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Maternal and Pediatric Nutrition



# The Mothers, Infants, and Lactation Quality (MILQ) Study: A Multi-Center Collaboration

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#### ABSTRACT

Little valid information is available on human milk nutrient concentrations, especially for micronutrients (MNs), and there are no valid reference values (RVs) across lactation. In this multi-center collaborative study, RVs will be established for human milk nutrients across the first 8.5 mo postpartum. Well-nourished, unsupplemented women in Bangladesh, Brazil, Denmark, and The Gambia (*n* = 250/site) were recruited during the third trimester of pregnancy. Milk, blood, saliva, urine, and stool samples from mothers and their infants are collected identically at 3 visits (1–3.49, 3.5–5.99, 6.0–8.49 mo postpartum). Milk analyses include macronutrients, selected vitamins, trace elements and minerals, iodine, metabolomics, amino acids, human milk oligosaccharides, and bioactive peptides. We measure milk volume; maternal and infant diets, anthropometry, and morbidity; infant development, maternal genome, and the infant and maternal microbiome. RVs will be constructed based on methods for the WHO Child Growth Standards and the Intergrowth-21st Project. This trial was registered at clinical trials.gov as NCT03254329. *Curr Dev Nutr* 2021;5:nzab116.

#### Keywords: human milk, macronutrients, micronutrients, composition, reference values

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Abbreviations used: DBS, dried blood spots; EBF, exclusive breastfeeding; MILQ Study, Mothers, Infants, and Lactation Quality Study; MN, micronutrient; MUAC, midupper arm circumference; RV, reference value.

#### Introduction

For the first 6 mo of life, the WHO recommends exclusive breastfeeding (EBF) (1). However, human milk is not only essential for optimal infant health and development during the first 6 mo, but it can also be an important source of nutrients and other factors for the following  $\approx 18$ mo (2), yet we lack valid information on its nutrient content (3, 4), especially for micronutrients (MNs). Many human milk MN concentrations are much lower where women consume poor diets (5, 6), including MNs of major public health importance, such as vitamin A, thiamin (B-1), B-12, and iodine (7–9). These low milk concentrations caused by maternal deficiency and/or low intake have documented adverse effects on infant health and development (10), e.g. growth faltering and developmental delays have been linked to low milk concentrations in conjunction with severe maternal-infant deficiencies of vitamin B-12, B-1, B-6, D, iodine, and choline (9, 11).

In 2006, the new WHO International Child Growth Standards exposed that growth faltering starts shortly after birth with a much higher prevalence during the first 6 mo than previously assumed (12). During the same timeframe for infancy, the high prevalence of MN deficiency has been reported in exclusively or predominantly breastfed infants (6), e.g. thiamin status in Cambodia, which expands to mothers and milk, and low status of several MNs in Bangladesh (13, 14), suggesting poor milk quality could contribute to the growth faltering and MN deficiency.

The last and only global data on milk composition with a focus on MNs were collected in the 1985 WHO Collaborative Study on Breast-Feeding, over 30 y ago, with limited data collection (15). Our recent literature review on human milk revealed the general lack of a systematic sample collection and of appropriate consideration of factors affecting milk nutrient concentrations, e.g. stage of lactation, maternal nutritional status and diet, or smoking and alcohol use, among others (4, 16, 17). Moreover, inadequate methods for analyzing nutrients in the human milk matrix and lack of reported validation data added to the unreliability of many past reports (3, 18). Therefore, the literature on human milk composition is very difficult to interpret and practically impossible to use for the purposes for which such information is needed.

Due to all these limitations, there are no adequately established reference values (RVs) for nutrient concentrations in milk, and recommended nutrient intakes of infants, young children, and lactating women are based on many unvalidated and incorrect estimates of milk MN concentrations. Given the lack of RVs there is no benchmark against which to evaluate human milk quality in different populations, or the possible need for or effects of nutrient interventions on human milk MNs.

The primary study outcome and analysis objective of the Mothers, Infants, and Lactation Quality (MILQ) Study is the construction of reference ranges for values of vitamin and mineral concentrations in human milk from well-nourished mothers and their healthy infants. The reference range curves, between the 2.5 and 97.5 percentiles, will be constructed between 1 and 8.5 mo by combining data from the 4 study sites, using nutrient and volume data from milk samples obtained from each mother during 3 visits postpartum. This article describes the methodologies and protocols implemented for conducting the MILQ Study.

# **Methods**

#### Study design and settings

The MILQ Study is a multi-center, collaborative project with data and sample collection in 4 countries, Bangladesh, Brazil, Denmark, and The Gambia. Samples and data (e.g. anthropometry, questionnaires) are collected from 250 mother-infant dyads per country site (n = 1000 mothers and 1000 infants) in a systematic, identical way across sites for colostrum (1-2 d postpartum), and at 3 subsequent time points: 1.0-3.49 mo (visit 2), 3.5-5.99 mo (visit 3), and 6.0-8.5 mo (visit 4) postpartum (n = 1000 dyads). Within each time window the collection of samples is randomized so that some samples and data are available for almost every day between 1 and 8.5 mo postpartum, enabling smoothing of reference curves. Visits 2 through 4 are divided into 3- or 4-wk time periods. If a participant's first visit is randomly assigned to the first week of a time period, then their second and third visits are also scheduled in the first week of the time block. To evaluate the effect of intraperson variability on outcome variables, a subset of 50 women in Bangladesh, Brazil, and The Gambia are providing a second milk sample on visits 2 and 3. Details about each study location and analytical site are provided in Tables 1 and 2.

# Inclusion criteria

Participant's inclusion and exclusion criteria vary across the study periods but are focused mainly on the nutritional status and health of the mother and infant, and the requirement for EBF during the first 3.5 mo postpartum. Detailed information about the inclusion and exclusion criteria are provided in **Table 3**.

## Data and sample collection

Data and biological samples are collected during recruitment (28 weeks of gestation), at 1–3 d postpartum (within 72 h of delivery), and at scheduled visits 2, 3, and 4 (1.0–3.49 mo, 3.5–5.99 mo, 6.0–8.5 mo postpartum). Samples collected include human milk, and maternal and infant blood, urine, and stool. Additional data collected includes maternal and infant anthropometry, dietary intakes, and morbidity. A detailed timeline of data and sample collection is provided in **Figures 1** and **2**. To estimate milk volume, 3 sites (Bangladesh, Brazil, and The Gambia) are collecting saliva samples as described below using deuterated water. In Denmark, where this method is generally not acceptable, the mothers are instructed by the research team to test weigh the infant as described below.

# Questionnaires.

The field staff use standardized questionnaires designed for the MILQ Study. Participant's information, pregnancy screening, maternal dietary data, and socioeconomic data is retrieved during recruitment (at visit 2 in Denmark). Each of the following visits is equipped with their unique set of questionnaires (Figure 1).

#### Anthropometry.

Anthropometry measures are especially important for excluding mothers who are under- or overweight and excluding malnourished infants with abnormal birth size and Z scores  $\leq 2$ . The methods that are used to perform accurate, precise, and standardized anthropometric measurements are carried out by standard methods, with regularly validated equipment. The anthropometric measurements include:

- Mother: weight, height, midupper arm circumference (MUAC) during pregnancy; (Denmark only at visit 2). Body composition using bioimpedance analysis (BIA) in Denmark and Brazil. BMI is calculated.
- Infant: weight, length, knee-heel length (in Denmark and The Gambia), MUAC, head circumference. Body composition is estimated from birth weight and length reported in health records, and by bioimpedance at visits 2, 3, and 4, in Brazil and Denmark, and by air displacement plethysmography until the infant reaches 8 kg, (PEAPOD, COSMED) in The Gambia. Z scores are calculated by the database system using WHO, 2006 growth standards

#### Milestones.

Developmental milestones are evaluated at visits 3 and 4 to identify any infants with poor development, and to support the evidence that the milk MN concentrations are adequate for normal development. The final assessments at visit 4 includes Milestones (also collected at visit 3) and the Ages and Stages Questionnaires (19) (also administered at visits 2 and 3 in Brazil). Failure to meet normal developmental milestones in  $\geq$ 2 domains will be considered abnormal during data analysis.

#### Dietary data.

Dietary assessment of the mother is made by 2, 24-h recalls per period (visits 2, 3, and 4), and during pregnancy in Brazil and Denmark. One is taken during the study visits and the other during the week ( $\leq$ 10 d) before or after the study visit but not on an adjacent day. Data is

	Bangladesh	Brazil	Denmark	The Gambia
		-		L - C
Location	Mirpur	Madureira and	Lopennagen	bakau and Fajara
		São Cristóvão		
	(periurban area of Dhaka)	(Rio de Janeiro)	Rigshospitalet	urban Gambia near Banjul
Average income, USD/mo	245	456	3120	247
Literacy, %	69	93	100	>65
Maternity leave, mo	6 <sup>1</sup>	4-6	9–12	6
EBF, %	1		!	1
2 mo		50	71	20
	I	)	07	0
4 mo	L L	I	00	Į
6 mo	55	I	I	47
Infant stunting at 6 mo, %	18.6	4.8	~	6
Usual diet	Rice	Rice and beans	Typical Western diet	Rice and maize
	Fruit and vegetables	Fruit and vegetables	Animal source foods, dairy	Fruit and vegetables
	Some animal source foods	Animal source foods, dairy		Groundnuts
		Pasta high-processed		Some animal source foods
		foods		
	2		- - - - -	-2
MIMN supplementation Government mandates or recommendations		No, unless prescribed	some, but not in this study	NO
Perinatal			Iron and folic acid	
	I			
Fregnancy	I	Iron and tolic acid		Iron and tolic acid
	I	I	Calcium (low cow milk	I
			intake)	
Fortification programs	No	Iron and folic acid (flour)	lodine (salt)	No
Facilities	icddr, b, Dhaka	Maternity Hospital	Hvidovre Hospital	SOS Mother and Child
		Herculano	-	Clinic,
		Pinheiro	Herlev Hospital	Bakoteh
		Municipal Center of Health	Rigshospitalet	
		Ernesto Zeterino Tibau Junir		
Type of informed consent		Written consent by study participant <sup>2</sup>	udy participant <sup>2</sup>	
Sample storage and use		Sample use is explained and consented to in the informed consent document	o in the informed consent document	
-	Samples are available for 5 y	Samples are available for	Samples are a	Samples are available for 5 y
		10 y		
	Samples may be used for oth	Samples may be used for other projects but may require additional		Samples may be used for
		consent		other projects
Deidentification and data	All data is collected k	All data is collected by study ID only into a secure database, at which point it cannot be linked back to the study participants	vhich point it cannot be linked back to	o the study participants

(Continued)

	Bangladesh	Brazil	Denmark	The Gambia
	Š	Separate, password-protected participant log is maintained by research team	nt log is maintained by research team	
Compensation	Free treatment of any conditions by study physician	Travel costs	Breastfeeding counseling by phone. if desired	Travel costs
		Breastfeeding counseling	A small gift for the infant	
Follow-up	If needed, referrals are provided	If needed, determined by	A medically trained person	If needed, determined by
-	to government specialized	study nurse, referrals are	is available for	study nurse, referrals are
	hospitals for further	provided to specialized	participants experiencing	provided to
	management	care facilities for further	any illness, or if any	MRCG@LSHTM, Fajara,
		management <sup>3</sup>	abnormal test results	clinic for further
				management

For government employees, varies for private entities.

<sup>3</sup>If women are unable to read the consent form, it is read to them in their mother tongue and full informed consent is confirmed by 1) the research staff (Bangladesh, Brazil), and 2) by a literate, independent witness (The Gambia).

<sup>1</sup>Issues related 1) to illness: public Basic Health Unit, 2) to breastfeeding issues: human milk bank at study site, 3) to depression: Psychiatric Institute at the University. EBF, exclusive breastfeeding; icddr, b, International principal investigator; Centre for Diarrhoeal Disease Research, Bangladesh; MRCG@LSHTM, The MRC Unit The Gambia at the London School of Hygiene and Tropical Medicine; MMN, multiple micronutrients; PI, JSD, US dollars collected about foods consumed the previous day based on a locally validated method. All sites have locally established procedures for food and nutrient intake assessment. Ideally, the total 6 d of recall data should reflect typical days of food intake, which may include weekend day(s). Data on the nutrient intakes of the mother enables classification of each woman based on the percent inadequate intakes (i.e. intakes below the Estimated Adequate Requirement) for each MN (calculated with the Intake Modeling, Assessment and Planning Program [IMAPP] software program), comparison between maternal intake and milk concentrations of MNs and other constituents, and comparison across sites.

# **Specimen collection**

All sample collection procedures have been developed with all field sites and collaborators at the analytical sites to ensure adequate sample collection (procedures and materials needed) for all planned analyses and are described in detail in the internal MILQ Study's "Manual of Operations and Procedures." This detailed document has been shared with all field sites as the sample collection protocol, which included a completed sample collection form that states the specimen collected, subject ID, day and time, time of last meal, time of last feed, and volume and aliquots collected. All specimen collections are carried out under dim light or reduced light exposure. All samples are processed immediately and stored at  $-80^{\circ}$ C until shipment to the analytical sites.

# Colostrum.

Colostrum is collected for antigen and MN analyses, with specifics to be determined based on volumes collected. At least 0.5 mL of colostrum is collected by manual expression from all women preferably 24–48 h after birth, but 48–72 h postpartum is acceptable. After cleaning the breast with mineral-free soap towelettes or deionized water, the sample is collected either into a collection tube or directly into a 2 mL amber storage tube. Samples collected by the mother at home are stored in the home freezer at  $-18^{\circ}$ C until the first clinic visit.

# Human milk and RNA.

Milk samples at visits 2, 3, and 4 are collected preferably from the breast opposite from the one that the infant last fed from, using the Symphony electronic hospital-grade breast pump (Medela) for a full-breast expression. In the event of pump failure, milk is expressed using a hand pump or with manual expression, and noted on the collection form. The breast used for collection is cleaned with a mineral-free soap towelette or deionized water prior to milk expression. If  $\geq$ 25mL of milk cannot be obtained from the first breast, the second breast is used to obtain a second full-breast expression, which will be noted on the collection form. A second milk sample from a subset of 50 women per study site is collected according to the protocol at visits 2, and 3, in Bangladesh, Brazil, and The Gambia, ~2 wk after the respective main visit study, in order to estimate intraindividual variability in milk composition.

Milk RNA is collected in Denmark and The Gambia; 10 mL of the full-breast expression is used for RNA extraction. Under sterile conditions, the milk is centrifuged to obtain the RNA pellet, which is washed with PBS and stored under TRIzol in a DNA/RNA-free tube at  $-80^{\circ}$ C. RNA is collected from 250 women at visit 2, and 50 women at visit 3 and visit 4, at both field sites.

**IABLE 1** (Continued)

Site	Analyses
USDA, ARS-WHRNC <sup>1</sup>	Macro- and micronutrients, and metabolomics
	Selected micronutrients in colostrum based on available volume
	Microbial community of infant stool by 16s rRNA sequencing
	Shotgun metagenomics sequencing of infant stool based on 16s rRNA
	Screening, DNA quality/quantity, and availability of other data
	Milk cell mRNA transcriptome
	Maternal genetic and epigenetic signatures (buffy coat)
UC Davis, USA <sup>2</sup>	Human milk oligosaccharides and proteomics
ETH Zürich <sup>3</sup>	lodine and iodine status biomarkers
St. John's Medical College <sup>4</sup>	Milk volume by $D_2O$ saliva analyses
University of Cambridge, UK <sup>5</sup>	Vitamin D in human milk

**TABLE 2** Analytical sites of the Mothers, Infant, and Lactation Quality (MILQ) Study

<sup>1</sup>USDA, Agricultural Research Service-Western Human Nutrition Research Center, Davis, CA, USA.

<sup>2</sup>University of California, Davis, CA, USA.

<sup>3</sup>Eidgenössische Technische Hochschule, Zürich, Switzerland.

<sup>4</sup>Bengaluru, India.

<sup>5</sup>Nutritional Biomarker Laboratory, MRC Epidemiology Unit, University of Cambridge, Cambridge, UK.

D<sub>2</sub>O, deuterated water.

#### Saliva and milk intake measurements.

In the same periods that milk is collected, milk volume is measured in all women at each site, to determine how usual milk volume, which varies greatly among women, is related to nutrient concentrations and to enable estimation of total daily nutrient intake by the infant. Three of the field sites (Bangladesh, Brazil, and The Gambia) are using the International Atomic Energy Agency (IAEA) mother to child deuterium (D2M) method (https://www-pub.iaea.org/MTCD/Publicatio ns/PDF/Pub1429\_web.pdf). Deuterated water (30 g) is administered to the mothers at visits 2, 3, and 4 and maternal and infant saliva samples collected at baseline, 1, 2, 3, 4, 13, and 14 d after dosing. Saliva samples are obtained from both mothers and infants, who are both weighed at baseline and day 14. Cotton balls are placed in the infant's mouth which are then squeezed by a syringe to retrieve the saliva. Denmark is estimating milk volume via 24-h test weighing of the infant before and after every feed for 24 h, plus 1 extra feed, to determine the 24-h intake. The infants are weighed with the same clothes and diaper at both weighings. Mothers are instructed by study personnel.

#### Blood.

A venous blood sample is collected into an EDTA vacutainer from the mother on visits 2, 3, and 4. Fasting overnight or for 3–4 h prior to blood collection was ideal if possible. The mother's recall of the time of her last meal is recorded. A venous sample is collected from all infants at visit 2, and to reduce the number of infant blood draws to a total of 2, from 50% of infants at visit 3 and the other 50% at visit 4, randomly selected. In The Gambia, however, 100% of infants are sampled at each visit as well as the mother during pregnancy. The mother's recall of the time of the infant's last meal is also recorded.

Plasma is obtained by centrifugation of the blood sample at  $1500 \times g$  for 10 min at 4°C. The plasma is stored in aliquots in amber tubes. The buffy coat is collected into a DNAse/RNAse-free, sterile tube. The volume of the remaining RBCs is determined in order to wash the RBCs with an equal volume of 0.9% saline solution. The RBCs are centrifuged again and the supernatant is removed. This procedure is repeated until the supernatant is clear. After diluting with equal amounts of deionized water, the washed RBCs are aliquoted into amber

tubes for storage. All aliquots are then frozen and stored at  $70^\circ\text{C}$  until analysis.

#### Dried blood spots.

Dried blood spots (DBS) are obtained for iodine status assessment. Four (infant) or 6 (mother) spots are prepared on filter paper cards using 50  $\mu$ L of whole blood from the EDTA vacutainer used for blood collection. If insufficient blood volume is obtained, 1 DBS is prepared. The blood is dispensed onto the filter paper without touching the paper, and after a drying period of 24 h at  $\leq 25^{\circ}$ C, the cards are stored in bags without the DBS touching each other. The DBS can be stored at  $-20^{\circ}$ C until shipment.

#### Urine.

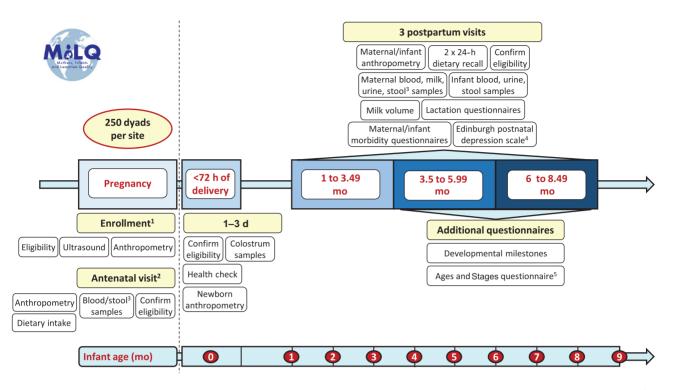
Maternal and infant urine is collected to assess iodine status across sites, which will likely vary depending on consumption of locally fortified foods, e.g. iodized salt. Population daily iodine intake will be estimated from spot urinary iodine and creatinine concentrations. A midstream clean catch specimen is collected from the mother at the first voiding of the bladder during each visit. Infant urine collection is enabled using a cotton ball, pads, or urine collection bags that are placed inside a disposable diaper. The wet but not soiled (with feces) cotton ball or pad is removed from the diaper and the urine is obtained by squeezing the cotton balls in a syringe, or by squeezing the pad in a plastic bag with a missing corner. The urine is collected into a beaker and 2 aliquots from both the mother and infant urine are stored for iodine assessment.

#### Stool.

Infant feces are used for microbiome profiling. Once the infant has a bowel movement, the stool is scored by the mother for consistency, color, and volume. About 1 g of the specimen is collected with the spoon (attached to the cap) of the feces collection tube. The spoon is placed into the collection tube and stored at  $-80^{\circ}$ C. If the specimen is collected outside the clinic (not applicable in The Gambia, all stool sample collection here is conducted at the study site), it must be stored in the home freezer and delivered frozen, on ice, for  $-80^{\circ}$ C storage as soon as

		brazil		
	<b>D</b>			
Recruitment	Directly by local research team	Invitation letter to meet	Directly by local research team	ch team
Enrollment	After delivery	X	During pregnancy	y
Mothers inclusion criteria				
Age, y		18–40		
Height, cm RMI bc/m <sup>2</sup>	>145		≥150	
	>18 5 to <30.0			>18 5 to <30.0
		_ >18 5 to <30 0	_ >18 5 to <30 0	
MUAC, GII Hemodobin <sup>1</sup> a/I	25 t0 2 33 100	/ 100		<pre></pre>
		ont or originat modical arablema (e.e. 20)		
		INUTEREVARIT PAST OF CURRENT INEGREAL PRODIERITS (E.G. GESTALIONAL MADELES, PRE-ECLARITIPSIA)	uauonai uiabetes, pre-eciampsia)	-4
	NO	NO	No C	NO
Alconol intake, mL	000	<40	06>	06>
Diet	Nonvegan or macrobiotic diet	>15 a of 4 of 8 food around/d	>15 a of >15 food aroune each/d	p/quee se
MMN use				
3rd trimester	Iron and folic acid			
			Vitamin D and calcium	
lactation	Iron and folic acid			
			Vitamin D and calcium	
Fortified foods		Low habitual intake of highly fortified foods, expect iodized salt	s, expect iodized salt	
Delivery		Singleton		
Weaks of destation		CV_75		
Breastfeeding				
		Current and		
2.3 Elo		Exclusive breastleeding		
Birthweight, g		2500-4200		
Infant health	No congen	No congenital malformations that interfere with feeding or growth and development	ng or growth and development	
Visit 2 exclusion criteria		Cessation of, or nonexclusive, breastfeeding	sastfeeding	
		Serious maternal illness	S	
		Infant length-, weight-for-age, or weight-for-length <-2 Z	t-for-length <-2 Z	
	Mat	Maternal MMN consumption during lactation other than stated above	other than stated above	
Visits 3 and 4 exclusion		Cessation of breastfeeding	bu	
criteria				
		Infant length-, weight-for-age, or weight-for-length <-2 Z	t-for-length <-2 Z	
	Abnormal intant developmer	Abnormal intant development based on Ages & Stages Questionnaire (abnormal performance on $\geq\!2$ of the / domains)	(abnormal performance on $\geq 2$ of the / of	domains)

**TABLE 3** Inclusion and exclusion criteria in the Mothers. Infants. and Lactation Ouality (MII O) Study



**FIGURE 1** Mothers, Infants, and Lactation Quality Study timeline, questionnaires, and data and sample collection form schedule. <sup>1</sup>3rd trimester. <sup>2</sup>Late 3rd trimester, not done in Bangladesh. <sup>3</sup>Maternal stool samples are not collected in Bangladesh. <sup>4</sup>Not collected in Denmark. <sup>5</sup>Only at visit 4 in The Gambia.

possible. All participants have been instructed by the field study staff and have confirmed the availability of a freezer for eligibility of at-home stool collection. All times of collection and storage are recorded.

# Laboratory analyses

All samples except those to be used for iodine and milk volume determination are shipped to the USDA, Agricultural Research Service-Western Human Nutrition Research Center (USDA, ARS-WHNRC), Davis, CA, USA. At each site, the research personnel prepare the aliquots for the following analyses (Table 3). Samples for human milk vitamin D analysis will be sent from the USDA/ARS-WHNRC to the Nutritional Biomarker Laboratory, MRC Epidemiology Unit, University of Cambridge, Cambridge, UK.

## Data management.

A dedicated, REDCap database was developed at a single site (The Gambia) to be used across all study settings. Unique  $\alpha$ -numeric study ID codes are generated and given to each participant. These will link all data collected from a specific individual. ID codes include a check letter to minimize ID errors. Data collection is carried out electronically, or on paper forms when necessary, by study staff. All forms in the field were designed specifically for the MILQ Study. Research assistants and supervisors review data on a daily basis before data from each field site is securely integrated into The Gambia REDCap database by a data manager. A copy of the final data obtained from the laboratory/analytical sites is reviewed by the study statistician before being securely integrated in a WHNRC-housed REDCap database. Both databases will be merged at

the end of the field site data collection and integration. Data management and security procedures, including assurance of confidentiality, adhere to the Collaborative Institutional Training Initiative (CITI) and the Canadian Tri-Council Policy Statement on Ethical Conduct for Research Involving Humans (TCPS2 CORE) guidelines, and are outlined in full in the protocol at clinicaltrials.gov (NCT03254329, 18 August, 2017).

## Statistical analysis

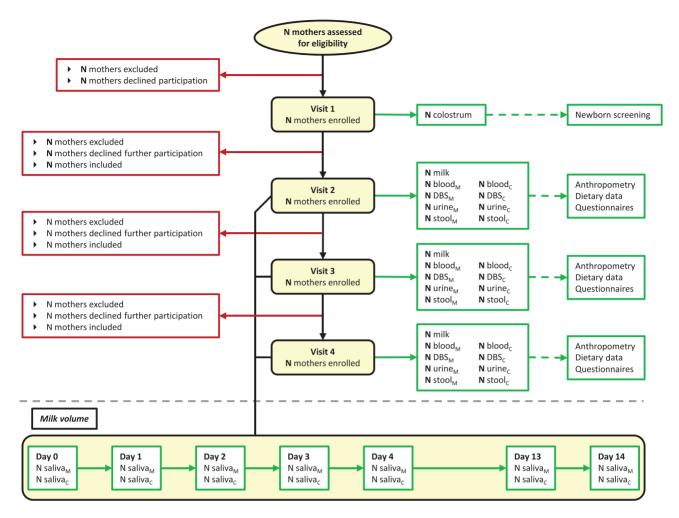
# Sample size calculations.

Since the primary outcome is estimated key centiles of the distributions of nutrient concentrations in human milk, and centiles are invariant to monotonic transformation, sample sizes for constructing RVs are based on: *1*) estimating the 50th and 5th percentiles of each breast milk nutrient across time periods within each site and *2*) being able to establish equivalency among study sites before pooling data.

# Estimating centiles.

We aim for a monotonic transformation for each variable across all study sites to attain normal (Gaussian) distribution, from which parametric estimates of the centiles will be constructed. Regression models will be used to estimate the parameters of the distribution at each time period, but since such models will not be determined until the data are available, the sample size is based on a generic estimate for the p<sup>th</sup> centile:

$$\bar{X} + Z_p \times s \times CF_s \tag{1}$$



**FIGURE 2** Mothers, Infants, and Lactation Quality Study participant flow chart, for all 4 study sites. <sup>1</sup>Denmark will use 24-h test weighing, instead of the Dose-to Mother deuterium method and saliva collections, for milk volume measurements.  $_{C}$ , child; DBS, dried blood spots;  $_{M}$ , mother.

where:

 $\bar{X}$  = sample mean (or estimated predicted value from the regression model)

 $Z_p = p^{\text{th}}$  centile of the standard normal distribution (negative for P < 0.50)

s =sample SD (or estimated SD from regression model)

 $CF_s$  = correction factor for bias of s as an estimator of population SD (negligible for n > 50).

The SE of this estimate, if *n* is > 50, is approximately:

$$\frac{\sigma}{\sqrt{n}}\sqrt{1+0.5\times Z_{\alpha}^2} \tag{2}$$

The desired width of the CI is based on the distance of the 50th and 5th centiles from neighboring centiles, which yields a sample size invariant to the location, scale, or shape of the underlying distribution, as long as the variable can be normalized. This requires an overall sample size of 255 per time point to avoid overlap between the 95% CIs around the 5th and 10th centiles, and to estimate the sample median within 5 centiles with 95% confidence (**Table 4**). Once the centiles of interest are estimated for the transformed variable, they will be back-transformed to be expressed in the original units.

For maternal and infant blood variables, it is likely that blood samples will not be available from all participants, so the sample size will be reduced by necessity. However, if it is determined that it's acceptable to pool information from  $\geq 2$  study sites to construct RVs for infant blood variables, the precision of the estimate will be acceptable.

## Establishing equivalency among study sites.

For each nutrient and time period, pairwise comparisons using the 2, 1-sided test (TOST) method will be used to compare the means of the 4 sites for equivalence (20). If differences >0.3 SD are detected, a followup test will be conducted to compare each mean to the combined means of the other 3 sites, to determine which sites can be pooled. A sample size of 200 per site in each time period is sufficient to detect differences of 0.25 SD with 80% power or differences of 0.30 SD with 90% power. Therefore, the planned sample size of 250 per site and time period will

multic caregoit	condimited and a second s	INIIK	DOOID	Urine	Stool	Method	Ket.
Fat-soluble vitamins and carotenoids	Vitamin A Vitamin E lpha-carotene eta-carotene eta-carotene	×	۳×			HPLC-MWL	(31)
	Lycopene Lutein/zeaxanthin Ergocalciferol (D <sub>2</sub> ) Cholecalciferol (D <sub>3</sub> ) 25-OH-D <sub>2</sub>	×	Å			UPLC-MS/MS CPBA	(32)
Water-soluble vitamins	25-OH-D <sub>3</sub> Thiamin (B-1) Thiamin monophosphate (B-1)	×	X <sup>R</sup>			HPLC-FLD	(33)
	Thiamin diphosphate (B-1) Riboflavin (B-2) FAD (B-2) FMN (R2)	×	X <sup>P,R</sup>			UPLC-MS/MS	(27)
	Nicotinacide (B-3) Nicotinacide (B-3) NAD (B-3) NAD (B-3) NAD(P) (B-3) Nicotinamide riboside (B-3) Tryptophan (amino acid, B-3) Partothenic acid (B-5) Pyridoxal (B-6) Pyridoxal (B-6) Pyridoxal 5-phosphate (B-6) Pyridoxal 5-phosphate (B-6) Biotin (B-7) Folic acid (B-9) 5-methyl tetrahydrofolate (B-9)	;	ę			CPBA2	(34)
Choline and related metabolites	Cobalamin (B-12) Choline Phosphocholine Glycerophospho choline Bettaine Carnitine Creatinine Dimethylglycine	××	×××			UPLC-MS/MS	(28, 35) (36)
B-12 biomarkers	Methionine Trimethylamine N-oxide Methylmalonic acid Homocysteine		<sup>d</sup> × <sup>d</sup> ×			UPLC-MS/MS HPLC-FLD	(38) (38)

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Mnerals and trace elements         Compet Selenion Solenion Solenion Solenion Solenion Solenion Solenion Solenion Calcum Dryord strutulared hormone Thyryord s	Analyte Category	Alialytes		2000		0000		
trace elements Zin Selenium Selenium Selenium Selenium Selenium Selenium Selenium Potastum Magnestum Magnestum Pota	Minerals and	Iron	×	Х <sup>Р</sup>			ICP-MS	(39)
Selenum Sodium Sodium Potssium Potssium Potssium Potssium Potssium Potssium Potsetium Calcium Thyropideulin Thyropideulin Protein Thyropideulin Protein Thyropideulin Protein	trace elements	Copper Zinc						
Potasium Porasium Magnesium Magnesium Magnesium Macronutrients     Potasium Potasium Magnesium Magnesium Magnesium Magnesium Macronutrients     Potasium Potasium Magnesium		Selenium						
Potassum     Potassum       Invorgiobulin     X       Acronutrients     Thyrogiobulin       Tyroid-stimulated hormone     Thyrogiobulin       Tyroid-stimulated hormone     Thyrogiobulin       Tyroid-stimulated hormone     Thyrogiobulin       Torsin     The Link       Matabolomics     The Link       Biocrates MarkPo CUANT 500 <sup>1</sup> X       Matabolomics     The Link       Biocrates MarkPo CUANT 500 <sup>1</sup> X       Careaction protein (CRP)     X <sup>2</sup> Matabolomics     The Link       Discrates MarkPo CORPA     Link       Torsin, Hou     The Link       Torsin, Hou     The Link       Torsin, Hou     The Link       Torsin     The Link       Torsin     Link       Torsin     The Link       Torsin     Link       Torsin     Link       Torsin     Link       Torsin     Link       Torsin </td <td></td> <td>Sodium</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>		Sodium						
Magnesum         Magnesum           Iodine         Tyroglobulin         ELSA         (4)           Tyroglobulin         Tyroglobulin         Tyroglobulin         (4)           Tyroglobulin         Tyroglobulin         Tyroglobulin         (4)           Tyroglobulin         Tyroglobulin         Tyroglobulin         (4)           Macronutrients         Protein         X         NRspectroscopy         (4)           Matronutrients         Protein         X         NRspectroscopy         (4)           Matronutrients         Proteomics         NRspectroscopy         (4)           Matronutrients         Proteomics         NRspectroscopy         (4)           Matronutrients         Proteomics         NRspectroscopy         (4)           Matronutrintin		Potassium						
Index     Calcum     XPBS     X     Clocking     (CP-MS     (CP-MS       Thyroglobulin     Thyroglobulin     Thyroglobulin     Thyroglobulin     (1)       Thyroglobulin     Thyroglobulin     Thyroglobulin     Thyroglobulin     (1)       Macronutrients     Total thyroxine     X     (2)       Macronutrients     Fortein     X     (2)       Fortein     Carbohydrates     (2)     (2)       Glycomics     Human milk oligosaccharides <sup>1</sup> X     (4)       Metabolomics     ElsA     (1)     (2)       Metabolomics     Biocrates MxP® QUANT 500 <sup>3</sup> X     (4)       Metabolomics     Biocrates MxP® QUANT 500 <sup>3</sup> (4)     (4)       Metabolomics     Biocrates MxP® QUANT 500 <sup>3</sup>		Magnesium						
Iodine status     Iodine Tryroglobul Tryroglobul     X     CP-NS     (41)       Tryroglobul     Tryroglobul     Tryroglobul     Tryroglobul     (41)       Tryroglobul     Tryroglobul     Tryroglobul     (41)       Tryroglobul     Protein     Nespectroscopy     (42)       Macronutrients     Protein     Nespectroscopy     (42)       Fat     Cabobydrates     Cabobydrates     Nespectroscopy     (43)       Glycomics     Human milk proteomics <sup>4</sup> X     Nespectroscopy     (43)       Metabolomics     Biocrates Number     Vertein     Nespectroscopy     (43)       Metabolomics     Creactive protein (ZPP)     X     Nespectroscopy     (43)       Metabolomics     Sociates Numpersonics <sup>4</sup> X     Nespectroscopy     (43)       Metabolomics     Globel transferin receptors     X*     Nespectroscopy     (43)       Inflammation markers     Sociates Numpersonics <sup>4</sup> X*     Nespectroscopy     (43)       Inflammation markers     Sociates Numpersonics <sup>4</sup>		Calcium						
Thyroglobulin     Thyroglobulin     ELISA     (41)       Thyroid-stimulated hormone     Thyroid-stimulated hormone     X     NIR-spectroscopy     (42)       Total thyroxine     X     Nin-spectroscopy     (42)       Protein     Carbohydrates     X     NIR-spectroscopy     (42)       Garbohydrates     X     Nin-spectroscopy     (43)       Mactonutrients     Earbohydrates     X     NIR-spectroscopy     (43)       Matbohomics     Biocrascharides <sup>3</sup> X     YP     UPLC.Chip/TOF-MS     (44)       Matbohomics     Biocrascharides <sup>3</sup> X     YP     UPLC.Chip/TOF-MS     (44)       Matbohomics     Biocrascharides <sup>3</sup> X     YP     UPLC.Chip/TOF-MS     (45)       Inflammation markers     c-1-acid glycoprotein (AGP)     X     YP     UPLC.Chip/TOF-MS     (45)       Inflammation markers     c-1-acid glycoprotein (AGP)     X     YP     UPLC.Chip/TOF-MS     (45)       Inflammation markers     c-1-acid glycoprotein (AGP)     XP     UPLC.Mip/TOF-MS     (45)       Inflammation markers     c-1-acid glycoprotein (AGP)     YP     UPLC.Mip/TOF-MS     (45)       Inflammation markers     c-1-acid glycoprotein (AGP)     YP     UPLC.Mip/TOF-MS     (45)       Inneatures     Soluble transferinin	lodine status	lodine	×		×		ICP-MS	(40)
Thyroid-stimulated hormone     Thyroid-stimulated hormone     NIR-spectroscopy     (42)       Total thyroxine     Tat     NIR-spectroscopy     (43)       Dist thyroxine     Eat     Carbohydrates     (44)       Clycomics     Human milk proteomics <sup>4</sup> X     (44)       Metabolonics     Biocrates MxP60 DJANT 500 <sup>3</sup> X     (44)       Metabolonics     Carbohydrates     (44)       Inflammation markers     carbohydrates     (45)       Creation for this     Carbohydrates     (44)       Inflammation markers     carbohydrates     (45)       Creation option (CRP)     X     (45)       Creation protein (CRP)     X     (45)       Creation option (CRP)     X     (45)       Information markers     carbohydrates     (45)       Creation option (CRP)     X     (45)       Creation option (CRP)     X     (45)       Information markers     Creation option (CRP)     (45)       Creation option (CRP)     X     (45)       Information markers     Creatin underse		Thyroglobulin		X <sup>DBS</sup>			ELISA	(41)
Total thyroxine     Total thyroxine     X     NIRspectroscopy     (42)       Fat     Eachohydrates     Carbohydrates     (43)       Glycomics     Human milk oligosaccharides <sup>3</sup> X     PPLC Chip/TOF-MS     (43)       Metabolomics     Biocrates Mx@OUANT 500 <sup>3</sup> X     PPLC Chip/TOF-MS     (43)       Metabolomics     Biocrates Mx@OUANT 500 <sup>3</sup> X     XP     PPLC Chip/TOF-MS     (43)       Inflammation markers     Carbohydrates     Carbohydrates     (43)     (44)       Inflammation markers     Carbohydrates     (44)     (45)     (45)       Inflammation markers     Carbohydrates     (45)     (45)     (45)       Inflammation     Carbohydrates     Carbohydrates     (45)     (45)       Inflamustion     Carbohy		Thyroid-stimulated hormone						
Macronutrients     Protein     X     NIR-spectroscopy     (42)       Carbohydrates     Carbohydrates     X     HPLC Chip/TOF-MS     (43)       Carbohydrates     Carbohydrates     Carbohydrates     (43)       Carbohydrates     Luman milk proteomics <sup>4</sup> X     HPLC Chip/TOF-MS     (43)       Metabolomics     Biocrates MxP® OUANT 500 <sup>3</sup> X     X     (43)       Metabolomics     Biocrates MxP® OUANT 500 <sup>3</sup> X     X     (43)       Inflammation markers     era-1-acid glycoprotein (AGP)     X     (43)       Creactive protein (CRP)     X     X     (45)     (45)       Inflammation markers     Evaluation     (45)     (45)     (45)       Inflammation     Evaluation     X     X     (45)     (45)       Inflammation     Evaluation     X     X     (45)     (45)       Inflammation     Evaluation     X     X     (45)		Total thyroxine						
Fat     Fat       Glycomics     HPLC Chip/TOF-MS       Human milk oligosaccharides <sup>3</sup> X       Human milk oligosaccharides <sup>3</sup> X       Human milk oligosaccharides <sup>3</sup> X       Human milk origosaccharides <sup>3</sup> X       Biocrates MxP® QUANT 500 <sup>3</sup> X       Biocrates MxP® QUANT 500 <sup>3</sup> X       Biocrates MxP® QUANT 500 <sup>3</sup> X       Retabolomics     Biocrates MxP® QUANT 500 <sup>3</sup> Biocrates MxP® QUANT 500 <sup>3</sup> X       Creactive protein (GP)     X       Creactive protein (CR)     X <sup>1</sup> II-16, II-2, II-3, II-3, X     X       TNF-a, IFN-y     MSD immunoassay       Icon status     Soluble transferrin receptors       Formin     X       Romones     Leptin, insulin, and adiponectin       Korobione     Leptin, insulin, and adiponectin       Korobione     K       Microbione     Microbione       Microbione     Microbione       Microbione     Microbione       Microbione     X <sup>1</sup> Microbione     Microbiol dood opots       Microbio	Macronutrients	Protein	×				NIR-spectroscopy	(42)
Glycomics     Carbohydrates     X     HPLC Chip/TOF-MS     (4)       Human milk oligosaccharides <sup>3</sup> X     X     UPLC.Cdip/TOF-MS     (4)       Metabolomics     Biocrates MxP® OUANT 500 <sup>3</sup> X     XP     UPLC.GqQ-MS     (4)       Metabolomics     Biocrates MxP® OUANT 500 <sup>3</sup> X     XP     UPLC.GqQ-MS     (4)       Metabolomics     Biocrates MxP® OUANT 500 <sup>3</sup> X     XP     UPLC.GqQ-MS     (4)       Inflammation markers     ar-1-acid glycoprotein (GP)     X     XP     (4)       Inflammation markers     ar-1-acid glycoprotein (GP)     X     (4)       Inflammation markers     ar-1-acid glycoprotein (GP)     (4)     (4)       Inflammation     The.u., Inflammation     (4)     (4)       Inflamones     Erotid     (4)     (4)		Fat					-	
Glycomics     Human milk oligosaccharides <sup>3</sup> X     HPLC Chip/TOF-MS     (43)       Metabolomics     Human milk porteomics <sup>4</sup> X     UPLC-CqQ-MS     (44)       Metabolomics     Biocrates MxP60 QUANT 500 <sup>3</sup> X     X     (45)       Metabolomics     Biocrates MxP60 QUANT 500 <sup>3</sup> X     X     (45)       Inflammation markers     c-lead by propretin (AGP)     X     X     (45)       C-reactive protein (AGP)     X     X     (45)       Inflammation markers     c-lead by propretin (AGP)     X     (45)       Drankers     c-lead by propretin (AGP)     X     (45)       Drankers     c-lead by propretin (AGP)     X     (45)       Drankers     c-lead by propretin (AGP)     X     (47)       Drankers     Solute transferin receptors     X     (47)       Drankers     Solute transferin receptors     X     (47)       Hormones     Leptin, insulin, and adjononectin     X     X       Genetics     Single nucleotide polymorphism     X <sup>B</sup> Solute By fixes       Cod Genet		Carbohydrates						
Human milk proteomics <sup>4</sup> X       X       UPLC-GqQ-MS       (4)         Metabolomics       Biocrates MxP® QUANT 500 <sup>3</sup> X       X       (4)         Inflammation markers       carcid glycoprotine (AGP)       X       (4)         Creactive protein (CRP)       C-reactive protein (CRP)       NSD immunoassay       (4)         Icn status       Collofle transferrin receptors       X       X       (4)         Icn status       Soluble transferrin receptors       X       X       (4)         Icn status       Soluble transferrin receptors       X       (4)       (4)         Icn status       Soluble transferrin receptors       X       (4)       (4)         Remain       Remain       X       X       (4)       (4)         Mornones       Leptin, insulin, and adiponectin       X       X       (4)       (4)         Genetics       Single nucleotide polymorphism       X       X       (4)       (4)         Microbione       Microbione       Microbial communy       (4)       (4)       (5)         Microbione       Microbial community       X       X       (4)       (5)         Microbiane       Microbial community       X       X       (5)	Glycomics	Human milk oligosaccharides <sup>3</sup>	×				HPLC Chip/TOF-MS	(43)
Metabolomics       Biocrates MxP® QUANT 500 <sup>3</sup> X       XP       UPLC-MS/MS       (45)         Inflammation markers       a-1-acid glycoprotein (AGP)       XP       CPBA       (45)         Inflammation markers       a-1-acid glycoprotein (AGP)       XP       (47)         Inflammation markers       a-1-acid glycoprotein (AGP)       XP       (47)         Inflammation markers       a-1-acid glycoprotein (AGP)       (47)         Inflammation markers       a-1-acid glycoprotein (AGP)       (47)         Inflammation markers       a-1-acid glycoprotein (CRP)       (47)         Inflammation markers       a-1-acid glycoprotein (CRP)       (49)         Inflammation markers       CPBA       (49)         Inflammation markers       Soluble transfer.In receptors       X       X         Inflammation markers       CPBA       (49)       (49)         Hornones       Leptin, insulin, and adiponectin       X       X       (49)         Hornones       Leptin, insulin, and adiponectin       X       X       (49)         Genetics       Single nucleoride polymorphism       X <sup>B</sup> (48)       (49)         Genetics       Single nucleoride polymorphism       X <sup>B</sup> (48)       (49)         Microbial co		Human milk proteomics <sup>4</sup>	×				UPLC-QqQ-MS	(44)
Inflammation markers     a-1-acid glycoprotein (AGP)     XP     CPBA     (46)       Creactive protein (CRP)     C-reactive protein (CRP)     (47)     (47)       TNF-a, IE-A, IL-A, IL-B, IL-6, IL-10, IL-33,     X     XP     (48)       Iron status     Soluble transferrin receptors     XP     (49)       Iron status     Soluble transferrin receptors     XP     (49)       Hormones     Leptin, insulin, and adiponectin     X     XP     (49)       Genetics     Single nucleotide polymorphism     XB     (48)     (49)       Microbial community     X     XP     (48)     (51)       Microbiane     Microbial community     XB     X     (48)       Microbiane     Microbial community     X     (48)       Microbiane     Microbial community     XB     (48)       Analyzed in "Buify cost," Buify cost," Plasma, RBCs, WNRNA     XRNA     (53)	Metabolomics	Biocrates MxP® QUANT 500 <sup>3</sup>	×	Ч×			UPLC-MS/MS	(45)
C-reactive protein (CR <sup>P</sup> ) LI-16, IL-4, IL-8, IL-6, IL-10, IL-33, X X P P Plates Plat	Inflammation markers	lpha-1-acid glycoprotein (AGP)		Х <sup>Р</sup>			CPBA	(46)
IL-1β, IL-4, IL-6, IL-10, IL-33,       X       MSD immunoassay       (48)         TNF-a, IFN-y       TNF-a, IFN-y       CPBA       (49)         Ton status       Soluble transferrin receptors       XP       CPBA       (49)         Ferritin       Soluble transferrin receptors       XP       CPBA       (49)         Hormones       Leptin, insulin, and adiponectin       X       X       (49)         Genetics       Single nucleotide polymorphism       X       X       (48)         Genetics       Single nucleotide polymorphism       X       X       (51)         CpG methylation patterns       XB       GWAS       (52)         Microbiome       Microbial community       X       X       Shotgun metagenomics       (52)         Transcriptomics       mRNA transcriptome       XRNA       X       Shotgun metagenomics       (55, 56)         Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dired blood spots, <sup>P</sup> Plasma, <sup>R</sup> BCS, <sup>RNA</sup> RNA       Analyzed in: <sup>Plant</sup> Analyzed in:		C-reactive protein (CRP)						(47)
TNF-w, IFN-y       TNF-w, IFN-y       plates         Iron status       Soluble transferrin receptors       XP       Plates       (49)         Hormones       Leptin, insulin, and adiponectin       X       NSD immunoassay       (50)         Hormones       Leptin, insulin, and adiponectin       X       NSD immunoassay       (50)         Genetics       Single nucleotide polymorphism       XB       Plates       (48)         Microbiome       Microbial community       XB       Plates       (51)         Microbial community       XB       YB       FWAS       (52)         Transcriptomics       mRNA transcriptome       XRNA       (55, 56)       (55, 56)		IL-1 <i>β</i> , IL-4, IL-8, IL-6, IL-10, IL-33,	×	×			MSD immunoassay	(48)
Iron status Soluble transferrin receptors XP CPBA (49) Ferritin Erritin Erritin Karalin, and adiponectin X X MSD immunoassay (48) Hormones Leptin, insulin, and adiponectin X 2B (50) Genetics Single nucleotide polymorphism XB CPBA (48) Genetics Single nucleotide polymorphism XB CPBA (51) CpG methylation patterns XB CPBA (51) CpG methylation patterns XB CPBA (51) EWAS (51) EWAS (52) Microbiome XRNA transcriptome XRNA Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dired blood spots, <sup>P</sup> plasma, <sup>R</sup> BCS, <sup>RNA</sup> ANA sequencing (53) Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dired blood spots, <sup>P</sup> plasma, <sup>R</sup> BCS, <sup>RNA</sup> ANA sequencing (53) Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dired blood spots, <sup>P</sup> plasma, <sup>R</sup> BCS, <sup>RNA</sup> ANA sequencing (53) Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dired blood spots, <sup>P</sup> plasma, <sup>R</sup> BCS, <sup>RNA</sup> ANA ANA Sequencing (53) Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dired blood spots, <sup>P</sup> plasma, <sup>R</sup> BCS, <sup>RNA</sup> ANA ANA ANA ANA ANA ANA ANA ANA ANA		TNF- $\alpha$ , IFN- $\gamma$					plates	
Ferritin       Formunoassay       (50)         Hormones       Leptin, insulin, and adiponectin       X       X       MSD immunoassay       (48)         Hormones       Leptin, insulin, and adiponectin       X       X       MSD immunoassay       (48)         Genetics       Single nucleotide polymorphism       X <sup>B</sup> CPG methylation patterns       (51)       EWAS       (51)         Microbiome       Microbial community       X <sup>B</sup> X       16s RNA sequencing       (53)         Transcriptomics       mRNA transcriptome       X <sup>RNA</sup> X       Shotgun metagenomics       (54)	Iron status	Soluble transferrin receptors		Х <sup>Р</sup>			CPBA	(49)
Hormones     Leptin, insulin, and adiponectin     X     MSD immunoassay     (48)       Genetics     Single nucleotide polymorphism     XB     plates     (51)       Genetics     Single nucleotide polymorphism     XB     EWAS     (51)       Genetics     Single nucleotide polymorphism     XB     EWAS     (51)       CpG methylation patterns     XB     EWAS     (52)       Microbiome     Microbial community     X     16s RNA sequencing     (53)       Transcriptomics     mRNA transcriptome     X <sup>RNA</sup> Shotgun metagenomics     (54)		Ferritin						(20)
Genetics     Single nucleotide polymorphism     XB     plates       GWAS     GWAS     GWAS       CpG methylation patterns     EWAS     (51)       Microbiome     Microbial community     X     16s RNA sequencing     (53)       Transcriptomics     mRNA transcriptome     X <sup>RNA</sup> (54)       Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dried blood spots, <sup>P</sup> plasma, RBCs, <sup>RNA</sup> RNA.     Microbiol     (55, 56)	Hormones	Leptin, insulin, and adiponectin	×	×			MSD immunoassay	(48)
Genetics     Single nucleotide polymorphism     X°     GMAS     (51)       CpG methylation patterns     X°     EWAS     (52)       Microbiome     Microbial community     X     16s RNA sequencing     (53)       Transcriptomics     mRNA transcriptome     X <sup>RNA</sup> (54)       Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dried blood spots, <sup>P</sup> plasma, <sup>R</sup> BCs, <sup>RNA</sup> RNA.     CPA     (55, 56)		-		q			plates	:
CpG methylation patterns     X <sup>B</sup> EWAS     (52)       Microbial community     X     16s RNA sequencing     (53)       Transcriptomics     RNA transcriptome     (54)       Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dried blood spots, <sup>P</sup> plasma, <sup>R</sup> RBCs, <sup>RNA</sup> RNA.     X <sup>RNA</sup> (54)	Genetics	Single nucleotide polymorphism		Ч			GWAS	(51)
Microbiome     Microbial community     X     16s RNA sequencing     (53)       Transcriptomics     RNA transcriptome     X     Shotgun metagenomics     (54)       Transcriptomics     RNA sequencing     (55, 56)     (54)       Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dried blood spots, <sup>P</sup> plasma, <sup>R</sup> RBCs, <sup>RNA</sup> RNA.     (55, 56)     (55, 56)		CpG methylation patterns		X <sup>B</sup>			EWAS	(52)
Transcriptomics       X       Shotgun metagenomics       (54)         Transcriptomics       mRNA transcriptome       X <sup>RNA</sup> (54)         Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dried blood spots, <sup>P</sup> plasma, <sup>R</sup> RBCs, <sup>RNA</sup> RNA.       (55, 56)	Microbiome	Microbial community				×	16s RNA sequencing	(23)
Transcriptomics mRNA transcriptome X <sup>RNA</sup> (55, 56) Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dried blood spots, <sup>P</sup> plasma, <sup>R</sup> RBCs, <sup>RNA</sup> RNA.						×	Shotgun metagenomics	(54)
Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> dried blood spots, <sup>P</sup> plasma, <sup>R</sup> RBCs, <sup>RNA</sup> RNA.	Transcriptomics	mRNA transcriptome	X <sup>RNA</sup>				RNA seq	(55, 56)
References that describe methods used for the analysis or on which the nonpublished methods are based on: UPBA, competitive protein binding assay, EVVAS, epidenome-wide association; FLU, fluorescence	Analyzed in: <sup>B</sup> buffy coat, <sup>DBS</sup> , <sup>1</sup> References that describe m	dried blood spots, <sup>p</sup> plasma, <sup>R</sup> RBCs, <sup>RNA</sup> RNA. ethods used for the analysis or on which the nonbub	olished methods are	e based on: CPBA, c	ompetitive protein k	oinding assay; EV	VAS, epigenome-wide association; F	LD, fluorescenc
detection; GWAS, genome-wide associations; ICP, inductively coupled plasma MS/MS; MSD, Meso Scale Discovery; MWL, multi-wavelength detection; NIR, near infra red; QqQ-MS, triple quad MS; TOF, time of flight;	detection: GWAS denome-w							

UPLC, ultra-performance-LC. <sup>2</sup>Plasma folate is analyzed by CPBA.

<sup>3</sup>https://lebrilla.faculty.ucdavis.edu/research/nutritional-glycomics/.

<sup>4</sup>https://biocrates.com/mxp-quant-500-kit/.

TABLE 5 Required	sample	sizes	for testing	equival	ence
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Differences (site means)	Sample size/site 80% power	Sample size/site 80% power
0.20 SD	310	429
0.25 SD	199	275
0.30 SD	139	191

provide adequate power to determine whether 1 site is different from the others (**Table 5**).

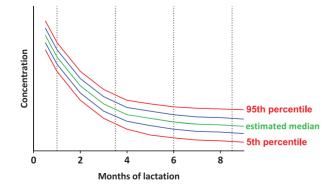
#### Sample size estimates for other analyses.

For some secondary outcome variables, e.g. free amino acids (FAA) in infant plasma, a complete sample set for analyses is not feasible or available. In such cases, only 100 samples for each site and time point are analyzed, assuming a priori that the values are similar enough to pool between  $\geq 2$  of the sites. If this is not possible, within-site percentiles will be presented with a caveat that these should not be considered as RVs, as precision is limited. However, 100 per time point is ample for looking at relations between secondary outcome variables and other outcomes, such as growth. Nevertheless, banked samples are available if initial data analyses show that analysis of >100 samples per point is needed.

Future exploratory analyses will also assess the relations between milk nutrient concentrations and other collected information, including milk volume, child growth, child developmental milestones, and maternal and infant plasma values. A sample size of 140 paired data points per site and time period is adequate to detect a correlation between continuous variables of 0.30 or higher with 95% power within each site, assuming a 2-sided alternative hypothesis, and therefore the planned sample size of 250 per site and time point is more than sufficient for these analyses.

# Data analysis plan.

The measured human milk nutrient concentrations and milk volume will be used to estimate daily nutrient intakes of the infants. The RVs will be based on these infant nutrient intakes, and developed and expressed as percentiles in these well-nourished, but nonsupplemented, population groups, following the principles used by the Intergrowth-21st Project (21), which are based on methods developed in the construction of the WHO Child Growth Standards (22) (illustrated in Figure 3). Cri-



**FIGURE 3** Hypothetical graphical illustration of reference values that may result for a micronutrient.

teria will be created for normal growth and development, and adequate nutritional status for each nutrient in question. Data from children who do not meet these criteria will be removed from the construction of the RVs.

For each nutrient and time period, the distribution of the human milk variable will be examined and extreme outliers will be investigated and, if needed, removed. Box–Cox and other transformations will be used to normalize each variable, preferably using the same transformation across study sites, and the Shapiro–Wilk statistic to assess normality; skewness, kurtosis. The general shape of the distribution will also be examined. Parameters will be compared among study sites as described above, and information from the sites will be combined if the sites are deemed to be equivalent, or if differences have no material effect on key percentiles; otherwise, different sets of RV centiles may be constructed per site, or with only 2 or 3 pooled sites.

Parameters from the identified distribution will be estimated from the data, and centiles of interest (5th, 25th, 50th, 75th, 95th) will be estimated for each nutrient based on the theoretical percentiles of the underlying distribution, as described above. Smoothing techniques such as cubic splines will be employed, assuming the centiles follow smooth and continuous functions. Tables and curves of estimated centiles will be created for each nutrient and time period. SAS for Windows Release 9.4 (SAS Institute) will be used for all analyses.

## Additional study outcomes

Besides the primary study outcome, the construction of the RVs, secondary study outcomes are included in the study design and categorized into outcomes that: 1) will be used for the construction of the RVs, 2) will not be used to construct RVs, and 3) country-specific measurements that are not used for RV development. These outcomes are summarized by category in **Table 6**.

# Ethics, dissemination, and trial status

Full ethical approvals at all study sites were obtained from:

- The Institutional Review Board of the University of California, Davis, CA, USA (IRB ID: 920618–1, Protocol HRP-503-MILQ IRB, Department of Health and Human Services FWA No: 00004557).
- The Internal Review Boards of the International Centre for Diarrhoeal Disease Research, Bangladesh (icddr, b; PR-17085).
- The National Commission for Research Ethics (2.086.708, 2.875.218, 4.865.685), the Research Ethics Committees of the Municipal Secretariat of Health and Civil Defense of the State of Rio de Janeiro and of the Maternity School of Rio de Janeiro Federal University, (1.948.992, 2.769.611, 4.449.007); and the Municipal Secretary of Health and Civil Defense of the State of Rio de Janeiro (2.100.255), Brazil; Project number: 64767717.4.0000.5275.
- The Committees on Biomedical Research Ethics for the Capital Region of Denmark (H-17015174).
- The joint Gambia Government/MRC The Gambia Ethics Committee (SCC 1572v1.1, Project ID/ethics ref: 22768).

The MILQ Study was registered at clinicaltrials.gov as NCT03254329 (18 August, 2017). Study progress was discussed among all field sites and the USDA/ARS-WHNRC, the main analytical site, and location of the Principal Investigator (PI), in biweekly

Outcome category <sup>1</sup>	Bangladesh	Brazil	Denmark	The Gambia
Used for	Macronutrients,	human milk oligosaccharides, p	eptides, and proteins, metabolo	mics
constructing RVs	Micronuti	ient status of mothers and infar	nts assessed in blood samples	
	Milk volume	by D <sub>2</sub> O	24-h infant test weighing	Milk volume by $D_2O$
	lo	dine and maternal and infant ic	odine status biomarker	
Not used for		Maternal and infant die	etary intakes	
constructing RVs		Maternal and infant an	thropometry	
		Materna	al body composition (bioimpedar	nce)
			ition (bioimpedance)	Infant body composition (air displacement plethysmography)
		Maternal and infant	morbidity	
		notor development assessment		
	Infant devel	opment using the Ages and Sta Infant microbic	iges questionnaire (6–8.5 mo pp) ome	
	Mat	ternal inflammatory markers (28	–30 weeks of gestation)	
Country-specific		Maternal nutrient intake a	t 35–37 weeks of gestation	
outcomes		Maternal microbiome <sup>3</sup>		Maternal microbiome <sup>3</sup>
			Maternal hemoglobin A1c <sup>4</sup>	
			Maternal markers of insulin sensitivity <sup>5</sup> Maternal lipid panel <sup>6</sup>	

## TABLE 6 Secondary outcomes of the Mothers, Infants, and Lactation Quality (MILQ) Study

<sup>1</sup>Unless otherwise stated, the time periods for the secondary outcome and analysis objectives are identical to those for the primary outcome (1–3.4, 3.5–5.9, 6–8.5 mo postpartum).

<sup>2</sup>Based on WHO's motor milestones.

<sup>3</sup>28–40 weeks of gestation, and 1 to 8.5 mo pp.

<sup>4</sup>28–30 weeks of gestation, measured in whole blood.

<sup>5</sup>28–30 weeks of gestation: insulin, C-peptide, and leptin (plasma).

<sup>6</sup>28–30 weeks of gestation: total cholesterol, HDL-cholesterol, LDL-cholesterol, very LDL-cholesterol, and triglycerides (plasma). D<sub>2</sub>O, deuterated water; pp, postpartum, RV, reference value.

conference calls. Updates of the study are provided monthly in conference calls between the PI and the funding agency, which are reported back to the field sites. Results are presented at national and international nutrition-related conferences, and in peer-reviewed journals. MILQ meetings will be hosted at 1 study site each year. Funding has been secured for students and researchers to visit other study sites for professional development. Deidentified data will be available in a public repository after full publication of the primary study outcomes (RVs) for milk MNs and country-specific approvals. The recruitment phase started between September 2017 (Denmark) and May 2018 (The Gambia) (Table 7). The analytical work began in May 2019 and is expected to be completed by December 2022.

## Discussion

Although the importance of human milk in maternal and infant nutrition has long been recognized, the MN intake recommendations for breastfeeding infants and lactating women are still based on old, uncertain, or even incorrect data (4) and are established mainly as Adequate Intakes for infants due to the lack of experimentally determined estimates of milk composition and nutrient requirements (23). Further, the dynamic changes in concentrations of many milk nutrients across lactation are not reflected in the current recommendations, which can lead to erroneous status evaluations (24, 25). Therefore, given the importance of adequate nutrient intake during the first 1000 d of life (26), which includes the WHO-recommended 6-mo period of EBF (1), this systematic, longitudinal assessment of the nutrient composition of human milk, in particular for MNs, using validated methodologies, is indispensable for correctly evaluating nutrient intake adequacy of breastfeeding infants.

We have developed and validated multiple novel methods for collecting and analyzing human milk (24, 27–30), which will be used or further optimized for the nutrient assessment in this study. The identical collection protocols and methods for analyzing human milk and blood samples minimize method bias and enable direct comparison between field sites. The inclusion and exclusion criteria, and collection of information on diets, anthropometry, and infant developmental milestones provide supportive information that participants are healthy and nutrient sufficient.

The 4 field sites are different in demographics, socioeconomic status, diets, and cultural practices which may preclude combining data across sites to establish the RVs, but the required number of participants per site allows for the creation of study-site-specific RVs if necessary.

# TABLE 7 Trial status

	Bangladesh	Brazil	Denmark	The Gambia	USA <sup>1</sup>
Recruitment start	04/2018	01/2018	09/2017	05/2018	_
Completion of field site work	Expected by 03/2022	Expected by 03/2022	12/2019	Expected by 03/2022	—
Sample analyses start	_	_	_	_	05/2019
Completion of sample analyses	—	—	—		Expected by 12/2022

<sup>1</sup>Additional analytical sites are Bengaluru, India (milk volume); Cambridge, UK (vitamin D in human milk); Zürich, Switzerland (iodine and iodine status in milk, dried blood spots, and urine).

The longitudinal data obtained by the staggered sample collection protocol will allow for the development of new dynamic RVs, providing percentile ranges for every stage of lactation covered in this study. These carefully developed RVs, in conjunction with the milk volume data, will be available for re-evaluating intake recommendations for infants, young children, and lactating women, and as benchmarks against which to evaluate human milk quality, and the effects of future nutrient intervention trials on milk composition.

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The authors' responsibilities were as follows—LHA, DH, SS, AD, KGE, MI, GK, KFM, and SEM: designed the research; DH, SSF, EB, AD, DF, KGE, SHC, FKK, and FN: conducted the research; JMP: performed the statistical analysis; LHA and DH: wrote the manuscript; LHA: had primary responsibility for final content; and all authors: read and approved the final manuscript.

#### References

- World Health Organization. Guidelines on HIV and Infant Feeding 2010: Principles and Recommendations for Infant Feeding in the Context of HIV and a Summary of Evidence. Geneva (Switzerland): WHO, 2010.
- World Health Organization [Internet]. Geneva (Switzerland): Breastfeeding c2021[ cited 7 Aug, 2021]. Available from: https://www.who.int/health-topic s/breastfeeding#tab=tab\_2.
- Hampel D, Dror DK, Allen LH. Micronutrients in human milk: analytical methods. Adv Nutr 2018;9(suppl\_1):313S–31S.
- Allen LH, Dror DK. Introduction to current knowledge on micronutrients in human milk: adequacy, analysis, and need for research. Adv Nutr 2018;9(suppl\_1):275S–7S.
- 5. Allen LH. Maternal micronutrient malnutrition: effects on breast milk and infant nutrition, and priorities for intervention. SCN News 1994;11:21–4.
- Allen LH. B vitamins in breast milk: relative importance of maternal status and intake, and effects on infant status and function. Adv Nutr 2012;3(3):362–9.
- Dror DK, Allen LH. Retinol-to-fat ratio and retinol concentration in human milk show similar time trends and associations with maternal factors at the population level: a systematic review and meta-analysis. Adv Nutr 2018;9(suppl\_1):332S-46S.
- Dror DK, Allen LH. Vitamin B-12 in human milk: a systematic review. Adv Nutr 2018;9(suppl\_1):358S–66S.
- 9. Dror DK, Allen LH. Iodine in human milk: a systematic review. Adv Nutr 2018;9(suppl\_1):347S-57S.
- Allen LH, Graham JM. Assuring micronutrient adequacy in the diets of young infants. In: Delange FM, West KP, editors. Micronutrient Deficiencies

in the First Months of Life. Basel (Switzerland): Nestec Ltd./S. Karger AG, 2003. p. 55–88.

- Dror DK, Allen LH. Effect of vitamin B12 deficiency on neurodevelopment in infants: current knowledge and possible mechanisms. Nutr Rev 2008;66(5):250–5.
- Victora CG, de Onis M, Hallal PC, Blössner M, Shrimpton R. Worldwide timing of growth faltering: revisiting implications for interventions. Pediatrics 2010;125(3):e473–e80.
- 13. Gallant J, Chan K, Green TJ, Wieringa FT, Leemaqz S, Ngik R, Measelle JR, Baldwin DA, Borath M, Sophonneary P, et al. Low-dose thiamine supplementation of lactating Cambodian mothers improves human milk thiamine concentrations: a randomized controlled trial. Am J Clin Nutr 2021;114(1):90–100.
- 14. Eneroth H, El Arifeen S, Persson LÅ, Lönnerdal B, Hossain MB, Stephensen CB, Ekström EC. Maternal multiple micronutrient supplementation has limited impact on micronutrient status of Bangladeshi infants compared with standard iron and folic acid supplementation. J Nutr 2010;140(3):618–24.
- World Health Organization. The quality and quantity of breast milk: report on the WHO collaborative study on breast-feeding. Geneva (Switzerland) 1985.
- Dror DK, Allen LH. Overview of nutrients in human milk. Adv Nutr 2018;9(suppl\_1):278S-94S.
- Allen LH, Donohue JA, Dror DK. Limitations of the evidence base used to set recommended nutrient intakes for infants and lactating women. Adv Nutr 2018;9(suppl\_1):295S-312S.
- Hampel D, Allen LH. Analyzing B-vitamins in human milk: methodological approaches. Crit Rev Food Sci Nutr 2016;56(3):494–511.
- Squires J, Bricker D. Ages and stages questionnaire [Internet]. Third edition. 2009. Brookes Publishing. Available from: www.brookespublishing.com.
- Walker E, Nowacki AS. Understanding equivalence and noninferiority testing. J Gen Intern Med 2011;26(2):192–6.
- Altman D, Ohuma E, Fetal I, Century NGCfts. Statistical considerations for the development of prescriptive fetal and newborn growth standards in the INTERGROWTH-21st project. BJOG: An International Journal of Obstetrics & Gynaecology 2013;120:71–76.
- 22. Borghi E, de Onis M, Garza C, Van den Broeck J, Frongillo EA, Grummer-Strawn L, Van Buuren S, Pan H, Molinari L, Martorell R. Construction of the World Health Organization child growth standards: selection of methods for attained growth curves. Stat Med 2006;25(2):247–65.
- 23. Institute of Medicine. Dietary Reference Intakes for Thiamin, Riboflavin, Niacin, Vitamin B6, Folate, Vitamin B12, Pantothenic Acid, Biotin, and Choline. Washington (DC): National Academies Press, 1998.
- 24. Hampel D, Shahab-Ferdows S, Gertz E, Flax VL, Adair LS, Bentley ME, Jamieson DJ, Tegha G, Chasela CS, Kamwendo D, et al. The effects of a lipid-based nutrient supplement and antiretroviral therapy in a randomized controlled trial on iron, copper, and zinc in milk from HIV-infected Malawian mothers and associations with maternal and infant biomarkers. Maternal & Child Nutrition 2018;14(2):e12503.
- 25. Whitfield KC, Shahab-Ferdows S, Kroeun H, Sophonneary P, Green TJ, Allen LH, Hampel D. Macro- and micronutrients in milk from healthy Cambodian mothers: status and interrelations. J Nutr 2020;150(6):1461–9.
- Oshin O, Hampel D, Idachaba F, Atayero A. The first 1,000 days: trends towards biosensing in assessing micronutrient deficiencies. J Phys Conf Ser 2019;1299:012136.

- 27. Hampel D, York ER, Allen LH. Ultra-performance liquid chromatography tandem mass-spectrometry (UPLC-MS/MS) for the rapid, simultaneous analysis of thiamin, riboflavin, flavin adenine dinucleotide, nicotinamide and pyridoxal in human milk. J Chromatogr B 2012;903: 7–13.
- Hampel D, Shahab-Ferdows S, Domek JM, Siddiqua T, Raqib R, Allen LH. Competitive chemiluminescent enzyme immunoassay for vitamin B12 analysis in human milk. Food Chem 2014;153:60–65.
- Hampel D, Shahab-Ferdows S, Islam MM, Peerson JM, Allen LH. Vitamin concentrations in human milk vary with time within feed, circadian rhythm, and single-dose supplementation. J Nutr 2017;147(4):603–11.
- Hampel D, Shahab-Ferdows S, Hossain M, Islam MM, Ahmed T, Allen LH. Validation and application of Biocrates Absolute*IDQ*<sup>®</sup> p180 targeted metabolomics kit using human milk. Nutrients 2019;11(8): 1733.
- 31. Turner T, Burri BJ. Rapid isocratic HPLC method and sample extraction procedures for measuring carotenoid, retinoid, and tocopherol concentrations in human blood and breast milk for intervention studies. Chromatographia 2012;75(5–6):241–52.
- 32. Roche Diagnostics. Vitamin D total II for cobase 411 analyzer Mannheim (Germany): Roche Diagnostics GmbH, 2017.
- 33. Hampel D, Shahab-Ferdows S, Adair LS, Bentley ME, Flax VL, Jamieson DJ, Ellington SR, Tegha G, Chasela CS, Kamwendo D, et al. Thiamin and riboflavin in human milk: effects of lipid-based nutrient supplementation and stage of lactation on vitamer secretion and contributions to total vitamin content. PLoS One 2016;11(2):e0149479.
- 34. Roche Diagnostics. Elecsys folate III. Mannheim (Germany): Roche Diagnostics GmbH, 2018.
- 35. Roche Diagnostics. Vitamin B12 II. Mannheim (Germany): Roche Diagnostics GmbH, 2017.
- 36. Hampel D, Shahab-Ferdows S, Nguyen N, Kac G, Allen LH. High-throughput analysis of water-soluble forms of choline and related metabolites in human milk by UPLC-MS/MS and its application. Frontiers in Nutrition 2021;7:604570.
- Pedersen TL, Keyes WR, Shahab-Ferdows S, Allen LH, Newman JW. Methylmalonic acid quantification in low serum volumes by UPLC-MS/MS. J Chromatogr B 2011;879(19):1502–6.
- Gilfix BM, Blank DW, Rosenblatt DS. Novel reductant for determination of total plasma homocysteine. Clin Chem 1997;43(4):687–8.
- Astolfi ML, Marconi E, Protano C, Vitali M, Schiavi E, Mastromarino P, Canepari S. Optimization and validation of a fast digestion method for the determination of major and trace elements in breast milk by ICP-MS. Anal Chim Acta 2018;1040:49–62.
- 40. Dold S, Baumgartner J, Zeder C, Krzystek A, Osei J, Haldimann M, Zimmermann M, Andersson M. Optimization of a new mass spectrometry method for measurement of breast milk iodine concentrations (BMIC) and an assessment of the effect of analytic method and timing of within-feed sample collection on BMIC. Thyroid 2016;26(2):287–95.

- 41. Stinca S, Andersson M, Weibel S, Herter-Aeberli I, Fingerhut R, Gowachirapant S, Hess SY, Jaiswal N, Jukić T, Kusic Z. Dried blood spot thyroglobulin as a biomarker of iodine status in pregnant women. The Journal of Clinical Endocrinology & Metabolism 2017;102(1):23–32.
- 42. Unity Scientific. Neonatal Analyzer Package. Milford (MA): Unity Scientific; 2017.
- 43. Wu LD, Ruhaak LR, Lebrilla CB. Analysis of milk oligosaccharides by mass spectrometry. In: Lauc G, Wuhrer M editors. High-Throughput Glycomics and Glycoproteomics. New York: Springer, 2017. p.121–9.
- 44. Huang J, Kailemia MJ, Goonatilleke E, Parker EA, Hong Q, Sabia R, Smilowitz JT, German JB, Lebrilla CB. Quantitation of human milk proteins and their glycoforms using multiple reaction monitoring (MRM). Anal Bioanal Chem 2017;409(2):589–606.
- 45. Hampel D, Shahab-Ferdows S, Kac G, Allen LH. Human milk metabolic profiling using biocrates mxp<sup>®</sup> quant 500 kit. Current Developments in Nutrition 2021;5(Supplement\_2):874.
- 46. Roche Diagnostics. AAGP2-Tina-quant alpha-acid glycoprotein Gen.2. Mannheim (Germany): Roche Diagnostics GmbH; 2017.
- 47. Roche Diagnostics. CRPHS-Cardiac C-reactive protein (Latex) high sensitivity. Mannheim (Germany): Roche Diagnostics GmbH; 2016.
- mesoscale.com [Internet]. Rockville (MD): Meso Scale Diagnostics, LLC; c2021[ cited 4 Aug, 2021]. Available from: https://www.mesoscale.com/.
- 49. Roche Diagnostics. STFR-Tina-quant Soluble Transferrin Receptor. Mannheim (Germany): Roche Diagnostics GmbH; 2021.
- 50. Roche Diagnostics. Elecsys® Ferritin. Rotkreuz (Switzerland): Roche Diagnostics International Ltd; 2014.
- 51. Marees AT, de Kluiver H, Stringer S, Vorspan F, Curis E, Marie-Claire C, Derks EM. A tutorial on conducting genome-wide association studies: quality control and statistical analysis. Int J Methods Psychiatr Res 2018;27(2):e1608.
- 52. Kim M, Costello J. DNA methylation: an epigenetic mark of cellular memory. Exp Mol Med 2017;49(4):e322.
- 53. Hald S, Schioldan AG, Moore ME, Dige A, Lærke HN, Agnholt J, Bach Knudsen KE, Hermansen K, Marco ML, Gregersen S. Effects of arabinoxylan and resistant starch on intestinal microbiota and short-chain fatty acids in subjects with metabolic syndrome: a randomised crossover study. PLoS One 2016;11(7):e0159223.
- Treiber ML, Taft DH, Korf I, Mills DA, Lemay DG. Pre- and post-sequencing recommendations for functional annotation of human fecal metagenomes. BMC Bioinformatics 2020;21(1):1–15.
- 55. Lemay DG, Hovey RC, Hartono SR, Hinde K, Smilowitz JT, Ventimiglia F, Schmidt KA, Lee JW, Islas-Trejo A, Silva PI. Sequencing the transcriptome of milk production: milk trumps mammary tissue. BMC Genomics 2013;14(1):1–17.
- 56. Westreich ST, Ardeshir A, Alkan Z, Kable ME, Korf I, Lemay DG. Fecal metatranscriptomics of macaques with idiopathic chronic diarrhea reveals altered mucin degradation and fucose utilization. Microbiome 2019;7(1):1–17.