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## Pomegranate juice alters the microbiota in breast milk and infant stool: a pilot study†

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Pomegranate juice (PomJ) contains ellagitannins (ETs) that are metabolized to ellagic acid (EA). Intestinal bacteria convert EA further to urolithins that are absorbed into the circulation and may provide health benefits. PomJ consumption by pregnant women was reported to be neuroprotective for their infants. In order to determine whether EA and metabolites are transferred from breast milk of mothers consuming PomJ to nursing infants, we performed an interventional pilot study and enrolled ten healthy women with full-term, exclusively breast-fed infants, consuming 8 oz. of PomJ daily for two weeks. Breast milk, plasma, urine and stool samples were collected from the mothers and the urine and stool samples from the infants before and after two weeks of PomJ consumption. Samples were analyzed using liquid chromatography-mass spectrometry to identify EA metabolites and 16S rRNA sequencing to determine changes in the microbiota. EA metabolite conjugates (dimethyl EA-glucuronide DMEAG and urolithin A-glucuronide UAG) were found in breast milk, plasma and urine from mothers and in urine of infants after 14 days of PomJ consumption. In addition, urolithin B-glucuronide (UBG) was found in breast milk, plasma and urine from two participants and urine from their infants. PomJ consumption was associated with a significant decrease in breast milk of *Lactococcus*, *Subdoligranulum*, and *Acinetobacter*, while the abundance of Firmicutes/*Faecalibacterium* increased significantly. In breast milk *Escherichia/Shigella* was inversely correlated to breast milk UAG. In infant stools, the abundance of *Lachnospirillum* and *Staphylococcus* was increased. Infant stool *Blautia* was positively correlated to breast milk and mother plasma UBG. This pilot study demonstrates that EA and its metabolites are absorbed by the nursing infant from breast milk, excreted in urine and impact the infant gut microbiome. The concentration of EA metabolites in breast milk increased over time. Phenolic compounds in breast milk could be a way to promote neuroprotective, antioxidant and anti-inflammatory health benefits in infants.

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### 1. Introduction

Pomegranate fruits have been used for centuries in ancient cultures for their medicinal purposes. The pomegranate antioxidant and anti-inflammatory effects led to studies of dietary supplementation with pomegranate juice (PomJ). These studies have shown protective effects of PomJ on cardiovascular disease, inflammatory bowel disease and prostate cancer.<sup>1–3</sup>

Most health benefits of PomJ have been attributed to the presence of ellagitannins (ETs), mainly punicalagins and ellagic acid (EA).<sup>4,5</sup> ETs are hydrolyzed in the intestine and absorbed as EA over six hours after administration of PomJ.<sup>6–8</sup>

After PJ consumption, the remaining ETs and EA are retained unabsorbed in the gut lumen where they interact with the complex intestinal bacterial microflora to form a family of urolithins in the gut.<sup>9,10</sup> There are inter-individual variations of ET metabolism producing either urolithin A (UA), urolithin B (UB), isourolithin or no urolithins (*i.e.* non-producers).<sup>11–13</sup> The term “metabotypes” is used to describe the differences among individuals in ET metabolism and these differences have also been related to biomarkers of risk for cardiovascular disease and other chronic diseases of aging.<sup>14,15</sup> These differences in metabolism have been associated with differences in the population of intestinal microflora.<sup>11</sup> There is growing interest in the potential of dietary supplements for the protection of placental oxidative stress and neuroprotection in the brain. In a study by Chen *et al.*, PomJ supplementation before delivery has been found to reduce oxidative stress and stimulus-induced apoptosis in the placenta.<sup>16</sup> In addition, supplementation of pomegranate in the drinking water of pregnant and nursing rodent dams was found to significantly

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decrease focal brain ischemia and periventricular white matter injury in either inflammatory or hypoxic-ischemic models.<sup>17–22</sup>

In a small pilot study, we found that EA and UA are present in breast milk and a study by Cortes-Martin *et al.* found urolithins in breast milk formed from mothers consuming walnuts.<sup>23,24</sup> As yet, whether pomegranate metabolites are absorbed by nursing infants and whether the metabolite of the mother is transferred to the infant has not been studied. The present pilot study has been designed to determine whether consumption of PomJ by the nursing mother will increase ET metabolites in the infants' urine and whether changes in the mothers' milk and stool microbiota are related to changes in the infants' stool microbiota.

## 2. Materials and methods

### 2.1. Study design

This was an interventional, open label pilot study. Twelve women and infants were recruited at the UCLA Center for Human Nutrition (Fig. 1). The clinical protocol was approved by the Institutional Review Board. The study was registered at clinicaltrials.gov (NCT04341961). All subjects gave written informed consent prior to the start of the study. The mothers were asked to consume 8 oz of PomJ daily for 2 weeks. A dose of 8 oz of PomJ was selected based on previous studies demonstrating health beneficial effects such as antioxidant, cholesterol lowering, anti-inflammatory and delaying prostate cancer recurrence resulting from consumption of 8 oz of PomJ or supplement with equivalent polyphenol content.<sup>3,25</sup> Dietary guidelines recommend consuming no more than 1 cup of juice daily.

At baseline and after 14 days of PomJ consumption, a blood sample, breast milk sample, 6 and 24 hours urine sample after PomJ consumption and stool sample were obtained from the mother and a spot urine and stool samples were collected

from the infants (Fig. 1). Urolithins and EA levels were determined in the maternal plasma, breast milk and urine, and in the infant urine. The stool microbiome was analyzed in breast milk and in the maternal and infant stool samples. Two weeks prior to the baseline visit, participants were asked to consume a low polyphenol diet. The low polyphenol diet includes no more than 3 servings of polyphenol rich fruits/vegetables and no more than 10 g of fiber. A list of foods containing high fiber and phenolic compounds was provided.

Following the pre-study washout phase, mothers consumed 8 oz of PomJ (PomWonderful Co, Los Angeles, CA) daily for two weeks. Participants were asked to return the unused study product to assess compliance. On the first day of PomJ consumption, mothers also collected their breast milk at 6 and 24 hours after consuming the first dose of PomJ. On the last day of PomJ consumption breast milk samples were collected at the same times. Stool samples were obtained from the mother and infants prior to PJ consumption and on the last day of PomJ consumption.

### 2.2. Participants

Healthy women who were postpartum less than 6 months with a full term baby born vaginally and who were exclusively breast feeding were enrolled in the study. Exclusion criteria included: (a) antibiotics or laxatives use during the 2 months before the study; (b) consuming pre- or probiotics or anti-inflammatory medication; (c) screening laboratory values outside of the normal range that were considered clinically significant for study participation by the investigator; and (d) allergy to pomegranate. There was one screen failure and one drop out for unknown reasons, and ten women completed the study.

### 2.3. Urine collection for infants

In female infants, the area around the urethra was thoroughly washed with soap or cleansing wipes. A plastic bag with an adhesive strip on one end was placed around the genital area.

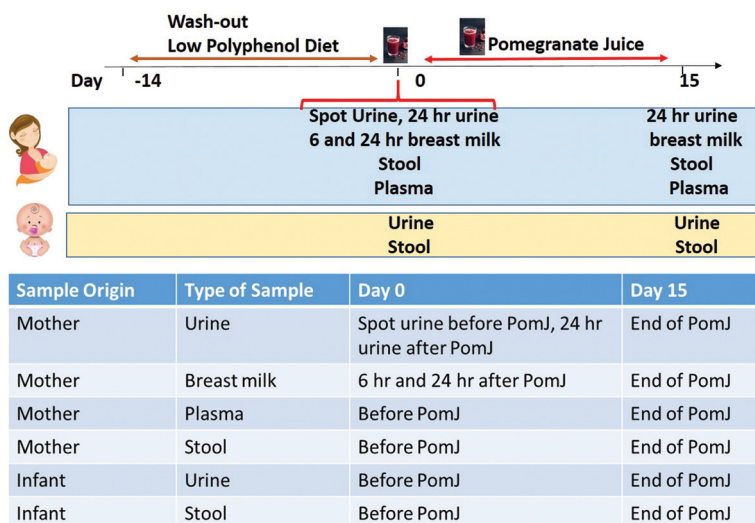


Fig. 1 Study design and samples collected.



For male infants, the bag was placed over the entire penis and the adhesive was attached to the skin. For females, the bag was placed over the labia on both sides. A diaper was placed over the urine collection bag to help prevent dislocation of the urine bag if the infant is active. Once the infant voided, the bagged specimen was transferred into a collection container for sample evaluation.

#### 2.4. Stool collection

An aliquot of a daily stool specimen was collected by the mothers using a standard procedure and stool from the infant diapers was collected. Stool samples were refrigerated and delivered to the UCLA Center for Human Nutrition in a cooler with ice pack (~4 °C) no later than 12 hours after collection. At the laboratory, stool was aliquoted into smaller vials and frozen immediately and stored at -80 °C. Approximately 1 g of stool was weighed and dried in a vacuum drying oven (15 in Hg) at 80 °C for 48 hours, then weighed again to establish the moisture content so that all counts could be corrected to dry weight.

#### 2.5. Reagents and instruments

All solvents were HPLC grade and purchased from Fisher Scientific (Tustin, CA, USA).

#### 2.6. Pomegranate metabolites in breast milk, plasma and urine

1 mL urine samples were processed as previously reported.<sup>6</sup> 5 mL of breast milk was mixed with 20 mL acetonitrile and mixed by vortex. The mixture was centrifuged at 4000 rpm for 10 min. The acetonitrile phase was dried using vacuum (speed-vac) and reconstituted in 1 mL water, and further purified using solid-phase extraction (SPE) on C18 cartridges (Waters WAT 036945) and eluted with 2 mL methanol. The elutant was placed in the SpeedVac to dry, reconstituted in 200 µL methanol:water 1:1 (v:v), and analyzed for ETs and ET metabolites, dimethylellagic acid glucuronide (DMEAG) and urolithin A-glucuronide (UAG) and urolithin B-glucuronide (UBG) by LC-MS/MS system to determine the presence and levels of as previously reported.<sup>6</sup> The concentrations were calculated based on UAG and UBG standards (Toronto Research Chemicals, Canada).

#### 2.7. Fecal 16S rRNA gene sequencing and taxonomic analysis

DNA extraction and sequencing of the 16S rRNA gene were performed as previously described by UCLA Microbiome Core.<sup>26</sup> Briefly, DNA from stool was extracted using the DNeasy power soil DNA isolation kit with bead beating (Qiagen, Valencia, CA) and from breast milk using the Qiagen DNeasy PowerFood Microbial Kit (Qiagen, Valencia, CA). For stool and breast milk samples, the 16S rRNA gene V4 variable region was amplified and barcoded using F515/R806 primers followed by 250 × 2 bp sequencing on an Illumina HiSeq 2500.<sup>26</sup>

Amplicon sequence variants (ASVs) were identified using DADA2 and annotated them against the SILVA v138 database. Alpha diversity metrics (Chao1 and Shannon index) were calculated after rarefaction to a depth of 15 000 sequences per sample and tested for significance using the Wilcoxon signed

ranked test at  $p < 0.05$ . Beta-diversity was calculated using Bray–Curtis dissimilarity. The relationships of samples across groups were determined by Permutational Multivariate Analysis of Variance (PERMANOVA) using the Adonis command provided by Vegan in R and were displayed via Principal Coordinate Analysis (PCoA) ordination.<sup>27</sup> DESeq2 was used to identify abundance changes at the genus level and differences that occurred between day 15 and day 29 samples following the PomJ intervention. *P*-Values were adjusted for multiple testing using Benjamini Hochberg false discovery rate correction in DESeq2. As Bonferroni correction is often considered overly conservative, we listed genera with *p*-value < 0.05 and marked those with adjusted *p* < 0.05 using \*. Spearman rank correlations were calculated and plotted as a heatmap between the Pom metabolites (breast milk and blood) and bacteria (breast milk and infant stool) at day 15 using R (version 3.5.2).<sup>28</sup> \*\*\*  $p < 0.001$ , \*\*  $p < 0.01$ , \*  $p < 0.05$ .

#### 2.8. Statistical analysis

This is the first study to examine the bioavailability of pomegranate metabolites in breast milk and potential impact on the baby. For sample size justification, we considered the formation of metabolites. Our published pilot study in 2 individuals demonstrated that breast milk concentration of dimethyl ellagic acid glucuronide (DMEAG) increased from  $0.2 \pm 0.3$  to  $0.7 \pm 0.3$  (nmol L<sup>-1</sup>) and urolithin A glucuronide (UAG) increased from  $3.0 \pm 4.2$  to  $28.9 \pm 10.4$  (nmol L<sup>-1</sup>) after two weeks of 8 oz. of PomJ consumption daily.<sup>23</sup> Assuming a correlation of 0.5 between baseline and 2 weeks PomJ consumption, a sample of sensitivity analysis demonstrated that  $N = 12$  participants would provide >80% power to detect the change in DMEAG and UAG.

Summary statistics (mean, SD, IQR and frequency distribution) and graphs were generated for baseline demographics and PomJ metabolites in mother urine, plasma, breast milk and infant urine using Graph Pad Prism 6 (San Diego, CA).

### 3. Results

#### 3.1. Demographic data

The demographic data are shown in Table 1. The maternal age ranged from 30–38 years. The infants' ages at the time of

**Table 1** Characteristics of the study participants

Baseline demographics	
Mother age (year)	34 ± 2.8
Infant age at start of PJ (month)	3.7 ± 1.4
Race/ethnicity of mother	
Asian	17%
Black/AA	0
White	83%
Hispanic	17%
Non-hispanic	83%

Data are means ± SD;  $N = 12$ .



screening ranged from 1 to 6 months. All mothers were in excellent health without any chronic medical conditions. All infants were delivered vaginally at an academic medical center without any complications. Compliance with PomJ consumption was 99%. Ten mothers with infants completed the study.

### 3.2. Pomegranate metabolite concentration

Following the low polyphenol-diet washout phase and prior to the first 8 oz of PomJ consumption, small amounts of Pom metabolites UAG and DMEAG were found in urine from 3 mothers (Fig. 2A). At baseline, 24 hours after the first 8 oz of PomJ consumption, urine samples from all mothers contained UAG (IQR:  $12.8 \mu\text{mol L}^{-1}$  [ $0.8\text{--}22.0 \mu\text{mol L}^{-1}$ ]) and DMEAG  $\mu\text{mol L}^{-1}$  (IQR:  $0.07 \mu\text{mol L}^{-1}$  [ $0.02\text{--}0.48 \mu\text{mol L}^{-1}$ ]) (Fig. 2B) and 24 h urine samples from two mothers contained UBG (Fig. 2B). After 14 days of consumption all Pom metabolite concentrations were increased compared to the baseline-24 h urine samples for UAG (IQR:  $15.9 \mu\text{mol L}^{-1}$  [ $10\text{--}117 \mu\text{mol L}^{-1}$ ]) and DMEAG (IQR:  $0.24 \mu\text{mol L}^{-1}$  [ $0.03\text{--}0.71 \mu\text{mol L}^{-1}$ ]). Urine samples from two mothers contained higher concentration of UBG.

At baseline breast milk from one mother contained a small amount of UAG 6 hours after PomJ consumption (Fig. 3) and breast milk from 6 mothers contained UAG 24 hours after PomJ consumption. At the end of the study, all breast milk samples contained UAG (IQR:  $10.7 \text{nmol L}^{-1}$  [ $3.2\text{--}28.7 \text{nmol L}^{-1}$ ]). The same two participants with UBG in urine, also showed UBG ( $0.5$  and  $11.9 \text{nmol L}^{-1}$ ) in breast milk (Fig. 3). Infant urine samples at baseline did not contain any EA metabolites (Fig. 3), while at end of the study all infant urine samples contained UAG (IQR:  $18.1 \text{nmol L}^{-1}$  [ $8.9\text{--}40.3 \text{nmol L}^{-1}$ ]). Infant urine from the same two mothers whose urine and breast milk samples contained UBG also contained UBG

( $1.8 \pm 5.5 \text{nmol L}^{-1}$  [ $0\text{--}17.5 \text{nmol L}^{-1}$ ]). Urine from five other infants contained DMEAG ( $0.8$  and  $17.5 \text{nmol L}^{-1}$ ) (Fig. 3).

### 3.3. Breast milk and stool microbiota

At the phylum level, breast milk microbiota contained Firmicutes > Proteobacteria > Verrucomicrobia > Deinococcus-Thermus > Actinobacteria (Fig. 4). Infant stool samples mainly contained Actinobacteria  $\geq$  Bacteroidetes  $\geq$  Firmicutes  $\geq$  Proteobacteria with large interindividual differences (Fig. 4). The microbiota in the mothers' stool samples contained mostly Firmicutes, Bacteroidetes and some Verrucomicrobia and Actinobacteria and minor amounts of Proteobacteria and Euryarchaeota (Fig. 4).

Alpha diversity was higher in mother stool samples compared to breast milk and infant stool. No change in alpha diversity was observed between baseline and 14 days of PomJ consumption (Fig. 5A and Fig. S2†).

Using Bray Curtis beta diversity analysis a distinct separation between breast milk, infant stool and mother stool clusters was observed (Fig. 5B). But no change in beta diversity was observed between samples collected at baseline and after 14 days of PomJ consumption (Fig. 5, ESI Table S1†).

In breast milk, changes in abundance of bacteria comparing before and after the PomJ intake showed a significant decrease in three bacteria from the phylum Firmicutes, genera *Lactococcus* ( $\log_2 -6.8$ ), *Subdoligranulum* ( $\log_2: -6.9$ ), unknown ( $\log_2 -24$ ) and one phylum Proteobacteria genus *Acinetobacter* ( $\log_2: -3.9$ ) (Fig. 6A), while the relative abundance of Firmicutes/*Faecalibacterium* increased significantly ( $\log_2: 6.1$ ). A difference in response to PomJ consumption was determined when analyzing the changes in abundance in breast milk and infant stool during the PomJ intake separately

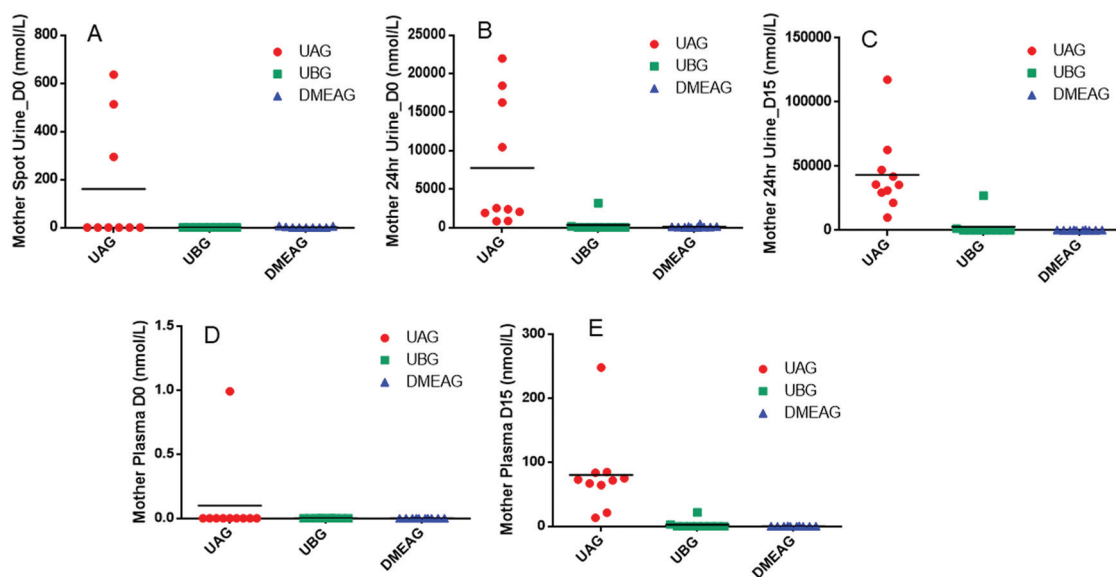
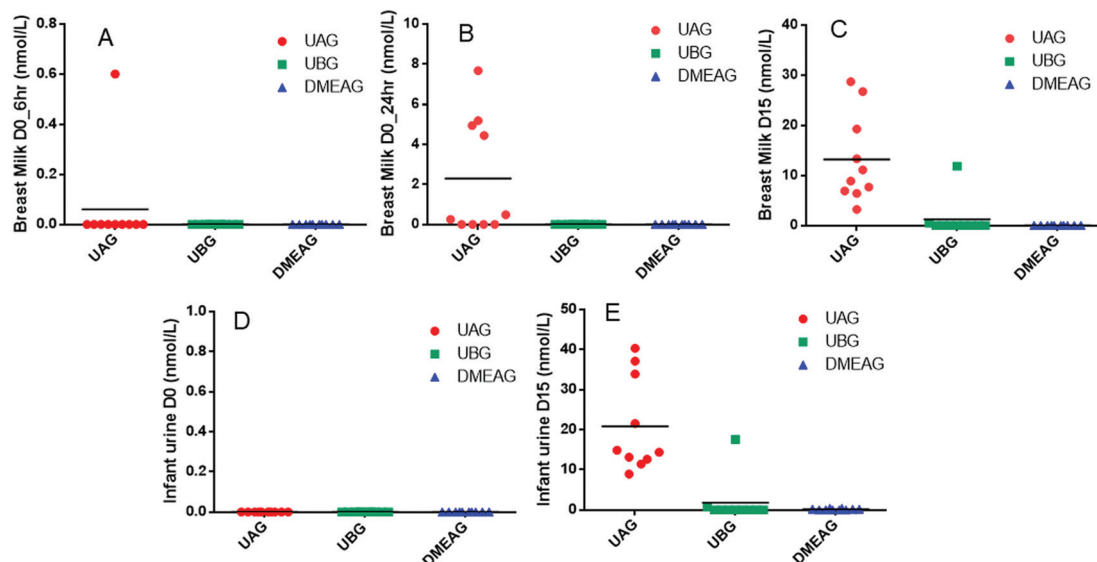
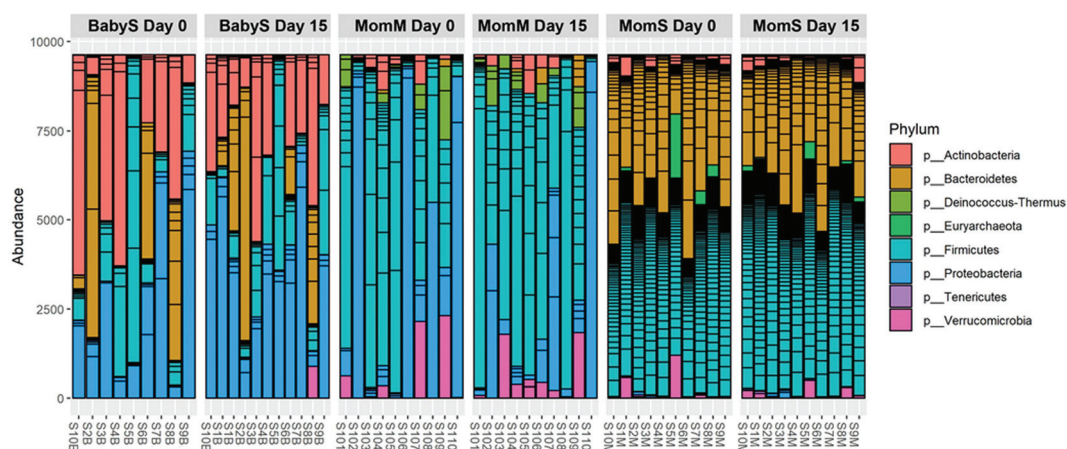


Fig. 2 Pomegranate metabolites in mother urine and plasma samples collected on day 0 and 15. (A) Spot urine collected on day 0 prior to PomJ consumption, (B) 24 h urine from day 0, (C) urine from day 15, (D) plasma from day 0 prior to PomJ consumption and (E) plasma from day 15.





**Fig. 3** Pom metabolite concentration in breast milk and infant urine samples collected on day 0 and 15. (A) Breast milk collected at 6 hours after PomJ consumption on day 0, (B) breast milk collected at 24 h on day 0, (C) breast milk collected at day 15, (D) infant urine collected on day 0, (E) infant urine collected on day 15. Breast milk  $N = 10$ . Infant urine day 15:  $N = 9$ , day 29:  $N = 10$ .



**Fig. 4** Relative abundance of microbiota on phylum level in breast milk, mother stool and infant stool samples collected on day 0 and 15. BabyS: baby stool, MomS: mother stool, MomM: mother breast milk.  $N = 10$ .

for the two metabolotypes (Fig. S3 and S4<sup>†</sup>). We performed a Spearman correlation analysis testing the correlation between the microbiota in breast milk (MomM) and infant stool (BabyS) and Pom metabolites in breast milk and mother blood (MomB) after 14 days of PomJ consumption (Fig. 7). Breast milk genus *Escherichia/Shigella* was negatively correlated with breast milk UAG (Fig. 7).

In the infant stool samples, changes in abundance of bacteria occurring during the mothers' PomJ consumption increased significantly for two bacteria from the phylum Firmicutes, genus *Lachnospirillum* ( $\log_2$  23.1) and *Staphylococcus* ( $\log_2$  4.8) (Fig. 6B). Infants with mothers from the metabolotype B showed significant differences after PomJ consumption for in 4 bacteria, with 2 bacteria from the

phylum Firmicutes (*Veillonella*  $\log_2$  23.9), 2 Bacteroidetes (*Bacteroides*  $\log_2$  25.4, *Parabacteroides*  $\log_2$  23.4) and 1 Actinobacter (*Bifidobacterium*  $\log_2$  25.0) (Fig. S4<sup>†</sup>). Spearman correlation analysis demonstrated that infant stool *Blautia* was positively correlated to breast milk and plasma UBG (Fig. 7). In addition infant stool *Enterococcus* was inversely related to breast milk UBG, plasma UBG and DMEAG.

In the maternal stool samples, changes were observed in the bacteria abundance occurring during the PomJ consumption of one bacterium from the phylum Firmicutes (*Anaerostipes*  $\log_2$  1.2) that was slightly increased. In the metabolotype B group one bacterium from the phylum Firmicutes (*Sutterella*  $\log_2$  8.9) was increased significantly (data not presented).



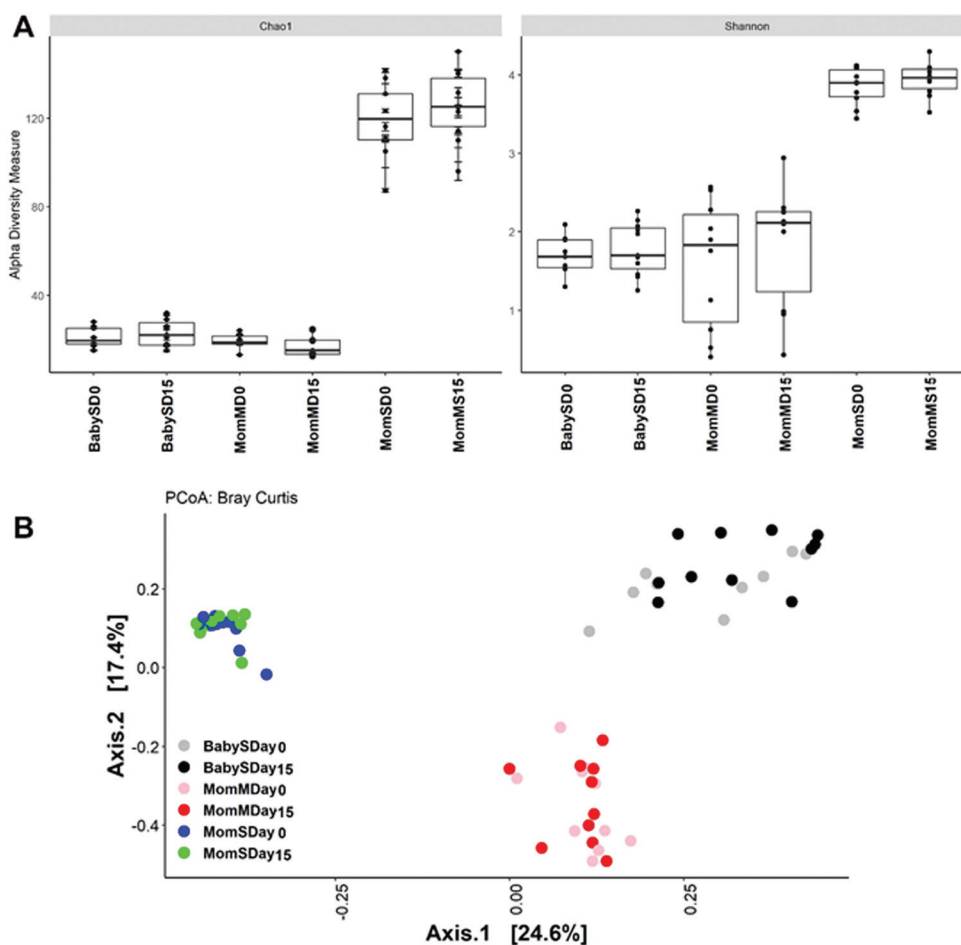


Fig. 5 (A) Alpha diversity and (B) beta diversity of breast milk and mother and infant stool samples collected at days 0 and 15. BabyS: baby stool, MomS: mother stool, MomM: mother breast milk.  $N = 10$ .

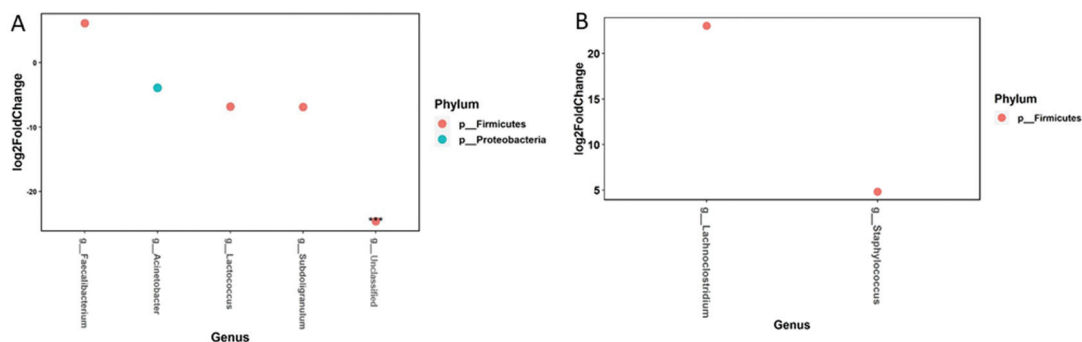


Fig. 6 Microbial genera identified to be significantly different in abundance between day 0 and day 15 after PomJ consumption in (A) breast milk and (B) infant stool.  $N = 10$ .

## 4. Discussion

The present study confirms findings from our previous pilot study demonstrating the presence of pomegranate metabolites in breast milk of mothers consuming PomJ. The novel finding in the present study is that Pom metabolites such

as UAG, UBG and DMEAG were absorbed by the infants from breast milk and excreted in the infants' urine. All mothers were UA producers, and two mothers had the intestinal microbiota to form both UA and UB with UBG present in their plasma, urine, breast milk and their children's urine.



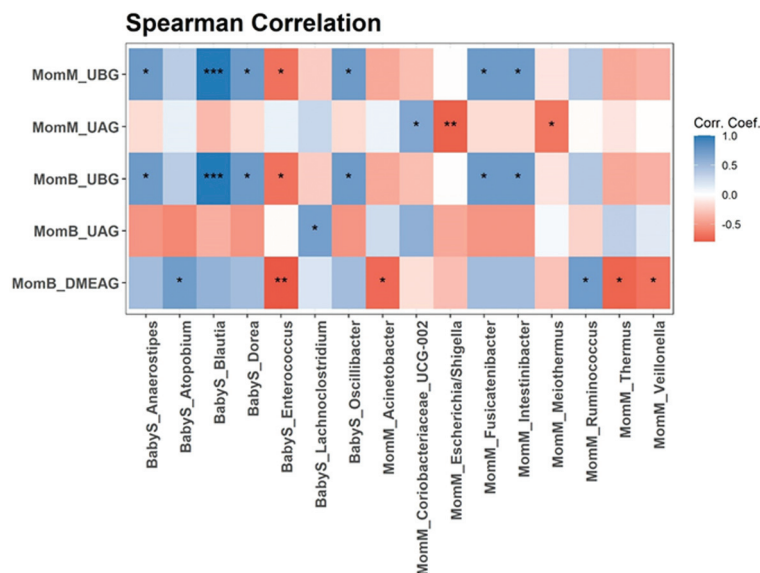


Fig. 7 Spearman rank correlations between Pom metabolites in breast milk and mothers' plasma and breast milk and infant stool microbiota at the genus level at day 15. \*\*\*  $p < 0.001$ , \*\*  $p < 0.01$ , \*  $p < 0.05$ . BabyS: baby stool, MomM: mother breast milk and MomB: mother blood.  $N = 10$ .

Pomegranate ellagitannins affect bacteria in the adult gut with acute and chronic exposure<sup>11</sup> and could also indirectly affect the microbiome of newborn infants. We propose that EA and urolithins ingested from breast milk are further metabolized by the infant's intestinal bacteria, leading to alteration in the infant's gut microbiota. In addition, there is evidence that gut associated anaerobes are transferred from the mother's intestine to neonates' intestines *via* breastfeeding.<sup>29–31</sup> In the present study, at the end of the washout phase, the relative abundance of bacteria were similar in breast milk and infant stool samples with abundance of Actinobacteria, Proteobacteria, and Firmicutes, while maternal stool samples were populated mainly with Firmicutes and Bacteroidetes. A published study by Pannaraj *et al.* showed similar results with main phyla in breast milk being Proteobacteria and Firmicutes and in infant stool being Proteobacteria and Actinobacteria.<sup>32</sup>

In the present study, changes in microbiota during the PomJ consumption in the breast milk including all participants demonstrated an increase in *Faecalibacterium* and a decrease in *Subdoligranulum* and *Acinetobacter*. In infant stool samples the relative abundance of *Lachnospirillum* ( $\log_2$  23.1) and *Staphylococcus* ( $\log_2$  4.8) were increased after 14 days of PomJ consumption (Fig. 6B). In addition, *Blautia* in infant stool, was positively correlated to breast milk and plasma UBG. *Lachnospirillum* and *Faecalibacterium* are common members of the mammary gland specific microbiome, which may colonize the mammary gland *via* physiological translocation before lactation.<sup>33</sup> They are able to colonize the gut of breastfeeding infants due to growth promotion by milk oligosaccharides.<sup>33</sup>

The anaerobic *Faecalibacterium*, *Lachnospirillum*, *Blautia* and *Acinetobacter* have been shown to transfer from the

maternal to infant gut during lactation.<sup>33</sup> *Faecalibacterium*, *Subdoligranulum* and *Lachnospirillum* belong to the class Clostridia and are anaerobic butyrate-producers.<sup>29</sup> These bacteria have been found in maternal fecal and breast milk samples.<sup>29</sup> Butyrate is a short-chain fatty acid (SCFA) which has beneficial effects on host physiology, including influencing the stability and physiology of the intestinal environment and serving as an energy source for colonocytes. Among several other roles, butyrate helps to maintain the integrity of the mucosa, protecting against cellular inflammation and promoting the removal of dysfunctional cells.<sup>34</sup> Formation of butyrate in breast milk might enhance butyrate in the infants' intestine and provide benefits to the infants as well.

A publication by Jost *et al.* showed that in breast milk and infant stool samples, the facultative anaerobic genus *Staphylococcus*, that increased in breast milk, was among the predominant members of the Firmicutes phylum.<sup>29,35</sup> *Lactococcus*, which decreased in breast milk belongs to the genus (<https://en.wikipedia.org/wiki/Genus>) of lactic acid bacteria ([https://en.wikipedia.org/wiki/Lactic\\_acid\\_bacteria](https://en.wikipedia.org/wiki/Lactic_acid_bacteria)) and is commonly found in breast milk.<sup>35</sup> The Spearman analysis showed an inverse correlation between breast milk UAG and breast milk *Escherichia/Shigella*. The abundance of *Escherichia/Shigella* in infants stool, has been associated with diarrhea in infants.<sup>36</sup> Our findings might demonstrate another benefit of PomJ consumption by nursing mothers.

Interindividual variation in urolithin formation has been observed in association with differences in microbiota populations. In a previous study from our laboratory, based on urinary and fecal content of the POM metabolite urolithin A (UA), we observed that 30% of individuals consuming Pom extract for 3 weeks did not form UA, while 40% formed UA and 30% experienced induction of UA formation during the 3



weeks of Pom extract consumption.<sup>11</sup> Selma *et al.* demonstrated that among healthy individuals 12–30% did not form urolithins and among the urolithin producers 57% formed urolithin A (UA), 31% urolithin B (UB).<sup>9,11</sup> In the present study, all participants formed urolithins at 24 h after PomJ consumption. This observation might be unique to breast feeding mothers, who also eat a healthy diet during pregnancy and breast feeding enabling their intestinal microflora to form urolithins. Only small changes were observed in relative bacterial abundance in the mother stool samples from day 0 to 15 demonstrating the stability of the microbiota in healthy individuals.

Two mothers were metatype B (UB producers), which was associated with a difference in the effects of PomJ consumption on microbiota composition in breast milk and infant stool samples (ESI Fig. S3 and S4†). Future studies with larger numbers of participants are necessary to analyze the difference in microbiota changes between metatype A and B.

Limitations of this intervention study include the lack of dietary assessment of food intake of the mothers. Knowing the dietary pattern of the participants would assist in interpreting the microbiota data. In addition, the small number of participants limited the number of participants with metatype B, which reduced the chance of finding differences in the microbiota between the metatypes.

In the present study the concentrations of urolithins in breast milk and mothers' urine samples match results from our previous pilot study.<sup>23</sup> EA and urolithins are bioavailable, which allows the absorption from breast milk by the infants. However, we found that infant urinary concentration of UAG was ~2000-fold lower compared to the mother urine.

More and more evidence has indicated that EA as well as its microbial metabolites urolithins are involved in many biological activities.<sup>10</sup> The antioxidant and anti-inflammation effects of EA have been studied.<sup>10,37</sup> The developing fetal brain is particularly vulnerable to the harmful effects of oxidative stress, with the result often similar to that of neurological injury caused by a hypoxic-ischemic event around the time of birth.<sup>38</sup> Neuroprotective potential in infants whose mothers consumed pomegranate juice demonstrated lower risk for brain injury, including any white or cortical grey matter injury compared to placebo.<sup>38,39</sup> Demonstrating health benefits of pomegranate consumption on infant health could provide a greater incentive for women to breast feed.

## 5. Conclusions

In this study, we showed for the first time that ellagic acid metabolites (DMEAG and UAG) can be absorbed by the infant from breast milk and are excreted in the infant urine. However, the Pom metabolite concentration in the infants' urine is much lower compared to the mothers' urine. The presence of PomJ metabolites in the infant's intestine was associated with an increase in the relative proportion of *Lachnoclostridium* (log<sub>2</sub> 23.1) and *Staphylococcus* (log<sub>2</sub> 4.8),

that could contribute to the health benefits of maternal PomJ consumption. This study provides important information demonstrating the need for future studies to investigate the impact of phenolic compounds from PomJ on neonatal health and development.

## Author contributions

SMH: Investigation, original draft; DH, ZL and MK: conceptualization, review & editing; JY formal analysis, review & editing; RPL, JH and DBN: methodology; GT: project administration.

## Conflicts of interest

There are no conflicts of interest to declare.

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