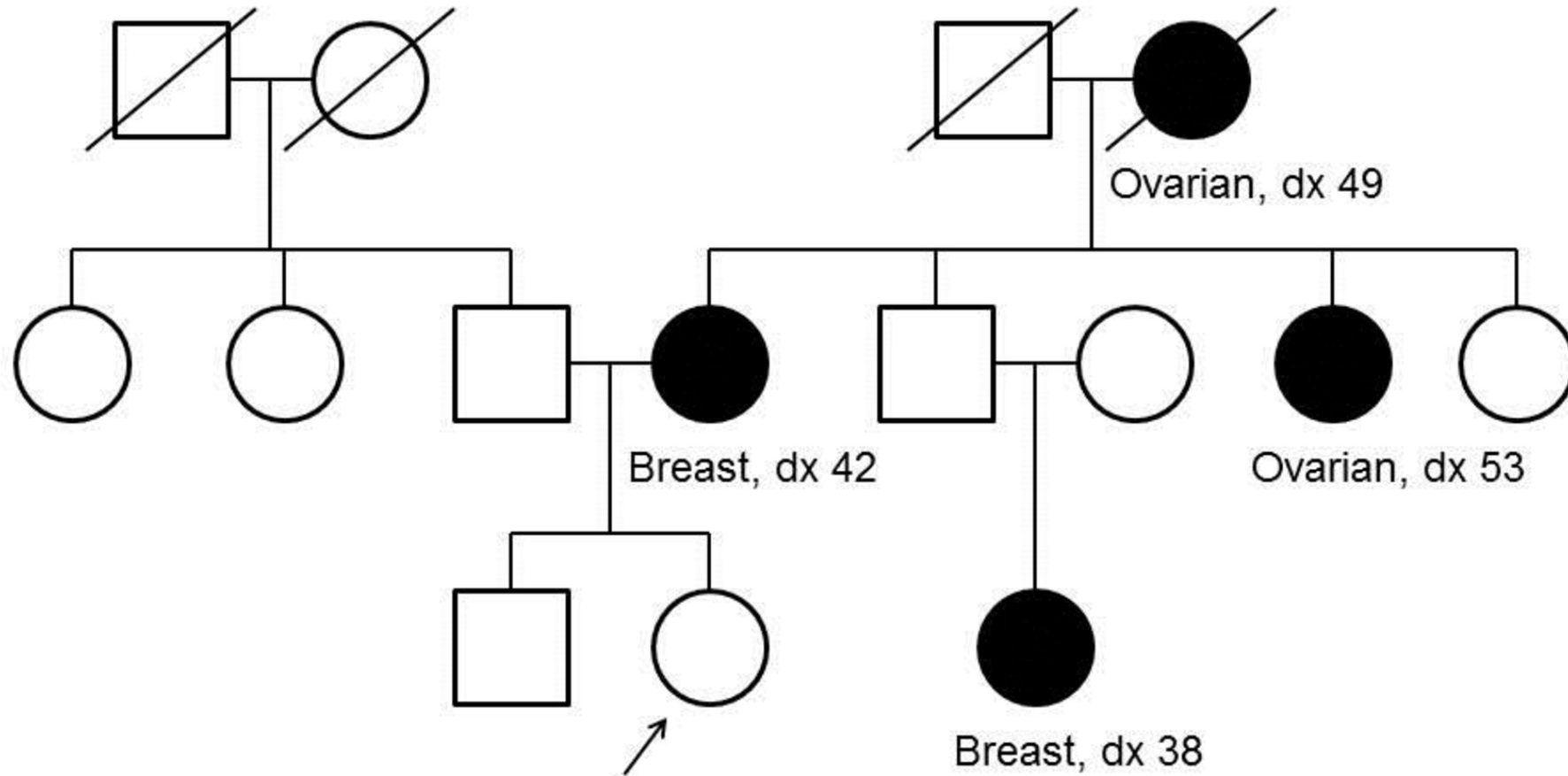
Genetics in Medicine (2017) 19, 484 | doi:10.1038/gim.2017.17

Why is this test different? Family





Classic *BRCA1* Pedigree



An ethical dilemma

- A newborn in the NICU with undiagnosed anomalies is sequenced
- A “known pathogenic” variant in BRCA1 is identified, Sanger confirmation reveals maternal inheritance
- What should the team do?
 - Research?
 - Clinical?

An ethical dilemma (adult onset conditions)

- Don't return
 - Respect autonomy of child
 - Right to an open future
 - Protect child from psychosocial harm
- Return
 - Obligation to benefit affected relative (mother)
 - Professional integrity
 - Health & life of parent is in child's best interest

What if this same variant were identified in the context of state-mandated or expanded newborn screening?

Why is this test different? Data Sharing

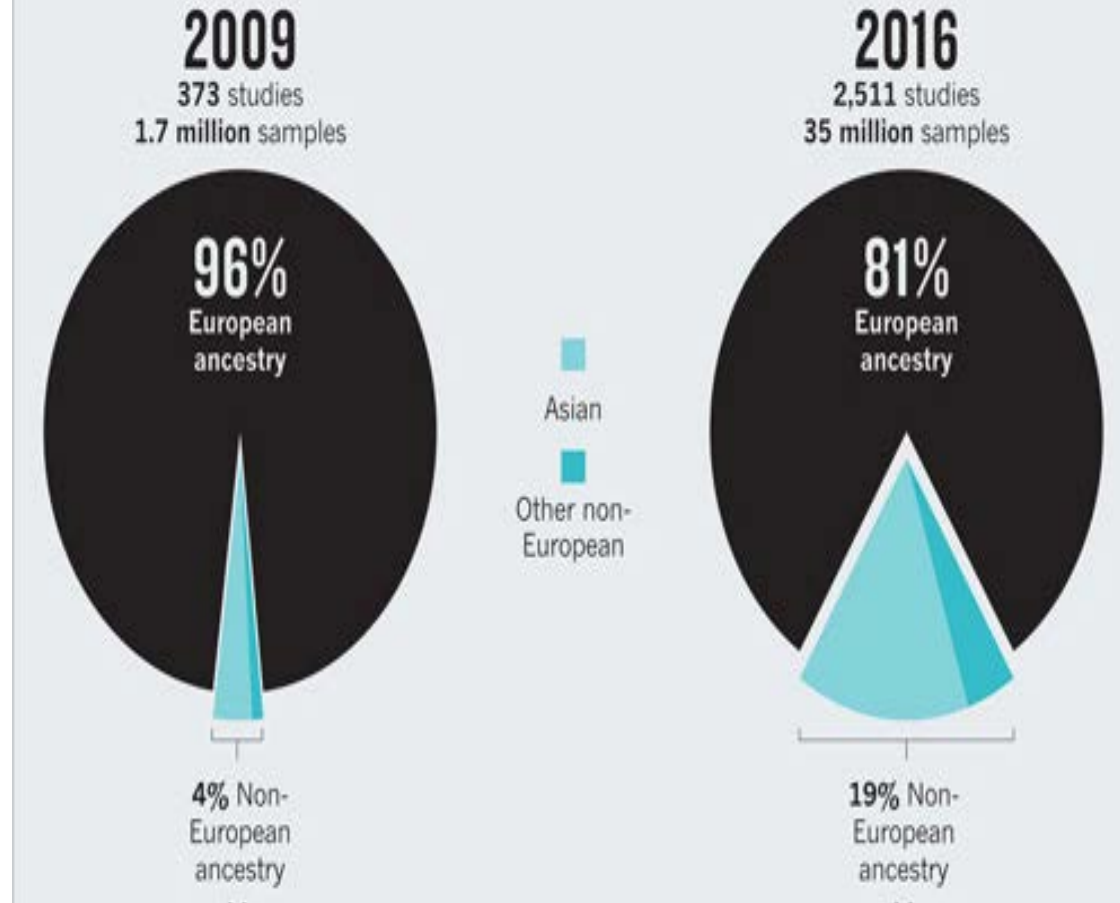


Data Sharing

- Variants can only be understood when compared with “reference” data bases, which can only work if data are broadly shared
- Uncertainty in interpretation (particularly for ancestrally diverse populations)
- Transformation in the demarcation between research vs. clinical care

PERSISTENT BIAS

Over the past seven years, the proportion of participants in genome-wide association studies (GWAS) that are of Asian ancestry has increased. Groups of other ancestries continue to be very poorly represented.



Popejoy, Alice B., and Stephanie M. Fullerton. "Genomics Is Failing on Diversity." *Nature* 538, no. 7624 (October 12, 2016): 161–64

Critical Need for Robust Community Engagement!

Social Justice: Insurance Coverage

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ARTICLE | **Genetics
inMedicine**

Private payer coverage policies for exome sequencing (ES) in pediatric patients: trends over time and analysis of evidence cited

Michael P. Douglas, MS¹, Stephanie L. Parker², Julia R. Trosman, PhD³, Anne M. Slavotinek, MD⁴ and Kathryn A. Phillips, PhD⁵

Social Justice & Access

- Study is the first in-depth review of private payer coverage policies for pediatric patients with neurodevelopmental disorders.
- In 2015, no payer (of 5) covered WES, but by 2017, three payers from the original registry payers did. And in 2017, 8 of the 15 payers covered WES.

Further Considerations

- Will identification of rare disorders not currently screened for by state NBS programs advance our understanding of conditions currently not recommended on the RUSP?
 - A difficult question!
 - Yes, but.

Further Considerations

- Eugenics Redux (sidebar by Diane Paul)
- Implications of adding “reproductive benefit” as a rationale for NBS
 - Moves beyond the best interest of the child
 - Will this be considered “state-sanctioned eugenics?”

Guiding Questions

- Which contextual forces shape our discussion of the utility of sequencing in newborns?
- Under what circumstances should newborns be sequenced?
- How should state-mandated newborn screening programs use sequencing?
- What role should parents play in determining how sequencing information about their infant is used and stored?
- Should sequencing be part of routine pediatric practice?