



In Our Own Time: Newborn Screening for Pompe Disease

Michele Caggana, Sc.D., FACMG June 22, 2017





Meyer Aviles @meyeravi · 1d Happens to the best of us



NHP Southern Command @NHPSouthernComm If the person who drove their car onto a rock leavi this morning wants their car they can come talk to Troopers to get it back!!!

♀ tì ♡ 10





If the person who drove their car onto a rock leaving **#edc** this morning wants their car they can come talk to NHP Troopers to get it back!!! Nevada Highway Patrol



Retweets Likes







Why Pompe Disease?

- Pilot study consented 3-4 NYC hospitals
- Dr. Melissa Wasserstein, PI NICHD
- MC recipient of contract from NICHD
- ACHDNC
- Universal screening in NY
- Substituted MPS1
- Added to regulation (when validated) along with ALD

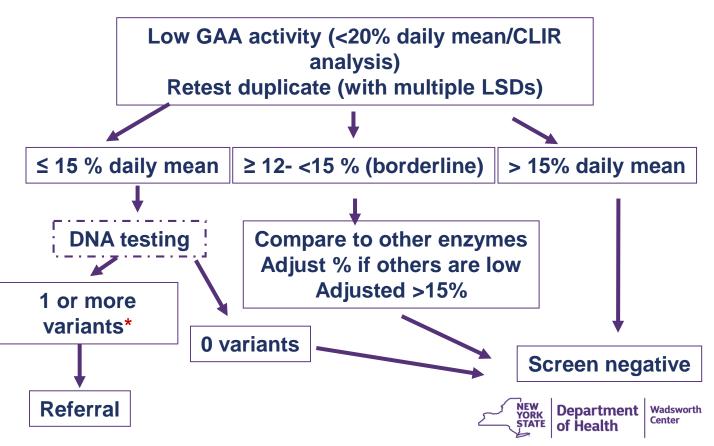


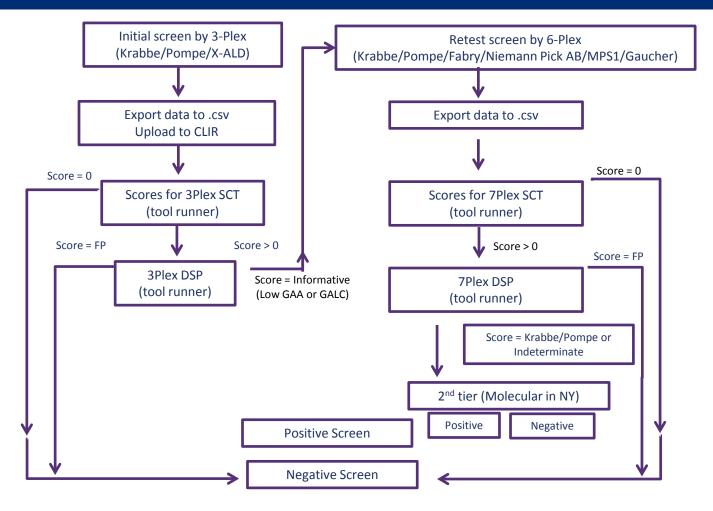
Pompe Disease

- Conservatively set cut-off at 15.0% for NIH pilot study after >5,000 tests
- MO provided screen positive specimens for interlaboratory comparison
- 12.0% was highest positive result on panel
- Much discussion on case definition for Pompe disease
- Issue with many LSDs having later onset
- Newborn screening???



Pompe Disease Screening Algorithm

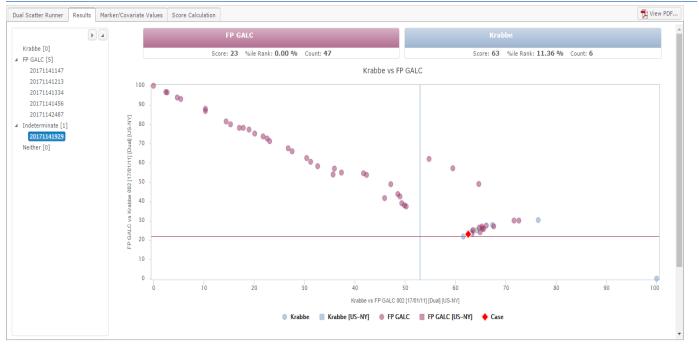




June 26, 2017



Dual Scatter Plot Runner: Krabbe vs FP GALC 002 [17/01/11] [Scatter] [US-NY]



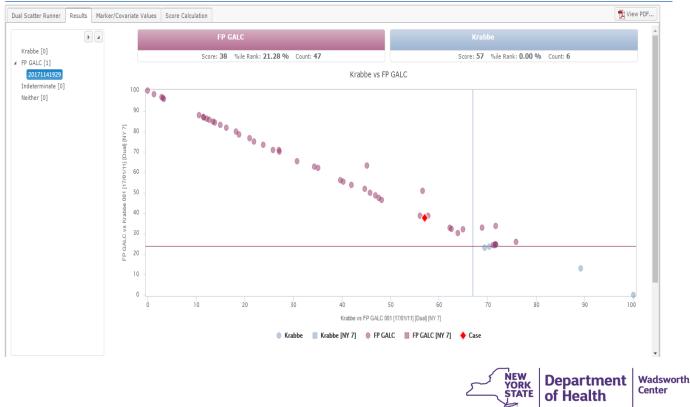
NEW YORK STATE Of Health Wadsworth Center

11

June 26, 2017

F	MAYO C	LINIC		CLIR - Collaborative Laboratory In	CLIR - Collaborative Laboratory Integrated Reports				
Applicat	on: KMPX	KMPX ►				monica.martin@health.ny.gov			
Home	Resources 🔻	Location Data 🔻	Post-Analytical Tools 🔻	Productivity Tools 🔻	Tasks 🔻		My Account *	Sign Out	

Dual Scatter Plot Runner: Krabbe vs FP GALC 001 [17/01/11] [Scatter] [NY 7]



Pompe Disease by the Numbers

Validation study 5,000 spring 2013 Pilot 30,358 May 2013 – Sept 2014 Live 701,332 since October 1, 2014 815 <20%

Samples from Missouri – all infantile <8% Of all affected specimens confirmed 12-13% Set cutoff at < 15% of mean



Pompe Disease by the Numbers

Live 701,332 October 1, 2014 131 specimens sequenced 10 common benign variants 19 pseudodeficiency / common benign

- **102** infants referred
 - 52 carriers
 - 4 infantile onset 3 classical / 1 non-classical
 - 17 2 pathogenic variants
 - 20 1 pathogenic / 1 VOUS
 - 9 2 VOUS



MPS1 Disease by the Numbers

- Added to pilot May 20, 2015 Total tested: 31,235
- 43 <20%; 31 <16%
- **10 positive infants referred for DNA**
- **5 were pseudodeficiency**
- 1 carrier
- 1 pseudodeficiency allele / VOUS
- 3 are being followed clinically/pending



Thank You !!



Joe Orsini, PhD Erin Hughes, MLS Patrick Hopkins Colleen Stevens, PhD



