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Young people's understanding, attitudes and involvement in decision-making about genome sequencing for rare diseases: A qualitative study with participants in the UK 100,000 genomes project

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Author statement

**Celine Lewis:** Conceptualization, funding acquisition , methodology, formal analysis, investigation, writing - original draft, project administration.

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1 **Young people's understanding, attitudes and involvement in decision-making about**  
2 **genome sequencing for rare diseases: A qualitative study with participants in the UK**  
3 **100,000 Genomes Project**

4

5 Running title: Young people's views about genome sequencing

6

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22

23 Conflict of interest: C.P. has been on a secondment with Genomics England as Clinical Lead

24 for Genetic Counselling since October 2016. The other authors declare no conflicts of

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26

27 Data Availability Statement: Excerpts of interview transcripts are available on request to the

28 corresponding author.

29 **Abstract**

30 Genome sequencing (GS) will have a profound impact on the diagnosis of rare and inherited  
31 diseases in children and young people. We conducted 27 semi-structured interviews with  
32 young people aged 11-19 having GS through the UK 100,000 Genomes Project. Participants  
33 demonstrated an understanding of the role and function of genes and DNA, however the  
34 terms 'genome' and 'genome sequencing' were less well understood. Participants were  
35 primarily motivated to take part to get a diagnosis or identify the gene causing their  
36 condition. The majority of participants understood they might not receive a diagnostic result.  
37 Most were unconcerned about data security or access, however anxieties existed around  
38 what the results might show and the potential for disappointment if the result was negative.  
39 Signing an assent form empowered young people, formalised the process and instilled a  
40 sense of responsibility for their choice to participate. Most young people ( $\geq 16$  years) had  
41 consented to receive secondary findings and had come to that decision without parental  
42 influence. Our research suggests that at least some young people are capable of making  
43 informed decisions about taking part in GS, and that involving them in discussions about  
44 testing can empower them to take responsibility over healthcare decisions that affect them.

45

46

47 Keywords: whole genome sequencing, secondary findings, young people, motivations,  
48 concerns, decision-making, rare disease

49

## 50 Introduction

51

52 The majority of rare diseases affect children and in many cases there is an underlying  
53 genetic cause for their condition (Wright et al., 2018). Many children with rare diseases,  
54 particularly those with developmental disorders, are undiagnosed (Firth and Wright, 2011).  
55 However, the advent of next generation sequencing technologies has revolutionised the way  
56 genetic testing can be conducted, enabling multiple genes or entire exomes or genomes to  
57 be sequenced simultaneously (Sun et al., 2015; Wright et al., 2018). Genome sequencing  
58 (GS) has been shown to increase diagnostic yield almost twofold compared to conventional  
59 panel testing (Lionel et al., 2017) and fourfold compared to chromosome microarray  
60 (Stavropoulos et al., 2016). The possible clinical benefits of a genetic diagnosis include  
61 ending the 'diagnostic odyssey' (Basel and McCarrier, 2017), access to information on  
62 management and therapy, a clearer prognosis, reproductive planning and opportunities to  
63 make contact with other families through disorder-specific support groups (Griffin et al.,  
64 2017; Thevenon et al., 2016). GS is therefore set to have a profound impact on children and  
65 young people with rare diseases and its implementation is being evaluated in a number of  
66 paediatric settings (Bowdin et al., 2016; Green et al., 2016; Turnbull et al., 2018).

67

68 Although a significant body of work has emerged in recent years exploring adult patients'  
69 experiences and attitudes towards GS, (Boeldt et al., 2017; Mackley et al., 2018; Roberts et  
70 al., 2018; Sanderson et al., 2015) very little empirical research in this area has included  
71 young people (Pervola et al., 2019; Raghuram Pillai et al., 2019) (sometimes referred to as  
72 'adolescents' and defined as aged 10-19 years by UNICEF (UNICEF, 2019)). To date, the  
73 limited work that has been done has primarily used hypothetical scenarios (Hufnagel et al.,  
74 2016; McGowan et al., 2018), or assessed adults' perspectives on sequencing in the  
75 paediatric setting (Fernandez et al., 2014; Levenseller et al., 2014). Young people with  
76 health-related issues are likely to face significantly different physical, psychological and  
77 social challenges from those of both young children and adults (Frederick, 2016). They may

78 have specific information and support needs including peer support, provision of age-  
79 appropriate information and healthcare providers who proactively raise salient issues  
80 (D'Agostino and Edelstein, 2013) Therefore, it is important to give them a voice regarding  
81 their understanding of the benefits and potential risks of GS as well as their preferences for  
82 involvement in decision-making.

83

84 The current legal position in the UK is that children under 16 years cannot make decisions  
85 about their healthcare without parental consent, unless they prove to have sufficient maturity  
86 and intellectual capacity (referred to as "Gillick competence") (Griffith, 2016). In other  
87 European countries, the age at which children can consent varies between 14-16, 18 or is  
88 dependent on maturity (European Union Agency for Fundamental Rights, 2017). In the  
89 United States of America, children's consent authority differs across states (Coleman and  
90 Rosoff, 2013). In some States no particular age is required, in some it is aged 14 and over  
91 and in others it is aged 18 and over. Studies have, however, shown that young people,  
92 particularly adolescents, do frequently have the capacity to be actively involved in  
93 discussions about their healthcare, including genetic testing (McGill et al., 2018; Pervola et  
94 al., 2019) and participating in research (Kuther and Posada, 2004) The American College of  
95 Medical Genetics and Genomics recently issued a statement in which they highlighted the  
96 importance of engaging young people in meaningful conversations about the goals and  
97 implications of genomic testing and potential findings, and consideration of its personal  
98 benefits and limitations (Bush et al., 2018). Engaging young people in medical decision-  
99 making has also been shown to be associated with lower decisional conflict (David et al.,  
100 2018).

101

102 In the 100,000 Genomes Project, a United Kingdom (UK) national programme charged with  
103 preparing the National Health Service (NHS) for the introduction of genomics into clinical  
104 practice, much attention was focused on involving young people in the decision-making  
105 process, including the development of age appropriate information materials and written

106 'assent' forms for participants under 16 years (Genomics England, 2015). Of the rare  
107 disease proband participants in the 100 000 Genomes Project, around a quarter of them  
108 were 15 years of age or under at the time of taking part (data accessed from the Genomics  
109 England Research Environment, 11th November, 2018). In that project, consent to take part  
110 included consenting to receive a clinical diagnosis where one is found, and allowing de-  
111 identified, individual clinical and genomic data to be used for research purposes (Turnbull et  
112 al., 2018). In addition, participants aged  $\geq 16$  years were able to opt in to receive clinically  
113 actionable 'secondary findings' such as hereditary breast and ovarian cancer (BRCA1/2) and  
114 hereditary colorectal cancer (Lynch syndrome) (Genomics England, 2015). Parents of  
115 children  $< 16$  years could also consent to receive secondary findings, which have symptoms  
116 which onset in childhood, to be looked-for in their child. These conditions include  
117 retinoblastoma, Von Hippel-Lindau syndrome, child onset multiple endocrine neoplasia types  
118 1 and 2, and childhood onset familial hypercholesterolaemia (Genomics England, 2015).

119

120 We sought to characterise the understanding, motivations, concerns and experiences of  
121 decision-making among young people having GS in relation to both the main findings and  
122 the secondary findings.

123

## 124 **Methods**

125

126 This was a qualitative study using a semi-structured interview format to enable in-depth  
127 exploration of young people's views.

128

### 129 ***Ethical approval***

130 NHS Research Ethics Committee approval for this study was obtained from West Midlands  
131 (15/WM/0258).

132

133

### 134 ***Sampling and Recruitment***

135 The study was conducted in the UK with young people affected by rare diseases taking part  
136 in the 100,000 Genomes Project. Participants were not eligible for the Project if they had a  
137 molecular diagnosis. For many recruitment categories, it was expected that patients had  
138 already undergone clinically appropriate genetic testing, but that no molecular diagnosis had  
139 been found (Genomics England, 2015).

140

141 Participants were recruited through a children's hospital in London specialising in rare  
142 diseases. Potential participants were identified by a member of the healthcare team  
143 recruiting participants into the rare disease arm of the 100,000 Genomes Project. The  
144 inclusion criteria comprised: young people aged between 11-19 years (including probands  
145 as well as siblings undergoing GS), not affected by intellectual disability, and able to read  
146 and communicate in English. Siblings were invited to take part in the study as they were  
147 participants in the 100,000 Genomes Project and assented/consented to take part. They  
148 also had the potential to learn about secondary findings. A cut-off of 11 years was chosen as  
149 this was the age from which young people were invited to sign an 'assent' form in the  
150 100,000 Genomes Project.

151

152 . At the end of the 100,000 Genomes Project consent discussion, potential participants were  
153 told about this interview study, and asked if they (and their parent(s) for participants aged  
154 11-15) were interested in taking part. If so, they were asked to complete a consent to contact  
155 form. CL (first author, behavioural scientist and research lead) then sent the potential  
156 participant or parent(s) a participant information sheet explaining the study and followed up  
157 via email or telephone a few days later to determine whether the young person was willing to  
158 participate and if so arrange an interview (telephone or face-to-face). Consent was required  
159 from both the parent and participant when the young person was aged under 16 years, but  
160 only the participant if over 16 years. None of the participants had received a GS result at the  
161 time of interview.

162

**163 Interviews**

164 Interviews were conducted by CL. The semi-structured interview guide was developed by an  
165 advisory team comprising genetic counsellors, a fetal medicine expert and genetic research  
166 scientists and explored the following topics: 1. knowledge and understanding of the term  
167 'genes and DNA', 'genomes', 'genome sequencing' as well as the study procedure (that it is  
168 voluntary, timeframes, data access etc), 2. motivations for assenting/consenting to GS, 3.  
169 concerns around GS, 4. Motivations and concerns regarding secondary findings, and 5.  
170 involvement in the decision-making process. Interviews were audio-recorded, transcribed,  
171 anonymised and participants were given pseudonyms.

172

**173 Data analysis**

174 An abductive approach for coding and analysis was employed starting with codes derived  
175 from the topic guide and allowing new codes to emerge from the data (Robert et al., 2015).  
176 Data analysis was conducted following the principles of thematic analysis (Braun and Clarke,  
177 2006). A draft codebook was devised by CL informed from the topic guide. Three transcripts  
178 were then independently read and coded by CL and SS and additional codes added. Coding  
179 was compared and a second codebook devised. Remaining transcripts were then coded by  
180 CL using this second codebook with a subset coded by SS to ensure inter-rater agreement.  
181 Once all transcripts had been coded, CL and JH reviewed and refined the themes and sub-  
182 themes (constant comparison). A Framework matrix was also created as a way of ordering  
183 the data to facilitate recognition of patterns such as contradictory findings (Gale et al., 2013).  
184 In particular, we were interested to see how frequently codes concerning participants'  
185 motivations and concerns occurred and explore whether they were influenced by factors  
186 such as age, gender or whether they had a 'working diagnosis'.

187

**188 Results**

189

## 190 ***Participant characteristics***

191 Between June 2016 and March 2018, 40 young people (and their parents) were approached  
192 about this study, and 27 agreed and participated (68% recruitment rate): 19 were female, 25  
193 were probands and two were unaffected siblings. Participants ages ranged from 11-18 years  
194 (mean = 14 years). The most common condition types for affected probands were skeletal  
195 (including osteogenesis imperfecta) (n=8) followed by renal (n=4) and dermatological (n=3).  
196 Fourteen probands had no diagnosis, 11 had a working diagnosis (e.g. epilepsy) but no  
197 known genetic aetiology (Table 1). Interviews lasted between 15 minutes and 49 minutes  
198 (median = 34); 25 were conducted by telephone, two were conducted face-to-face.

199

200

## 201 ***Qualitative findings***

202

### 203 *Theme 1: Knowledge*

#### 204 *1.1 The terms 'gene' and 'DNA' are well understood*

205 Participants frequently described the function of genes and DNA using analogies including  
206 “an instruction manual or an encyclopaedia of you” (Rowena, 13 years) and “like a  
207 fingerprint” (Alice, 13 years). Genes and/or DNA were described as “what makes you, you”  
208 (Laura, 13 years,) and “control how your body performs” (Craig, 16 years). Around half of  
209 participants understood that genes and DNA are “passed down”, and nearly all expressed an  
210 understanding that genes can cause health problems:

211

212 “I know that I’ve got a fault somewhere in there, I got told it was like spelling. If the  
213 specific gene, it’s like a letter, if that’s not in the right place the spelling is wrong so  
214 that means my genes for that specific thing would be wrong.” (Harry, 13 years.)

215

216 Some participants displayed more advanced knowledge. For example, two spoke about  
217 inheriting “two sets of genes, one from each parent” (Emma, 13 years), two participants,  
218 aged 16 years and 18 years, referenced the letters A, G, T and C, and two participants (13  
219 and 17 years) mentioned the terms recessive and dominant inheritance, although only the  
220 older participant (Martin, 17 years) was able to articulate how these genes functioned in  
221 practice: “there are loads of genes that are recessive, which don’t show but they’re still  
222 there”. This participant also expressed an understanding of gene-environment interaction;  
223 “Certain things with your genes you can’t help, but it’s still a lot about your lifestyle decisions  
224 as well”. In most cases, participants commented that their knowledge of genetics had been  
225 acquired at school, but in some cases had been reinforced through the 100,000 Genomes  
226 Project. A few of the younger participants (11-13 years) had not heard of terms such as  
227 DNA and gene prior to the consent appointment.

228

229 *1.2: The terms ‘genome’ and ‘genome sequencing’ are less well understood*

230 Only a quarter correctly referred to the term ‘genome’ as being “all the genes” (Kathryn 16  
231 years) or “all the DNA letters” (James, 18 years), and these participants were generally older  
232 (15-18 years). Regarding the term “genome sequencing”, half spoke of looking at the “order”  
233 (Ash, 14 years) or “pattern” (Craig, 16 years) of the genes, ten participants explicitly stated  
234 they did not know what the term genome sequencing meant (median age 13.5 years), and  
235 five did not remember hearing the term during the consent appointment.

236

237 When asked why their parents were also asked to provide their DNA for the study, four  
238 participants (13-16 years) understood that it was for comparative purposes. One participant,  
239 aged 13 years, articulated how her unique DNA sequence would be compared to her  
240 parents’ DNA and also potentially other people’s with the same condition;

241

242 “Everyone’s got their individual sequence so everyone is different, so you can look at  
243 your own [genome] and compare it to other people’s. So they might compare mine to

244 my mum or other people with JDM [juvenile dermatomyositis] to see what the links  
245 are” (Elli, 13 years).

246

247 Notably, when asked whether they would definitely get a result from having their genome  
248 sequenced, most correctly understood that “some people get a diagnosis but not  
249 everybody.” (Emma, 13 years).

250

251 *Theme 2: Motivations*

252 *2.1: Young people cited multiple practical benefits*

253 All participants in the study were motivated to take part in the 100,000 Genomes Project  
254 because there was, potentially, a perceived benefit to them. These motivations included  
255 wanting to get a diagnosis, to identify the gene causing their condition, or to find out if the  
256 condition was genetic.

257

258 When exploring the importance of a diagnosis, some spoke of wanting to know if they had  
259 inherited the condition, or whether they might pass the condition on to their own children, a  
260 concern notably raised by some of the younger participants in the study:

261

262 “Also, if I ever have children when I’m older, will they get it and will the doctors be  
263 able to help them?” (Rowena, 13 years).

264

265 A prognosis was raised as being important by around a third of participants, for example,  
266 Mazie (13 years) spoke about wanting to know “if I will develop anything else”. Some thought  
267 a diagnosis would “help doctors to know what medication might be better than others” (Elli,  
268 13 years). A couple spoke of wanting a diagnosis to “end all of the testing” and a few  
269 participants discussed that an important practical benefit of a diagnosis was being “able to  
270 explain to people what’s actually wrong” (Louisa, 13 years).

271

272 Participants were realistic about the limitations of GS, with around half articulating that a  
273 diagnosis was unlikely to have a significant impact. For example, Elliott commented that “it’d  
274 be nice, but I don’t think it’ll change my life” (Elliott 15 years). Only a few participants (aged  
275 13, 15 and 16) spoke of being motivated because they wanted to “cure” their condition.

276

277 *2.2: Potential emotional benefits were also important*

278 A third of participants cited motivations of a psychological nature. These included wanting an  
279 “answer...to put a few questions to rest” (Elliott, 15 years), to “stop me from keep on  
280 wondering how I got it” (Elli, 13 years), to gain “closure” (Emma, 13 years), and for  
281 reassurance “that it’s not something I’ve done to cause it” (Katrina, 16 years). Amy spoke of  
282 the importance of a diagnosis in validating to others that she did have a genetic condition:

283

284 “I’d like to put a label on it, because it’s hard to explain to other people and it’s almost  
285 like people think ‘Oh, she hasn’t got a diagnosis so she hasn’t got anything wrong’”  
286 (Amy, 16 years).

287

288 *2.2: Young people are also motivated to help other people and contribute to science*

289 Almost all participants cited altruistic motivations. This included the potential benefits that  
290 taking part could have for others with the same condition, such as treatment or a quicker  
291 diagnosis. Rowena reflected on the research that had gone before which had subsequently  
292 benefited her:

293

294 “The reason I have been given the medication so quickly, is because they’ve done  
295 this sort of thing on other people which has helped me to be served in this way.”  
296 (Rowena, 13 years).

297

298 When comparing the motivations for taking part in the 100,000 Genomes Project, age  
299 appeared to be an important factor. Younger participants (11 to 13 years) cited nearly twice  
300 as many benefits directly related to them compared to benefits to others. Older participants  
301 (14 to 18 years) also cited more benefits to themselves compared to others, but the  
302 difference was less pronounced than that apparent among younger participants. No  
303 differences were observed when comparing across whether participants had a 'working  
304 diagnosis' or no diagnosis.

305

306 *Theme 3: Concerns*

307 *3.1: Some participants were anxious about what the result might show and the potential for*  
308 *disappointment if the result was negative*

309 When prompted, most participants commented that they did not have any concerns about  
310 having GS. However, a few participants did raise concerns about the potential emotional  
311 impact of the result, such as the potential for the result to reveal their condition was more  
312 serious than expected:

313

314 "Maybe if it's life threatening, like if something comes back that might shock me or  
315 something I never knew before which would scare me" (Claire, 17 years).

316

317 Similarly, Kathryn (16 years) spoke of having concerns that the results might show "I've got  
318 another problem that I need to manage". A few spoke of potentially being disappointed if  
319 they didn't get a result, for example, Laura (13 years) said that "if they can't find it, like it's  
320 going to be a bit sad because you want to know".

321

322 *3.2: Most participants did not have concerns about data security or access*

323 Most participants felt reassured by the data being deidentified so "they can't trace it back to  
324 me" (Kathryn, 16 years), and made comments signalling their trust the NHS: "I'm quite  
325 confident that they're going to keep it safe" (Emma, 13 years). Some older participants were

326 unclear how their data could be used against them, even if it was accessed without their  
327 permission.

328

329 Regarding data access, a number of participants articulated that the involvement of for-profit  
330 companies in research was “a good thing [because] medicines [are] produced from that”  
331 (Elliott, 15 years). Two participants were, however, ambivalent about ‘for-profit’ companies  
332 having access to their data, although both made comments in which they acknowledged the  
333 role of such companies in “help[ing] research, they can fund developing a cure” (Craig, 16  
334 years).

335

336 Only one person raised concerns about health insurance companies accessing his genomic  
337 data. In this case, the participants had been reassured by his father who had “assured him  
338 that for now, at least until 2019 I think they said health insurance companies wouldn’t be  
339 able to access any of that information” (James, 18 years).

340

#### 341 *Theme 4. Decision-making*

342 *4.1 Most young people felt the decision to take part in the 100,000 Genomes Project had*  
343 *been patient-led or a joint decision with parents.*

344 All participants were aware that taking part in the study was voluntary. Half of participants,  
345 and in particular the older participants, felt that the decision to have GS had been *their*  
346 decision: “My dad was there at the appointment but I think it was my decision because I  
347 wanted to try and find out what it was that was causing my problems” (James, 18 years).

348 These participants frequently spoke about making their own decisions about many aspects  
349 of their healthcare. For example, Kathryn (16 years), spoke about how her mum had “taken  
350 a step back from dealing with hospital appointments” and that she now “manage[d] my own  
351 medication”. For her, the decision to take part in the 100,000 Genomes Project was a

352 continuation of being responsible for her own health: “the genomes thing, is kind of just a  
353 continuation of that, just managing like my own condition and stuff”.

354

355 In around a third of cases, the decision to take part in the project was a joint decision  
356 between the participant and their parent(s). Despite parents being the ones who ultimately  
357 signed the consent form for their child to take part, participants reflected on the importance  
358 of being involved in those conversations. For example, Emma commented:

359

360 “I think ultimately it’s my parents’ decision but I should get a lot of say in it...a thing  
361 like that is going to impact me more than it’s going to impact them, so I think it is very  
362 important for me to be involved in conversations like that.” (Emma, 13 years)

363

364 In five instances (which included participants aged between 12 and 15 years), the decision  
365 to take part was made primarily by the parents. However, in these cases, the participant had  
366 agreed with that decision. Rowena, 13 years, spoke of not wanting to make the decision on  
367 her own, and was reassured that her parents were involved, suggesting that younger  
368 participants still relied on their parents to make important health-related decisions on their  
369 behalf:

370

371 “I wouldn’t want to make the decision on my own without knowing that it was the right  
372 thing to do. My parents said I think this is a good idea for you to do this and knowing  
373 my parents they would generally always make good decisions and they know what  
374 they’re doing and I trust them.” (Rowena, 13 years).

375

376 None of the participants described not wanting to take part and their parents having exerted  
377 pressure on them to participate.

378

379 *Theme 4.2: Involving young people in decision-making is empowering.*

380 Involving young people the discussion about genome sequencing, including asking them to  
381 sign an assent form empowered young people, formalised the process and instilled a sense  
382 of responsibility for their choice to participate. This is highlighted through comments such as:  
383 “it made me feel important, not just a blood source” (Elliott, 15 years), “I feel like I have a  
384 responsibility in some way” (Charlotte, 11 years), and “I think it shows that it’s not just about  
385 how old you are, it matters if you think you want to do this” (Rowena, 13 years). Notably,  
386 Fiona (11 years) commented that she hadn’t been asked to sign anything but would have  
387 valued the opportunity to do so as it might have made her more inclined to understand the  
388 study: “If I signed it, the questions that were being asked on the form, I might have  
389 understood more what was going to happen”. Moreover, where young people weren’t being  
390 involved in the consent discussion, it resulted in them “just zon[ing] out” and not “really  
391 pay[ing] much attention...even though it was primarily about me”. (Ash, 14 years)

392

393 *Theme 5.1: Young people are motivated to receive secondary findings so they can take*  
394 *action and be prepared*

395 Participants were primarily motivated to receive secondary findings for reasons related to  
396 taking action and “to be prepared”, but also acknowledged that “just because there’s a  
397 possibility, doesn’t mean it will happen.” (Rowena, 13 years). Other motivating factors  
398 included wanting to regain a sense of control over one’s health when so much was outside  
399 of their control “The suddenness and unexpectedness of my tumours have caused a few  
400 mental health issues...it would be helpful to know if something like [cancer/heart disease]  
401 could happen.” (Katrina, 16 years). Some participants who had consented to receive adult  
402 onset secondary findings, envisaged that they would adapt their behaviour e.g. “stop  
403 smoking...stop eating sugary foods” (Martin, 17 years) if they were found to be at increased  
404 risk. Two participants were in part motivated because there was a family history of cancer.  
405 One participant linked her motivation to being part of the ‘information age’; “the age in which  
406 I live, everybody wants to know as much as they can about themselves” (Claire, 17 years).

407

408 Of the eleven participants eligible to consent to adult onset secondary findings, only one  
409 participant declined to receive these. His decision was in part related to the advice he had  
410 been given by the health professional consenting him into the 100,000 Genomes Project,  
411 that he would be “too young to do anything about it”, and that he could receive secondary  
412 findings results at a later date.

413

414

415 When participants were questioned as to how they might feel if they were to find out they  
416 were at increased risk for certain conditions, six participants reflected that they would find  
417 the information “worrying”, with Kathryn (16 years) raising concerns that “it would be another  
418 thing wrong” that she would have to deal with alongside her current genetic condition.  
419 Nevertheless, all six who did articulate concerns, commented that they still wanted to know.

420

421 *Theme 5.2: Young people are making independent decisions about adult onset secondary*  
422 *findings without parental influence*

423 Most participants who had consented to receive adult-onset secondary findings described  
424 making decisions without parental influence, and justified this approach with comments in  
425 which they were keen to exert their autonomy around decisions related to their health e.g. “I  
426 feel like I was responsible enough to make that decision myself” (Amy, 16 years) and “in the  
427 end it’s about my body” (Seeta, 17 years). Two participants had, however, included their  
428 parents in the decision-making process. In one instance, there were divergent views  
429 amongst family members, with a father raising concerns “that if something comes up and it’s  
430 really bad” he didn’t want his daughter to “have to deal with it yet.” (Claire, 17 years).  
431 Nevertheless, despite her father’s reluctance, Claire had exerted her agency over the  
432 situation; “it’s up to me and I wanted the information to come back”.

433

434 *Theme 5.3: Young people under 16 years of age want to be involved in decisions around*  
435 *childhood onset secondary findings*

436 When it came to decision-making about childhood onset secondary findings, around a third  
437 of participants under 16 years did not recall the discussion. Of those that did remember,  
438 some had been actively involved in the discussion and clearly valued the opportunity to be  
439 involved in those decisions as highlighted by Rowena (13 years) who said “I wanted to know  
440 and so I said to my parents ‘yes I do want to do this’”. Those that hadn’t been involved  
441 commented that they would have liked to have been involved in such discussions as  
442 highlighted by Emma (13 years) who commented that “It does affect me the most...I should  
443 get a lot of say in it”.

444

445

446

#### 447 **Discussion**

448

449 To our knowledge, this is the first study to explore the attitudes of paediatric rare disease  
450 patients being offered GS. In the new UK NHS Genomic Medicine Service, around half of  
451 the rare and inherited disorders for which GS will be routinely available are conditions that  
452 affect young people (NHS England, 2019). Collecting empirical evidence about young  
453 people’s understanding, attitudes and preferences regarding decision-making can inform  
454 recommendations and best-practices.

455

456 Young people in our study demonstrated an understanding of the role and function of genes  
457 and DNA including a basic understanding of inheritance. However the terms ‘genome’ and  
458 ‘genome sequencing’ were less well understood, particularly amongst younger participants.  
459 These findings echo those from our quantitative survey study examining knowledge of  
460 genetics and genomics amongst 554 school pupils (Lewis et al., 2020). This finding is likely  
461 to reflect the National Curriculum in England where concepts such as genetics and DNA are

462 introduced from age 11 and the concept of genomics from around age 15 (Department for  
463 Education, 2015). The majority of participants in the present study understood that a  
464 limitation of GS is that they might not receive a diagnostic result. This is important given that  
465 currently around only 40% of paediatric patients get a result from GS (Lionel et al., 2017).  
466 Our data suggest that in the new NHS Genomic Medicine Service, it is important that health  
467 professionals check young people's understanding, particularly around what GS is and the  
468 current limitations of the technology to ensure they do not have unrealistic expectations  
469 about what results they might receive. Educational resources such as animations may be an  
470 effective way of supporting and enhancing young people's understanding during the in-  
471 person appointment.

472

473 A notable finding from our study is that young people were able to project how they might  
474 respond to a diagnostic result or a negative result and articulate their potential emotional  
475 reaction (fear, anxiety, disappointment etc). Such concerns may be realistic: Werner-Lin et  
476 al. found that parents and adolescents who had received non-actionable paediatric exome  
477 sequencing results initially experienced emotions including frustration, disappointment and  
478 fear (Werner-Lin et al., 2018). Giving young people the opportunity to discuss the potential  
479 emotional impact of GS findings in more depth including the option to discuss these  
480 separately from other family members, might be good practice going forward.

481

482 One area where our findings differ to research conducted with adults (McCormack et al.,  
483 2016; Robinson et al., 2016) is that young people did not have concerns about data security  
484 or insurance. A similar finding was reported by Pillai et al. who found that parents were more  
485 likely than adolescents to indicate that concerns around privacy and confidentiality  
486 influenced their decision to learn secondary findings results about their children (Raghuram  
487 Pillai et al., 2019). Young people had confidence that the NHS would protect their data and  
488 did not know how their data could be used against them. This is perhaps not surprising given  
489 most young people in this age group have not yet had to think about insurance, but may also

490 reflect a lack of awareness regarding the potential for genomic data to be used to  
491 discriminate against them in the future (e.g. employment). In other contexts, research has  
492 also shown that in the context of online personal information, young people feel they have  
493 “nothing to hide” and therefore do not consider privacy relevant for them (Adorjan and  
494 Ricciardelli, 2019). Further research could further explore whether this mindset applies to  
495 young people in the context of genomic data.

496

497 The majority of young people in this study felt that *they* had made the decision to take part in  
498 the 100,000 Genomes Project and receive main findings related to their condition, or that it  
499 had been a joint decision with their parents. This reflects the ethos of the project which  
500 emphasised the importance of inclusive decision-making (Genomics England, 2015). Our  
501 findings also shed light on the choices, justifications and parental involvement in young  
502 people’s decisions about secondary findings. Notable findings include that 1) participants  
503 (under 16 years) were keen to be involved in discussions around whether to find out about  
504 childhood onset conditions, and 2) most older participants (16 years and over) wanted to  
505 receive adult onset secondary findings, had made that decision independently of their  
506 parents, and expressed justifications regarding these independent choices that related to  
507 notions of autonomy and independence. Similar themes emerged in a previous study with  
508 adolescents aged 13 to 17 years old without a clinical indication for genomic testing in the  
509 USA (Pervola et al., 2019).

510

511 Four capacities have been described that are required for (medical) decision-making; these  
512 are (1) communicating a choice, (2) understanding, (3) reasoning, and (4) appreciation  
513 (Grootens-Wiegers et al., 2017). In this study we found that participants understood that  
514 participating was voluntary and were communicative and expressed a choice (capacity 1);  
515 they understood why they were undergoing GS (capacity 2); they were able to apply logical  
516 reasoning and weigh up the potential benefits and risks of taking part e.g. getting a  
517 diagnosis vs. not getting a diagnosis (capacity 3); and they were able to appreciate the

518 relevance of taking part for them as well as others (capacity 4). Thus, our findings suggest  
519 that many of the participants in our study *are* likely to have had the capacity to make an  
520 informed decision and felt empowered by being actively included in the decision-making and  
521 assent processes. This is an important finding as it has implications for clinical practice in  
522 that it underscores the importance of health professionals actively involving young people in  
523 the discussion and decision-making around GS. The finding that young people valued the  
524 opportunity to be involved in the decision-making process and in particular provide written  
525 assent is also notable and we recommend that this practice should continue.

526

#### 527 Strengths and limitations

528 This study adds much-needed empirical data on a topic that has received relatively little  
529 attention to-date, namely the views and experiences of young people having GS. A strength  
530 of this study is the diverse range of condition-types that affected participants in the sample.  
531 As with all qualitative studies, participants were self-selecting; participants with negative  
532 experiences may have been less willing to take part. In addition, this study did not include  
533 participants with intellectual disability which makes up a sizable number of children who  
534 might be offered GS (Wright et al., 2015). Finally, no demographic data on the parents  
535 (socioeconomic background or education level) were collected and thus we are unable to  
536 comment on the background of the participants. Participants' background may have had an  
537 impact on their level of understanding and/or attitudes towards genome sequencing.

538

#### 539 Conclusion

540 Young people understood the potential benefits of GS for both themselves and others, as  
541 well as the limitations of the technology. Our research provides evidence to show that there  
542 will be some young people with rare diseases that 1. are *capable* of making informed-  
543 decisions to take part in testing, and 2. that involving them in testing decisions *empowers*  
544 them to take responsibility over healthcare decisions that affect them. Further research with  
545 young people after they receive GS results will add to understanding of their overall

546 experience of this technology. In addition, future research could focus on the experiences of  
547 young people with intellectual disability, in particular whether and how to facilitate  
548 empowerment and inclusivity.

549

550

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563 References

564

565 Adorjan, M., Ricciardelli, R., 2019. A New Privacy Paradox? Youth Agentic Practices  
566 of Privacy Management Despite "Nothing to Hide" Online. *Canadian review of  
567 sociology = Revue canadienne de sociologie* 56(1), 8-29.

568 Basel, D., McCarrier, J., 2017. Ending a Diagnostic Odyssey: Family Education,  
569 Counseling, and Response to Eventual Diagnosis. *Pediatric clinics of North America*  
570 64(1), 265-272.

571 Boeldt, D.L., Cheung, C., Ariniello, L., Darst, B.F., Topol, S., Schork, N.J., Philis-  
572 Tsimikas, A., Torkamani, A., Fortmann, A.L., Bloss, C.S., 2017. Patient perspectives  
573 on whole-genome sequencing for undiagnosed diseases. *Personalized medicine*  
574 14(1), 17-25.

575 Bowdin, S.C., Hayeems, R.Z., Monfared, N., Cohn, R.D., Meyn, M.S., 2016. The  
576 SickKids Genome Clinic: developing and evaluating a pediatric model for  
577 individualized genomic medicine. *Clinical genetics* 89(1), 10-19.

578 Braun, V., Clarke, V., 2006. Using thematic analysis in psychology. *Qualitative  
579 Research in Psychology* 3(2), 77-101.

- 580 Bush, L.W., Bartoshesky, L.E., David, K.L., Wilfond, B., Williams, J.L., Holm, I.A.,  
581 2018. Pediatric clinical exome/genome sequencing and the engagement process:  
582 encouraging active conversation with the older child and adolescent: points to  
583 consider—a statement of the American College of Medical Genetics and Genomics  
584 (ACMG). *Genetics in medicine : official journal of the American College of Medical*  
585 *Genetics* 20(7), 692-694.
- 586 Coleman, D.L., Rosoff, P.M., 2013. The legal authority of mature minors to consent  
587 to general medical treatment. *Pediatrics* 131(4), 786-793.
- 588 D'Agostino, N.M., Edelstein, K., 2013. Psychosocial challenges and resource needs  
589 of young adult cancer survivors: implications for program development. *Journal of*  
590 *psychosocial oncology* 31(6), 585-600.
- 591 David, J.G., Lo, M.S., Langer, D.A., 2018. Adolescent and parent perspectives on  
592 medical decision-making for chronic illness. *Families, systems & health : the journal*  
593 *of collaborative family healthcare* 36(2), 248-251.
- 594 Department for Education, 2015. National curriculum in England: science  
595 programmes of study. [https://www.gov.uk/government/publications/national-](https://www.gov.uk/government/publications/national-curriculum-in-england-science-programmes-of-study)  
596 [curriculum-in-england-science-programmes-of-study](https://www.gov.uk/government/publications/national-curriculum-in-england-science-programmes-of-study). (Accessed Accessed 30th  
597 January 2019).
- 598 European Union Agency for Fundamental Rights, 2017. Minimum age requirements  
599 related to rights of the child in the EU. [https://fra.europa.eu/en/publications-and-](https://fra.europa.eu/en/publications-and-resources/data-and-maps/minag?dataSource=MINAG_en_62756&media=png&width=740&topic=group05&question=MINAG_HE01&plot=MAP&subset=NONE&subsetValue=NONE&answer=MINAG_HE01&year=2017)  
600 [resources/data-and-](https://fra.europa.eu/en/publications-and-resources/data-and-maps/minag?dataSource=MINAG_en_62756&media=png&width=740&topic=group05&question=MINAG_HE01&plot=MAP&subset=NONE&subsetValue=NONE&answer=MINAG_HE01&year=2017)  
601 [maps/minag?dataSource=MINAG\\_en\\_62756&media=png&width=740&topic=group0](https://fra.europa.eu/en/publications-and-resources/data-and-maps/minag?dataSource=MINAG_en_62756&media=png&width=740&topic=group05&question=MINAG_HE01&plot=MAP&subset=NONE&subsetValue=NONE&answer=MINAG_HE01&year=2017)  
602 [5&question=MINAG\\_HE01&plot=MAP&subset=NONE&subsetValue=NONE&answer](https://fra.europa.eu/en/publications-and-resources/data-and-maps/minag?dataSource=MINAG_en_62756&media=png&width=740&topic=group05&question=MINAG_HE01&plot=MAP&subset=NONE&subsetValue=NONE&answer=MINAG_HE01&year=2017)  
603 [=MINAG\\_HE01&year=2017](https://fra.europa.eu/en/publications-and-resources/data-and-maps/minag?dataSource=MINAG_en_62756&media=png&width=740&topic=group05&question=MINAG_HE01&plot=MAP&subset=NONE&subsetValue=NONE&answer=MINAG_HE01&year=2017).
- 604 Fernandez, C.V., Bouffet, E., Malkin, D., Jabado, N., O'Connell, C., Avar, D.,  
605 Knoppers, B.M., Ferguson, M., Boycott, K.M., Sorensen, P.H., Orr, A.C., Robitaille,  
606 J.M., McMaster, C.R., 2014. Attitudes of parents toward the return of targeted and  
607 incidental genomic research findings in children. *Genetics in medicine : official*  
608 *journal of the American College of Medical Genetics* 16(8), 633-640.
- 609 Firth, H.V., Wright, C.F., 2011. The Deciphering Developmental Disorders (DDD)  
610 study. *Developmental medicine and child neurology* 53(8), 702-703.
- 611 Frederick, C., 2016. Psychosocial Challenges/Transition to Adulthood. *Pediatric*  
612 *clinics of North America* 63(4), 735-749.
- 613 Gale, N.K., Heath, G., Cameron, E., Rashid, S., Redwood, S., 2013. Using the  
614 framework method for the analysis of qualitative data in multi-disciplinary health  
615 research. *BMC medical research methodology* 13, 117.
- 616 Genomics England, 2015. The 100,000 Genomes Project Protocol.  
617 [https://www.genomicsengland.co.uk/wp-](https://www.genomicsengland.co.uk/wp-content/uploads/2015/03/GenomicEnglandProtocol_030315_v8.pdf)  
618 [content/uploads/2015/03/GenomicEnglandProtocol\\_030315\\_v8.pdf](https://www.genomicsengland.co.uk/wp-content/uploads/2015/03/GenomicEnglandProtocol_030315_v8.pdf). (Accessed  
619 Accessed 12th January 2017).
- 620 Green, R.C., Goddard, K.A., Jarvik, G.P., Amendola, L.M., Appelbaum, P.S., Berg,  
621 J.S., Bernhardt, B.A., Biesecker, L.G., Biswas, S., Blout, C.L., Bowling, K.M.,  
622 Brothers, K.B., Burke, W., Caga-Anan, C.F., Chinnaiyan, A.M., Chung, W.K.,  
623 Clayton, E.W., Cooper, G.M., East, K., Evans, J.P., Fullerton, S.M., Garraway, L.A.,  
624 Garrett, J.R., Gray, S.W., Henderson, G.E., Hindorff, L.A., Holm, I.A., Lewis, M.H.,  
625 Hutter, C.M., Janne, P.A., Joffe, S., Kaufman, D., Knoppers, B.M., Koenig, B.A.,  
626 Krantz, I.D., Manolio, T.A., McCullough, L., McEwen, J., McGuire, A., Muzny, D.,  
627 Myers, R.M., Nickerson, D.A., Ou, J., Parsons, D.W., Petersen, G.M., Plon, S.E.,  
628 Rehm, H.L., Roberts, J.S., Robinson, D., Salama, J.S., Scollon, S., Sharp, R.R.,  
629 Shirts, B., Spinner, N.B., Tabor, H.K., Tarczy-Hornoch, P., Veenstra, D.L., Wagle, N.,

- 630 Weck, K., Wilfond, B.S., Wilhelmsen, K., Wolf, S.M., Wynn, J., Yu, J.H., 2016.  
631 Clinical Sequencing Exploratory Research Consortium: Accelerating Evidence-  
632 Based Practice of Genomic Medicine. *American journal of human genetics* 98(6),  
633 1051-1016.
- 634 Griffin, B.H., Chitty, L.S., Bitner-Glindzicz, M., 2017. The 100 000 Genomes Project:  
635 What it means for paediatrics. *Archives of disease in childhood. Education and*  
636 *practice edition* 102(2), 105-107.
- 637 Griffith, R., 2016. What is Gillick competence? *Human vaccines &*  
638 *immunotherapeutics* 12(1), 244-247.
- 639 Grootens-Wiegers, P., Hein, I.M., van den Broek, J.M., de Vries, M.C., 2017. Medical  
640 decision-making in children and adolescents: developmental and neuroscientific  
641 aspects. *BMC pediatrics* 17(1), 120.
- 642 Hufnagel, S.B., Martin, L.J., Cassedy, A., Hopkin, R.J., Antommaria, A.H., 2016.  
643 Adolescents' preferences regarding disclosure of incidental findings in genomic  
644 sequencing that are not medically actionable in childhood. *American journal of*  
645 *medical genetics. Part A* 170(8), 2083-2088.
- 646 Kuther, T.L., Posada, M., 2004. Children and adolescents' capacity to provide  
647 informed consent for participation in research. *Advances in psychology research* 32,  
648 163-173.
- 649 Levenseller, B.L., Soucier, D.J., Miller, V.A., Harris, D., Conway, L., Bernhardt, B.A.,  
650 2014. Stakeholders' opinions on the implementation of pediatric whole exome  
651 sequencing: implications for informed consent. *Journal of genetic counseling* 23(4),  
652 552-565.
- 653 Lewis, C., Sanderson, S.C., Hammond, J., Hill, M., Searle, B., Hunter, A., Patch, C.,  
654 Chitty, L.S., 2020. Development and mixed-methods evaluation of an online  
655 animation for young people about genome sequencing. *European journal of human*  
656 *genetics : EJHG*.
- 657 Lionel, A.C., Costain, G., Monfared, N., Walker, S., Reuter, M.S., Hosseini, S.M.,  
658 Thiruvahindrapuram, B., Merico, D., Jobling, R., Nalpathamkalam, T., Pellecchia, G.,  
659 Sung, W.W.L., Wang, Z., Bikangaga, P., Boelman, C., Carter, M.T., Cordeiro, D.,  
660 Cytrynbaum, C., Dell, S.D., Dhir, P., Dowling, J.J., Heon, E., Hewson, S., Hiraki, L.,  
661 Inbar-Feigenberg, M., Klatt, R., Kronick, J., Laxer, R.M., Licht, C., MacDonald, H.,  
662 Mercimek-Andrews, S., Mendoza-Londono, R., Piscione, T., Schneider, R., Schulze,  
663 A., Silverman, E., Siriwardena, K., Snead, O.C., Sondheimer, N., Sutherland, J.,  
664 Vincent, A., Wasserman, J.D., Weksberg, R., Shuman, C., Carew, C., Szego, M.J.,  
665 Hayeems, R.Z., Basran, R., Stavropoulos, D.J., Ray, P.N., Bowdin, S., Meyn, M.S.,  
666 Cohn, R.D., Scherer, S.W., Marshall, C.R., 2017. Improved diagnostic yield  
667 compared with targeted gene sequencing panels suggests a role for whole-genome  
668 sequencing as a first-tier genetic test. *Genetics in medicine : official journal of the*  
669 *American College of Medical Genetics* 20(4), 435-443.
- 670 Mackley, M.P., Blair, E., Parker, M., Taylor, J.C., Watkins, H., Ormondroyd, E., 2018.  
671 Views of rare disease participants in a UK whole-genome sequencing study towards  
672 secondary findings: a qualitative study. *European journal of human genetics : EJHG*  
673 26(5), 652-659.
- 674 McCormack, P., Kole, A., Gainotti, S., Mascalzoni, D., Molster, C., Lochmuller, H.,  
675 Woods, S., 2016. 'You should at least ask'. The expectations, hopes and fears of  
676 rare disease patients on large-scale data and biomaterial sharing for genomics  
677 research. *European journal of human genetics : EJHG* 24(10), 1403-1408.
- 678 McGill, B.C., Wakefield, C.E., Vetsch, J., Barlow-Stewart, K., Kasparian, N.A.,  
679 Patenaude, A.F., Young, M.A., Cohn, R.J., Tucker, K.M., 2018. Children and young

680 people's understanding of inherited conditions and their attitudes towards genetic  
681 testing: A systematic review. *Clinical genetics* 1, 10-22.

682 McGowan, M.L., Prows, C.A., DeJonckheere, M., Brinkman, W.B., Vaughn, L.,  
683 Myers, M.F., 2018. Adolescent and Parental Attitudes About Return of Genomic  
684 Research Results: Focus Group Findings Regarding Decisional Preferences. *Journal*  
685 *of empirical research on human research ethics : JERHRE* 13(4), 371-382.

686 NHS England, 2019. National Genomic Test Directory.  
687 <https://www.england.nhs.uk/publication/national-genomic-test-directories/>. (Accessed  
688 5th August 2019).

689 Pervola, J., Myers, M.F., McGowan, M.L., Prows, C.A., 2019. Giving adolescents a  
690 voice: the types of genetic information adolescents choose to learn and why.  
691 *Genetics in medicine : official journal of the American College of Medical Genetics*  
692 21(4), 965-971.

693 Raghuram Pillai, P., Prows, C.A., Martin, L.J., Myers, M.F., 2019. Decisional conflict  
694 among adolescents and parents making decisions about genomic sequencing  
695 results. *Clinical genetics*.

696 Robert, G., Cornwell, J., Locock, L., Purushotham, A., Sturmey, G., Gager, M., 2015.  
697 Patients and staff as codesigners of healthcare services. *Bmj* 350, g7714.

698 Roberts, J.S., Robinson, J.O., Diamond, P.M., Bharadwaj, A., Christensen, K.D.,  
699 Lee, K.B., Green, R.C., McGuire, A.L., 2018. Patient understanding of, satisfaction  
700 with, and perceived utility of whole-genome sequencing: findings from the MedSeq  
701 Project. *Genetics in medicine : official journal of the American College of Medical*  
702 *Genetics* 20(9), 1069-1076.

703 Robinson, J.O., Carroll, T.M., Feuerman, L.Z., Perry, D.L., Hoffman-Andrews, L.,  
704 Walsh, R.C., Christensen, K.D., Green, R.C., McGuire, A.L., 2016. Participants and  
705 Study Decliners' Perspectives About the Risks of Participating in a Clinical Trial of  
706 Whole Genome Sequencing. *Journal of empirical research on human research*  
707 *ethics : JERHRE* 11(1), 21-30.

708 Sanderson, S.C., Linderman, M.D., Suckiel, S.A., Diaz, G.A., Zinberg, R.E.,  
709 Ferryman, K., Wasserstein, M., Kasarskis, A., Schadt, E.E., 2015. Motivations,  
710 concerns and preferences of personal genome sequencing research participants:  
711 Baseline findings from the HealthSeq project. *European journal of human genetics :*  
712 *EJHG* 24(1), 14-20.

713 Stavropoulos, D.J., Merico, D., Jobling, R., Bowdin, S., Monfared, N.,  
714 Thiruvahindrapuram, B., Nalpathamkalam, T., Pellecchia, G., Yuen, R.K.C., Szego,  
715 M.J., Hayeems, R.Z., Shaul, R.Z., Brudno, M., Girdea, M., Frey, B., Alipanahi, B.,  
716 Ahmed, S., Babul-Hirji, R., Porras, R.B., Carter, M.T., Chad, L., Chaudhry, A.,  
717 Chitayat, D., Doust, S.J., Cytrynbaum, C., Dupuis, L., Ejaz, R., Fishman, L., Guerin,  
718 A., Hashemi, B., Helal, M., Hewson, S., Inbar-Feigenberg, M., Kannu, P., Karp, N.,  
719 Kim, R., Kronick, J., Liston, E., MacDonald, H., Mercimek-Mahmutoglu, S.,  
720 Mendoza-Londono, R., Nasr, E., Nimmo, G., Parkinson, N., Quercia, N., Raiman, J.,  
721 Roifman, M., Schulze, A., Shugar, A., Shuman, C., Sinajon, P., Siriwardena, K.,  
722 Weksberg, R., Yoon, G., Carew, C., Erickson, R., Leach, R.A., Klein, R., Ray, P.N.,  
723 Meyn, M.S., Scherer, S.W., Cohn, R.D., Marshall, C.R., 2016. Whole Genome  
724 Sequencing Expands Diagnostic Utility and Improves Clinical Management in  
725 Pediatric Medicine. *NPJ genomic medicine* 1(15012).

726 Sun, Y., Ruivenkamp, C.A., Hoffer, M.J., Vrijenhoek, T., Kriek, M., van Asperen,  
727 C.J., den Dunnen, J.T., Santen, G.W., 2015. Next-generation diagnostics: gene  
728 panel, exome, or whole genome? *Human mutation* 36(6), 648-655.

729 Thevenon, J., Duffourd, Y., Masurel-Paulet, A., Lefebvre, M., Feillet, F., El  
730 Chehadeh-Djebbar, S., St-Onge, J., Steinmetz, A., Huet, F., Chouchane, M.,  
731 Darmency-Stamboul, V., Callier, P., Thauvin-Robinet, C., Faivre, L., Riviere, J.B.,  
732 2016. Diagnostic odyssey in severe neurodevelopmental disorders: toward clinical  
733 whole-exome sequencing as a first-line diagnostic test. *Clinical genetics* 89(6), 700-  
734 707.

735 Turnbull, C., Scott, R.H., Thomas, E., Jones, L., Murugaesu, N., Pretty, F.B., Halai,  
736 D., Baple, E., Craig, C., Hamblin, A., Henderson, S., Patch, C., O'Neill, A.,  
737 Devereaux, A., Smith, K., Martin, A.R., Sosinsky, A., McDonagh, E.M., Sultana, R.,  
738 Mueller, M., Smedley, D., Toms, A., Dinh, L., Fowler, T., Bale, M., Hubbard, T.,  
739 Rendon, A., Hill, S., Caulfield, M.J., 2018. The 100 000 Genomes Project: bringing  
740 whole genome sequencing to the NHS. *Bmj* 361, k1687.

741 UNICEF, 2019. <https://data.unicef.org/topic/adolescents/overview>. (Accessed 10  
742 February 2020).

743 Werner-Lin, A., Zaspel, L., Carlson, M., Mueller, R., Walser, S.A., Desai, R.,  
744 Bernhardt, B.A., 2018. Gratitude, protective buffering, and cognitive dissonance:  
745 How families respond to pediatric whole exome sequencing in the absence of  
746 actionable results. *American journal of medical genetics. Part A* 176(3), 578-588.

747 Wright, C.F., Fitzgerald, T.W., Jones, W.D., Clayton, S., McRae, J.F., van  
748 Kogelenberg, M., King, D.A., Ambridge, K., Barrett, D.M., Bayzatinova, T., Bevan,  
749 A.P., Bragin, E., Chatzimichali, E.A., Gribble, S., Jones, P., Krishnappa, N., Mason,  
750 L.E., Miller, R., Morley, K.I., Parthiban, V., Prigmore, E., Rajan, D., Sifrim, A.,  
751 Swaminathan, G.J., Tivey, A.R., Middleton, A., Parker, M., Carter, N.P., Barrett, J.C.,  
752 Hurles, M.E., FitzPatrick, D.R., Firth, H.V., 2015. Genetic diagnosis of developmental  
753 disorders in the DDD study: a scalable analysis of genome-wide research data.  
754 *Lancet (London, England)* 385(9975), 1305-1314.

755 Wright, C.F., FitzPatrick, D.R., Firth, H.V., 2018. Paediatric genomics: diagnosing  
756 rare disease in children. *Nature reviews. Genetics* 19(5), 253-268.

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Table 1: Participant characteristics

<b>Participant characteristics</b>	
<b>Age</b>	11-18 years, mean=14 years
11-13	12
14-16	10
17-19	5
<b>Gender</b>	
Female	19
Male	8
<b>Proband or sibling</b>	
Proband	25
Sibling	2
<b>Condition type (probands)</b>	
Skeletal	8
Renal	4
Dermatological	3
Autoimmune	2
Hearing	2
Ophthalmological	2
Congenital heart disorder	2
Neurological	1
Endocrine	1
<b>Diagnosis (probands)</b>	
No diagnosis	14
Working diagnosis but aetiology unknown	11