

Establishing Quality Assurance Protocols for Low Frequency Events in a High Throughput SCID NBS Laboratory

2011 Newborn Screening and Genetics Testing Symposium San Diego, CA

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202,298 infants screened for SCID

4 SCID

~1:50,000

Surveillance: No known false negative

Through guthrie date 10/25/2011

Assay and Laboratory Development



Implementing SCID NBS with Multiplexed Assays in an Integrated Program Approach CDC National Center for Environmental Health Grant # IV01-EH000362-03

New England Newborn Screening Program Staff

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GRANT COMPONENTS

- ASSAY DEVELOPMENT
- SCREENING IMPLEMENTATION

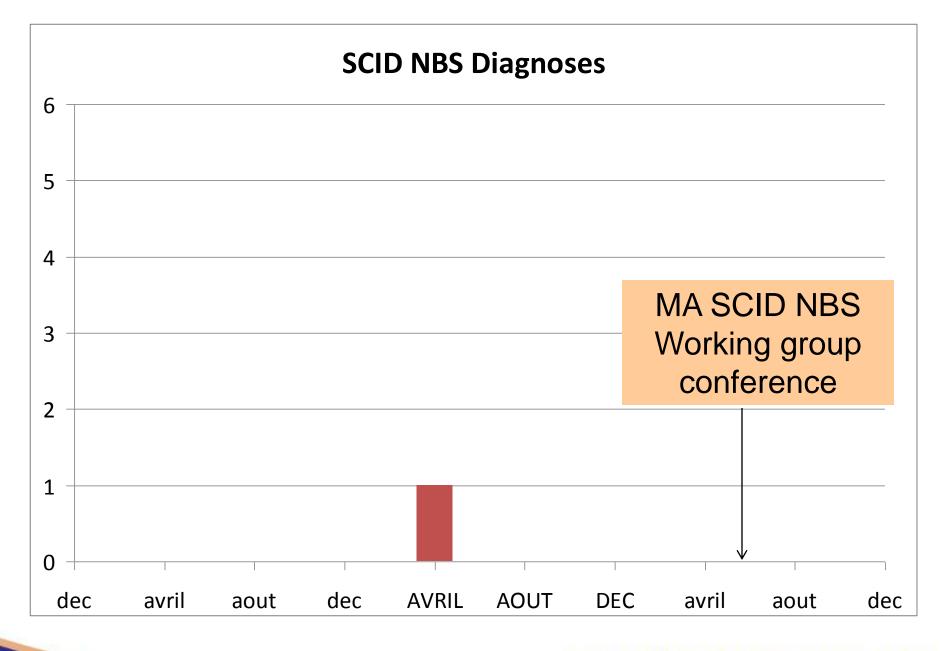
• ALGORITHM REFINEMENT

- TECHNOLOGY TRANSFER
- PUBLICATIONS

US PILOT STUDIES INTERIM FINDINGS

Combined data from CDC and NICHD-funded pilots (through April 2011)

State	WI	MA	NY	CA	PR	LA	TOT
screened	243,707	161,707	136,635	358,000	29,000	31,464	960,513
SCID	4	1	4	5	0	0	14
rate (1/x)	61,000	161,707	34,000	72,000			69,000



Are we generating false negative results?

If so, what could be the cause?

Review data:

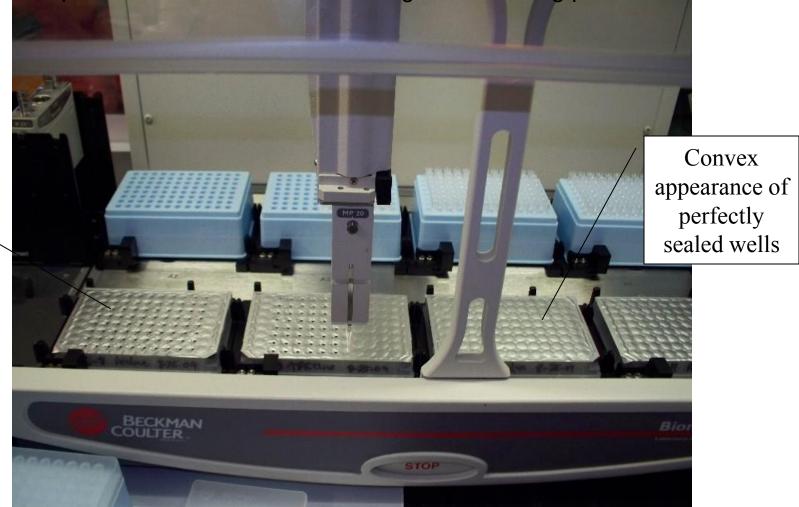
What is the potential for

Contamination?

Imprecision of the assay?

96-Well DNA Sample Plates on Biomek Deck

Individual wells remain hermetically sealed until punctured with the pipette tip for transfer to the 384-well reagent-containing plate.



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Pierced wells

Laboratory Algorithm

Components that might fail a run if copies detected

- No template controls, DNA from "no TREC" samples
- "blank spots" that are prepped with clinical specimens

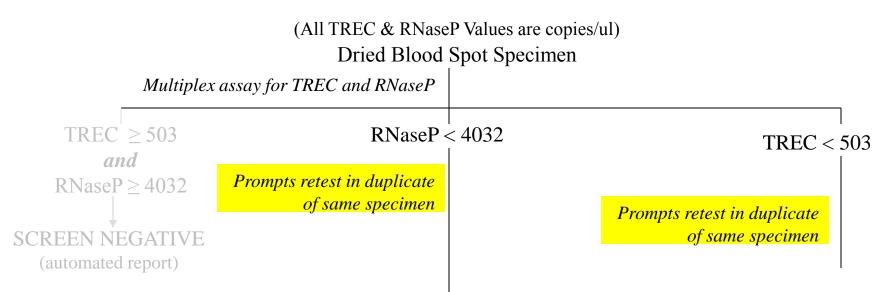
Laboratory Algorithm

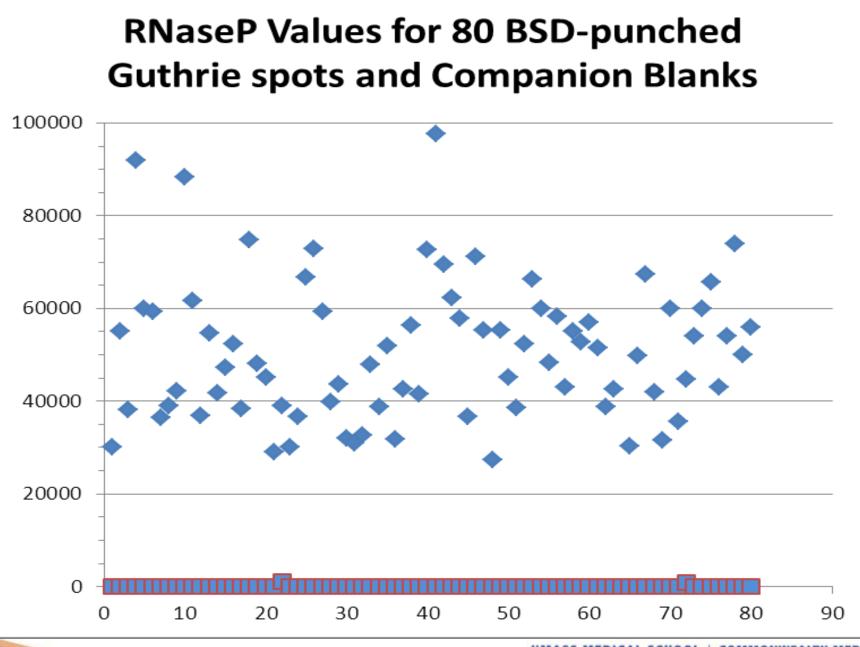
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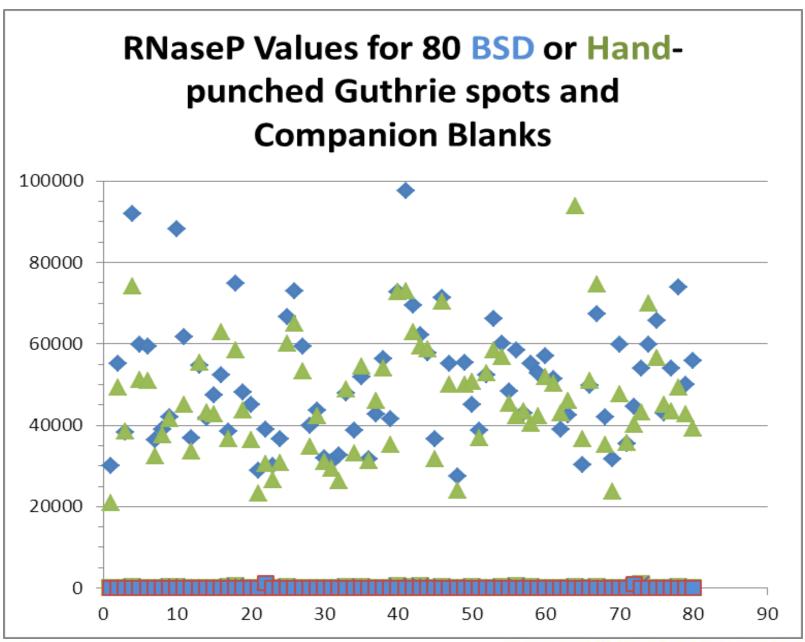
Other "blank spots"

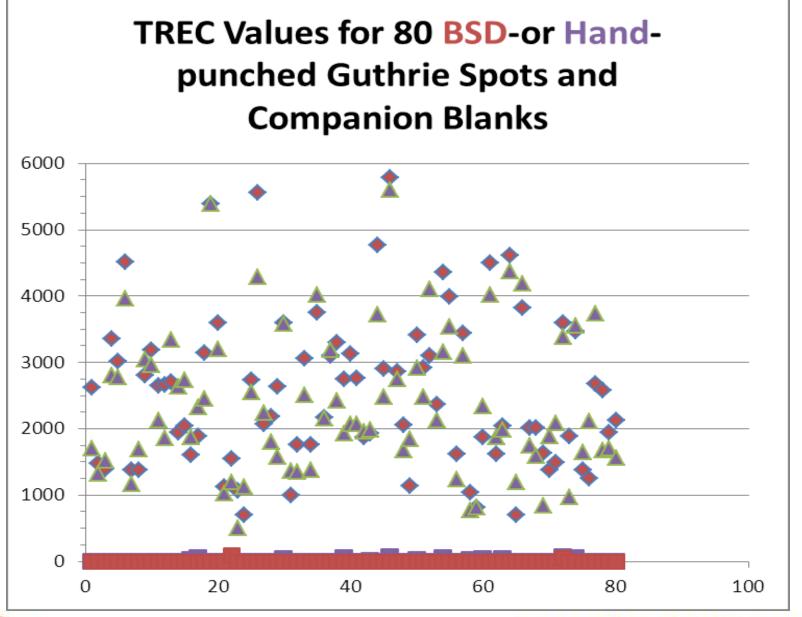
SCID NBS Algorithm



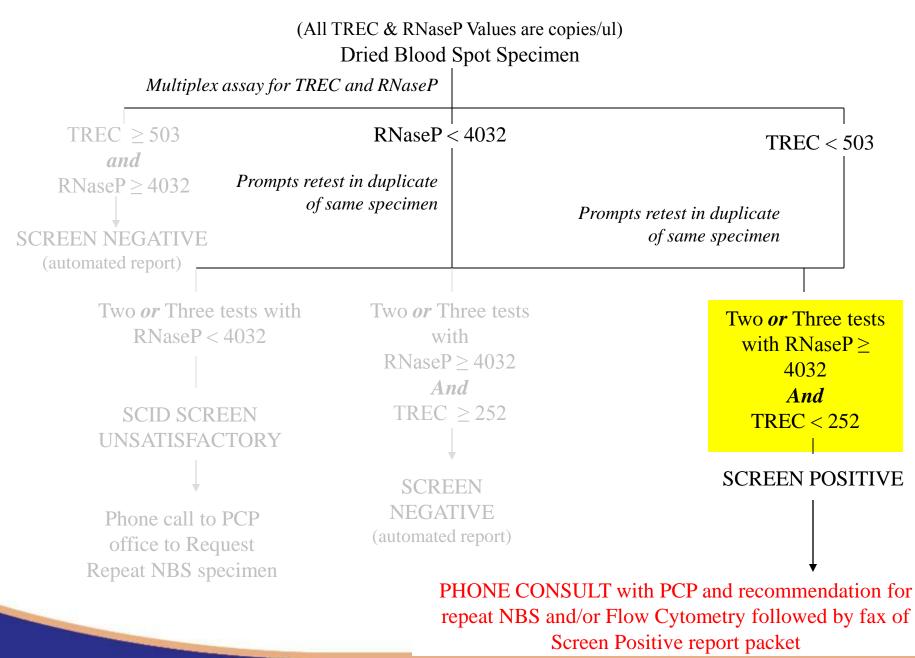


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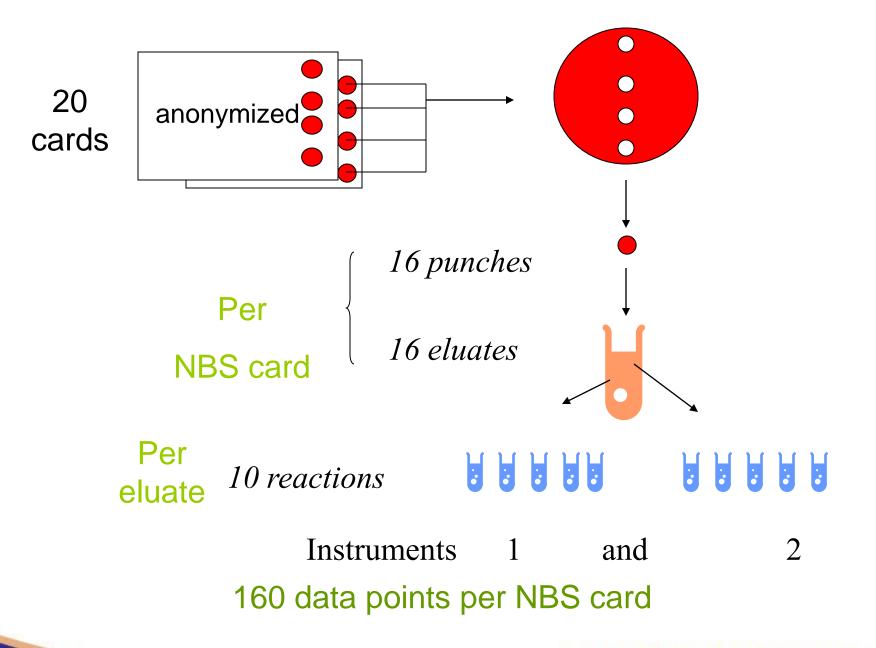
SCID NBS Algorithm



Precision of the Assay

Considerations

- intrinsic to the assay
 - within eluate
 - Between instruments, calibration curves
- intrinsic to the sample
 - Across a spot; DNA preparation



Precision of the Assay – eluates

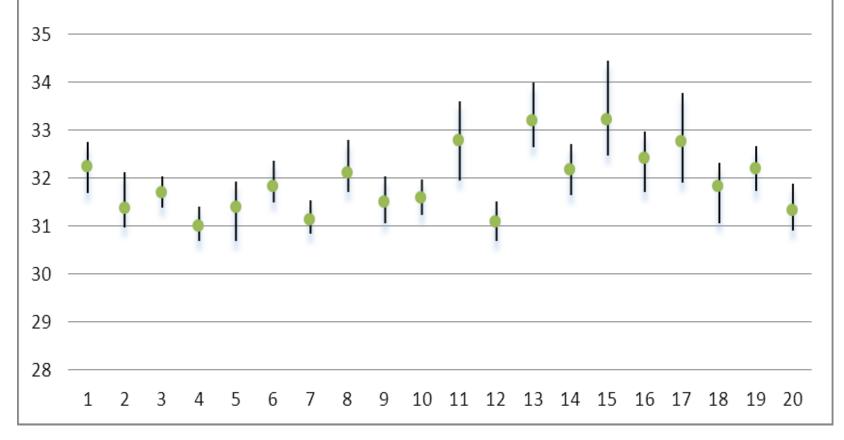
(sample from 320 eluates)

	Cq values	Cq values	
	31.51	31.27	
	31.14	31.11	
471 copies /ul	31.13	31.24	463 copies/ul
	30.88	30.96	
	31.07	30.82	

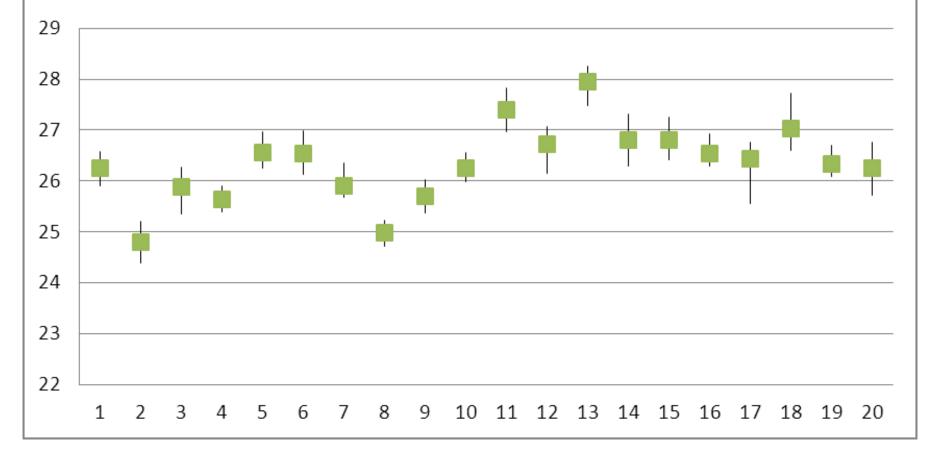
Mean SD all eluates 0.21

Precision of the Assay – across a spot

TREC Cq Values from Sampling across 20 Guthrie Spots Min, Max, Mean for 16 Punches



RNase P Cq Values from Sampling across 20 Guthrie Spots Min, Max, Mean for 16 Punches



Thoughts

Confirms High Quality Robust Assay

It's always a good thing to check...for things that might be reasonably preventable.

Punching considerations Use of replicate data within algorithm

