



 Wisconsin Newborn Screening Laboratory

# Molecular Technology in Newborn Screening: SCID and Beyond

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**Mei Baker, M.D., FACMG**

Associate Professor, Department of Pediatrics

Co-Director, NBS Laboratory at WSLH

University of Wisconsin School of Medicine and Public Health



DEPARTMENT OF  
**Pediatrics**

UNIVERSITY OF WISCONSIN SCHOOL OF MEDICINE AND PUBLIC HEALTH

# History of NBS Molecular Testing

- **1994**
  - Washington: hemoglobin second tier testing (Hb S, C, and E) by RELP
  - Wisconsin: *CFTR* mutation analysis for F508del
- **1998**
  - New England: 2 *GALT* mutations (Q & N) by RFLP
- **1999**
  - New England: *MCADD* (c.985A>G) by RFLP
- **2005**
  - Wisconsin: *MSUD* (p.Y438N) by Tetra-primer ARMS-PCR

# History of NBS Molecular Testing

- **2006**
  - New York: Krabbe disease (3 polymorphisms & 5 mutations) by DNA sequencing
- **2008**
  - Wisconsin: TREC assay for SCID screening by Real-time PCR
    - *1<sup>st</sup> use of molecular test as a primary test for population screen*
- **2010**
  - 36 NBSPs in US use molecular testing for CF

# Severe Combined Immunodeficiency (SCID)

Then...



Now...

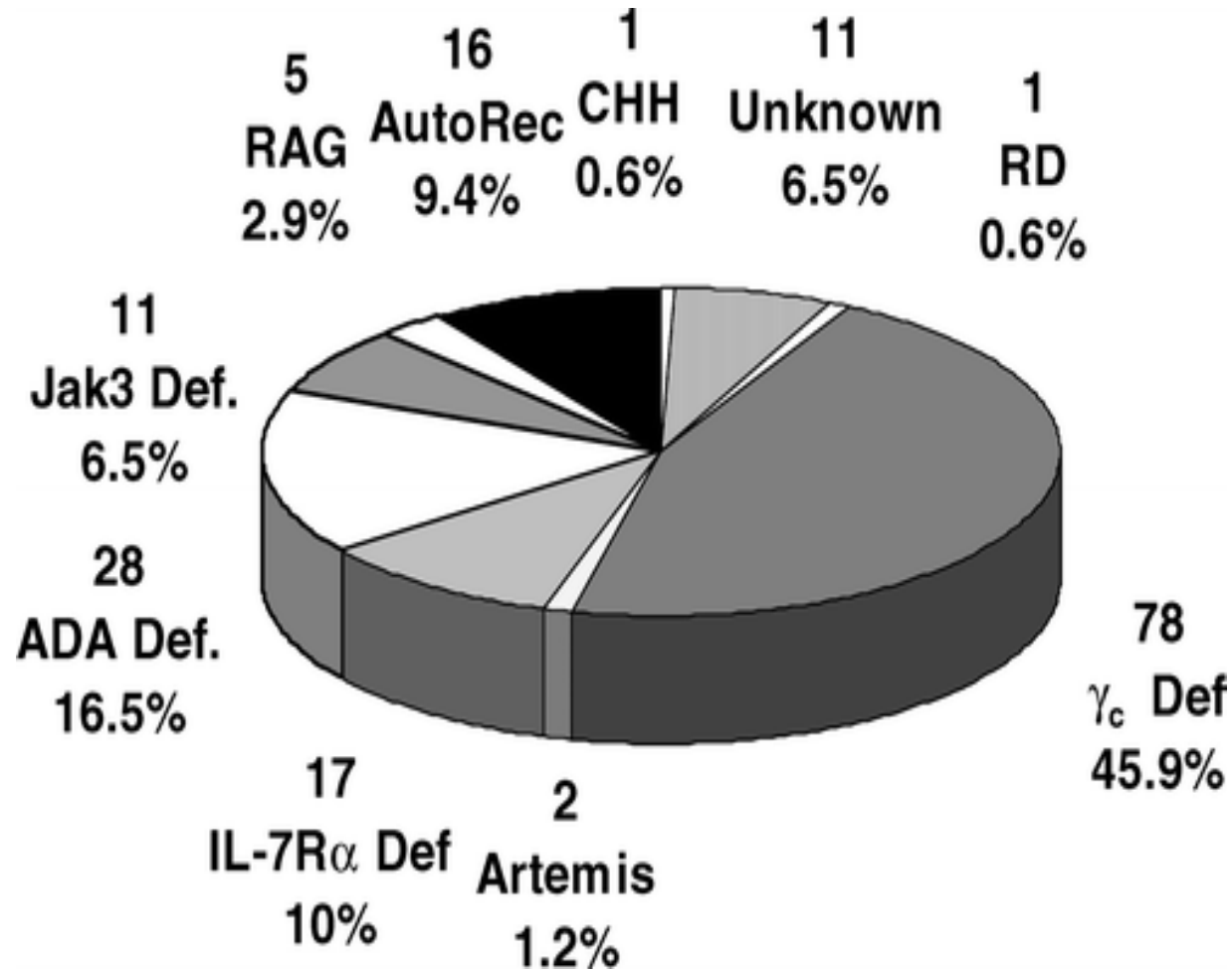


# Severe Combined Immunodeficiency (SCID)

- **Infections in first year of life**
  - recurrent, etiology bacterial, viral and fungal
  - persistent despite routine treatment
  - severe--including sepsis, meningitis
  - opportunistic pathogens, such as PCP (pneumonia)
- **Failure to thrive, chronic diarrhea**
- **T cells decreased or absent**
  - poor proliferation *in vitro* to mitogens
- **B cells absent or non-functional**
  - low Ig's after maternal IgG wanes; no specific antibody responses
- **Fatal without immune reconstitution**

# SCID Genetic Analysis

- X-linked SCID is most common form (males)
- Specific gene defect can be found in 80% of cases (15 genes known)
- Clinical applications:
  - Carrier and prenatal dx
  - Predict response to BMT
  - Gene therapy



# Available Curative Treatment Modalities for SCID

- Bone Marrow Transplantation
- Gene Therapy (X-linked and adenosine deaminase deficiency SCID)

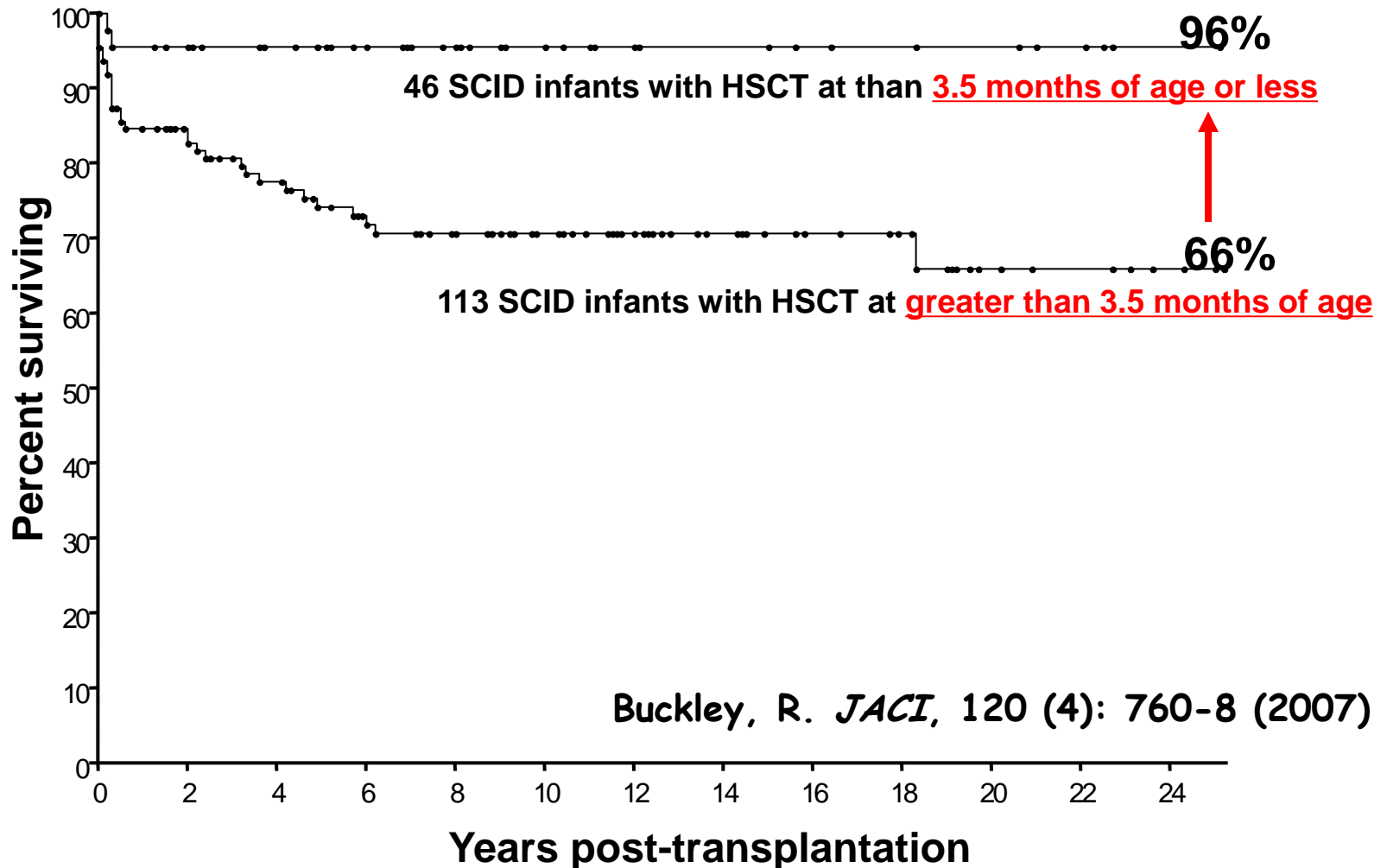
# Does SCID fulfill NBS criteria?

- Prevalence of the disease (1:100,000 or greater)
  - SCID: 1:66,000 (conservative estimate)
- Can the disorder be detected by routine physical exam?
  - SCID: No, SCID baby appears normal at birth.
- Does the disorder have a short asymptomatic period after birth?
  - SCID: Yes, SCID baby can be protected by passive maternal immunity.
- Does the disease cause serious medical complications?
  - SCID: Yes, universally fatal within the first year of life
- Is there potential for successful treatment?
  - SCID: Yes, hematopoietic stem cell transplantation
- Is there a confirmatory test?
  - SCID: Yes, lymphocyte subpopulation analysis (flow cytometry)
- Does early intervention leads better outcome?
  - SCID: Yes!
- Is there a screening test?
  - SCID: Yes, measurement of TRECs using real-time qPCR



# SCID: Benefits of Early Diagnosis

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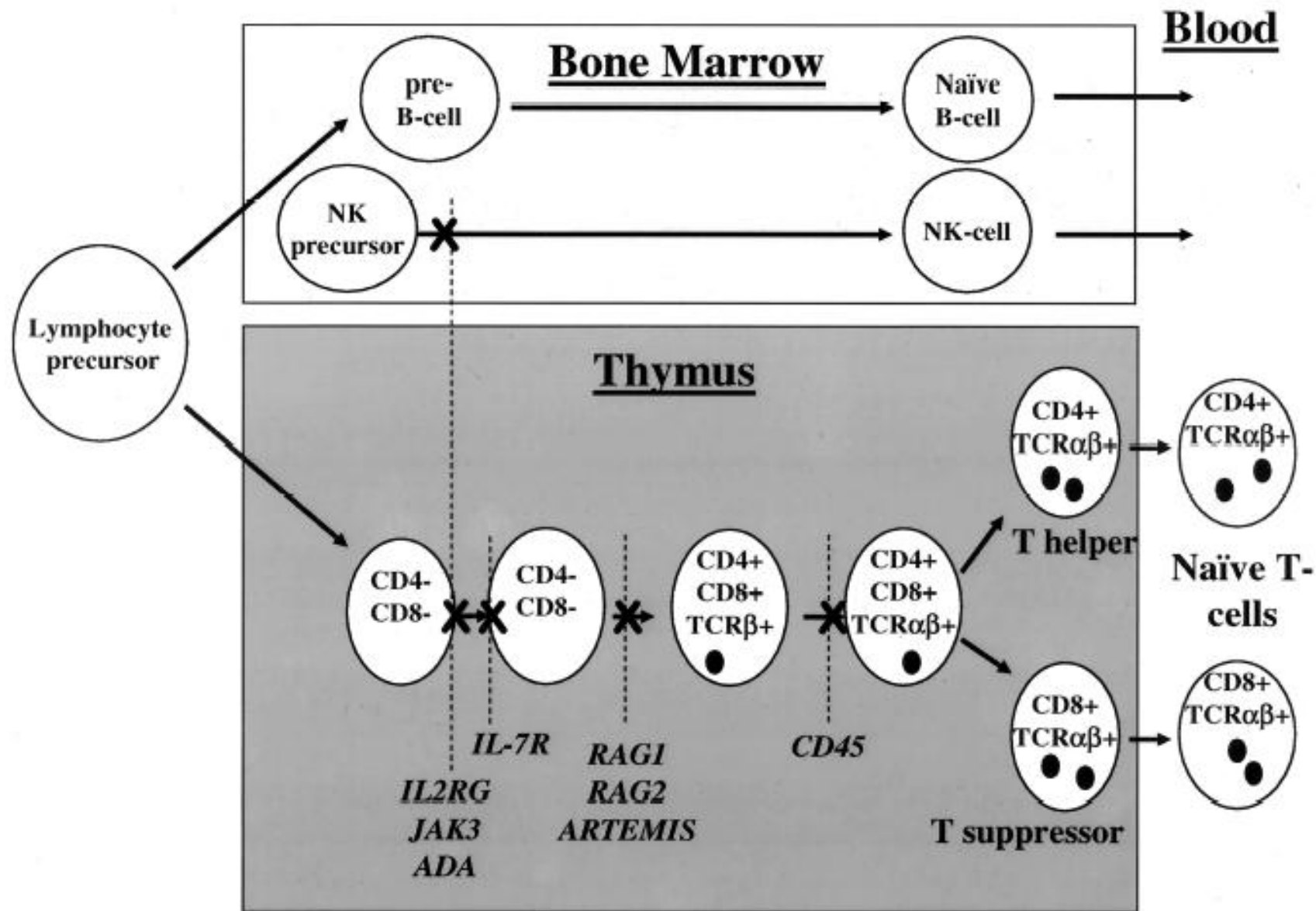


# Screening for SCID in Newborns

## Considerations

- Many genes
- Many mutations in each known gene
- Some genotypes still not known

# TRECs are reduced in nearly ALL forms of SCID



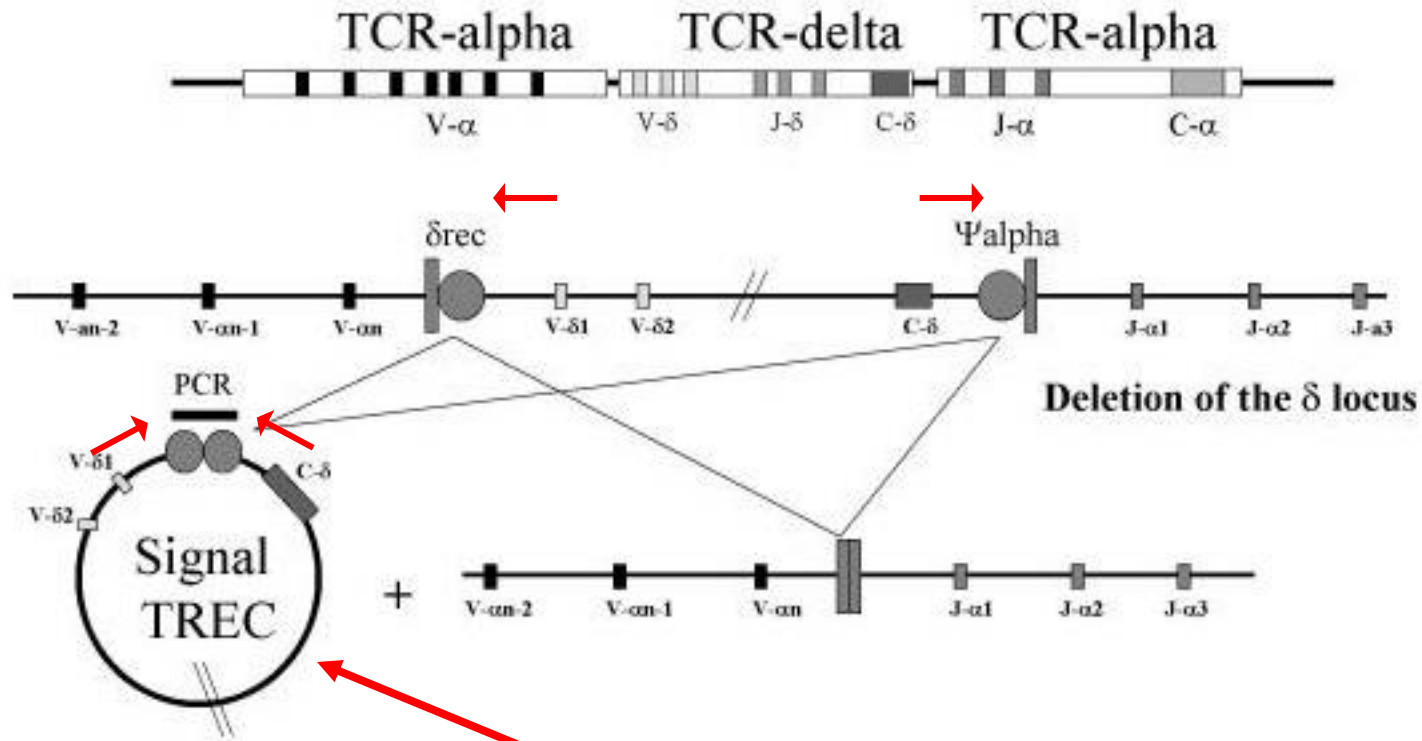
# T-cell Generation in Newborns

- **Two mechanisms:**
  - Thymic output
  - Postthymic T-cell proliferation
  
- **Consequences:**
  - Majority of T-cells are naïve T cells in newborns.
  - TREC s are diluted out, and 10% T cells contain TRECs in newborns.

Schönland et al. *Blood*.2003; 102: 1428-1434

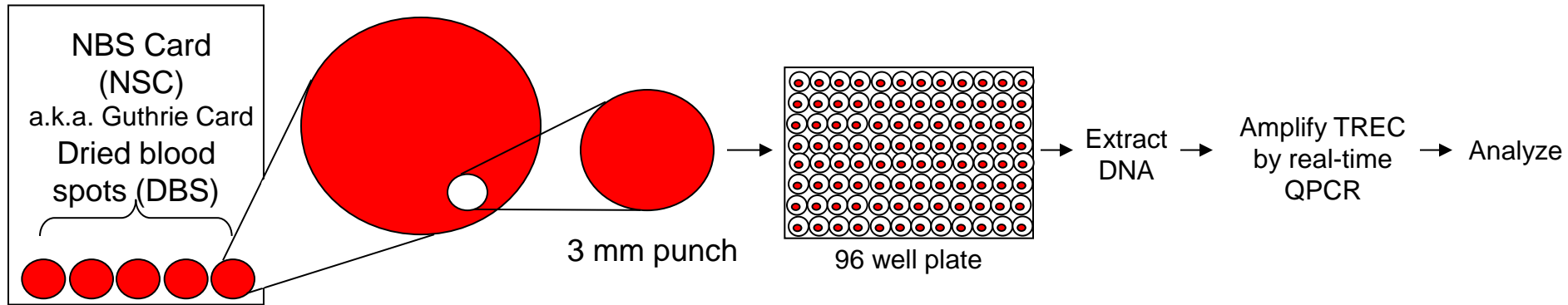
Gent et al, *Clinical Immunology*. 2009; 133: 95–107

# T Cell Receptor Recombination During Development in the Thymus



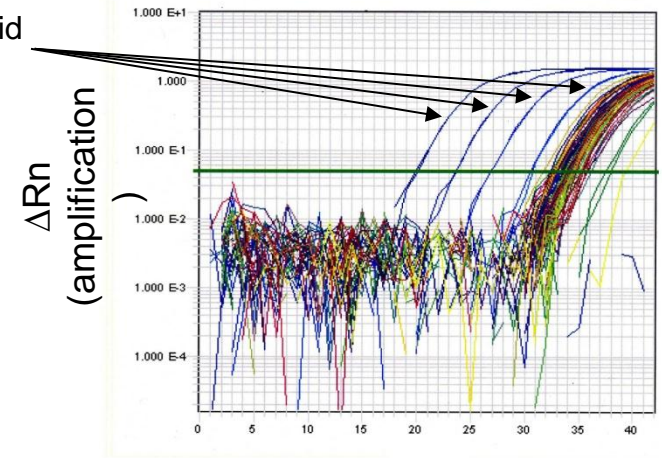
Generation of T cell receptor excision circles (TRECs) occur in >70% of all new (naïve) T cells and can be detected by PCR

# Overall Analysis Scheme

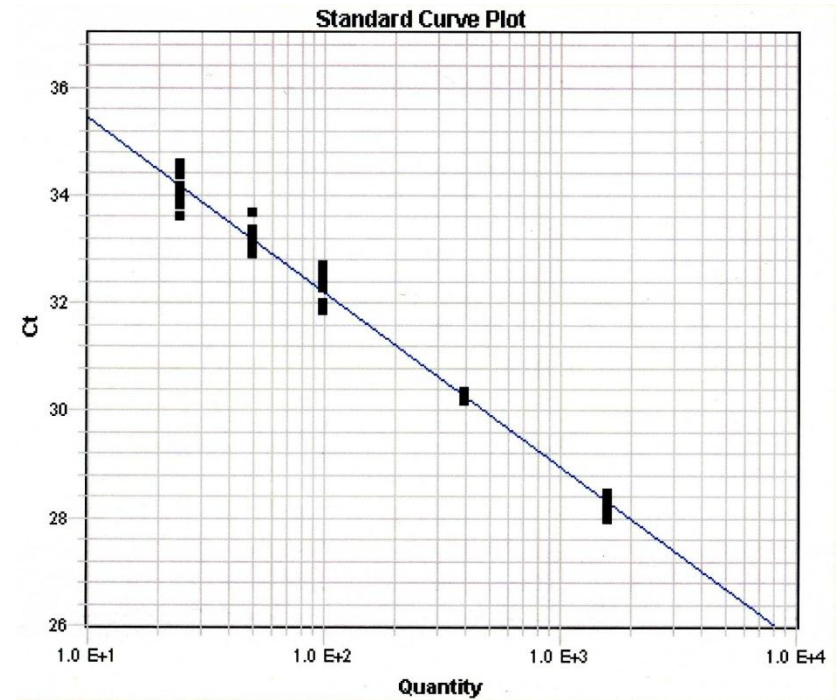
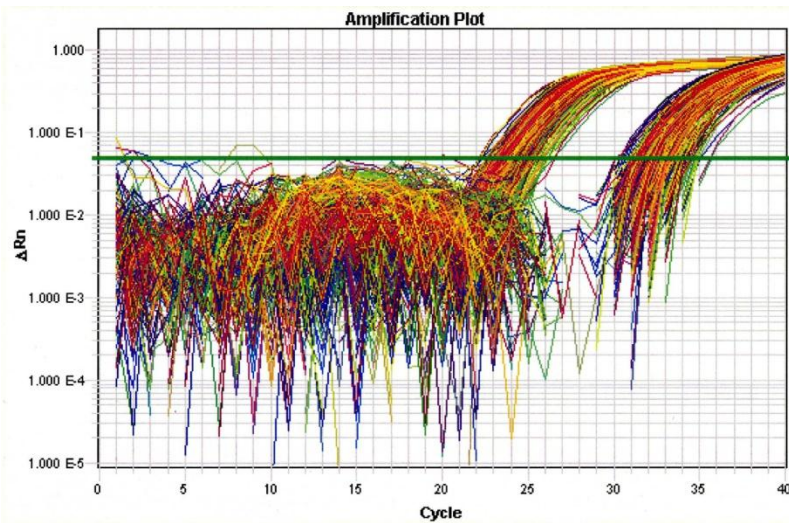


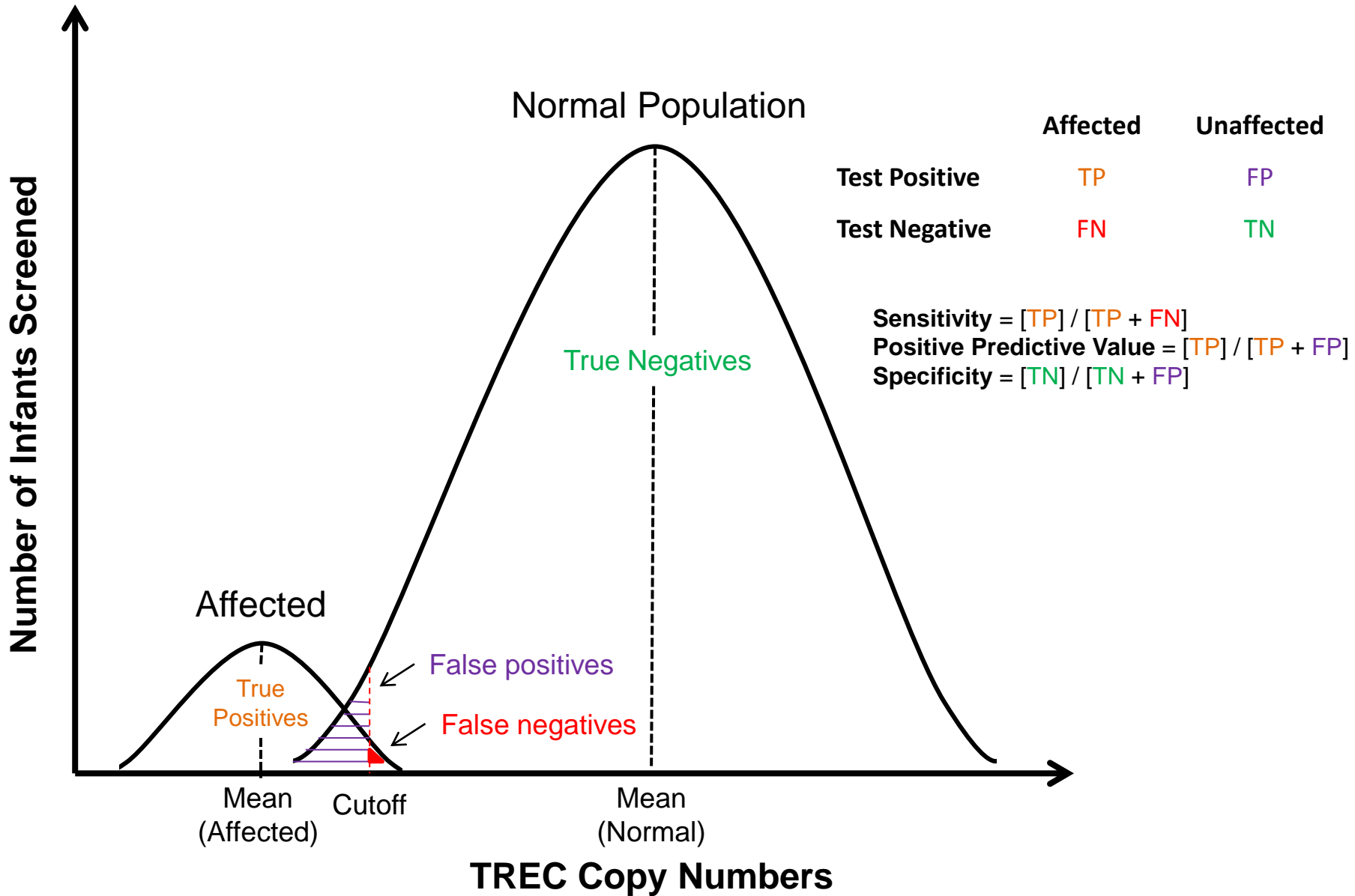
ABI 7900HT Fast Real-Time PCR System

TREC plasmid calibrators



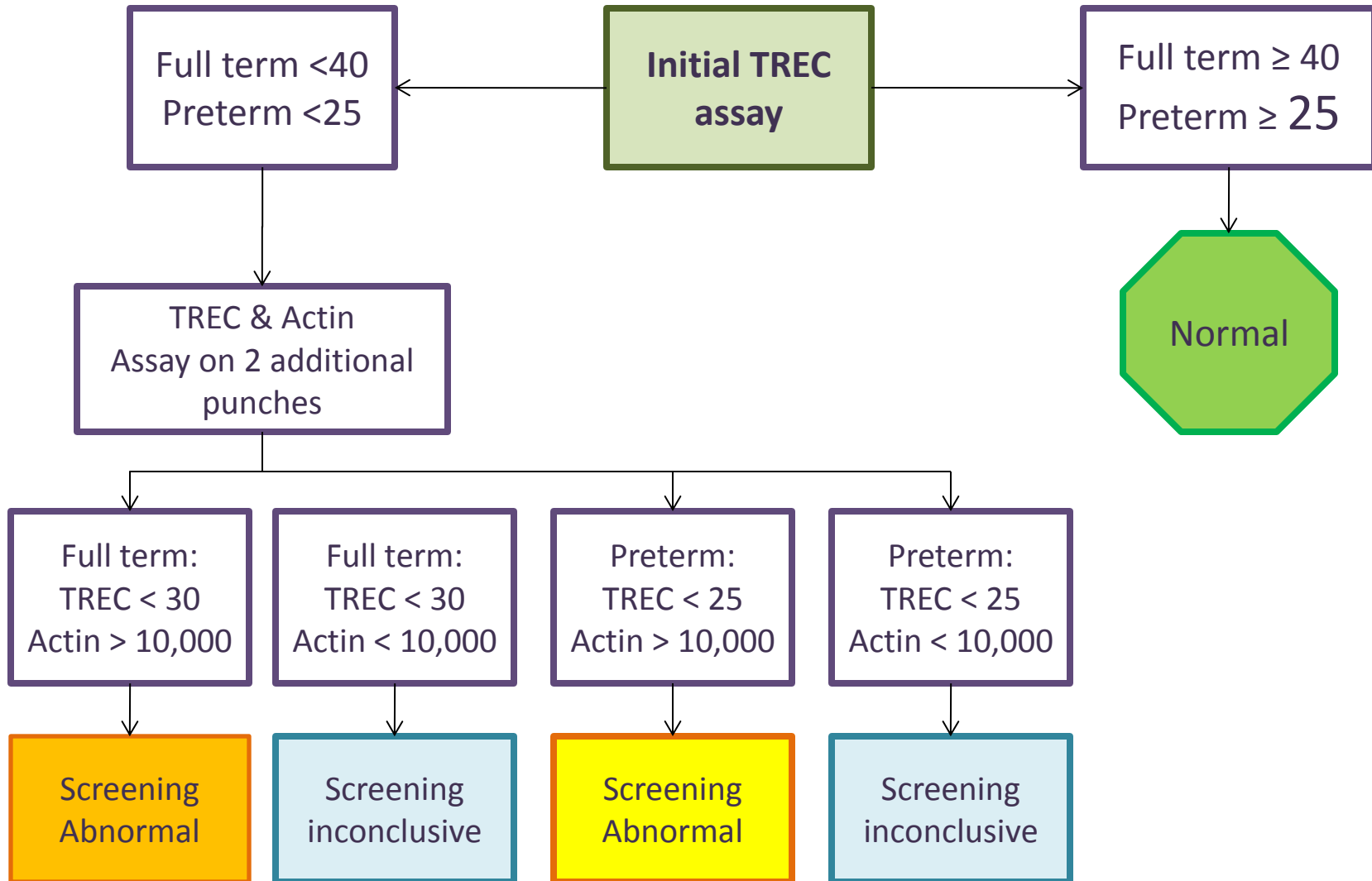
# Multiplexing \_384-well Plate



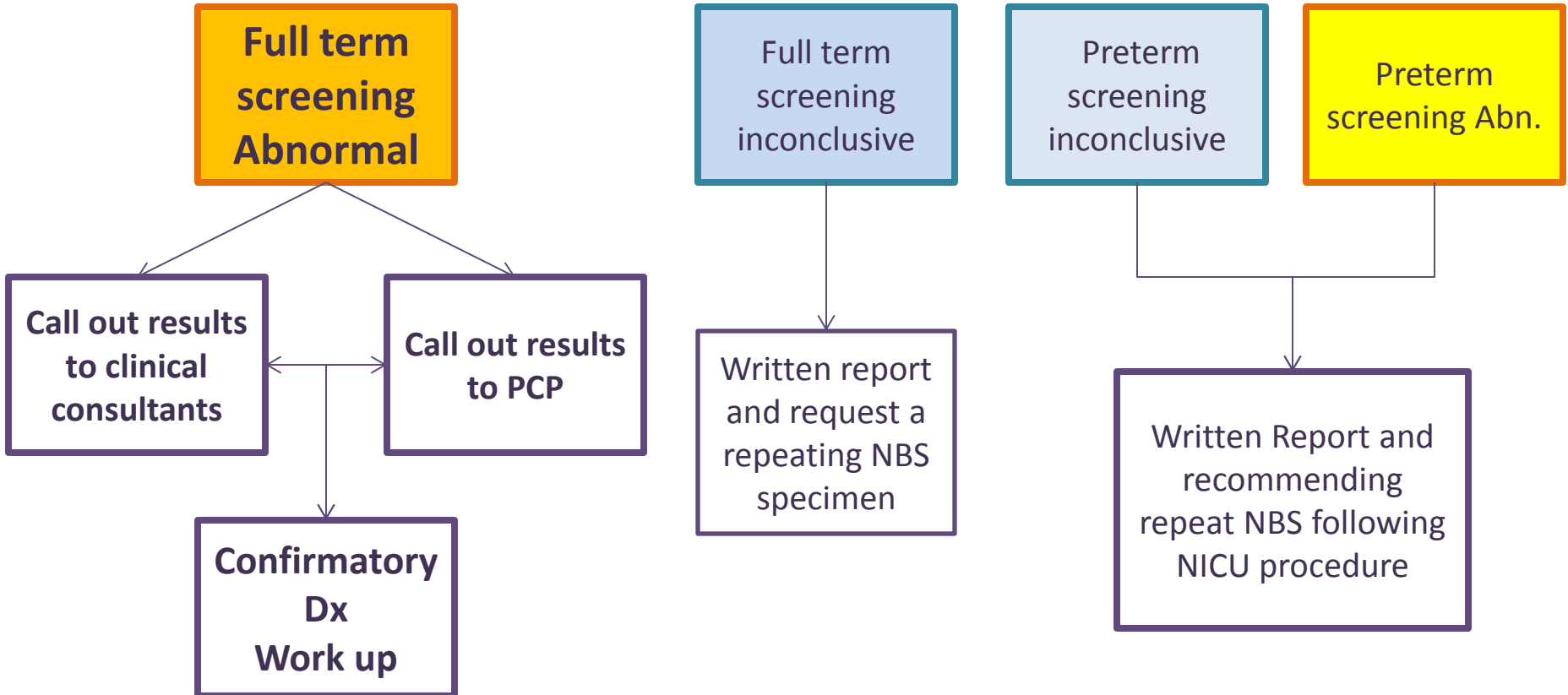




# SCID Testing Algorithm



# SCID Reporting Algorithm



# Confirmatory testing

- Flow cytometry  
lymphocyte subset enumeration for T , B and NK cell  
quantitation
- Lymphocyte (T and/or B) proliferation tests
- Quantitative immunoglobulin assessment (IgG, IgA, IgM and IgE)
- HIV testing (to rule out secondary causes of T-cell lymphopenia)
- Genetics testing
- Others: enzymes, Fluorescence *in situ* hybridization (FISH)

# Special Considerations

- TREC copy numbers
  - Measurement units
  - DNA extraction
  - Calibrators
- TREC assay platform
  - Multiplexing vs. single target
  - 384-well vs. 96-well
- Automation
- QA/QC issues
- Premature Newborns

# Wisconsin Experience

*(January 1, 2008- December 31, 2012)*

<b>Infants Screened:</b>	<b>340,037</b>
- Premature (< 37 wks)	30,664
- Full term	309,373
<b>Abnormal results:</b>	<b>246</b>
- Premature (<37 wks)	147 (0.04%)
- Full term	99 (0.03%)
<b>Inconclusive Results:</b>	<b>472</b>
- Premature (<37 wks)	382 (0.11%)
- Full term	90 (0.03%)

**Total number of flow cytometry referral: 108**

# Wisconsin Experience

*(January 1, 2008- December 31, 2012)*

## *Severe T cell Lymphopenia Cases*

- Rac 2 mutation
- ADA SCID
- T-B-NK+ SCID
- T-B+NK+ (3)
- RAG 1 SCID

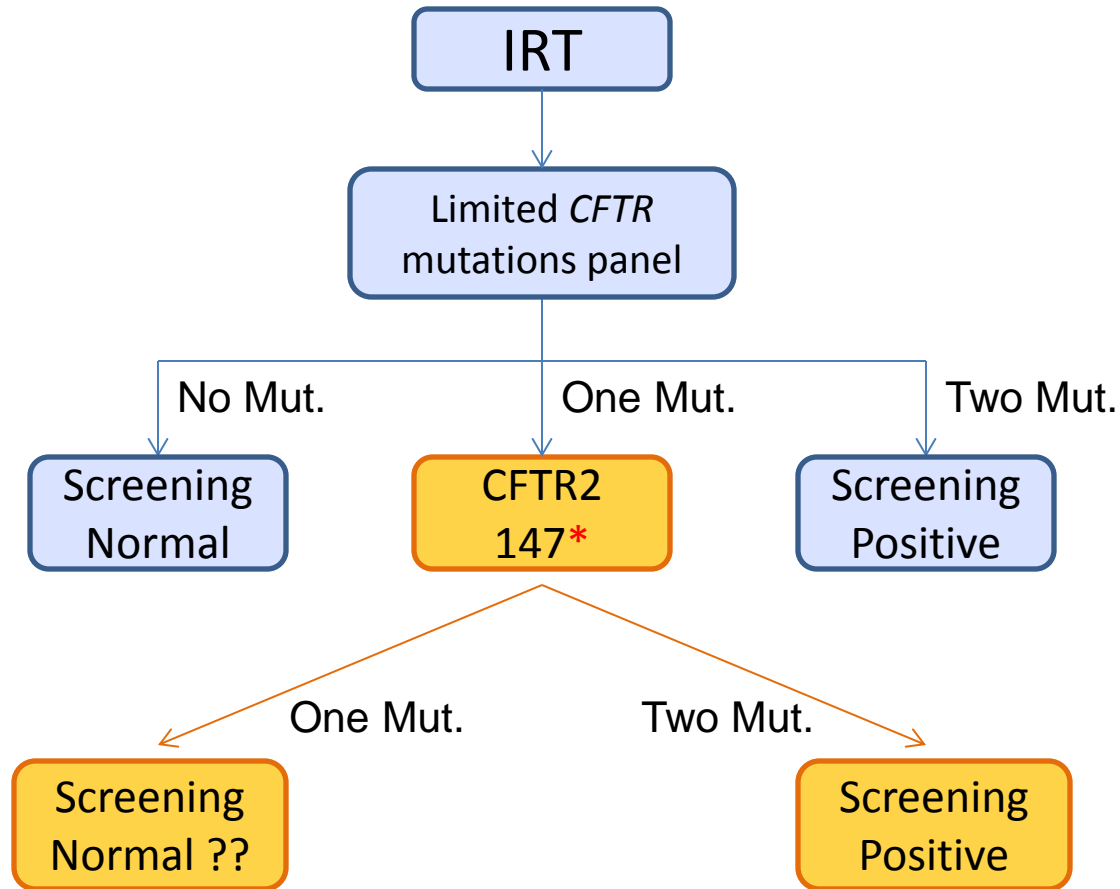
# Wisconsin Experience

*(January 1, 2008- December 31, 2012)*

## Other T cell Lymphopenia Cases

- Chromosomal abnormalities
  - 22q11.2 deletion (11)
  - Trisomy 21
- Syndromes with T cell impairment
  - Jacobsen syndrome
  - Tar syndrome
  - Ectrodactyly ectodermic dysplasia
  - Ataxia Telangiectasia
- Idiopathic T-cell lymphopenia

# Improving NBS for CF to Reduce Screening false positives using Next Generation sequencing Technology



\*Disease-causing mutations and mutations with varying consequences. (Sosnay et al, *Nature Genetics*, 2013)



# Specific Aims

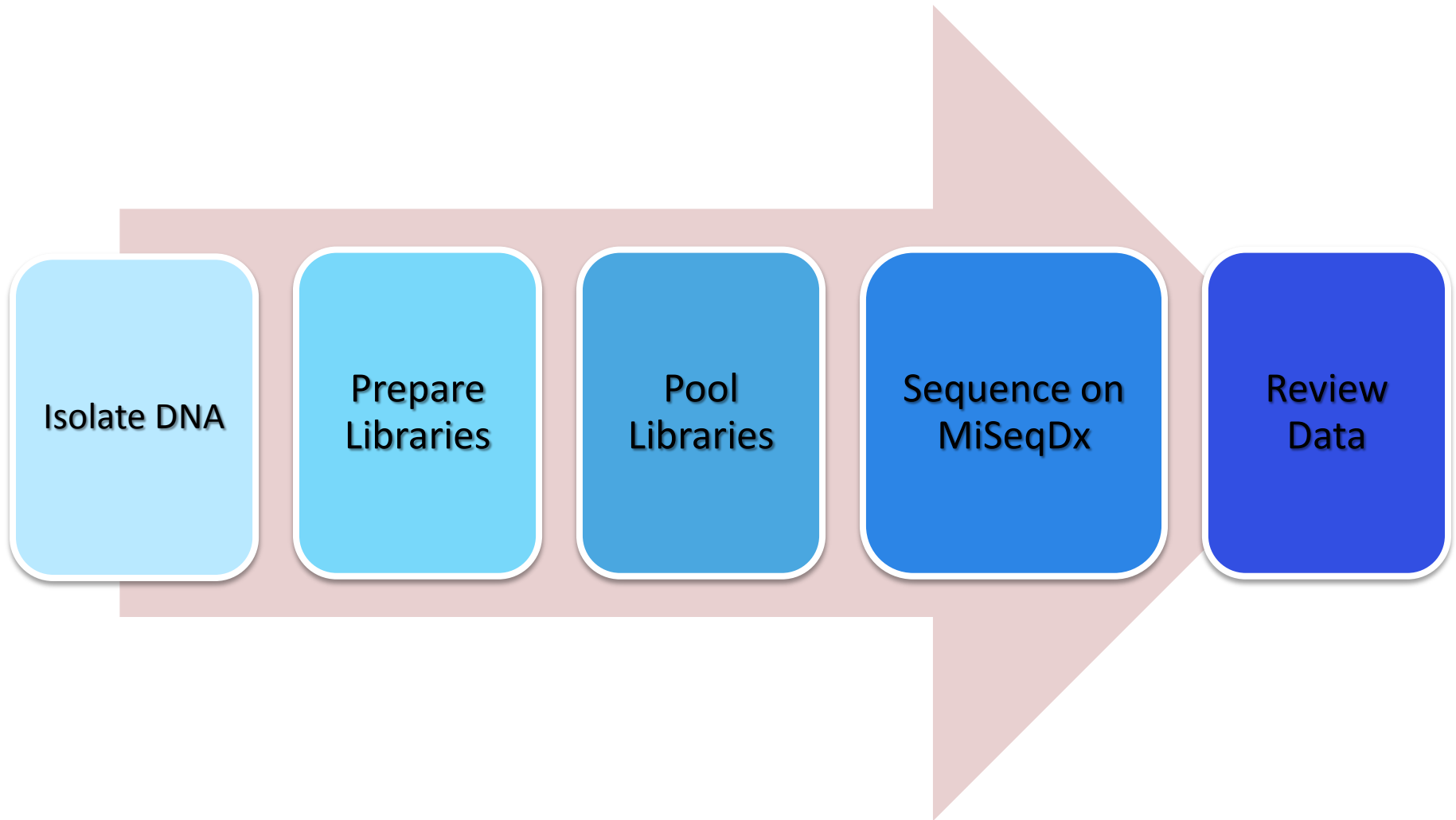
1. Establish a method of simultaneously detecting 162 *CFTR* mutations/gene variants *using dried blood spot routine newborn screening specimens to create IRT/DNA/DNA CF screening opportunity.*
2. Demonstrate that the three-tier IRT/DNA/DNA CF screening protocol would significantly reduce false positive screening results caused by identification of CF heterozygote carrier infants.
3. Demonstrate that it is cost effective to implement the three-tier IRT/DNA/DNA CF screening protocol into routine NBS for CF.

# MiSeqDx Cystic Fibrosis System

- 162 *CFTR* mutations/variants (IUO version\*)
  - 127 single nucleotide mutations/variants
  - 32 insertion/deletion mutations
  - 2 large deletions
  - PolyTG/PolyT region

\*Product is currently under FDA review.

# Cystic Fibrosis System Workflow

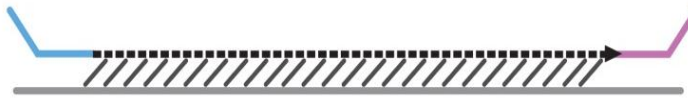


# Sequencing Library Generation

CF variant-specific probes hybridize to flanking regions of interest in unfragmented gDNA



Extension/Ligation between CF variant-specific probes across regions of interest



PCR adds indices and sequencing primers



Uniquely tagged library ready for cluster generation and sequencing



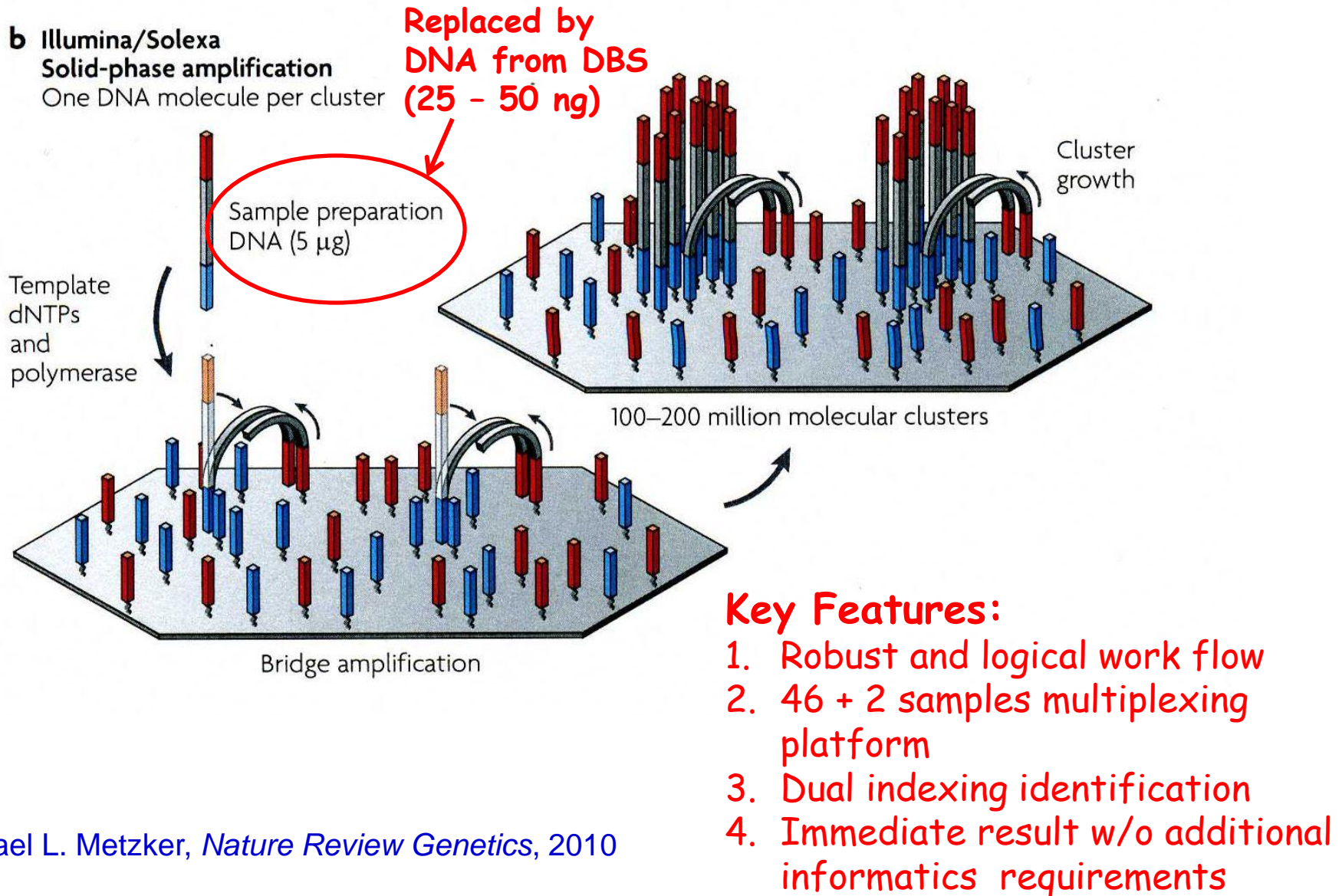
Hybridization

Extension/Ligation

PCR Primers  
Anneal

PCR amplification

# Genotyping-by-sequencing





# Funding Support for SCID



Wisconsin Newborn Screening Laboratory

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Children's Specialty Group™

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