

Measurement of intestinal fatty-acid binding protein and intestinal barrier function biomarkers in DBS using ELISA

Elizabeth Kim, M.S.



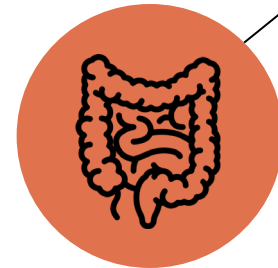
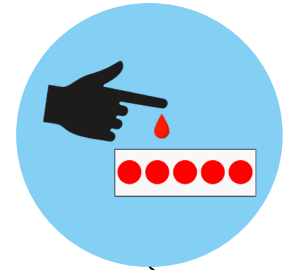
BAYLOR
UNIVERSITY



HUMAN EVOLUTIONARY
BIOLOGY & HEALTH LAB



National Institutes of Health



Human Evolutionary Biology and Health Lab

Baylor University



- Investigate bio-energetic pathways shaping variation in human development, metabolism, and health.
- Combine diverse field and laboratory approaches, focusing on participant-friendly measurements and the use of **minimally invasive biomarkers**.



DBS



Hair



Saliva



Fecal



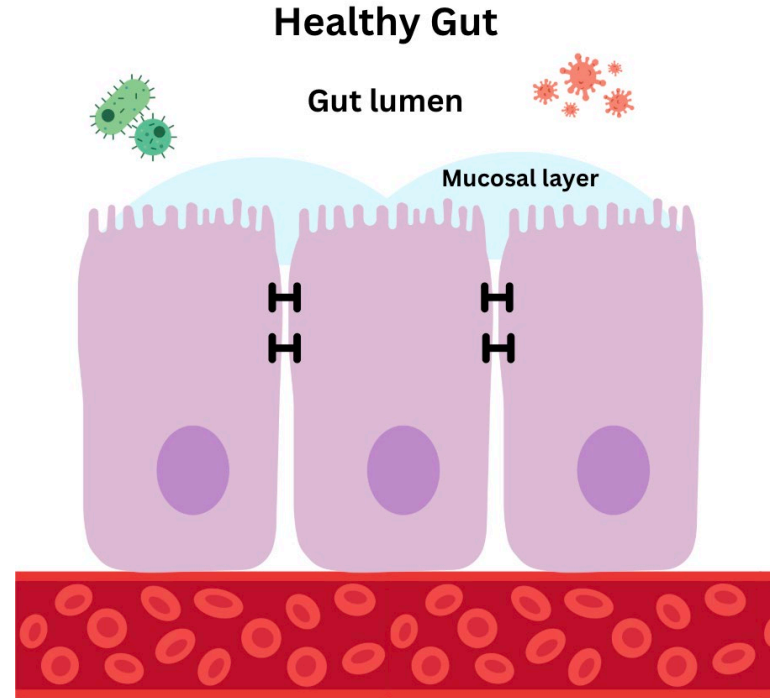
Urine

Dr. Samuel
Urlacher
(PI)

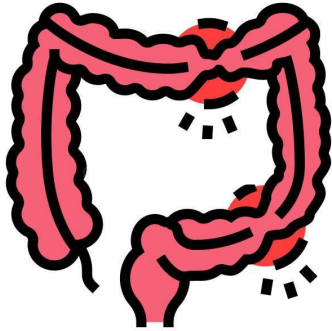


Intestinal function is critical for regulating homeostasis and health

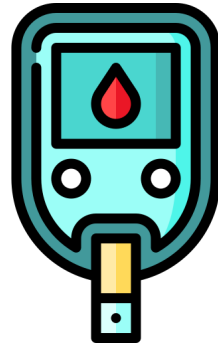
- Nutrient digestion and absorption
- Barrier protection from the external environment
- Immune regulation
- Host-microbe interactions



Intestinal function is critical for regulating homeostasis and health



Irritable bowel syndrome,
Crohn's disease



Type I diabetes

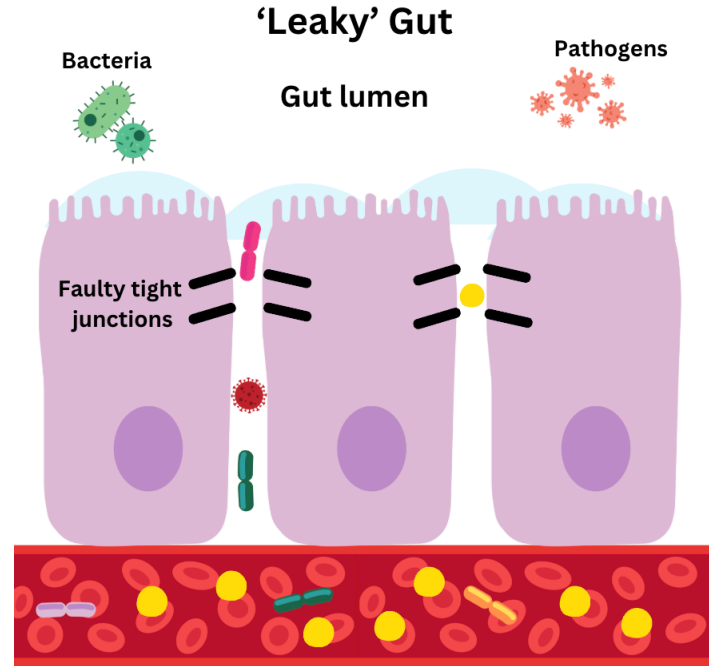


Non-alcoholic fatty liver
disease

Environmental and lifestyle factors shape variation in intestinal function and health

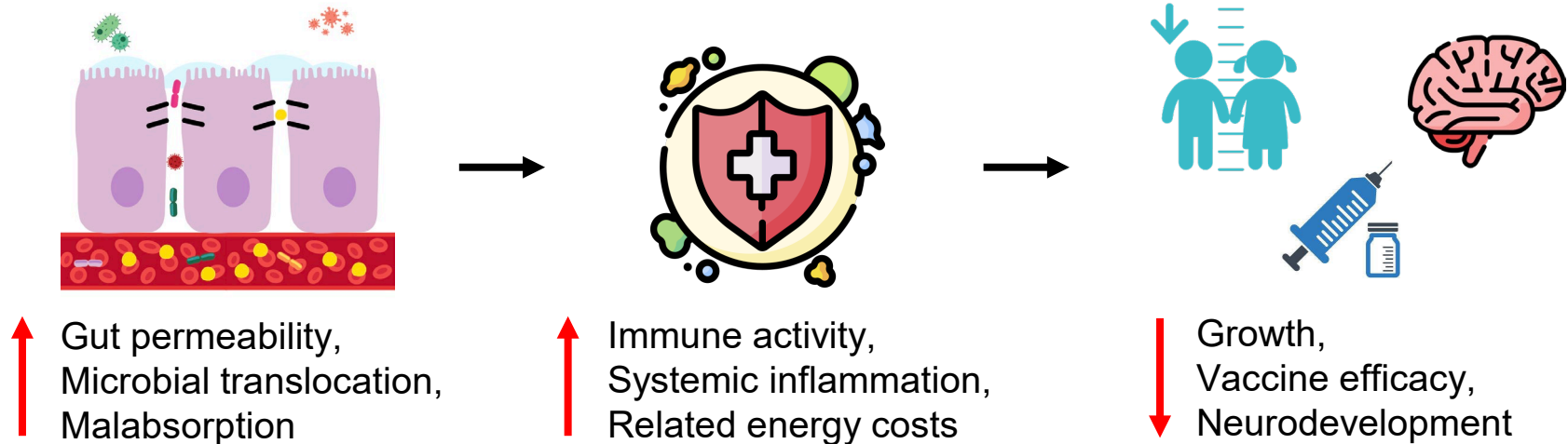
- Systemic inflammation
- Microbial translocation
- Malabsorption

Environmental enteric dysfunction (EED)



Environmental Enteric Dysfunction (EED)

- Acquired subclinical condition of small intestine
- Pathogen exposure from unsanitary living conditions
- Nearly ubiquitous in low-resource and disadvantaged contexts



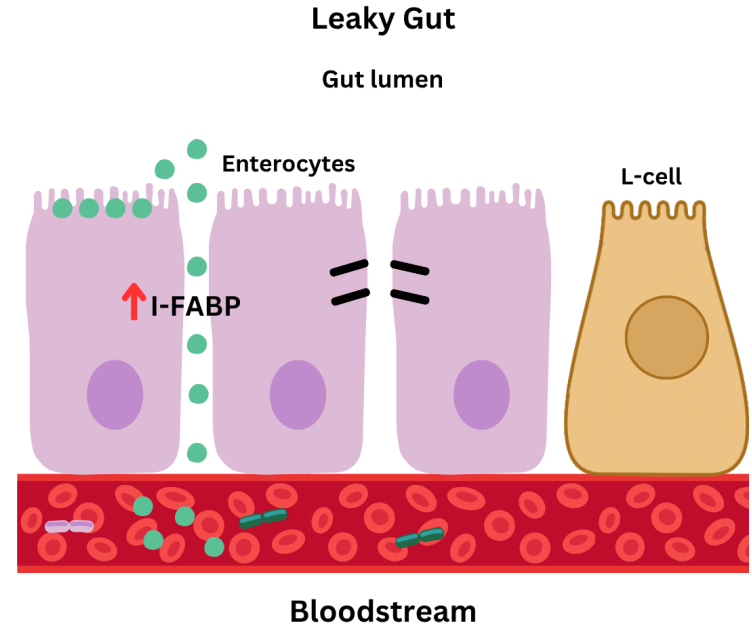
**Standardized and objective measures
for EED assessment are limited
and rely on **invasive sampling methods****

Objective: validate EED biomarkers in finger-prick dried blood spots (DBS) via ELISA

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1) Intestinal fatty-acid binding protein (I-FABP)

- Cytosolic protein, transports long-chain fatty acids
- ↑ circulating I-FABP → ↑ intestinal barrier damage



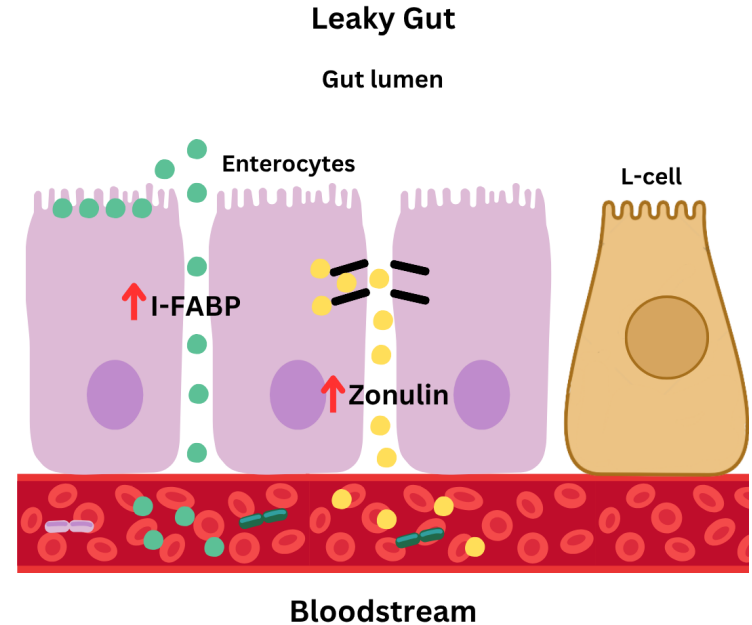
Objective: validate EED biomarkers in finger-prick dried blood spots (DBS) via ELISA

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2) Zonulin

- Regulator of tight junctions
- ↑ circulating zonulin → ↑ intestinal permeability



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1) Intestinal fatty-acid binding protein (I-FABP)

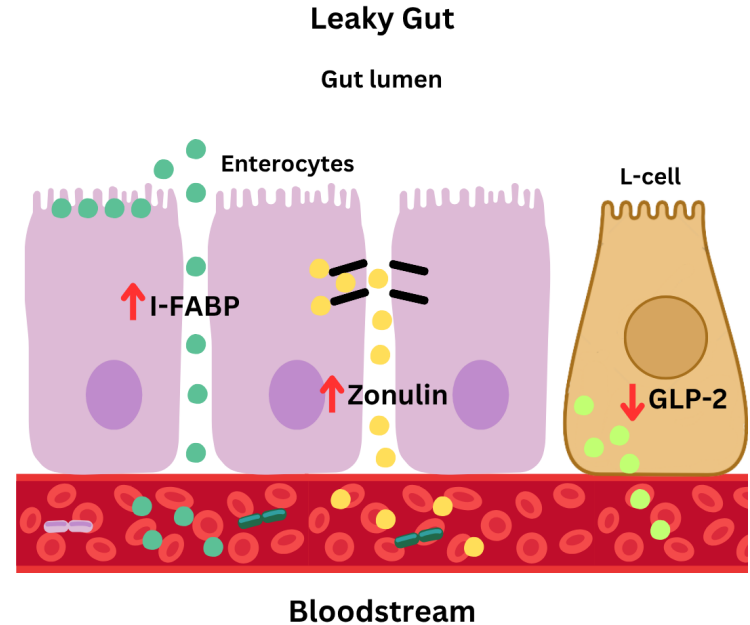
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2) Zonulin

- Regulator of tight junctions
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3) Glucagon-like peptide 2 (GLP-2)

- Repair mucosa and promote nutrient absorption
- ↓ circulating GLP-2 → ↑ intestinal barrier damage, ↓ nutrient absorption



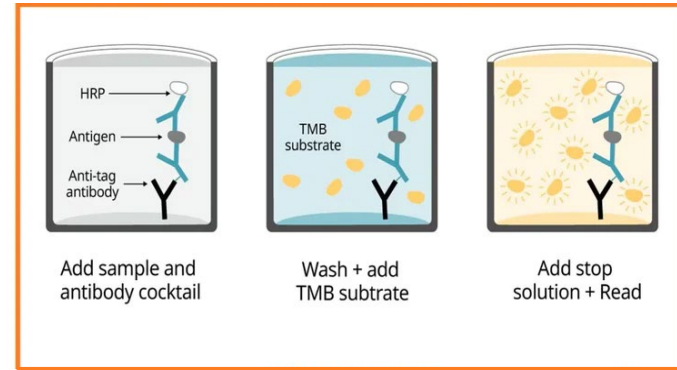
Samples: 'Waco100' matched DBS and serum

- Matched finger-prick DBS and serum samples
- Convenience sample of **117 adults** residing in McLennan County, Waco, Texas
- Standard collection procedures
 - Protein saver filter paper cards (Whatman #903)
 - Stored at -80°C until analysis
- Sample collection as part of the Waco COVID Survey



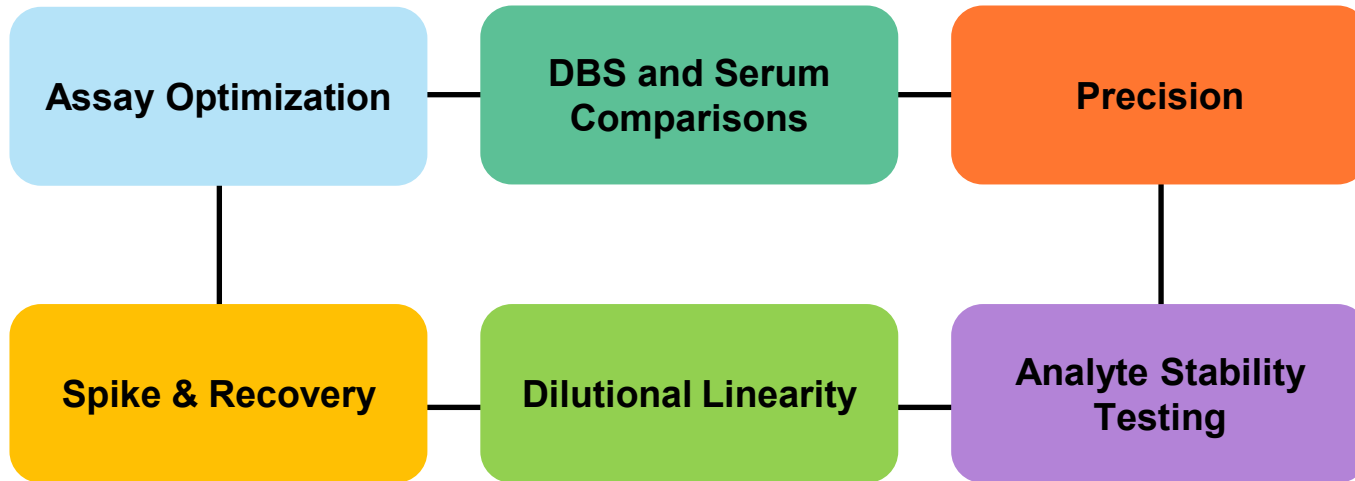
Kit selection: Abcam Human FABP2 ELISA (ab234566)

- Abcam Human FABP2 ELISA Kit (ab234566)
 - Quantitative calorimetric sandwich ELISA
 - Validated for urine, serum, plasma, and cell culture samples
- Assay range: 12.5 pg/mL to 800 pg/mL
- Low sensitivity: 1.7 pg/mL
- Relatively minimal sample requirement (50uL/well)
- Short total incubation time: 70 minutes



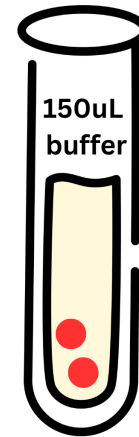
Validation Testing and Methods

- Following established protocols developed by McDade



Assay Optimization

- Determine ideal DBS sample elution parameters
- DBS elution strategy: **2 small (3.2mm²) punches eluted in 150uL** of Sample Diluent NS
 - Eluted overnight for 12 hours at 4°C
- Serum samples diluted 1:19 with Sample Diluent NS



DBS and Serum Comparisons

I-FABP was quantified in all DBS samples ($n = 117$)

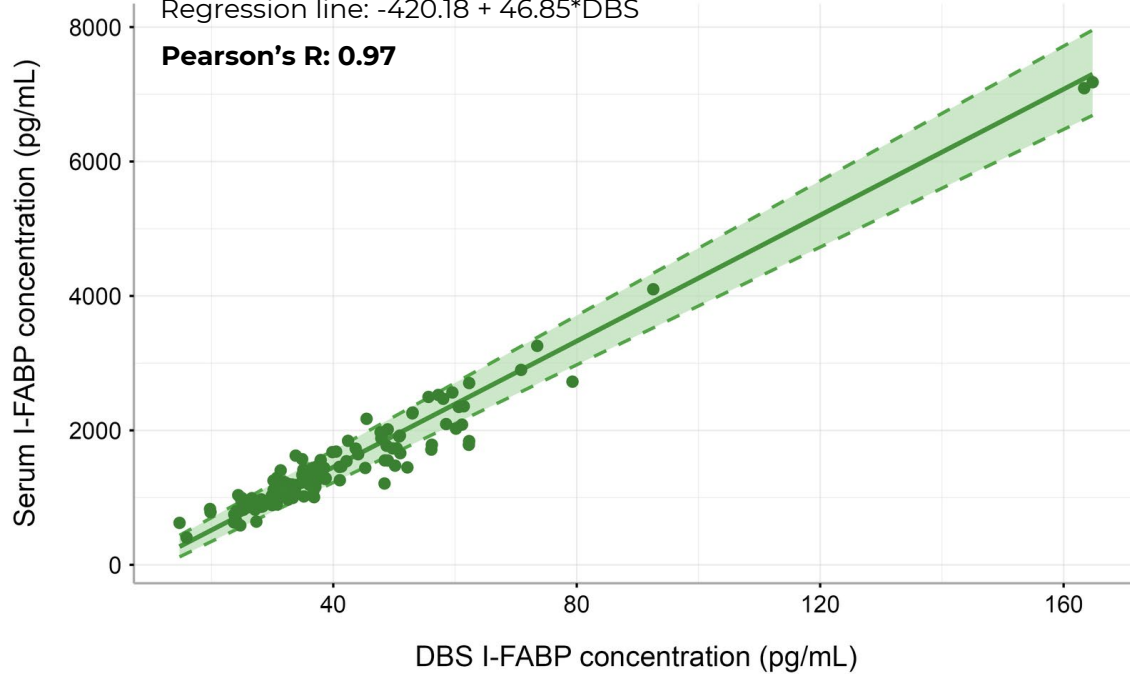
Mean I-FABP in DBS:
41.97pg/mL \pm 21.2 (SD)

Strong linear relationship between DBS and serum I-FABP
(Pearson's R: 0.97)

Passing-Bablok Regression Plot

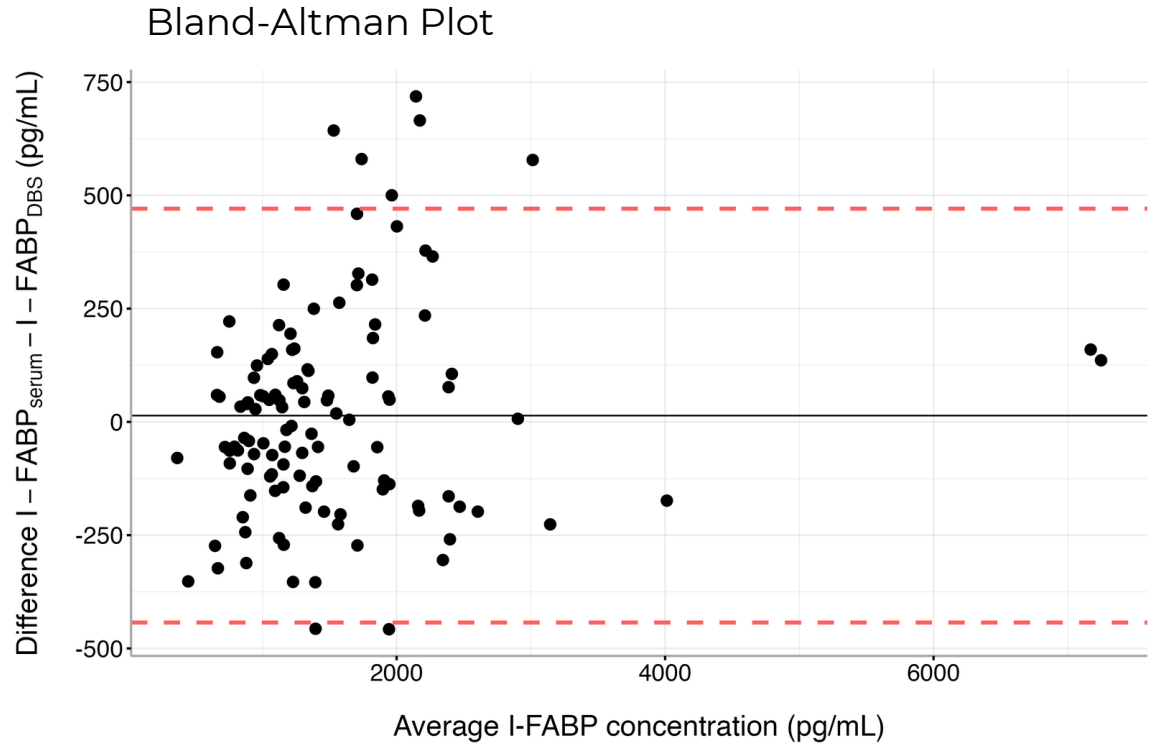
Regression line: $-420.18 + 46.85 \cdot \text{DBS}$

Pearson's R: 0.97



DBS and Serum Comparisons

Minimal bias observed
between serum
I-FABP and DBS I-FABP
levels (**bias = 13.9**)



Precision

Intra-assay CV

Average CV for duplicate measures of DBS samples

4.0 %

Inter-assay CV

Average CV for DBS control included on each plate

3.9 %

Limit of Detection

Calculated as the concentration of the mean OD value for all blank wells + 2 SDs

5.27 pg/mL

Spike and Recovery

Average % recovery: **101.8%**

Low Spike

+208 pg/mL
I-FABP

101.9%
recovery

Med Spike

+385 pg/mL
I-FABP

100.5%
recovery

High Spike

+536 pg/mL
I-FABP

102.8%
recovery

Dilutional Linearity

Average % recovery: **111.2%**

1:2 Dilution

106.8%
recovery

1:4 Dilution

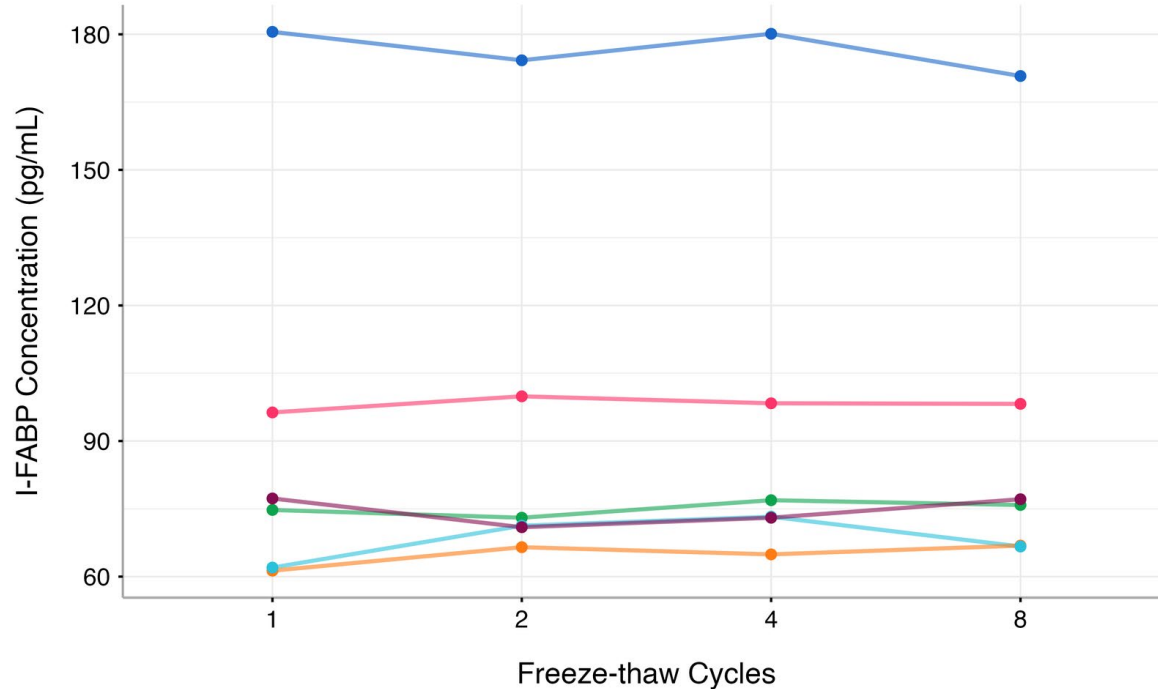
118%
recovery

1:8 Dilution

108.9%
recovery

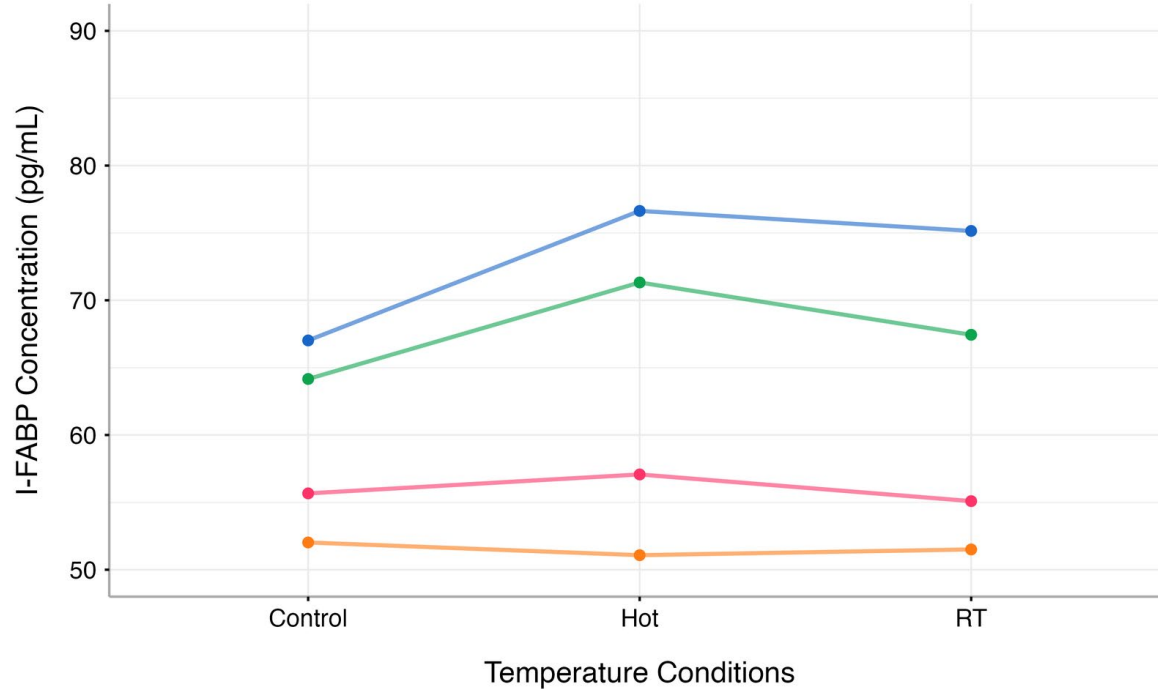
Analyte Stability

No significant differences in I-FABP concentrations across 1, 2, 4, and 8 F/T cycles ($p > 0.05$)



Analyte Stability

No significant differences in I-FABP concentrations across control, hot, and RT conditions ($p > 0.05$) for at least 72 hours



I-FABP can be reliably quantified in DBS using ELISA

Abcam Human FABP2 ELISA Kit (ab234566)

Assay Optimization

DBS elution:
2 small (3.2mm²)
punches in 150uL diluent

DBS and Serum Comparisons

Strong linear relationship
(Pearson's R: 0.97)
No obvious bias
(bias = 13.9)

Precision

Intra-assay CV: **4.0%**
Inter-assay CV: **3.9%**
LOD: **5.27pg/mL**

Spike and Recovery

Average percent
recovery:
101.8%

Dilutional Linearity

Average percent
recovery:
111.2%

Analyte Stability

I-FABP **stable** for up to
8 F/T cycles and at
least **72 hours** at **RT**
and **hot** conditions
($p > 0.05$).

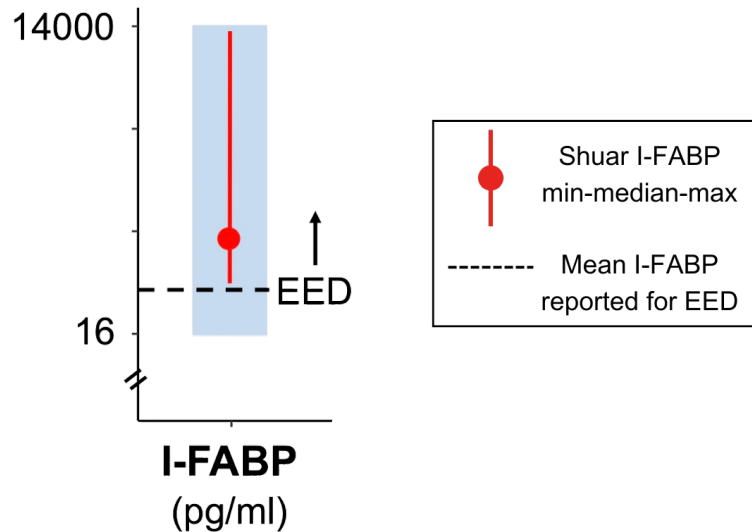
DBS I-FABP measurement in Amazonian Shuar Children

- Indigenous population living in Amazonian Ecuador
- Experience varying rates of market integration
- High immune activity and pathogen burden
- Intestinal inflammation and parasites
- 41% of Shuar children are stunted

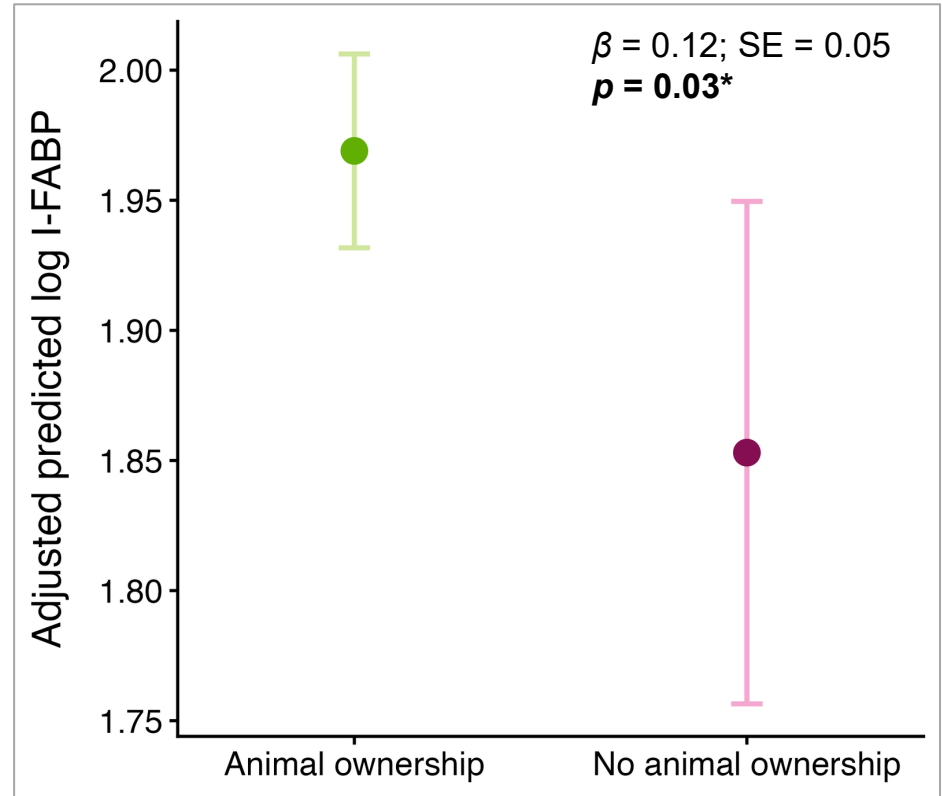
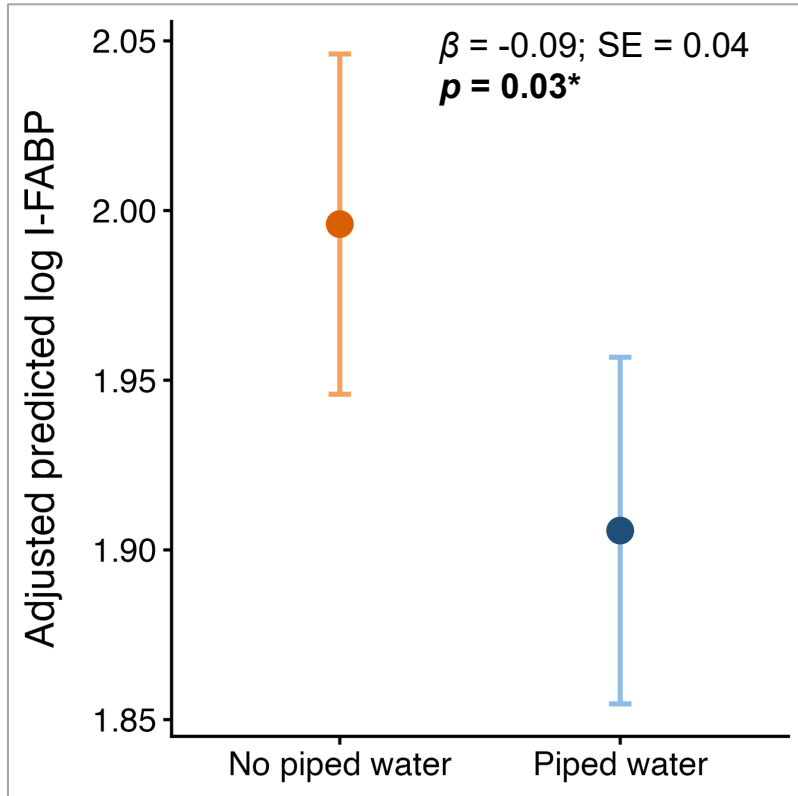


DBS I-FABP measurement in Amazonian Shuar Children

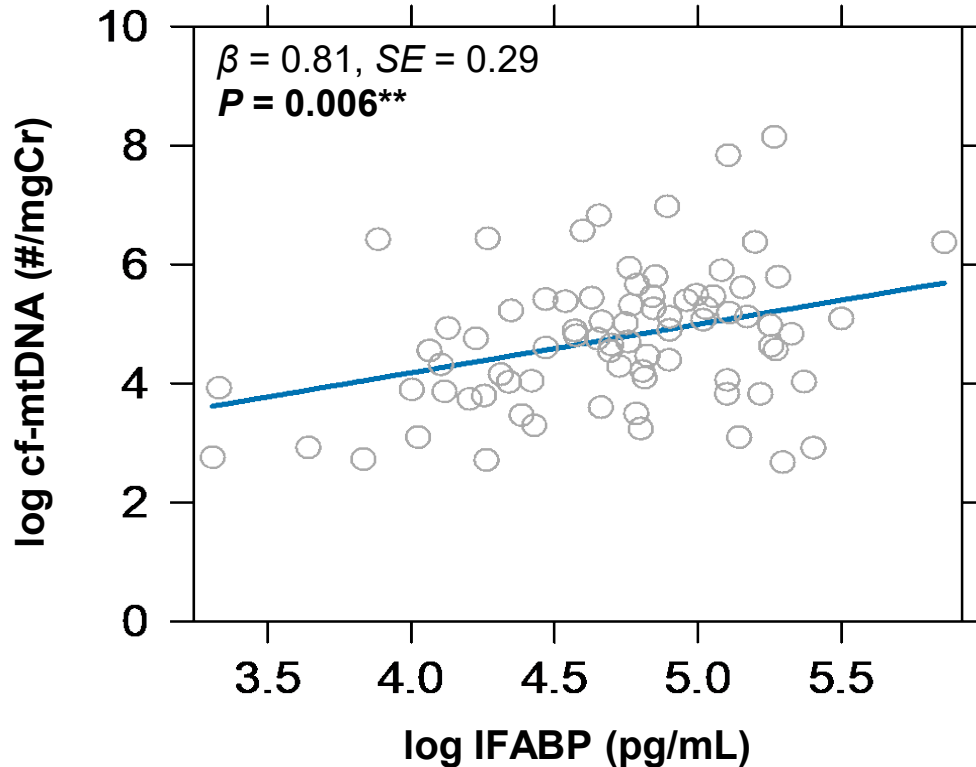
- 312 Shuar children, ages 4 – 12 years old
- Demonstrate **high levels of I-FABP** indicative of **EED**



Shuar children's IFABP levels are significantly associated with household living conditions



Shuar children's IFABP levels are significantly associated with cell-free mitochondrial DNA



Thank you!

Questions?

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We thank the Waco COVID participants and Shuar families for their participation.



HEBHL Team

