



60 Years of Newborn Screening in a Nutshell



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Learning Objectives

- **Demonstrate a basic understanding of newborn screening (NBS)**
- **Identify some differences between NBS programs**
- **Recall available tools to react appropriately to abnormal NBS results**

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Newborn Screening

A public health program:

- Aimed at identification of conditions for which early intervention can prevent
 - mortality
 - morbidity
 - disabilities
- Performed by analysis of diagnostic markers in blood spots collected on filter paper on the second day of life (exception: hearing loss and congenital cyanotic heart disease)

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Newborn Screening Pioneers

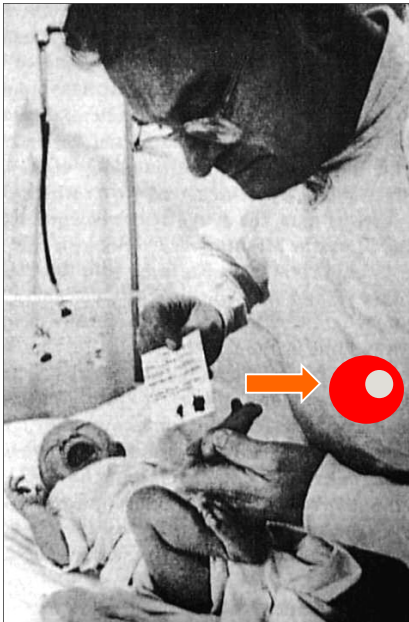


Horst Bickel
1918-2000

Robert Guthrie
1916-1995

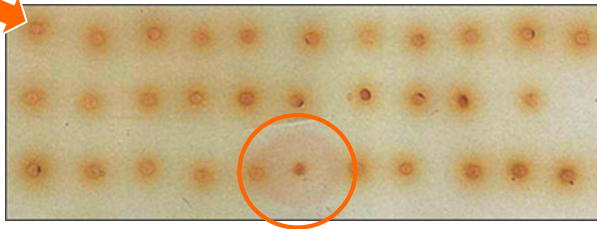
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Newborn Screening: The Early Years



1958: Bacterial inhibition assay (BIA) for PKU

1961: Newborn screening for PKU



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History of Newborn Screening

- 1962 JFK promotes a 20-state trial of the “Guthrie test”
- 1963 - Massachusetts mandates NBS for PKU
- Oregon adds Galactosemia to NBS for PKU

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The Traditional NBS Model: BIA

- One disease
- One test 
- One marker
- One cut-off (N/Abn)



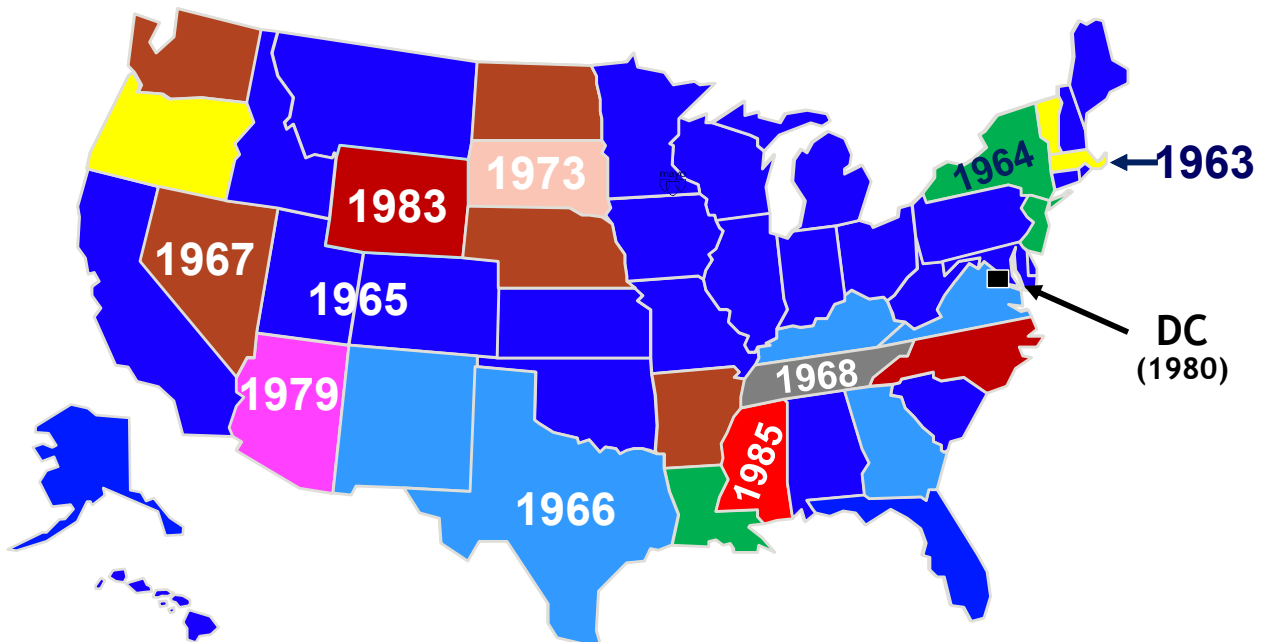
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History of Newborn Screening

- 1962 JFK promotes a 20-state trial of the “Guthrie test”
- 1963 MA mandates NBS for PKU; OR adds Galactosemia
- from 1963 NBS extends to all 50 states and includes metabolic and non-metabolic disorders

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Newborn Screening Implementation in the USA



adapted from: Therrell BL, Adams J. *J Inherit Metab Dis.* 2007; 30: 447-65 with addtl info from Kathryn Tullis (DE) and Sydney Williamson-White (VT).

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History of Newborn Screening

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- 1967 Massachusetts begins urine NBS using paper chromatography; stopped in 1991. Quebec began 1971, continues (in Sherbrooke)
- 1989 McCabe* et al describe **molecular genetic analysis** as a 2nd tier test for sickle cell disease screening

*Baylor College of Medicine, Houston, TX

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MCAD Deficiency

First described by Kolvraa et al in 1982

Incidence: ~1 : 10,000 live births

Gene: ACADM (1p31) (common mutation 985A→ G)

Symptoms: - Hypoketotic hypoglycemia
 - Reye-like syndrome
 - Sudden unexpected death

Treatment: Avoidance of fasting, IV glucose during stress

Prognosis: - Excellent when treated before onset of symptoms
 - 30-50% of mortality during first acute episode

Diagnosis: Acylcarnitine profile by tandem mass spectrometry (MS/MS)

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Tandem Mass Spectrometry Systems (MS/MS)



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Application of MS/MS to NBS

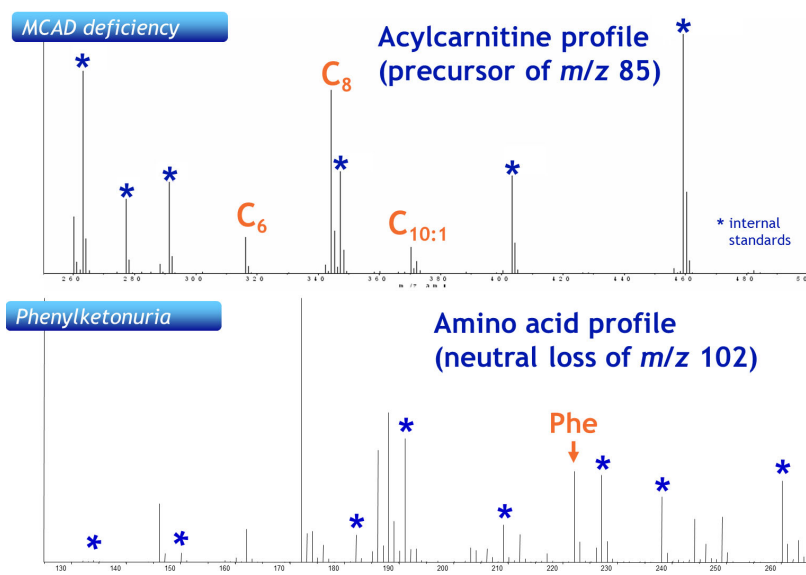
■ Primary screening (multiplex platform)

- Acylcarnitines, amino acids, succinylacetone, creatine*, creatinine*, guanidinoacetate*
- Lysosomal enzyme activities, lysophosphatidylcholines[^]
- (Bile acohols for Cerebrotendinous xantomathosis, bile acids for Niemann-Pick C disease, hemoglobin for Hemoglobinopathies)

*for creatine deficiency disorders
[^]for X-adrenoleukodystrophy (ALD)


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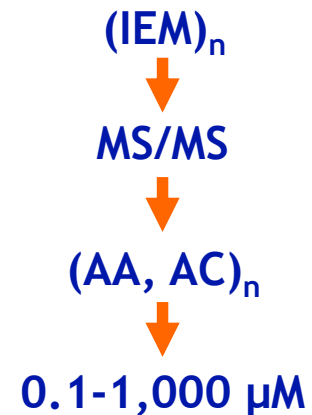
NBS by MS/MS



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A New NBS Model: MS/MS

- Many conditions
- One test 
- Many markers
- Many cut offs



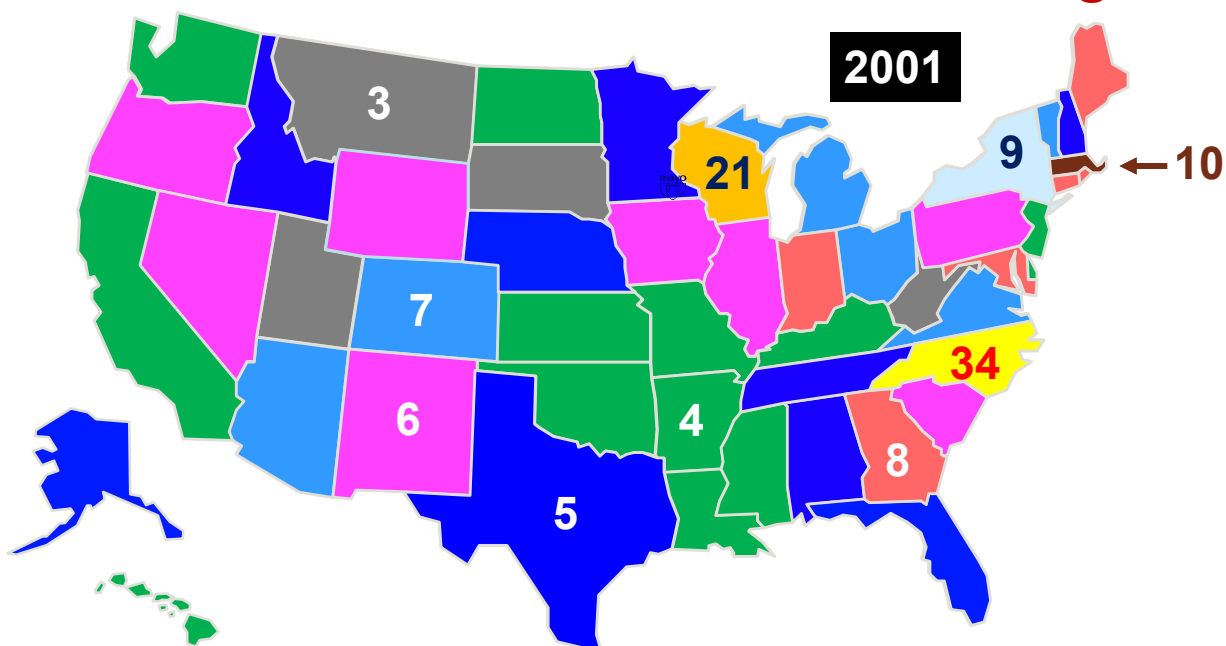
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- 1994: Molecular genetic analysis applied to CF screening (2nd tier test)
- 1996: Naylor starts using MS/MS for NBS in a private lab (Neogen)

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Number of Conditions included in NBS Programs



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Expansion of Newborn Screening: Uniform Screening Panel

May 2006 • Vol. 8 • No. 5, Supplement

executive summary

Michael S. Watson, PhD, Marie Y. Mann, MD, MPH, Michele A. Lloyd-Puryear, MD, PhD, Piero Rinaldo, MD, PhD, and R. Rodney Howell, MD, editors



The Maternal and Child Health Bureau commissioned the American College of Medical Genetics to outline a process for the standardization of outcomes and guidelines for state newborn screening programs and to define responsibilities for collecting and evaluating outcome data, including a recommended uniform panel of conditions to include in state newborn screening programs. The expert panel identified 29 conditions for which screening should be mandated. An additional 25 conditions were identified because they are part of the differential diagnosis of a condition in the core panel, they are clinically significant and revealed with screening technology but lack an efficacious treatment, or they represent incidental findings for which there is potential clinical significance. The process of identification is described, and recommendations are provided. *Genet Med* 2006;8(5, Supplement): 1S-11S.

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Newborn Screening Criteria

1968 - WHO (Wilson & Jungner)

- Treatable illness
- Detectable in newborn period
- Presymptomatic initiation of treatment is beneficial
- Available resources for diagnosis/treatment/follow-up
- Availability of a simple method for sample collection
- Evidence of substantial public benefit & acceptance
- Suitable and simple test methods
- Acceptable costs

2006 - ACMG Criteria

- Clinical characteristics (e.g., incidence, burden of disease if not treated, phenotype in the newborn)
- Analytical characteristics of the screening test (e.g., availability, features of the platform)
- Diagnosis, treatment and management of the condition in both acute and chronic forms (includes the availability of health professionals experienced in diagnosis, treatment, and management)

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ACMG Panel: Final Scoring

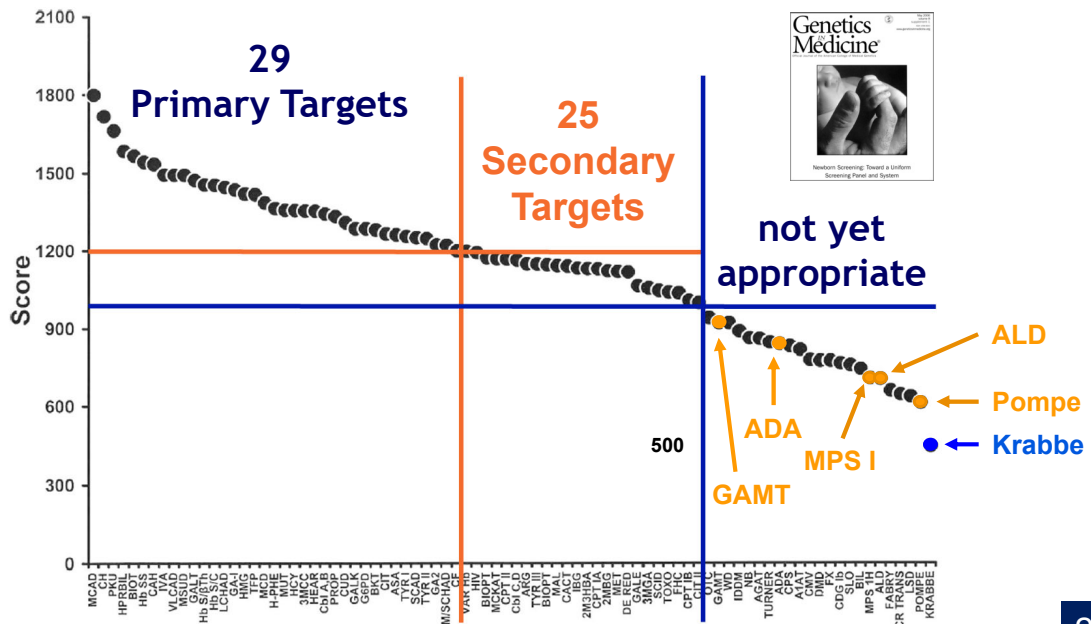


Fig. 1. Scoring by test availability (separates out those conditions that have an acceptable, validated, population-based screening test from those that do not).

2006

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ACMG's Recommended NBS Panel

29 Core Conditions

25 Secondary Targets

Table 2
Newborn screening panel: core panel and secondary targets

MS/MS					SECONDARY TARGETS				
Acylcarnitines		Amino acids			6 OA	8 FAO	8 AA	1 Hb Pathies	2 Others
9 OA	5 FAO	6 AA	3 Hb Pathies	6 Others					
CORE PANEL									
IVA	MCAD	PKU	Hb SS*	CH	Cbl C,D*	SCAD	HYPER-PHE	Var Hb*	GALK*
GAI	VLCAD	MSUD	Hb S/ β Th*	BIOT	MAL	GA2	TYR II		GALE
HMG	LCHAD	HCY*	Hb S/C*	CAH*	IBG	M/SCHAD	BIOPT (BS)		
MCD	TFP	CIT		GALT	2M3HBA	MCKAT	ARG		
MUT*	CUD	ASA		HEAR	2MBG	CPT II	TYR III		
3MCC*		TYR I*		CF	3MGA	CACT	BIOPT (REG)		
Cbl A,B*						CPT IA	MET		
PROP						DE RED	CIT II		
BKT									

NOTE: Codes are as follows: OA, disorders of organic acid metabolism; FAO, disorders of fatty acid metabolism; AA, disorders of amino acid metabolism; Hb Pathies, hemoglobinopathies.
* Identifies conditions for which specific discussions of unique issues are found in the main report.

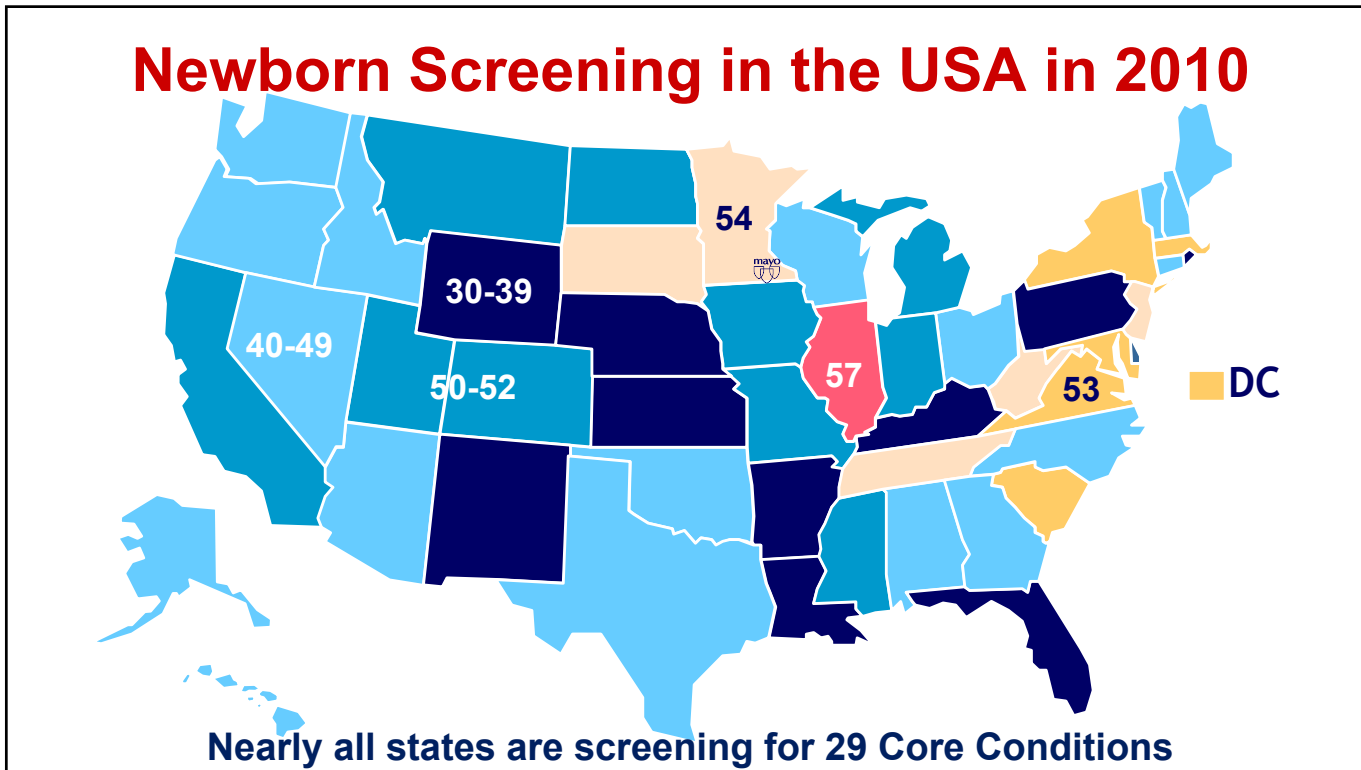
Watson MS et al. *Genet Med.* 2006; 8(5, Suppl 1): 1S-252S

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- 1993: Chace et al describe NBS for PKU using MS/MS
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- 1996: Naylor starts using MS/MS for NBS in a private lab (Neogen)
- 2005: ACHDNC recommends adoption of ACMG recommended screening panel
- 2010: HHS Secretary adopts ACMG recommendation as "Recommended Uniform Screening Panel" (RUSP)

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HRSA's ACHDNC

https://www.hrsa.gov/advisory-committees/heritable-disorders

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- [Newborn Screening Timeliness Goals](#)
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Advisory Committee on Heritable Disorders in Newborns and Children

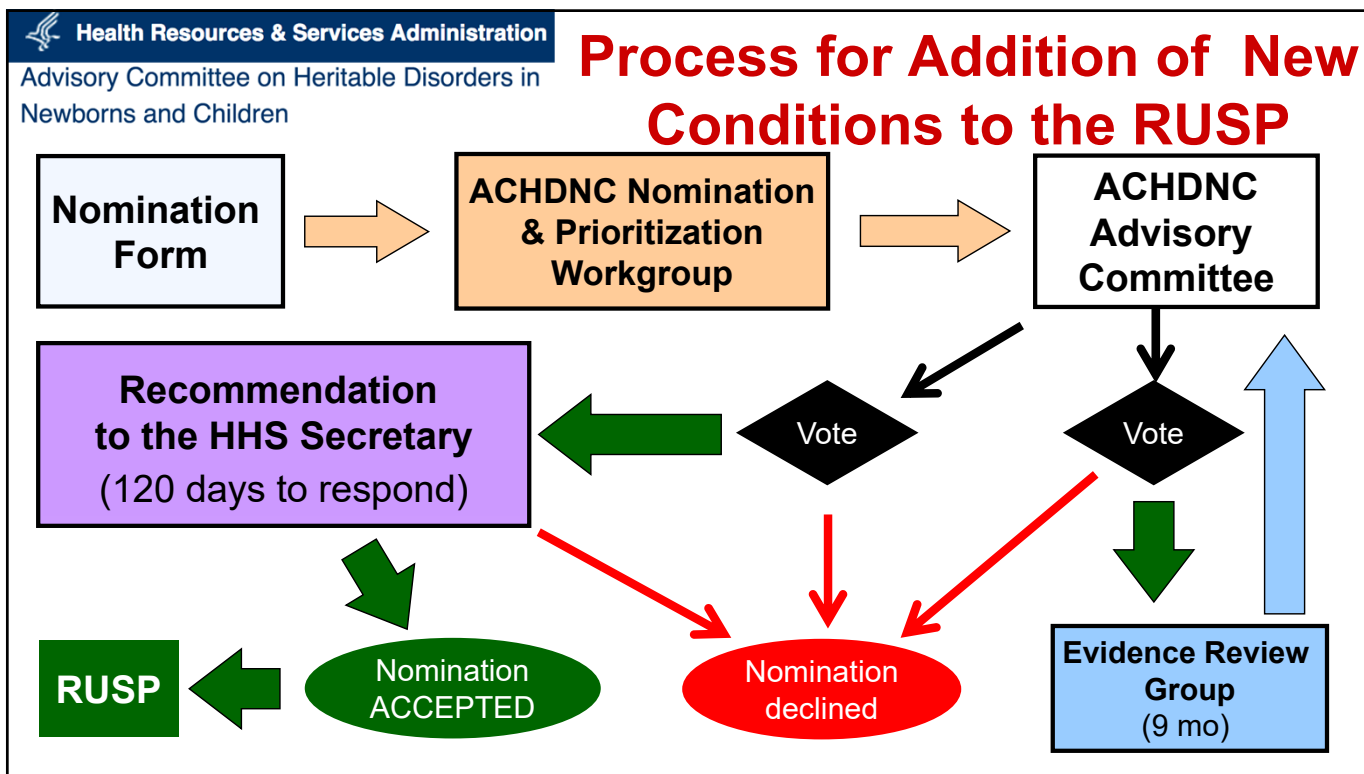
Upcoming Meeting

The Committee will meet via in-person and via webinar on May 4-5, 2023. The address for the meeting is 5600 Fishers Lane, Rockville, Maryland 20857. Please continue to check this website for the registration link and additional meeting information.

If you are a non-U.S. citizen who would like to attend the May meeting in-person, please contact ACHDNC@hrsa.gov by April 12, 2023.

The Advisory Committee on Heritable Disorders in Newborns and Children (Committee) was established under the [Public Health Service \(PHS\) Act, 42 U.S.C. 217a: Advisory councils or committees](#) (PDF - 214 KB), and [Title XI § 1111 \(42 U.S.C. § 300b-10\)](#) (PDF - 208 KB).

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- ACHDNC History

09/2007: SCID → added to RUSP: 5/2010

10/2007: Pompe disease

12/2007: Niemann-Pick A/B disease

12/2007: Fabry disease

01/2008: Krabbe disease

06/2008: Spinal muscular atrophy (SMA)

04/2009: Hemoglobin H disease

07/2008: Hyperbilirubinemia/Kernicterus

10/2009: Critical Congenital Heart Disease → added to RUSP: 9/2011

01/2011: 22q11 deletion syndrome

02/2012: Pompe disease → added to RUSP: 3/2015

02/2012: MPS I → added to RUSP: 2/2016

02/2012: X-Adrenoleukodystrophy

09/2013: X-Adrenoleukodystrophy → added to RUSP: 2/2016

05/2016: Guanidinoacetate Me-Transferase (GAMT) deficiency

02/2017: SMA → added to RUSP: 7/2018

05/2021: MPS II → added to RUSP: 8/2022

08/2021: GAMT → added to RUSP: 1/2023

08/2021: Krabbe disease → again NOT added to RUSP: 2/2023

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Recommended Uniform Screening Panel

Recommended Uniform Screening Panel
Core Conditions
(As of January 2023)

X: Condition is in this category - Condition is not in this category

Core Condition	Metabolic Disorder - Organic acid condition	Metabolic Disorder - Fatty acid oxidation disorder	Metabolic Disorder - Amino acid disorder	Endocrine Disorder	Hemoglobin Disorder	Other Disorder
Propionic Acidemia	X					
Methylmalonic Acidemia (methylmalonyl-CoA mutase)	X					
Methylmalonic Acidemia (Cobalamin disorders)	X					
Isovaleric Acidemia	X					
3-Methylcrotonyl-CoA Carboxylase Deficiency	X					
3-Hydroxy-3-Methylglutaryl-CoA Synthase Deficiency	X					
Holocarboxylase Synthase Deficiency	X					
β-Ketothiolase Deficiency	X					
Glutaric Acidemia Type I	X					
Carnitine Uptake Defect/Carnitine Transport Defect		X				
Medium-chain Acyl-CoA Dehydrogenase Deficiency		X				
Very Long-chain Acyl-CoA Dehydrogenase Deficiency		X				
Long-chain L3-Hydroxyacyl-CoA Dehydrogenase Deficiency		X				
Trifunctional Protein Deficiency		X				
Argininosuccinic Aciduria			X			
Citrullinemia, Type I			X			
Maple Syrup Urine Disease			X			
Homocystinuria			X			
Classic Phenylketonuria			X			
Tyrosinemia, Type I			X			
Guandinoacetate Methyltransferase Deficiency			X			
Primary Congenital Hypothyroidism				X		
Congenital adrenal hyperplasia				X		
S.S Disease (Sickle Cell Anemia)					X	
S, Beta-Thalassemia					X	
S,C Disease					X	
Biotinidase Deficiency						X
Critical Congenital Heart Disease						X
Cystic Fibrosis						X
Classic Galactosemia						X
Glycogen Storage Disease Type II (Pompe)						X
Hearing Loss						X

Recommended Uniform Screening Panel
Core Conditions
(As of January 2023)

Core Condition - continued

Core Condition - continued	Metabolic Disorder - Organic acid condition	Metabolic Disorder - Fatty acid oxidation disorder	Metabolic Disorder - Amino acid disorder	Endocrine Disorder	Hemoglobin Disorder	Other Disorder
Severe Combined Immunodeficiencies						X
Mucopolysaccharidosis Type I						X
X-linked Adrenoleukodystrophy						X
Spinal Muscular Atrophy due to homozygous deletion of exon 7 in SMN1						X
Mucopolysaccharidosis Type II						X

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Core
Conditions

Spinal Muscular Atrophy due to homozygous deletion of exon 7 in SMN1

<https://www.hrsa.gov/advisory-committees/heritable-disorders/rusp> (last accessed: 4/2/2023)

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Recommended Uniform Screening Panel

Recommended Uniform Screening Panel
Core Conditions
(As of January 2023)

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Propionic Acidemia	X					
Methylmalonic Acidemia (methylmalonyl-CoA mutase)	X					
Methylmalonic Acidemia (Cobalamin disorders)	X					
Isovaleric Acidemia	X					
3-Methylcrotonyl-CoA Carboxylase Deficiency	X					
3-Hydroxy-3-Methylglutaryl-CoA Synthase Deficiency	X					
Holocarboxylase Synthase Deficiency	X					
β-Ketothiolase Deficiency	X					
Glutaric Acidemia Type I	X					
Carnitine Uptake Defect/Carnitine Transport Defect		X				
Medium-chain Acyl-CoA Dehydrogenase Deficiency		X				
Very Long-chain Acyl-CoA Dehydrogenase Deficiency		X				
Long-chain L3-Hydroxyacyl-CoA Dehydrogenase Deficiency		X				
Trifunctional Protein Deficiency		X				
Argininosuccinic Aciduria			X			
Citrullinemia, Type I			X			
Maple Syrup Urine Disease			X			
Homocystinuria			X			
Classic Phenylketonuria			X			
Tyrosinemia, Type I			X			
Guandinoacetate Methyltransferase Deficiency			X			
Primary Congenital Hypothyroidism				X		
Congenital adrenal hyperplasia				X		
S.S Disease (Sickle Cell Anemia)					X	
S, Beta-Thalassemia					X	
S,C Disease					X	
Biotinidase Deficiency						X
Critical Congenital Heart Disease						X
Cystic Fibrosis						X
Classic Galactosemia						X
Glycogen Storage Disease Type II (Pompe)						X
Hearing Loss						X

Recommended Uniform Screening Panel
Core Conditions
(As of January 2023)

Core Condition - continued

Core Condition - continued	Metabolic Disorder - Organic acid condition	Metabolic Disorder - Fatty acid oxidation disorder	Metabolic Disorder - Amino acid disorder	Endocrine Disorder	Hemoglobin Disorder	Other Disorder
Severe Combined Immunodeficiencies						X
Mucopolysaccharidosis Type I						X
X-linked Adrenoleukodystrophy						X
Spinal Muscular Atrophy due to homozygous deletion of exon 7 in SMN1						X
Mucopolysaccharidosis Type II						X

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Core
Conditions

26+
Secondary
Conditions

Recommended Uniform Screening Panel¹
SECONDARY CONDITIONS²
(As of January 2023)

Secondary Condition	Metabolic Disorder - Organic acid condition	Metabolic Disorder - Fatty acid oxidation disorder	Metabolic Disorder - Amino acid disorder	Endocrine Disorder	Hemoglobin Disorder	Other Disorder
Methylmalonic acidemia with homocystinuria	X					
Malonic acidemia	X					
Isobutyrylglycinuria	X					
3-Methylcrotonylglycinuria	X					
3-Methylglutaconic aciduria	X					
2-Methyl-3-hydroxybutyric aciduria	X					
Short-chain acyl-CoA dehydrogenase deficiency		X				
Medium/short-chain L3-hydroxyacyl-CoA dehydrogenase deficiency		X				
Medium-chain ketacyl-CoA thiolase deficiency		X				
2,4-Diacyl-CoA reductase deficiency		X				
Carnitine palmitoyltransferase type I deficiency		X				
Glutaric acidemia type II		X				
Medium-chain ketacyl-CoA thiolase deficiency		X				
Carnitine palmitoyltransferase type II deficiency		X				
Carnitine acylcarnitine transferase deficiency		X				
Arginemia			X			
Citrullinemia, type II			X			
Hypermethioninemia			X			
Banigan hyperphenylalaninemia			X			
Biopterin defect in cofactor biosynthesis			X			

Secondary Condition - Continued

Secondary Condition - Continued	Metabolic Disorder - Organic acid condition	Metabolic Disorder - Fatty acid oxidation disorder	Metabolic Disorder - Amino acid disorder	Endocrine Disorder	Hemoglobin Disorder	Other Disorder
Biopterin defect in cofactor regeneration			X			
Tyrosinemia, type II			X			
Tyrosinemia, type III			X			
Various other hemoglobinopathies					X	
Galactosemia deficiency						X
Galactokinase deficiency						X
T-cell related lymphocyte deficiencies						X

1. Selection of conditions based upon "Newborn Screening: Towards a Uniform Screening Panel and System." Genetics Med. 2006; 6(5) Suppl. S12-S22. as endorsed by the American College of Medical Genetics (ACMG) and cosponsored by the Health Resources and Services Administration (HRSA).
2. Disorders that can be detected in the differential diagnosis of a core disorder.
3. Nonrecurrant for conditions based upon "Naming and Coding Disorders (Conditions) Included in Newborn Screening Panels." Pediatrics. 2006; 117 (3) Suppl. S308-S314.

<https://www.hrsa.gov/advisory-committees/heritable-disorders/rusp> (last accessed: 4/2/2023)

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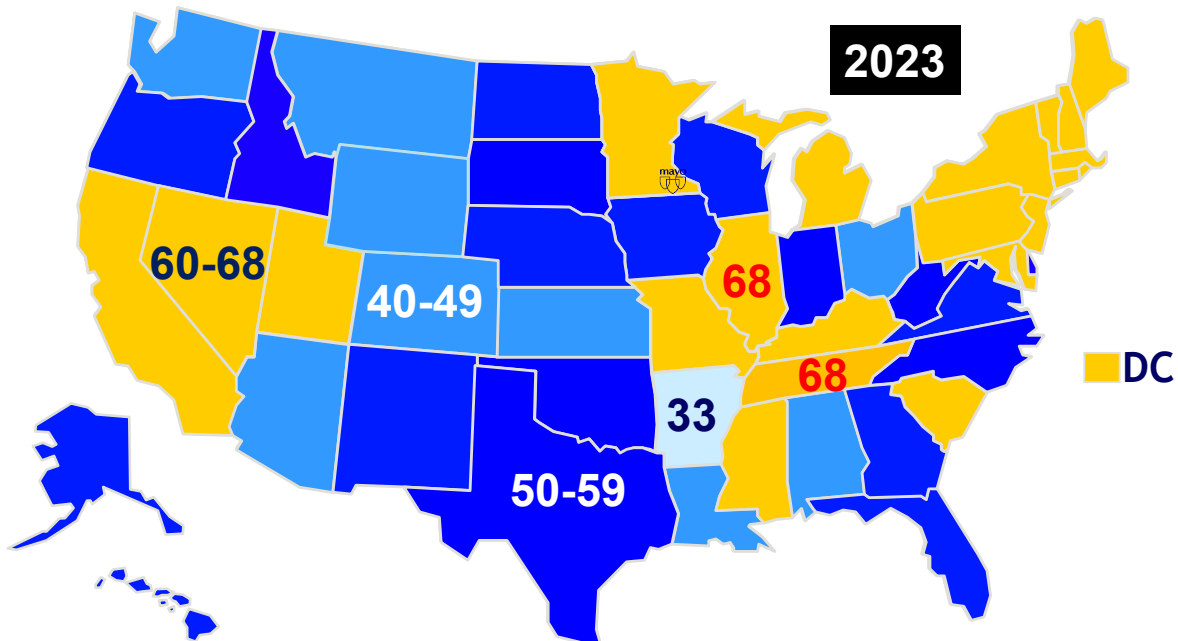
Newborn Screening

A public health program:

- Aimed at identification of conditions for which early intervention can prevent mortality, morbidity and disabilities.
- Performed by analysis of diagnostic markers in blood spots collected on filter paper on the second day of life (exception: hearing loss and congenital cyanotic heart disease).
- Extent of program is determined by each state independently.

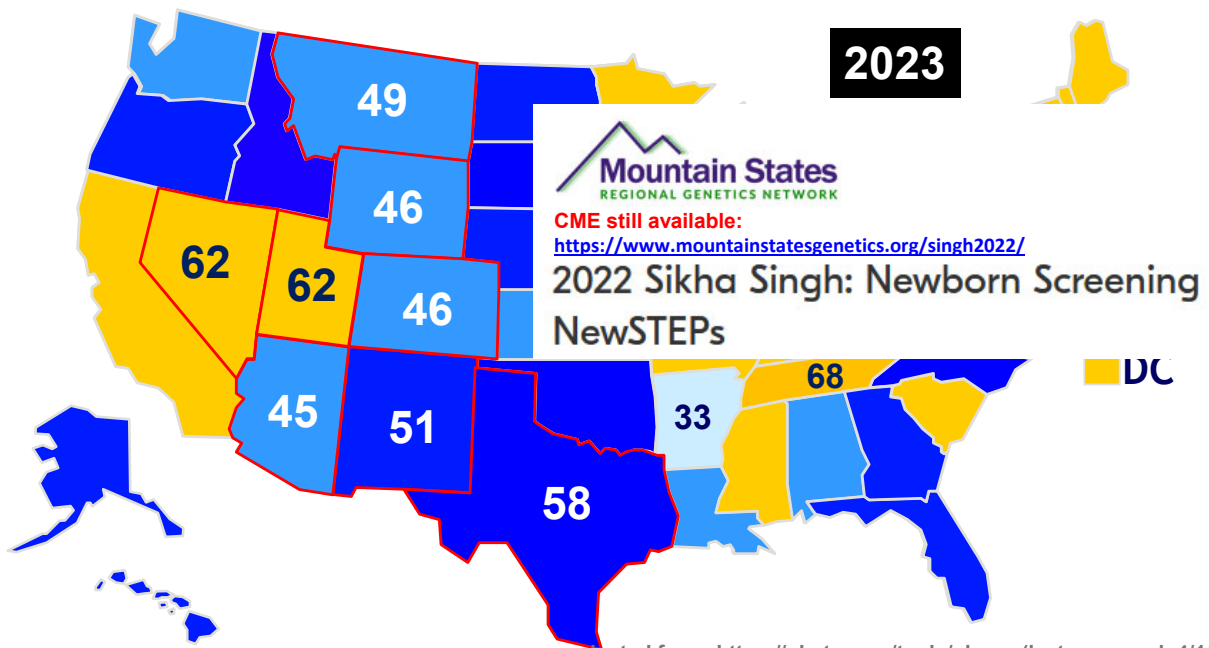
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Number of Conditions included in NBS Programs



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Number of Conditions included in NBS Programs



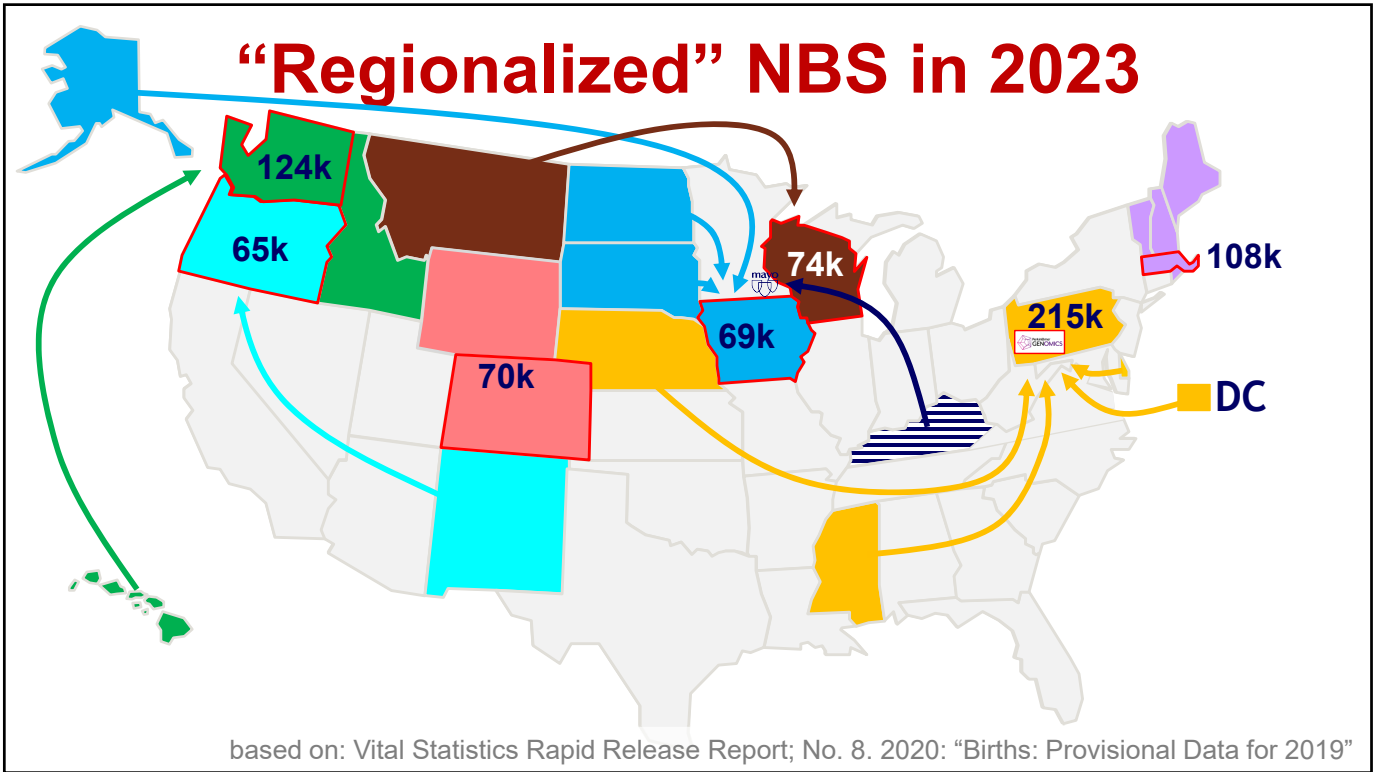
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Newborn Screening

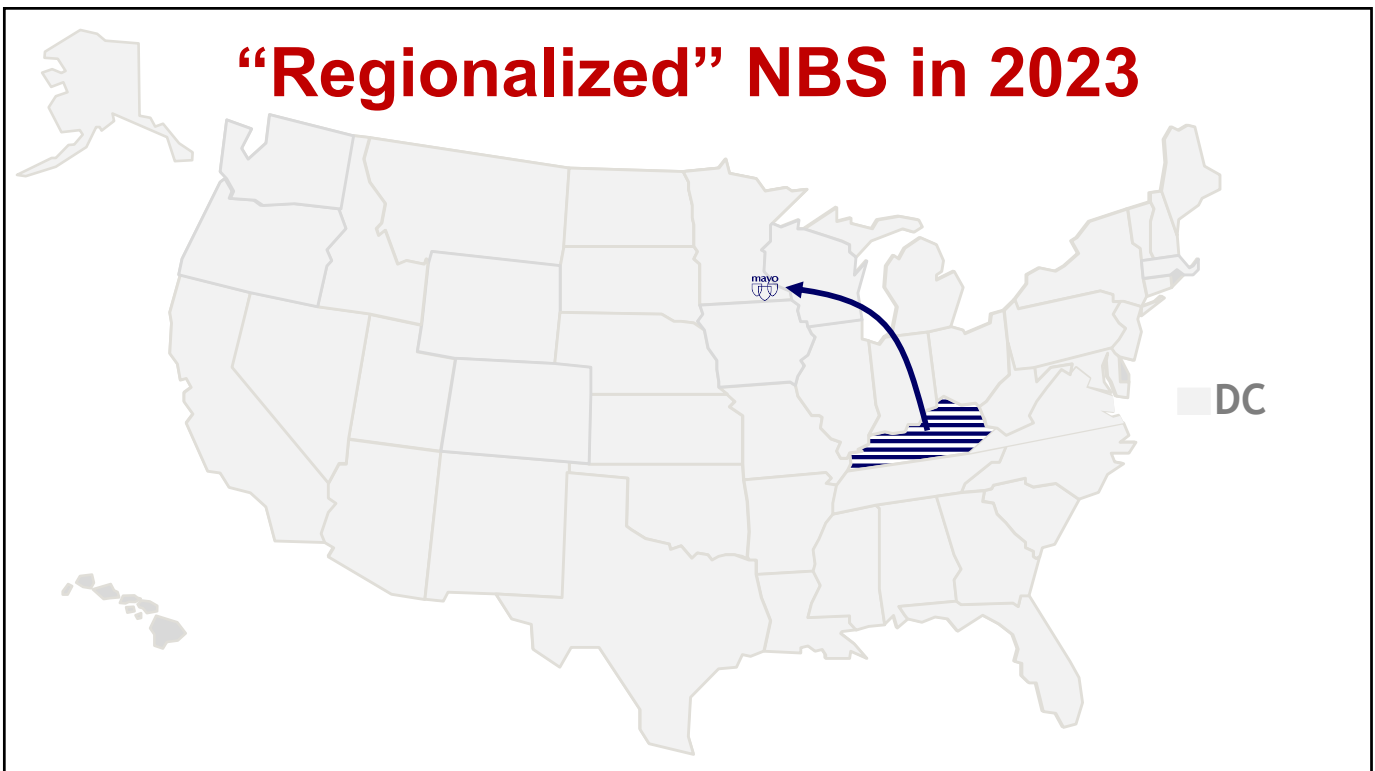
A public health program:

- Aimed at identification of conditions for which early intervention can prevent mortality, morbidity and disabilities.
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- Extent of program determined by each state independently.
- Administered by each state but testing may be out of state

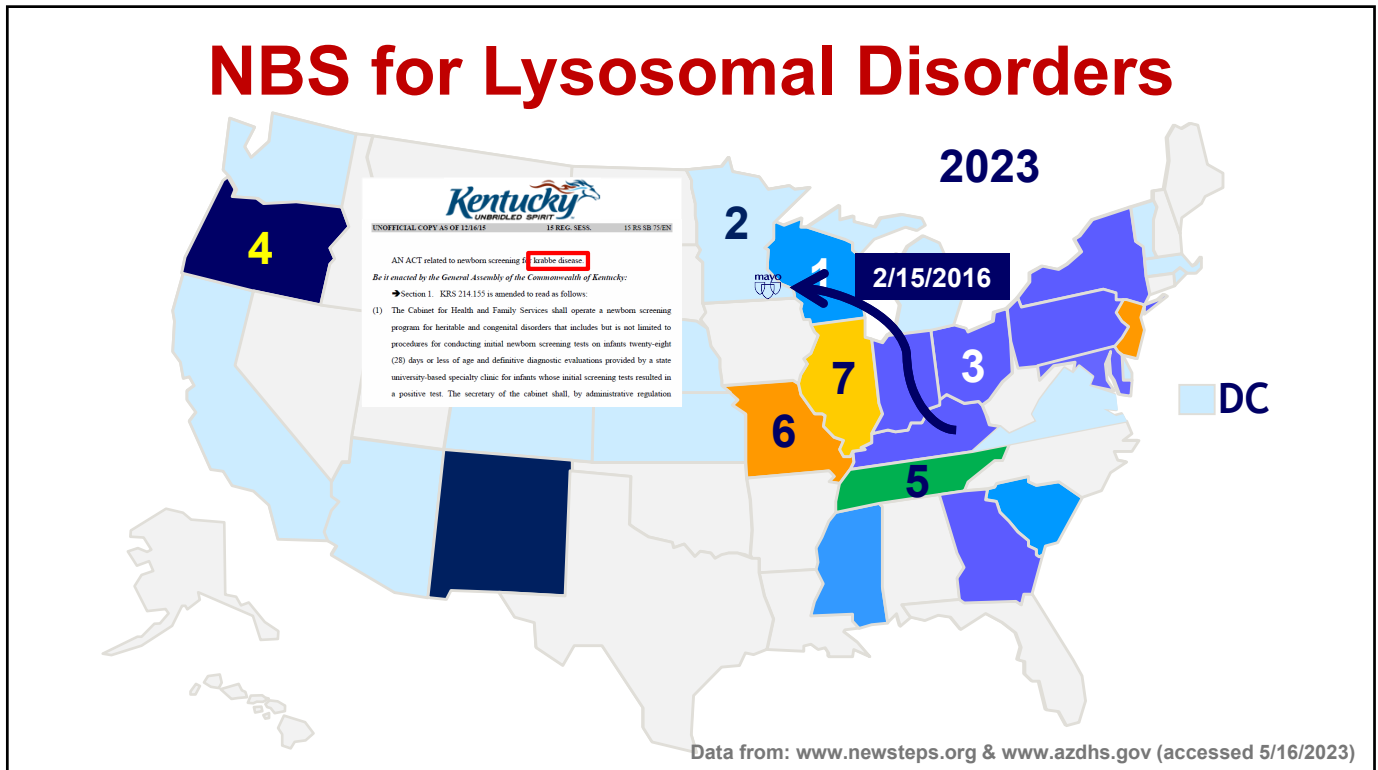
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Krabbe Disease

- **Infantile Krabbe disease (IKD) presents in first 12 months:**
 - Extreme irritability
 - Spasticity
 - Developmental regression
 - Median survival: <2 yrs of age
- **Late Infantile KD (LIKD):**
 - onset of irreversible and progressive symptoms between 1-3 yrs
 - Median survival: 7 yrs
- **Juvenile KD (JKD):** progressive symptoms as of 4-17 yrs
- **Adult KD (AKD):** progressive symptoms as of 18 or more yrs
- **JKD and LKD can present with weakness, spasticity, ataxia, vision loss, and/or as neuropsychiatric disease in adults**



Krabbe K. *Brain*, 1916; 39: 74-114

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Why NBS for Krabbe Disease

Clinical Chemistry 50:10
1785-1796 (2004)

Automation and
Analytical Techniques

Direct Multiplex Assay of Lysosomal Enzymes in Dried Blood Spots for Newborn Screening

YIHUN LI,¹ C. RONALD SCOTT,² NESTOR A. CHAMOLES,³ AHMAD GHAYAMI,⁴ B. MARIO PINTO,⁴ FRANTISEK TURECEK,¹ and MICHAEL H. GELB^{1,5*}

Background: Newborn screening for deficiency in the lysosomal enzymes that cause Fabry, Gaucher, Krabbe, Niemann-Pick A/B, and Pompe diseases is warranted because treatment for these syndromes is now available or anticipated in the near future. We describe a multiplex screening method for all five lysosomal enzymes that uses newborn-screening cards containing dried blood spots as the enzyme source.

Methods: We used a cassette of substrates and internal standards to directly quantify the enzymatic activities, and tandem mass spectrometry for enzymatic product detection. Rehydrated dried blood spots were incubated with the enzyme substrates. We used liquid-liquid extraction followed by solid-phase extraction with octyl gel to separate the enzyme products from the inhibitors and neuropeptides.

Results: with Gaucher, 5 with Niemann-Pick A/B, 11 with Pompe, 5 with Fabry, and 12 with Krabbe disease, and in all cases the enzyme activities were below the minimum activities measured in a collection of heterozygous carriers and healthy noncarrier individuals. The enzyme activities measured in 5-9 heterozygous carriers were approximately one-half those measured with 19-32 healthy individuals, but there was partial overlap of each condition between the data sets for carriers and healthy individuals.

Conclusions: For all five diseases, the affected individuals were detected. The assay can be readily automated,

and the anticipated reagent and supply costs are well within the budget limits of newborn-screening centers. © 2004 American Association for Clinical Chemistry

The application of mass spectrometry in disease diagnostics is rapidly advancing. Major applications include (a) quantification of metabolites in dried blood spots (DBS)¹ by tandem mass spectrometry (1); (b) identification and quantification of protein biomarkers in serum by surface-enhanced laser desorption/ionization mass spectrometry (2); (c) analysis of PCR-amplified DNA by MassArray to identify mutated genes (3); and (d) quantification of enzyme activities in cell lysates by affinity capture/

of disorders that affect both somatic organs and the central nervous system. Depending on the disorder, they can manifest from birth to adulthood. Until recently, there has been minimal therapeutic intervention available to alter the natural course of the disorders, but with the recent availability of commercial preparations of recombinant enzymes, selective lysosomal disorders are now amenable to therapeutic intervention (7-9). There is excellent documentation that enzyme replacement therapy for Gaucher disease [acid β -glucocerebrosidase (ABG) deficiency] and Fabry disease [acid α -galactosidase A (GLA) deficiency] may alter the natural progression of the disorders and improve the clinical phenotype. Clinical trials are currently underway for Hunter syndrome and Pompe disease [lysosomal acid α -glucosidase (CAA) deficiency], disor-

Departments of ¹Chemistry, ²Pediatrics, and ³Biochemistry, University of Washington, Seattle, WA; ⁴Laboratory of Neurochemistry, Buenos Aires, Argentina; ⁵Department of Chemistry, Simon Fraser University, Burnaby, British Columbia, Canada; *Address correspondence to the author at Department of Chemistry, Campus Box 357070, University of Washington, Seattle, WA 98195. Fax: 206-685-8663; e-mail: gelb@chem.washington.edu.
Received April 20, 2004; accepted June 29, 2004.
Previously published online at DOI: 10.1373/jclinchem.2004.05947

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Transplantation of Umbilical-Cord Blood in Babies with Infantile Krabbe's Disease

Maria L. Escolar, M.D., Michele D. Poe, Ph.D., James M. Provenzale, M.D., Karen C. Richards, M.D., June Allison, R.N., Susan Wood, P.N.P., David A. Wenger, Ph.D., Daniel Pietryga, M.D., Donna Wall, M.D., Martin Champagne, M.D., Richard Morse, M.D., William Kivitit, M.D., Ph.D., and Joanne Kurtzberg, M.D.

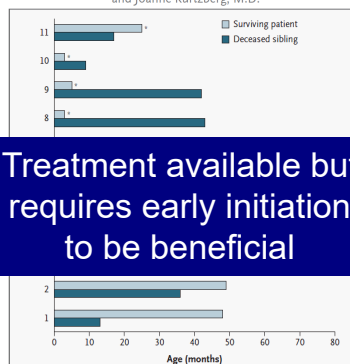


Figure 2. Survival of Patients and Siblings.

The age at death of untreated siblings of patients with Krabbe's disease who underwent transplantation is shown, as is the age of newborns who underwent transplantation and who were alive as of January 28, 2005. Six patients had outlived their siblings by 8 to 48 months. An asterisk indicates patients who have not yet reached the age of death of their siblings.

N ENGL J MED 352:20 WWW.NEJM.ORG MAY 19, 2005

Mayo Clinic's Supplemental NBS for Kentucky

- Public/private partnership after law to add Krabbe disease to KY NBS panel became effective on 6/24/2015.
- KY NBS Lab started sending DBS to Mayo (MN) for screening for Krabbe disease, Pompe disease and MPS I on 2/15/16; ALD was added on 7/9/18.
- DBS are separated in KY & shipped to Mayo; demographic data transmitted electronically.
- KY operates 6 days a week, Mayo operates 7 days a week.
- Samples received are analyzed overnight; results reported next day.
- Leftover DBS are returned to KY every other week.

Mayo Clinic's NBS for Kentucky

10-plex Assay* → **CLIR for KD, MPS I, PD, ALD**

- **INFORMATIVE Score for KD**
- **INFORMATIVE Score for PD**
- **INFORMATIVE Score for MPS I**
- **INFORMATIVE Score for ALD**
- **NO Informative Score**

*6 enzymes + 4 lysophosphatidylcholines by FIA-MS/MS

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ORIGINAL RESEARCH ARTICLE

Open

Precision newborn screening for lysosomal disorders

Melissa M. Minter Baerg, MBA¹, Stephanie D. Stoway, MPH¹, Jeremy Hart, MD^{2,3}, Lea Mott, MT², Dawn S. Peck, MS, CGC¹, Stephanie L. Nielt, MTASCP⁴, Jason S. Eckerman, BS¹, Jean M. Lacey, BS¹, Coleman T. Turgeon, MS¹, Dimitar Gavrilov, MD, PhD¹, Devin Oglesbee, PhD¹, Kimiyo Raymond, MD¹, Silvia Tortorelli, MD, PhD¹, Dietrich Matern, MD, PhD¹, Lars Morkrid, MD, PhD⁴ and Piero Rinaldo, MD, PhD¹

Purpose: The implementation of newborn screening for lysosomal disorders has uncovered overall poor specificity, psychosocial harm experienced by caregivers, and costly follow-up testing of false-positive cases. We report an informatics solution proven to minimize these issues.

Methods: The Kentucky Department of Public Health outsourced testing for mucopolysaccharidosis type I (MPS I) and Pompe disease, conditions recently added to the recommended uniform screening panel, plus Krabbe disease, which was added by legislative mandate. A total of 55,161 specimens were collected from infants born over 1 year starting from February 2016. Testing by tandem mass spectrometry was integrated with multivariate pattern recognition software (Collaborative Laboratory Integrated Report), which is freely available to newborn screening programs for selection of cases for which a biochemical second-tier test is needed.

Results: Of five presumptive positive cases, one was affected with infantile Krabbe disease, two with Pompe disease, and one with MPS I. The remaining case was a heterozygote for the latter condition. The false-positive rate was 0.0018% and the positive predictive value was 80%.

Conclusion: Postanalytical interpretive tools can drastically reduce false-positive outcomes, with preliminary evidence of no greater risk of false-negative events, still to be verified by long-term surveillance.

Genet Med advance online publication 9 November 2017

Key Words: collaborative laboratory integrated report; Krabbe disease; mucopolysaccharidosis type I; newborn screening; Pompe disease

INTRODUCTION

Newborn screening is a public health program aimed at the identification of conditions for which early intervention can prevent mortality, morbidity, and disabilities,¹ but is not without its challenges.² In recent years, expansions of testing panels have been proposed and adopted either as national standards³ or as nonbinding recommendations⁴ by the US Department of Health and Human Services (HHS). Currently, the recommended uniform screening panel encompasses 34 conditions. The HHS Secretary's Advisory Committee on Heritable Disorders in Newborns and Children is tasked with overseeing the process⁵ of adding emerging conditions^{6,7} to the recommended panel. The most recent additions are acid α-glucosidase (GAA) deficiency (Pompe disease),⁸ α-iduronidase (IDUA) deficiency (MPS I),⁹ and X-linked adrenoleukodystrophy.¹⁰ Other lysosomal disorders, particularly galactocerebrosidase (GALC) deficiency (Krabbe disease),¹¹ have been turned down by the committee because they lacked evidence of net benefits. However, advocacy efforts and legislative mandates have propelled six states to

disorders,^{12–14} but reports of outcomes and performance have not been encouraging.^{15–17}

In 2015, the legislature of the Commonwealth of Kentucky passed bill KRS 214.155, mandating screening for Krabbe disease. This action was driven by an advocacy campaign led by the parents of a child affected with the infantile form of the disease. To accelerate implementation, the Kentucky Department for Public Health reached out to the Biochemical Genetics Laboratory at Mayo Clinic (Rochester, Minnesota) to negotiate outsourcing of screening for Krabbe disease by measuring only GALC activity. This assay had to be performed in parallel to local testing for all other conditions and be completed within 24 hours to avoid any further delays in the care of early-onset cases.¹⁸ This request was not fulfilled because conventional interpretation methods of a single marker (cutoff percent of daily mean) are not suited to differentiation of affected patients from individuals who are either heterozygous or carry pseudo-deficiency alleles.^{19,20} The laboratory made a counter-proposal to perform testing of six lysosomal enzyme activities using available substrates, inclusive of the previously mentioned disorders, to better differentiate affected patients from unaffected individuals.

Biomedical Genetics Laboratory, Department of Laboratory Medicine and Pathology, Mayo Clinic, Rochester, Minnesota, USA; Division of Laboratory Medicine, Department of Public Health, Frankfort, Kentucky, USA; Department of Pathology & Laboratory Medicine, University of Kentucky, Lexington, Kentucky, USA; Medical Biotechnology, Oslo University Hospital, Rikshospitalet, Oslo, Norway; Correspondence: Piero Rinaldo (rinaldo@mayo.edu)

Submitted 5 July 2017; accepted 28 September 2017; advance online publication 9 November 2017; doi:10.1038/gim.2017.194

GENETICS IN MEDICINE | Volume 20 | Number 8 | August 2018

847

Screening NEGATIVE

41

Mayo Clinic's NBS for Kentucky

10-plex Assay → **CLIR for KD, MPS I, PD, ALD**

- **INFORMATIVE Score for KD**
- **INFORMATIVE Score for PD**
- **INFORMATIVE Score for MPS I**
- **INFORMATIVE Score for ALD**
- **NO Informative Score**

→ **INFORMATIVE Score for KD**

→ **INFORMATIVE Score for PD**

→ **INFORMATIVE Score for MPS I**

→ **INFORMATIVE Score for ALD**

Second Tier Test

Screening NEGATIVE

42

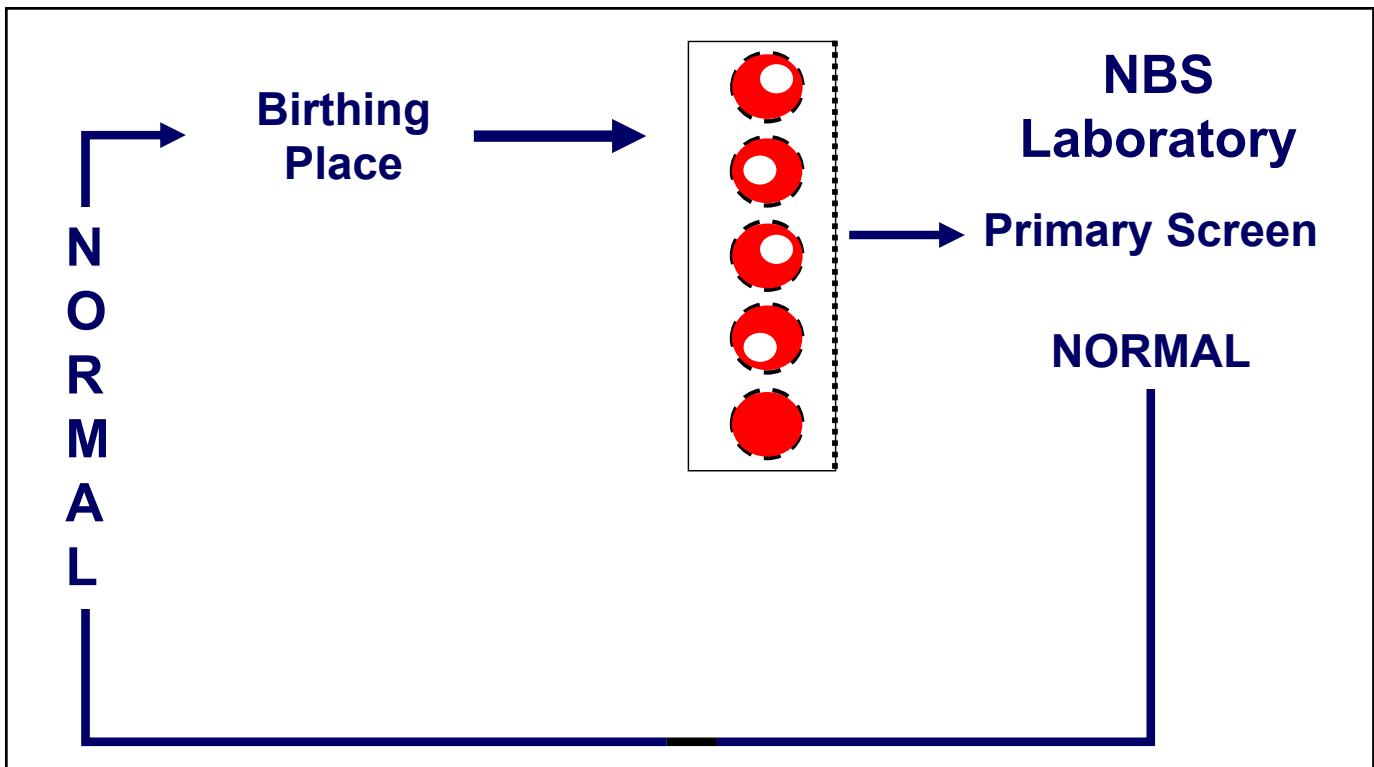
21

What are 2nd Tier Tests?

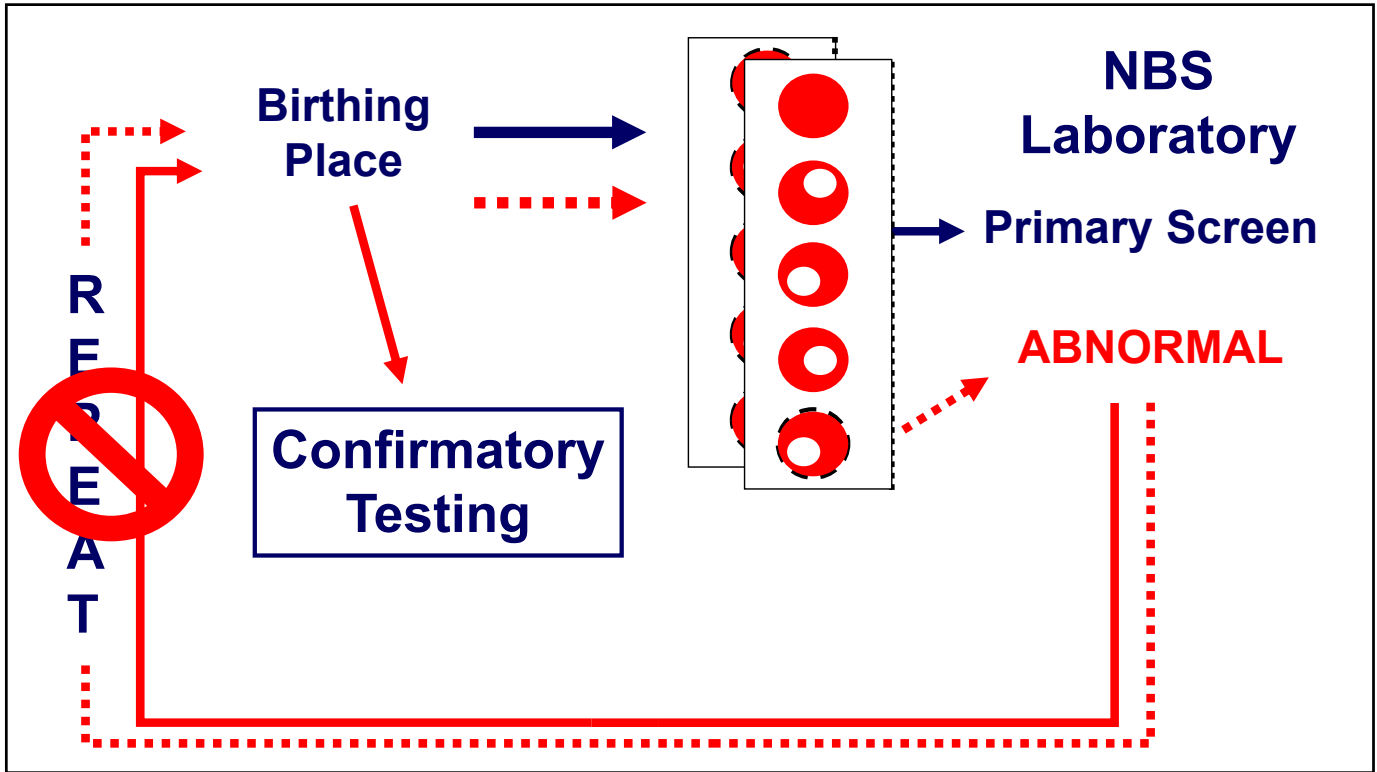
- A cost effective approach to reduce false positive results when normal population and disease range overlap (poor specificity)
- After primary screen
- Same specimen, no additional patient contact
- Normal 2nd tier test result overrules primary screen
 - ➔ reduction of false positive results
- Examples: - biochemical (CAH, MSUD, PA, MMA, RMD, HCU, SCAD, GA I, GA II, Pompe, Krabbe, MPS I, MPS II),
- molecular (CF)



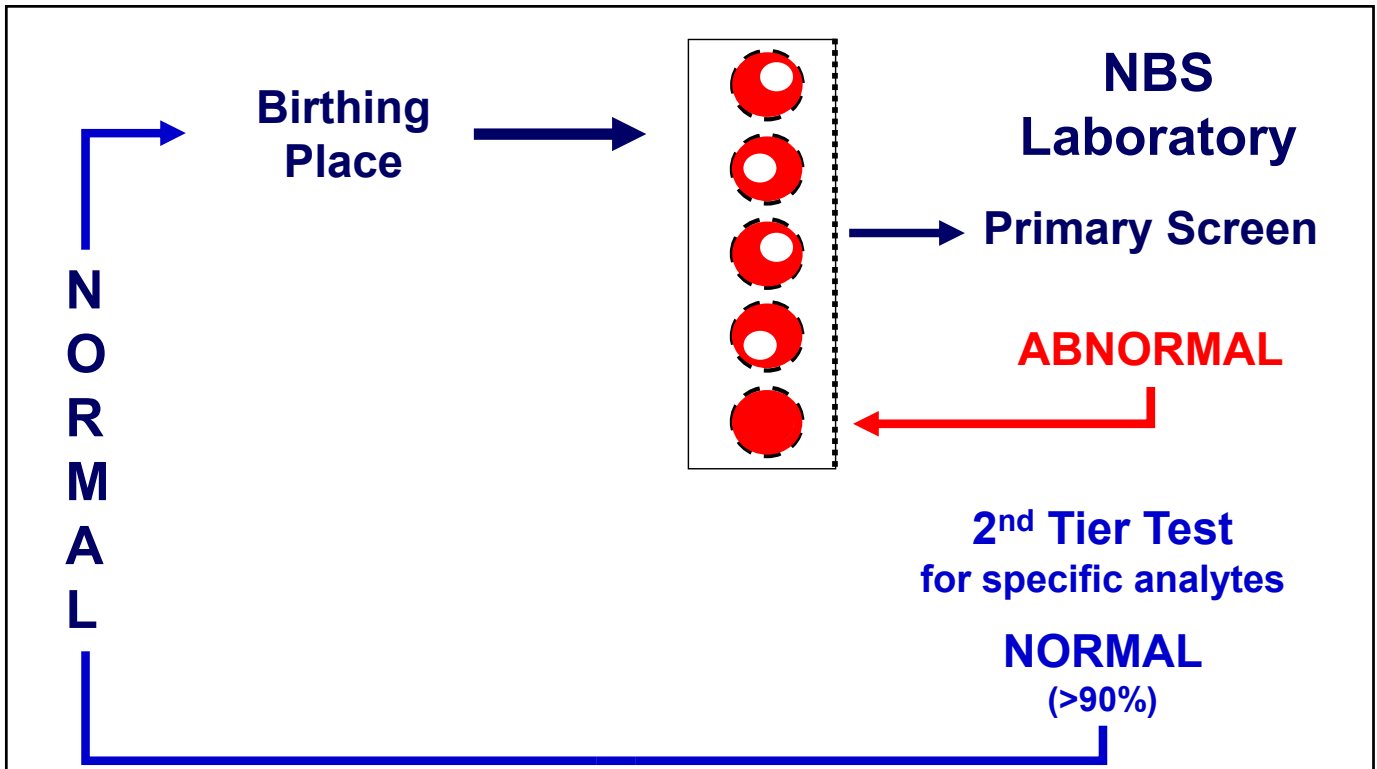
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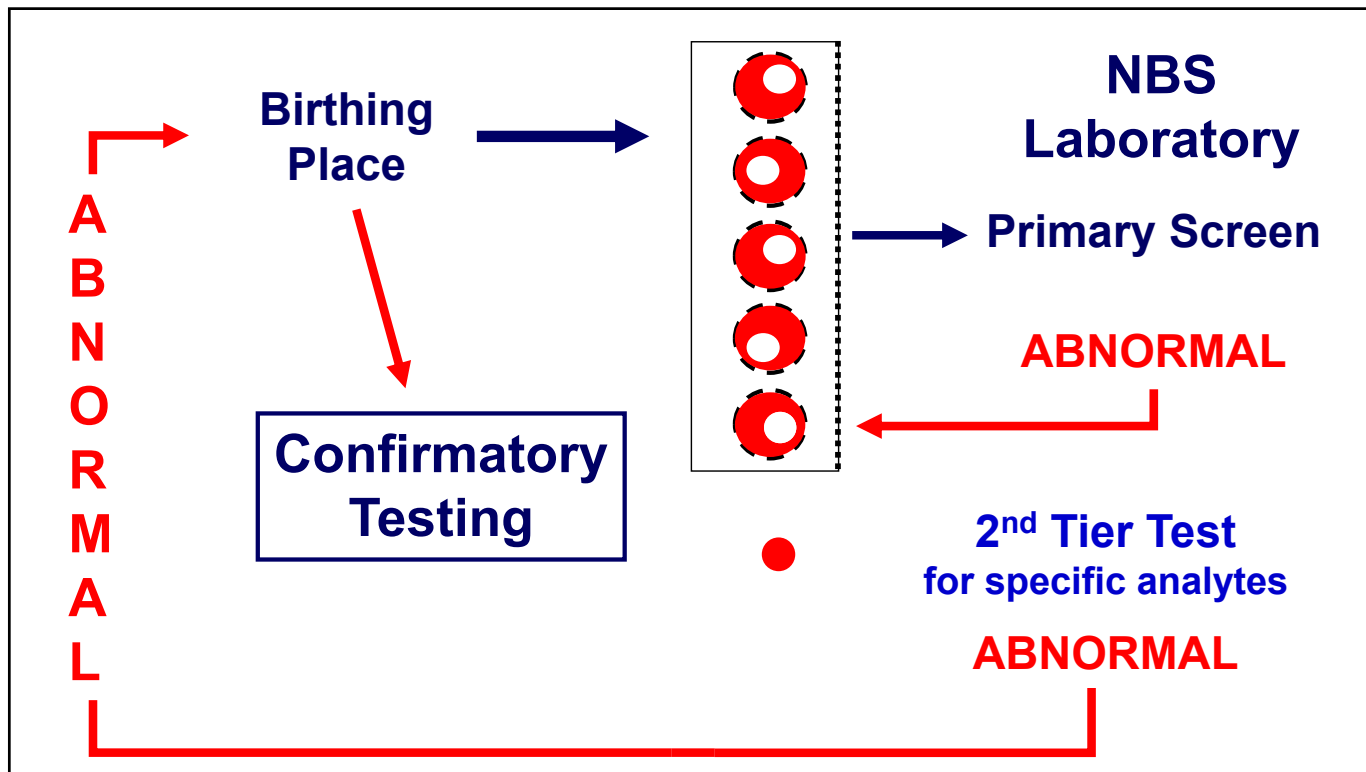
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Krabbe

J Inher Metab Dis
DOI 10.1007/s10545-015-9122-z

ORIGINAL ARTICLE

Measurement of psychosine in dried blood spots — a possible improvement to newborn screening programs for Krabbe disease

Coteman T, Furgson J, Orsini J, Karen A, Sanders M, Mark J, Magner T, Thomas J, Langan P, Maria L, Essner P, Patricia Duffner D, Devin Oglesbe D, Dmitar Gavrilov S, Silvia Tortorelli P, Piero Rinaldo P, Kimyo Raymond P, Dietrich Mattern P

Received: 3 December 2014 / Revised: 30 January 2015 / Accepted: 3 February 2015
© SSIEM 2015

Abstract
Background: Newborn screening (NBS) for Krabbe disease (KD) in New York and Missouri is conducted by measuring galactosylceramidase (GALC) activity using tandem mass spectrometry (MS/MS). These NBS efforts have shown that the incidence of KD is unexpectedly low (1,400,000) while many individuals (ca. 1,600,000) with reduced GALC activity and genotypes of uncertain significance are detected and subjected to follow-up testing. Psychosine (PSY) is a putative marker of KD progression and can be measured in dried blood spots (DBS). We sought to determine the role that PSY levels play in NBS for KD, follow-up, and treatment monitoring. **Methods:** PSY was cloned from DBS with methanol containing N,N-dimethyl-D-cylophosphorimide as internal standard (IS). Liquid chromatography-MS/MS was conducted over 17 minutes in the multiple reaction monitoring positive mode to follow the precursor to product species transitions for PSY and IS. Separation of the structural isomers PSY and galactosylpsychosine was accomplished by hydrophilic interaction liquid chromatography. **Results:** Pre-analytical and analytical factors were studied and revealed satisfactory results. PSY was also measured in DBS collected from controls (range: $3-112$, $N=220</math>), KD patients at various disease stages (range: 3–112, $N=26</math>), and GALC mutation carriers (range: $0.15-10.6$, $N=18</math>). **Conclusions:** PSY measurement in DBS could serve as a 2nd tier assay in NBS for KD, simplify and reduce the cost of follow-up protocols, help determine disease progression, and be used to monitor KD patients following hematopoietic stem cell transplantation. However, additional chronological measurements of PSY in KD patients are required to confirm these possibilities.$$$

Abbreviations
GALC Galactosylceramidase
DBS Dried blood spots
ISCT Hematopoietic stem cell transplantation
KD Krabbe disease
LC-MS/MS Liquid chromatography-tandem mass spectrometry
NBS Newborn screening
PSY Psychosine

Introduction
Krabbe disease (KD, OMIM #245200), also known as globoid cell leukodystrophy, is a devastating neurodegenerative condition caused by lysosomal galactosylceramidase (GALC, EC 3.2.1.46) deficiency. Patients with the classic, early-infantile phenotype of this autosomal-recessive condition present in the first few months of life with irritability,

MPS I II

ORIGINAL ARTICLE

Incorporation of Second-Tier Biomarker Testing Improves the Specificity of Newborn Screening for Mucopolysaccharidosis Type I

Dawn S. Peck^{1,2}, Jean M. Laey³, Amy L. White^{1,3}, Gisèle Pino¹, April L. Studinski¹, Rachel Fisher², Ayesha Ahmad², Linda Spencer², Sarah Viall⁴, Natalie Shallow⁵, Amy Siemsen⁶, J. Austin Haman⁷, Brianna K. Murray⁸, Kelly L. Jones^{9,10}, Dmitar Gavrilov¹, Devin Oglesbe¹⁰, Kimyo Raymond², Dietrich Mattern², Piero Rinaldo² and Silvia Tortorelli^{1*}

Abstract
Background: Newborn screening (NBS) for Mucopolysaccharidosis Type I (MPS I) is conducted by measuring dermatan sulfate (DS) and heparan sulfate (HS) in dried blood spots (DBS) using tandem mass spectrometry (MS/MS). These NBS efforts have shown that the incidence of MPS I is unexpectedly low (1,400,000) while many individuals (ca. 1,600,000) with reduced DS and HS activity and genotypes of uncertain significance are detected and subjected to follow-up testing. Psychosine (PSY) is a putative marker of MPS I progression and can be measured in DBS. We sought to determine the role that PSY levels play in NBS for MPS I, follow-up, and treatment monitoring. **Methods:** PSY was cloned from DBS with methanol containing N,N-dimethyl-D-cylophosphorimide as internal standard (IS). Liquid chromatography-MS/MS was conducted over 17 minutes in the multiple reaction monitoring positive mode to follow the precursor to product species transitions for PSY and IS. Separation of the structural isomers PSY and galactosylpsychosine was accomplished by hydrophilic interaction liquid chromatography. **Results:** Pre-analytical and analytical factors were studied and revealed satisfactory results. PSY was also measured in DBS collected from controls (range: $3-112$, $N=220</math>), MPS I patients at various disease stages (range: 3–112, $N=26</math>), and GALC mutation carriers (range: $0.15-10.6$, $N=18</math>). **Conclusions:** PSY measurement in DBS could serve as a 2nd tier assay in NBS for MPS I, simplify and reduce the cost of follow-up protocols, help determine disease progression, and be used to monitor MPS I patients following hematopoietic stem cell transplantation. However, additional chronological measurements of PSY in MPS I patients are required to confirm these possibilities.$$$

Abbreviations
GALC Galactosylceramidase
DBS Dried blood spots
ISCT Hematopoietic stem cell transplantation
KD Krabbe disease
LC-MS/MS Liquid chromatography-tandem mass spectrometry
NBS Newborn screening
PSY Psychosine

Introduction
Krabbe disease (KD, OMIM #245200), also known as globoid cell leukodystrophy, is a devastating neurodegenerative condition caused by lysosomal galactosylceramidase (GALC, EC 3.2.1.46) deficiency. Patients with the classic, early-infantile phenotype of this autosomal-recessive condition present in the first few months of life with irritability,

Genetics in Medicine | ORIGINAL RESEARCH ARTICLE

Moonlighting newborn screening markers: the incidental discovery of a second-tier test for Pompe disease

Silvia Tortorelli, MD, PhD¹, Jason S. Eckerman, BS¹, Joseph J. Orsini, PhD², Colleen Stevens, PhD², Jeremy Hart, MD^{3,4}, Patricia L. Hall, PhD^{5,6}, John J. Alexander, PhD^{7,8}, John J. Alexander, PhD^{7,8}, Dmitar Gavrilov, MD, PhD¹, Devin Oglesbe, PhD¹⁰, Kimyo Raymond, MD², Dietrich Mattern, PhD², Piero Rinaldo, PhD² and Silvia Tortorelli, MD, PhD^{1*}

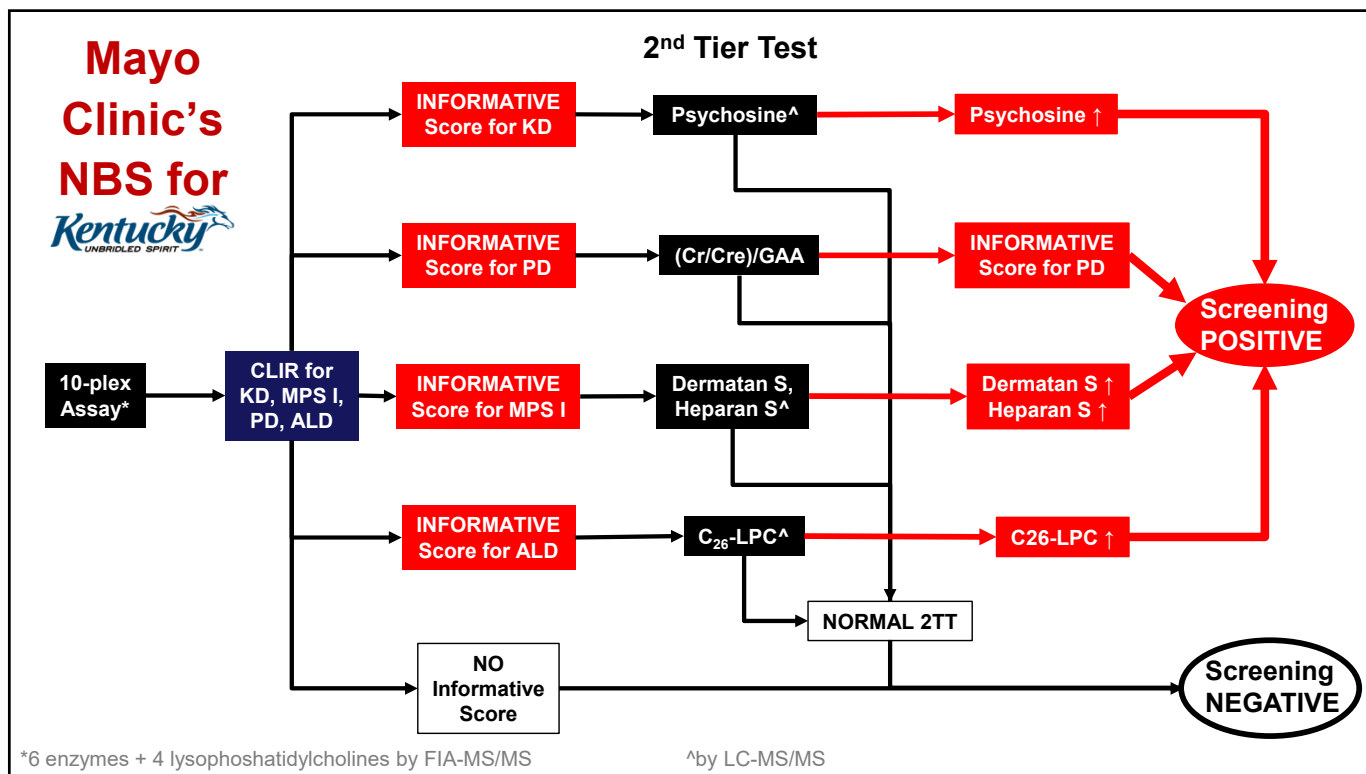
Abstract
Background: Newborn screening (NBS) for Pompe disease (PD) is conducted by measuring alpha-glucosidase (GAA) activity using tandem mass spectrometry (MS/MS). These NBS efforts have shown that the incidence of PD is unexpectedly low (1,400,000) while many individuals (ca. 1,600,000) with reduced GAA activity and genotypes of uncertain significance are detected and subjected to follow-up testing. Psychosine (PSY) is a putative marker of PD progression and can be measured in DBS. We sought to determine the role that PSY levels play in NBS for PD, follow-up, and treatment monitoring. **Methods:** PSY was cloned from DBS with methanol containing N,N-dimethyl-D-cylophosphorimide as internal standard (IS). Liquid chromatography-MS/MS was conducted over 17 minutes in the multiple reaction monitoring positive mode to follow the precursor to product species transitions for PSY and IS. Separation of the structural isomers PSY and galactosylpsychosine was accomplished by hydrophilic interaction liquid chromatography. **Results:** Pre-analytical and analytical factors were studied and revealed satisfactory results. PSY was also measured in DBS collected from controls (range: $3-112$, $N=220</math>), PD patients at various disease stages (range: 3–112, $N=26</math>), and GALC mutation carriers (range: $0.15-10.6$, $N=18</math>). **Conclusions:** PSY measurement in DBS could serve as a 2nd tier assay in NBS for PD, simplify and reduce the cost of follow-up protocols, help determine disease progression, and be used to monitor PD patients following hematopoietic stem cell transplantation. However, additional chronological measurements of PSY in PD patients are required to confirm these possibilities.$$$

Abbreviations
GAA Alpha-glucosidase
DBS Dried blood spots
ISCT Hematopoietic stem cell transplantation
PD Pompe disease
LC-MS/MS Liquid chromatography-tandem mass spectrometry
NBS Newborn screening
PSY Psychosine

Introduction
Krabbe disease (KD, OMIM #245200), also known as globoid cell leukodystrophy, is a devastating neurodegenerative condition caused by lysosomal galactosylceramidase (GALC, EC 3.2.1.46) deficiency. Patients with the classic, early-infantile phenotype of this autosomal-recessive condition present in the first few months of life with irritability,

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Kentucky Status

Newborns screened: 379,507 (2/15/2016 – 2/28/2023)*

Condition	2nd Tier Test	True Positive (TP)	False Positive (FP)	False Positive Rate	Positive Predictive Value	Final Diagnosis
Krabbe	116 (0.031%)	2	0	0%	100%	Infantile KD
Pompe	188 (0.049%)	24	14	0.004%	63%	Late onset PD
MPS I	149 (0.039%)	10	7	0.002%	59%	MPS I Hurler/Scheie
ALD*	1,368* (0.556%*)	8	0	0%	100%	6 ALD, 2 Zellweger
TOTAL		44	21	0.006%	68%	

*Newborns screened for ALD (7/9/2018 – 2/28/2023): 246,137

False Positive Rate: (FP/Total)×100%; Positive Predictive Value: (TP/[TP+FP])×100%

50

A Case of Krabbe Disease from KY

NBS collected	DBS arrives in KY Lab	DBS arrives at Mayo			Patient admitted at Duke		
		Sample prep and incubation	Informative for Krabbe → initiate rpt + 2TTs (PSY & 30kbDel)	GALC ↓ PSY ↑ 30kbDel -/- → initiate follow up, initiate GALC sequencing	GALC in WBC, PSY, Parental DNA, HLA typing, Neurologic exam, MRI brain with DTI, BAER, VEP, EEG, NCV, Neurocognitive testing, CSF protein, Psy in CSF	Diagnosis confirmed by: GALC in WBC PSY in DBS and CSF Genotype: 1 MUT + 1 VUS + 2 Pseudo	HSCT
		Sat	Sun	Mon			
2 nd DOL	3 rd DOL	4 th DOL	5 th DOL	6 th DOL	7 th DOL	9 th DOL	24 th DOL

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1st KY Case with Krabbe Disease



2 yrs old

- PSY not normalized (at least not by 2 y/o)
- Sitting, but not walking
- Non-verbal, but expressive
- recurrent autoimmune hemolytic anemia

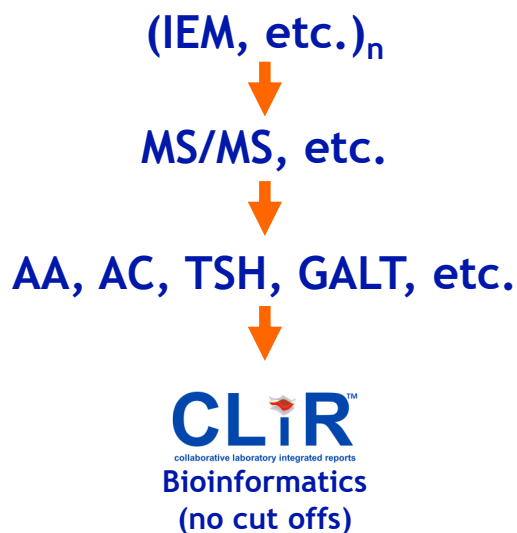


4 years old

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The Newest NBS Model

- Many conditions
- Many tests 
- Many markers
- Pattern recognition (disease risk)



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The Newest NBS Model

- Many *Open*
- Many
- Many
- *(dise*

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ORIGINAL RESEARCH ARTICLE | Genetics in Medicine

Postanalytical tools improve performance of newborn screening by tandem mass spectrometry

Patricia L. Hall, PhD¹, Gregg Marquardt, MSS¹, David M.S. McHugh¹, Robert J. Currier, PhD², Hao Tang, PhD², Stephanie D. Stoway, BS¹ and Piero Rinaldo, MD, PhD¹

Purpose: The purpose of this study was to compare performance metrics of postanalytical interpretive tools of the Region 4 Stork collaborative project to the actual outcome based on cutoff values for individual target conditions were deemed informative when equal or greater to the value representing the first percentile rank of known true-positive cases (17,099 cases in total).

Results: In the study period, the actual false-positive rate and positive predictive value were 0.26 and 10%, respectively. Utilization of the Region 4 Stork tools, simple interpretation rules, and second-tier tests

Conclusion: Region 4 Stork interpretive tools, second-tier tests, and other evidence-based interpretation rules could have reduced false-positive cases by up to 90% in California.

Key Words: cutoff values; newborn screening; postanalytical interpretive tools; second-tier test; tandem mass spectrometry

n

:

T, etc.

(no cut offs)



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Newborn Screening

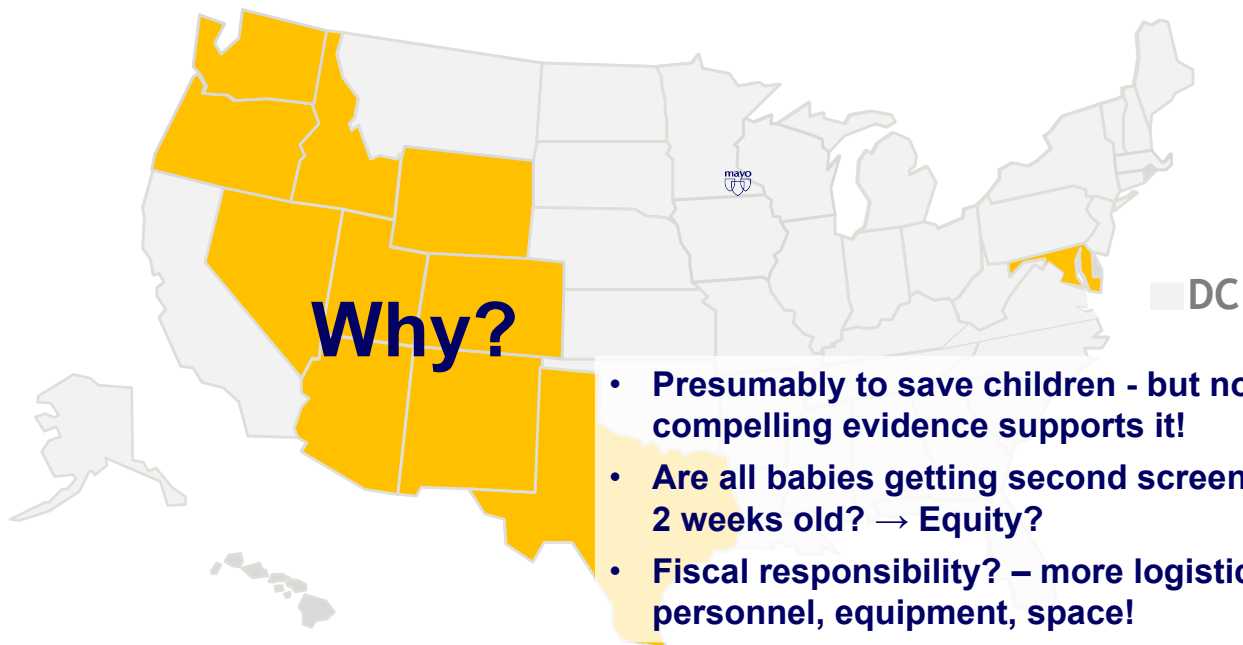
A public health program:

- Aimed at identification of conditions for which early intervention can prevent mortality, morbidity and disabilities.
- Performed by analysis of diagnostic markers in blood spots collected on filter paper on the second day of life (exception: hearing loss and congenital cyanotic heart disease).
- Extent of program determined by each state independently.
- Administered by each state but testing may be out of state
- States may screen twice: by 2 days old + 1-2 weeks old



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“Two Screen” States in 2023



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Newborn Screening

What do screening laboratories like to do on Friday afternoons?

Inform you of a presumptive positive result on one of your newborn patients!



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What to do when you get a call about an abnormal "PKU test"?

- A. I tell myself not to panic!**
- B. I take the necessary information but make sure they mean an elevated Phe, not an abnormal NBS for Pompe disease or something else.**
- C. I ask around if any of my colleagues knows what to do.**
- D. I call the colleague who did not fall asleep during Dr. Matern's presentation.**
- E. I'm self-sufficient and Google for ideas.**
- F. I'm self-sufficient and go to the NCCRCG or ACMG website.**

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The screenshot shows the website for the National Coordinating Center for the Regional Genetics Networks (NCCRCG). The browser address bar displays 'https://nccrcg.org'. A navigation menu is open, highlighting 'ACMG ACT Sheets and Algorithms' with a red arrow. The website header includes a banner for the '2023 Genetics Workforce Survey is Now Live!' and a 'LEARN MORE' button. The main content area features a map of the United States with regional labels: MBRGN, Heartland, Midwest, NYMAG, and SERN. A blue button at the bottom of the main content area says 'CONNECT WITH YOUR RGN >'. The Mayo Clinic logo is visible in the bottom left corner.

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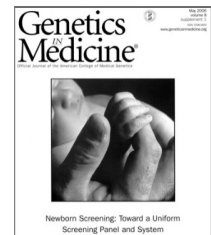
The ACMG Newborn Screening Diagnosis & Follow-up Work Group

- **Geneticists, other specialists and primary care providers involved in NBS for endocrine, hematologic, genetic and metabolic diseases**
- **ACT sheets and diagnostic algorithms**
- **ACT sheets include:**
 - **Information about the analytes and their clinical significance**
 - **Links to informational resources, if needed**
 - **Links to websites that allow identification of regional subspecialists for consultation and referral if desired**

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Development of ACT Sheet Concept

- Began when NBS expansion was apparent
- Too many “fact” sheets
- Need was for tools to support clinical decision-making for all NBS conditions
 - Point of care education and decision support
 - EMR compatibility
 - Pop-up education
 - Clinical decision support (CDS) (i.e. directive support)



From: Michael Watson, PhD; American College of Medical Genetics and National Coordinating Center for Regional Genetics and Newborn Screening Collaboratives; [May 7, 2007](#)

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NBS ACT(ion) Sheets and Diagnostic Algorithms

- For all conditions in uniform panel
- ACMG Board approved and AAP Board endorsed
- Posted on ACMG website
 - Genetics Home Reference, NNSGRC, and many others link to these
- Distributed to NBS labs and programs
 - To accompany all “screen positive” lab reports
- Distributed to RCs to coordinate use with local and regional plans
- Survey of utility presently being conducted

From: Michael Watson, PhD; American College of Medical Genetics and National Coordinating Center for Regional Genetics and Newborn Screening Collaboratives; [May 7, 2007](#)

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The ACMG Newborn Screening Diagnosis & Follow-up Work Group

Harvey Levy, MD (Chair)

Endocrinology

Stephen LaFranchi, MD (OR)
Phyllis Speiser, MD (NY)
Kelly Leight, JD (CARES Found.)

Hematology

James Eckman, MD (GA)
Peter Lane, MD (GA)
Carolyn Hoppe, MD (CA)

Genetics

Gary Cutting, MD (MD)
Cynthia Morton, PhD (MA)
Richard Smith, MD (IA)

HRSA

Marie Mann, MD, MPH (DC)
Michelle Lloyd-Puryear, MD, PhD (DC)

Michael Watson, PhD (Project Dir.)

Metabolism

Gerard Berry, MD (PA)
Stephen Goodman, MD (CO) ←
Harvey Levy, MD (MA)
Deborah Marsden, MD (MA)
Dietrich Matern, MD, PhD (MN)
William Nyhan, MD (CA)

Primary Care

Danielle Laraque, MD (NY)
Barbara Yawn, MD (MN)

Newborn Screening

Julie Miller, MS (NE)
Kenneth Pass, PhD (NY)
Bradford Therrell, PhD (TX) ←

From: Michael Watson, PhD; American College of Medical Genetics and National Coordinating Center for Regional Genetics and Newborn Screening Collaboratives; May 7, 2007

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Conclusions

- **ACT sheets and algorithms were designed to aid in the most appropriate and timely evaluation and treatment of newborns with abnormal newborn screening results**
- **Algorithms probably work best when clinician and laboratorians “talk”**
- **Constructive feedback to the ACMG is encouraged for continued improvement**
- **ACT sheets and algorithms are freely available for adaptation to regional needs (practice, hospital, screening program)**
- **Updates must be communicated effectively to users and adopters**

From: Michael Watson, PhD; American College of Medical Genetics and National Coordinating Center for Regional Genetics and Newborn Screening Collaboratives; May 7, 2007

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ACMG
American College of Medical Genetics and Genomics

Newborn Screening ACT Sheet
[Decreased galactocerebrosidase, elevated psychosine]
Krabbe Disease (infantile form)

Differential Diagnosis: Saposin A deficiency.

Condition Description: Krabbe disease (globoid cell leukodystrophy) is a lysosomal galactocerebrosidase, resulting in impaired turnover of myelin with subsequent dys oligodendrocytes and Schwann cells. The infantile form usually presents before the age asymptomatic and, if untreated, survival beyond age 2 years is uncommon.

You Should Take the Following IMMEDIATE Actions:

- Inform family of the newborn screening result.
- Ascertain clinical status (newborns are asymptomatic).
- Consult with or refer to pediatric metabolic and transplant specialist the sa
- Evaluate the newborn (perform physical examination, newborns are expc
- Initiate confirmatory/diagnostic testing, as recommended by the specialist.
- Provide the family with basic information about Krabbe disease and its ma
- Report final diagnostic outcome to newborn screening program.

Diagnostic Evaluation: Leukocyte galactocerebrosidase enzyme assay and meas psychosine concentration. Decreased enzyme activity is suggestive of Krabbe disease. C galactocerebrosidase activity and psychosine concentration predict the phenotype (t Krabbe disease). Molecular genetic testing can confirm the diagnosis.

Clinical Considerations: The clinical presentation of Krabbe disease ranges from a form to more slowly progressive later-onset variants. All forms of Krabbe disease an but the age of onset and rate of progression vary widely. The only available therapy transplantation that is most effective if performed before 30 days of life in patients t the onset of clinical symptoms in the late-onset forms. Gene therapy and other clinic A deficiency has been described in <10 patients, is clinically very similar to Krabbe d newborn screening.

Additional Information:

[How to Communicate Newborn Screening Results](#)
[Gene Reviews](#)
[Medline Plus](#)
[Condition Information for Families- HRSA Newborn Screening Clearinghou](#)

Referral (local, state, regional, and national):

[Find a Genetics Clinic Directory](#)
[Genetic Testing Registry](#)

ACMG
American College of Medical Genetics and Genomics

Newborn Screening ACT Sheet
[Decreased galactocerebrosidase, mildly elevated psychosine]
Krabbe Disease (late-onset form)

Differential Diagnosis: Saposin A deficiency.

Condition Description: Krabbe disease (globoid cell leukodystrophy) is a lysos galactocerebrosidase, resulting in impaired turnover of myelin with subsequen oligodendrocytes and Schwann cells. There is variability in severity and age of .

You Should Take the Following Actions:

- Inform family of the newborn screening result.
- Ascertain clinical status (newborns are asymptomatic).
- Consult with a pediatric metabolic specialist.
- Evaluate the newborn (perform physical examination, newborns are e
- Initiate confirmatory/diagnostic testing, as recommended by the speci
- Provide the family with basic information about Krabbe disease and it
- Report final diagnostic outcome to newborn screening program.

Diagnostic Evaluation: Leukocyte galactocerebrosidase enzyme assay and a psychosine concentration. Decreased enzyme activity is suggestive of Krabbe d exclude pseudodeficiency, which causes decreased enzyme levels without disea galactocerebrosidase activity and psychosine concentration predict the phenoty Krabbe disease). Molecular genetic testing can confirm the diagnosis.

Clinical Considerations: This screening result is more likely associated with t but all forms of Krabbe disease are associated with leukodystrophy with age of widely. The only available therapy is hematopoietic stem cell transplantation tl of clinical symptoms. Gene therapy and other clinical trials may be available. S in <10 patients, is clinically very similar to Krabbe disease, and may be detecta

Additional Information:

[How to Communicate Newborn Screening Results](#)
[Gene Reviews](#)
[Medline Plus](#)
[Condition Information for Families- HRSA Newborn Screening Clearing](#)

Referral (local, state, regional, and national):

[Find a Genetics Clinic Directory](#)
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ACMG
American College of Medical Genetics and Genomics

Krabbe disease: galactocerebrosidase deficiency

Key:

- Actions are shown in shaded ovals; results are in the unshaded ovals
- Diagnostic outcomes are shown in boxes
- Dashed line reflects an optional test.

Abbreviations:
GALC = galactocerebrosidase
RBC = red blood cells

This practice resource is designed primarily as an educational resource for medical geneticists and other clinicians to help them provide quality medical services. Adherence to this practice resource is completely voluntary and does not necessarily assure a successful medical outcome. This practice resource should not be considered inclusive of all proper procedures and tests or exclusive of other procedures and tests that are reasonably directed to obtaining the same results. In determining the propriety of any specific procedure or test, the clinician should apply his or her own professional judgment to the specific clinical circumstances presented by the individual patient or specimen. Clinicians are encouraged to document the reasons for the use of a particular procedure or test, whether or not it is in conformance with this practice resource. Clinicians also are advised to take notice of the date this practice resource was adopted, and to consider other medical and scientific information that becomes available after that date. It also would be prudent to consider whether intellectual property interests may restrict the performance of certain tests and other procedures.

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Newborn Screening Going Forward

OPINION

Neonatal screening by DNA microarray: spots and chips

Nancy S. Green and Kenneth A. Pass

2005

NATURE REVIEWS | GENETICS

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NEWS RELEASES

Wednesday, September 4, 2013

2013

NIH program explores the use of genomic sequencing in newborn healthcare

Can sequencing of newborns' genomes provide useful medical information beyond what current newborn screening already provides? Pilot projects to examine this important question are being funded by the *Eunice Kennedy Shriver* National Institute of Child Health and Human Development (NICHD) and the National Human Genome Research Institute (NHGRI), both parts of the National Institutes of Health. Awards of \$5 million to four grantees have been made in fiscal year 2013 under the Genomic Sequencing and Newborn Screening Disorders research program. The program will be funded at \$25 million over five years, as funds are made available.

Each of the new awards will consist of three parts: Genomic sequencing and analysis; research related to patient care; and the ethical, legal and social implications of using genomic information in the newborn period. Teams of researchers will work to further the understanding of disorders that appear in newborns and to improve treatments for these diseases using genomic information.

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Parental Views on Expanded Newborn Screening Using Whole-Genome Sequencing

Galim Joseph, PhD, Flavia Chen, MPH, Julie Harris-Wal, PhD, MPH,¹ Jennifer M. Puck, MD, Charlotte Young, BS, Barbara A. Koenig, PhD*Pediatrics*. 2016;137(s1):e20153731H

abstract **BACKGROUND AND OBJECTIVE:** The potential application of whole-genome sequencing (WGS) to state-mandated standard newborn screening (NBS) challenges the traditional public health approach to NBS and raises ethical, policy, and clinical practice issues. This article examines the perspectives and values of diverse healthy pregnant women and parents of children diagnosed with a primary immunodeficiency disorder about traditional NBS and expanded NBS with the use of WGS.

METHODS: We conducted 4 focus groups (3 in English and 1 in Spanish) with socioeconomically and ethnically diverse pregnant women ($n = 26$), and a comparison group with parents of children diagnosed with a primary immunodeficiency disorder ($n = 5$).

RESULTS: Pediatric policy-relevant themes that emerged from our analysis of the focus group data are presented within 4 categories: (1) perspectives on traditional NBS, (2) informed consent, (3) return of results, and (4) storage and retrieval of results. Analyses indicate that study participants desired greater inclusion in the NBS process. Despite an optimistic orientation to the potential benefits and limited harms likely to result from genomic applications of NBS, parents voiced concerns about privacy and control over test results, limited trust in the medical system and the state-run NBS program, informed these concerns.

CONCLUSIONS: Expanded NBS with WGS for pediatricians may require management of more genetic conditions, including mutations that convey risk to both the child and parents for adult-onset disorders, and an informed-consent process to manage the genomic data and storage of blood spots. Attention to how these technologies are understood in diverse populations is needed for effective implementation.



Interpretation of Genomic Sequencing Results in Healthy and Ill Newborns: Results from the BabySeq Project

Ozge Ceyhan-Birsoy,^{1,2} Jaelyn B. Murry,^{2,4} Kalolina Machini,^{2,4} Matthew S. Lebo,^{2,3,4,11} Timothy W. Yu,^{5,6,9} Shawn Fayer,⁷ Cassie A. Genetti,⁸ Talia S. Schwartz,⁸ Pankaj B. Agrawal,^{4,5,8} Richard B. Parad,^{4,9} Ingrid A. Holm,^{4,5} Amy L. McGuire,¹⁰ Robert C. Green,^{4,7,11} Heidi L. Rehm,^{2,3,4,11,12} Alan H. Beggs,^{4,5,*} and The BabySeq Project Team

Genomic sequencing provides many opportunities in newborn clinical care, but the challenges of interpreting and reporting newborn genomic sequencing (nGS) results need to be addressed for its broader and effective application. The BabySeq Project is a pilot randomized clinical trial that explores the medical, behavioral, and economic impacts of nGS in well newborns and those admitted to a neonatal intensive care unit (NICU). Here we present childhood-onset and actionable adult-onset disease risk, carrier status, and pharmacogenomics findings from nGS of 493 newborns in the BabySeq Project. nGS revealed a risk of childhood-onset disease in 13/493 (2.6%) newborns; none of the disease risks were anticipated based on the infants' known clinical or family histories. nGS also revealed actionable adult-onset disease risk in 3/85 (3.5%) newborns whose parents consented to receive this information. Carrier status for recessive diseases and pharmacogenomics variants were reported in 88% and 5% of newborns, respectively. Additional indication-based analyses were performed in 29/32 (91%) NICU newborns and 6/127 (5%) healthy newborns who later had presentations that prompted a diagnostic analysis. No variants that sufficiently explained the reason for the indications were identified; however, suspicious but uncertain results were reported in five newborns. Testing parental samples contributed to the interpretation and reporting of results in 13/159 (8%) newborns. Our results suggest that nGS can effectively detect risk and carrier status for a wide range of disorders that are not detectable by current newborn screening assays or predicted based on the infant's known clinical or family history, and the interpretation of results can substantially benefit from parental testing.

The American Journal of Human Genetics 104, 76–93, January 3, 2019

Perceived Benefits, Risks, and Utility of Newborn Genomic Sequencing in the BabySeq Project

Stacey Pereira, PhD,¹ Jill Oliver Robinson, MA,¹ Amanda M. Gutierrez, BA,¹ Devan K. Petersen, MPH,¹ Rebecca L. Hsu, BA,¹ Caroline H. Lee,² Talia S. Schwartz, BA,³ Ingrid A. Holm, MD, MPH,⁴ Alan H. Beggs, PhD,^{5,*} Robert C. Green, MD, PhD,^{4,6,7} Amy L. McGuire, JD, PhD,⁸ on behalf of The BabySeq Project Group*Pediatrics* 2019;143;S6

abstract **BACKGROUND AND OBJECTIVES:** There is interest in applying genomic sequencing (GS) to newborns' clinical care. Here we explore parents' and clinicians' attitudes toward and perceptions of the risks, benefits, and utility of newborn GS compared with newborn screening (NBS) prior to receiving study results.

METHODS: The BabySeq Project is a randomized controlled trial used to explore the impact of integrating GS into the clinical care of newborns. Parents ($n = 493$) of enrolled infants ($n = 309$) and clinicians ($n = 144$) completed a baseline survey at enrollment. We examined between-group differences in perceived utility and attitudes toward NBS and GS. Open-ended responses about risks and benefits of each technology were categorized by theme.

RESULTS: The majority of parents (71%) and clinicians (51%) agreed that there are health benefits of GS, although parents and clinicians agreed more that there are risks associated with GS (35%, 70%) than with NBS (19%, 39%; all $P < .05$). Parents perceived more benefit and less risk of GS than did clinicians. Clinicians endorsed concerns about privacy and discrimination related to genomic information more strongly than did parents, and parents anticipated benefits of GS that clinicians did not.

CONCLUSIONS: Parents and clinicians are less confident in GS than NBS, but parents perceive a more favorable risk/benefit ratio of GS than do clinicians. Clinicians should be aware that parents' optimism may stem from their perceived benefits beyond clinical utility.

Also identified one patient with:

- nonsyndromic hearing loss
 - nonclassic CAH
 - partial Biotinidase deficiency.
- NBS was normal in all 3 infants.**

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LETTERS

https://doi.org/10.1038/s41591-022-0964-5



Check for updates

The role of exome sequencing in newborn screening for inborn errors of metabolism

Aashish N. Adhikari, Renata C. Gallagher, Yaogong Wang, Robert J. Currier, George Amatuni, Laia Bassaganya, Flavia Chen, Kunal Kundu, Mark Kvale, Sean D. Mooney, Robert L. Nussbaum, Savanna S. Randi, Jeremy Sanford, Joseph T. Shieh, Rajgopal Srinivasan, Uma Sunderam, Hao Tang, Dedeepya Vaka, Yanguan Zou, Barbara A. Koenig, Pul-Yan Kwok, Neil Risch, Jennifer M. Puck, and Steven E. Brenner

Public health newborn screening (NBS) programs provide population-scale ascertainment of rare, treatable conditions that require urgent intervention. Tandem mass spectrometry (MS/MS) is currently used to screen newborns for a panel of rare inborn errors of metabolism (IEMs)...

newborns. Yet, population-scale studies to establish performance characteristics of sequencing for NBS have not been reported. NBS IEMs provide an ideal model for evaluating the role of sequencing in population screening because most are Mendelian disorders affecting well-understood biochemical pathways...

Effective population-level NBS must rapidly identify the few individuals at risk of disease with extraordinary sensitivity, high specificity and limited manual review...

Genomic sequencing, now commonly used for diagnosis of rare disorders, has been recommended for nearly all seriously ill children in intensive care units, proposed for all newborns to personalize their medical care and marketed for screening

parameters on the validation set to derive a customized, robust, sensitive pipeline for reporting potential disease-causing variants in IEM genes in a screening context...

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LETTERS

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ARTICLE

A genome sequencing system for universal newborn screening, diagnosis, and precision medicine for severe genetic diseases

Stephen F. Kingsmore, Laurie D. Smith, Chris M. Kurnard, Matthew Bainbridge, Sergey Batokov, Wendy Benson, Eric Bilincov, Sara Caylor, Christina Chambers, Guillermo Del Angel, David P. Dimmock, Yan Ding, Katarzyna Ellsworth, Annette Feigenbaum, Erwin Frise, Robert C. Green, Lucia Guidugli, Kevin P. Hall, Christian Hansen, Charlotte A. Hobbs, Scott D. Kahn, Mark Kiel, Lucia Van Der Knaap, Chad Kriwof, Yong H. Kwon, Lakshminarasimha Madhavara, Jennie Le, Sebastien Lefebvre, Rebecca Mardach, William R. Mowrey, Danny Oh, Mallory J. Owen, George Powley, Gunter Scharer, Seth Shelmut, Mari Tokita, Shyamal S. Mehta, Albert Ortol, Stavros Papadopoulos, James Perry, Edwin Rosales, Erica Sanford, Steve Schwartz, Duke Tran, Martin G. Reese, Meredith Wright, Narayan Veeraghavan, Kristen Wigby, Mary J. Willis, Aaron R. Wolen, and Thomas Defay

and informed by root cause analysis. In 119 affected children who had been diagnosed by rWGS, 87% were positive for NBS-rWGS. The diagnostic sensitivity of NBS-rWGS can be further increased by inclusion of variants identified by

Cost effectiveness studies of NBS-rWGS have not yet been performed. While NBS-rWGS is intended to supplement NBS-MS, not replace it, the current cost of NBS-MS for the 35 core disorders on the RUSP provides a reference point

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for what is likely to be acceptable for public-health-funded NBS-rWGS. Most states publish the fees charged for NBS-MS, which represent part of the total cost. The highest such fee is \$220 per newborn. Diagnostic rWGS costs RCGM ~\$8,500 per newborn. However, the interpretation burden of NBS-rWGS is about one thousandth that of Dx-rWGS and several biotechnology companies have indicated that \$100 rWGS will be possible in the relatively near future.

we achieved by combining screening, diagnosis, large genome-phenotype datasets, and learning feedback loops. We invite groups worldwide to join the BeginNGS (Newborn Genomic Sequencing) consortium in implementation studies of NBS-rWGS in diverse populations.

Data and code availability

Consented proband and parent data analyzed in this study and non-human subjects data generated during this study are available at the Longitudinal Pediatric Data Resource (LPDR) under access-informed methods for newborn screening, diagnosis, and virtual, acute management guidance for 388 diseases with effective treatments and report analytic performance and clinical utility in large retrospective datasets.

Screening Panel (RUSP)—increased from 27 to 35, and the number of affected infants identified increased from 6,439 to 6,466. However, there are ~7,000 known genetic diseases and hundreds of targeted treatments that

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Genomic Newborn Screening?

- Who should do this? Public Health NBS Labs? Contracted academic or private labs? Partnership: NBS lab does test, consultant or AI interprets results?
- What to report?
 - Actionable genotype* for disease with onset in infancy only?
 - Actionable genotype* for disease with onset at any age (e.g. cancer predisposition, pharmacogenomics)?
 - Any 'pathogenic' genotype* for a disease treatable or not?
 - Any genotype (pathogenic vs. variants of uncertain significance)?
- What platform or reagent to use (performance not uniform)?

*likely not equitable for most genes and for several more years!

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Realistic NBS Forecast to the 2030s

- Conditions nominated/added to the RUSP and the relevant screening strategies will be (better) defined, also paying attention to equity.
- Conditions could be added to RUSP at a pace of 1 – 2 per year (<20 by 2032) facilitated by grouping 'like' conditions (e.g. Mucopolysaccharidoses), but current ACHDNC voting members seem to put the brakes on.
- More (gene) therapies will become available, but at moderate pace.
- Genetic/genomic testing will become more commonplace in NBS, but biochemical testing will not be replaced.
- Rational (!) regionalization of NBS in the US, despite advantages, will not happen.

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- Rational (!) regionalization of NBS in the US, despite advantages, will not happen.

But remember: addition to RUSP ≠ state implementation

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SUMMARY

- Newborn screening is one of the most successful public health programs.
- In the US, mechanisms are in place to expand the RUSP based on evidence (although evidence usually emerges only once screening has started).
- The most efficient and effective approach to newborn screening relies primarily on biochemical genetic assays – for 1st and 2nd tier testing.
- Molecular genetic testing/genomics for newborn screening is currently of limited value given high cost, time to complete analysis, and the frequency of genotypes of uncertain significance.
- The biggest limitation to expansion of newborn screening is not laboratory testing and result interpretation but the lack of treatment options.

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