

1 **The Pregnancy and EARly Life study (PEARL) - A longitudinal study to understand**  
2 **how gut microbes contribute to maintaining health during pregnancy and early life.**

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19

20 **1 Abstract**

21 **1.1 BACKGROUND:**

22 The early life period represents the first step in establishing a beneficial microbial  
23 ecosystem, which in turn affects both short and longer-term health. Changes during  
24 pregnancy influence the neonatal microbiome; through transmission of maternal  
25 microbes during childbirth, and beyond, through nutritional programming. However, in-  
26 depth exploration of longitudinal maternal-infant cohorts, with sampling of multiple body  
27 sites, complemented by clinical and nutritional metadata, and use of cutting-edge  
28 experimental systems are limited.

29 The PEARL study will increase our knowledge of; how microbes (including  
30 viruses/phages, bacteria, fungi and archaea) change in composition and functional  
31 capacity during pregnancy; transmission pathways from mother to infant; the impact of  
32 various factors on microbial communities across pregnancy and early life (e.g. diet), and  
33 how these microbes interact with other microbes and modulate host processes, including  
34 links to disease onset.

## 35 **1.2 METHODS:**

36 PEARL is a longitudinal observational prospective study of 250 pregnant women and  
37 their newborns, with stool and blood samples, questionnaires and routine clinical data  
38 collected during pregnancy, labour, birth and up to 24 months post birth.

39 Metagenomic sequencing of samples will be used to define microbiome profiles, and  
40 allow for genus, species and strain-level taxonomic identification and corresponding  
41 functional analysis. A subset of samples will be analysed for host (immune/metabolite)  
42 molecules to identify factors that alter the host gut environment. Culturing will be used to  
43 identify new strains of health-promoting bacteria, and potential pathogens. Various *in*  
44 *vitro* and *in vivo* experiments will probe underlying mechanisms governing microbe-  
45 microbe and microbe-host interactions.

## 46 **1.3 DISCUSSION:**

47 Longitudinal studies, like PEARL, are critical if we are to define biomarkers, determine  
48 mechanisms underlying microbiome profiles in health and disease, and develop new  
49 diet- and microbe-based therapies to be tested in future studies and clinical trials.

#### 50 **1.4 TRIAL REGISTRATION:**

51 This study is registered in the ClinicalTrials.gov Database with ID: NCT03916874

#### 52 **1.5 KEYWORDS:**

53 Microbiome, Pregnancy, Early-life, Gut health, Health

## 54 **2 Background**

55 The human gut microbiota comprises a complex microbial ecosystem (including  
56 viruses/phages, bacteria, fungi and archaea) which benefit their host through acquisition of  
57 additional nutrients and energy from dietary components, optimised development of the  
58 immune system, and resistance against pathogens [1,2]. As the neonatal gut is essentially  
59 sterile, these beneficial microbes and their associated functions, must be acquired during  
60 and after birth; this is achieved through initial colonisation by pioneer bacteria, successive  
61 diversification, and changes in microbial population densities over time, until a climax or  
62 'stable' microbiome is established during infancy and early childhood [3,4].

63 The initial colonisation and establishment of the gut microbiome during early life influences  
64 physiological and immune development [4]. Critically, disturbances within this pioneering  
65 microbial community (both in mother and infant) have the potential to increase the risk of  
66 developing diseases such as autoimmune conditions, allergic-type disorders, infections and  
67 chronic intestinal diseases [4,5,6]. Factors such as antibiotic usage, diet (i.e. breast versus  
68 formula milk) and birth mode (i.e. Caesarean section versus vaginal) can affect the gut  
69 microbiota during this time [7,8,9].

70 Transmission of microbial species from the maternal microbiome to the infant occurs at birth,  
71 with subsequent waves of colonisation occurring as the infant ages [10,11,12]. There is also  
72 an increasing awareness of the importance of maternal health during pregnancy for infant  
73 development and health, both before and after birth [13]. Importantly, these life stage events  
74 appear to be governed by particular interactions (immune and dietary) that 'select' beneficial  
75 microbes such as *Bifidobacterium* [4,14,15]. Species in this genus are prevalent in the  
76 gastrointestinal tract of mothers in the later stages of pregnancy and can represent up to  
77 80% of the total microbiome in healthy infants [14,16,17]. These pioneering microbial  
78 species and strains contribute to ecosystem structuring, which is heavily influenced by their  
79 ability to metabolise complex sugars presents in breast milk – human milk oligosaccharides  
80 [17].

81 Although several studies have looked at these key factors at these critical timepoints, there  
82 are still many unknowns with respect to this key developmental window including; how the  
83 microbiome changes in response to different phases of pregnancy across different body  
84 sites; whether microbes from mothers are directly passed to infants during birth, and how  
85 birth mode affects this (i.e. vaginal birth vs. C-section); how factors like diet and antibiotics  
86 influence the maternal microbiome, and what impact this has on developing infant microbial  
87 communities; how different feeding regiments (e.g. breast vs. formula milk) influence specific  
88 microbial populations in the infant (e.g. *Bifidobacterium*); how these microbes influence  
89 immune and metabolic health.; and if in-depth mechanistic studies using *in vitro* and *in vivo*  
90 models can determine how specific microbes and communities contribute to healthy  
91 development and prevent disease incidence.

92 Longitudinal pregnancy and infant studies like PEARL, will provide an invaluable resource to  
93 study the importance of the early life microbiome. Gathering these data is critical for  
94 identification and development of new therapies and health- associated practices to improve  
95 health, both in the short- term and across the life course.

96        **2.1 OBJECTIVES:**

97        The primary objective of the PEARL study is to describe the determinants, function and  
98        composition of maternal and neonatal microbiomes throughout pregnancy and early  
99        childhood in a cohort of mothers and babies without serious health conditions.

100       Other objectives of this study are;

101            1) to link or correlate microbiome profiles to routinely collected clinical information in  
102            mothers and infants (e.g. antibiotic usage, birth mode) and dietary information via  
103            questionnaires (e.g. breast vs. formula milk), and to host metabolites and immune  
104            markers (e.g. cytokines) measured in samples collected;

105            2) to use microbiome samples in preclinical models to describe the role that the  
106            early- life microbiome plays in immune and metabolic development and resistance to  
107            diseases;

108            3) to use analysis and microbiome samples (using *in vitro* and preclinical models) to  
109            characterise the cause, effect and consequences of disturbances in the early-life  
110            microbiome due to external factors (e.g. antibiotics, birthing method, diet) on  
111            subsequent health outcomes;

112            4) to define and characterise early-life associated microbiome species (e.g.  
113            *Bifidobacterium* species and *Enterococcus* species) from samples for their  
114            probiotic/pathogenic traits (using *in vitro* and preclinical models).

115        An additional key output, which will be available to the wider research community, is a  
116        comprehensive collection of microbiome samples and associated clinical and diet/lifestyle  
117        information that will be accessible via requests to Professor Lindsay Hall and the Norwich  
118        Research Park (NRP) Biorepository, with all microbiome sequence data publicly available  
119        after deposition in public repositories.

120 **3 METHODS:**

121 PEARL is a longitudinal study of 250 pregnant women and their new born(s). Biological  
122 samples and data will be collected in pregnancy, labour, birth and up to 24 months post  
123 birth. In the event of multiple live births, each infant will be included. Figure 2 provides an  
124 overview of the overall design of the study and shows which samples and data are collected  
125 at each timepoint. There are up to 12 collection timepoints, as well as 4 optional blood  
126 sample donations which will be collected at routine maternity care appointments. The study  
127 is divided into three collection phases: pregnancy, birth to four months, and four months to  
128 24 months post birth.

129 Pregnant women can be recruited to PEARL at any time up to 22 weeks gestation. During  
130 phase 1 participants will be asked to collect samples and complete questionnaires during  
131 each of the three trimesters of pregnancy if they are involved in the study at that point.  
132 Participants are also able to donate blood samples at weeks 12, 16, 20, 28.

133 During phase 2, participants will collect samples at three timepoints from themselves and  
134 their infant(s); birth, at 1 week post birth, and at 3 weeks post birth.

135 Phase 3 collections will be every 4 months (4, 8, 12, 16, 20 and 24 months). With the  
136 exception of month 4, further collections of samples will be from the infant only.

137 At each collection time point, there is a window of 7 days within which the participant can  
138 collect their samples. This is to provide as much flexibility for participants as possible.

139 Participants will receive a reminder message via text message or email when their samples  
140 are due to be taken, and again towards the end of the timepoint window.

141 Most samples will be collected at home using packs provided to the participants. The  
142 participants will be provided with a small freezer where they will store their samples until  
143 collected by a member of the research team. This freezer will be dedicated to holding study

144 samples to minimize any cross-contamination risks to the participant. Collections from the  
145 participants' homes will occur at 4 points throughout the study after phase 1, after phase two  
146 and at month 12 and month 24 during phase 3.

147 At birth, or any time the participant or her infant is an inpatient at the Norfolk and Norwich  
148 University Hospitals NHS Foundation Trust (NNUH), participants can collect their samples at  
149 the NNUH and give them to their midwife/obstetrician who will put these in the PEARL study  
150 freezer located at the hospital.

151 The sample collection protocol has been designed using significant optimisation information  
152 from our previous study, BAMBI [18], with additional information gathered from other  
153 longitudinal cohort studies including the Flemish Gut Project [19], and we are confident that  
154 our current protocol will allow participants to easily collect and store samples at -20°C until  
155 collection.

156 After collection from participants' homes, all samples will be appropriately aliquoted and will  
157 be stored in QIB at -80°C prior to analysis. Additional aliquots may also be stored at the  
158 NRP Biorepository at -80°C for future research if the participant has provided this additional  
159 consent.

160 Participants will complete 3 questionnaires during the study (timepoints detailed in figure 2)  
161 including questions on health, lifestyle, and diet. The questionnaires include:

- 162 • Centre for Disease Control and Prevention (CDCP) questionnaires for pregnancy,  
163 breastfeeding and infant feeding practices [20]. Modified, with permission, for the  
164 purposes of this study (supplementary files 1 - 7).
- 165 • A health questionnaire to provide information on lifestyle, diet and  
166 medication/medical history, developed for the study (supplementary files 8-9).
- 167 • A dietary perceptions and preferences questionnaire, developed for the study  
168 (supplementary files 10-12).

169 The research team will also collect; clinical information from GP and hospital records that  
170 relate to any change in the participants or her infants' medical history during the course of  
171 the study; routine data collected by health visitors (developmental Ages & Stages data)  
172 which includes information on fine and gross motor skills and problem solving; and routine  
173 pregnancy scans and tests.

## 174 **COVID19 Mitigation Strategies**

175 In response to the COVID-19 pandemic, it is important to implement a risk mitigation  
176 strategy to protect the health and safety of both participants and researchers. Participants  
177 will be advised that they should not collect samples if they or anyone in their household has  
178 COVID19 symptoms, are self-isolating, are awaiting test results, or have in the last 70 days  
179 tested positive for COVID19. The research team will send collection packs and arrange the  
180 collection of participants' samples following Standard Operating Procedures (SOPs)/  
181 COVID-19 risk-assessments.

## 182 **3.1 PARTICIPANT SELECTION**

### 183 **3.1.1 Sample Size**

184 PEARL will recruit 250 pregnant women. Our primary aim is to detect associations between  
185 aspects of neonatal and maternal microbiomes, and their associations with external factors.  
186 As an indicative power calculation, if a potential risk factor has a prevalence of 25% (recent  
187 data suggests that 25% of mothers who deliver at our recruitment hospital [Norfolk and  
188 Norwich University Hospitals NHS Foundation Trust – NNUH] do not initiate breast feeding;  
189 the UK Caesarean section rate is also currently around 25%). Then with 200 observations  
190 we would have power of 80% to detect a difference of 0.46 standard deviations in any given  
191 outcome measure between those with and without the feature at a p-value of 0.05. This is  
192 commonly understood to correspond to a 'moderate' size of effect between an exposure and  
193 an outcome.



194 For continuous outcomes or correlations between 250 mother and baby pairs, we would be  
195 able to detect correlations of  $r=0.2$  with 80% at  $p<0.05$ . This corresponds to a continuous  
196 feature accounting for  $r^2=4\%$  of the population variation in any given characteristic of the  
197 microbiome.

198 For secondary analyses, if the p-value threshold for statistical significance is corrected to  
199  $p<0.005$  to account for multiplicity then the corresponding smallest detectable differences  
200 are 0.6 standard deviations between groups defined by a binary factor (e.g. mode of delivery  
201 or breastfeeding) or correlation of 0.25 between continuous variables.

202 Previously, Azad et al (2015) [21] explore effects of maternal antibiotic use, mode of delivery  
203 and breastfeeding on diversity and major taxon abundance among a cohort of 198 Canadian  
204 neonates at 3 and 12 months of age. Differences in the three most abundant taxa between  
205 elective Caesarean section ( $n=19$ ) and vaginal births ( $n=96$ ) were detected at three months  
206 with p-values of  $<0.001$  and  $<0.01$ , corresponding to effect sizes of  $>0.63$  and  $>0.49$   
207 standard deviations.

### 208 **3.1.2 Cohort**

209 250 healthy pregnant women will be recruited from across Norfolk. We will be recruiting  
210 participants from the maternity unit at a single major regional hospital, the Norwich and  
211 Norfolk University Hospital Foundation Trust (NNUH) which provides care for a population of  
212 approximately 1,000,000. The vast majority of women in the region will use the maternity  
213 services of this hospital. Participants will all be planning to give birth at NNUH or have a  
214 home birth with their maternity care provided by the NNUH. The total number of live births  
215 per year at NNUH is  $\sim 6000$ . Our primary objective is to describe the determinants, function  
216 and composition of maternal and neonatal microbiomes throughout pregnancy and early  
217 childhood in a cohort of mothers and their babies without serious health conditions.

218 Therefore, we will be seeking a 'healthy' cohort of mothers, with the following inclusion and  
219 exclusion criteria applied.

220 ***Inclusion criteria***

- 221 • Confirmed pregnancy and aged at least 18 years.
- 222 • Must be able to understand the requirements of the study and provide signed and  
223 dated informed consent for herself and her unborn child.
- 224 • Planning to give birth at NNUH or at home under the care of NNUH.
- 225 • At the point of study consent, be  $\leq 22$  weeks pregnant.
- 226 • Body Mass Index (BMI) between 18 and 35kg/m<sup>2</sup>
- 227 • Must be willing to provide biological samples over a period of 31 months (urine  
228 samples, stool samples, low vaginal swabs, skin swabs and breast milk (if  
229 breastfeeding this is optional). Blood samples are optional.
- 230 • Must be willing to accommodate a small freezer to store frozen samples for the  
231 duration of the study.
- 232 • Must be willing to complete study questionnaires over a period of 31 months.

233 ***Exclusion criteria***

234 The participant may not enter the study if ANY of the following apply:

- 235 • Pregnancy is for surrogacy purposes.
- 236 • Planned adoption, fostering of baby or baby not planned to be living with  
237 biological mother.
- 238 • Currently taking part in an interventional study.
- 239 • Living with or related to a member of the research team.
- 240 • Current smoker.
- 241 • Taken antibiotics or antifungals or antivirals within the last 3 months.

- 242 • Taken steroids within the last 6 months.
- 243 • Currently taking more than a daily dose of probiotics.
- 244 • History of polyps within the gut.
- 245 • Long-standing gastrointestinal or liver function abnormality requiring on-going  
246 medical management or medication.
- 247 • Current or history of cancer except for squamous or basal cell carcinomas of the  
248 skin that have been medically managed by local excision.
- 249 • Unstable dietary history as defined by major changes in diet during the previous  
250 month, where the participant has stopped or significantly increased a major food  
251 group in the diet, for example changed to vegan, vegetarian or stopped eating  
252 red meat.
- 253 • History of alcohol, drug or substance abuse.
- 254 • History of Hepatitis B or Hepatitis C.
- 255 • Any confirmed or suspected pre-existing condition/state of immunosuppression or  
256 immunodeficiency (primary or acquired), for example Rheumatoid Arthritis, Type  
257 1 Diabetes, Multiple Sclerosis, Asthma, Eczema and Psoriasis. (Participants who  
258 are asymptomatic of Asthma, Eczema and Psoriasis in the last 5 years can be  
259 included in the study).
- 260 • Major surgery of the gastrointestinal tract, apart from gall bladder or appendix  
261 removal, in the past five years.
- 262 • Any major bowel resection at any time.
- 263 • History of Ulcerative Colitis or Crohn's Disease or Diverticulitis.
- 264 • Persistent, infectious gastroenteritis, gastritis, persistent or chronic diarrhoea of  
265 unknown cause, Clostridium difficile infection (recurrent) or Helicobacter pylori  
266 infection (untreated).
- 267 • Chronic constipation.

268        **3.2 RECRUITMENT**

269        Recruitment began in May 2019 and will continue until May 2022. Recruitment will be mainly  
270        opportunistic at the NNUH NHS Foundation Trust antenatal clinics and midwife led  
271        community clinics in GP surgeries and health clinics.

272        In response to the COVID19 pandemic, digital appointments have been put in place for  
273        pregnant women that replace the community clinics in GP surgeries and health clinics. Due  
274        to the efficiencies that this brings, it is anticipated that these appointments will remain  
275        common practise even after measures put in place to control the pandemic have been lifted.  
276        We, therefore, also intend to include virtual recruitment as a routine recruitment method for  
277        this study.

278        Following the first appointment a pregnant woman has as part of her routine maternity care  
279        at NNUH, eligible, patients considered to have a low risk pregnancy will be contacted by a  
280        research midwife by telephone to introduce the study and ask if the participant is interested  
281        in taking part. If the participant is interested, they will be provided with the participant  
282        information sheet (PIS) via email and a video/telephone call will be arranged to continue the  
283        informed consent process if they so wish. Participants may also contact the study researcher  
284        directly by entering their interest into the expression of interest form on the PEARL study  
285        web page, by telephone, or via email. Participants may request a face to face consent  
286        appointment which will be held at the Quadram Institute Clinical Research Facility, however,  
287        this may not be possible if there are local restrictions in place to manage the COVID19  
288        pandemic. Written informed consent will be obtained following Good Clinical Practice (GCP)  
289        guidelines by the research midwife or a member of the research study team.

290        The participant will then be assigned their unique PEARL study participant number which will  
291        provide pseudonymisation for the participant throughout the study.

292 It will be clearly stated that the participant is free to withdraw from the study at any time for  
293 any reason without prejudice to future care, without affecting their legal rights, and with no  
294 obligation to give the reason for withdrawal.

### 295 **3.3 SAMPLE BANKING AT THE NRP BIOREPOSITORY**

296 A key aspect of the PEARL study involves creating biobank of samples and data which  
297 can be used in future ethically approved research. To enable this, and following  
298 Informed Consent for study participation, all participants will be asked if they would be  
299 willing for their samples to be stored anonymously at the Norwich Research Park (NRP)  
300 Biorepository during and after the end of the study for long term storage. The NRP  
301 Biorepository is a tissue bank licensed by the UK's Human Tissue Authority (HTA) with  
302 appropriate ethics approval which was granted by the NHS Health Research Authority  
303 (HRA), East of England-Cambridge East Research Ethics Committee (REC). Samples  
304 stored at NRP Biorepository will be used in future ethically approved research.

305 Participants consenting to the long-term storage of their samples at the NRP  
306 Biorepository are given the option to provide explicit consent for their samples to  
307 potentially be used in future ethically approved animal, cloning, or commercial studies,  
308 however, these aspects are optional and not required for samples to be stored long term  
309 in the NRP Biorepository. Data will be recorded by returning a copy of the consent form  
310 to the Biorepository. The participant will be asked to read the current version of the NRP  
311 Biorepository Information Sheet and to sign this consent if they wish to participate in this  
312 aspect of the study. If the participant declines long term storage at the NRP  
313 Biorepository, all samples at the end of the study will be destroyed appropriately in  
314 accordance with the Human Tissue Act requirements.

### 315 **3.4 STATISTICS AND ANALYSIS**

#### 316 **3.4.1 Interim Analyses**

317 Interim reports will include the number of accruals to the study, the quality of the samples  
318 returned, the completeness and validity of the questionnaire data returned. The report will  
319 also detail any missing data and any participants who are lost to follow-up.

### 320 **3.4.2 Analysis of Samples**

321 Samples will be subjected to microbiome analysis using high-throughput sequencing  
322 following standardised protocols validated and quality controlled from BAMBI SOPs,  
323 including: shotgun metagenomics for species-level taxonomic identification; functional  
324 profiling; and analysis of the gut-associated antimicrobial resistance 'resistome' using open  
325 access bioinformatics pipelines and new pipelines.

326 Bioinformatics pipelines are constantly being improved/updated, thus we will use the most  
327 appropriate ones available at time of analysis which may include e.g. MetaPhlAn 3 [22].

328 Downstream statistical analyses will also be performed by using various packages in R.

329 Stool samples will also be analysed for host (immune/metabolite) molecules to determine  
330 which factors (e.g. antibiotics, diet) alter the host gut environment, which will be carried out  
331 using standard lab protocols (e.g. multi-plex cytokine assays, and NMR metabolomics).

332 Samples will also be cultured (single and complex [i.e. model colon systems]) to identify new  
333 strains of beneficial or 'probiotic' bacteria such as *Bifidobacterium* species, and also any  
334 potential pathogens, for use in additional *in vitro* and *in vivo* studies to answer specific  
335 questions about how early life microbes modulate host functions.

### 336 **3.5 DESCRIPTION OF STATISTICAL METHODS**

337 A main aim of this study is to establish a cohort and repository of samples for future  
338 research, hence there are many possible future analyses that may be done using the  
339 samples collected as a part of this study.

340 For primary analyses, the composition of the microbiome in each sample (as defined by  
341 participant, site and time point) will be described in terms of individual taxon abundance and  
342 overall measures of diversity. Several key aspects of composition of the maternal and  
343 neonatal microbiome will be identified as candidates for association with measures of health  
344 status based on review of previous studies and theoretical considerations.

345 The dynamics of microbiome development and transmission from mother to baby will be  
346 determined by describing and correlating the distribution of key measures of structure and  
347 composition between samples.

348 The effect of factors that might affect microbiome composition, such as breastfeeding  
349 initiation and mode of delivery, will be assessed using regression models.

350 For exploratory (hypothesis generating) analyses, multivariate methods such as partial least  
351 squares regression modelling will be used to explore internal correlation structure within  
352 each sample and test whether any set of microbiome characteristics identified correlates  
353 with a hypothesised clinical factor.

354 Throughout, careful attention will be paid to the possibility of false positive results occurring  
355 through multiplicity given the large number of hypotheses being tested and the large number  
356 of parameters to be set in bioinformatics and statistical techniques. This will be mitigated as  
357 far as possible for each hypothesis by clearly pre-specifying each individual analysis in a  
358 statistical analysis plan detailing the coding of exposures, outcomes and covariates and  
359 primary statistical methods to be employed including any subgroup analyses. These  
360 statistical analysis plans will be developed and will be pre-registered following data collection  
361 so that the distributions of key variables are known but before relevant analysis of  
362 microbiome and health outcome information is undertaken. Exploratory and secondary  
363 analyses will be clearly reported as such in all outputs, and results from all analyses will be  
364 published irrespective of whether findings are predominantly positive or negative.

365 **3.6 STUDY MANAGEMENT PLAN**

366 **3.6.1 Trial Management Group**

367 The Trial Management Group including the Chief Investigator (Prof Lindsay Hall), the study  
368 research team, NHS Principal Investigators and Clinical Research Network's (CRN)  
369 Research Nurses will be responsible for the day-to-day management of the trial. They will  
370 monitor all aspects of the conduct and progress of the trial, ensure that the protocol is  
371 adhered to and take appropriate action to safeguard participants and the quality of the trial  
372 itself.

373 **3.6.2 Trial Management Oversight Group**

374 Management of the study will be overseen by a Trial Management Oversight Group whose  
375 membership is made up of representatives from;

- 376 • Health, Safety, Environment and Quality Assurance.
- 377 • QIB Sponsor Representative.
- 378 • Gut Microbes and Health QIB Programme Manager.
- 379 • QIB Statistician.
- 380 • Information Technology Security Specialist.
- 381 • NRP Biorepository.
- 382 • Patient and Public Involvement.
- 383 • QIB Bioinformatician.

384  
385 The Trial Management Oversight Group will be responsible for overseeing the running of the  
386 study. They will ensure the monitoring and facilitating the progress of the study, and ensure  
387 the study is delivered within the projected timelines. Recruitment targets, success of data  
388 collection, and any specific issues arising will be addressed.

389 **3.7 DATA MANAGEMENT PLAN**



390 **3.7.1 Description of the Data**

391 Our data collected for this study will be the following: -

392 • Clinical data and measurements taken as part of routine care (Antenatal and Postnatal  
393 Data Collection Case Report Forms).

394 • Participant-reported general health and lifestyle data (Participant Trimester 2 and  
395 Participant 24 months Post Birth Health Questionnaires – Supplementary files 8 and 9) and  
396 Participant Dietary Preferences and Perceptions Questionnaire (Participant Trimester 2, 4  
397 months post birth and 24 months post birth – supplementary files 10 - 12).

398 • CDCP questionnaires, (participant-reported data, dietary intake and allergies data –  
399 supplementary files 1 - 7).

400 • Biological Sample data (blood, stool, urine, colostrum/breast milk, cord blood, low vaginal  
401 swabs, skin swabs and meconium samples).

402 • Data collected from NNUH Integrated Clinical Environment (ICE) system (routine  
403 pregnancy results), other relevant databases within the Trust and hospital notes where it is  
404 relevant to the study.

405 All of the above collected data will be anonymised and stored on a secure IT (Information  
406 Technology) system which only the study research team will have access to.

407 A data monitoring committee is not needed because the trial is not testing a drug or device  
408 and any safety concerns associated with the trial have been reviewed by the ethics  
409 committee.

410 **3.7.2 Specific Management of Samples**

411 All biological samples and data collected as part of the study will be pseudonymised.  
412 Laboratory results will be maintained in a database and will be in file formats that can be  
413 shared internally. Sample management at QIB and the NRP biorepository will be enabled by  
414 use of a locally acquired Laboratory Information Management System. Only anonymised  
415 individual-level data will be shared within study team members (QIB and NHS investigators).

### 416 **3.7.3 Data Collection / Generation**

417 Data from medical records (hospital notes and GP records) will be extracted onto study-  
418 specific, ethically approved, forms and then uploaded into secure databases which has  
419 shared access with appropriately authorised research staff working on the project in  
420 compliance with International Good Clinical Practice (GCP) standards.

### 421 **3.7.4 Data Sharing and Access**

422 The research protocol is registered at ClinicalTrials.gov (NCT03916874). The study team  
423 will ensure full compliance with the standards required for deposition of information in any  
424 relevant public databases. Only anonymised individual-level datasets will be shared outside  
425 the team. Consent forms clearly state the data sharing procedures for data generated from  
426 this study.

427 All data will be managed, protected and shared in accordance with the requirements the  
428 Biotechnology and Biological Sciences Research Council (BBSRC) Data Sharing Policy.

429 Direct access will be granted to authorised representatives from the Sponsor and host  
430 institution for monitoring and/or audit of the study to ensure compliance with regulations.

## 431 **ETHICAL AND REGULATORY CONSIDERATIONS**

### 432 **3.7.5 Declaration of Helsinki and Relevant Regulations**

433 The Investigator will ensure that this study is conducted in accordance with the principles of  
434 the Declaration of Helsinki. The proposed research will be conducted in accordance with the

435 conditions and principles of the International Conference on Harmonisation Good Clinical  
436 Practice, and in compliance with national law. The research will meet the requirements of the  
437 new EU General Data Protection Regulation (GDPR), UK Data Protection Act 2018 and  
438 relevant sponsor's policies.

### 439 **3.7.6 Expenses and Benefits**

440 There are no planned in-person study visits for this study. However, if a participant requests  
441 a face to face consent appointment at the Quadram Institute Clinical Research Facility,  
442 provisions will be made to reimburse the participant for car parking, and travel expenses at  
443 45p per mile. As a thank you, participants will be sent a £20 shopping voucher once the 24-  
444 month samples and data have been received. At the end of the study, the participant will  
445 also have the option of keeping the freezer if they wish.

446 It is not anticipated that any post-trial care will be required. If any participant is harmed  
447 whilst taking part in this clinical research study as a result of negligence on the part of a  
448 member of the study team, QIB holds liability insurance for such circumstances.

## 449 **4 Discussion**

450 Only by conducting in-depth longitudinal studies of large cohorts of hosts (i.e. mothers and  
451 their babies in the case of PEARL) can we identify the factors that are responsible for  
452 shaping and sustaining the microbiome in health, or for causing disturbances in disease.  
453 Longitudinal studies, like PEARL, are critical if we are to define biomarkers and develop new  
454 diet- and microbe-based therapies to promote health during pregnancy and early life in  
455 future studies and clinical trials [23]. Our pregnant mother and baby cohort will provide a  
456 unique collection of samples robustly linked to detailed clinical information and early-life diet  
457 information, which will be available to the wider research community, thus providing a  
458 significant UK-based resource.

459 A significantly novel aspect of PEARL, is to input clinical samples and data into experimental  
460 model systems to show how the microbiome modulates specific early life developmental

461 programming, and to 'collect' novel early life associated microbiome species (e.g.  
462 *Bifidobacterium*, *Ruminococcus*, *Enterococcus*) that can be used in these systems, and why  
463 may form the basis of for future microbiota modulation clinical trials.

464 Our protocol for PEARL from the first trimester of pregnancy and foetal development through  
465 to early childhood has been designed to complement other UK-based cohort studies looking  
466 at pregnancy and early life such as the BabyBiome Study [24] and COCO90s [25]. However,  
467 to the best of our knowledge, no study has attempted to undertake a comprehensive  
468 microbiome analysis throughout pregnancy and into infancy, and none have linked this to  
469 detailed patient information that we are gathering through routine clinical data collection,  
470 lifestyle, and dietary information.

471 The major unanticipated operational issue impacting the PEARL study during 2020-2021 is  
472 the COVID-19 pandemic. With this, came implementation of national and local restrictions to  
473 reduce transmission of the virus. This included stay at home orders, social distancing and  
474 requirements to work from home where possible. The impacts on research studies such as  
475 PEARL have been significant.

476 Early in the pandemic, the PEARL study had to temporarily pause research activities while  
477 risk assessments could be urgently revised and adaptations to processes could be established  
478 to ensure safety of both participants and staff. We developed a remote process to recruit  
479 participants and introduced a reduced collection schedule from participants' homes to  
480 reduce any face to face contact involved in taking part in the study. All sample collections  
481 adhere to local and national guidelines on COVID-19 restrictions set out by the government.

482 During a second wave of the pandemic and increased pressure on local NHS trusts, we had  
483 to temporarily pause all recruitment activity to release any NHS resource that we were using  
484 for recruitment activities.

485 We have seen an increased withdrawal rate to the study at times when local restrictions for  
486 COVID-19 have been implemented which could be attributed to the increased stress brought  
487 on by these 'locked down' environments, such as increased workload of working families to  
488 home school children. Therefore, not wishing to continue in the research environment.

489 We have made several amendments to the study to increase safety of participants and  
490 research staff during the pandemic, with the hope that these actions will mean the PEARL  
491 study will not be affected in the long term.

492 It is likely COVID-19 will have implications on future studies and risk assessments should be  
493 made prior to set up.

## 494 **5 List of abbreviations**

495 BAMBI study - Baby-Associated MicroBiota of the Intestine study

496 BBSRC - Biotechnology and Biological Sciences Research Council

497 BMI - body mass index

498 CDCP – Centre for Disease Control and Prevention

499 CI – chief investigator

500 CRN – Clinical Research Network

501 GCP - Good Clinical Practice (GCP)

502 GDPR - General Data Protection Regulation

503 GP – General Practitioner

504 HRA – Health Research Authority

- 505 HRGC – Human Research and Governance Committee
- 506 ICH GCP - International Conference on Harmonisation Good Clinical Practice
- 507 ICE – Integrated Clinical Environment
- 508 IT – Information Technology
- 509 NHS – National Health Service
- 510 NIHR – National Institute for Health Research
- 511 NNUH – Norfolk and Norwich University Hospitals NHS Foundation Trust
- 512 NRP – Norwich Research Park
- 513 PIS – Participant Information Sheet
- 514 QI CRF – Quadram Institute Clinical Research Facility
- 515 REC – Research Ethics Committee
- 516 SOPs - standard operating procedures

517 **6 Declarations**

518 **6.1 ETHICS APPROVAL AND CONSENT TO PARTICIPATE**

519 The PEARL study has been reviewed and agreed by the Human Research Governance  
520 Committee at the Quadram Institute Bioscience and the London-Dulwich Research Ethics  
521 Committee (reference 18/LO/1703) and received written ethical approval by the Human  
522 Research Authority. IRAS project ID number 241880.

523 All participants in the PEARL study must be able to understand the requirements of the  
524 study and provide signed and dated informed consent for themselves and their unborn child.

525 An annual progress report will be submitted to the REC committee, HRA (and sponsor). In  
526 addition, an end of study notification and final report will be submitted to the same parties.

527 All amendments to the study protocol will be submitted to the London-Dulwich Research  
528 Ethics Committee for their approval and are reported to the sponsor and investigators.

529 Solicited and spontaneously reported adverse events and other unintended effects of the  
530 trial interventions and trial conduct will be recorded regularly in line with ethics  
531 requirements. The sponsor may conduct random audit processes to ensure compliance to  
532 the current protocol.

## 533 **6.2 CONSENT FOR PUBLICATION**

534 Not Applicable

## 535 **6.3 AVAILABILITY OF DATA AND MATERIALS**

536 Data generated from the PEARL study will adhere to the Biotechnology and Biological  
537 Sciences Research Council (BBSRC) Data Sharing Policy. Anonymised datasets will be  
538 kept indefinitely and available to other researchers. Access to such data should be  
539 requested through the Chief Investigator, Professor Lindsay Hall.

540 There will be no limitations to the dissemination of the results. It is anticipated that the  
541 results of this research project will be published and/or presented in a variety of forums,  
542 including peer reviewed journal publications, conference presentations, communications  
543 and media releases. In any publication and/or presentation, information will be provided  
544 in such a way that participants cannot be identified.

## 545 **6.4 COMPETING INTERESTS**

546 The authors declare that they have no competing interests.

547        **6.5 FUNDING**

548        The study has been funded by BBSRC through an Institute Strategic Programme (ISP)  
549        award to the QIB Gut Health and Food Safety Programme (BB/R012490/1), and its  
550        constituent projects BBS/E/F/000PR10353 and BBS/E/F/000PR10356. The study is into the  
551        NIHR CRN Central Portfolio Management System (CPMS, add study number and speciality)  
552        portfolio which provides additional support in terms of hospital infrastructure and staff  
553        support. George Savva is funded through the BBSRC Core Capability Grant BB/CCG1860/1  
554        at the Quadram Institute Bioscience. The Achiever Medical Laboratory Information  
555        Management System was procured using the BBSRC Capital Grant Award for the  
556        enhancement of the NRP Biorepository.

557        The primary sponsor of the project is the Quadram Institute Bioscience, responsible for:  
558        study design; data collection, management, analysis and interpretation; and writing of  
559        reports for publication. The funding bodies played no role in study protocol development,  
560        ethics, or have any involvement in ongoing activities.

561        **6.6 AUTHORS' CONTRIBUTIONS**

562        LJH conceived and designed the PEARL study, provided insights from previous research  
563        studies, was a major contributor to the writing of the manuscript and is the Chief Investigator  
564        of the PEARL study. SP wrote the manuscript, provided substantial contributions to the  
565        design of this protocol and is the Principal Investigator of the PEARL study. RW was a major  
566        contributor to the writing of the manuscript, provided substantial contributions to the design  
567        of this protocol and provides oversight and technical support to the PEARL study. TA was a  
568        major contributor in writing the manuscript and provides technical support to the PEARL  
569        study. GS provided statistical expertise and related protocol oversight and contributed to  
570        final manuscript drafting. AH provided substantial contribution to protocol design and



571 contributed to final drafting of the manuscript. All authors read and approved the final  
572 manuscript.

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578 Information Management System for the PEARL study team.

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580 Jackson and Rosalinde Bailey for their involvement in the PEARL study Trial  
581 Management Oversight Group and their invaluable contributions and advice regarding  
582 participant involvement in this clinical research.

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660

661 **8 Figures, tables and additional files**

662 **8.1 FIGURES**

663 Figure 1 - Overview of development of the microbiome during early life and the factors that  
664 influence this.

665 Figure 2 - PEARL study overview

666 **8.2 Supplementary files**

667 Supplementary File 1 - CDCP Participant Pregnancy Questionnaire Trimester 2

668 Supplementary File 2 - CDCP Newborn 3 Weeks Post Birth Questionnaire

669 Supplementary File 3 - CDCP Newborn 4 Months Post Birth Questionnaire

670 Supplementary File 4 - CDCP Newborn 8 Months Post Birth Questionnaire

671 Supplementary File 5 - CDCP Newborn 16 Months Post Birth Questionnaire

672 Supplementary File 6 - CDCP Newborn 20 Months Post Birth Questionnaire

673 Supplementary File 7 - CDCP Newborn 24 Months Post Birth Questionnaire

674 Supplementary File 8 - Participant Trimester 2 Health Questionnaire

675 Supplementary File 9 - Participant 24 Months Post Birth Health Questionnaire

676 Supplementary File 10 - T2 Participant Dietary Preferences and Perceptions Questionnaire

677 Supplementary File 11 - 4 Months Post Birth Participant Dietary Preferences and  
678 Perceptions Questionnaire

679 Supplementary File 12 - 24 Months Post Birth Dietary Preferences and Perceptions  
680 Questionnaire