

The gut microbiota and SES in preterm infants in the Chicago area

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Prematurity

- What is full term? 37-40 weeks gestation
- What is prematurity? < 37 weeks gestation
- What is the limit of viability? 22-23 weeks gestation

Prematurity



Full Term, 40
weeks



Premature, 23 weeks

Smallest Survivor

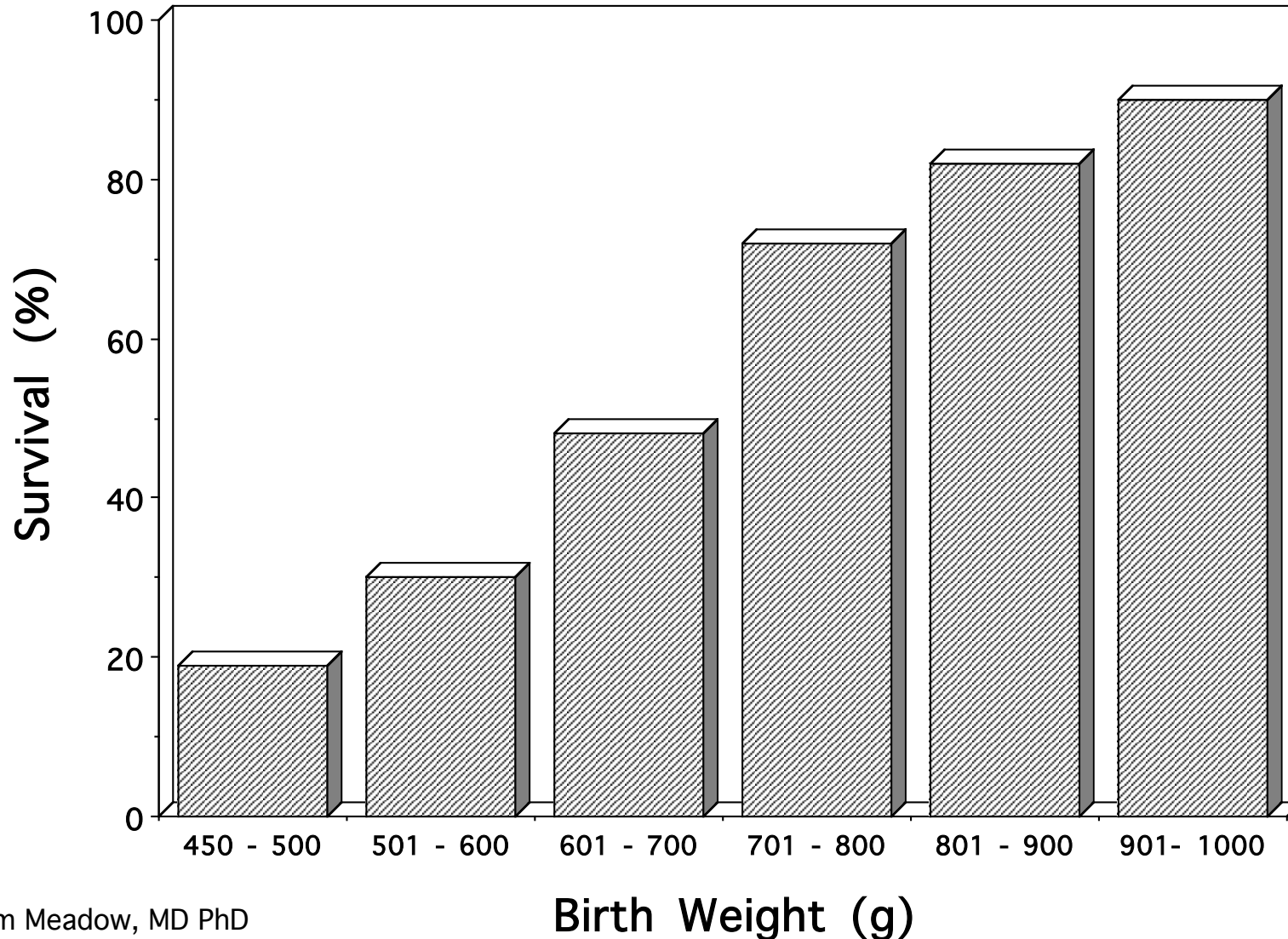


Birth weight 8.6 oz

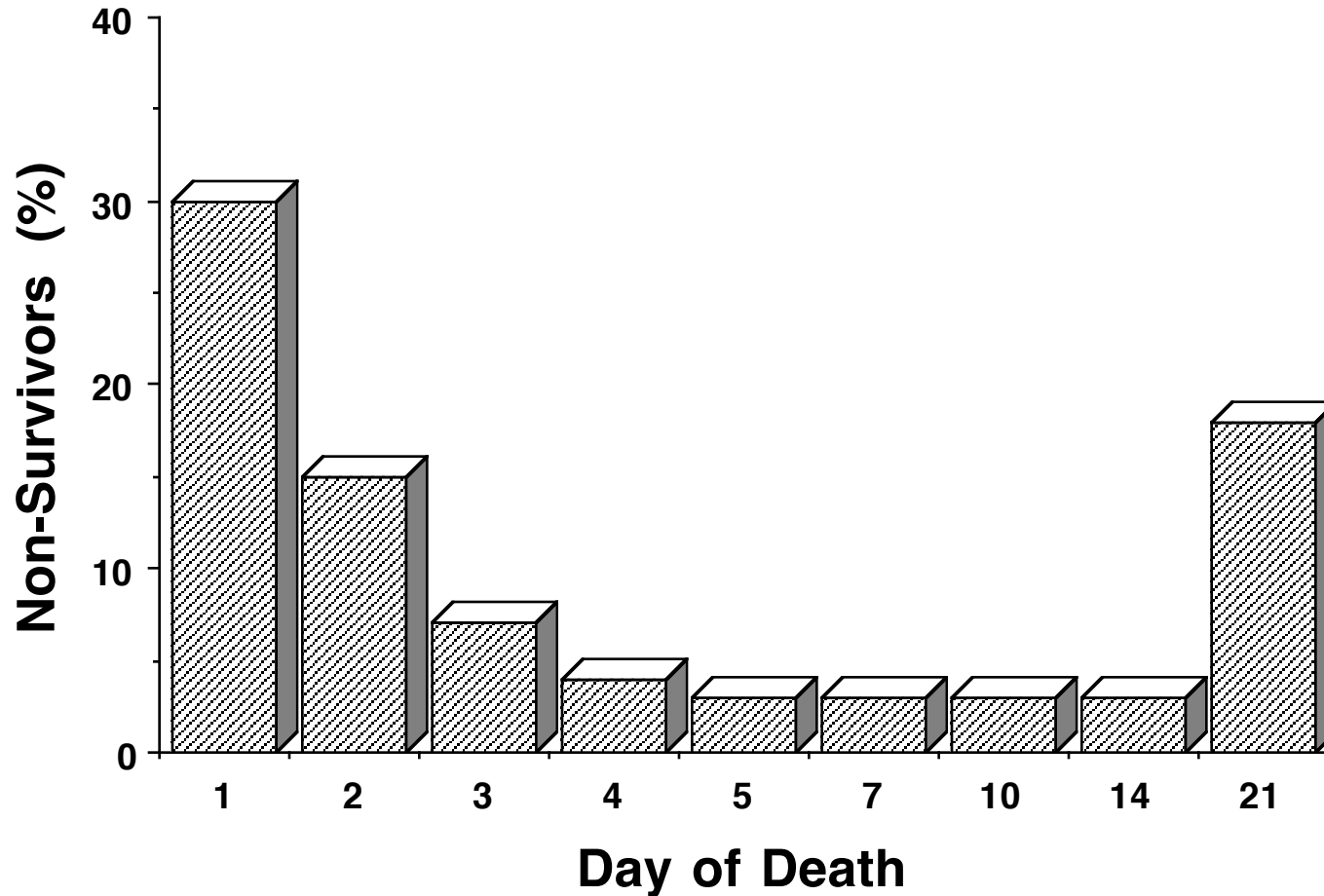


12 oz.

Survival vs Birth Weight for ELBW Infants

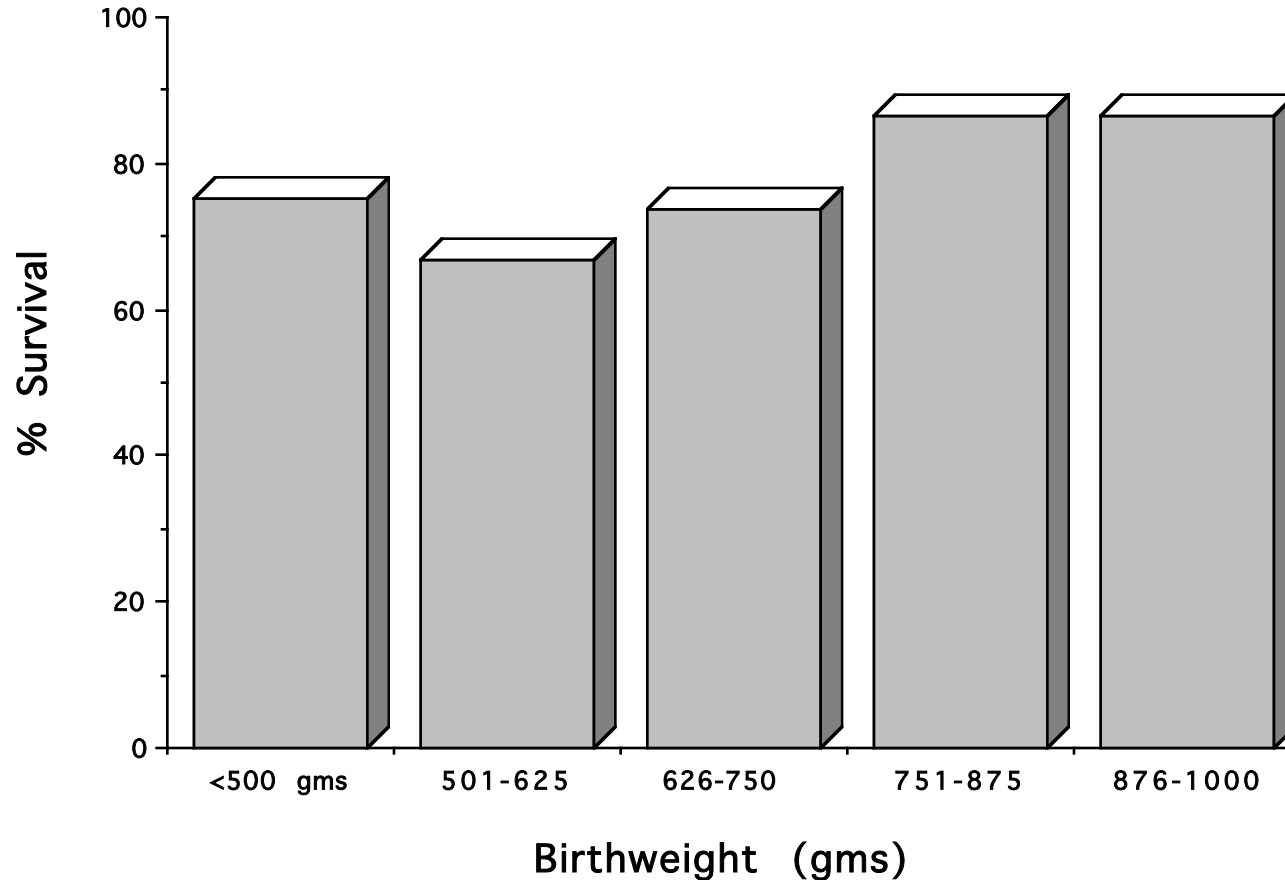


Day of Death for ELBW Non-Survivors



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The University of Chicago

Survival as a function of birthweight for all patients alive on Day 4 (n = 249)



The Preterm Infant - 12% of births



Disproportionately account for:

40% of children who have cerebral palsy (CP)

25% of children with hearing impairment

35% of those with vision impairment.

The NICU

NICU

Antibiotics

Delayed Feeding

Breast Milk vs. Formula

H2 Blockers

Opioids

Instrumentation



The Value of Preterm Infant Environmental Health Cohorts The Canary in the Coal Mine

Annemarie Stroustrup, MD, MPH^{1,2}; Susan L. Teitelbaum, PhD²; Judy L. Aschner, MD³

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Like the canary in the coal mine—or asthmatics in air pollution studies—children born preterm may serve as a sentinel population owing to increased susceptibility to the sometimes modest effects of common toxicants, improving study power and decreasing necessary sample size.

Necrotizing Enterocolitis (NEC)-

Inflammatory bowel necrosis that primarily afflicts premature infants after the initiation of enteral feeding.

Risk Factors

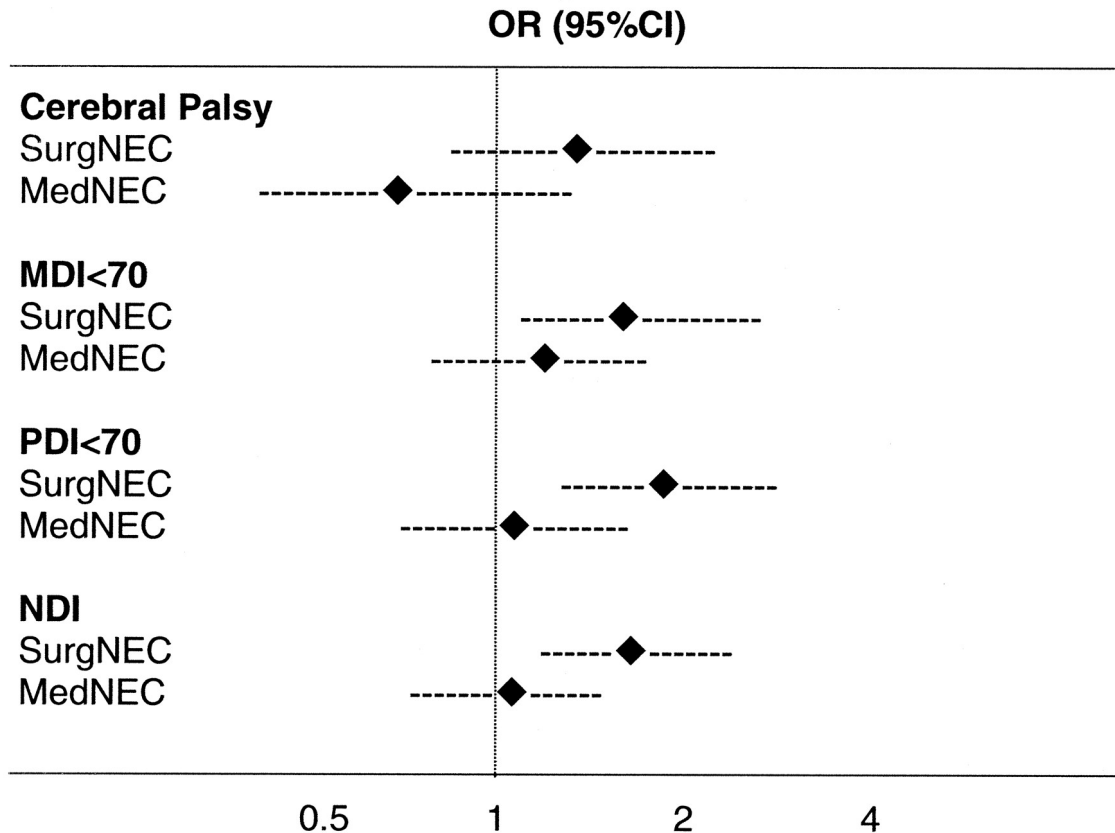
- Prematurity
- Bacterial Colonization
- Enteral Feeding
- Hypoxia/Altered intestinal blood flow

NEC and Neurologic Outcome

Neurodevelopmental and Growth Outcomes of
Extremely Low Birth Weight Infants after NEC Hintz,
et al Pediatrics 2005

Multicenter, retrospective analysis 1995-1998
Infants in NICHD Neonatal Network <1000gm
5553 ELBW entered into registry
2948 infants evaluated at 18 and 22 months
124 – surgically managed NEC
121 – Medically managed NEC

Neurodevelopmental Outcome associated with NEC



SurgNEC

PVL 14% vs. 7%
 BPD 57% vs. 43%
 CP 24% vs 15%
 Decreased growth
 all parameters

Hypothesis:

Enteral feeding results in colonization of the uniquely susceptible premature intestine with pathogenic bacteria, resulting in an exaggerated inflammatory response.

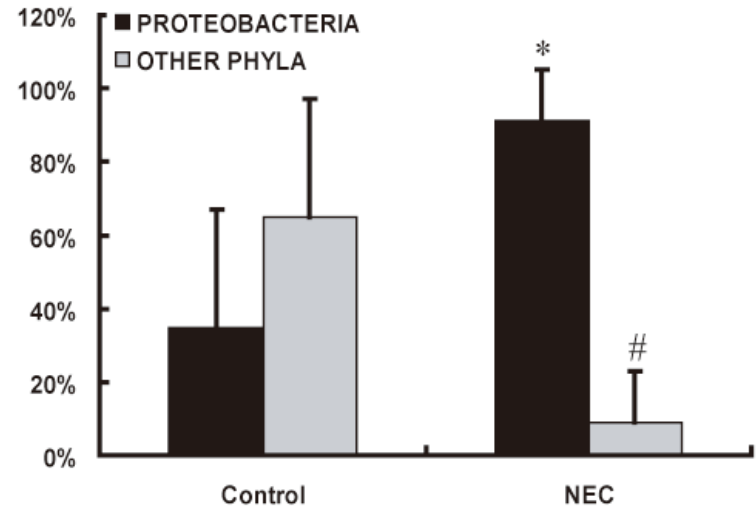
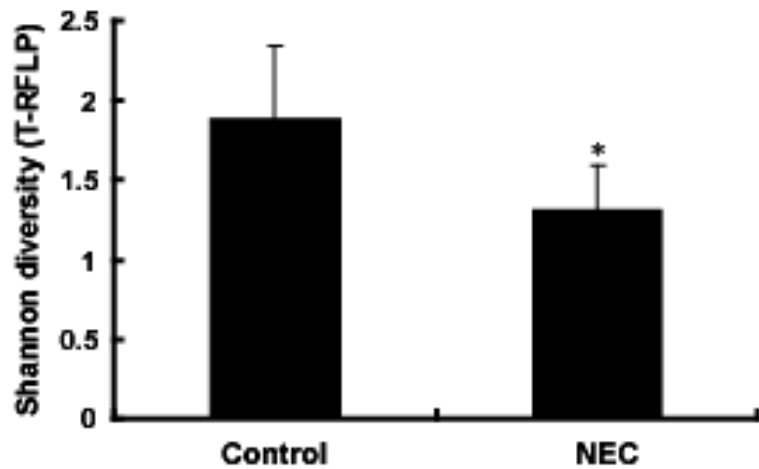
Question:

- Can microbiome analysis be used to identify the pathogenic bacteria associated with NEC?

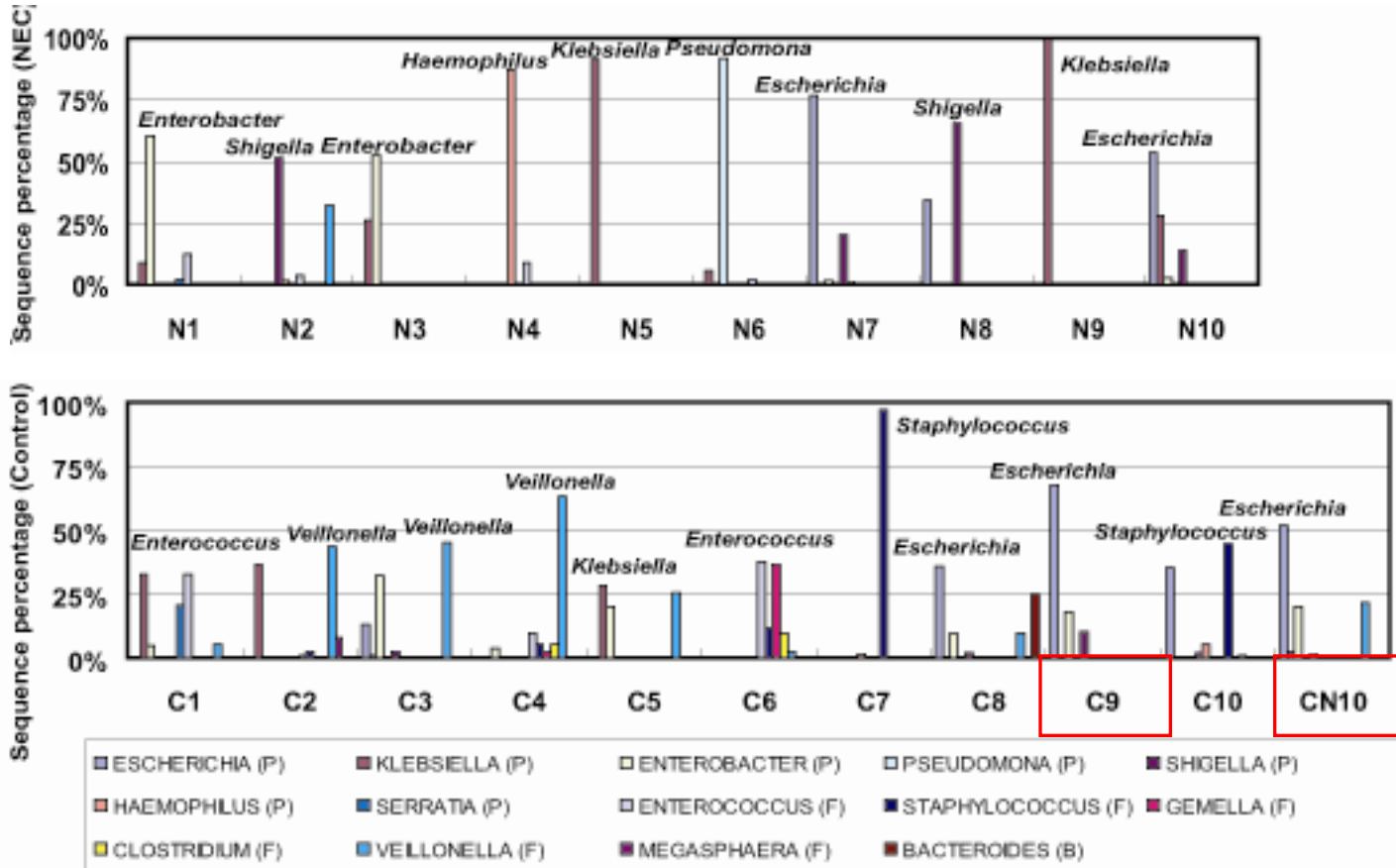
Methods:

- 20 patients - 10 with NEC 10 control
 - 4 sets of twins
- Analysis of fecal samples prior to onset of NEC by 16S rRNA sequencing

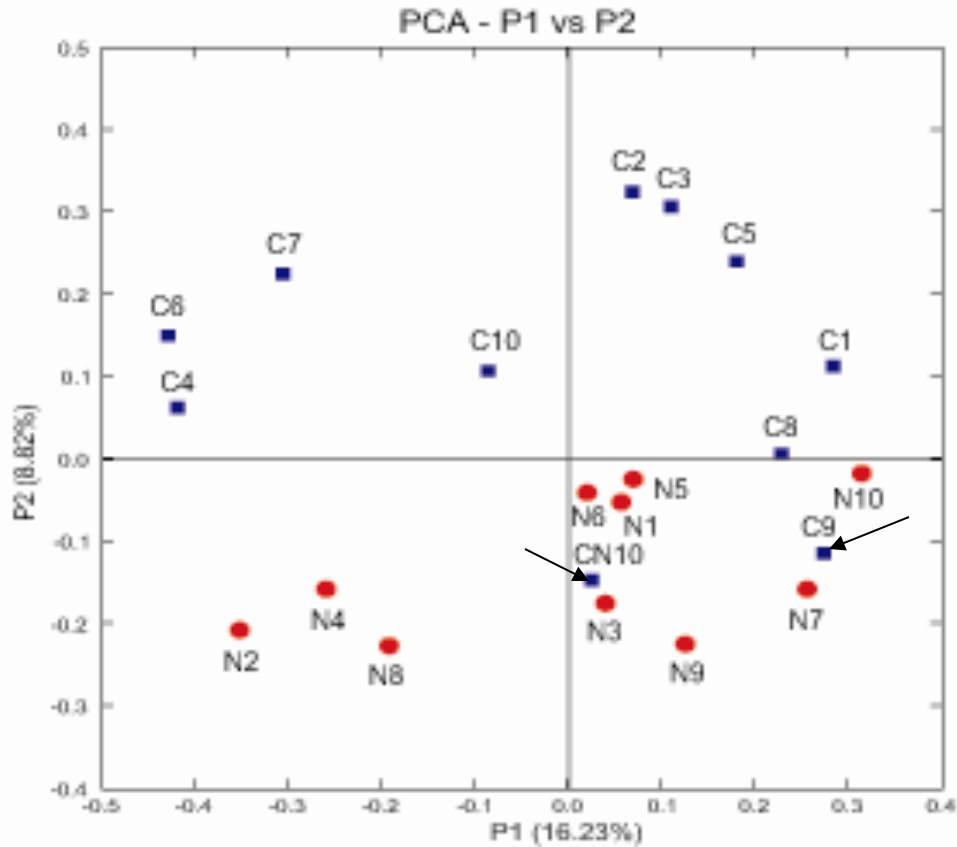
Bacterial Diversity and NEC



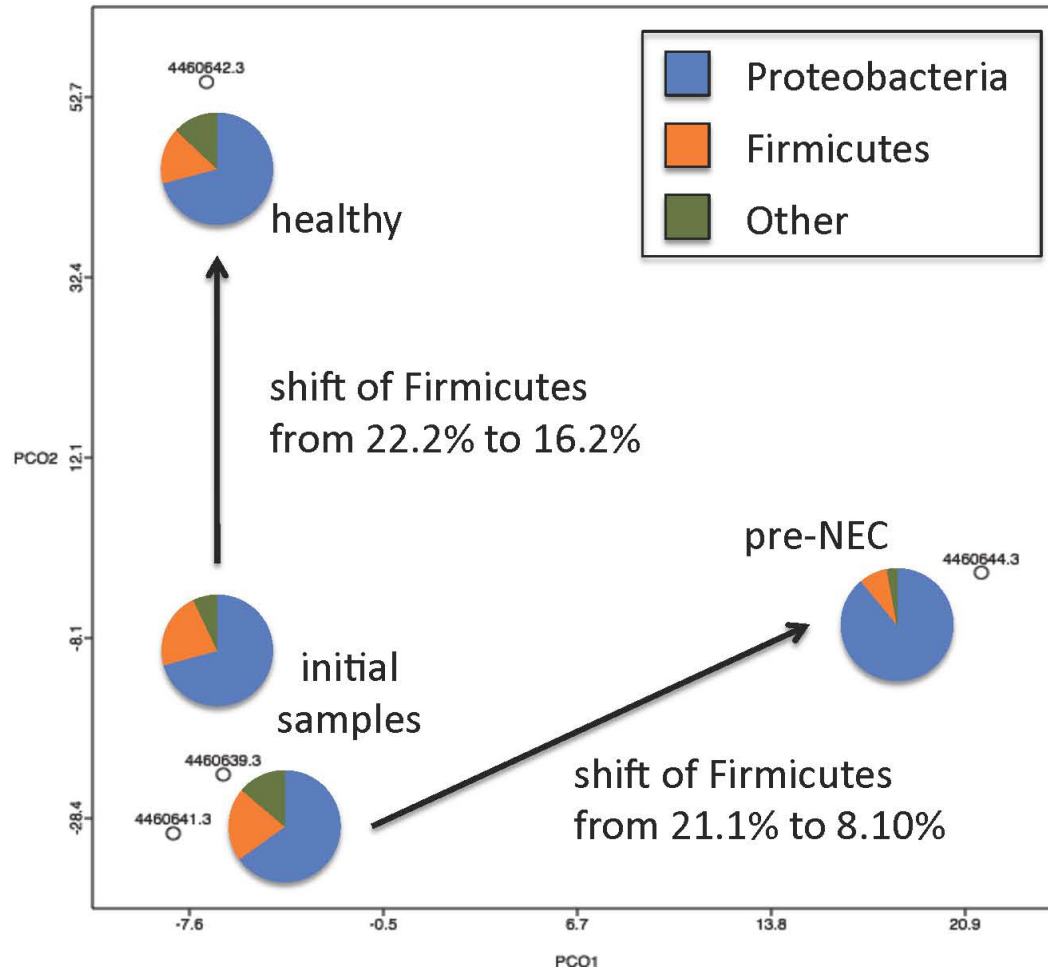
Bacterial Colonization and NEC



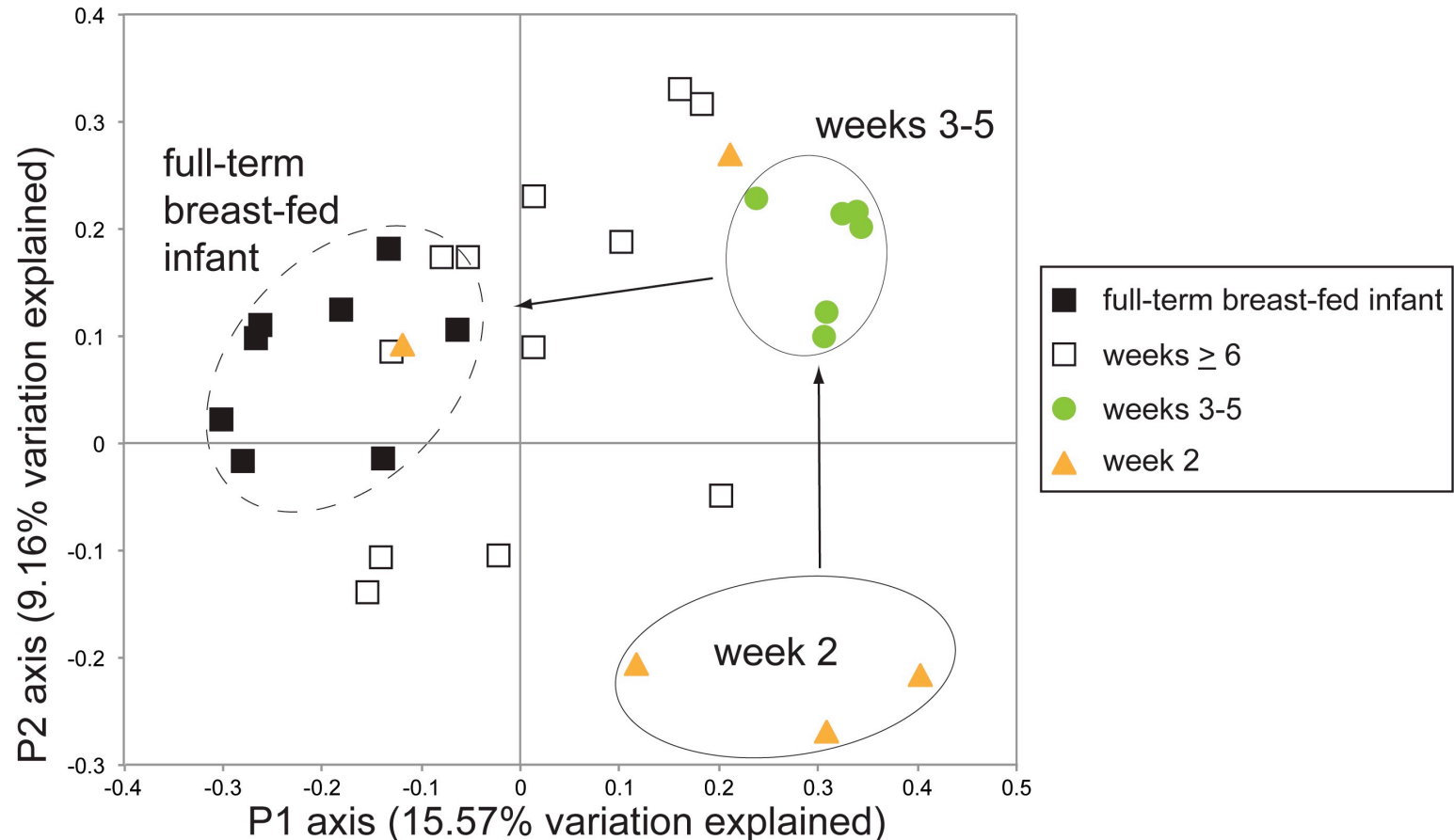
Bacterial Colonization and NEC



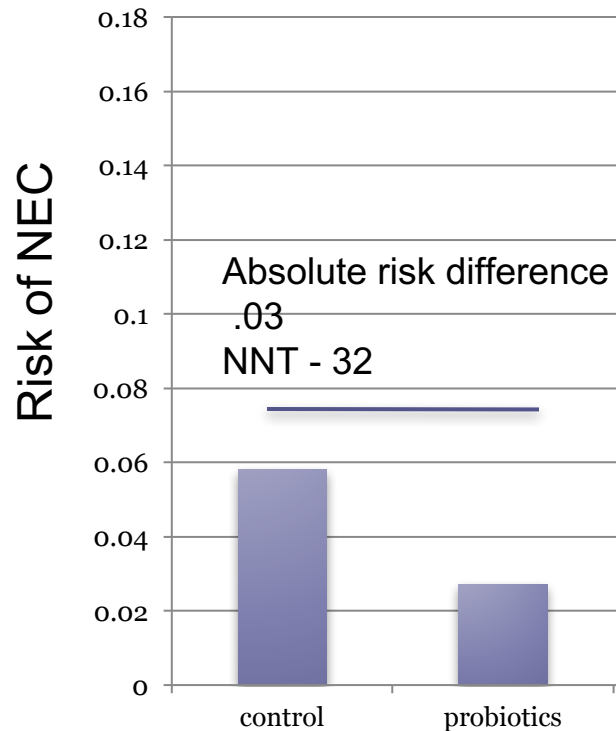
Shift in Microbiome



Temporal progression of the preterm infant microbiota



Modification of the early microbiome and NEC



AlFaleh K and Anabrees J The Cochrane Collaboration 2014;
Cotten et al Pediatrics 2009 January; 123: 58-66;
Meinzen-Derr et al J Perinatol 2009 January; 29 (1): 57-62

Prematurity



Full Term, 40 weeks



Premature, 23 weeks

Host development coincides with microbiome development

Growth in the Neonatal Intensive Care Unit Influences Neurodevelopmental and Growth Outcomes of Extremely Low Birth Weight Infants

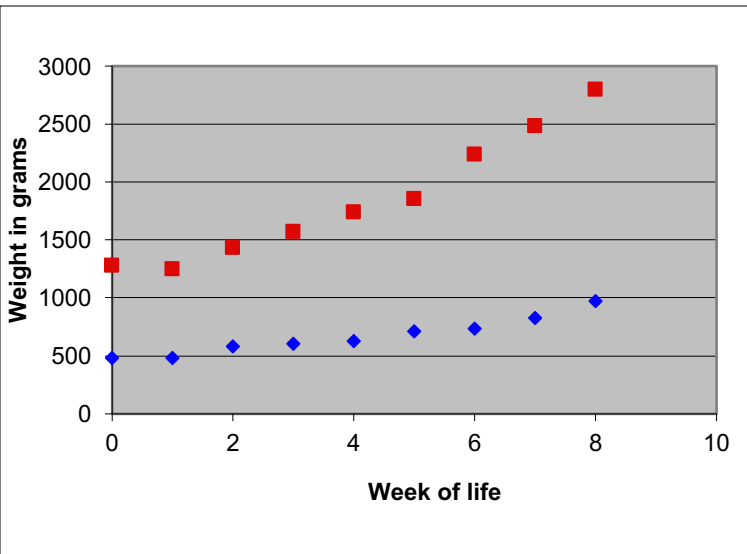
Richard A. Ehrenkranz, MD^a, Anna M. Dusick, MD^b, Betty R. Vohr, MD^c, Linda L. Wright, MD^d, Lisa A. Wrage, MPH^e, W. Kenneth Poole, PhD^e, for the National Institutes of Child Health and Human Development Neonatal Research Network

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CONCLUSIONS. These analyses suggest that growth velocity during an ELBW infant's NICU hospitalization exerts a significant, and possibly independent, effect on neurodevelopmental and growth outcomes at 18 to 22 months' corrected age.

Preterm Infant Microbiota (M_{PI})

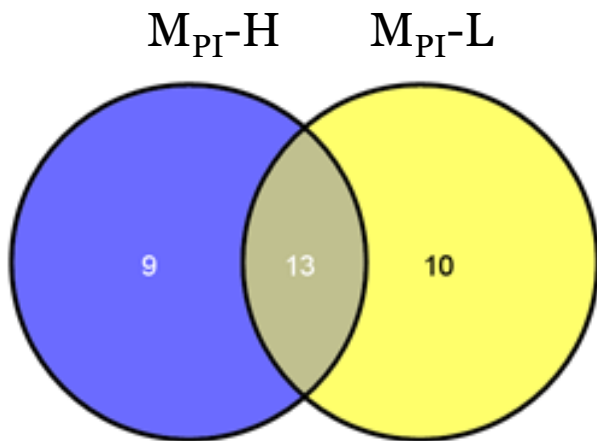


Good weight gain M_{PI-H}
12 do

Pyrosequencing ID	% of Total Flora
Comamonas sp	21
Weissella confusa	13
Leuconostoc citreum	10
Comamonas testosteroni	10
Lactococcus sp	10
Acinetobacter sp	3
Citrobacter sp	3
Lactococcus lactis	3
Delftia sp	2
Polaromonas sp	2
Alcaligenes sp	2
Enterococcus italicus	1
Streptococcus sp	1
Chryseobacterium bovis	1
Lactococcus raffinolactis	1
Aeromonas sp	1
Arcobacter butzleri	0.5
Brevundimonas sp	0.5
Fusobacterium sp	0.5
Bacteroides sp	0.5
Veillonella sp	0.5
Weissella cibaria	0.5

Poor weight gain M_{PI-L}
6 do

Pyrosequencing ID	% of Total Flora
Comamonas sp	16
Weissella confusa	15
Leuconostoc citreum	12
Lactococcus sp	12
Comamonas testosteroni	11
Citrobacter sp	6
Delftia sp	4
Lactococcus lactis	4
Acinetobacter sp	3
Streptococcus sp	2
Enterobacter sp	2
Lactococcus raffinolactis	1
Leuconostoc sp	1
Flavobacterium sp	1
Stenotrophomonas maltophilia	1
Ralstonia sp	0.5
Moraxella osloensis	0.5
Veillonella sp	0.5
Pseudomonas sp	0.5
Acidovorax sp	0.5
Achromobacter sp	0.5
Weissella cibaria	0.5
Kordia sp	0.5



Transfer of infant microbiome to germ free mice

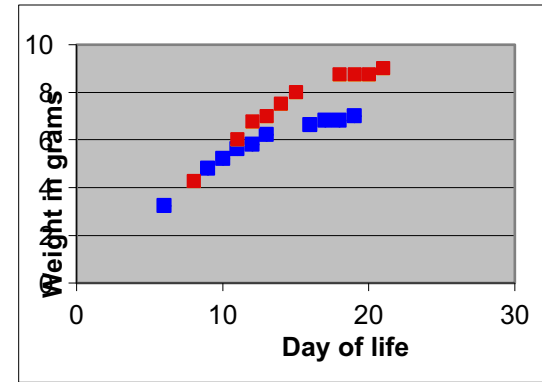
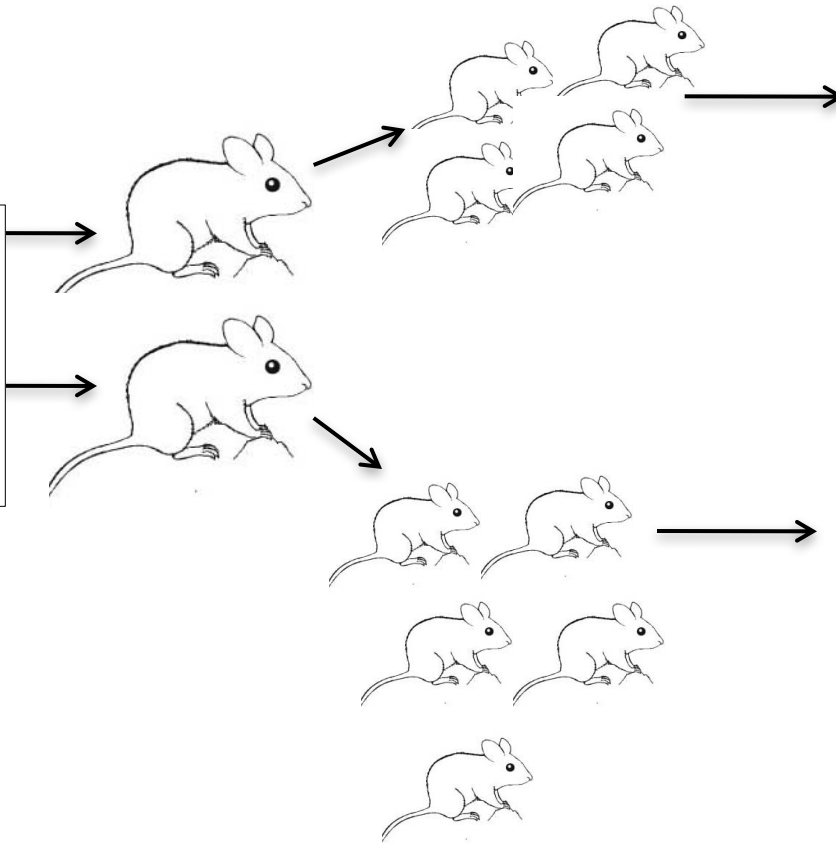
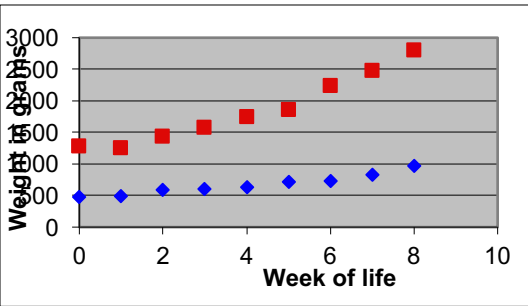
Fecal lysate
from infant

Germ free pregnant
dam C57/Bl6

Litter

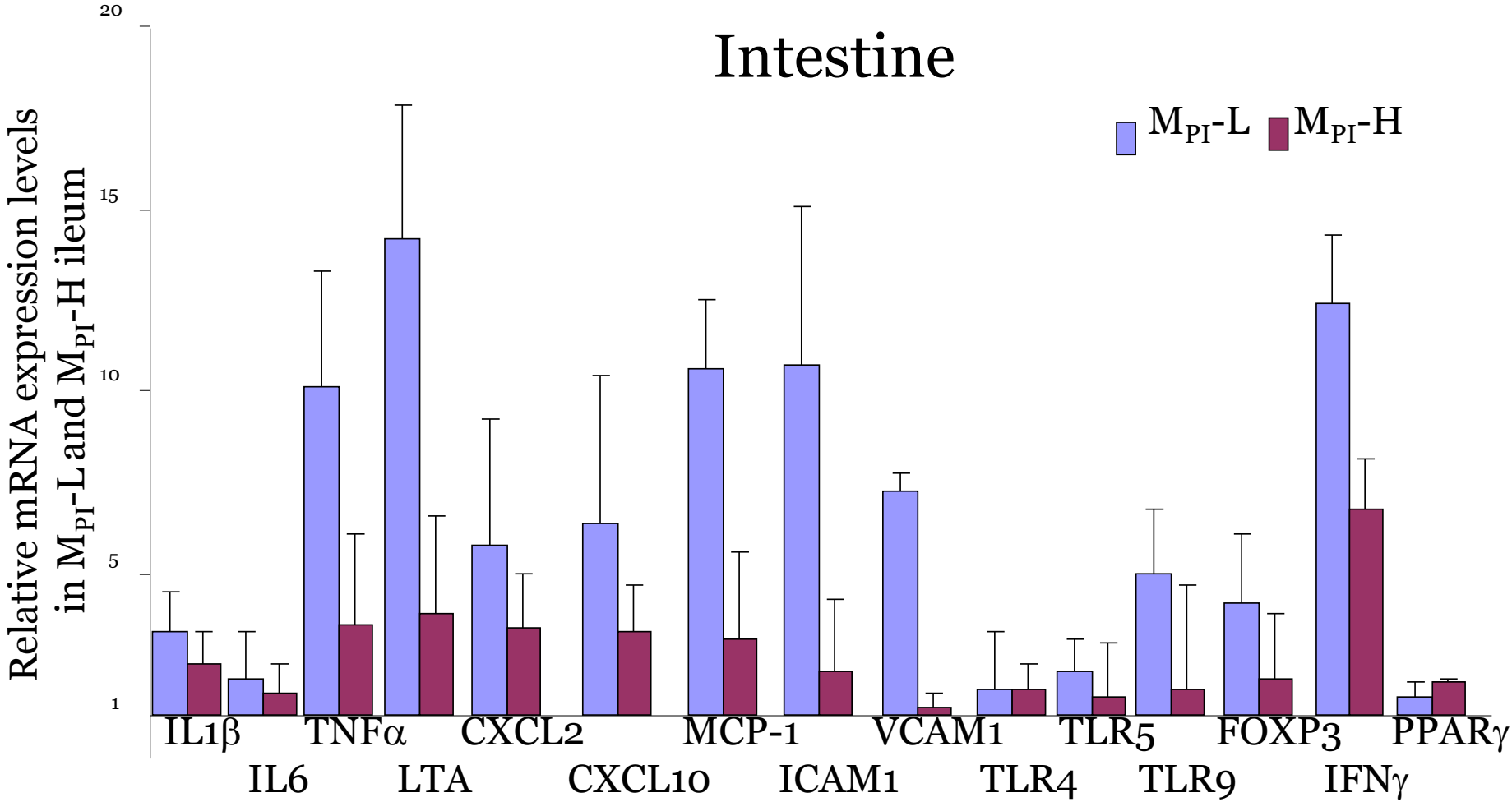
Daily weight

Wean



PCR validation of Microarray

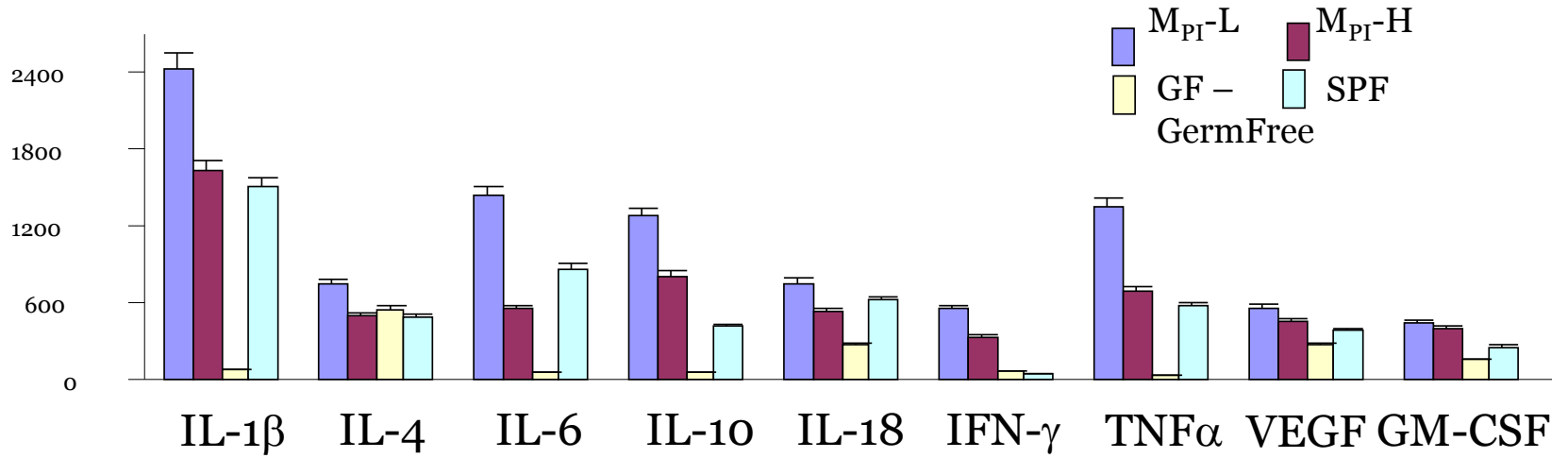
Intestine



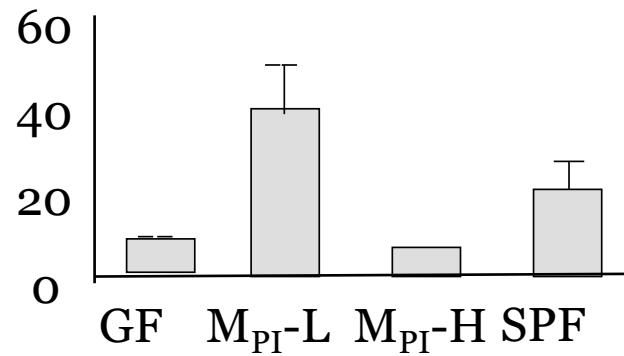
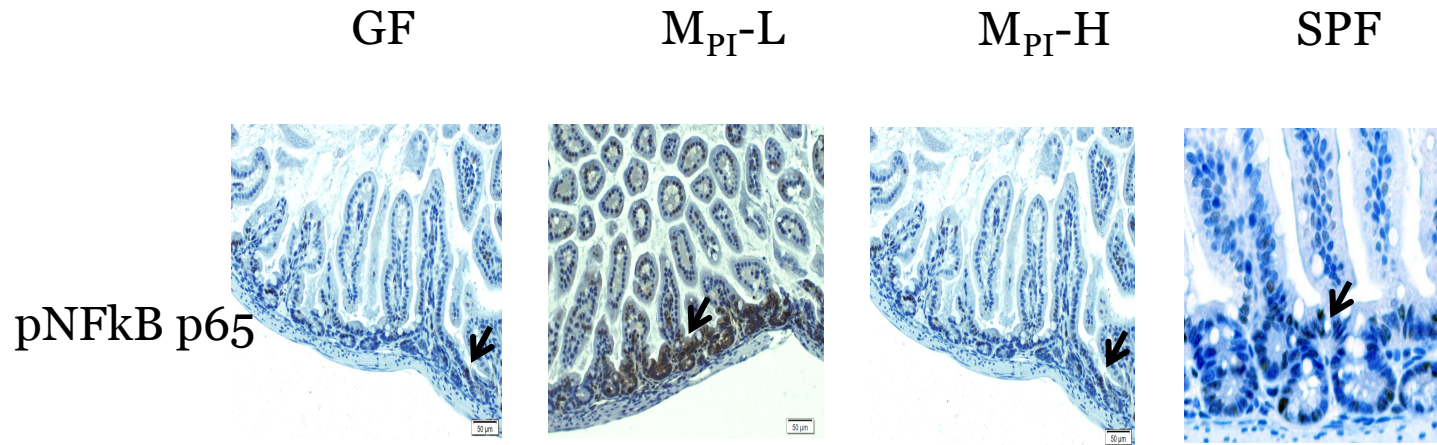
Cytokine Expression

Cytokine unit/ 0.1ml serum

Serum

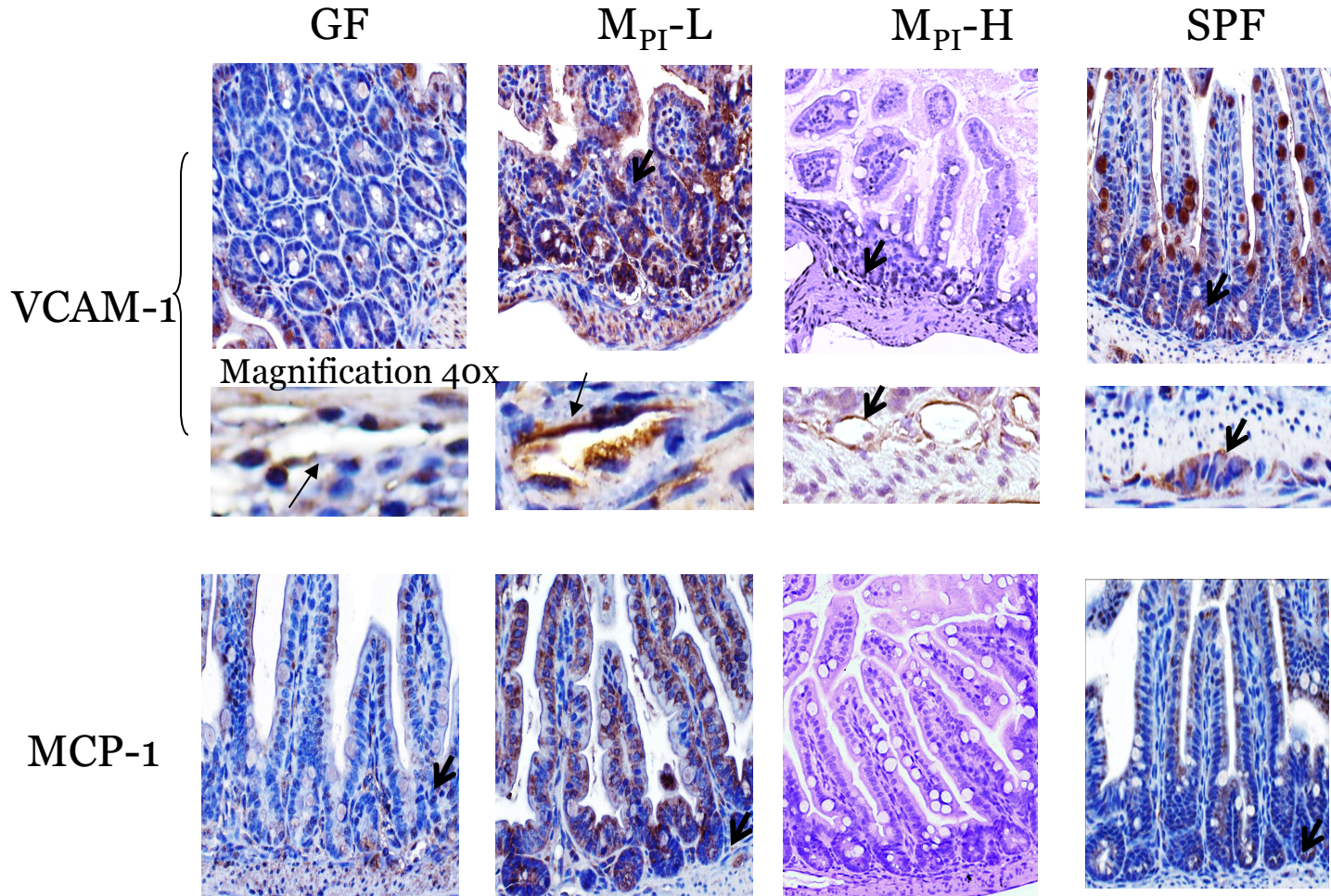


NF- κ B Activation



Labeling index for pNF κ B p65 nuclear translocation

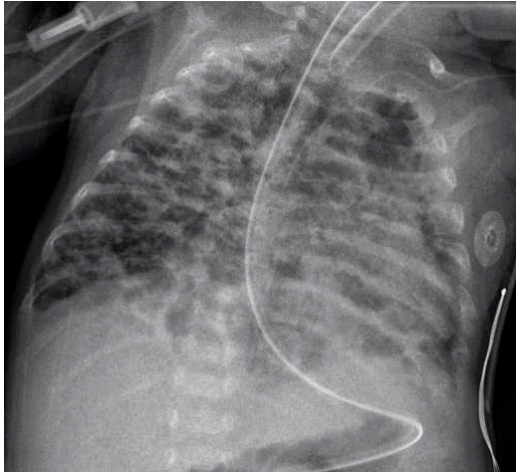
NF- κ B Dependent Cytokines



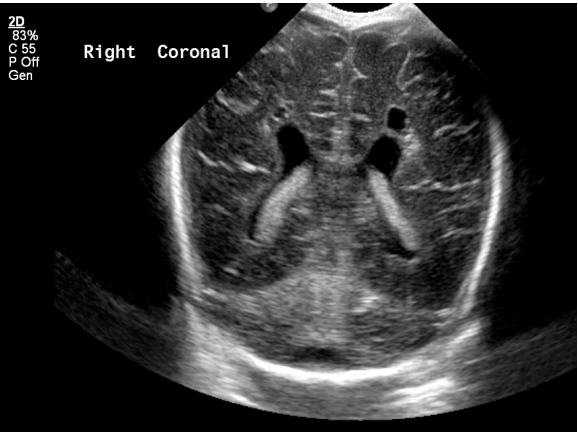
Inflammation and Prematurity



NEC



BPD



PVL

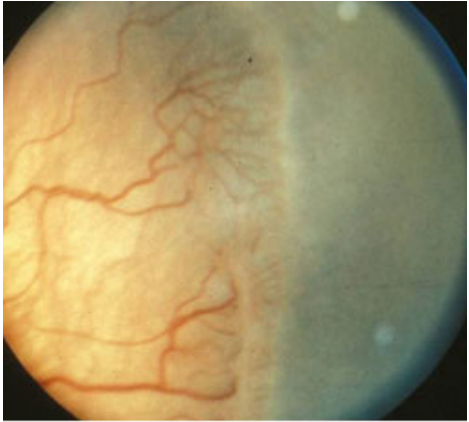


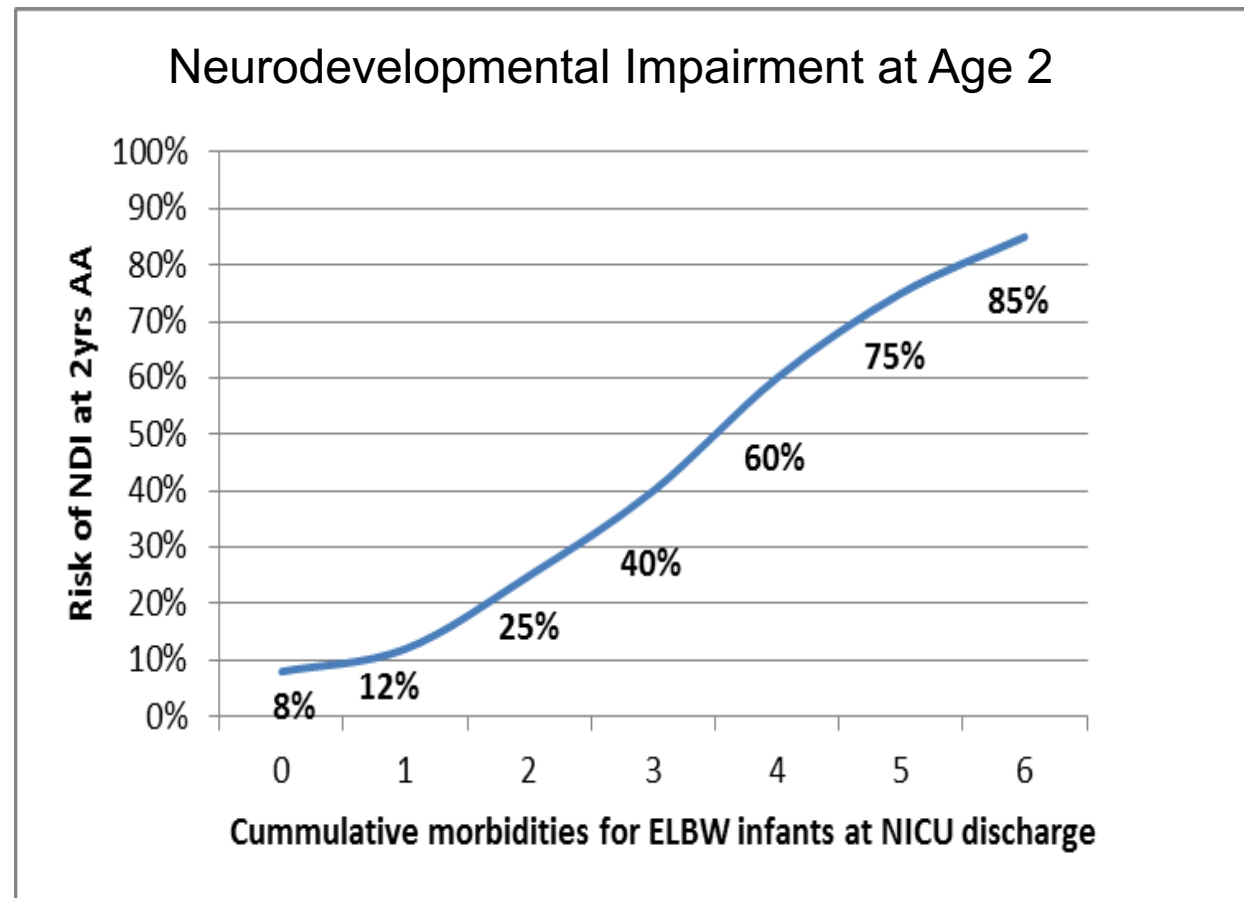
Fig. 3 Stage 3 ROP.

ROP

Neurodevelopmental Outcome in Preterm infants

Morbidities

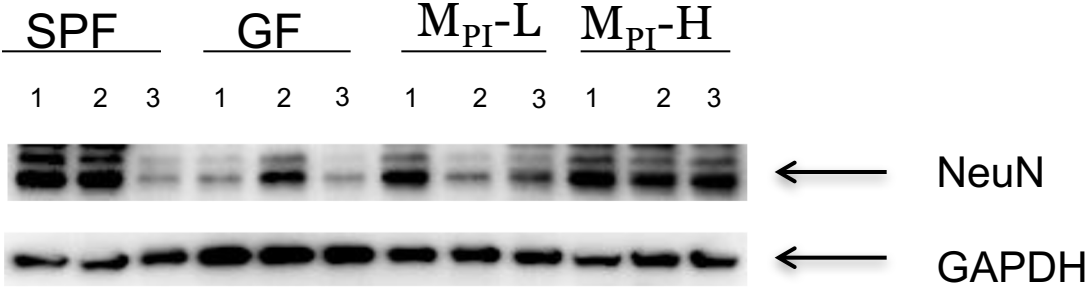
1. Bronchopulmonary dysplasia
2. Necrotizing Enterocolitis
3. Intraventricular Hemorrhage
4. Periventricular Leukomalacia
5. Retinopathy of Prematurity
6. Sepsis



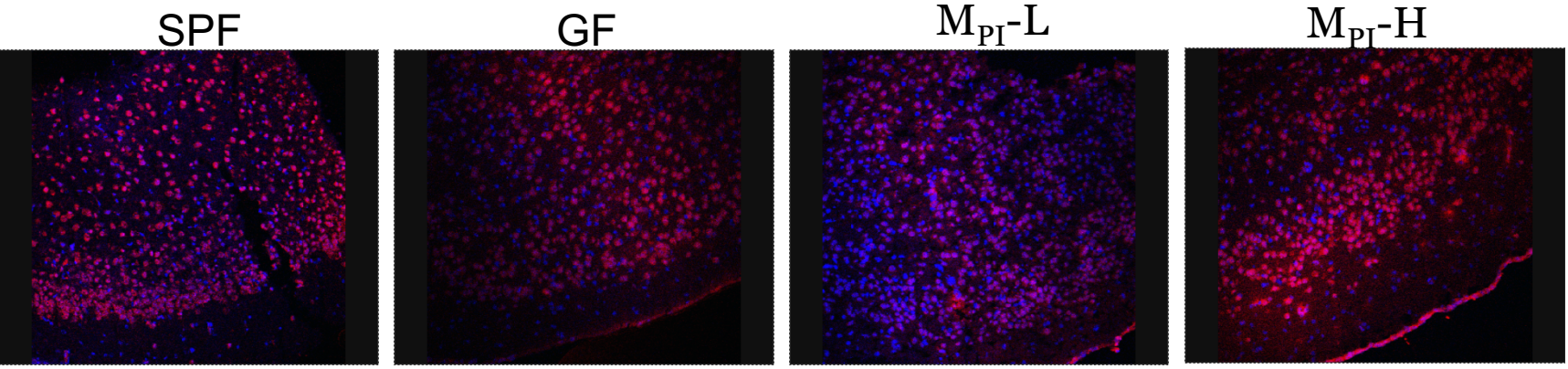
Regulation of cortex neuronal development by gut microbiota.

11/16/17

A.



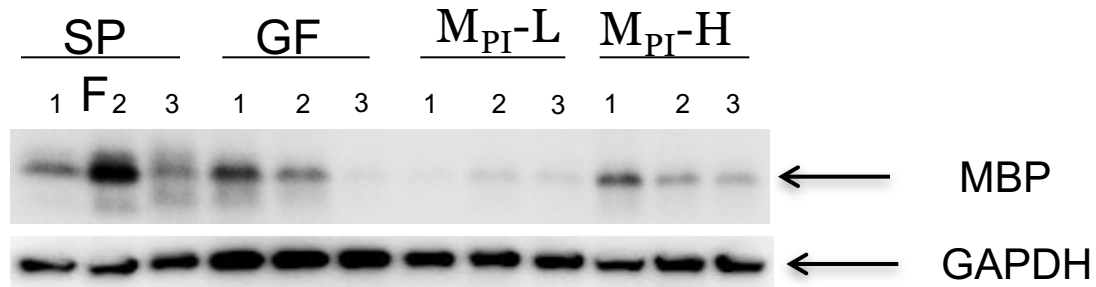
B.



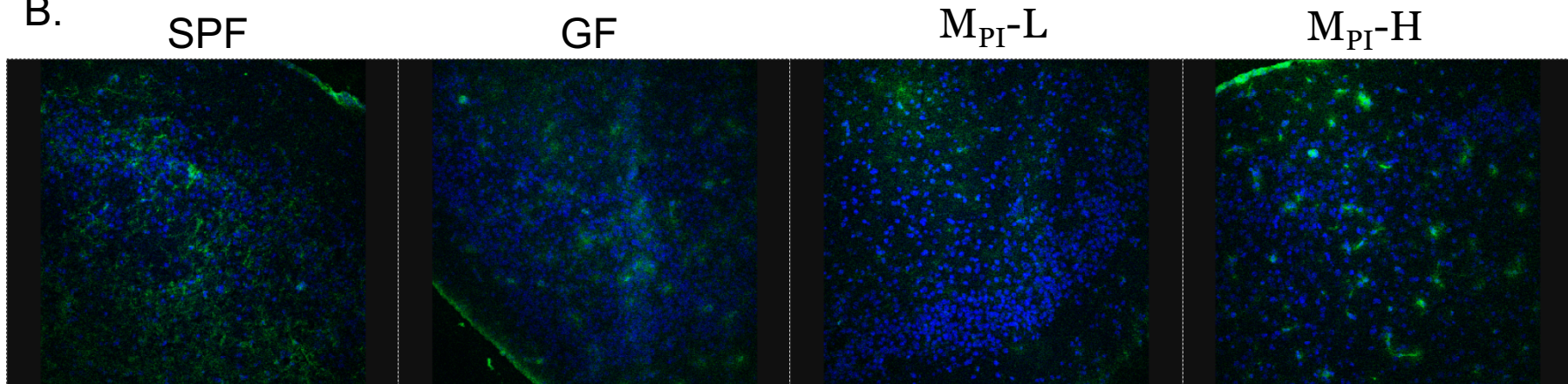
DAPI
NeuN

Regulation of cortex myelination by gut microbiota. 11/16/17

A.



B.



DAPI

MBP

IGF1

- We have established that certain microbiota colonization normalized the growth in GF mice (M2).
- Mutation(s) in the *igf-1* gene or in the *igf1r* gene are found to be associated with severe body growth failure, microcephaly, and developmental delay.
- In rodents, *igf-1* gene disruption results in reduced brain size, CNS hypomyelination and loss of hippocampal granules and striatal parvalbumin-containing neurons.
- GF mice have lower circulating IGF-1 comparing to SPF mice.
- IGF1 crosses the blood brain barrier

Hypothesis

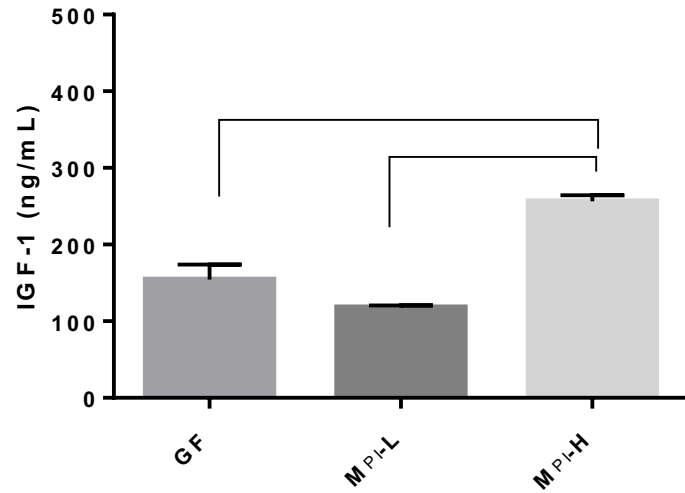
- Microbial colonization can modulate brain development through regulation of IGF1.

Microbiome influences serum IGF-1

11/16/17

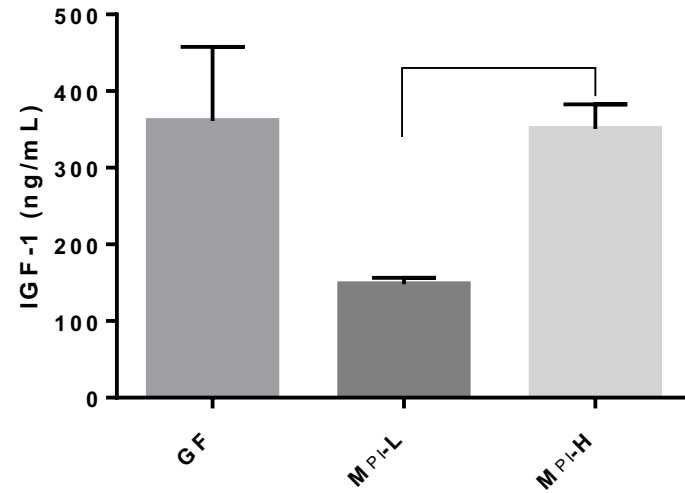
A

Serum IGF-1 2 weeks



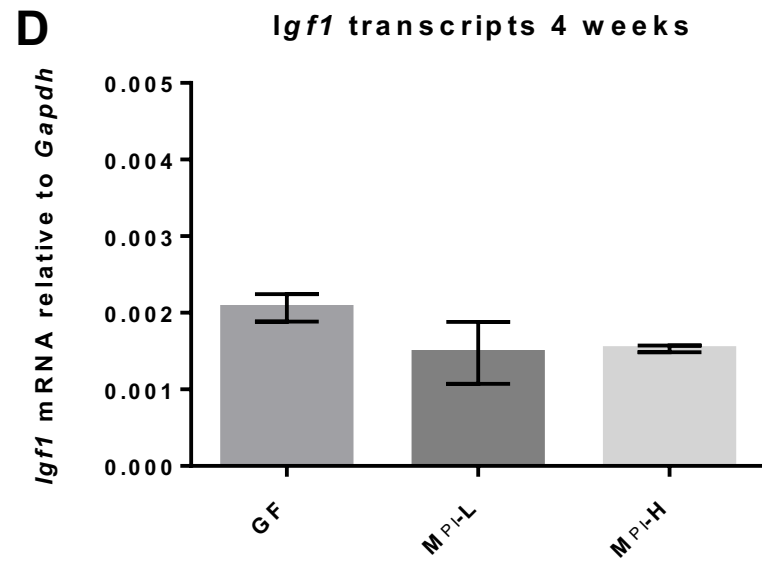
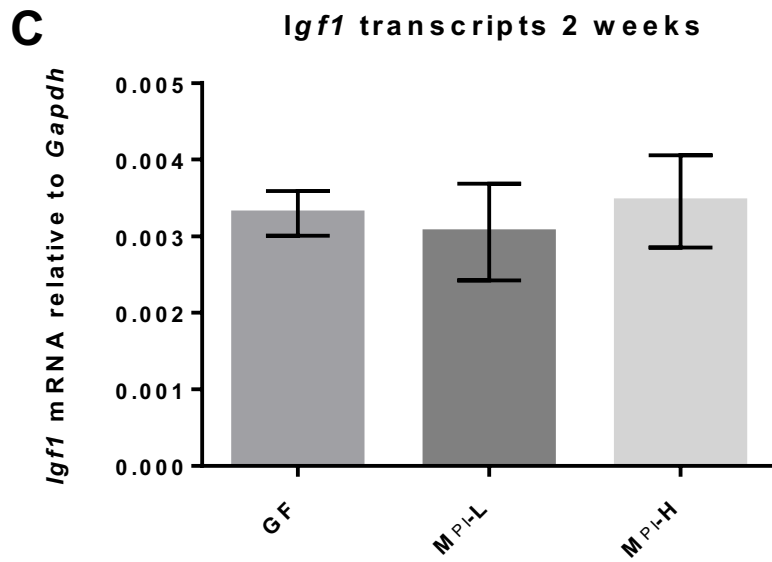
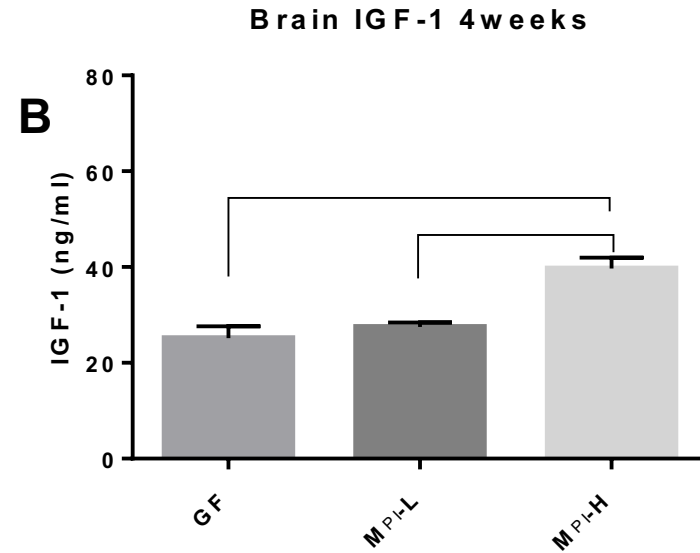
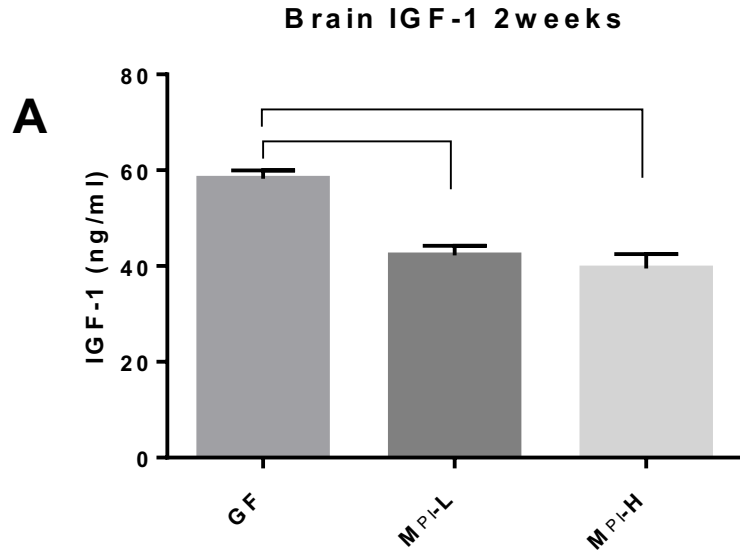
B

Serum IGF-1 4 weeks

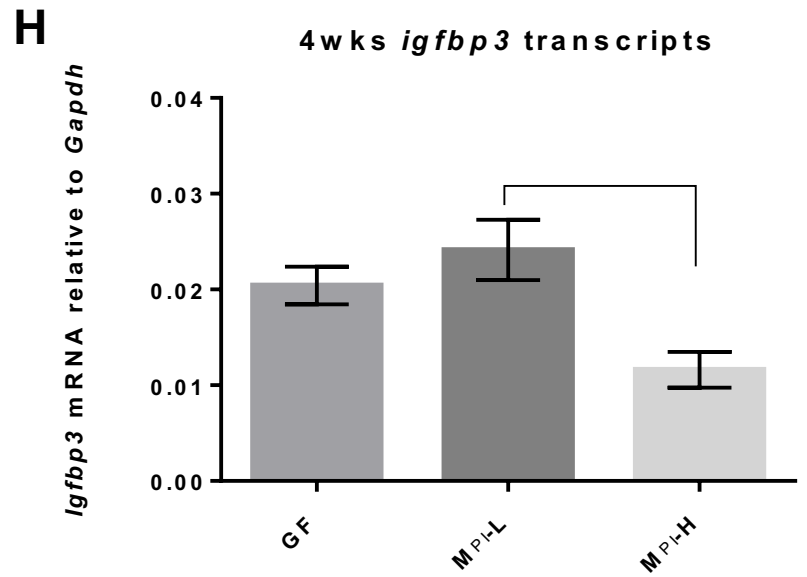
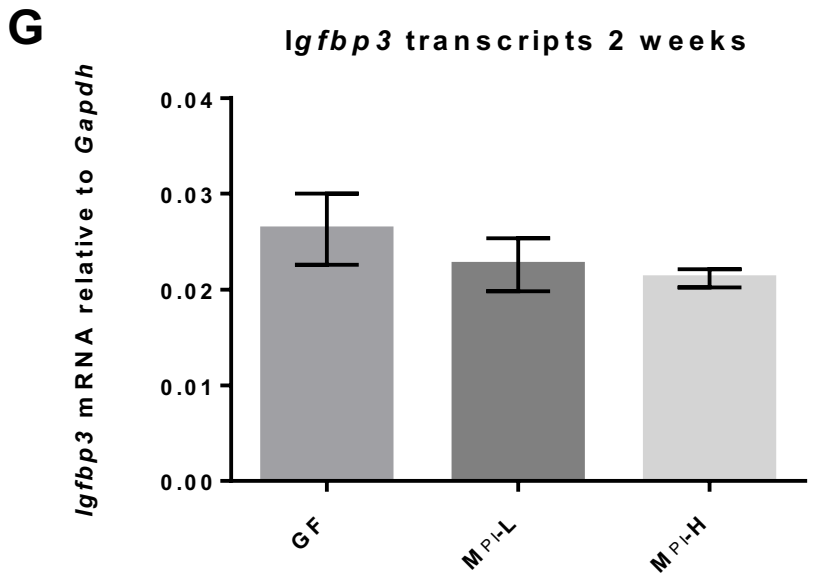
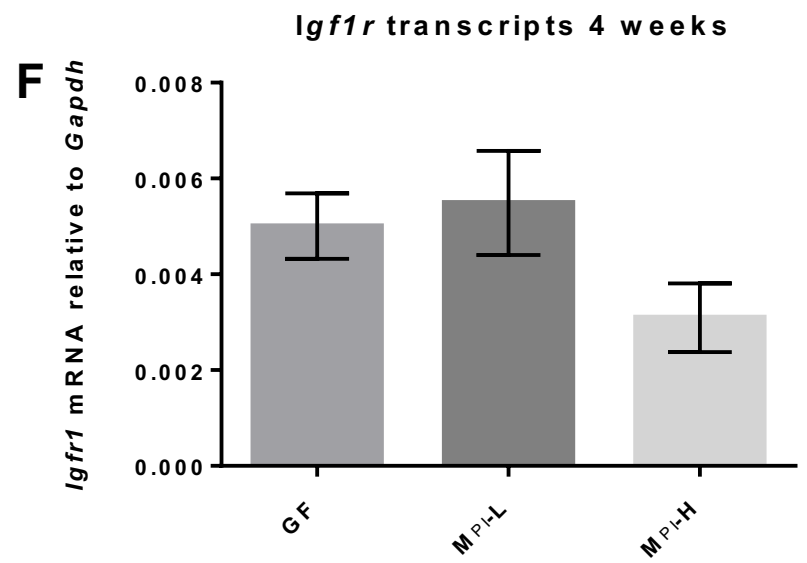
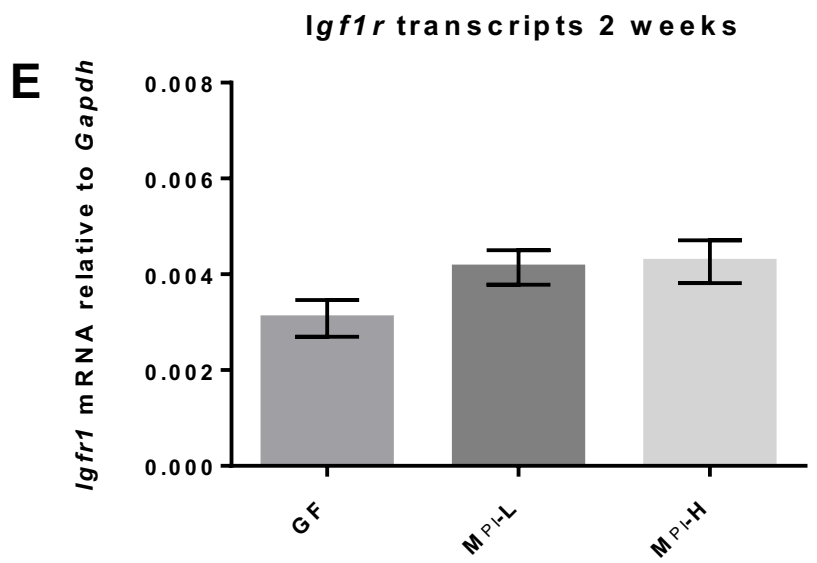


Alterations in brain IGF-1

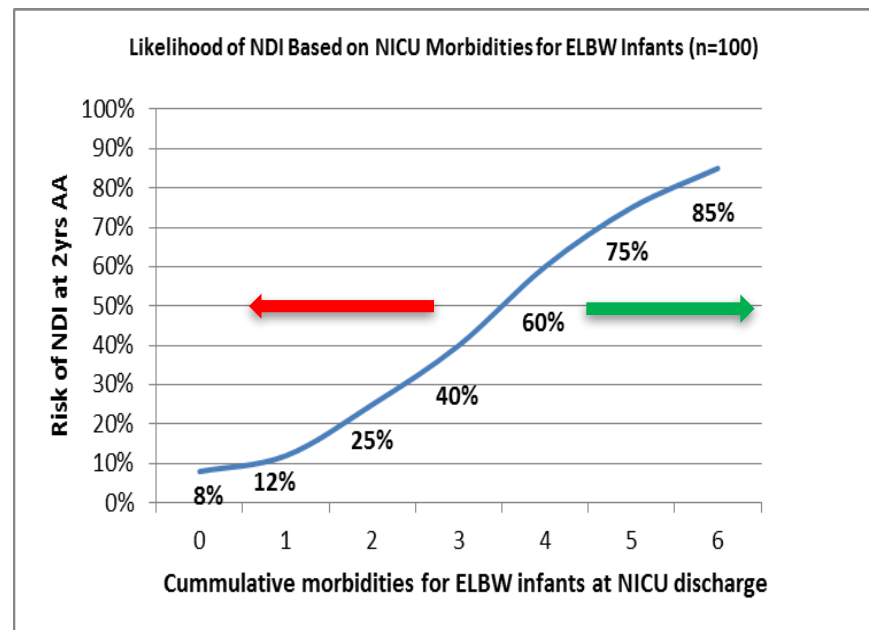
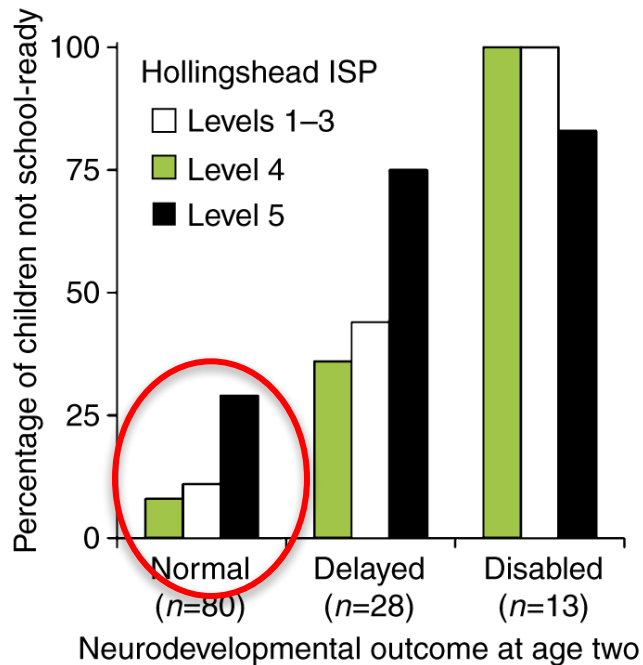
11/16/17



Alterations in brain IGF1r and IGFBP3



Effect of socioeconomic status on Neurodevelopment



Patrianakos-Hoobler et al Dev Med Child Neurol.
2010 Apr;52(4):379-85

Microbiome?

ENVIRONMENTAL INFLUENCES ON CHILD HEALTH OUTCOMES (ECHO) PROGRAM

Environmental influences on Child Health Outcomes (ECHO) Program

[Director's Page](#)

[Planning](#)

[Governance](#)

[Program Components](#)

[Funding](#)

[Announcements](#)



NIH officially launches the ECHO program with more than \$150 million in awards



Learn about the new ECHO program components and awardees

About the ECHO Program

The means by which poverty alters neurodevelopment are unknown.

The microbiome is influenced by environment and in turn influences brain development.

We hypothesize that the microbiome is a biologic effector of the influence of SES and environment on neurodevelopment.

Related Information

ECHO
Environmental Influences on Child Health Outcomes

NIH plans to support multiple studies using existing cohorts groups of diverse U.S. children who have already participated in other research studies to answer questions about the effects of a broad range of environmental factors on child health & development. This approach will allow the field to combine data and maximize the use of existing research on its diverse questions that could not be addressed by each study alone.

The basics

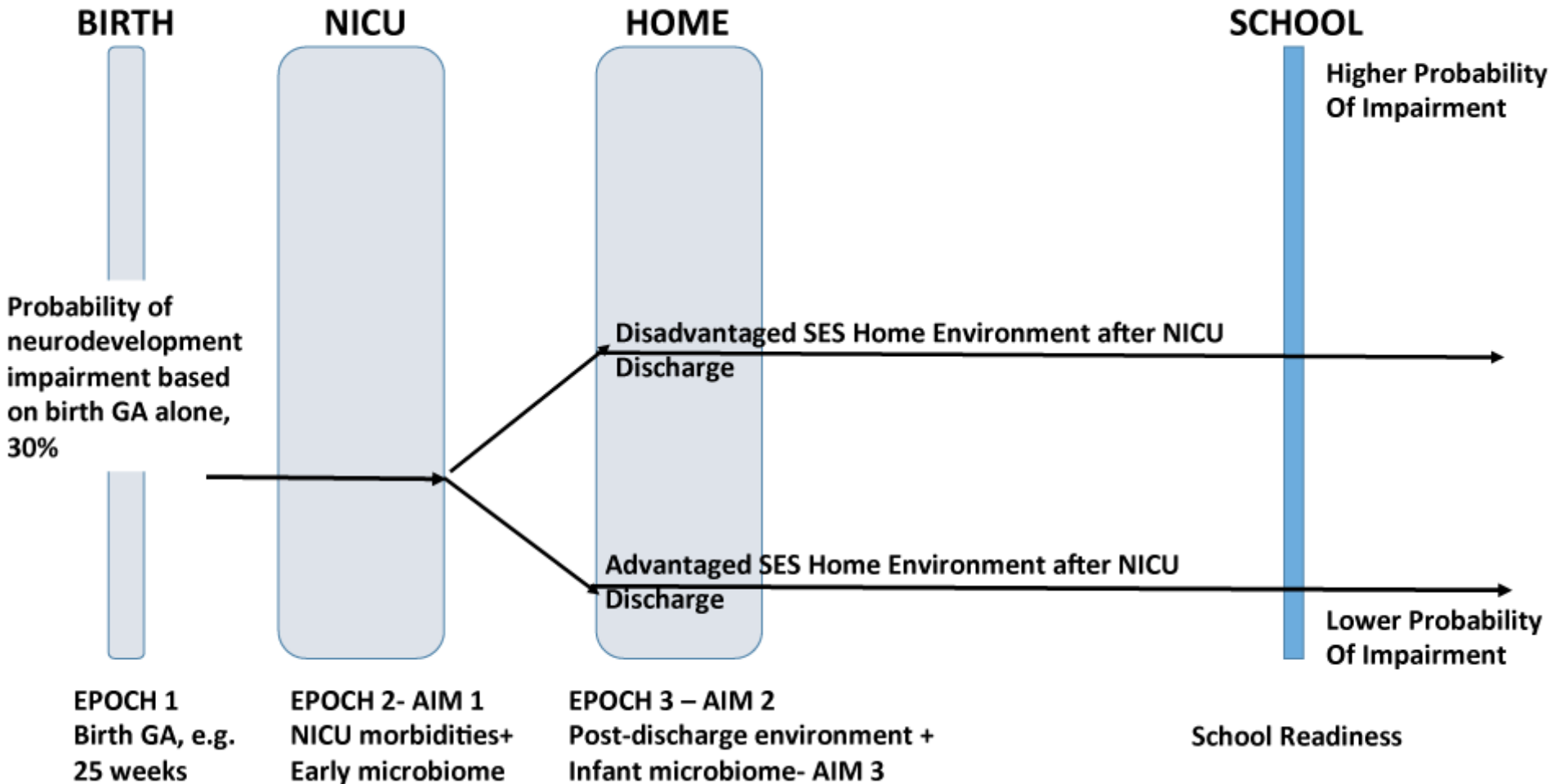
The ECHO program will allow researchers to maximize the use of existing resources such as collections of biological tissues collected during pregnancy and delivery, leverage existing datasets by linking to additional datasets, develop a repository on the trajectory of health, development, and genetic/clinical health to predict disease development, and test new tools and approaches for environmental and pediatric monitoring.

What we hope to learn

In addition to each investigator's specific research questions, all of the ECHO studies will be expected to collect standardized information (Core Elements on):

- Genetic health and development
- Genetic influences on child health and development
- Environmental factors
- Patient/parent-reported outcomes (PROs)

Influences on Preterm Infant Neurodevelopment Potential From Birth To School Age



Impact of Microbiome is Modifiable



Unborn



Baby



Toddler



Adult



Elderly



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