

# Newborn Screening for Cystic Fibrosis

## Three States' Experience with IRT/IRT/DNA

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# Colorado's CF Newborn Screening History with IRT/IRT

- Historic false negative rate (*non Meconium Ileus*)
  - $18/327 = 5.5\%$  (3.5 – 8.5, 95% CI) – *Sontag et al J Peds 2005*
  - Approximately 1 infant every 1-2 years
  - Suggestions other infants may have been missed
- How can we improve the sensitivity of the CF Newborn Screen?
  - Can we improve the sensitivity without increasing the burden of the screen?

# Our Goals for a new screen in Colorado

- Minimize false negatives
- Reduce the number of false positives
- Provide a more specific diagnosis, ie DNA
- Minimize the need for genetic counseling for detection of carriers
- Reduce parental stress
  - Reduce the time to a diagnosis
  - Reduce the number of children/parents recalled for testing
- Reduce costs of screening and follow-up

# IRT/IRT<sub>1↑</sub>/DNA in Colorado

- Decrease 1<sup>st</sup> screen cutoff
  - 105ng/ml (99.7 %ile) to 97<sup>th</sup> %ile (60ng/mL)
- **Link** 1<sup>st</sup> and 2<sup>nd</sup> screen specimens for each baby
  - SpecimenGate
- Test 2<sup>nd</sup> screen **ONLY** if first screen > 60ng/mL
- Mutation analysis if **BOTH** first and second screen results > 60ng/mL

# Implementation of IRT/IRT/DNA

- Currently being used in 3 states:
  - Colorado (June 2008) – Appx 70,000 births/yr
  - Utah (January 2009) – Appx 60,000 births/yr
  - Texas (January 2010) – Appx 400,000 births/yr

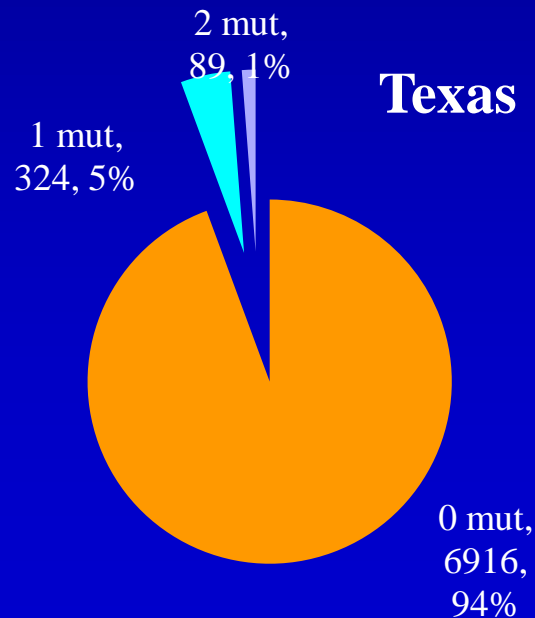
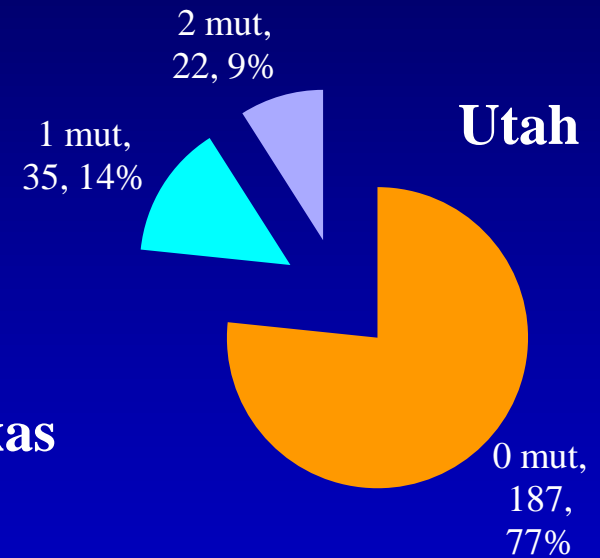
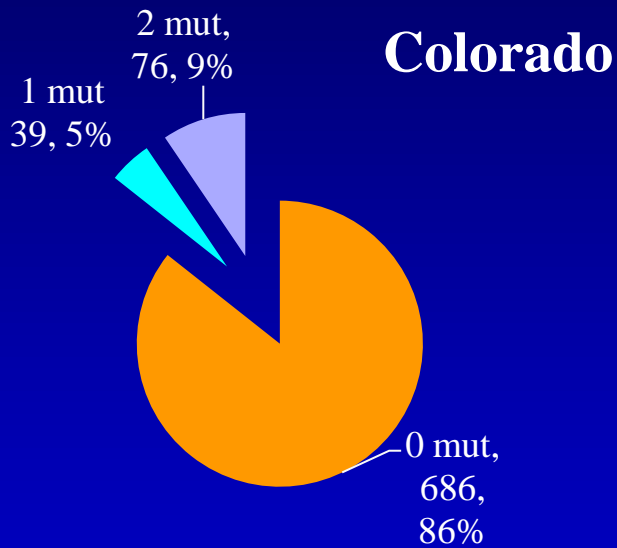
# Basic Algorithms – similarities and differences

	Colorado	Texas	Utah
Cutoffs (1 <sup>st</sup> /2 <sup>nd</sup> )	60/60ng/mL	60/60ng/mL	60/60ng/mL
Other cutoffs		46.5ng/mL – over 30 days	>97%ile prior to 6/1/10
Testing all 2 <sup>nd</sup> Screens	NO	YES	NO
Number of mutations	32	40	32
Ultra-high cutoffs	>150 ng/mL	>150 ng/mL	NONE
Dates reported	7/1/08 – 8/1/11	12/1/09 – 8/31/11	1/1/09 – 3/31/11

# Results of IRT/IRT/DNA algorithm

State	Colorado	Texas	Utah
Total Screened	213,770	675,882	120,385
Positive on first screen	4191 (2.0%)	15,154 (2.3%)	2362 (2.0%)
Persistently elevated IRT	318 (0.15%)	1,555 (0.23%)	230 (0.19%)
Number of mutation analyses	801 (0.37%)	7,329 (1.1%)	244 (0.21%)

# Results of mutation analyses performed

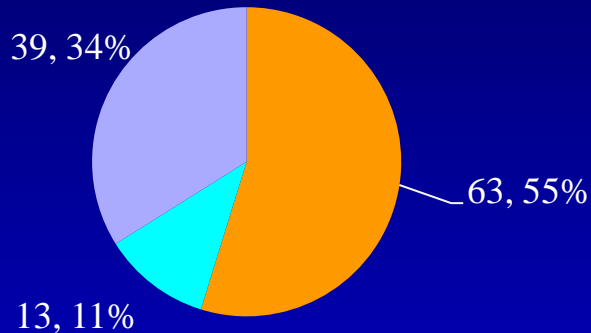




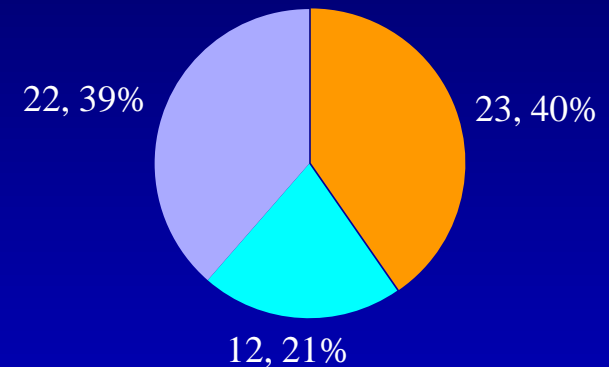
# Of those with at least one mutation....

*PPV of  $\geq 1$  CFTR mutation*  
*CF: Carrier Ratio*

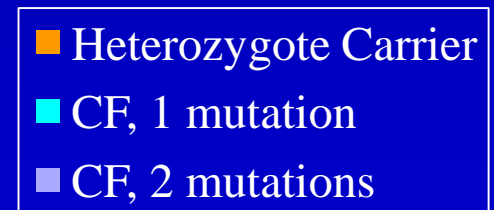
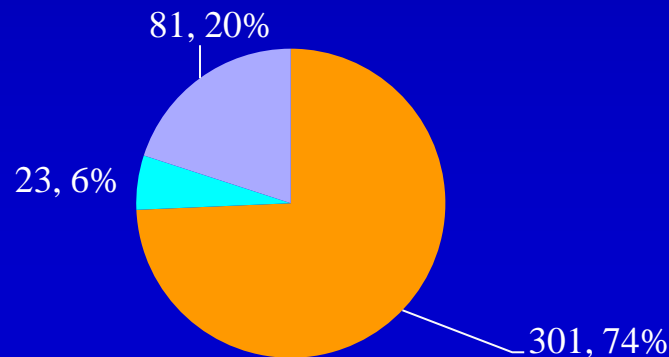
**Colorado**  
**PPV = 45%**  
**1: 1.2**



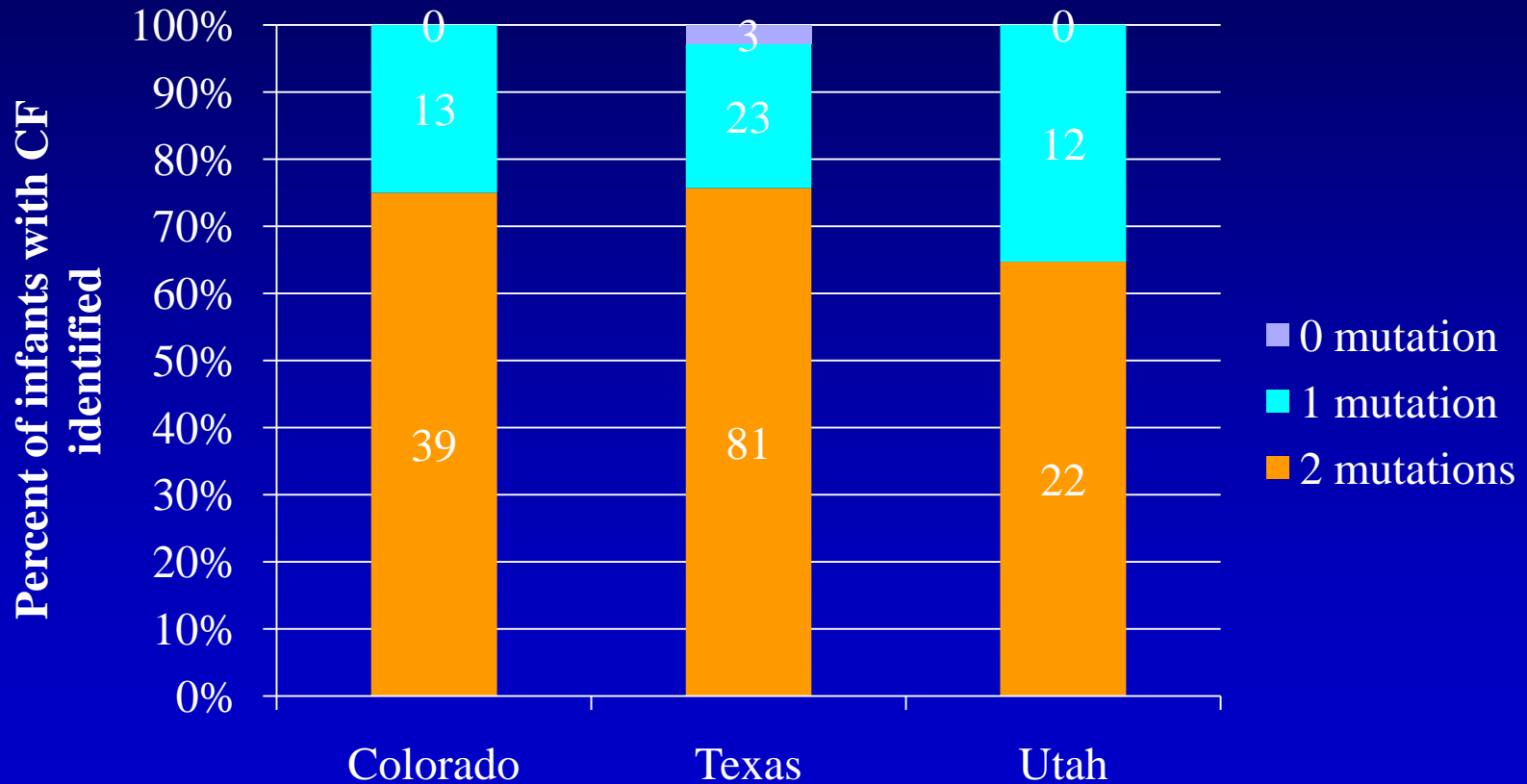
**Utah**  
**PPV = 59.6%**  
**1:0.67**



**Texas**  
**PPV = 25.7%**  
**1:2.9**



# Over 60% of CF Cases identified had 2 mutations on the panels



# Missed Cases

State	Colorado	Texas	Utah
Missed on screen	2	6	1
Missed by IRT	2 (52, 55ng/mL)	4 (19, 42, 43 and 48 ng/mL)	N/A
Missed by DNA	0	1	1 (Q943X)
False Negative Rate	4.2% (1.3-14.0)	4.7% (2.1-10.5)	2.9% (0.7-14.5)

# The new algorithm in Colorado is accurately identifying more babies with CF

- 5/52(9.6%) Babies with IRTs <105 ng/mL and >60 ng/mL have been identified cases that would not have been identified by an IRT/IRT algorithm
- Two other babies were missed by Colorado's program: - hypothetical total missed case to date (had cutoff not changed): 7/52 (13.5%, 95% CI: 6.7 – 25.3%)

# 80% of cases in the new cutoff window of IRT/IRT/DNA (60-105ng/mL) in Colorado were pancreatic insufficient

	First IRT (ng/mL)	Second IRT (ng/mL)	Genotype	Pancreatic Status
Would have been missed with 105 ng/mL cutoff	98	87	R347P/UNK	Sufficient
	98	99	F508/G542X	Insufficient
	68	127	F508/F508	Insufficient
	78	79	F508/F508	Insufficient
	76		663delT/G551D	Insufficient
Missed with 60 ng/mL cutoff	55		N1303K/2789+5G->A	Sufficient
	50		F508/R117H	Sufficient

# Two IRTs (Always vs. Selective)

- Colorado and Utah test IRTs on second specimens if the first specimen is above threshold
- Texas tests IRTs on ALL first and second specimens
  - Has identified 2/107 (1.8%) babies with CF on second screen that had normal first IRTs (<60ng/mL)
- Balancing cost of additional tests with sensitivity of tests

# Ultra-high algorithm

- Recalling infants with persistently *extremely* elevated IRTs (>99.9%ile) and no mutations on panel
- Texas and Colorado
  - 150ng/mL w/no mutations=> Sweat test
  - Texas has identified 2/107 babies (1.9%) with ultra-high algorithm
- Especially useful if mutation panel may miss some race/ethnicity groups

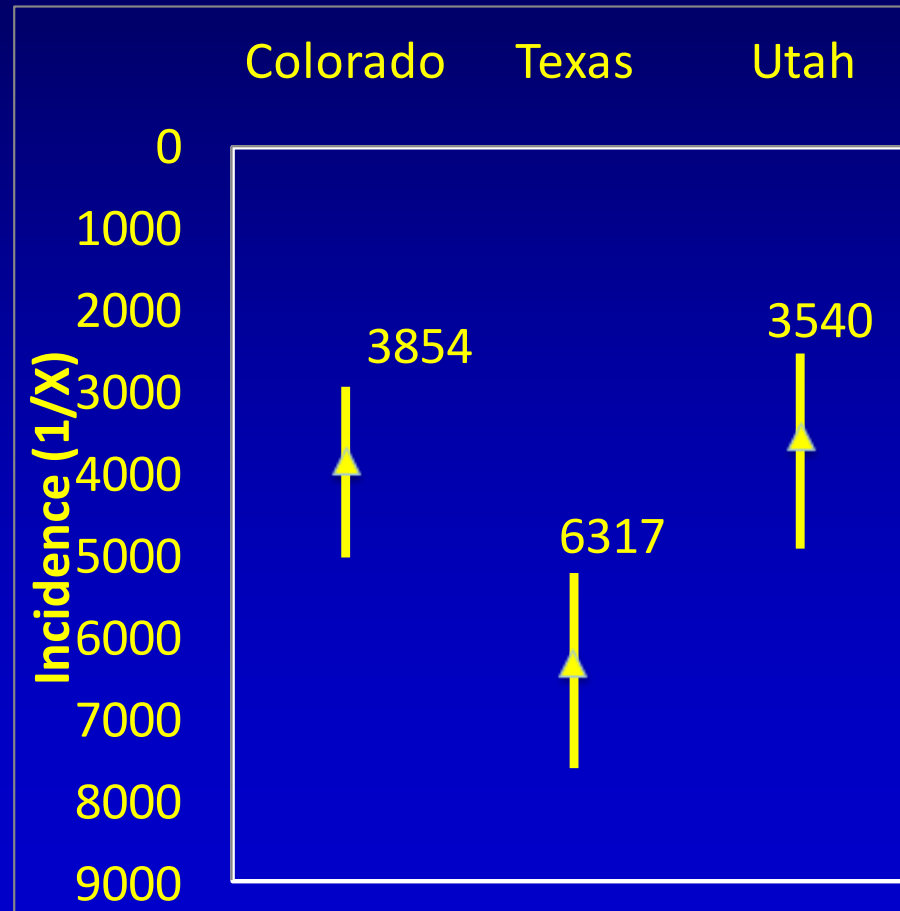
# Unlinked samples

- If first sample has elevated IRT AND second sample is not received, or not 'linkable' the 1<sup>st</sup> sample can be tested for CFTR mutations
- Colorado identified 2/52 (3.8%) cases by testing DNA on first sample.

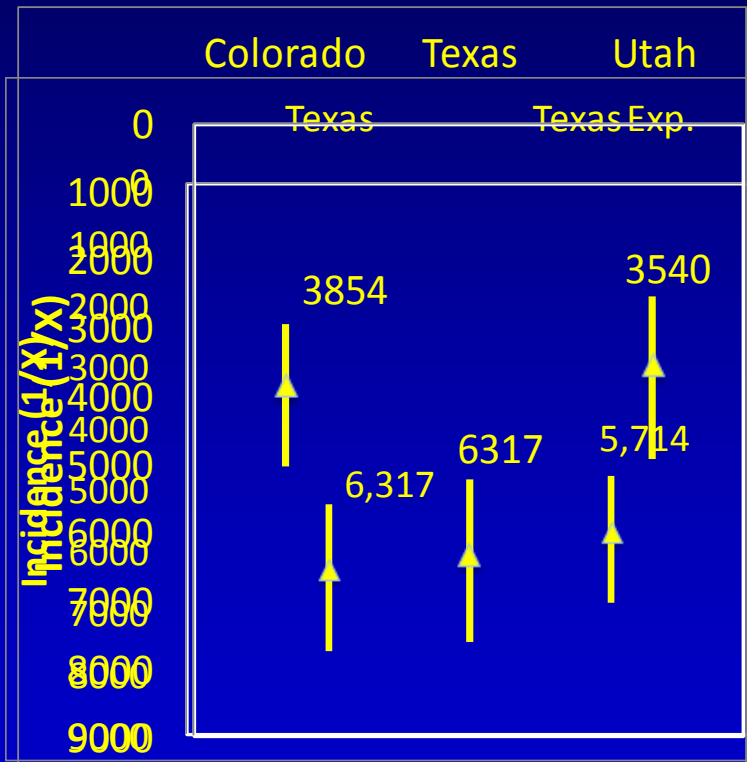


# Incidence of CF

*Newborn screening across 3 states*

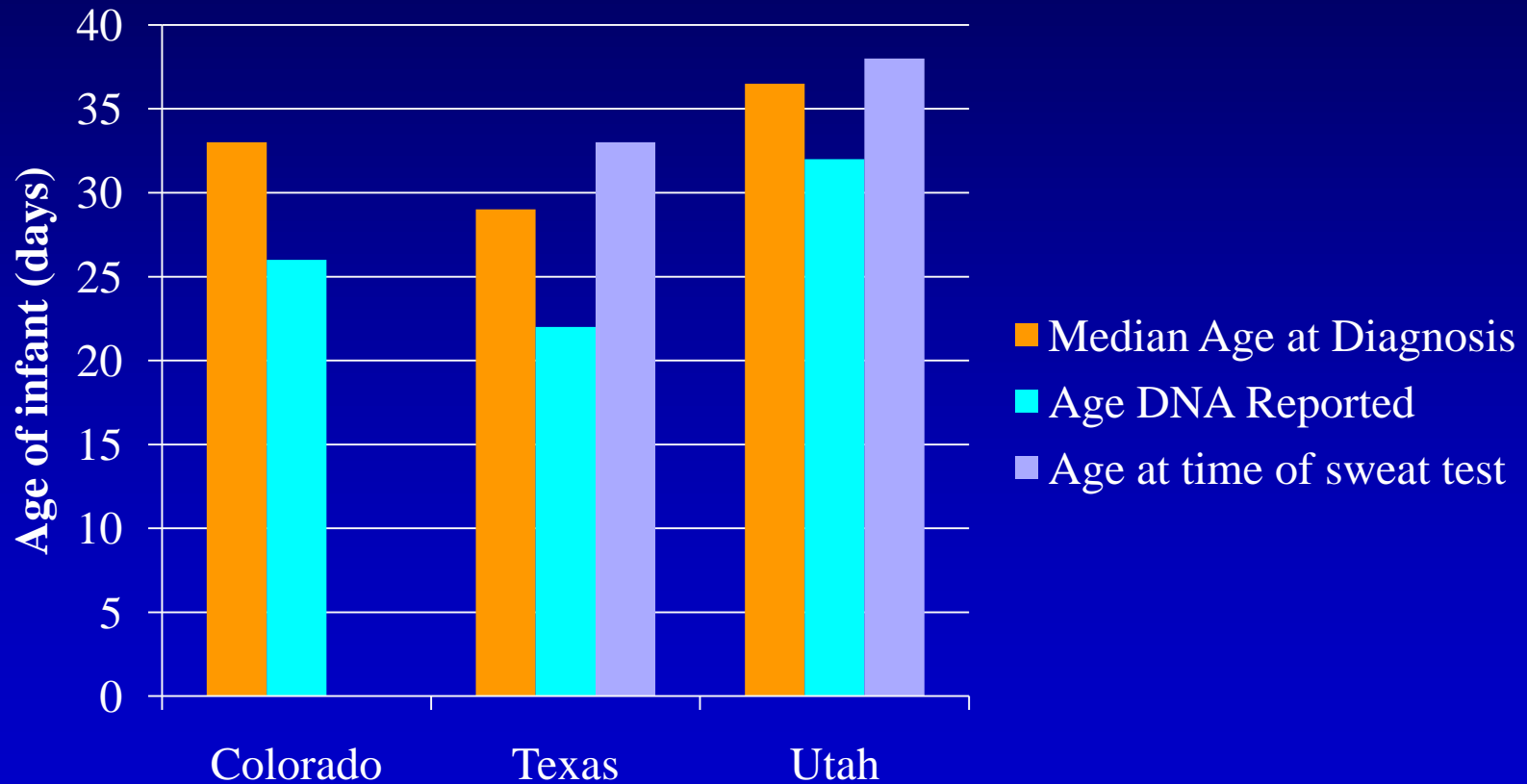


# What would the *Expected* number of children with CF be in Texas?



- Applying published disease frequencies:
  - Caucasians: 1/3,600
  - Hispanics: 1/6,500
  - African American: 1/29,000
  - Native American: 1/2,700
  - Colorado disease frequencies from: Sontag et al *J Pediatr* 2005;147
- Texas Vital Statistics (2008)
  - 34% Caucasian
  - 50% Hispanic
  - 11% African American
  - 5% Other
- **675, 882 births => Expected 118 babies**

# Age of infant at time of diagnosis, sweat test



Median age of diagnosis in Colorado has changed from 35 days (2008) to 31 days (2009) to 28 days (2010)

# Challenges to IRT/IRT/DNA analysis

## Specific to IRT/IRT/DNA

- Requires 2 samples
- May increase time to diagnosis
  - There are many steps that can be taken to shorten time to diagnosis
- Linking the samples

## DNA challenges alone

- Clinicians ‘trust’ DNA
  - Need to educate clinicians that mistakes can happen in all tests
- Identification of carriers requires counseling
- May miss individuals with rare mutations

# Advantages to IRT/IRT/DNA

## Specific to IRT/IRT/DNA

- More specific test
  - Lower number of false positives identified and referred to sweat testing
- More sensitive test than IRT/IRT
  - Lower cutoffs
- May be more sensitive than IRT/DNA in some situations
  - TX – repeating IRT on infants x2 identified infants

## DNA advantages alone

- Offers a more specific result in many cases
  - >60% of CF cases identified had 2 mutations.
- Can provide additional genetic information
  - Allow genetic counseling of parents of carriers

# Conclusion

- IRT/IRT/DNA is a sensitive algorithm for the identification of CF
- Identifies fewer carriers than reported values for programs employing IRT/DNA
- May have longer time to diagnosis but can still be achieved <1 month
- Should be considered by states with 2 DBS collections

# Thank You for Sharing Data

- Texas
  - Lynette Borgfeld
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- Colorado
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- Utah
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  - Barbara Chatfield



The new face of the infant with cystic fibrosis