# UNIVERSITY<sup>OF</sup> BIRMINGHAM

**Research at Birmingham** 

# Prenatal whole exome sequencing

Quinlan-Jones, Elizabeth; Kilby, Mark; Greenfield, Sheila; Parker, Michael; McMullan, Dominic; Hurles, Matthew E; Hillman, Sarah C

DOI: 10.1002/pd.4916

License: Other (please specify with Rights Statement)

Document Version Peer reviewed version

Citation for published version (Harvard):

Quinlan-Jones, E, Kilby, M, Greenfield, S, Parker, M, McMullan, D, Hurles, ME & Hillman, SC 2016, 'Prenatal whole exome sequencing: the views of clinicians, scientists, genetic counsellors and patient representatives', Prenatal Diagnosis. https://doi.org/10.1002/pd.4916

Link to publication on Research at Birmingham portal

#### Publisher Rights Statement:

This is the peer reviewed version of the following article: Prenatal whole exome sequencing: the views of clinicians, scientists, genetic counsellors and patient representatives, which has been published in final form at 10.1002/pd.4916. This article may be used for non-commercial purposes in accordance with Wiley Terms and Conditions for Self-Archiving.

#### **General rights**

Unless a licence is specified above, all rights (including copyright and moral rights) in this document are retained by the authors and/or the copyright holders. The express permission of the copyright holder must be obtained for any use of this material other than for purposes permitted by law.

• Users may freely distribute the URL that is used to identify this publication.

• Users may download and/or print one copy of the publication from the University of Birmingham research portal for the purpose of private study or non-commercial research.

User may use extracts from the document in line with the concept of 'fair dealing' under the Copyright, Designs and Patents Act 1988 (?)
Users may not further distribute the material nor use it for the purposes of commercial gain.

Where a licence is displayed above, please note the terms and conditions of the licence govern your use of this document.

When citing, please reference the published version.

#### Take down policy

While the University of Birmingham exercises care and attention in making items available there are rare occasions when an item has been uploaded in error or has been deemed to be commercially or otherwise sensitive.

If you believe that this is the case for this document, please contact UBIRA@lists.bham.ac.uk providing details and we will remove access to the work immediately and investigate.

2	Counsellors and Patient Representatives
3	
4	Elizabeth Quinlan-Jones <sup>1</sup> , Mark D. Kilby <sup>1,2,5</sup> , Sheila Greenfield <sup>3</sup> , Michael
5	Parker <sup>4</sup> , Dominic McMullan <sup>6</sup> , Matthew Hurles <sup>7</sup> , Sarah Christine Hillman <sup>1,5</sup> and
6	for the Prenatal Assessment of Genomes and Exomes (PAGE) Study
7	Collaborative Group
8	<sup>1</sup> Fetal Medicine Centre, Birmingham Women's NHS Foundation Trust, Edgbaston,
9	Birmingham, B15 2TG, UK
10	<sup>2</sup> Birmingham Centre for Women's and New-born Health, College of Medical and
11	Dental Sciences, University of Birmingham, Edgbaston, Birmingham, B15 2TT, UK
12	<sup>3</sup> Institute of Applied Health Research, College of Medical and Dental Sciences,
13	University of Birmingham, Edgbaston, Birmingham, B15 2TT, UK
14	<sup>4</sup> The Ethox Centre, Nuffield Department of Population Health, University of Oxford,
15	Old Road Campus, Oxford, OX3 7LF, UK
16	<sup>5</sup> Institute of Metabolism and Systems Research, College of Medical and Dental
17	Sciences, University of Birmingham, B15 2TT, UK
18	<sup>6</sup> West Midlands Regional Genetics Laboratory, Birmingham Women's NHS
19	Foundation Trust, Edgbaston, Birmingham, B15 2TG, UK
20	<sup>7</sup> Wellcome Trust Sanger Institute, Wellcome Trust Genome Campus, Hinxton,
21	Cambridge, CB10 1SA, UK
22	Running title: Views on prenatal whole exome sequencing (WES)

Prenatal Whole Exome Sequencing; the Views of Clinicians, Scientists, Genetic

#### 23 Manuscript word count: 3496

- 24 Corresponding author: Professor Mark D. Kilby, Birmingham Centre for Women's
- and New-born Health, College of Medical and Dental Sciences, University of
- Birmingham, Edgbaston, Birmingham, B15 2TT, UK, Tel: 0121 627 2778, Fax: 0121
- 27 623 6875, Email: M.D.Kilby@Bham.ac.uk

# 28 Funding

- 29 The PAGE project is funded by the Wellcome Trust/Department of Health Innovation
- 30 Challenge Fund (HICF R7-396).
- 31 We are grateful for the participation of the following charities in this work Antenatal
- 32 Results and Choices (ARC), Genetic Alliance UK, SWAN (Syndromes Without A
- Name) and Unique (the rare chromosome disorder support group).

#### 34 **Conflict of interest**

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

#### 36 What is already known about this topic?

- Prenatal WES generates variants of uncertain significance (VUS) and
- 38 incidental findings (ICFs).

#### 39 What does this study add?

- Consent-takers require training.
- An overview of the findings that will/won't be reported should be provided.
- Patient Representative Groups (PRGs) felt women want access to all
- 43 information and re-interpretation of results over time.

44

 Clinical Professionals (CPs) felt that interpretation should be at the point of testing only.

46

45

#### 47 Abstract

# 48 **Objective**

- 49 Focus groups were conducted with individuals involved in prenatal diagnosis to
- 50 determine their opinions relating to WES in fetuses with structural anomalies.

#### 51 Method

- 52 Five representatives of patient groups/charities (PRGs) and eight clinical
- 53 professionals (CPs) participated. Three focus groups occurred (the two groups
- separately and then combined). Framework analysis was performed to elicit themes.

55 A thematic coding frame was identified based on emerging themes.

#### 56 **Results**

57 Seven main themes (consent, analysis, interpretation/reinterpretation of results,

58 prenatal issues, uncertainty, incidental findings, and information access) with sub-

themes emerged. The main themes were raised by both groups, apart from

60 'analysis' which was raised by CPs only. Some subthemes were raised by PRGs

- and CPs (with different perspectives). Others were raised either by PRGs or CPs,
- 62 showing differences in patient/clinician agendas.

#### 63 **Conclusions**

64 Prenatal consent for WES is not a 'perfect' process but consent takers should be

- 65 fully educated regarding the test. PRGs highlighted issues involving access to
- results feeling that women want to know all information. PRGs also felt that patients

want re-interpretation of results over time whilst CPs felt that interpretation should be
performed at the point of testing only.

69 **Key words:** Prenatal; genetic testing; whole exome sequencing.

70

#### 71 Introduction

Standard chromosome testing (G-band karyotyping) undertaken prenatally has been 72 largely superseded by the use of chromosomal microarrays (CMAs)<sup>1</sup> identifying sub-73 microscopic rearrangements undetectable by conventional cytogenetic methods<sup>2</sup>. 74 Next generation sequencing (NGS) technologies represent a further development in 75 terms of the quantity of genetic data obtainable and the bioinformatics needed to 76 fully utilise and interpret results<sup>3</sup>. NGS offers knowledge, but comes with 77 challenges<sup>4</sup>. Genome wide testing produces huge quantities of information, some of 78 which may be uncertain and/or unanticipated, raising ethical concerns about 79 disclosure and stimulating debate regarding how best to integrate such testing into 80 prenatal clinical practice<sup>5</sup>. 81

Within the postnatal/paediatric setting parents value being able to choose the types of genetic information they wish to receive and their understanding of the different options for the return of findings (and the implications of receiving different kinds of results) can be facilitated by the consent process<sup>6</sup>. Parents do not express desire to know any and all genetic findings<sup>7</sup>, rather they prefer to receive information that they consider to be actionable, allowing them to balance the possible benefits and harms of learning their children's genetic results<sup>8</sup>. Parents can sometimes find themselves in uncharted territory needing to decide which types of findings (beyond primary
variants) to receive<sup>7</sup>.

There is little guidance relating to the process and content of informed consent for 91 whole genome sequencing (WGS) and whole exome sequencing (WES) in the 92 prenatal setting or the means by which results should be reported back to families<sup>9</sup>. 93 Despite uncertainties, sequencing technologies are being introduced to clinical 94 practice and reduction in cost is focusing the need to evaluate the balance of 95 potential benefits and harms for patients undergoing prenatal genetic diagnosis<sup>10</sup>. A 96 significant barrier to the integration of WES/WGS into clinical care involves the 97 management of incidental findings (results that are not related to the patient's clinical 98 indication for testing)<sup>9</sup>. The issue is compounded by the biological time-frame of 99 pregnancy, which creates a sense of time pressure<sup>11</sup>. It is essential that we seek to 100 understand the impact WGS/WES (and the uncertainty associated with it) has on 101 families, if not we risk, potentially incorrectly, assuming families are making properly 102 informed decisions<sup>12</sup>. 103

As a first step to gain insight into the opinions of individuals with personal or 104 professional experience of WES within the prenatal setting, focus group sessions 105 were conducted with representatives of patient groups/charities (PRGs) that support 106 families undergoing genetic testing and genetic diagnosis, and clinical professionals 107 and clinical genetic scientists (CPs) involved in prenatal diagnosis to discuss the 108 issues. The aim of the focus group sessions with PRGs and CPs reported here was 109 110 to gain information to subsequently inform ethical guidance relating to prenatal genetic sequencing and to help design an interview schedule to be used to 111

understand the experiences of families undergoing prenatal WES as a further phaseof the work.

114

### 115 Method

To identify participants for the PRG and CP focus groups members of the Prenatal 116 Assessment of Genomes and Exomes (PAGE) Study working group used 117 convenience sampling<sup>13</sup> to contact individuals known to be experts in their field. 118 Three groups were held in succession during the afternoon of 9<sup>th</sup> October 2014. 119 The first focus group consisted of five PRGs from the charities; Antenatal Results 120 and Choices (ARC), Genetic Alliance UK, SWAN and Unique. The second focus 121 group consisted of eight CPs (two fetal medicine consultants, two genetic 122 counsellors, two consultant clinical geneticists and two clinical genetic scientists). 123 The third focus group combined all thirteen participants of the first and second focus 124 groups. The focus groups were conducted by SH, EQJ and MP using a topic guide; 125 the main areas covered are shown in Table 1. We held separate focus groups of 126 PRGs and CPs first in order to allow for any topics to be discussed that might not be 127 discussed in the presence of the other group. The third group used the same topic 128 guide, but focused on areas that had been felt by the facilitators to be areas of 129 differences of opinion (between the CPs and PRGs) during focus groups one and 130 two. The size of the focus groups was limited by the number of professionals we 131 could assimilate geographically on the same day. All participants gave written 132 consent. Ethical approval for the focus groups was provided by the NRES 133 Committee West Midlands - South Birmingham (14/WM/0150). 134

#### 136 Analysis

The focus groups were voice recorded and then transcribed verbatim. Data was 137 analysed using a thematic approach<sup>14,15</sup>. To gain familiarization with the data the 138 transcripts were read and re-read by SH and EQJ. Throughout this process key 139 ideas and recurrent themes were noted. A coding frame was then identified based 140 on the emerging themes. The coding frame was refined as transcripts were added. 141 This was agreed between three authors (SH, EQJ and SG). All text was indexed 142 numerically, with numbers placed in the margin beside the text. The original pieces 143 of data were charted using Excel (©Microsoft Office 2010). Charts were developed 144 using themes and subthemes. 145

146

#### 147 **Results**

The thirteen participants came from four different charities and six healthcare sites 148 within three geographical areas of the UK (Table 2). Seven main themes with sub-149 themes were identified. With the exception of theme two, 'Analysis', which was 150 raised by CPs only (FG1) all themes were discussed by both PRGs and CPs (FG1 151 and FG2). Within those seven main themes some similar subthemes were either a) 152 raised by both groups (with similar or different opinions) or b) different subthemes 153 were raised by the separate groups, showing a difference in the patient and clinician 154 agenda (Table 3). Quotations with their focus group identifier (FG1, FG2, and FG3) 155 are used to reflect the themes and sub-themes. 156

#### 157 **Theme One: Consent**

The first theme, consent, was discussed by PRGs and CPs. Much of the discussion focused on the problem of consenting for a complicated test and the time/resources that facilitating informed consent would require. In addition, the possibility of an 'opt in' consent form was discussed whereby patients could give different levels of consent depending on the results they wanted to receive.

Both PRGs and CPs expressed concern about how much detailed informationshould be given in the consent process:

165 CP "There seems to be a variation and divergence of opinion between clinical

166 geneticists and the clinicians that deal with the parents as to how much information

167 needs to be provided about problems that are clearly not related to the indication or

reason for testing and I think that is my major concern" FG2

Both expressed concern about who would obtain consent and the possibility of an 'education gap' if those taking consent did not have a full understanding of the testing:

172 PRG "Who is going to be doing all this counselling? It can't possibly be Geneticists,

it's going to be non-genetics professionals and I think there is a huge education gap

174 there which needs filling" FG1

Both also expressed concern about adding pressure to 'overstretched services' and the time it would take to consent for prenatal WES given the scope of what it could report:

PRG "I think there is a worry too about the pressure it puts on genetics, pressure on genetic counsellors, because it is all, certainly in the first instance, going to be

180 focused on them and they are already stretched" FG1

Finally both PRGs and CPs discussed the option of an 'opt in' consent form whereby patients could choose to receive findings of incidental significance in addition to results relating to the primary reason for testing. PRGs felt this was something patients would welcome. CPs however felt that this type of consent would need to be taken by a clinical geneticist or genetics counsellor.

An area discussed by PRGs only was motivation for testing. PRGs felt the most common motivation for testing was reassurance. Other motivators were recurrence risk, 'for extra information' and wanting a genetically perfect baby:

189 PRG "maybe there would be pressure for people to make sure their baby is

190 perfect...it is a bit of a nightmare really" FG1

#### 191 **Theme Two: Analysis**

This theme was only discussed by the CP group. This is not surprising given that the CP group contained clinical scientists. The potential to 'target' the testing to relevant genes was discussed. It was felt this would negate the problem of incidental findings but in practice would be difficult to achieve given the current limits of genetic knowledge:

197 CP "I kind of assume that you are going to do a target interpretation of that data and 198 what you target is going to affect how you consent so if you are not going to look at 199 BRCA1 and BRCA2 then you don't need to consent about it" FG2

#### 200 Theme Three: Interpretation/reinterpretation of results

Although this was discussed by both PRGs and CPs, there was a difference of

202 opinion between the groups regarding reinterpretation of results over time. CPs felt

results should be reviewed at the time of testing only. PRGs felt that patients wouldwant information as and whenever it became available:

PRG "Our families that we support, they live without knowing for years and years,
some of them, and their need for that diagnosis never goes away...if something five
years down the line came up and suddenly they could link that then those families
would most definitely want to know" FG3

209 *CP* "it is a unique situation in medicine where we might have to reinterpret a test that 210 was done for an entirely different reason five years ago in the context of what is 211 known now...if the mother or father had not reported it [a medical concern] and the 212 child hasn't been presented to a medical practitioner, do we have a right to go along 213 [contact the family] and say okay we found this relationship [genetic variation] exists 214 and disrupt this family when they have perceived no medical problem at all?" FG3 215 Only PRGs discussed access to the generated genetic data:

PRG "a high percentage of families said if you had knowledge about me, my child or
my baby, that is my knowledge and I want it, even to the point of wanting the raw
sequencing data" FG1

PRGs felt that women and their families wanted 'all' the information possible but that
when the test became a reality fewer may choose to receive results of uncertain
significance or incidental findings:

PRG "Experience from when the Huntington's test was made available on the NHS
was that the community wanted it and everyone would go for it and then in practice I
think it's about a third go for it...we think maybe this (WES) is the same thing again"
FG1

#### 226 Theme Four: Issues specific to prenatal WES

The reasons that prenatal exome sequencing is different from postnatal sequencing were explored by both PRGs and CPs. Both agreed that pregnancy is a uniquely stressful situation with a 'biological timeframe':

230 PRG "Your mind is jelly. It takes you weeks to get your mind working properly, even

if you are in the business, so God help people who have not even got the basic

knowledge of what genetic testing is and what it means" FG1

The PRGs alone discussed non-agreement between partners. They also discussed the difficulties that couples have prenatally making an 'imaginary leap' as to what they would do with results:

236 PRG "a lot of people they will nod their heads and make the right noises but they

237 might not have thought the consequences through and they are the ones when

something anomalous is picked up who will need the most time and concentration in

helping them to work out what the result means to them" FG1

The CPs group raised the issue that there is a more ambiguous phenotype

antenatally, for instance you cannot see neurodevelopment, and this is an obvious

242 limitation to counselling.

#### 243 Theme Five: Uncertainty

There is often uncertainty in prenatal counselling for structural fetal anomalies as the full phenotype may not be detectable on scan and a genetic diagnosis maybe associated with variable penetrance. Additionally WES detects variants where there is not enough definitive information to say that the genetic difference is the cause of the scan findings. These variants of uncertain significance (VUS) present difficultiesin the counselling of women if they are reported.

Both the CPs and PRGs agreed that reporting VUS to patients can have a negative

impact on the patient and potentially the doctor-patient relationship:

252 CP "The time that I have had patients really angry has been when I have been

reporting back uncertainty. They are in the middle of this situation where they are

trying to make a decision and I tell them something and then say "but I don't know

what that means" and I have had really angry reactions" FG2

However both groups also agreed that VUS should not be withheld:

257 PRG "there is a tremendous pressure when they (CPs) are giving information for

which they can give no real certainty...but I would not want that to take away from

the autonomy of that woman from making a decision to end the pregnancy if that is

what they [she would] want because the potential we have at the moment is to

261 potentially be paternalistic about the information given because of what might be

262 done with it" FG1

There was also consensus between the groups that VUS should be recorded in

databases to build up a picture of whether the variants are benign or pathological.

#### 265 **Theme Six: Incidental findings and prenatal WES**

WES is capable of detecting 'incidental findings' which are mutations which can sometimes associate with pathology. These findings are incidental because they are unrelated to the reason for testing. Reporting incidental findings will have implications for CPs' time and healthcare resources, and there was a difference between the views of CPs and PRGs. Some CPs felt that incidental findings shouldnot be reported:

CP "We don't have a national screening program [to identify incidental findings] (for
adults) so why are we doing screening by subterfuge [to detect such findings]
through the fetus" FG2

275 Other CPs discussed that there appears to be a progression to the reporting of

incidental findings postnatally if there is treatment for the condition available.

277 PRGs highlighted the potential injury to the relationship between patients and

278 medical professionals if an incidental result was revealed subsequently and it was

279 felt this information had been withheld.

#### 280 Theme Seven: Access to prenatal WES information

Both the PRGs and CPs agreed on the need for clear detailed written information to take away after the consultation. The PRGs suggested more detailed signposting or information sharing, particularly in relation to patient charities that could provide focused support to families. CPs also highlighted the need for national reporting guidance:

286 CP "There should be some written information. Ideally in this day and age and 287 definitely in 10 years there should be a dedicated website that they (parents) can 288 access and find out information" FG2

289 *CP "I think the ideal scenario would be to have national or even better international* 290 *criteria for what is a definite [pathological variant] and what is a VUS and therefore* 291 *you minimise the possibility [of uncertainty] for the parents" FG2*  292

#### 293 Conclusions

All themes, with the exception of 'Analysis', were discussed by both CPs and PRGs. Both groups generally had similar opinions. The process of consent for prenatal WES was considered and concerns were raised regarding the current lack of clinical geneticists/counsellors available to facilitate consent in prenatal clinical practice within the UK National Healthcare System.

They also discussed the depth of the consent prior to the test, particularly when 299 300 taken under stressful circumstances. Previously authors have commented "that it is virtually impossible to counsel in these circumstances"<sup>16</sup>. When pregnant women 301 find themselves in a stressful position, they may cope by complying with what they 302 believe is the health professional's recommendations<sup>17</sup>. It was generally agreed that 303 clinicians should do the best job possible pre-test but understand that the process 304 will not be perfect and that more detailed information should be provided to families 305 306 when genetic anomalies are found.

The issue of access to results was highlighted by the PRGs who felt that women would want to know all information generated as it was '*their genome*'. PRGs also felt that patients would ideally want reinterpretation of genetic information over time, for instance if a VUS was recorded and was later found out to be pathological. Conversely some CPs felt that interpretation should be performed at the point of testing only and that on-going review was unsustainable. This is in contrast to the views of Yu et al that propose "*results should be viewed as a dynamic, sustained*  resource of information that is available to an individual not only at a single point in
time, but over many years and even possibly a lifetime<sup>"18</sup>.

It was felt that conveying uncertain information could create tension in the doctor 316 patient relationship. In these circumstances patients require rapid follow up with a 317 consultant clinical geneticist. Even when this has occurred people may make 318 incorrect conclusions to fit with their own schemata<sup>12</sup>. Bernhardt et al interviewed 319 women with VUS. Many of them considered uncertainty to be information that they 320 wished they did not have ("toxic knowledge")<sup>2</sup>. Women were left feeling anxious, 321 and these concerns lingered into worries about their child's development. This 322 would be in opposition to recent research showing that patients consider all 323 information very important<sup>19</sup>. 324

Some CPs felt that we should not report genes relating to adult onset conditions and allow the sequencing to become a screening test. However there has been progression towards reporting of adult onset conditions in the postnatal arena (as per guidance by the ACMG<sup>20</sup>) and it seems possible that this may transfer into the prenatal setting. Srebniak et al found that 55% of future parents want to be informed about adverse health effects at an adult stage but did not make a distinction between treatable and non treatable conditions<sup>21</sup>.

The potential contrast in views of the CPs and PRGs is also highlighted in the recent publication of views of nearly 7000 people on the return of incidental results from genetic sequencing<sup>22</sup>. Here compared with the public, genetic health professionals were five times more likely to think that incidental findings should not be returned. Participants were more interested in learning about conditions that were preventable and less interested in receiving information that is uncertain and cannot be interpreted at the moment. It maybe that genetic health professionals anticipate a
vast increase in workload with the seemingly rapid progression towards the use of
sequencing in the prenatal and postnatal setting<sup>22</sup>. Recently Kalynchuk et al
surveyed parental attitudes to WES and found that 83% felt it should be offered and
54% would potentially accept it. Only 2.2% were opposed to the testing. However
over 70% reported an increased risk of adult onset conditions or a variant of
uncertain significance would cause them anxiety<sup>23</sup>.

345

### 346 Limitations

The number of focus group participants was limited by the number of CPs and PRGs 347 who could be brought together geographically. Therefore this is a relatively small 348 study. However it has been suggested that in qualitative work a small sample can 349 provide useful information about participants' experience <sup>24</sup>. The number of focus 350 groups we carried out accords with guidance for a 'small' study<sup>25</sup>, in which we were 351 seeking information to inform further work and we did not aim or claim to reach data 352 saturation<sup>26</sup>. We cannot comment on the extent to which the views expressed reflect 353 those of CPs and PRGs as a whole, and further themes such may have arisen had 354 we carried out further focus groups. There are a number of stakeholder groups 355 involved in WES. This paper has presented the views of two such groups and 356 although the patients' opinions themselves were not included in this study, we were 357 able to gain useful insights into the topic area to inform further work to explore 358 families' experiences. Using the themes which emerged from our focus groups a 359 semi-structured interview has been designed and patients will be interviewed to 360 determine their opinions on prenatal exome sequencing as part of the PAGE project 361

(http://www.pageuk.org). The opinions of obstetricians and gynaecologists, who are 362 not specialists in fetal medicine, were not explored in this research and the 363 perspectives of this particular group of clinical professionals may well have revealed 364 some additional insights. There may also be details of significance that participants 365 might have been willing to share more privately rather than in a focus group setting 366 <sup>27</sup>. As private feedback was not sought from participants following the focus groups 367 368 we are unable to comment on this, and as such the authors accept this as limiting aspect of this research. 369

It is premature to make concrete recommendations from these qualitative data but our findings suggest that consent in the prenatal arena is not a 'perfect' process.
Consent-takers should be fully educated regarding the test. This work did not seek to fully explore the characterisation of the information that should be conveyed to make consent valid. We feel that further qualitative work needs to explore this and in particular capture the views of women and their families.

376

#### 377 **References**

1 Hillman SC, Pretlove S, Coomarasamy A, *et al.* Additional information from array
comparative genomic hybridization technology over conventional karyotyping in
prenatal diagnosis: a systematic review and meta-analysis. Ultrasound Obstet
Gynecol 2011; 37:6-14.

382	2 Bernhardt BA, Soucier D, Hanson K, et al. Women's experiences receiving
383	abnormal prenatal chromosomal microarray testing results. Genet Med 2013; 15:
384	139-145.

3 Mardis ER. Next-generation DNA sequencing methods. Annu Rev Genomics Hum
Genet 2008; 9:387-402.

4 Brandt DS, Shinkunas L, Hillis SL, *et al.* A closer look at the recommended criteria
for disclosing genetic results: perspectives of medical genetic specialists, genomic
researchers and institutional review board chairs. J Genet Couns 2013; 22:544-553.

5 Reiff M, Mueller R, Mulchandani S, *et al.* A qualitative study of healthcare

<sup>391</sup> providers' perspectives on the implications of genome-wide testing in paediatric

clinical practice. J Genet Couns 2014; 23:474-488.

6 Bollinger JM, Scott J, Dvoskin R, *et al.* Public preferences regarding the return of
individual genetic research results: findings from a qualitative focus group study.
Genet Med 2012; 14:451-457.

7 Sapp JC. Dong D, Stark C, et al. Parental attitudes, values, and beliefs toward the
return of results from exome sequencing in children. Clin Genet 2014; 85(2):120126.

8 Bradbury AR, Patrick-Miller L, Egleston B. Parent opinions regarding the genetic
testing of minors. Genet Med 2010; 28:3498-3505.

9 Levenseller BL, Soucier DJ, Miller VA, *et al.* Stakeholders' opinions on the
implementation of paediatric whole exome sequencing: implications for informed
consent. J Genet Couns 2014; 23:552-565.

404 10 Fisher A, Bonner C, Biankin AV, *et al.* Factors influencing intention to undergo
405 whole genome screening in future healthcare: a single-blind parallel-group
406 randomised trial. Prev Med 2012; 55:514-520.

11 Scully JL, Porz R, Rehmann-Sutter C. 'You don't make genetic test decisions
from one day to the next' – using time to preserve moral space. Bioethics 2007; 21:
208-217.

12 Hillman SC, Skelton J, Quinlan-Jones E, *et al.* 'If it helps...' The use of microarray
technology in prenatal testing: patient and partners reflections. Am J Med Genet A
2013; 161A:1619-1627.

413 13 Patton MQ. Qualitative Evaluation and Research Methods (2<sup>nd</sup> Edition) Newbury
414 Park, California, Sage. 1990.

14 Ritchie J, Spencer L. Applied policy research (Ch 9) In: Analyzing qualitative data.
London: Routledge. 1994:177-194.

417 15 Srivastava A, Thomson SB. Framework Analysis: A Qualitative Methodology for
418 Applied Policy Research. Journal of Administration and Governance 2009; 4:72-79.

419 16 Vetro A, Bouman K, Hastings R, *et al.* The introduction of arrays in prenatal
420 diagnosis: A special challenge. Hum Mutat 2012a; 33:923-9.

17 Aune I, Moller A, 'I want a choice, but I don't want to decide' a qualitative study of
pregnant women's experiences regarding early ultrasound risk assessment for
chromosomal anomalies. Midwifery 2012; 28:14-23.

18 Yu JH, Jamal SM, Tabor HK, *et al.* Self-guided management of exome and
whole-genome sequencing results: changing the results return model. Genet Med
2013; 15(9):684-90.

19 Waler SA, Kellom KS, Palmer SC, *et al.* Comparing genetic counselor's and
patient's perceptions of needs in prenatal chromosomal testing. Prenat Diagn
2015;35:870-878.

20 Green RC, Berg JS, Grody WW, *et al.* ACMG recommendations for reporting of
incidental findings in clinical exome and genome sequencing. American College of
Medical Genetics and Genomics. Genet Med 2013; 15:565-574.

21 Srebniak M, Boter M, Oudesluijs G, *et al.* Application of SNP array for rapid
prenatal diagnosis: implementation, genetic counselling and diagnostic flow. Eur J
Hum Genet 2011; 19:1230-7.

436 22 Middleton A, Morley KI, Bragin E, *et al.* Attitudes of nearly 7000 health
437 professionals, genomic researchers and publics toward the return of incidental
438 results from sequencing research. Eur J Hum Genet 2015; 24(1):21-29.

- 439 23 Kalynchuk EJ, Althouse A, Parker LS, *et al.* Prenatal whole-exome sequencing:
  440 parental attitudes. Prenat Diagn 2015; 35:1030-6.
- 24 Sandelowski M, Sample size in qualitative research. Research in Nursing and
  Health 1995; 18:179-183.
- 25 Braun V, Clark V. Successful qualitative research: A practical guide for beginners.
  London, Sage, 2013.
- 26 Baker SE, Edwards R. How many qualitative interviews is enough? Expert voices
- and early career reflections on sampling and cases in qualitative research. NCRM,
- 447 National Centre for Research Methods Review Paper
- eprints.ncrm.ac.uk/2273/4/how\_many\_interviews.pdf [accessed on 12 April 2016].
- 449 27 Jayasekara RS, Focus groups in nursing research: Methodological perspectives.
- 450 Nursing Outlook 2012; 60(6):411-416

451

452