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Parental experiences of prenatal whole exome sequencing (WES) in cases of ultrasound diagnosed fetal structural anomaly

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Parental Experiences of Prenatal Whole Exome Sequencing (WES) in Cases of

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27 Conflict of interest

28 The authors are unaware of any potential conflict of interest.

29 What is already known about this topic?

- Prenatal WES for genetic diagnosis is possible, but little is known regarding
- 31 parental experiences of prenatal sequencing

32 What does this study add?

- Parents require specific information to help them decide whether to undergo
- 34 WES for prenatal diagnosis
- Appropriate counselling is essential for informed consent
- Parents require explanation about what WES might identify, and how and
- 37 when findings are returned

38 Abstract

39 **Objective**

- 40 To explore parental experiences of WES for prenatal diagnosis, and ascertain what
- 41 influenced their decision-making to undergo testing.
- 42 Method

43 Twelve women comprised a purposeful sample in a series of semi-structured

44 interviews. All had received a fetal anomaly diagnosis on ultrasound. A topic guide

45 was used, and transcripts were thematically analysed to elicit key themes.

46 **Results**

47 Five main themes (parental experiences of prenatal WES, need for information,

48 consent/reasons for prenatal WES, sources of support for prenatal WES, and return

49 of WES findings to families) emerged, some with multiple sub-themes.

50 **Conclusions**

Parents desired as much information as possible and appreciated information being 51 repeated, and provided in various formats. Many struggled with clinical uncertainty 52 relating to the cause and prognosis following a fetal anomaly diagnosis, and found it 53 difficult to balance the risks of invasive testing against their need for more definitive 54 information. Parents trusted their clinicians and valued their support with decisions 55 in pregnancy. Testing was sometimes pursued to reassure parents that their baby 56 was 'normal' rather than to confirm an underlying genetic problem. Parents were 57 motivated to undergo WES for personal and altruistic reasons but disliked waiting 58 times for results, and were uncertain about what findings might be returned. 59

60 Key words: Prenatal; whole exome sequencing; parents views

61 Introduction

Structural anomalies are diagnosed by ultrasound in up to 3% of pregnancies^[1].
Fetal outcome is variable depending on the type of abnormalities identified, and the
underlying genetic aetiology^[2]. Determining the potential cause of fetal anomalies
enables a more accurate diagnosis and provides prognostic information relating to

the pregnancy and the likely risk of recurrence^[2]. Genetic testing is available for 66 parents following the identification of a fetal anomaly and recent advances in 67 molecular genetics are enabling increasingly detailed prenatal genetic 68 investigation^[3]. Prenatal genetic diagnosis is of significant value to parents and can 69 assist with prospective planning for optimal perinatal management^[4]. It may also 70 provide a means to inform parental decisions regarding the continuation or 71 termination of an affected pregnancy. Currently in the UK prenatal genetic testing 72 involves increasingly routine QF-PCR (Quantitative Fluorescence-Polymerase Chain 73 Reaction) and CMA (Chromosomal Microarray) to identify chromosomal differences 74 and variations in copy number (CNVs). Targeted genetic sequencing of exonic 75 regions is used to detect single nucleotide variants (SNVs) associated with various 76 single gene disorders, but this modality has limited potential to identify CNVs. 77 Whole genome sequencing (WGS) approaches are beginning to be used and have 78 the ability to detect CNVs. 79

80 Next-Generation Sequencing (NGS) and the Prenatal Application of Whole

81 Exome Sequencing (WES) Approaches

NGS applications are broadening the scope of prenatal diagnosis to identify the 82 genetic aetiology of sporadic and inherited disease^[5] and are revolutionising current 83 practice in prenatal diagnostics^[6]. Sequencing analysis of trio (fetal and biparental) 84 DNA can identify genetic alterations that are potentially causative of fetal 85 abnormalities, but this technology is only recently being evaluated within prenatal 86 medicine. WES captures the majority of regions that encode proteins to identify 87 SNVs and small insertions and deletions (indels)^[5]. As a technique it has proved 88 useful to the diagnosis of known genetic disease and to the discovery of novel 89

disorder genes,^[7] and is increasingly being used to diagnose rare Mendelian 90 conditions (when standard tests are uninformative)^[8]. The use of WES in prenatal 91 diagnosis is potentially advantageous as its accuracy enables personalised care, 92 prospective risk assessment and preventative fertility treatment, reproductive genetic 93 counselling and family planning^[9]. As such, if a definitive diagnosis is made this 94 testing may aid understanding of aetiology, potential co-morbidities and risk of 95 recurrence. However, NGS, in the prenatal setting, presents potential challenges 96 around the interpretation of results, especially if positive results are not thought to be 97 'causative' or are of unknown significance. The detection of these secondary and/or 98 incidental findings (ICFs), may have significant and morbid emotional effects on the 99 parents and also impact negatively on parental decision making in the prenatal 100 setting. 101

Several studies involving WES in patients with Mendelian disease have 102 demonstrated a diagnostic yield in the order of 25%^[10-11]. This indicates that WES is 103 complementary to conventional prenatal diagnostic techniques^{[12].} Research relating 104 to the use of genetic sequencing for prenatal diagnosis in on-going pregnancy is 105 limited,^[4] however, the feasibility of WES in prenatal diagnostics has been 106 demonstrated in small case series^[3,12,13]. Survey data involving 186 expectant 107 parents in the USA demonstrated that 83% thought that prenatal WES should be 108 offered,^[14] and research into the views of fifteen women with non-continuing 109 anomalous pregnancies found that they had high hopes and expectations of WES, 110 despite testing enabling a diagnosis in only 1 in 3 (30% of cases)^[15]. 111

Successful implementation of WES for prenatal diagnosis would require rigoroushealth economic assessment, and would be dependent upon the development of

rapid analytical and interpretation pipelines^[12]. Sequencing findings would need to 114 be available within a timeframe that would assist parents to make informed decisions 115 relating to the affected pregnancy, and this will only be possible when the knowledge 116 base relating to the genetic causes of prenatal structural anomalies is significantly 117 developed^[12]. The challenge of prenatal WES will be the integration of sequencing 118 analysis into prenatal diagnostics as part of a responsible and ethical framework for 119 clinical practice^[2]. Currently, the PAGE consortium project funded by the Department 120 of Health/Wellcome Trust is prospectively recruiting parent/fetus trios across the UK 121 to investigate the prenatal use of WES as a diagnostic tool in structurally abnormal 122 fetuses^[16]. The study will analyse ~1000 trio whole exomes with the aim to elucidate 123 the relative contribution of different forms of genetic variation to prenatal structural 124 anomalies. 125

As the use of WES increases, and transfers from the research setting to routine 126 care, it will be important to ensure a streamlined approach to the integration of 127 genomic analysis to existing prenatal care pathways. This transition will require an 128 129 understanding of parental acceptability and expectations around sequencing analysis for prenatal diagnosis following discussion with parents who have personal 130 experience of this type of genetic testing in pregnancy. These parents will provide a 131 unique perspective on their experiences as it is important to ensure that this 132 technology is translated into clinical care because parents consider it to be of value. 133 The views of parents who have undergone genetic sequencing for prenatal diagnosis 134 have not been formally explored using qualitative interview methodology. The aim of 135 this research was to gain insight into the experience of parents who have undergone 136 prenatal WES following a fetal anomaly diagnosis, to understand more about what 137

influenced their decision-making to have testing, and elicit their beliefs around how
they perceived WES to be of potential benefit. Qualitative methods allow for
exploration of parental experiences, beliefs and feelings around the use of prenatal
WES in a way that quantitative methodology cannot. It is important to understand
parental views around prenatal sequencing to inform the routine use of these
technologies in the future.

144 Method

A purposeful sample^[17] (i.e. parents who had undergone WES for prenatal diagnosis 145 following enrolment in the PAGE Study^[16]) was selected to participate in this 146 research. All parents had received focussed pre-test counselling for approximately 147 one hour from a fetal medicine specialist regarding standard invasive prenatal testing 148 options (QF-PCR and CMA), non-invasive prenatal testing (NIPT) for common 149 aneuploidy, and WES as part of the PAGE Study. All were informed prior to testing 150 151 that trio analysis (biparental/fetus) would be performed and that results would not be available within the timeframe of their pregnancy. It was also explained that only 152 pathological findings considered to have contributed to the fetal phenotype would be 153 returned, and that no uncertain, secondary or incidental information would be 154 reported. They were also told that WES could potentially detect up to 10% more 155 causes for fetal structural anomalies above standard testing based on exiting 156 evidence. Fifteen women were approached at random by EQJ (research midwife) 157 during their appointments at the Birmingham Women's Hospital Fetal Medicine 158 Centre and asked to participate in an interview, three of whom declined without 159 giving a reason, thus the study sample composed twelve women (Figure 1). It was 160 161 anticipated that if data saturation was not reached after twelve interviews then more

interviews would continue until data saturation was achieved. Interviews were 162 undertaken either at the hospital or at home depending on parental preference. A 163 topic guide was used to guide guestioning, and interviews were carried out by EQJ 164 with each interview lasting approximately 30 to 45 minutes. Development of the 165 topic guide was informed by related focus group research with stakeholders 166 undertaken by EQJ and others^[18]. Women were interviewed alone, or with their 167 168 partner/or other close family member. All women spoke English although this was not a criterion for inclusion. The timing of interviews varied, but all were carried out 169 within two weeks of parents giving consent for WES. Issues explored with parents 170 included their personal experience of prenatal genetic testing and diagnosis, and 171 what they remembered and understood regarding WES. Parents were asked about 172 their expectations and concerns relating to prenatal genetic sequencing, and about 173 the factors that influenced them to undergo testing, including the information they 174 required to inform their decision. Interviews were digitally recorded and transcribed. 175 176 National Research Ethics Service approval to undertake this study was granted by West Midlands - South Birmingham Committee (REC Reference 14/WM/0150). 177

178 Analysis

Analysis of the interview data followed a standard thematic approach^[19]. Transcripts were read by EQJ to enable familiarisation. Using an inductive process^[20] the transcripts were then coded for similarities and differences in content to develop a coding frame. Encompassing key themes with underpinning sub-themes were produced by combining the identified codes. Two transcripts were independently read by SCH (clinical co-facilitator for aforementioned focus groups and interview design) who similarly used thematic analysis to elicit themes^[19]. The coding frame

developed by EQJ was shared with SCH and was subsequently modified. The 186 coding frame and agreed themes were shared with SMG (medical sociologist). Two 187 further transcripts were analysed by SMG using the established coding frame. 188 Further amendments to the coding frame were not thought necessary as a result of 189 this analysis. All three researchers met to reach a consensus that the themes 190 identified were indeed reflective of the accounts provided. A rapid analysis of the 191 192 interview transcripts was then undertaken by EQJ to ensure completeness and assess for data saturation^[19]. A consensus decision by the three researchers was 193 made that data saturation had occurred and that no further interviews were required. 194

195 **Results**

Participants were diverse with regard to age, ethnicity, parity and gestation, and had 196 varying diagnoses of both isolated and multiple fetal structural anomalies (Figure 1). 197 Women were aged between 21 and 38, and identified themselves as Caucasian, 198 199 Black African or Asian, with Caucasian women comprising 75% of the sample. Of the 12 women interviewed 7 (58%) were multiparous and gestational ages ranged 200 from 12 to 38 weeks. There was an equal split between isolated and multiple 201 structural abnormalities and the prognosis for fetuses were variable and sometimes 202 uncertain. 203

Five main themes emerged some with multiple sub-themes (Figure 2).

205 Theme One: Parental experiences of prenatal WES

206 Parents sometimes struggled to balance the risks of invasive testing against the

207 perceived benefit of receiving a genetic diagnosis, particularly if there was

208 uncertainty relating to the ultrasound features and the prognosis for the baby (this

was especially true if there was a previous history of miscarriage and any associatedtraumatic memories)

"It was more the risk factor because I had a miscarriage last year and it was
really horrible so we didn't want to go through that again, especially as I was
well over 20 weeks and into my second trimester and the baby was fully
formed, so that was quite worrying but it wasn't so much for the results"
(Interview 4 – Mother)

Parents felt shocked when first told that their baby had a congenital difference, but 216 this initial shock was often replaced with on-going anxiety. Some said it felt as 217 though a 'heavy weight' had been placed upon them, and found the experience to be 218 extremely scary. Parents appeared to worry more about the uncertain prognosis for 219 the baby and less about the genetic findings that testing might identify. Many 220 remembered feeling overwhelmed by the different tests available, and felt that their 221 222 worries and concerns were compounded because they had so much to think about 223 at the time:

"It was scary to be honest with you, all the different tests and constant worry.
It was worrying because we didn't know what she (baby) would look like or
anything like that" (Interview 2 – Father)

Self-blame that they had done something to have caused the fetal anomaly was a common parental concern, thus a desire for reassurance that this was not the case was reported. All parents described that they trusted their clinicians and valued receiving their clear explanations. Parents described that they were assisted in their decision-making when they felt supported by clinicians, and believed that any

prenatal testing options discussed by the consultant overseeing their care would berelevant and useful which reassured them:

"We thought that it would give us some reassurance and help us plan and
prepare for the future" (Interview 4 – Father)

Some parents described how they tried to remain hopeful for a good pregnancy outcome, but also felt that they would love the baby regardless of any disability they may have. Some remembered consciously blocking out their concerns in an attempt to keep positive, believing that searching out more information would only serve to exacerbate their worry. When faced with various options, parents felt that they could make difficult decisions if they were not pressurised and were given enough time:

"I think we've tried to blank quite a lot of it because we don't want to be
negative. When she is here we will cross that path won't we?" (Interview 2 –

244 Mother)

245 **Theme Two: Need for information**

A desire for information to understand more about the anomaly affecting their baby and the different testing and treatment options available was universally reported by parents. Parents needed to ask lots of questions of their clinicians as they tried to balance the pros and cons of testing:

250 *"More information is all good because it helps us understand whatever it is.*"

- 251 You can prepare yourself and your family and do what you possibly can with
- the information that you are given" (Interview 1 Father)

A need for repetition of complex information was also evident as parents found it
difficult to fully understand everything that they were told at the initial consultation.
Discussion and explanation on more than one occasion was found to be helpful, and
parents appreciated receiving clinical details in written format relating to the specific
anomalies identified:

258 "Some things you don't understand, some of the things the doctor says"

259 (Interview 2 – Mother)

- 260 "But when they break it down into smaller (pieces), all these big words like,
- 261 and obviously we don't know what they mean, but they do break it down"

262 (Interview 2 – Father)

263 Theme Three: Consent and reasons for prenatal WES

Desiring more information and a wish to rule out as much as possible was a key motivator for parents to undergo prenatal testing. Parents perceived WES as a more detailed assessment to find out additional genetic causes for the anomalies affecting their baby that are not tested for routinely. They considered more information to be the best thing for parents and the baby and this was often the main reason for testing:

"It was going to test for more than everything else, and if there was anything
rare that it is more likely to pick that up, and he explained that it will take much

272 *longer" (Interview 10 – Mother)*

Parents were aware that testing involved looking for differences and similarities
between their individual DNA and the DNA of their baby. It was understood that the
testing would not benefit the current pregnancy (because results would not be

276 reported back within the timeframe of pregnancy), but thought that it may be helpful
277 for the baby when older, or if it could provide information for future pregnancy
278 planning:

"It was to try and work out if there is anything between us (parents) that has
caused the anomalies. I do not know whether it searches for one or both or
whatever, but just that it is trying to find out if there is anything that is within
either of us that has made these things happen in the baby" (Interview 3 –
Mother)

Parents were sometimes uncertain about what was actually being tested for or ruled out and would have appreciated hearing about some example conditions. Most felt that it was better to know about any genetic causes and hoped that the testing would provide answers which would be reported back to them:

"I would like to know about what other things they test for because I asked
them and they said they would test for over 200 things but I would have liked
examples because that was still worrying me" (Interview 5 – Mother)

Parents described their decision to have prenatal WES as an opportunity to help others in the future. Altruistic motivations involved feeling that it was important to gather more information on the genetic causes of fetal anomalies, and viewed their participation as a means to contribute to research and the progression of medical knowledge:

296 *"I was kind of contributing to something really, to help others in the future. It is*297 *the only way you are going to learn and evolve in the medical field. If you can*

298

299

achieve anything with it then I would be more than happy" (Interview 7 – Mother)

300 Theme Four: Sources of support for prenatal WES

Electronic and written sources of support were helpful to parents when faced with the decision of whether to undergo prenatal sequencing. Many opted to avoid the internet due to a perceived risk of inaccurate information. Parents felt that some internet sources showed the extremes of disease and were not always relevant. Some accessed NHS websites and Wikipedia feeling that these were more trustworthy sources. Information leaflets on specific conditions were generally found to be helpful:

- We got advice before we came here as well to steer clear of the internet
 because obviously you get a lot of misinformation, so I kind of took that to
- 310 heart as it sounded quite sensible so I have not really been googling"

311 *(Interview 1 – Father)*

Interactive sources of support were reported to be helpful and parents valued being able to ask questions directly. One couple described that they would have liked to speak to other parents with similar experiences, suggesting that a workshop where they could find out more information and ask questions could be a forum for this:

316 *"Maybe a workshop held by the hospital or midwife that is solely dedicated to*

- 317 this as part of their job, where they would have all the knowledge and can
- 318 educate families, and where parents can come together and share their

319 *experiences" (Interview 4 – Father)*

320 Theme Five: Return of prenatal WES findings to families

Presently, the prenatal WES 'clinical pipeline' within the PAGE Study^[16] takes up to 321 twelve months. The delay in receiving results was felt by some parents to have 322 prolonged their worry and anxiety. Parents still wished to have WES even though 323 324 they knew that there would be a significant wait for results believing that they would still rather know than not know about any relevant genetic findings. Many described 325 that having this information eventually would help their understanding and better 326 327 equip them to cope and prepare for any challenges ahead. Some felt that more information in time relating to the risk of recurrence was worth waiting for and would 328 possibly assist them with future pregnancy planning: 329

330 *"That was what I hated, just waiting (for results)" (Interview 5 – Mother)*

Some parents were uncertain regarding the process by which results would be returned and would have appreciated having this better explained to them. Some parents preferred to return to the hospital and have the results explained by familiar clinicians face-to-face. All were happy for their information to be stored and shared with other clinicians and researchers involved in prenatal diagnosis, and although some said that they preferred their personal information to be anonymised, others were less concerned about protecting their identity:

"If there was anything (genetic results) we would like to come back here
(hospital) and sit down and discuss it face-to-face with you guys (medical
team) because we are comfortable with you" (Interview I – Mother)

341 Conclusions

This is the first qualitative interview study exploring parental experiences of WES for prenatal diagnosis. The findings are important because they are novel in this context

and contribute to a limited body of evidence relating to parental experiences of 344 prenatal sequencing in structurally abnormal fetuses. Given the potential for NGS 345 techniques to detect genetic alterations that are causative of various developmental 346 fetal anomalies, it is likely that prenatal sequencing will be integrated into existing 347 prenatal care pathways in the foreseeable future. Transition from the research 348 setting to the clinic will require an assessment of the acceptability of prenatal 349 350 sequencing for genetic diagnosis to evaluate if testing is desirable to parents. This research has highlighted the views of parents who have undergone WES for prenatal 351 diagnosis, and provided insight in to their decision-making to proceed with testing, 352 and what they perceived the potential benefits of WES to be. Facilitating appropriate 353 consent for testing was highlighted by parents as extremely important, who felt that 354 they needed clearer information regarding what WES might identify, and what, when 355 and how results would be returned. If WES is to be routinely available for prenatal 356 diagnosis this will require the development of national and international guidance 357 358 that encompass the consent procedure, as well as the option for parents to opt in or out of receiving information which is not directly related to the prenatal findings (i.e. 359 the primary indication for testing) both for the unborn baby and for themselves^[13]. 360 361 Inevitably, prenatal WES and the interpretation of results will become more rapid and clinical usefulness will be significantly improved. Likewise, contribution of parental 362 views around prenatal WES will assist with streamlining the clinical use of the 363 technology for diagnostic purposes. However, CMA research indicates that variants 364 of uncertain significance (VUS) may continue, in a small number of cases, to have 365 morbid emotional consequences^[21,22]. The need for public debate around the use 366 and potential benefits, as well as the drawbacks of prenatal genetic diagnosis is 367

clear, to facilitate the general acceptance and integration of sequencing techniques
 into routine prenatal care^[23].

370 Limitations

It is acknowledged that the views expressed by some parents (such as the need for 371 more information to balance risks, feelings of self-blame, and consciously blocking 372 out concerns to remain positive), are likely to be applicable to any couple whose 373 baby has ultrasonographically detected fetal anomalies irrespective of whether they 374 decide to undergo prenatal testing (including WES). As such, these particular 375 376 findings are not necessarily unique in this context. This research explored the experience of parents who underwent WES for prenatal diagnosis at one large UK 377 fetal medicine centre and parents at other centres (within the UK or internationally) 378 may have different views. The opinions of parents who declined WES are similarly 379 not well represented. It cannot be assumed therefore that the findings are applicable 380 381 to all parents; moreover they may not reflect the views of parents who decline genetic diagnosis using invasive methods. Further research that considers the 382 opinions of parents who decline prenatal sequencing is needed. Ethical approval for 383 the PAGE Study^[16] only permitted the return of results to families considered to be 384 pathogenic and contributing to the prenatal phenotype, thus it was not possible to 385 explore parental views around the return of VUS and ICFs. Parental opinions 386 regarding the return of VUS and ICFs will be explored in a planned further phase of 387 work. 388

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393 References

- ^[1] Springett A. Congenital anomaly statistics. In: Morris J. (Ed) Congenital Anomaly
 Statistics: England and Wales. London, British Isles Network of Congenital Anomaly
 Registers, 2010
- ^[2] Hillman SC. Williams D. Carss KJ. *et al.* Prenatal exome sequencing for fetuses
- 398 with structural abnormalities: the next step. Ultrasound in Obstetrics and
- 399 Gynaecology 2015 (45):4-9
- ^[3] Pangalos C. Hagnefelt B. Lilakos K. First applications of a targeted exome
- 401 sequencing approach in fetuses with ultrasound abnormalities reveals an important
- 402 fraction of cases with associated gene defects. Peer Journal 2016 4:e41955;DOI
- 403 10.7717/peerj.1955
- ⁴⁰⁴ ^[4] Van den Veyver IB. Eng CM. Genome-wide sequencing for prenatal detection of
- 405 fetal single-gene disorders. Cold Spring Harbour Perspectives in Medicine 2015
- 406 (5):a023077
- ^[5] Babkina N. Graham JM. New genetic testing in prenatal diagnosis. Seminars in
 Fetal & Neonatal Medicine 2014 (19):214-219
- ^[6] Rabbani B. Tekin M. Mahdieh N. The promise of whole exome sequencing in
- 410 medical genetics. Journal of Human Genetics 2014;(59):5-15

^[7] Sawyer SL. Hartley T. Dyment DA. *et al.* Utility of whole-exome sequencing for
those near the end of the diagnostic odyssey: time to address gaps in care. Clinical
Genetics 2016 (89):275-284

^[8] Volk A. Conboy E. Wical B. *et al.* Whole-exome sequencing in the clinic: Lessons
from six consecutive cases from the clinician's perspective. Molecular Syndromology
2015 (6):23-31

^[9] Peters DG. Svetlana A. Yatsenko MD. *et al.* Recent advances of genomic testing
in perinatal medicine. Seminars in Perinatology 2015 Feb;39(1):44-54

^[10] Yang Y. Muzny DM. Reid JG. *et al.* Clinical whole-exome sequencing for the

diagnosis of Mendelian disorders. New England Journal of Medicine 2012

421 (369):1502-1511

422 ^[11] Yang Y. Muzny DM. Xia F. et al. Molecular findings among patients referred for

clinical whole-exome sequencing. JAMA 2014 November 12;312(18):1870-1879

^[12] Carss KJ. Hillman SC. Parthiban V. *et al.* Exome sequencing improves genetic

425 diagnosis of structural fetal abnormalities revealed by ultrasound. Human Molecular

426 Genetics 2014 23(12):3269-3277

^[13] Drury S. Williams H. Trump N. et al. Exome sequencing for prenatal diagnosis of

428 fetuses with sonographic abnormalities. Prenatal Diagnosis 2015 (35):1010-1017

^[14] Kalynchuk EJ. Althouse A. Parker LS. *et al*. Prenatal whole-exome sequencing:

430 parental attitudes. Prenatal Diagnosis 2015 (35):1030-6

431	^[15] Vora NL. Brandt A. Strande N. <i>et al.</i> Prenatal exome sequencing in anomalous
432	fetuses: New opportunities and challenges. 2017 Genetic Medicine in press
433	^[16] Prenatal Assessment of Genomes and Exomes (PAGE) Study[homepage on the
434	internet]. Cambridge, Wellcome Trust Sanger Institute; 2017 [cited 07 Mar 2017].
435	Available from: www.pageuk.org
436	^[17] Patton MQ. Qualitative research and evaluation methods, 3 rd Edition, Sage
437	Publications, Thousand Oaks; 2002
438	^[18] Quinlan-Jones E. Kilby MD. Greenfield S. et al. Prenatal whole exome
439	sequencing: the views of clinicians, scientists, genetic counsellors and patient
440	representatives. Prenatal Diagnosis 2016 Oct;36(10):935-941
441	^[19] Boyatzis RE. Transforming Qualitative Information. Thematic Analysis and Code
442	Development. London, Sage Publications; 1998
443	^[20] Robinson A. Phenomenology. In: Cluett ER. Bluff R. editors. Principles and
444	practice of research in midwifery. Philadelphia, Bailliere Tindall; 2000
445	^[21] Bernhardt BA. Soucier D. Hanson K. et al. Women's experiences receiving

abnormal prenatal chromosomal microarray testing results. Genetic Medicine 2013

447 (15):139-145

^[22] Hillman SC. Skelton J. Quinlan-Jones E. *et al.* 'If it helps...' The use of microarray

technology in prenatal testing: patient and partners reflections. American Journal of

450 Medical Genetics A 2013 (161A):1619-1627

^[23] Vermeesch JR. Voet T. Devriendt K. Prenatal and pre-implantation genetic

diagnosis. Nature Reviews Genetics 2016 (17):643-656