

# The Infant Microbiome and Necrotizing Enterocolitis

St. Louis Infant Microbiome Team

HMP

1 September 2010

St. Louis, MO

# Baby Boy

## 26 Weeks gestation, C-section, BW= 720 g

### Antibiotics:

- DOLs 1-4: ampicillin, gentamicin (prematurity, choriamnionitis)
- DOLs 5-10: cefotaxime (CSF)
- DOLs 5-7: vancomycin, fluconazole (pathology report, failure to wean from ventilator)
- DOLs 9-18: vancomycin, cefotaxime, fluconazole (leukocytosis, failure to wean from ventilator)

### Feeds:

- DOLs 8-21: formula (24 kcal, premature formula)

### DOL 21:

- Transfusion, ventricular tap (intraventricular hemorrhage)  
Increasing ventilator settings: increased opacity; vancomycin, clindamycin

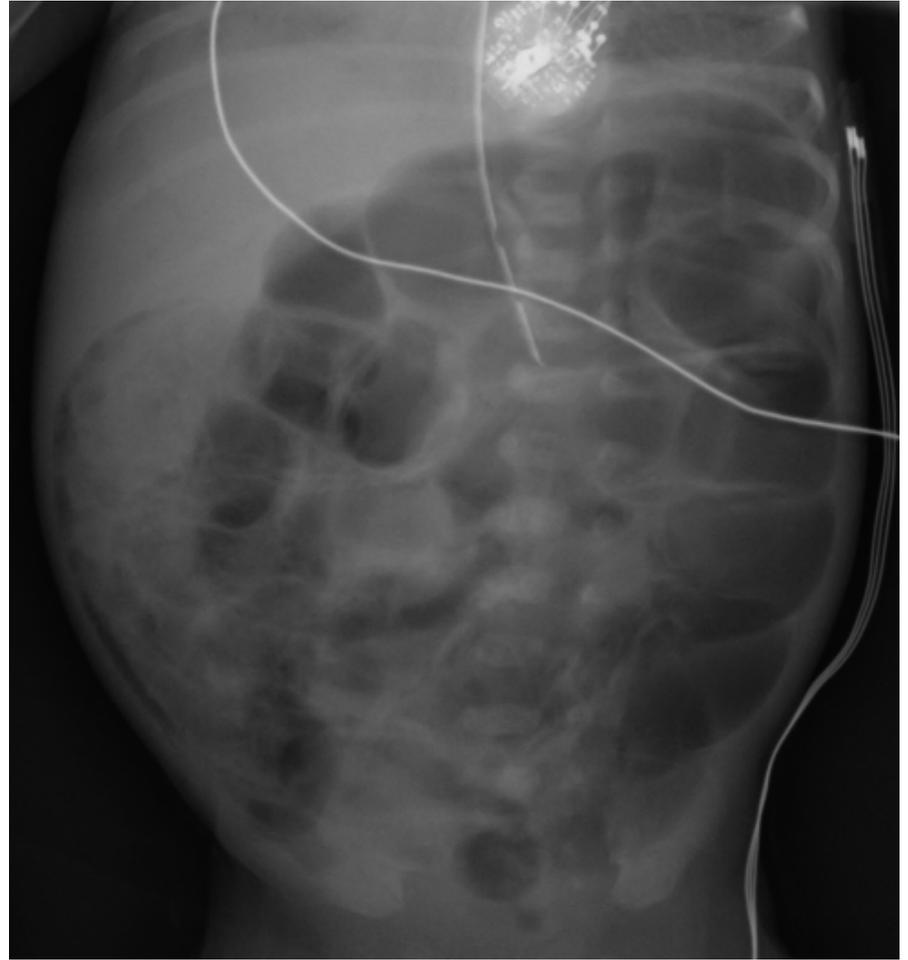
### DOL 22:

- Bloody stool

DOL 22



2:58 AM  
Gaseous distension



8:15 AM  
Extensive right flank  
pneumatosis.

# Exploratory Operation

“... panintestinal necrosis beginning just distal to the Ligament of Treitz and extending to the beginning of the cecum ...”

“... this was not compatible with life and not amenable to any surgical intervention.”

# Necrotizing Enterocolitis (NEC)

Common: 7-12% of all prematures (<1500 g BW)

Devastating: 20-30% case fatality rate

Almost all cases occur by DOL 35, patients are hospitalized, identifiable risk cohort (BW < 1500)

Occurs postnatally (i.e, after microbial colonization), probiotics might protect, antibiotics might increase risk

# The Hypothesis

- One or more identifiable components of the enteric biomass (including bacteria, their genes, transcripts, and gene products)

# Study Structure

7 day/week consent

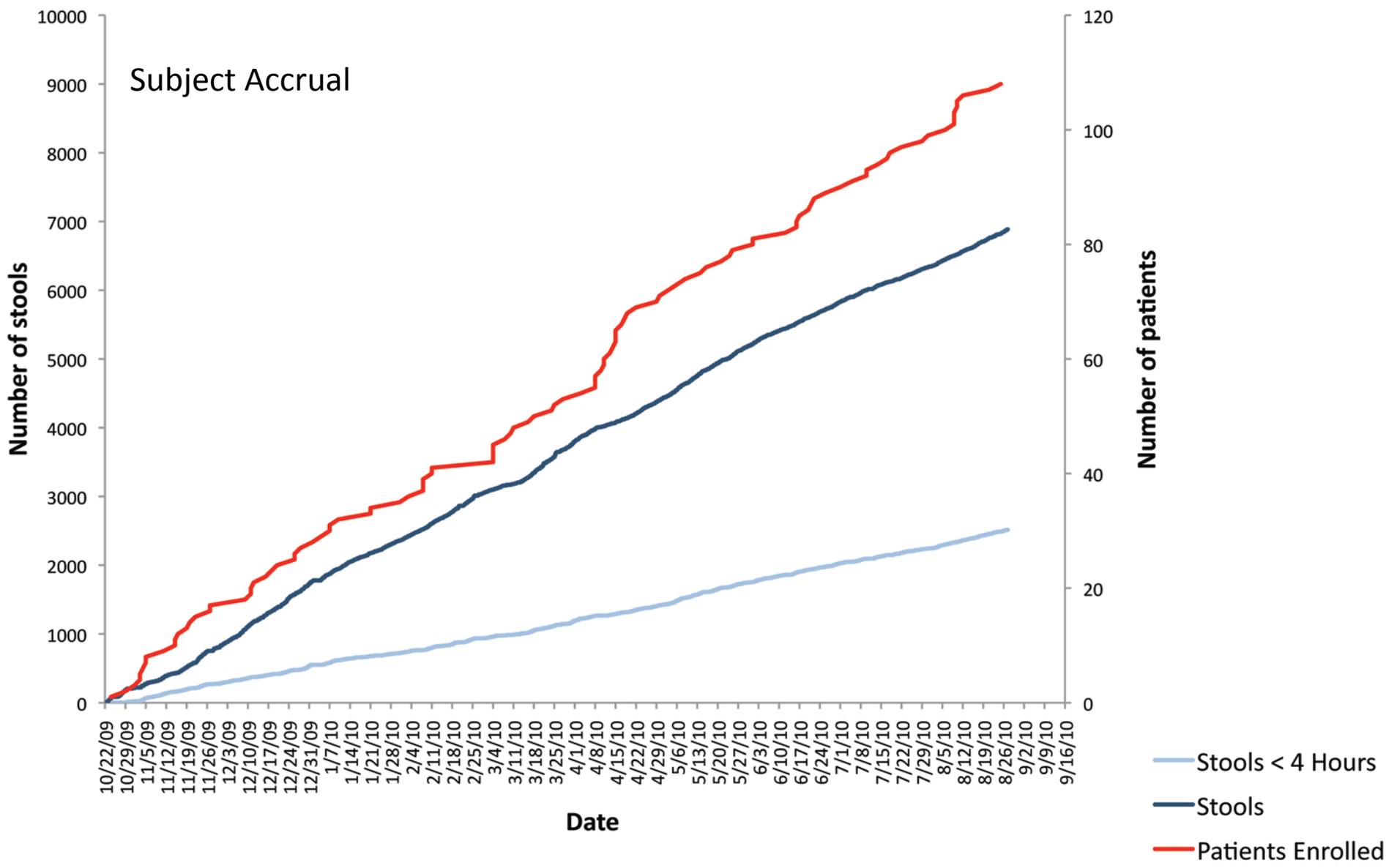
All stools sought (prioritized)

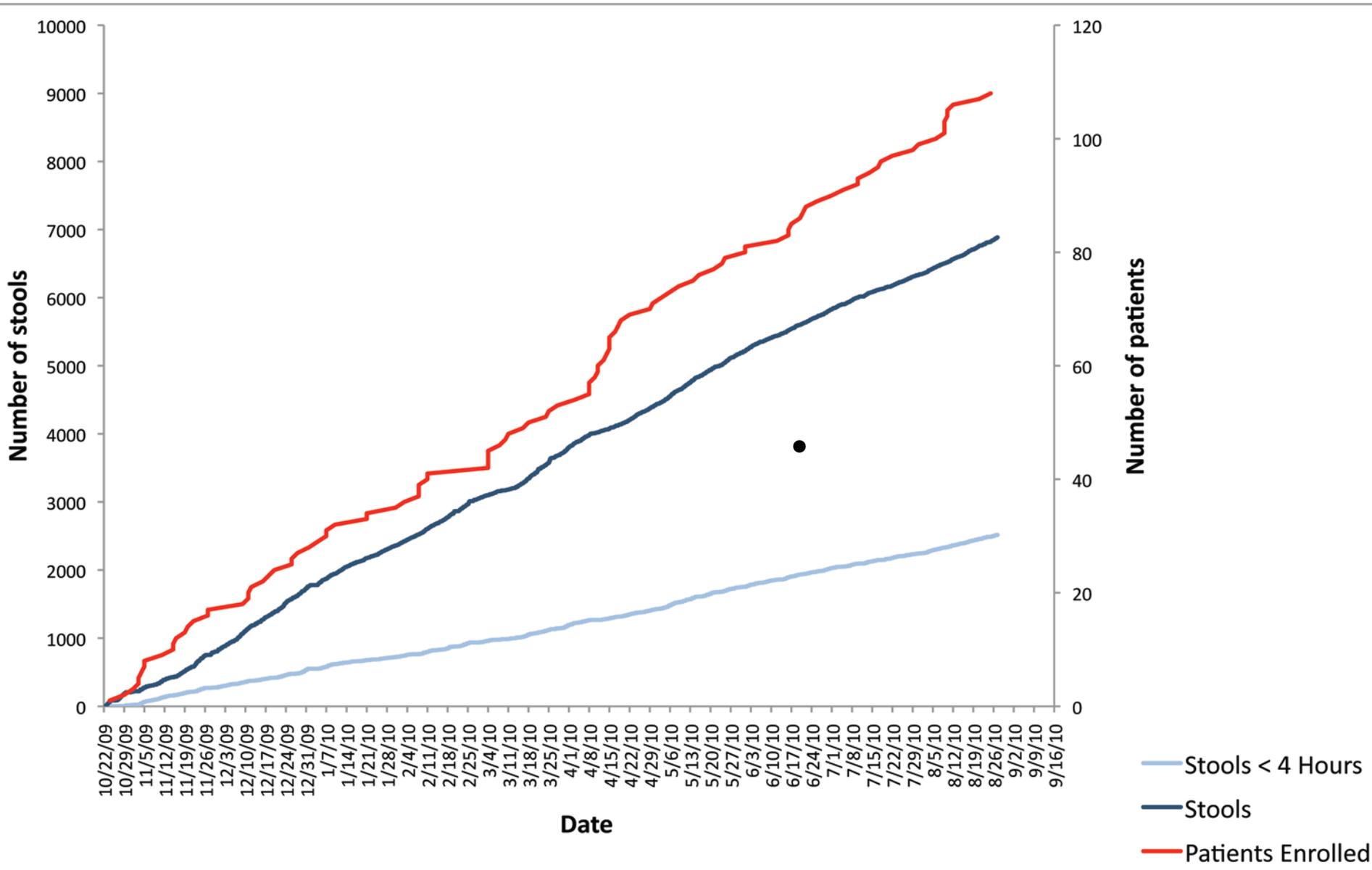
Weekly bloods

Daily 'real time' data

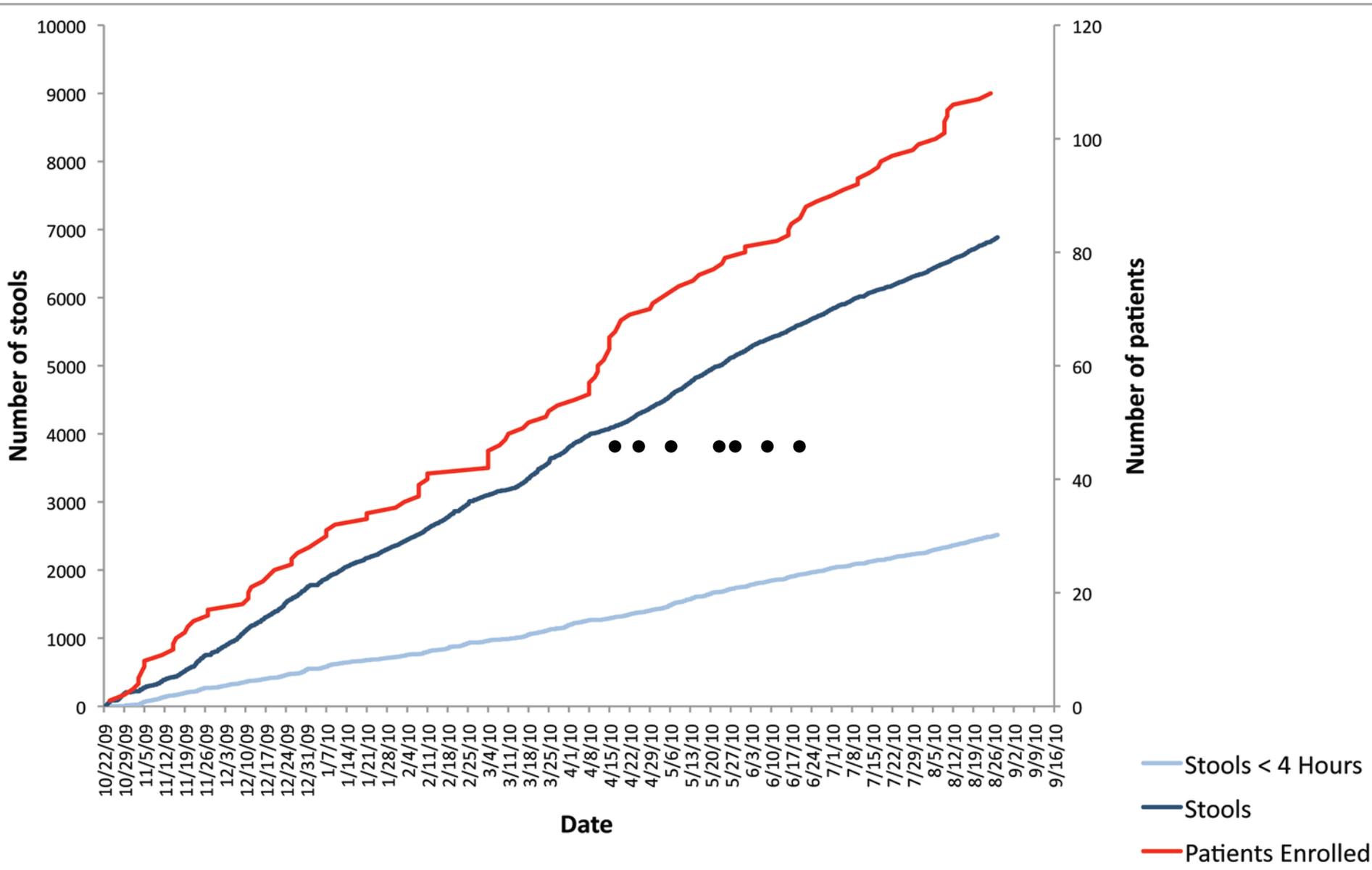
(esp. feeds and antibiotics)

Specimen provenance and curation

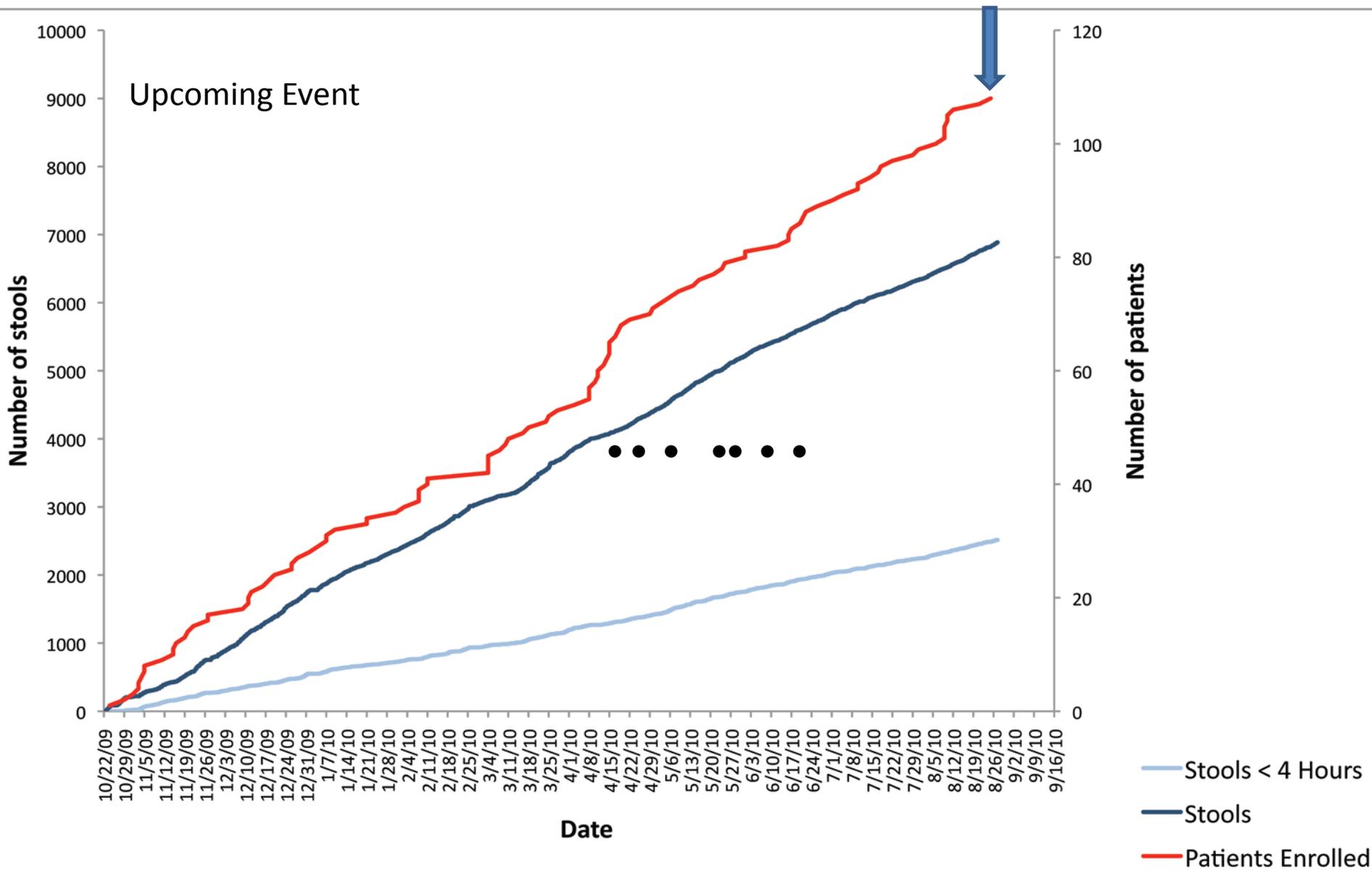




- Stools < 4 Hours
- Stools
- Patients Enrolled

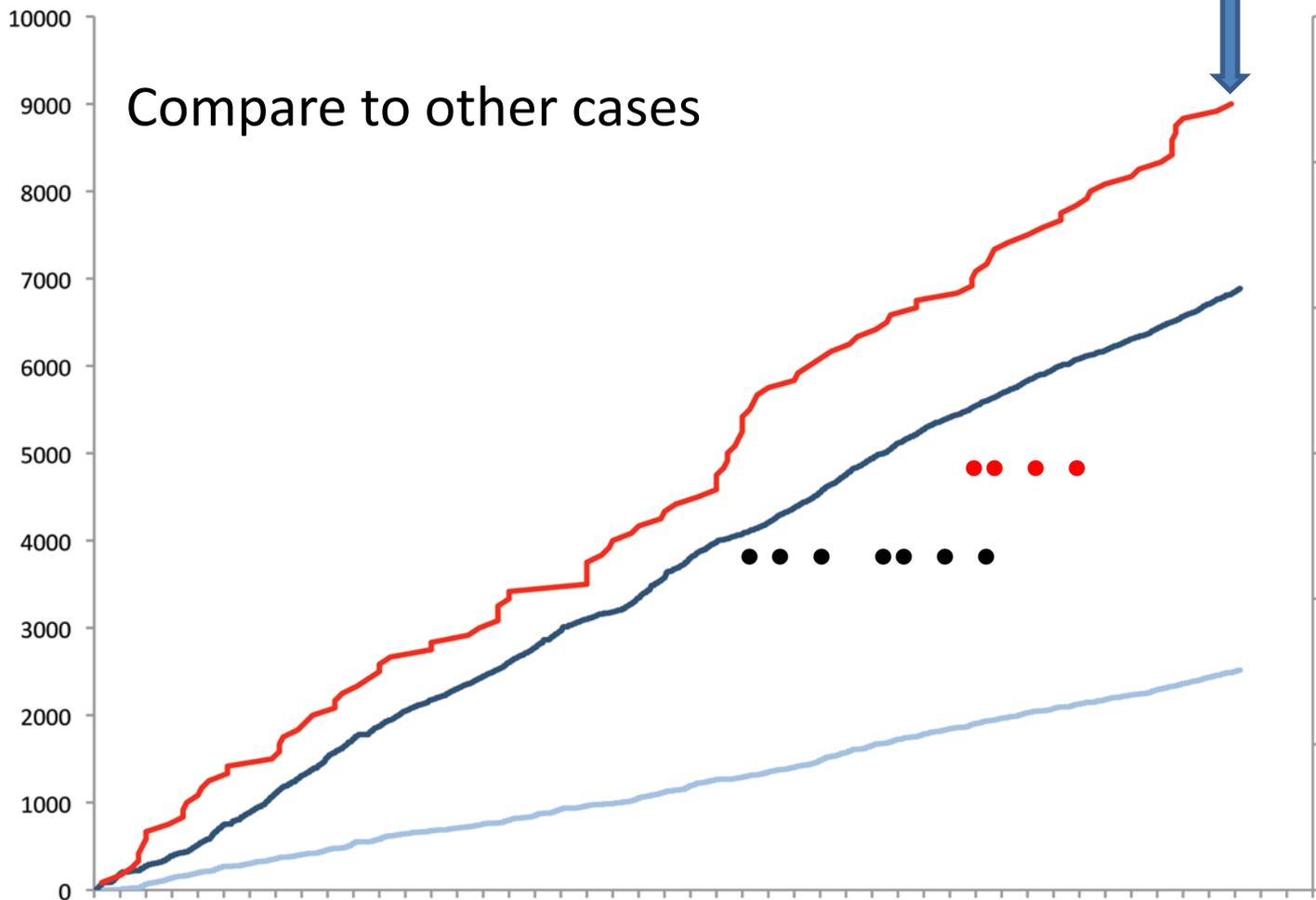


— Stools < 4 Hours  
 — Stools  
 — Patients Enrolled



Number of stools

Compare to other cases



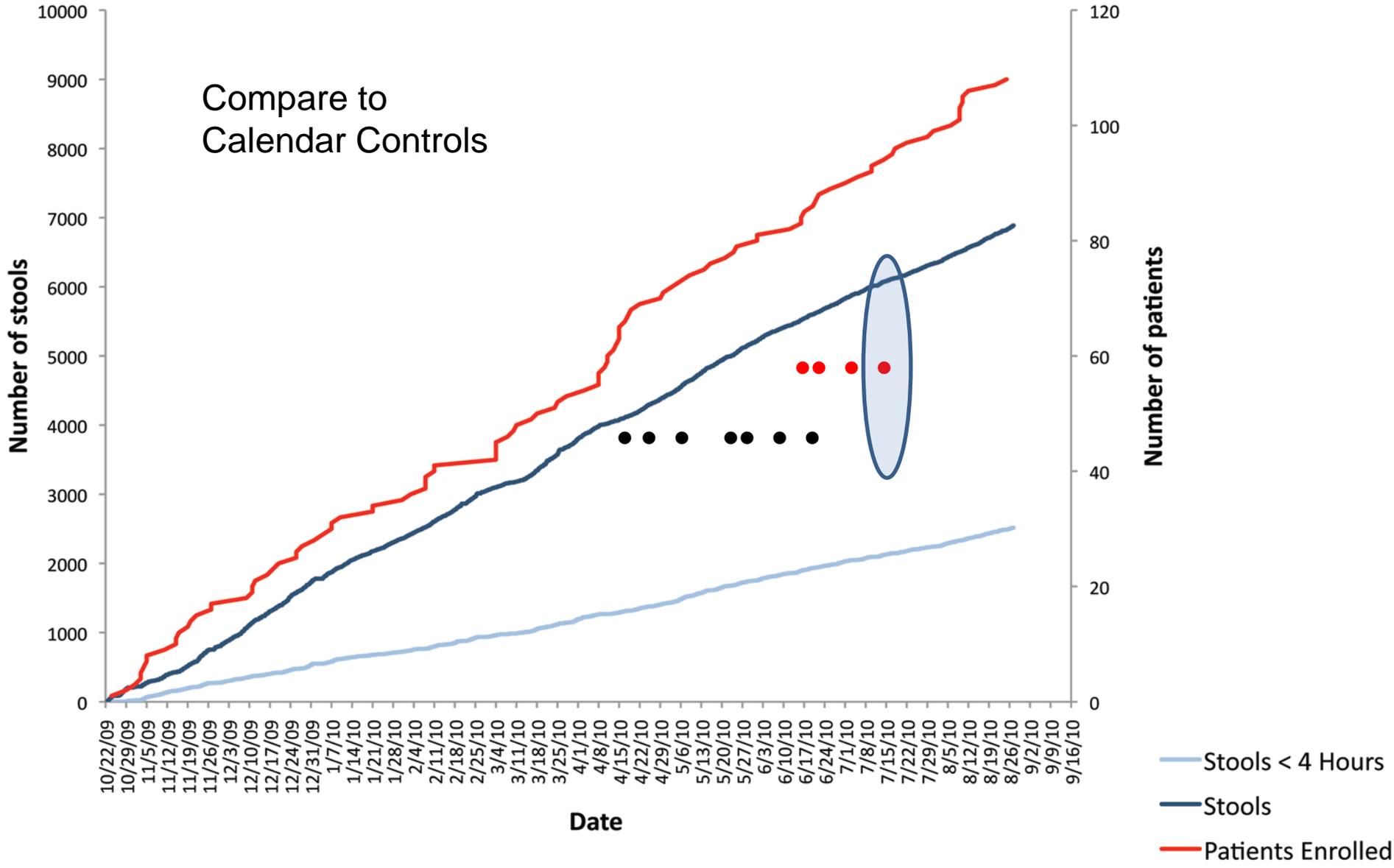
Number of patients

10/22/09 10/29/09 11/5/09 11/12/09 11/19/09 11/26/09 12/3/09 12/10/09 12/17/09 12/24/09 12/31/09 1/7/10 1/14/10 1/21/10 1/28/10 2/4/10 2/11/10 2/18/10 2/25/10 3/4/10 3/11/10 3/18/10 3/25/10 4/1/10 4/8/10 4/15/10 4/22/10 4/29/10 5/6/10 5/13/10 5/20/10 5/27/10 6/3/10 6/10/10 6/17/10 6/24/10 7/1/10 7/8/10 7/15/10 7/22/10 7/29/10 8/5/10 8/12/10 8/19/10 8/26/10 9/2/10 9/9/10 9/16/10

Date

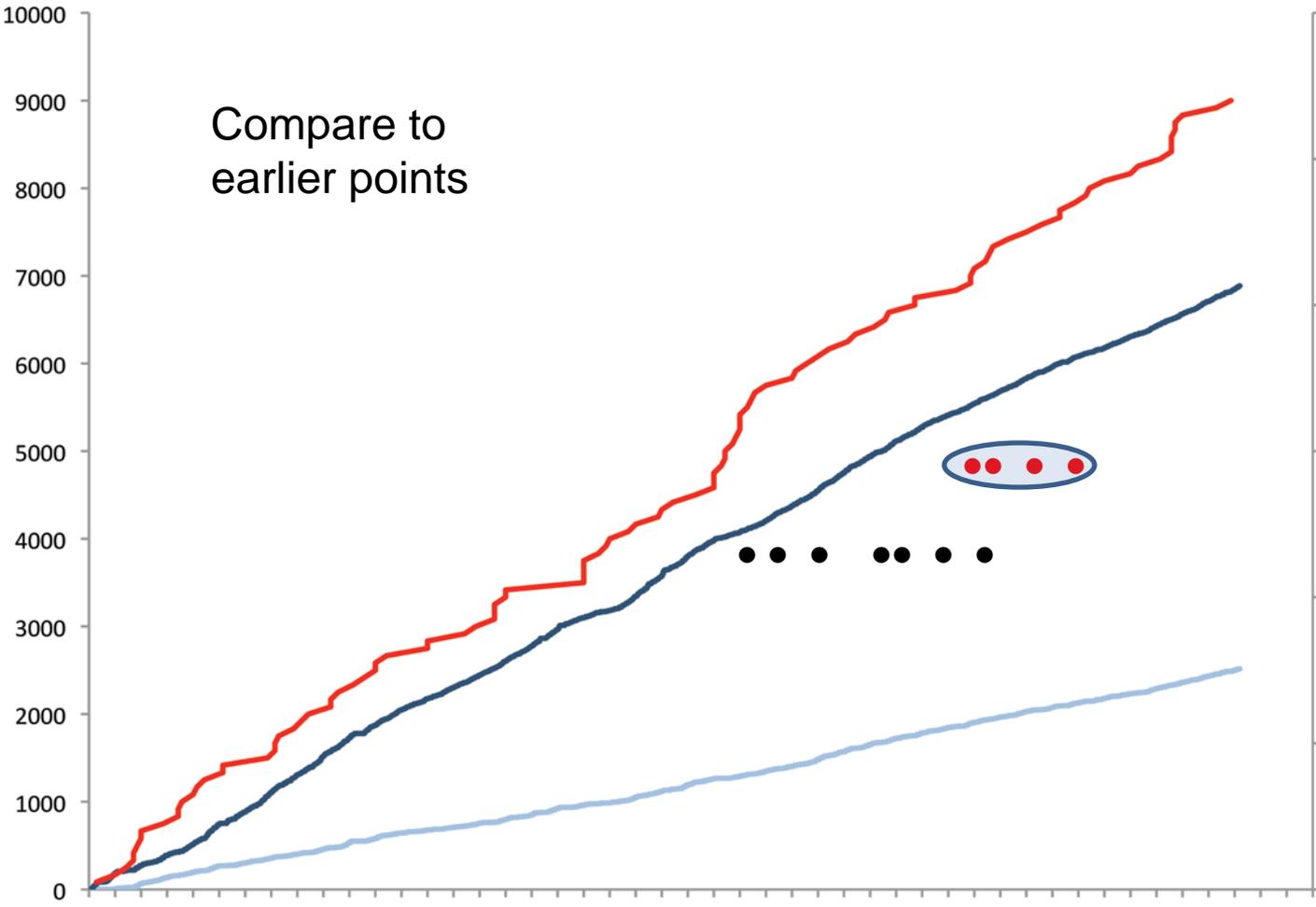
- Stools < 4 Hours
- Stools
- Patients Enrolled

Compare to  
Calendar Controls



Number of stools

Compare to  
earlier points

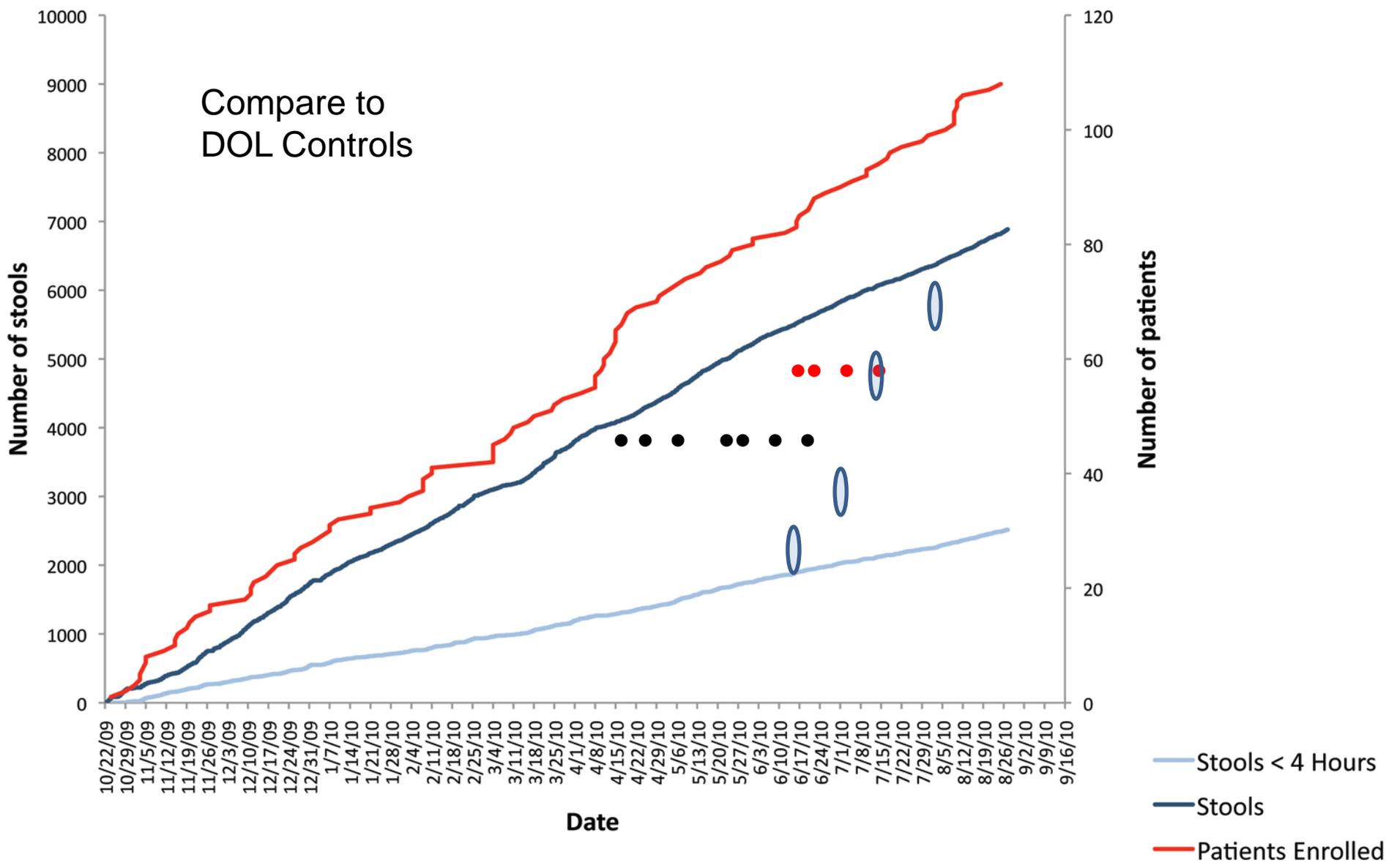


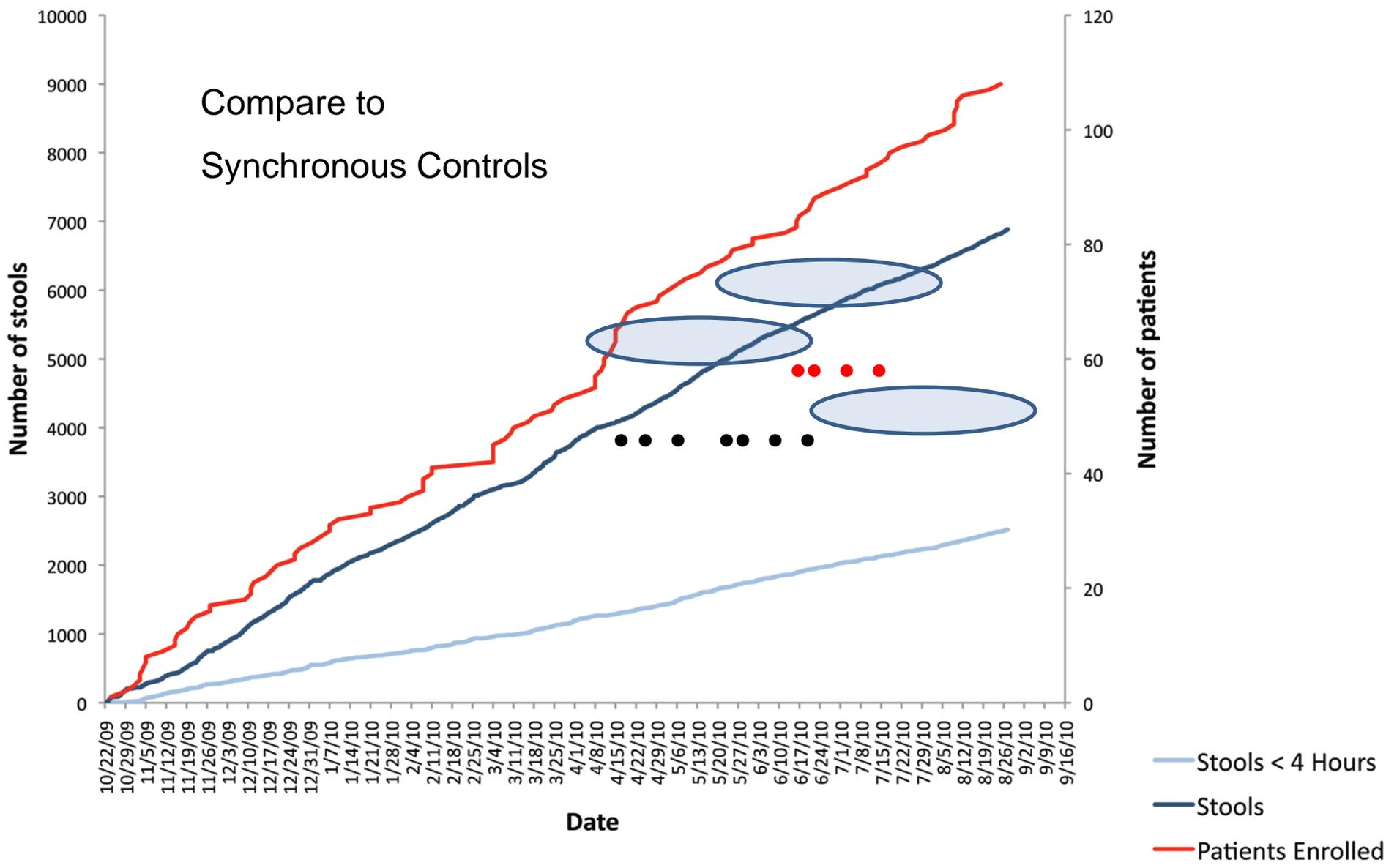
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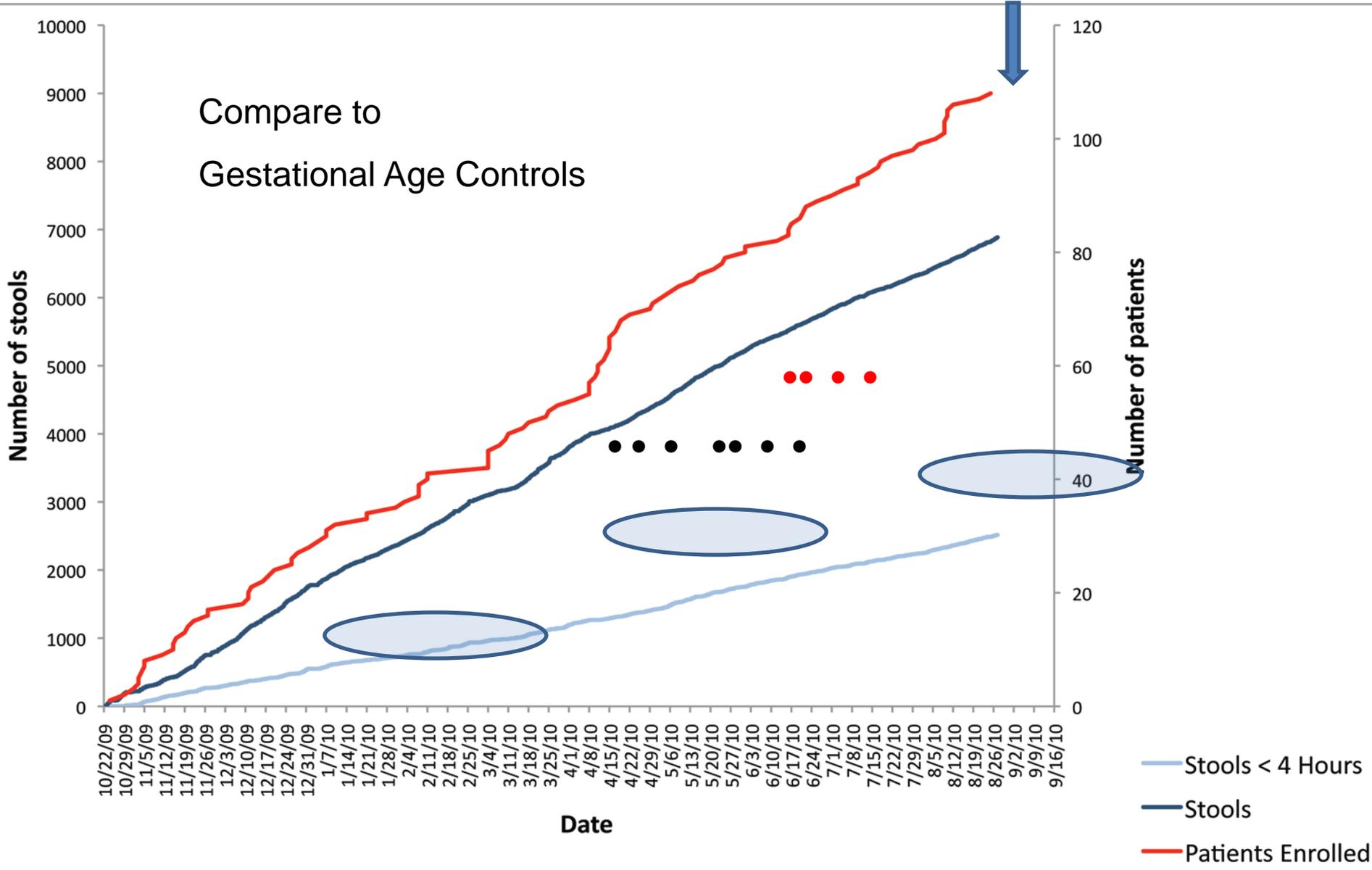
Date

- Stools < 4 Hours
- Stools
- Patients Enrolled





- Stools < 4 Hours
- Stools
- Patients Enrolled



# Diminished Progression towards Bacterial Diversification

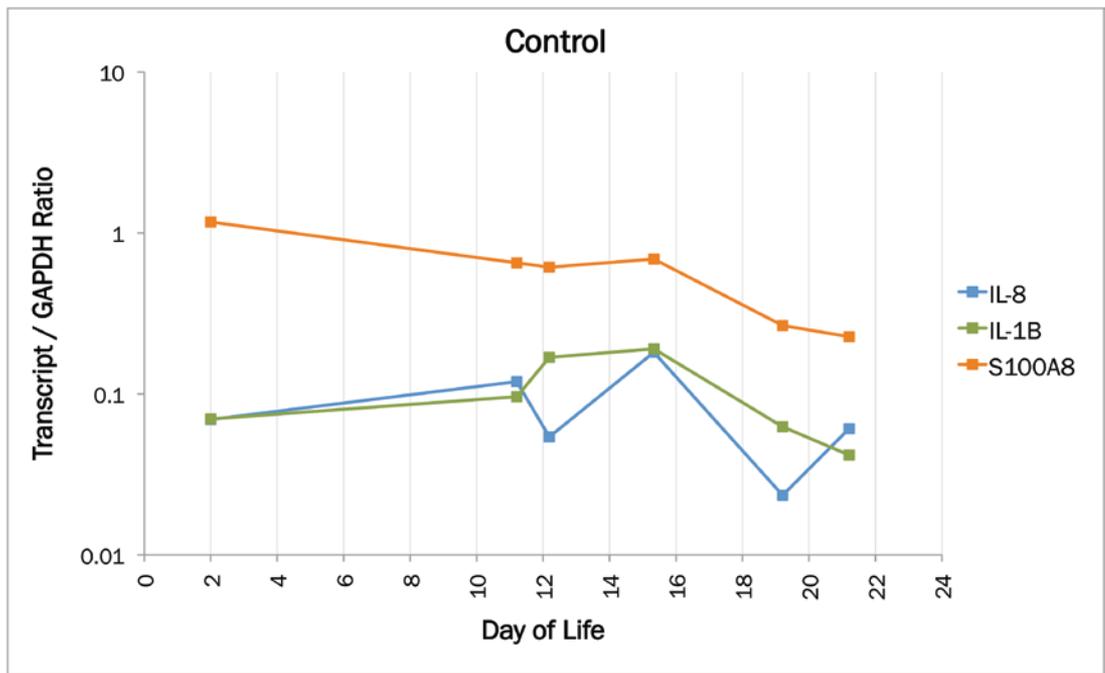
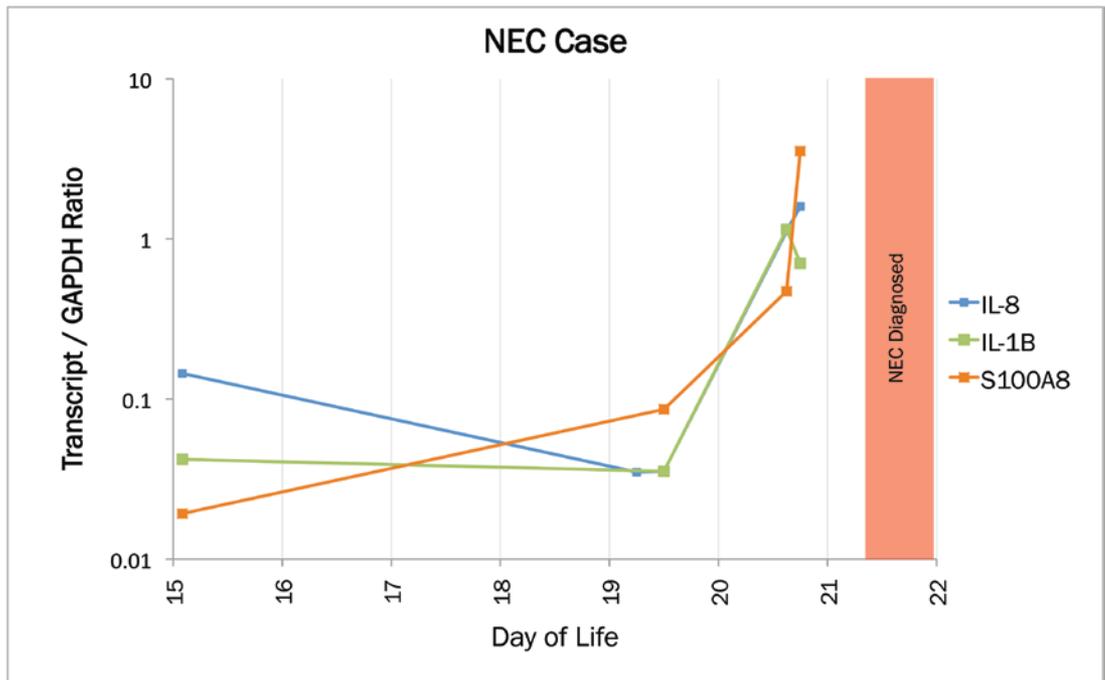
October – November 2009

Outcome	Model	Day of Life		% of days on antibiotics	
		est	P	est	p
<b>Shannon Index</b>	Continuous predictor, mixed model	0.0096	0.0013	-0.40713	0.0363
<b>Simpson Index</b>	Continuous predictor, mixed model	0.0033	0.0082	-0.2423	0.0027
<b>Gram negative per 1000 reads</b>	GEE Model, Poisson, distribution, continuous predictor	ns	ns	ns	ns
<b>Cocci per 1,000 reads</b>	GEE Model, Poisson distribution	-0.011	0.034	ns	ns
<b>Aerobic per 1000 reads</b>	GEE Model, Poisson distribution	ns	ns	ns	ns

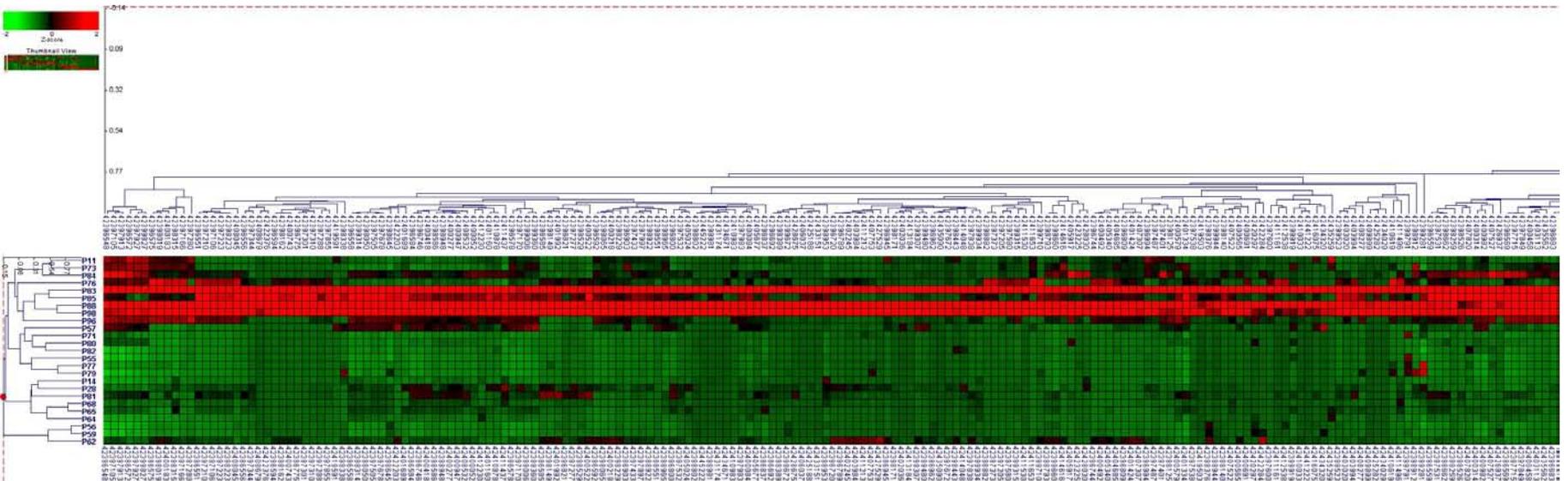
February – June 2010

Day of Life		% of days on antibiotics	
est.	p	est.	p
-0.015 + .0049 (D15)	<.0001	ns	ns
- 0.0472+ 0.00174 (D12)	<.0001	ns	ns
-0.067	0.0001	ns	ns
-0.041	0.0008	ns	ns
0.0367	0.003	ns	ns

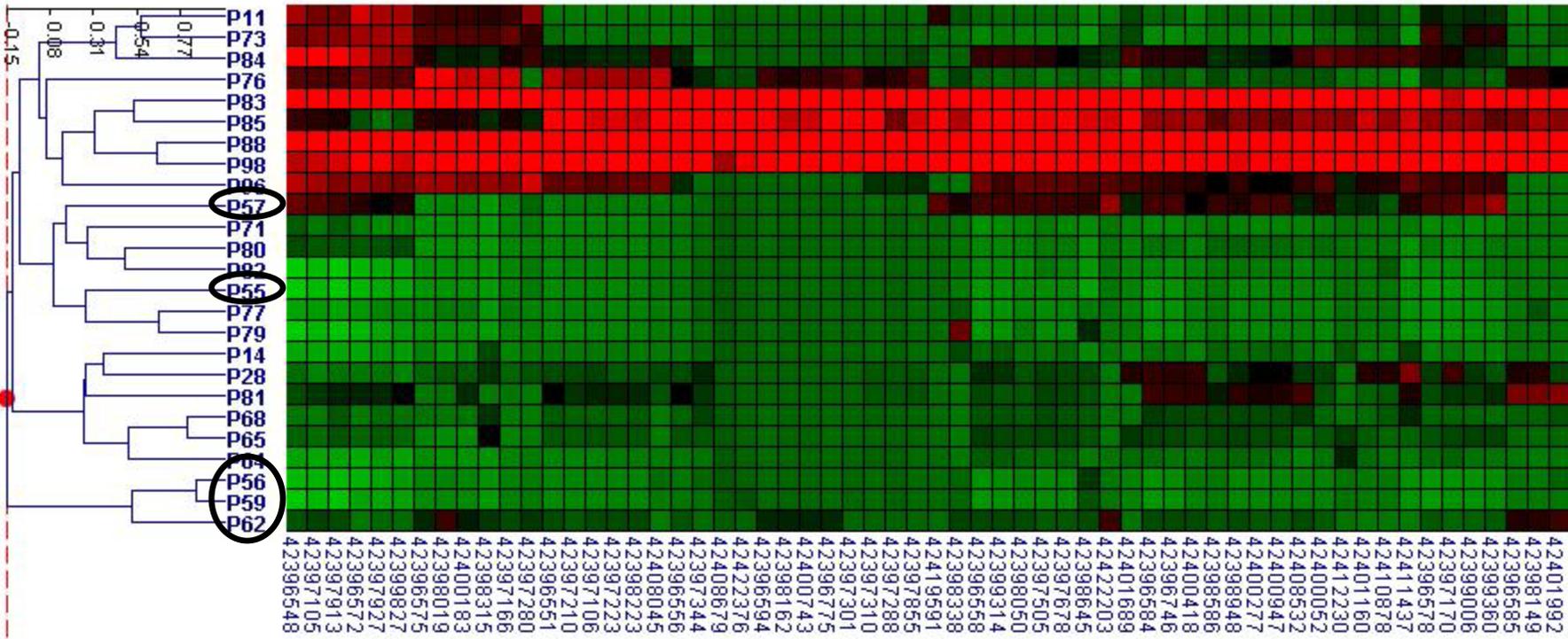
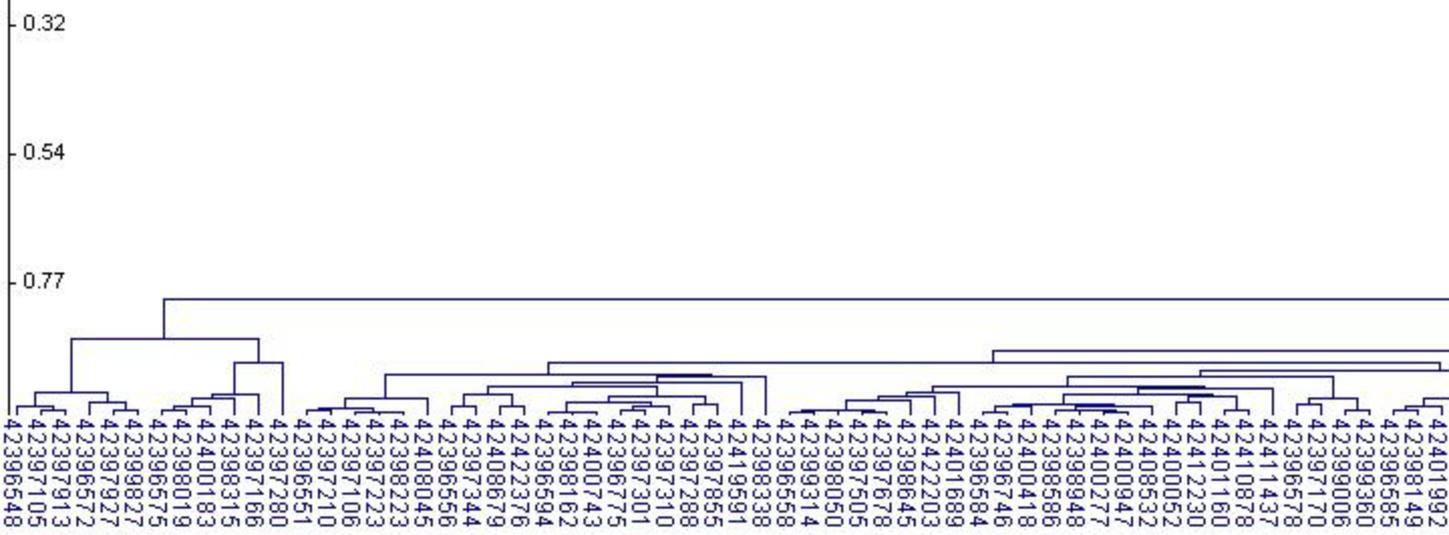


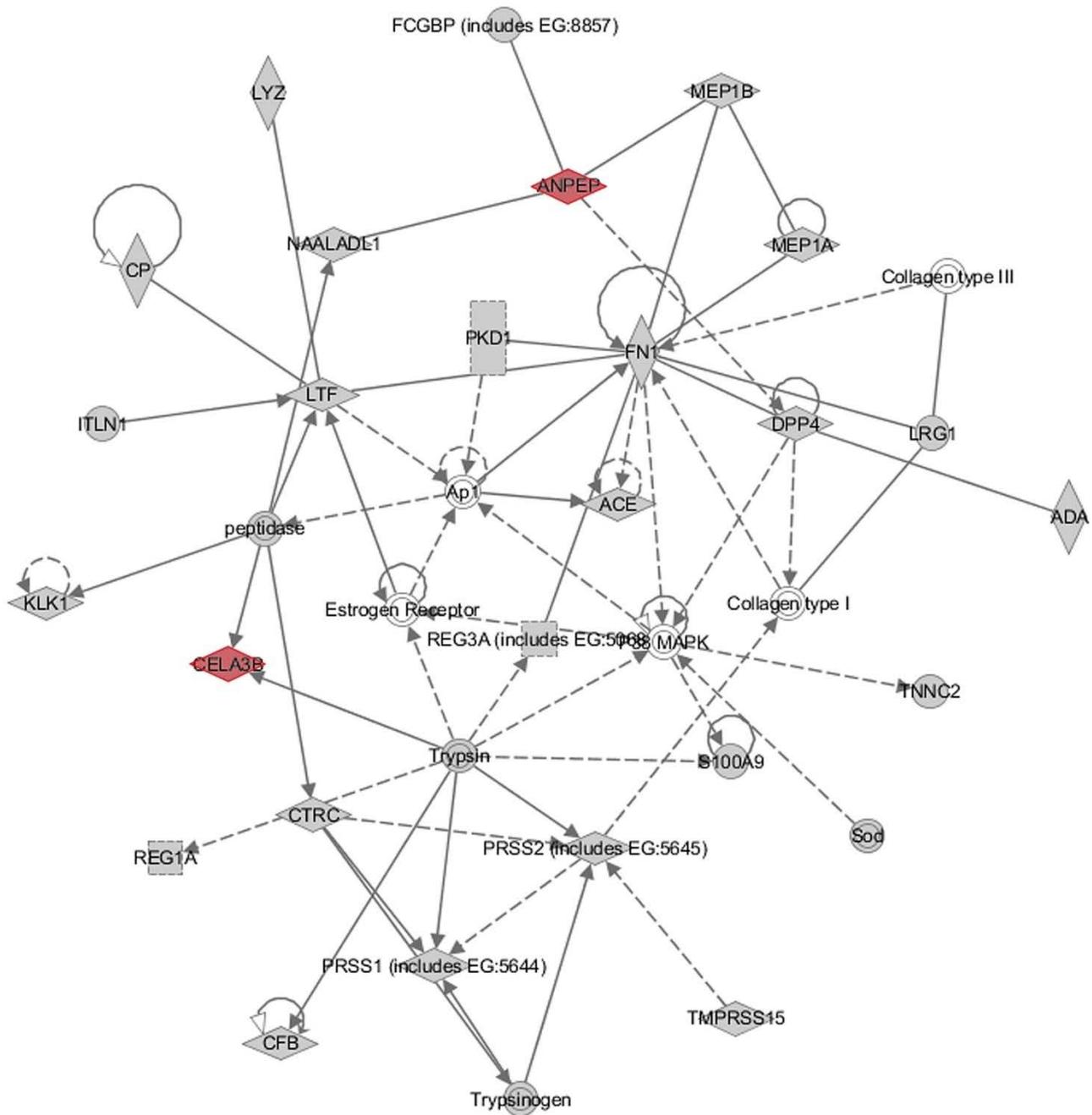


# Is there a NEC protein signature?











# VRE

Boy born after 31 weeks gestation (BW = 1250 g)

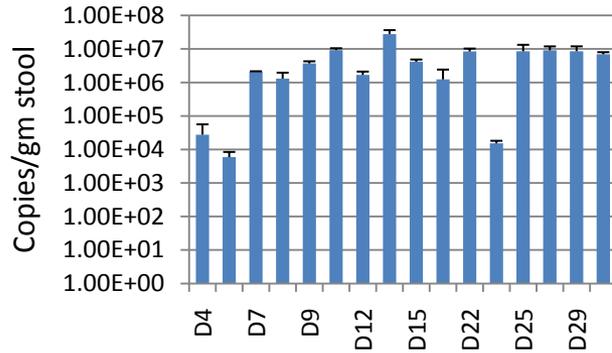
Received antibiotics on days 1-3 of life, gaining and growing

On DOL 21, stool culture screen:

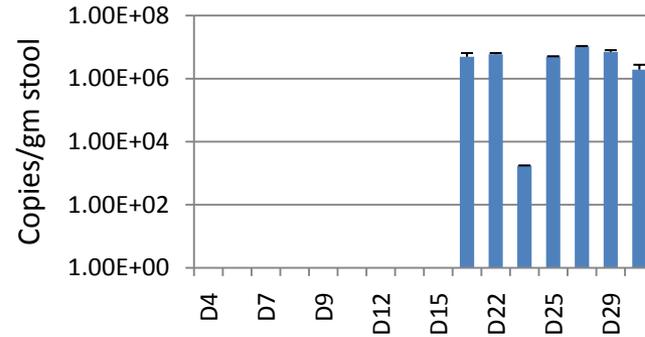
vancomycin resistant enterococci (VRE)

Nine pre-positive and seven post-positive stools available

### Ent16s

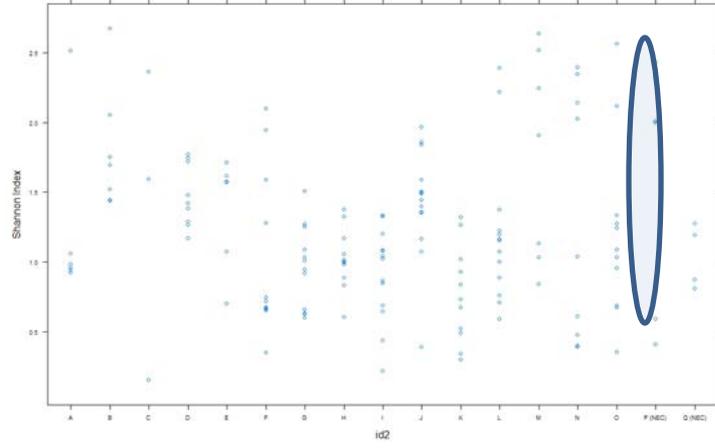


### vanA

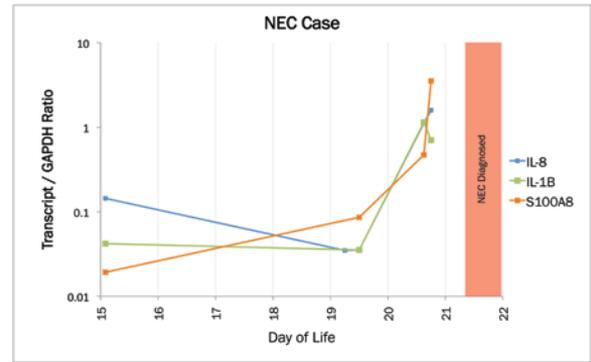


Specimen	Subjects (n)	Specimens (n)	Enterococci (log10)/g (median, range)	vanA percent of enterococci
Pre VRE	1	9	6.1 (2.7-6.9)	0
Post VRE	1	7	6.3 (4.2-7.1)	71% (11-100%)
All-patient	1	16	6.1 (2.7-7.1)*	0
Controls	8	59	4.5 (2.2-7.1)*	0
			* p<0.0001	

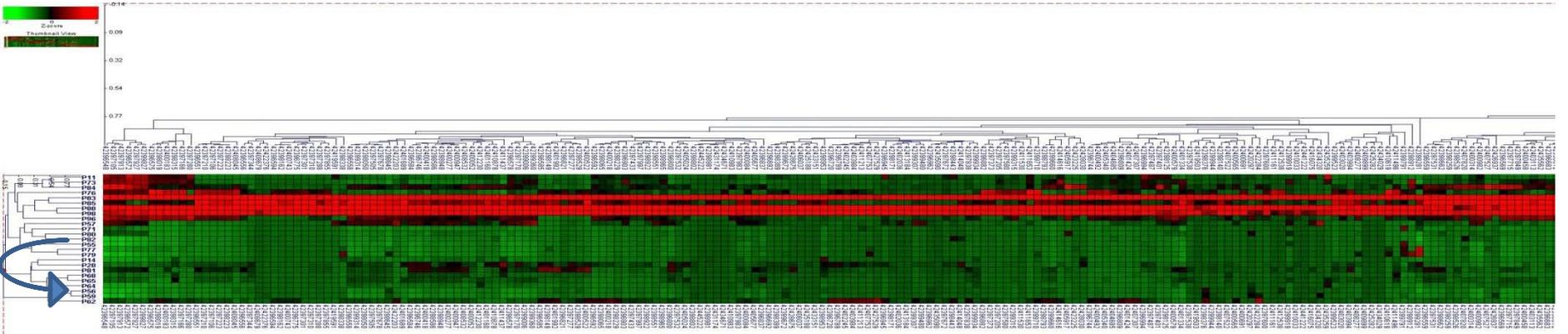
1.



2.



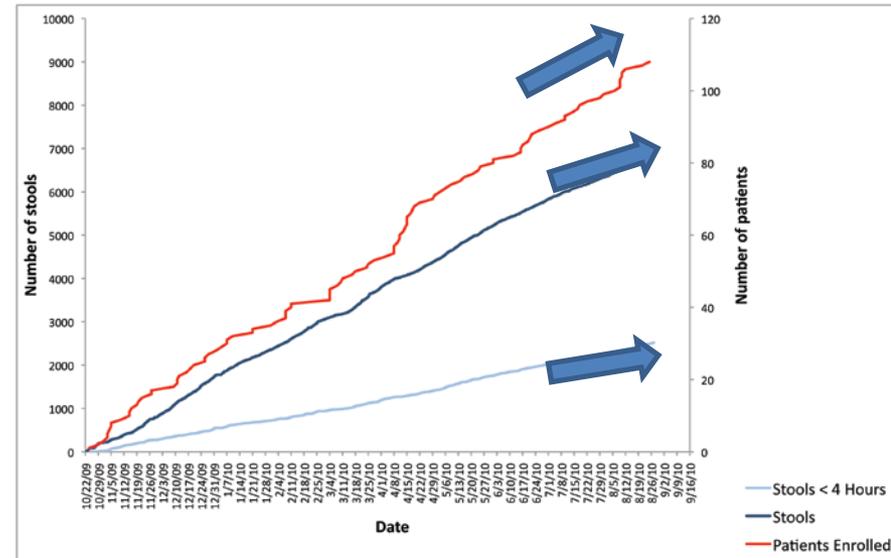
3.





# Future Directions

- Extended accrual
- Shotgun sequencing
- Metatranscriptomics
- Viral discovery
- Proteomics



# Thanks!

NICU Families and Nurses

St. Louis Infant Microbiome Team

Subjects: Barb Warner, Julie Hoffman, Laura Linneman, Malick Ndao, Aaron Hamvas

Laboratory: Harry Stevens, Billy Bennett, Nur Shaikh, Katie Farrell, Raffi Lev-Tzion, Katie Hoffman, Katie Becker

Genome Center: Erica Sodergren, George Weinstock, David Wang, Lucinda Fulton, Otis Hall, Kathy Mihindukulasuriya, Michelle O'Laughlin, Vince Magrini

Statistics: Bill Shannon, Elena Deych, Jia Wang

Proteomics: Reid Townsend, Petra Gilmore, Jim Malone,  
Alan Davis

NIAID: Melody Mills and UH2 AI083265