

Published in Chokoshvili, D., Vears, D., Borry, P. (2018). Expanded carrier screening for monogenic disorders: Where are we now? *Prenatal Diagnosis*, 38 (1), Art.No. 10.1002/pd.5109, 59-66.

1 **Title:** Expanded carrier screening for monogenic disorders: Where are we now?

2 **Running head:** Expanded carrier screening for monogenic disorders

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19 **Funding sources**

20 None

21 **Conflict of interest**

22 None declared

23

24 **Word count:** 3814

25 **Table count:** 4

26 **Figure count:** 0

27

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29 **What is already known about this topic?**

- 30 • Expanded carrier screening for reproductive purposes has been available to prospective
31 parents since 2010, primarily through direct-to-consumer genetic testing companies
- 32 • The market of expanded carrier screening is rapidly growing, with more than 200000
33 tests being performed annually in the US alone

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35 **What does this study add?**

- 36 • This study provides a comprehensive review of the expanded carrier screening tests
37 currently available, focusing on test characteristics such as the number and nature of
38 disorders included and variant interpretation strategies used
- 39 • Appreciable differences have been identified across providers, with the number of
40 disorders screened for ranging from 41 to 1700. In addition, providers differed
41 considerably in terms of the mutations screened and/or variant interpretation/reporting
42 strategies.

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53 **ABSTRACT**

54 **Background**

55 Expanded carrier screening (ECS), which can identify carriers of a large number of recessive
56 disorders in the general population, has grown in popularity and is now widely accessible to
57 prospective parents. This article presents a comprehensive overview of the characteristics of
58 currently available ECS tests.

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60 **Methods**

61 To identify relevant ECS providers, we employed a multi-step approach, which included online
62 searching, review of the recent literature, and consultations with researchers familiar with the
63 current landscape of ECS.

64
65 **Results**

66 As of January 2017, there were sixteen providers of ECS tests: 13 commercial companies, two
67 medical hospitals and one academic diagnostic laboratory. We observed drastic differences in
68 the characteristics of ECS tests, with the number of conditions ranging from 41 to 1700. Only
69 three conditions (Cystic fibrosis, Maple syrup urine disease 1b, and Niemann-Pick disease)
70 were screened for by all providers. Where the same disease gene was included by multiple
71 providers, substantial differences existed in the mutations screened and/or variant
72 interpretation/reporting strategies.

73
74 **Conclusion**

75 Given the importance of carrier screening results in reproductive decision-making, the observed
76 heterogeneity across ECS panels is concerning. Efforts should be made to ensure that clear and
77 concrete criteria are in place to guide the development of ECS panels.

78
79 **Key words:** Expanded carrier screening, genetic testing, reproductive genetics, consumer
80 genomics, direct-to-consumer testing

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85 INTRODUCTION

86 Carrier screening is a form of reproductive genetic testing which aims to identify healthy
87 individuals with a single mutated copy of a gene associated with a recessive (autosomal or X-
88 linked) disorder. Such individuals, known as carriers, are typically unaffected, but they may be
89 at risk of having a child with the disorder. In particular, there is a one-in-four chance in each
90 pregnancy that a couple will conceive an affected child where both members of the couple carry
91 a faulty copy of a gene associated with the same autosomal recessive disorder or where the
92 female member is the carrier of an X-linked recessive disorder.¹ Due to the recessive pattern of
93 inheritance, most carrier couples have no family history suggestive of the disorder and are
94 therefore unaware of their reproductive risk.^{2,3} As a consequence, it is common for couples to
95 only find out about their carrier status after giving birth to an affected child.⁴ Such couples
96 could benefit from carrier screening, ideally before conception, to identify their reproductive
97 risks and inform decisions.⁵ If identified preconceptionally, couples can choose to pursue
98 artificial reproduction through pre-implantation genetic diagnosis or using egg or sperm from a
99 non-carrier donor. They can also decide to forego pregnancy with the current partner and, for
100 example, adopt a child instead. Where carrier status is identified in an ongoing pregnancy, the
101 options are limited to deciding on the prenatal diagnosis and termination of pregnancy if the
102 fetus is found to be affected.⁶ Alternatively, some carrier couples may choose not to alter their
103 reproductive plans and use their test results for information purposes, such as to emotionally
104 and logistically prepare for the possibility of having an affected child. Where effective
105 therapeutic interventions for the identified condition exist, awareness of carrier status may also
106 allow for reducing morbidity and mortality in an affected child, by initiating medical treatment
107 early on in life.⁷

108 Traditionally, carrier screening was available for a small number of recessive disorders with a
109 relatively high prevalence in the general population or among select ethnic groups, such as the
110 Ashkenazi Jewish.^{8,9} However, the continuous improvements in molecular genetics and
111 increasing characterization of genetic disorders, has lead to the development of expanded
112 carrier screening (ECS) tests for a large number of conditions.¹⁰ Adding genes to ECS tests is
113 now technically easy and inexpensive, which has resulted in commercially available ECS
114 panels screening for hundreds of recessive disorders.¹¹

115 There are two major advantages of ECS, compared to the traditional ethnicity-based approach
116 to carrier screening. First, ECS identifies carriers in the general population, thus potentially
117 benefiting at-risk couples regardless of their ethnic background. Second, ECS is a more cost-
118 effective approach to identifying carriers, with the cost of a typical ECS not exceeding that of
119 carrier screening tests for specific monogenic disorders.^{12,13} Due to these advantages, ECS has
120 grown in popularity and the number of ECS tests performed annually has been estimated at
121 200 000 in the US.¹⁴ However, to date, the expansion of ECS panels has mainly been driven by
122 technology, which has been met with considerable criticism. Authors have argued that
123 screening for the highest number of recessive disorders may not necessarily promote
124 reproductive autonomy, as carrier status information may be less helpful for couples at risk of
125 conceiving a child with a mild, low-penetrant or phenotypically variable disease.^{5,15,16} Because
126 of this, professional medical organizations have advocated for clear medical criteria to be used
127 for the development of ECS panels, as opposed to including as many disorders as technically
128 possible.^{5,12,16} For example, in a 2015 joint statement, several US genetic and reproductive
129 health organizations recommended limiting ECS to serious health conditions and discouraged
130 screening for disorders predominantly manifesting themselves in adulthood (such as alpha 1
131 antitrypsin deficiency) or conditions characterized by low penetrance and variable expressivity

132 (e.g. MTHFR deficiency and hereditary hemochromatosis).¹² Similarly, professional
133 recommendations emphasize the importance of well-established genotype-phenotype
134 correlations for disease-associated variants and require the existence of sufficient clinical
135 evidence to report a variant as pathogenic.^{5,12}

136 While recent professional recommendations addressing ECS panel composition offer valuable
137 guidance in the development of ECS tests, it is unclear whether ECS providers follow these
138 recommendations in practice. Moreover, some recommendations lack specificity and can be
139 subject to interpretation. Providers of ECS may hold different views on whether a particular
140 disorder should be characterized as severe, or how to define whether a genotype-phenotype
141 correlation is clear. This complexity means that at present, decisions regarding the inclusion of
142 disorders on ECS, and whether to categorize a particular variant as pathogenic, primarily lie
143 with individual ECS providers.¹¹ Given the subjective nature of decisions surrounding the
144 development of ECS tests, the aim of this study was to comprehensively review and compare
145 ECS tests, including: the list and nature of recessive disorders screened for by different
146 providers, characteristics of disease-specific mutation panels and/or novel variant interpretation
147 and reporting practices.

148

149 **METHODS**

150 In order to identify potentially relevant providers of ECS, three complementary strategies were
151 employed during the period of August-December 2016. First, an online search was performed
152 using the following search string: “(carrier OR reproductive OR preconception OR
153 preconceptional) AND (screening OR test OR testing) AND (purchase OR buy OR order)”.
154 The search was performed independently by two researchers (DC and DV) using a de-
155 personalized Google search to execute the query. Both researchers reviewed the first 30 pages

156 (300 entries) of the results, beyond which results were deemed to be repetitive and of no
157 additional value. This process identified 27 providers of ECS services.

158 Second, we identified five additional ECS providers by reviewing recent academic literature on
159 carrier screening and consumer genetic testing. These providers were either based in non-
160 English speaking countries, or did not actively advertise their ECS services through their
161 websites and thus were not identified in the online search. Third, we consulted with colleagues
162 from VU University Medical Centre Amsterdam, who had carried out a systematic review of
163 cystic fibrosis carrier screening offers, which identified two additional providers.

164 Using these combined search strategies, we identified 34 ECS providers. However, as the goal
165 of this study was to review the characteristics of ECS tests, we excluded intermediary
166 companies and medical clinics offering ECS services developed by external laboratories. We
167 also excluded carrier screening offers primarily targeted at specific ethnic groups (e.g.
168 Ashkenazi Jewish screening programs), and research-oriented ECS initiatives (e.g.
169 preconception carrier screening offer from University of Groningen, the Netherlands), as these
170 tests are currently not accessible to the general public. Finally, we excluded one testing
171 company whose website did not contain information regarding the genes on the company's ECS
172 panel. This resulted in 16 ECS providers. The available information about their ECS tests, web-
173 based content, information brochures and/or hypothetical test reports was archived and
174 downloaded for subsequent analysis. Archiving took place on 19 January 2017.

175 Data from the 16 providers were input in a Microsoft Excel document and analyzed using
176 descriptive statistics. As some providers used different names for the same disorder and,
177 occasionally, denoted the same gene by different gene aliases, entries in the Excel document
178 were manually reviewed for accuracy. Where synonymous diseases or gene aliases were
179 identified, entries were aggregated to ensure that they were treated as the same.

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181

182 **RESULTS**

183 **Comparison of ECS Panels among Various Providers**

184 The general overview of the current ECS landscape is summarized in Table 1. The majority of
185 the ECS providers (13/16) were private genetic testing companies, with only two medical
186 hospitals and one diagnostic laboratory offering internally-developed ECS tests. Additionally,
187 most (13/16) providers of ECS tests were based in the US, with only three operating elsewhere.
188 The number of recessive disorders included on ECS panels was between 41 and 1792, while
189 the number of genes ranged from 40 to 1556. Of note, several providers offered an option to
190 screen for a subset of disorders. For example, consumers of *Natera*, together with the ordering
191 physician, could choose to purchase one of the five *Horizon*TM panels screening for up to 270+
192 disorders, with *Horizon*TM 274 being the most comprehensive test option. Screening for a subset
193 of disorders was also possible with other providers, including *EGL Genetics*, *Genaware*, *Good*
194 *Start Genetics*, *GenPath*, *Integrated Genetics*, *Mount Sinai Hospital*, and *Progenity*. In most
195 cases, providers offered smaller, standardized panels of so-called “(Ashkenazi) Jewish”
196 disorders. However, some used other criteria to differentiate among subpanels, such as GenPath
197 Diagnostics who screened for a subset of ~70 disorders described as ‘clinically manageable’
198 (i.e. where effective therapeutic interventions exist). Additionally, several providers had
199 standard panels of <20 disorders, consisting exclusively of disorders for which carrier screening
200 has been recommended by a major professional organization, such as the American College of
201 Medical Genetics and Genomics (ACMG) or the American College of Obstetricians and
202 Gynecologists (ACOG).

203 Table 2 shows the list of recessive genes and their associated disorders screened for by the
204 highest number of ECS providers (see supplementary material for a complete list of genes
205 screened for by at least one provider). Although the providers collectively screened for
206 approximately 1700 recessive genes (supplementary material), only three genes, *BCKDHB*,
207 *CFTR*, and *SMPD1* were included by all 16 providers (Table 2). Only 167 genes (approximately
208 10%) were included by at least half of the providers (8/16), while more than 1000 genes were
209 screened for by a single provider (supplementary material). Among the genes screened for by
210 the highest number of providers, most were associated with autosomal recessive disorders.
211 However, some X-linked disorders, such as fragile X syndrome and ornithine transcarbamylase
212 deficiency, were also commonly included. Clinical characteristics of the included disorders
213 varied substantially. For example, the same number of providers (15) screened for Tay-Sachs
214 disease, a lethal disorder typically resulting in death during infancy, as phenylketonuria, a
215 chronic condition treatable through dietary interventions. (Table 2). To a lesser extent,
216 providers also screened for disorders inherited in an autosomal dominant manner. For example,
217 familial hypercholesterolemia (*LDLR* gene) and factor V Leiden thrombophilia (*F5* gene), both
218 of which typically result in morbidity in homozygous individuals, were included on ECS panels,
219 by four and three providers, respectively (Supplementary material).

220 We also looked at whether providers screened for the three disorders (alpha 1 antitrypsin
221 deficiency, hereditary hemochromatosis, and MTHFR deficiency) for which carrier screening
222 is currently considered of unclear clinical value and therefore discouraged by several
223 professional organizations.¹² The majority of providers (12/16) included at least one of the three
224 disorders on their standard panels, while two providers routinely screened for all three (Table
225 3). The approach adopted by *Counsyl* was different from other providers, as the company had
226 included alpha 1 antitrypsin deficiency on their Family Prep™ ECS test, but performed

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227 screening for MTHFR deficiency and hereditary hemochromatosis only upon consumer
228 request.

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231 **Comparison of Screening Strategies and Mutation Panels among ECS Providers**

232 We observed that even where the same gene was screened for by multiple providers, there were
233 significant differences in the list of included pathogenic variants and/or approaches to
234 interpreting novel variants, if non-targeted sequencing was used. To illustrate these differences,
235 we compared mutation panels and/or variant interpretation strategies, where available, for the
236 three recessive disorders screened for by all 16 providers of ECS (Maple syrup urine disease
237 type 1B, cystic fibrosis and other CFTR-related disorders, and Niemann-Pick disease A/B).

238 Providers differed in their approaches to screening, with some employing targeted genotyping
239 and others using sequencing as the primary screening strategy (Table 4). Some providers (e.g.
240 *Counsyl* and *Natera*) used a combination of both strategies, or offered a choice between these
241 strategies to the ordering healthcare professional (e.g. *EGL Laboratory*). Where mutation
242 panels were used, the number of variants screened for varied considerably, with differences
243 being the most prominent in *CFTR*, where the number of variants included ranged from 28 to
244 more than 600. Among those providers who employed sequencing, we identified different
245 approaches to reporting of novel variants. For example, *Counsyl* had a policy to limiting
246 reporting to pathogenic and likely pathogenic variants, whereas *Baby Genes* also communicated
247 variants of uncertain significance ‘that result in non-synonymous protein changes but have no
248 known clinical association’.

249

250 **DISCUSSION**

251 In this review of currently available ECS offers, we have found considerable discrepancies
252 across various ECS tests. One of the most striking differences among ECS panels was in the
253 number of recessive genes screened, which ranged from 40 to above 1500. Only three recessive
254 disease genes were screened for by all sixteen providers and the majority of genes were included
255 by less than half of them. This suggests that the providers may have used quite different criteria
256 to develop their ECS panels. Reducing the differences to achieve greater harmonization among
257 ECS panels would therefore require that the providers use consistent disease inclusion criteria.
258 One approach that has been gaining acceptance in the context of ECS is inclusion of recessive
259 disorder genes based on the severity of their associated phenotypes. In 2014, Lazarin and
260 colleagues proposed a method for categorizing disorders on the spectrum of severity ranging
261 from ‘mild’ to ‘profound’.¹⁷ Using severity as a key criterion for including disorders in ECS
262 may be a sound strategy for several reasons. First, as demonstrated in recent studies, it is feasible
263 to achieve a severity-based taxonomy for genetic conditions, with medical professionals and
264 laypersons holding similar views regarding the severity of a particular disorder.^{18,19} Second,
265 couples at risk of having an affected child often decide not to alter their reproductive plans
266 when they perceive the disorder to be mild, indicating that screening for mild disorders may be
267 of limited utility in the context of reproductive decision-making.²⁰ Third, taking steps to prevent
268 the birth of a child with a mild disorder, for example, through prenatal diagnosis and termination
269 of an affected pregnancy, may be refused by providers of reproductive healthcare on legal or
270 moral grounds, making it problematic for couples who wish to utilize these options.²¹ Although,
271 in principle, carrier couples may still benefit from their carrier status information by
272 emotionally and logistically preparing for the possibility of having an affected child, these
273 benefits may be less prominent where the identified disorder is of a mild nature.²² However,

274 categorizing recessive disorders by severity can also be complicated by the fact that some
275 disorders are associated with multiple phenotypes that may vary in their clinical
276 characteristics.²³ Among the ECS panels analyzed in our study, we observed that smaller panels
277 tended to be designed in the way that maximizes the reliability of positive test results. For
278 example, *23andMe* and *VU University Medical Centre Amsterdam*, who screen for 40 and 50
279 genes, respectively, utilized targeted genotyping for known pathogenic variants in relatively
280 common recessive disorders. Additionally, neither provider screened for the three disorders
281 currently considered of unclear clinical value due to their variable expressivity and low
282 penetrance (alpha 1 antitrypsin deficiency, MTHFR deficiency, and hereditary
283 hemochromatosis). In contrast, larger ECS panels typically aimed at maximizing the reliability
284 of negative test results, employing more comprehensive targeted mutation panels and/or non-
285 targeted sequencing. The principal advantage of more comprehensive ECS tests is that couples
286 receiving a negative test result can be highly confident that they are definitely not at risk of
287 having a child with one of the screened disorders.²³ However, this approach may also lead to
288 an increased proportion of false positive results, where some couples identified as carriers
289 through the test will not be at risk of having an affected child. For example, in hereditary
290 hemochromatosis, genotype-phenotype correlations are very low, with the penetrance estimated
291 at <10%.¹² This means that the majority of couples where both members are identified as
292 ‘carriers’ of hereditary hemochromatosis do, in effect, receive false positive results.
293 Importantly, the issue of low penetrance is not unique to disorders with unclear clinical value
294 in carrier screening. For example, in cystic fibrosis (*CFTR* gene), where screening is widely
295 recommended, there is conflicting evidence regarding the pathogenicity of some variants, such
296 as L997F (c.2991G>C).²⁴ Studies have suggested that a substantial proportion of individuals
297 who are compound heterozygotes for L997F and a ‘classical’ pathogenic mutation may never
298 present with symptoms, while others may develop a phenotype ranging from a mild *CFTR*-

299 related disorder to classical cystic fibrosis.²⁵ Consequently, carrier screening for L997F has
300 been contested, as reproductive couples in which one partner screens positive for L997F and
301 the other partner is found to be a carrier of another CFTR-related mutation will be confronted
302 with uncertain results.^{25,26} While couples with uncertain carrier screening results may benefit
303 from post-test genetic counseling to assist them to make reproductive choices with the
304 information at hand, genetic counseling is not routinely offered by all providers of ECS tests.¹¹

305 Owing to the challenges raised by unclear clinical significance of some variants, professional
306 medical organizations recommend that carrier screening be limited to variants with clearly
307 established genotype-phenotype correlations.^{5,12} Accordingly, the use of non-targeted
308 sequencing in the context of carrier screening has been controversial, as this approach may
309 routinely identify novel variants, many of which could be of unclear clinical significance.^{22,27,28}

310 On the other hand, some novel variants, such as those occurring at canonical splice sites, could
311 be clearly pathogenic with a highly predictable health impact.²⁹ These may be particularly
312 common in individuals from geographically isolated areas or with ethnic backgrounds where
313 genotype data is sparse. Therefore, when applied under rigorous laboratory protocols and
314 manual curation of all novel potentially relevant pathogenic variants, non-targeted sequencing
315 can reliably identify some additional carrier couples.²³ Among the ECS providers analyzed in
316 our study, we found that half of them (8/16) employed non-targeted sequencing, either as the
317 primary or as a complementary strategy to identify carriers. In this regard, we found substantial
318 differences in variant reporting practices, with some limiting reporting to clearly/likely
319 pathogenic variants, while others routinely reporting variants of uncertain significance. These
320 differences have implications for the results received by consumers of ECS services. An
321 individual may receive positive, negative, or indeterminate carrier status result for the same

322 variant in a recessive disease gene, depending on the provider through which they pursue
323 testing.³⁰

324 Finally, the present review of ECS panels also revealed that some providers offered tests that
325 may identify individuals who are undergoing screening as either being affected, or being at risk
326 of developing a genetic disease. This primarily refers to screening for X-linked recessive
327 disorders which may identify affected women, and to autosomal recessive disorders where
328 affected (homozygous or compound heterozygous) individuals may not have come to medical
329 attention due to the late-onset or relatively mild nature of the disorder. Furthermore, some ECS
330 panels also included a handful of autosomal dominant disorders, such as familial
331 hypercholesterolemia and factor V Leiden thrombophilia, where routine identification of
332 potentially affected individuals is even more likely. These findings suggest that the traditional
333 distinction between carrier screening and diagnostic or predictive genetic testing has become
334 blurred. It is important to ensure that individuals undergoing ECS are adequately informed of a
335 potential health-related finding and have access to quality genetic counseling, as well as follow-
336 up diagnostic tests and clinical care, where necessary.¹⁵

337 Given the significance of carrier status information it is essential that test results communicated
338 to prospective parents are reliable, correctly estimating their reproductive and/or health risks.
339 Of particular importance is devising criteria for interpreting and reporting novel variants, since
340 an increasing number of providers are moving away from using targeted genotyping panels and
341 are adopting non-targeted sequencing. Additionally, efforts should be made to reduce
342 discrepancies among ECS panels by applying consistent criteria for including genetic disorders.
343 The professional medical community has an important role to play by providing continuous
344 guidance and updated recommendations regarding ECS test characteristics. However, this alone
345 may not be sufficient, since some providers of ECS may choose not to follow such

346 recommendations, as evidenced in the present study. Therefore, closer collaboration between
347 the professional medical community and the providers of ECS might be necessary in order to
348 develop best testing practices.

349

350

351 **LIMITATIONS**

352 Despite our multi-step approach to identifying all the relevant providers of ECS, it is possible
353 that some providers were not included in the present study. This is primarily due to the fact that
354 the market of ECS tests is still in its early stages and is undergoing expansion, with a growing
355 number of new providers entering the field. Some of these providers may be operating in
356 specific geographical areas and could have their websites in a language other than English,
357 which makes their identification particularly challenging. As providers of ECS tests often used
358 different disease names and, to a lesser extent, different gene aliases in the description of their
359 ECS panels, it is possible that the data used in our study contained some errors. However, we
360 made efforts to minimize the possibility of such errors by carefully reviewing the entries for the
361 genes/diseases and comparing ambiguous items with the entries in authoritative databases such
362 as ClinVar and Online Mendelian Inheritance in Man (OMIM).

363

364 **CONCLUSION**

365 In this study, we have compared ECS tests offered by sixteen providers. We found substantial
366 differences in terms of both panel size and the lists of recessive disorder genes included on ECS
367 panels. Furthermore, where multiple providers screened for the same gene, their approaches

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368 varied in the lists of specific disease-associated variants (in targeted genotyping) and variant
369 interpretation/reporting practices (in non-targeted sequencing).

370 Given the significance of ECS test results in reproductive decision-making and, increasingly,
371 in personal medical care, such drastic differences among providers are concerning. Efforts
372 should be made to achieve a greater harmonization of ECS panels by using consistent criteria
373 for the inclusion of genes and disorders. Additionally, guidance is needed towards developing
374 clear standards for variant interpretation and reporting practices. This can be best accomplished
375 through a close collaboration between the professional genetic community and the providers of
376 ECS tests.

377 **ACKNOWLEDGEMENTS**

378 We would like to thank our colleagues who assisted us in the identification of relevant ECS
379 providers

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381 **Ethics approval**

382 Not applicable

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Table 1. An overview of the available ECS offers*

Provider	Type of provider	Country	Number of conditions screened**	Number of genes screened**
23andMe	Genetic testing company	USA	41	40
Baby Genes	Genetic testing company	USA	71	92
Igenomix	Genetic testing company	Spain	633	549
Counsyl	Genetic testing company	USA	113	111
EGL Genetics	Diagnostic laboratory	USA	147	148
VU University Medical Centre Amsterdam	Medical hospital	The Netherlands	50	50
GeneAware	Genetic testing company	USA	158	159
GenPath Diagnostics	Genetic testing company	USA	166	166
Good Start Genetics	Genetic testing company	USA	252	281
Integrated genetics	Genetic testing company	USA	135	136
Macrogen	Genetic testing company	South Korea	1792	1556
Mount Sinai Hospital	Medical hospital	USA	256	279
Natera	Genetic testing company	USA	272	273
Pathway Genomics	Genetic testing company	USA	73	74
Progenity	Genetic testing company	USA	230	226
Recombine	Genetic testing company	USA	314	301

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* Accurate as of 19 January 2017. Excludes ethnicity-specific and research-oriented ECS offers.

466 ** The highest number of genes/conditions screened for by any given provider. In some cases, testing can be performed for a
467 subset of these genes and/or disorders.

468 NB: The number of genes may be a more accurate measure for comparing the sizes of ECS panels than the number of
469 conditions. This is because providers may use different approaches to defining a 'condition'. For example, providers A and B
470 may screen for the same 6 genes associated with Neuronal Ceroid-Lipofuscinosis. However, provider A may consider this as
471 a single entry on the list of conditions, whereas provider B may include it as 6 separate entries. On the other hand, the number
472 of genes included on an ECS test, is not subject to the provider's interpretation.
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Table 2. Recessive disorder genes screened for by the highest number of ECS providers

Screened by	Gene (Associated disorder(s))
16 providers	<i>BCKDHB</i> (Maple syrup urine disease type 1B); <i>CFTR</i> (Cystic fibrosis and other CFTR-related disorders); <i>SMPDI</i> (Niemann-Pick disease A/B)
15 providers	<i>ACADM</i> (Medium chain acyl-CoA dehydrogenase deficiency, MCAD); <i>ASPA</i> (Canavan disease); <i>CBS</i> (Homocystinuria); <i>FAH</i> (Tyrosinemia type I); <i>FANCC</i> (Fanconi anemia group C); <i>G6PC</i> (Glycogen storage disease Type Ia); <i>GAA</i> (Glycogen storage disease, type II); <i>HBB</i> (Hemoglobinopathies, sickle cell disease); <i>HEXA</i> (Tay-Sachs disease); <i>IKBKAP</i> (Familial dysautonomia); <i>MCOLNI</i> (Mucopolipidosis type IV); <i>NPCI</i> (Niemann-Pick disease type C1); <i>PAH</i> (Phenylketonuria)
14 providers	<i>ACADVL</i> (Very long-Chain acyl-CoA dehydrogenase deficiency); <i>ASSI</i> (Citrullinemia, type I); <i>BLM/RECQL3</i> (Bloom syndrome); <i>CLN5</i> (Neuronal ceroid lipofuscinosis type 5); <i>DLD</i> (Dihydrolipoamide dehydrogenase deficiency or Maple syrup urine disease type 3); <i>FCYT/PKHD1</i> (Polycystic kidney disease, autosomal recessive); <i>GALC</i> (Krabbe disease); <i>GALT</i> (Galactosemia); <i>GBA</i> (Gaucher disease); <i>GCDH</i> (Glutaric acidemia type 1); <i>HADHA</i> (Long-Chain 3-Hydroxyacyl-CoA dehydrogenase deficiency); <i>Trifunctional protein deficiency</i> ; <i>IDUA</i> (Mucopolysaccharidosis type I or Hurler syndrome); <i>PEX7</i> (Rhizomelic chondrodysplasia punctata, Type 1); <i>PMM2</i> (Congenital disorders of glycosylation type 1a); <i>PPT1</i> (Congenital disorders of glycosylation type 1b); <i>SACS</i> (Autosomal recessive spastic ataxia of Charlevoix-Saguenay or ARSACS)
13 providers	<i>AGA</i> (Aspartylglucosaminuria); <i>ALDH3A2</i> (Sjogren-Larsson syndrome); <i>ARSA</i> (Metachromatic leukodystrophy); <i>ASL</i> (Argininosuccinic aciduria); <i>BCSIL</i> (GRACILE syndrome; Mitochondrial complex III deficiency, nuclear type 1); <i>CLRN1</i> (Usher syndrome type 3A); <i>CPT2</i> (Carnitine palmitoyltransferase deficiency, type 2); <i>DHCR7</i> (Smith-Lemli-Opitz syndrome); <i>GRHPR</i> (Primary hyperoxaluria type 2); <i>HSD17B4</i> (D-bifunctional protein deficiency); <i>LAMB3</i> (Herlitz junctional epidermolysis bullosa, LAMB3-related); <i>NBN</i> (Nijmegen breakage syndrome); <i>NEB</i> (Nemaline myopathy); <i>PCDH15</i> (Usher syndrome type 1F); <i>PEXI</i> (Zellweger spectrum disorder); <i>SLC12A6</i> (Andermann syndrome); <i>SLC17A5</i> (Sialic acid storage disease or Salla disease); <i>SLC37A4</i> (Glycogen storage disease Type Ib); <i>SMNI</i> (Spinal muscular atrophy); <i>TMEM216</i> (Joubert syndrome type 2); <i>TPPI</i> (Neuronal ceroid-lipofuscinosis, TPP1-Related); <i>TTPA</i> (Ataxia with vitamin E deficiency)
12 providers	<i>AGL</i> (Glycogen storage disease, type III); <i>AGXT</i> (Primary hyperoxaluria, Type 1); <i>ALDOB</i> (Hereditary fructose intolerance); <i>ALPL</i> (Hypophosphatasia); <i>ATP7B</i> (Wilson disease); <i>BCKDHA</i> (Maple syrup urine disease type 1a); <i>BTD</i> (Biotinidase deficiency); <i>CLN8</i> (CLN8-related Neuronal ceroid lipofuscinosis); <i>CPT1A</i> (Carnitine palmitoyltransferase type 1A deficiency); <i>CTNS</i> (Cystinosis); <i>FKTN</i> (Walker-Warburg syndrome; Fukuyama congenital muscular dystrophy); <i>GJB2</i> (Non-syndromic hearing loss); <i>HEXB</i> (Sandhoff disease); <i>HMGCL</i> (HMG-CoA Lyase Deficiency); <i>IVD</i> (Isovaleric acidemia); <i>MAN2B1</i> (Alpha-mannosidosis); <i>MEFV</i> (Familial mediterranean fever); <i>MMAA</i> (Methylmalonic aciduria, cblA type); <i>MMACHC</i> (Methylmalonic aciduria and homocystinuria, type cblC); <i>MUT</i> (MUT-related Methylmalonic aciduria); <i>NPHS1</i> (Nephrotic syndrome, type I); <i>PCCA</i> (Propionic acidemia type 1); <i>PCCB</i> (Propionic acidemia type 2); <i>PDCN/NPHS2</i> (Steroid-resistant nephrotic syndrome); <i>SGCA</i> (Limb-girdle muscular dystrophy type 2D); <i>SGCB</i> (Muscular dystrophy, limb-girdle, type 2E); <i>SLC26A2</i> (Skeletal dysplasias; Achondrogenesis type 1B); <i>SLC26A4</i> (Pendred syndrome)
11 providers	<i>ABCC8</i> (Familial hyperinsulinism/permanent neonatal diabetes); <i>ACATI</i> (Beta-ketothiolase deficiency); <i>ATM</i> (Ataxia-telangiectasia); <i>BBS10</i> (BBS10-related Bardet-Biedl syndrome); <i>CTSK</i> (Pycnodysostosis); <i>GLB1</i> (Mucopolysaccharidosis, type IVB / GM1 Gangliosidosis); <i>FMRI</i> (Fragile X syndrome); <i>GNE</i> (Inclusion body myopathy 2, Nonaka type); <i>GNPTAB</i> (Mucopolipidosis type 2/type 3); <i>LAMA3</i> (LAMA3-related Herlitz junctional epidermolysis bullosa); <i>LAMC2</i> (LAMC2-related Herlitz junctional epidermolysis bullosa); <i>LRRPRC</i> (Leigh syndrome, French-Canadian type); <i>MLC1</i> (Megalencephalic leukoencephalopathy); <i>MPI</i> (Congenital disorders of glycosylation type 1b); <i>NPC2</i> (Niemann-Pick disease type C2); <i>OTC</i> (Ornithine transcarbamylase deficiency); <i>SLC22A5</i> (Primary carnitine deficiency)

10 providers

ADA (Adenosine deaminase deficiency; ADA-related severe combined immunodeficiency); *ADAMTS2* (Ehlers-Danlos syndrome: Type VIIC); *BBS1* (BBS1-related Bardet-Biedl Syndrome); *CAPN3* (Limb-girdle muscular dystrophy, type 2A); *CLN3* (CLN3-related neuronal ceroid lipofuscinosis or Batten disease); *ETHE1* (Ethylmalonic encephalopathy); *HGSNAT* (Mucopolysaccharidosis type IIIC, Sanfilippo type); *HLCS* (Holocarboxylase synthetase deficiency); *IL2RG* (X-Linked severe combined immunodeficiency); *MMAB* (Methylmalonic aciduria, cblB type); *MPL* (Congenital amegakaryocytic thrombocytopenia); *MTTP* (Abetalipoproteinemia); *OPA3* (3-methylglutaconic aciduria, type III); *POMGNT1* (POMGNT1-related muscle-eye-brain disease); *PYGM* (Glycogen storage disease, type V); *RMRP* (Cartilage-hair hypoplasia); *TH* (Segawa syndrome)

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Table 3. Providers' practices regarding the inclusion of Alpha 1 antitrypsin deficiency, MTHFR deficiency, and hereditary hemochromatosis on their ECS panels

Provider	Alpha 1 Antitrypsin deficiency (<i>SERPINA1</i> gene)	MTHFR deficiency (<i>MTHFR</i> gene)	Hereditary Hemochromatosis (<i>HFE</i> , <i>HFE2</i> , <i>TFR2</i> genes)
23andMe	Not included	Not included	Not included
Baby Genes	Not included	Included	Not included
Igenomix	Included	Included	Included (HFE, TFR2)
Counsyl	Included	<i>Not included *</i>	<i>Not included * (HFE)</i>
EGL Genetics	Included	Not included	Included (HFE)
VU University Medical Centre Amsterdam	Not included	Not included	Not included
GeneAware	Included	Not included	Not included
GenPath Diagnostics	Not included	Not included	Not included
Good Start Genetics	Not included	Included	Included (HFE2, TFR2)
Integrated Genetics	Not included	Not included	Not included
Macrogen	Not included	Included	Included (HFE)
Mount Sinai Hospital	Not included	Included	Included (HFE2, TFR2)
Natera	Not included	Included	Included (HFE2, TFR2)
Pathway Genomics	Included	Not included	Included (HFE)

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Progenity	Included	Not included	Included (HFE, HFE2, TFR2)
Recombine	Included	Included	Included (HFE2, TFR2)

477 * Not part of the standard screening panel, but can be included if specifically requested by the consumer.
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Table 4. Screening strategies and the size of mutation panels for the three genes screened for by all 16 providers

Provider	Maple Syrup Urine Disease Type 1B (BCKDHB gene)	Cystic fibrosis and other CFTR-related disorders (CFTR gene)	Niemann-Pick Disease A/B (SMPD1 gene)
23andMe	TG (2 variants)	TG (28 variants)	TG (3 variants)
Baby Genes	Seq.	Seq.	Seq.
Igenomix	TG (24 variants)	TG (146 variants)	TG (42 variants)
Counsyl	TG (3 variants) + Seq.	TG (99 variants) + Seq.	TG (4 variants) + Seq.
EGL Genetics	TG or Seq.	TG or Seq.	TG or Seq.
VU University Medical Centre Amsterdam	TG	TG	TG
GeneAware	Seq.	CNV + Seq.	Seq.
GenPath Diagnostics	TG (3 variants)	TG (220 variants)	TG (6 variants)
Good Start Genetics	TG + Seq.	TG + Seq.	TG + Seq.
Integrated Genetics	TG	TG (609 variants)	TG
Macrogen	TG (1 variant)	TG (102 variants)	TG (14 variants)
Mount Sinai Hospital	Seq.	TG + Seq.	Seq.
Natera	TG (21 variants) + Seq.	TG (579 variants) + Seq.	TG (50 variants) + Seq.

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Pathway Genomics	TG (3 variants)	TG (82 variants)	TG (5 variants)
Progenity	TG (3 variants)	TG (656 variants)	TG (4 variants)
Recombine	TG (6 variants)	TG (150 variants)	TG (9 variants)

- 481 TG - Targeted genotyping
- 482 Seq. - (non-targeted) sequencing
- 483 CNV - Copy number variation analysis
- 484