

Screening for three lysosomal storage diseases in a NBS laboratory, and the potential to expand to a nine-plex assay

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Potential candidates for newborn screening of Lysosomal disorders.

Disorder	Rx	Requires early detection
Gaucher	ERT/SM	
Fabry	ERT/SM	+/-
MPS-I	ERT/BMT	+
Pompe	ERT/SM	+
MPS-II	ERT	+
MPS-IVA	ERT	+
MPS-VI	ERT	+
Krabbe	BMT	
Niemann-Pick B	?	
MLD	BMT	+

Product Description

Substrates and internal standards are available without charge through the CDC

Each vial contains the optimized ratio of substrate : internal standard for 1200 tests for:

Gaucher disease

Pompe disease

Fabry disease

Niemann-Pick A/B

Krabbe disease (600 tests)

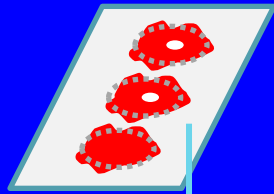
MPS-I

Each box contains 6000 tests

Reagents for each disorder are packaged separately to allow end-user to choose menu-style

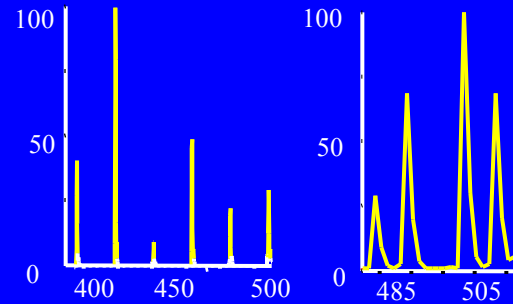
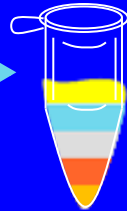


Triplex Procedure



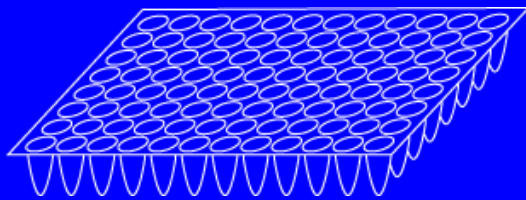
Anonymous blood spots

Fabry, Pompe, MPS-1

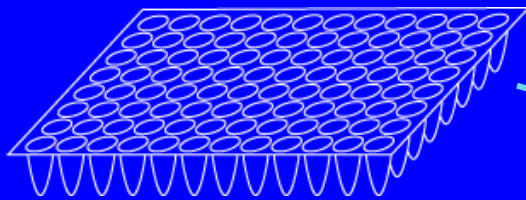


MS/MS

normal

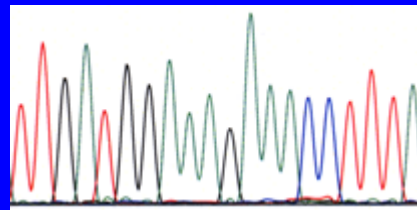


Assay plate



Duplicate plate

abnormal



unaffected

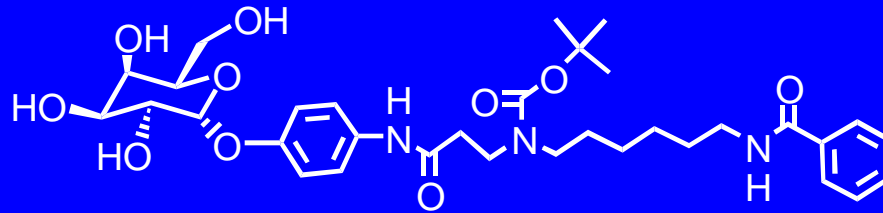
affected

DNA sequencing (GLA, GAA, IdUA) for genotype

Fabry Disease

- X-linked
- Deficiency of acid α -galactosidase
- Shortened life expectancy from:
 - Renal failure at 30-40 years
 - Hypertrophic cardiomyopathy
 - CNS strokes
- Childhood symptoms of:
 - Peripheral pain
 - Fatigue

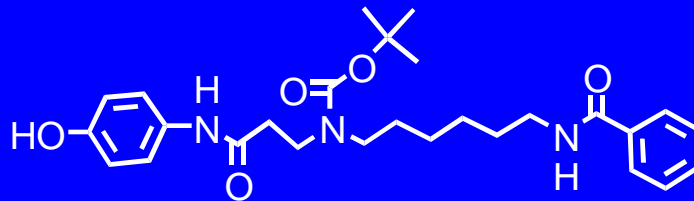
Fabry Assay



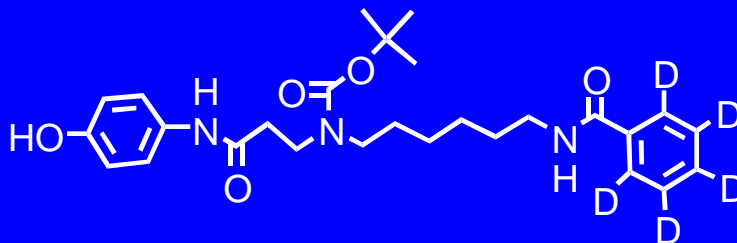
Substrate



**Acid alpha-Galactosidase (α-Gal)
(Fabry)**



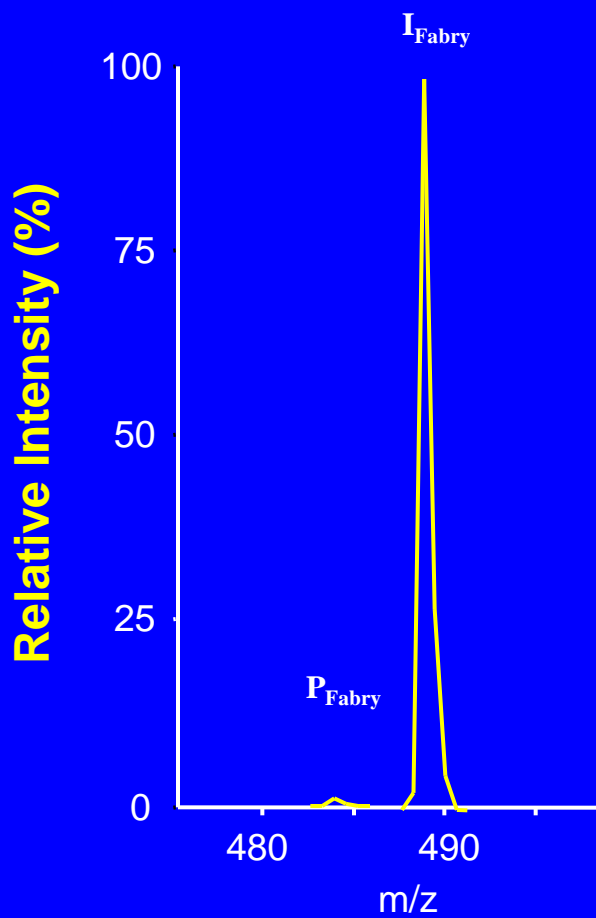
Product



Internal standard

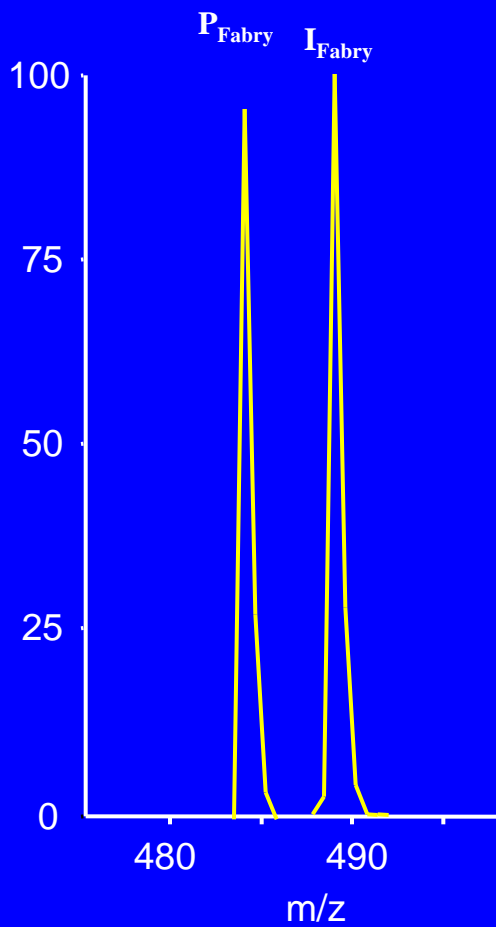
Fabry

$$P_{\text{Fabry}} / I_{\text{Fabry}} = 1.06\%$$



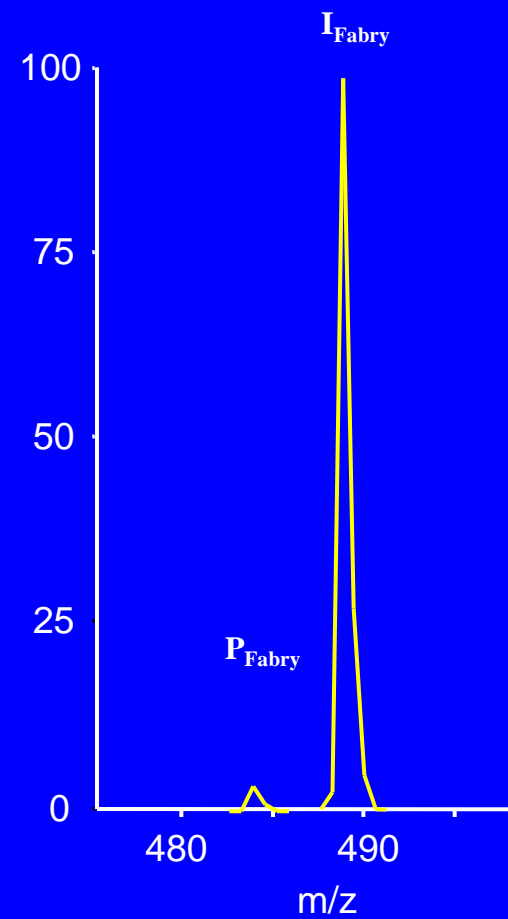
Blank

$$P_{\text{Fabry}} / I_{\text{Fabry}} = 95.7\%$$



Unaffected

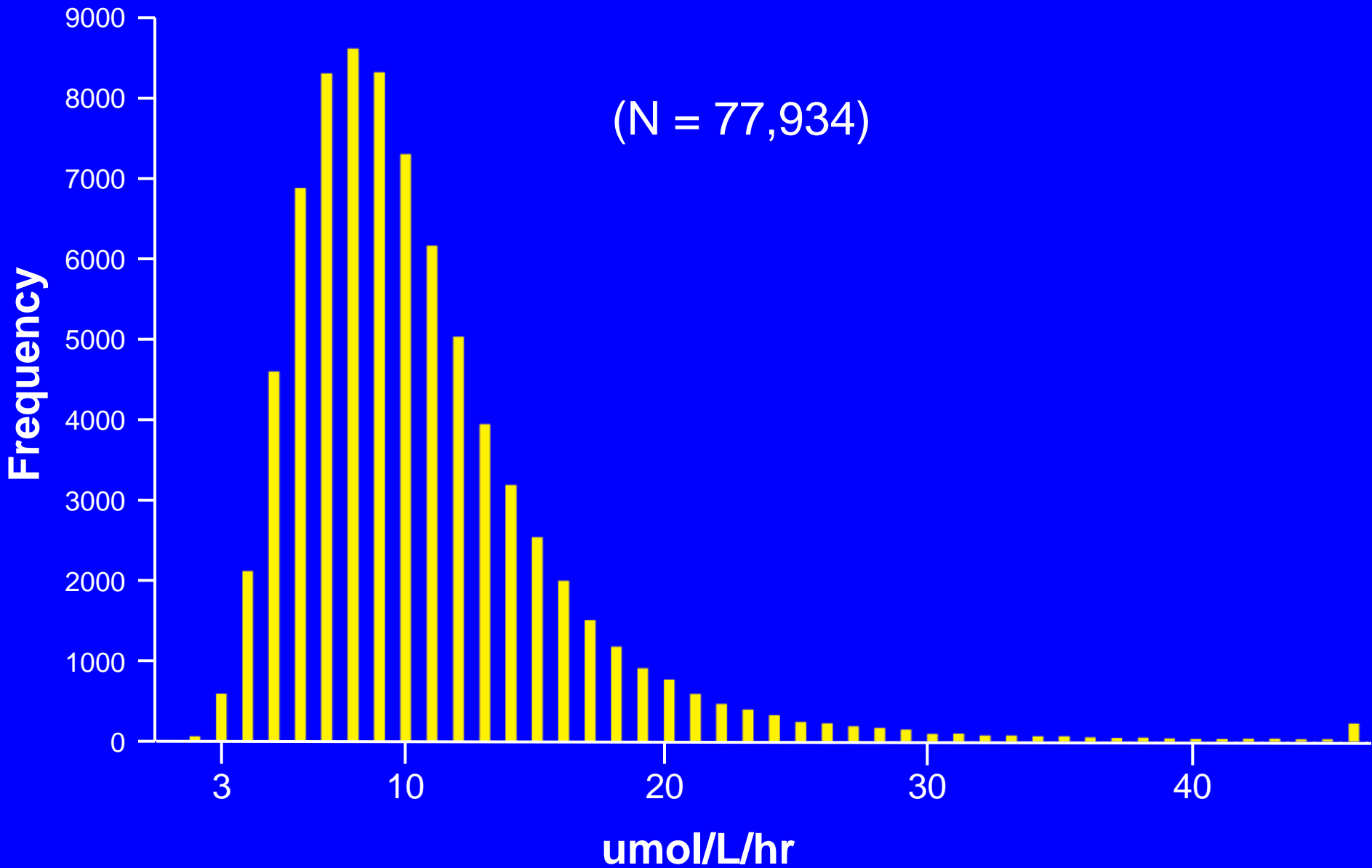
$$P_{\text{Fabry}} / I_{\text{Fabry}} = 3.5\%$$



Affected

GLA activity (Fabry)

mean = 10.7 ; 15% of mean = 1.6



Fabry Disease

Xq22

13 kb

7 exons

429 AA's



400+ pathogenic mutations

- Prevalence: 1/17,000 - 1/50,000
- Taiwan population: 1/16,000 with hypertrophic cardiomyopathy (IVS4 + 919g>a)
- Italian newborn screening: 1/3,100

Table 1: Fabry Disease

Cut off at 15% of assay mean: 1.5 $\mu\text{mole/hr/L}$	Enzyme activity ($\mu\text{mole/hr/L}$)	% of mean $\bar{X}=10.5$	Mutations
	1.72	16.1	p. Ser 334Asn/wt (F)
	1.50	14.0	wt
	1.38	12.9	wt
	1.32	12.3	p.Ala143Thr (M)
	1.12	10.5	wt
	0.89	8.3	p.Asp313Gly (M)
	0.58	5.4	p.Asn215Ser (M)
known affected (n=5)	0.13 – 0.50	1.2 – 4.7	

(M) = male
(F) = female

Positive predicted Value = 0.42
Specificity ~1.0
Sensitivity ~ ?

clinical prevalence: 1/ 40,000 males
NBS prevalence: 1/ 12,000 males

Pompe

Deficiency of lysosomal acid
 α –glucosidase

Clinical symptoms:

- Progressive muscle weakness with variable onset; 3 mo to adulthood
- Cardiac failure in infancy

In adults:

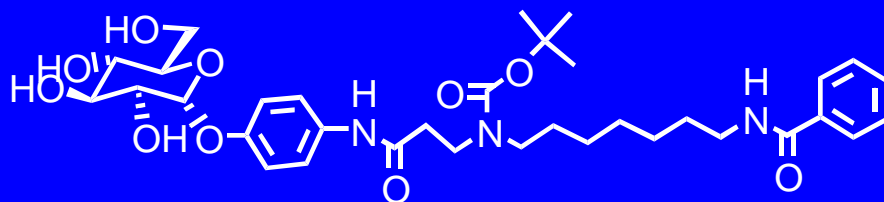
- Progressive muscle weakness
- Respiratory failure

Infantile-Onset Pompe Disease

Head Lag



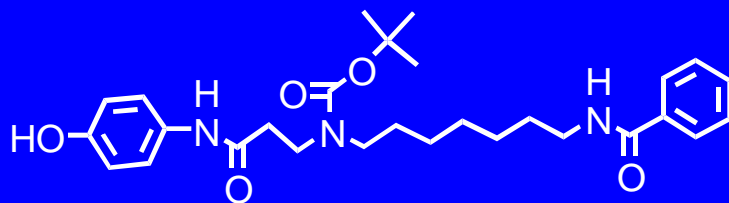
Pompe Assay



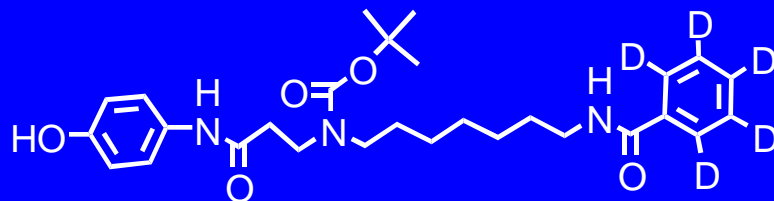
Substrate



**Acid alpha-Glucosidase (α -Glu)
(Pompe)**



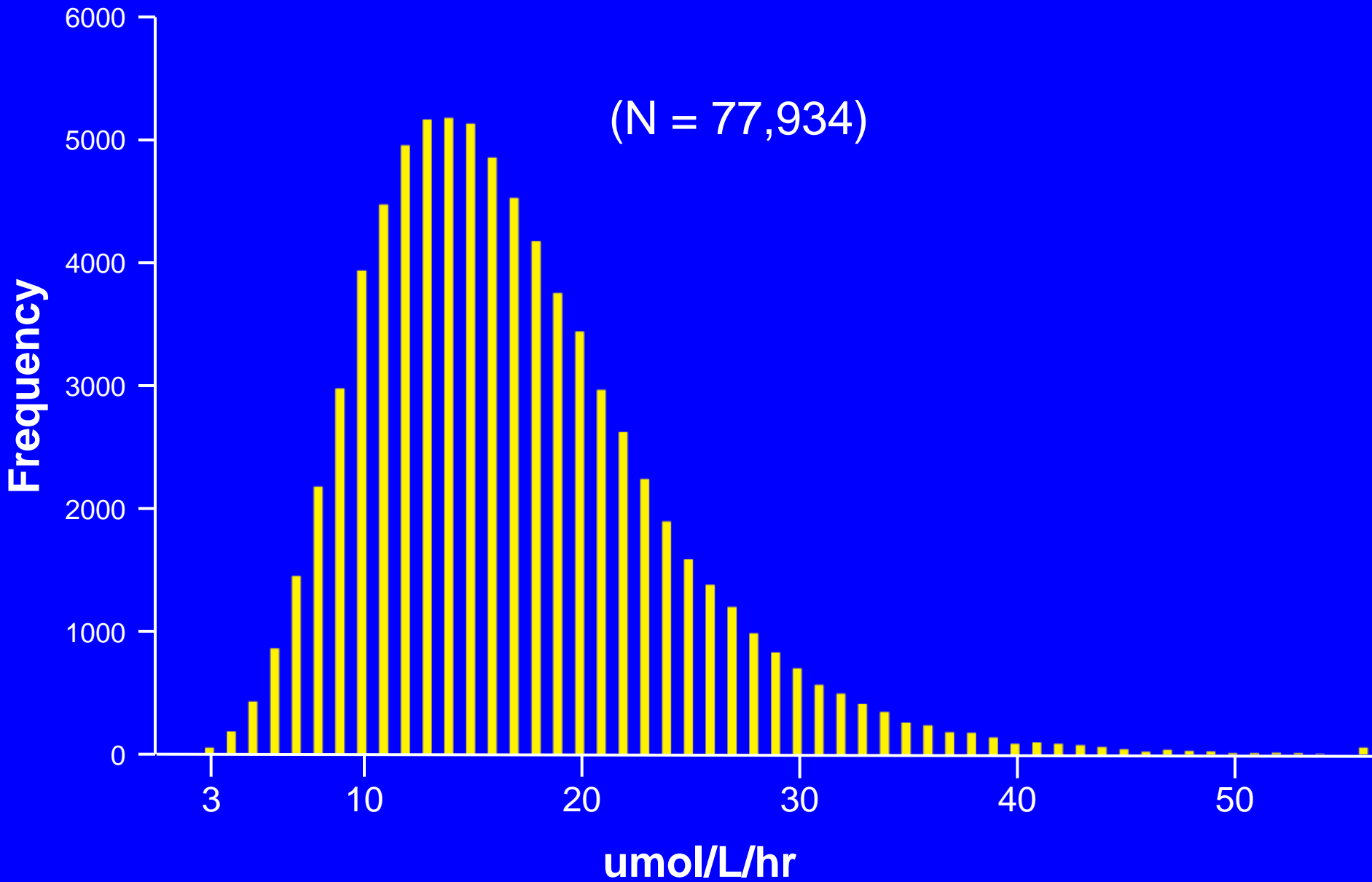
Product



Internal standard

GAA activity (Pompe)

mean = 17.3 ; 15% of mean = 2.6



Pompe Disease

17q 25.2

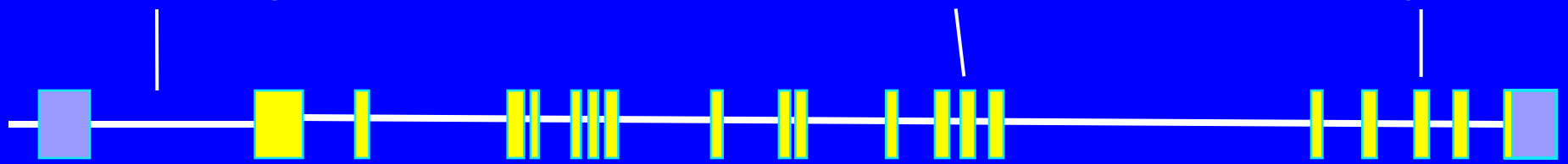
25 kb

20 exons

c.IVS1-13t>g

p.Asp645Glu

p.Arg854X



Prevalence:

Af. American:	1/14,000	(infant)
US population:	1/40,000	(infant & adult)
European:	1/100,000	(infant)
	1/60,000	(adult)

Table 3: Pompe Disease

Cut off at 15% of assay mean: <2.6 μmole/hr/L	Enzyme activity (μmole/hr/L)	% of mean - X=17.3	Mutations
	2.53	14.4	*IVS1-13t>g/IVS1-13t>g
	2.46	14.0	wt
	2.44	13.9	p.Gly576Ser;p.Glu689Lys/wt
	2.38	13.5	IVS1-13t>g/p.Glu689Lys
	2.21	12.6	IVS1-13t>g/p.Glu689Lys
	2.2	12.5	p.Gly576Ser;p.Glu689Lys/p.Glu945Lys
	1.77	10.1	p.Glu689Lys/?exon9
	1.70	9.7	p.Met122Lys / p.Val642Asp + psdef
	1.70	9.7	unable to sequence
	1.57	8.9	*IVS1-13t>g/IVS1-13t>g
	1.42	8.1	wt (?)
known affected (n=5)	0.13 – 0.50	1.2 – 4.7	

PPV=0.27

Prevalence=1/27,000

MPS-1 disease

Hurler phenotype:

early diagnosis with:

Coarse features

Dyostosis multiplex

Progressive intellectual loss

Cloudy cornea

‘attenuated’ phenotype:

Mild somatic changes

Slow neurological progression

Stiff joints

Thickened dura

Cloudy cornea

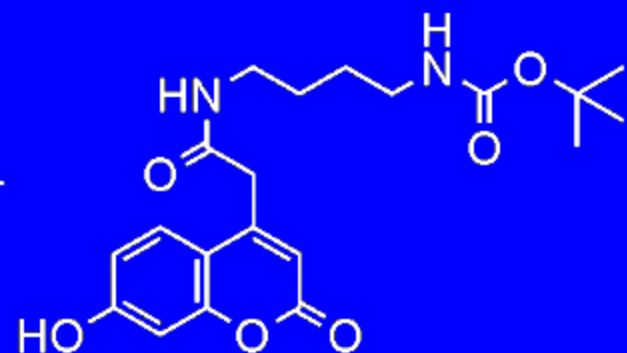


MPS-I Assay

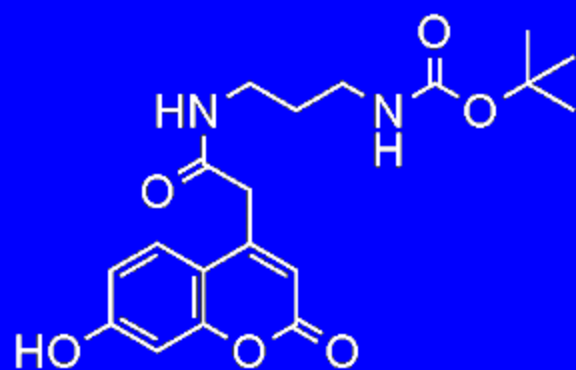


IdA-S (m/z 567.2 for M+H⁺)

IdA in DBS $\xrightarrow{\hspace{1cm}}$



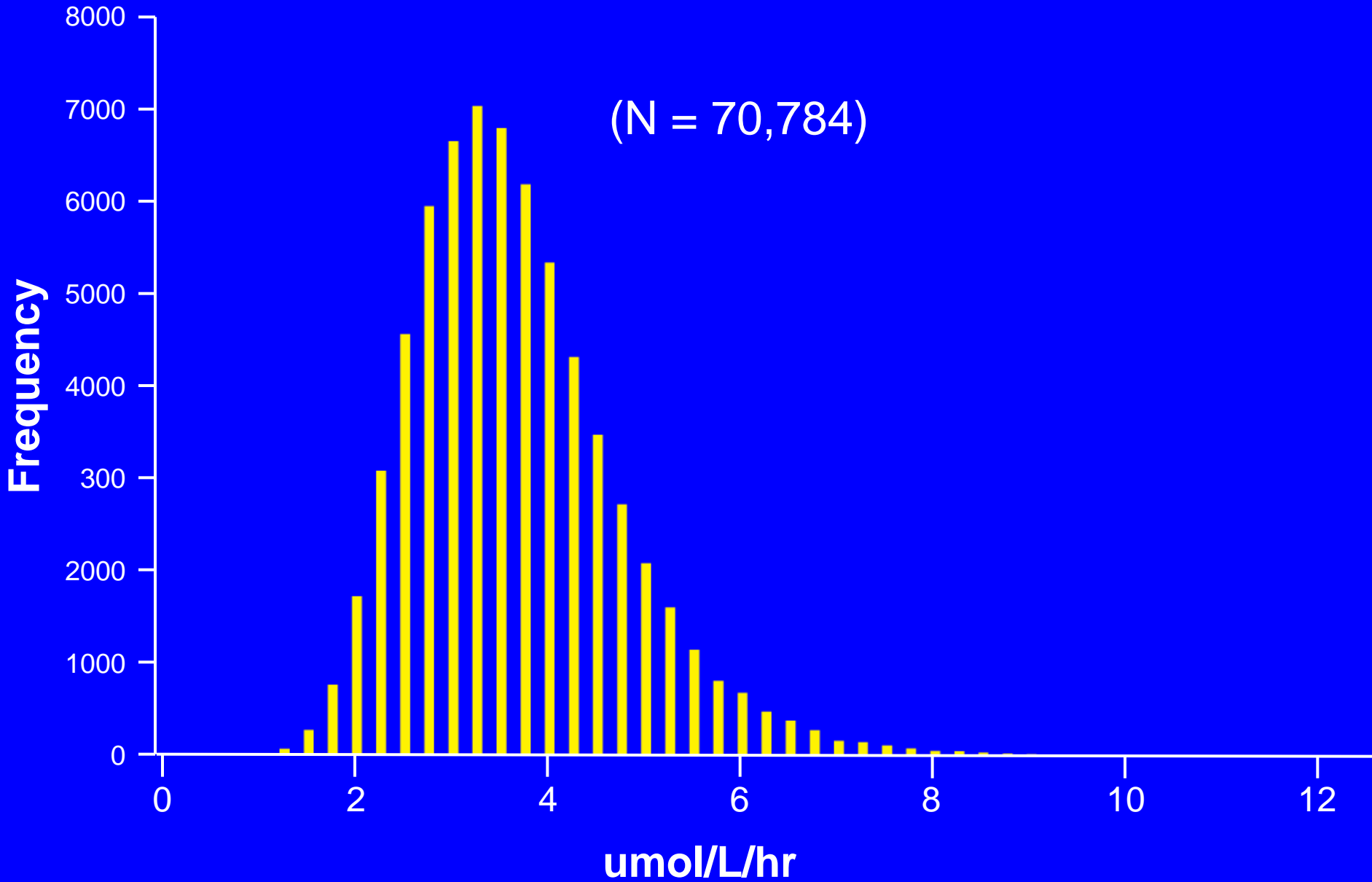
IdA-P (m/z 391.2 for M+H⁺)



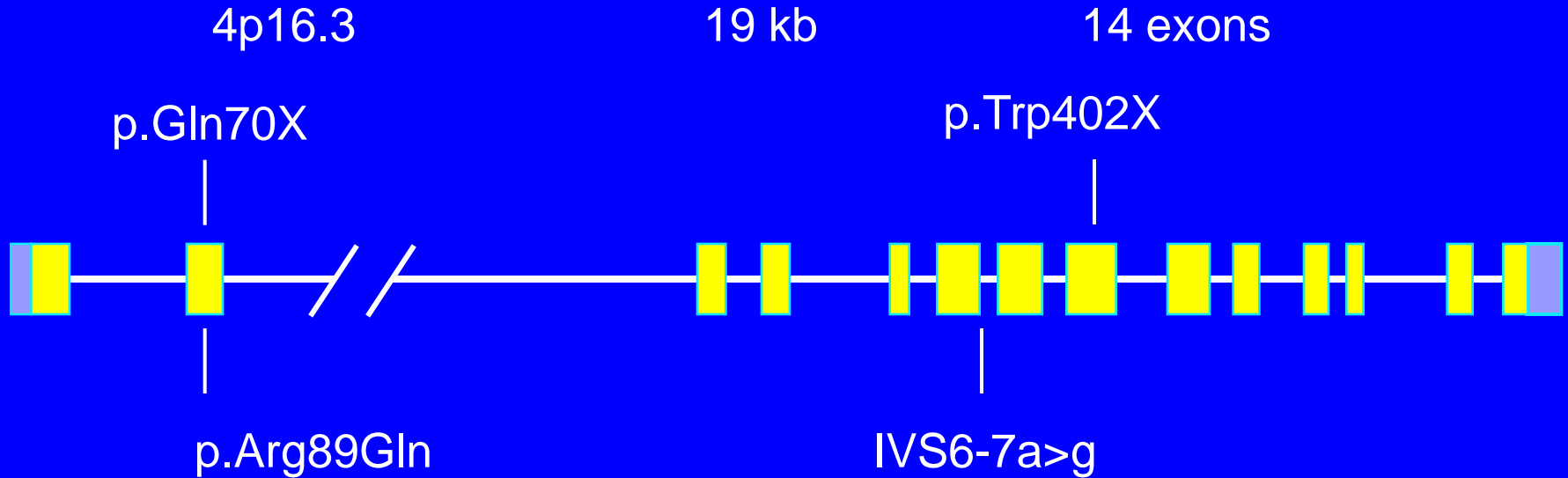
IdA-IS (m/z 377.2 for M+H⁺)

IDUA activity (MPS-I)

mean = 3.6 ; 30% of mean = 1.09



MPS-1 Disease



Clinical prevalence: ~1/100,000

Table 2: MPS-I

Cut off at 30% of assay mean: < 1.07	Enzyme activity ($\mu\text{mole/hr/L}$)	% of mean $\bar{X}=3.6$	Mutations
	1.08	29.2	wt
	1.06	28.6	p.Asp119Tyr / p.Glu84Ser
	1.05	28.4	p.Gln70X / p.Gln70X
	1.02	27.6	p.Trp402X / wt
	0.94	25.4	wt (poor punch)
known affected (n=5)	0.58 – 0.83	18.4 – 22.5	
known carriers (n=4)	1.44 – 3.02	39 – 81	

Positive predictive value = 0.4

Clinical prevalence: $\sim 1/100,000$

NBS prevalence : $\sim 1/35,000$

Summary

- MS/MS assay can easily be adapted to a NBS screening laboratory
- Multiplexing multiple enzymes simplifies the procedure and is a built-in control for sample integrity
- In the first 70,000+ samples:
 - 8 positive newborns identifies with LSD
- Overall prevalence: $\sim 1/9,600$

Potential Diseases for LSD Screening By Multiplex Analysis

Fabry

Krabbe

MPS-II

Pompe

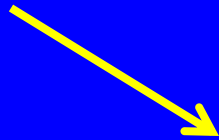
Gaucher

MPS-IVA

MPS-I

Niemann-Pick

MPS-VI



organic layer



injection into MS/MS

In queue

MPS-IVB, MLD

CNL1, CNL2

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