



Blood Phenylalanine Measurement in Phenylketonuria (PKU): A Comparison Between Plasma (PAA) vs. Dried Blood Spot (DBS)

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Background

- Phenylalanine (Phe) monitoring is essential for managing phenylketonuria (PKU)
- Phe levels can be measured using plasma or dried blood spot (DBS) samples
- Prior studies report inconsistent differences between specimen types
- Variability depends on methodology and laboratory techniques

Objectives

- Compare plasma and DBS phenylalanine levels using same methodology LC-MS/MS
- Evaluate agreement and systematic bias between methods
- Assess implications for clinical interpretation

Methods

Study Design

- Systematic literature review for meta-analysis of different lab methods – Ion exchange chromatography (IEC), tandem mass spectrometry (MS/MS), liquid chromatography tandem mass spectrometry (LC-MS/MS), high-performance liquid chromatography tandem mass spectrometry (HPLC-MS/MS), amino acid analyzer (AAA) – and different specimen types – plasma, dried blood spot from capillary (DBSC) and from vein (DBSV).
- Comparative analysis of plasma vs dried blood spot (DBS) samples

Measurement:

- Phenylalanine quantified using LC-MS/MS

Sample Collection

- Paired plasma and DBS samples collected at Emory MNT4P Metabolic Camp 2024
- Samples analyzed under the same conditions

Statistical Analysis

- Bland–Altman analysis → bias & agreement
- Concordance correlation coefficient (CCC)
- Regression models for bias correction

Total campers with PKU who attended Emory Metabolic Camp 2024
(N = 31)

Campers with pre- and post-camp plasma DBS available
(N = 18)

Campers with complete/valid pre- and post-camp plasma and DBS
(N = 17)

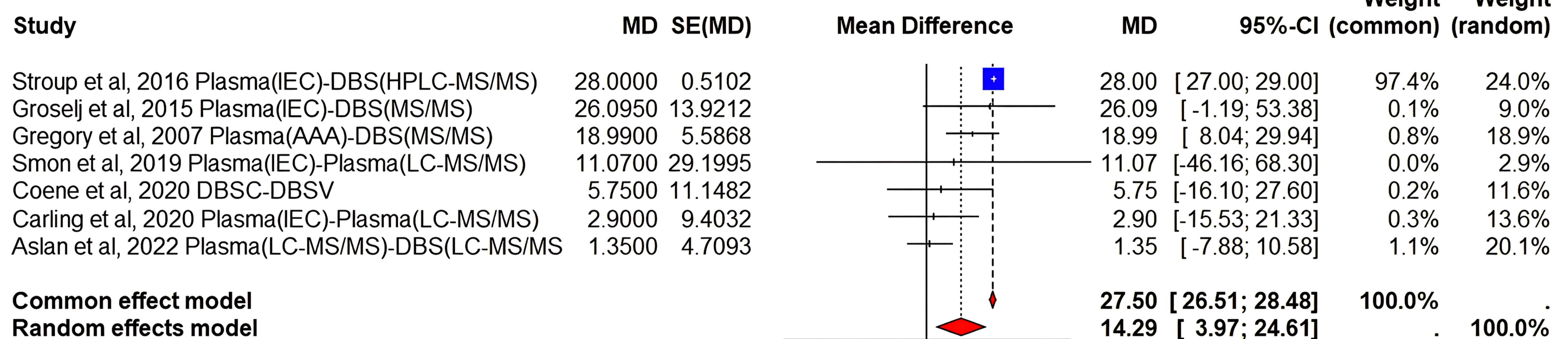
Pre- and Post-camp data were combined/pooled

Total sample size for plasma (N = 34) and DBS (N = 34)

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Results

Figure 1. Meta-analysis of Plasma vs DBS Phenylalanine Differences



Heterogeneity: $I^2 = 86.7\%$, $\tau^2 = 115.4063$, $p < 0.0001$

Meta-analysis Shows High Variability: Pooled mean difference: **27.5%** (fixed-effects) **14.3%** (random-effects); High heterogeneity: $I^2 = 86.7\%$, $p < 0.000$; Differences likely driven by methodological variation. Differences between plasma and DBS are consistent but highly variable across studies.

Table 1. Sample Collected during MNT4P Metabolic Camp 2024

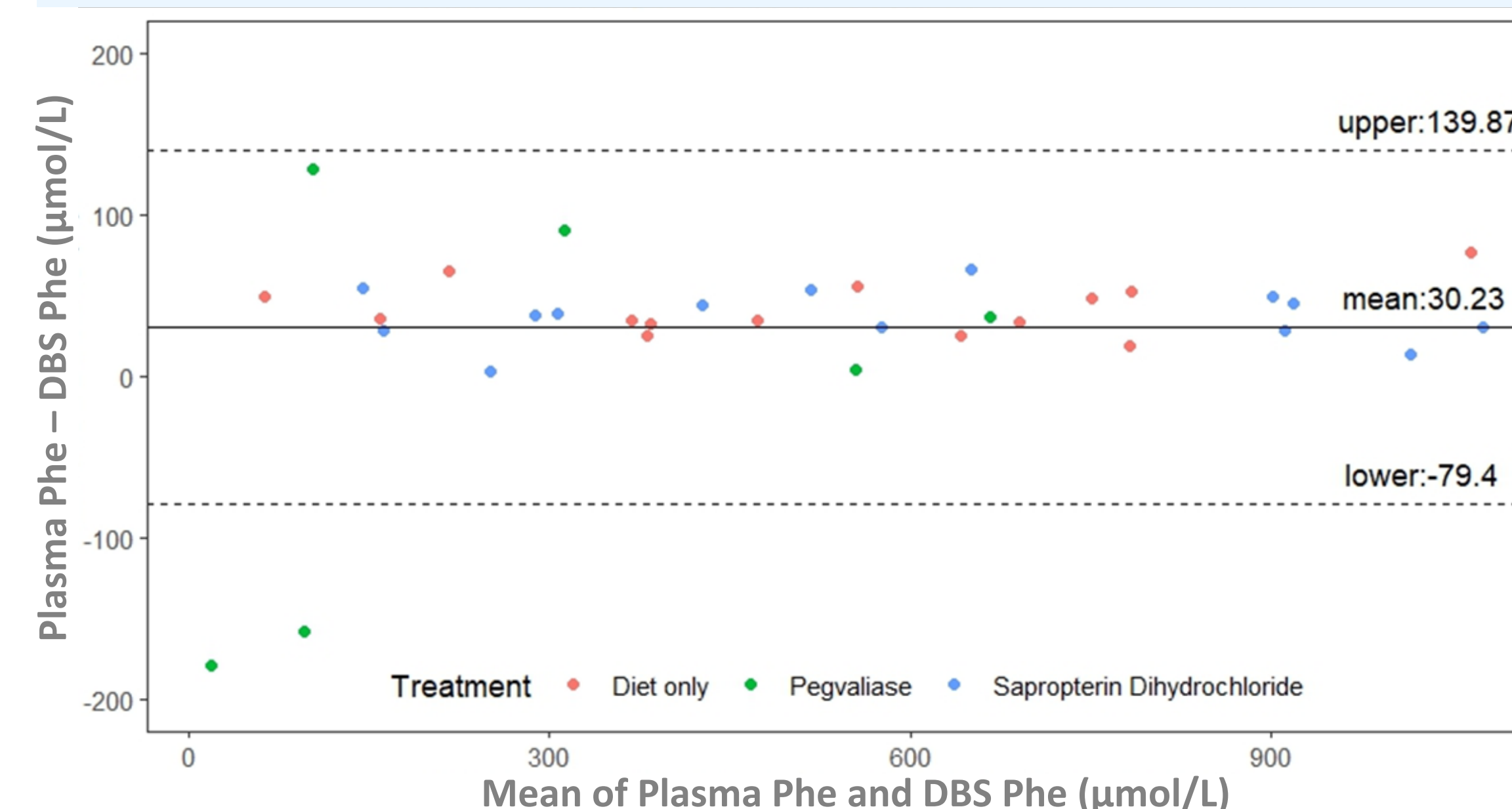
Specimen Types	N	Mean	SD	p
Dried Blood Spots (DBS)	34	408	252	***
Plasma Amino Acid (PAA)	34	604	380	***

*Statistical significance was calculated with Wilcoxon signed-rank test. *** $p < .001$.

^bPhe concentration was noted as (μM)

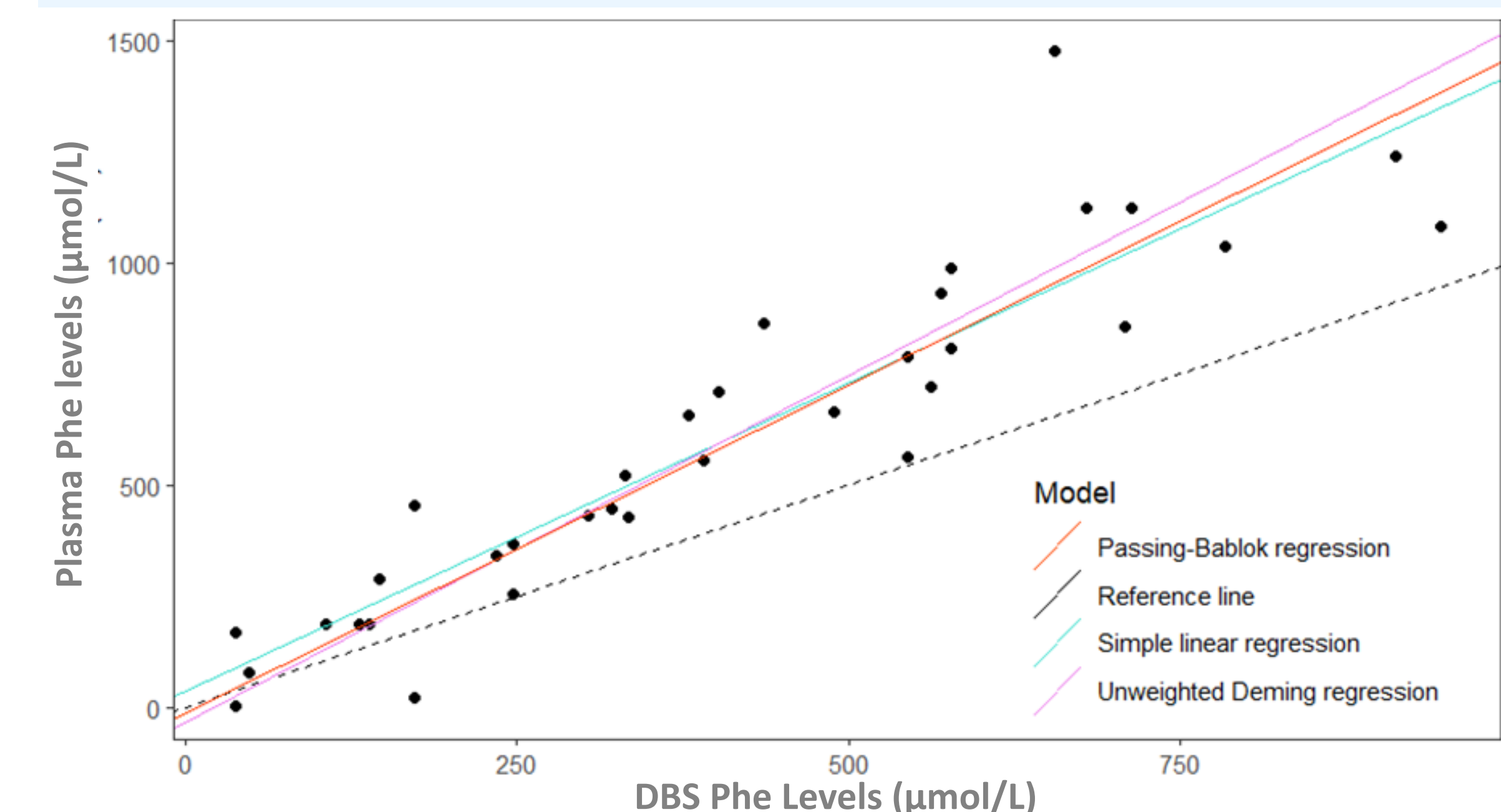
Plasma phenylalanine levels were significantly higher than DBS despite measurement using the same LC-MS/MS method ($p < 0.001$).

Figure 2. Bland-Altman Analysis of PAA vs. DBS



Bland–Altman analysis shows a ~30% positive bias, with plasma values generally higher than DBS and wide limits of agreement (-79.5% to 139.9%) indicating substantial variability. Variability appears greater at lower end.

Figure 3. Systematic and Proportional Bias Corrections



Model	Estimate	Lower CI.	Upper CI.
Raw data	0.7121	0.5753	0.8101
Simple linear	0.917	0.8428	0.957
Deming unweighted	0.9196	0.8458	0.9589
Passing-bablok	0.9194	0.8453	0.9588

*Agreement rates was calculated with Concordance Correlation Coefficient.

^bCoefficients were transformed with Fisher's Z transformation.

^cCI. indicates 95% confidence interval for Z distribution.

Plasma phenylalanine correlates with DBS phenylalanine but is systematically higher, requiring correction for comparison. The concordance correlation coefficient (CCC) was 0.71 (95% CI: 0.58–0.81), indicating moderate agreement, and improved to strong agreement after bias correction using regression models.

Conclusions

- Plasma phenylalanine levels are higher than DBS across the measurement range.
- A consistent systematic bias (~30%) affects comparability
- Adjustment methods improve agreement
- Clinical interpretation of DBS values should account for this bias

Standardization or correction factors are needed for accurate PKU monitoring.