

Challenges and Opportunities for Precision and Personalized Nutrition

A Workshop Series | August 10-12, 2021

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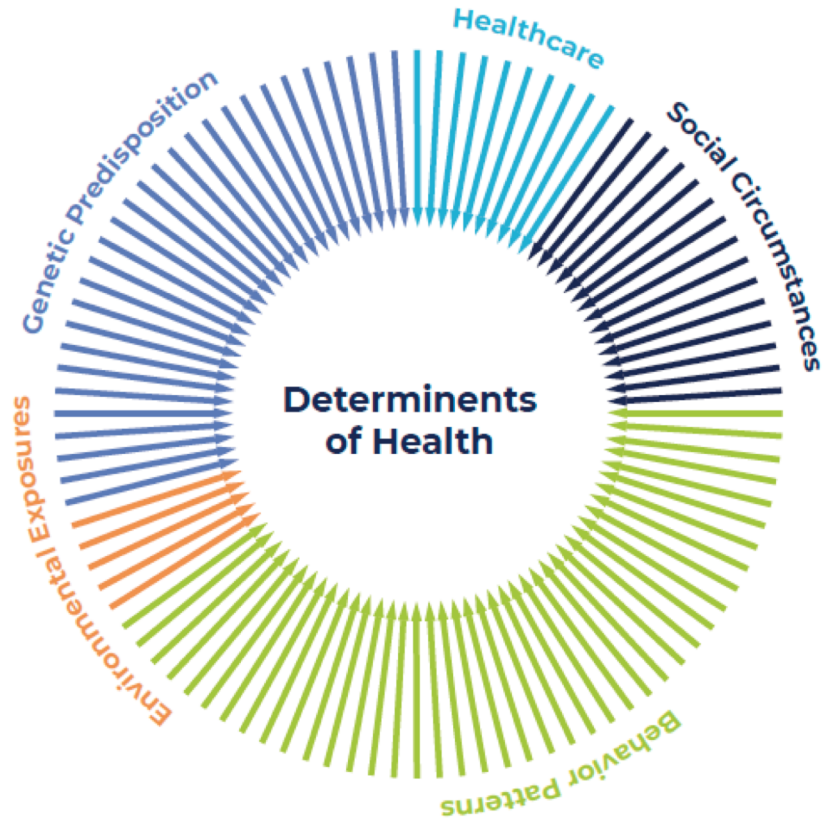
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Vydiant is a for-profit company

Health is Affected by Many Factors



nutrition, lifestyle, genetics

physical, mental, social, environmental determinants

physiological and molecular processes

drugs, disease conditions and co-morbidities

For this session, we focus on genetic contribution

Genetic Makeup: Vitamin¹ Baseline and Intervention Response

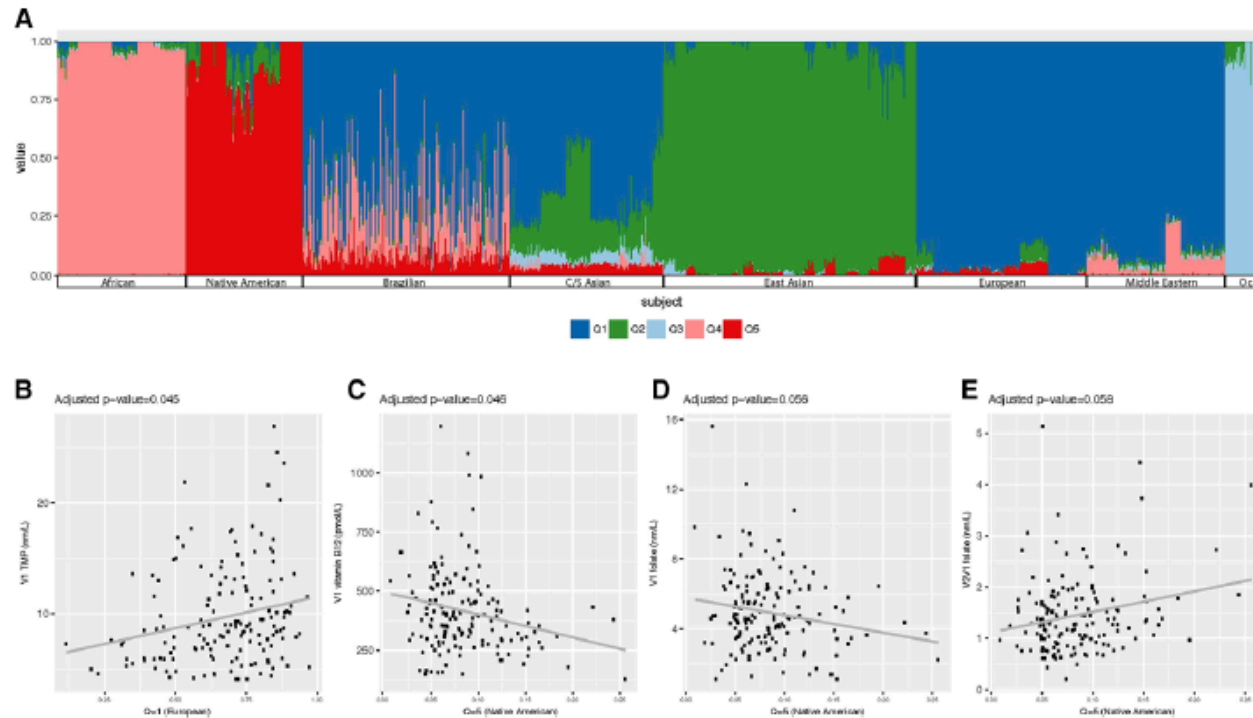


Figure 2. Admixture analysis. Influence of genetic ancestry on baseline vitamin levels. Ancestry markers from the Human Genome Diversity Project (HGDP) reference populations were used A) to identify admixture in data from unrelated participants from both years as per methods. To test whether linear regression between the ancestral components and baseline vitamin levels existed, a $k = 5$ model was used to the following covariates: trial year, sex, age, fat mass, and tanner score. Adjusted p -value of 0.05 was used as significance threshold. B) Baseline TMP and Q1 (Europe) with estimate of regression coefficient (ERC) 4.57, C) baseline vitamin B12 and Q5 (Native American) ERC = 186.53, D) baseline folate and Q5 (Native American) ERC = 2.13, and E) folate response as ratio of V2/V1 and k5 (Native American) with ERC = 0.77.

Primary Results

Low baseline TMP associated with % European ancestry (Q1)

Low baseline B12 associated with % Native Am (Q5)

Low baseline folate associated with % Native Am (Q5)

Folate responds to intervention (V2/V1) associated with NA (Q5)

¹ Measured 21 vitamins in plasma

Some Studies of Polygenic Risk Scores and Diet

Process	PRS	Reference	Process	PRS	Reference
BMI	10	<i>NEJM</i> 367 , 15 (2012)	Starch metabolism	9	<i>Diabetes</i> 69 , 1917 (2020)
BMI	32	<i>BMJ</i> 348 , g1610 (2014)	Adiposity	159	<i>EJN</i> 60 , 249 (2021)
Lipid metabolism	7	<i>Genes&Nutrition</i> 10 , 45 (2015)	Lean body mass	5	<i>Diab Ob Metab</i> 22 , 2305 (2020)
T2 Diabetes	31	<i>AJCN</i> 104 , 198 (2016)	CHO intake	37	<i>Scientific Rep</i> 11 , 13180 (2021)
BMI	77	<i>BMC Med</i> 15 , 97 (2017)	Energy intake	44	<i>Scientific Rep</i> 11 , 13180 (2021)
Vitamin D	9	<i>Hum Genet</i> 138 , 1155 (2019)	Fat intake	19	<i>Scientific Rep</i> 11 , 13180 (2021)
Vitamin D	3	<i>AJCN</i> 108 , 1129 (2019)	Exercise onset	25	<i>Scientific Rep</i> 11 , 13180 (2021)
Blood pressure	63	<i>Hypertension</i> 74 , 1460 (2019)	SCFA	7	<i>AJCN</i> 114 , 42 (2021)

Genetic Makeup and Vitamin Baseline and Intervention Response

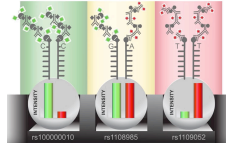
Participant
DNA

RESEARCH ARTICLE
Molecular Nutrition
Food Research
www.mnf-journal.com
Clinical and Vitamin Response to a Short-Term
Multi-Micronutrient Intervention in Brazilian Children and
Teens: From Population Data to Interindividual Responses

Human Genome
Diversity Project



Genotype

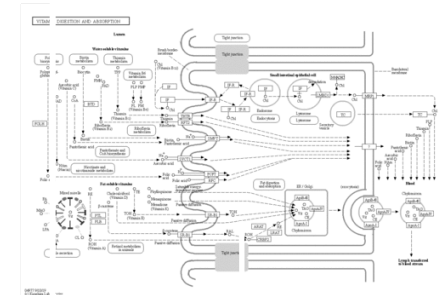


Exome Sequencing



5 million + Exon (coverage 70 – 100x)

Vitamin B12 Metabolism



76307 SNPs

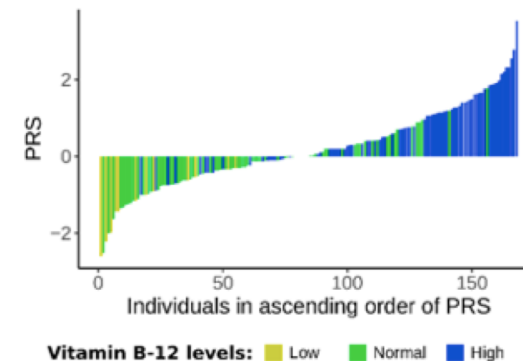
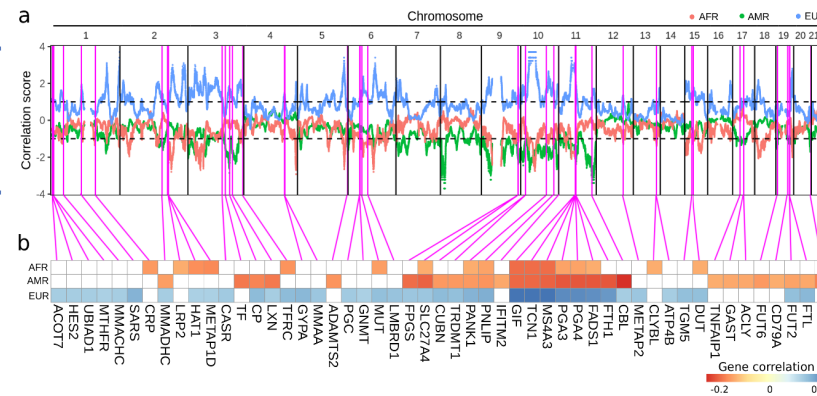
2621297 SNPs

6699 B12 SNPs in 90 genes

Ancestry
analyses

Map B12 genes to
ancestry regions

36 SNP B12 PRS
Associations

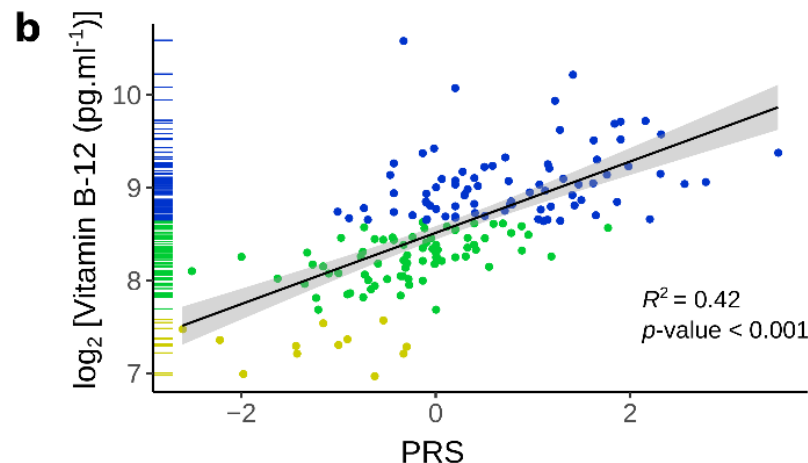
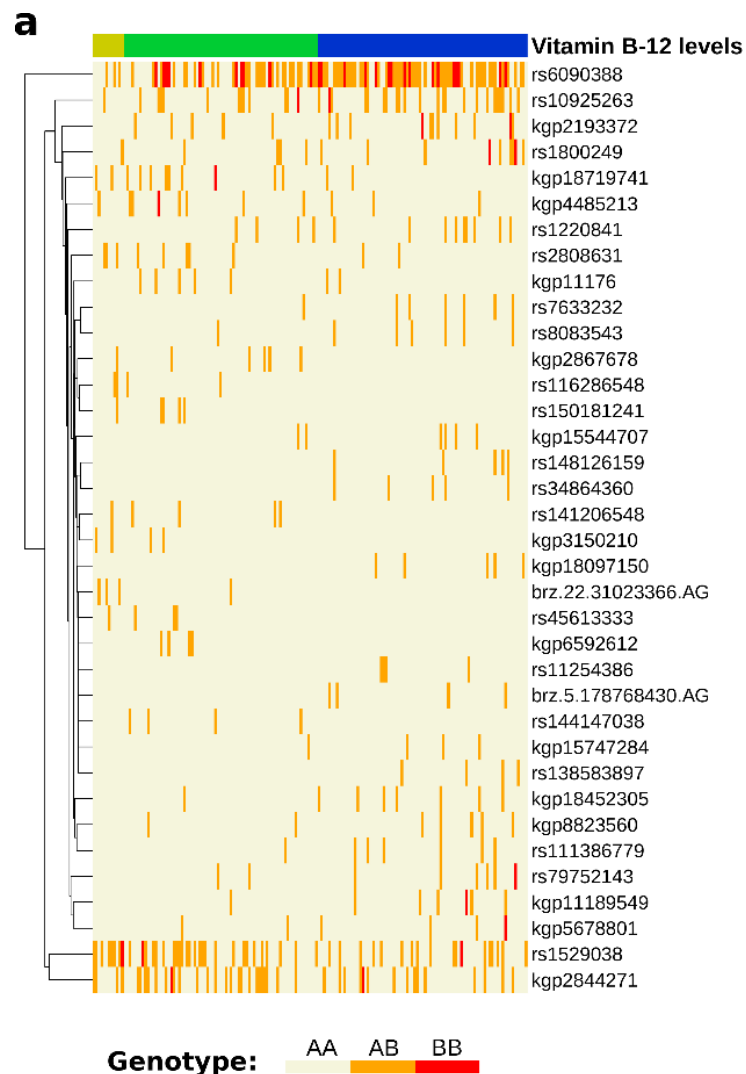


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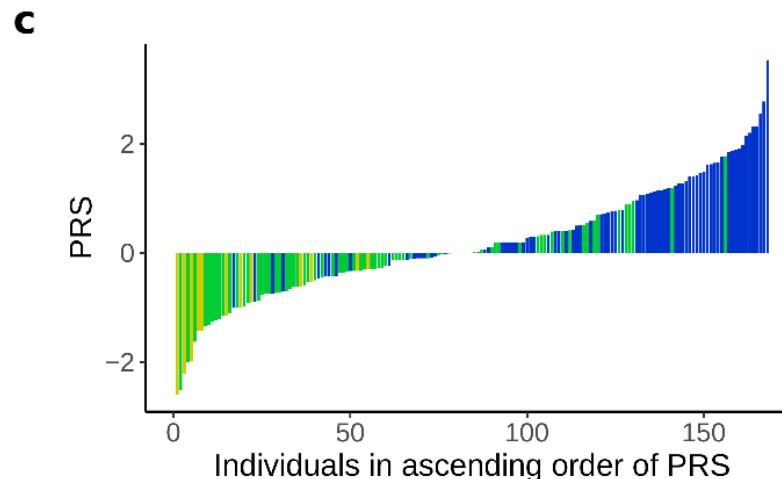
Scientific Reports 11, 11992 (2021)

Polygenic Risk Score for B12 Levels @ Baseline

Scientific Reports 11, 11992

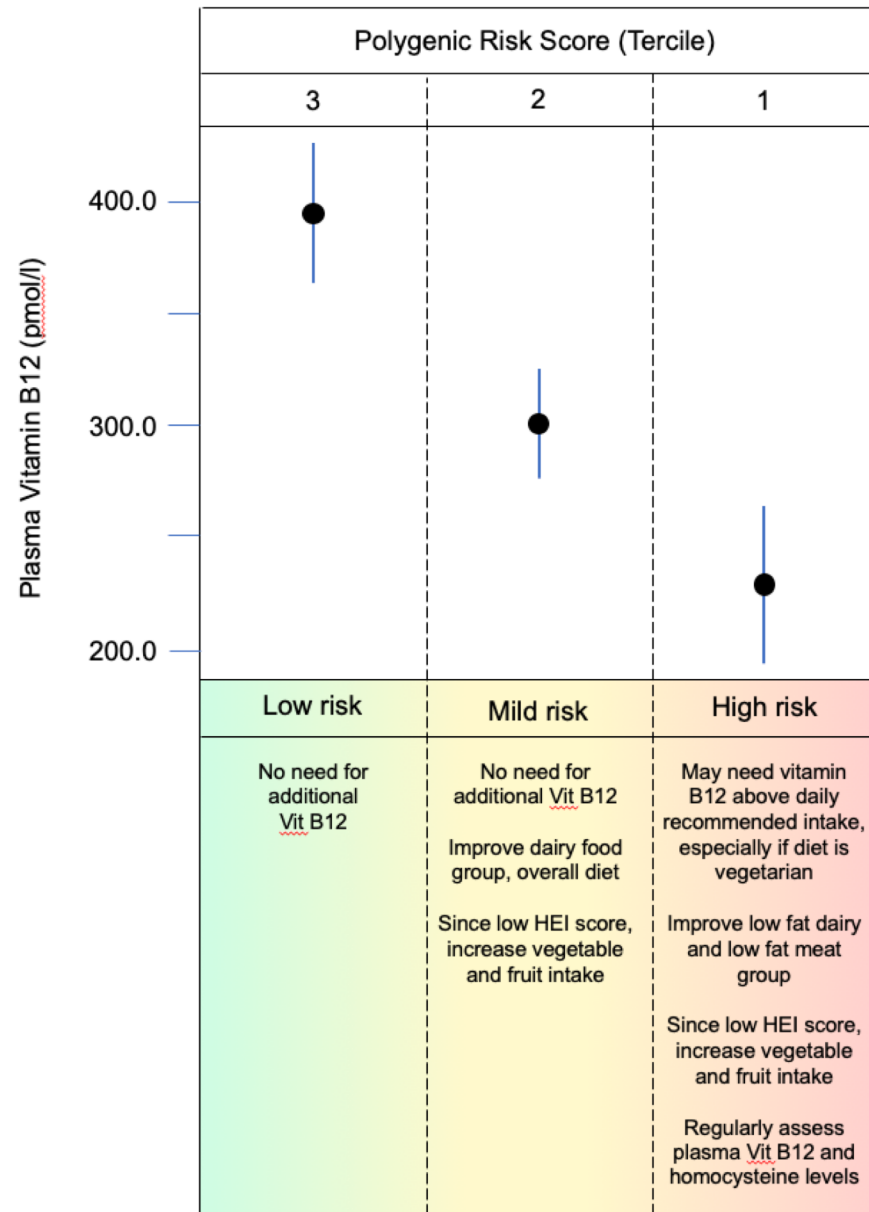


PRS of SNPs in 36 B12 genes explained 42% of phenotype



Individuals in ascending order of PRS versus baseline B12 levels.

Covariates: sex, BMI, age, total HEI and mean ancestry components (AFR, AMR, and EUR) per each individual.



Proof of Concept for Use of PRS in Personalized Diets

Individuals were stratified into terciles based on their PRS based on 36 SNPs in 26 B12 genes

Analyzed how age, sex, body mass index, total HEI, meat and milk intake (HEI components), socioeconomic status and levels of vitamin B12, homocysteine, folate, pyridoxal and riboflavin are distributed across the terciles.

Kruskal-Wallis was employed to evaluate the significance of differences among terciles within a significance level of 0.05

Path for Assessing Genetic Contributions to Phenotype

Assessing multiple variants provides more validity than single SNP

Genetic architecture is important: we corrected for AMI, EUR, AFR ancestry

Analysis of associations corrected for factors that may influence metabolite levels

Focus on “pathway” variants allowed analysis of small population

Results need to be replicated including co-variates used in this study

Thank You & Discussion

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Jacqueline Pontes Monterio PhD (co-principal investigator)

Clinical and Vitamin Response to a Short-Term Multi-Micronutrient Intervention in Brazilian Children and Teens: From Population Data to Interindividual Responses

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Mol Nutr Food Res. 62(6):e1700613 (2018)

Contribution of genetic ancestry and polygenic risk score in meeting vitamin B12 needs in healthy Brazilian children and adolescents

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