REVIEW ARTICLE



Barriers and facilitators for cascade testing in genetic conditions: a systematic review

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Received: 2 April 2020 / Revised: 21 August 2020 / Accepted: 1 September 2020 / Published online: 18 September 2020 © The Author(s), under exclusive licence to European Society of Human Genetics 2020

Abstract

Cascade testing is the process of offering genetic counseling and testing to at-risk relatives of an individual who has been diagnosed with a genetic condition. It is critical for increasing the identification rates of individuals with these conditions and the uptake of appropriate preventive health services. The process of cascade testing is highly varied in clinical practice, and a comprehensive understanding of factors that hinder or enhance its implementation is necessary to improve this process. We conducted a systematic review to identify barriers and facilitators for cascade testing and searched PubMed, CINAHL via EBSCO, Web of Science, EMBASE, and the Cochrane Library for articles published from the databases' inception to November 2018. Thirty articles met inclusion criteria. Barriers and facilitators identified from these studies at the individual-level were organized into the following categories: (1) demographics, (2) knowledge, (3) attitudes, beliefs, and emotional responses of the individual, and (4) perceptions of relatives, relatives' responses, and attitudes toward relatives. At the interpersonal-level, barriers and facilitators were categorized as (1) family communication-, support- and dynamics-, and (2) provider-factors. Finally, barriers at the environmental-level relating to accessibility of genetic services were also identified. Our findings suggest that several individual, interpersonal and environmental factors may play a role in cascade testing. Future studies to further investigate these barriers and facilitators are needed to inform future interventions for improving the implementation of cascade testing for genetic conditions in clinical practice.

Introduction

Cascade testing is the process of offering genetic testing to at-risk relatives of an individual who has been diagnosed with a genetic condition (i.e., the index patient or proband).

Supplementary information The online version of this article (https://doi.org/10.1038/s41431-020-00725-5) contains supplementary material, which is available to authorized users.

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This process is critical for timely initiation of riskmanagement strategies such as surveillance and prophylactic strategies, and is widely used in autosomal dominant conditions such as familial cancers and familial hypercholesterolemia. Clinical guidelines for these diseases recommend that cascade testing be offered to relatives of probands to identify cases [1–3]. Despite this, studies suggest that several genetic conditions remain underdiagnosed in the population [4–6], indicating that the implementation of cascade testing in clinical practice needs to be optimized.

To date, few studies have systematically reviewed the uptake of cascade testing in relatives of probands and the effectiveness of such interventions across various diseases. In a scoping review focused on the delivery of cascade testing for hereditary conditions, Roberts et al. provide a broad overview of cascade testing interventions, policy considerations, barriers and facilitators to their use and research gaps [7]. Based on their findings, several research gaps remain in the literature on cascade testing, including limited use of rigorous methods to test the efficacy of cascade testing programs and interventions. Understanding factors that influence whether probands disclose genetic information and whether relatives pursue genetic testing will be critical to design effective interventions to improve cascade testing. To address this need, we conducted an indepth systematic review of the research literature to identify barriers and facilitators that may affect the uptake of cascade testing.

Methods information sources and search strategy

The Preferred Reporting Items for Systematic Reviews and Meta-analyses (PRISMA) reporting guidelines [8] were followed for this study (Supplemental Material, Appendix A). The following databases: PubMed, CINAHL via EBSCO, Web of Science, EMBASE, and the Cochrane Library were electronically searched for articles published from the database inception date to 18 November, 2018 using keywords and appropriate subject headings that captured the range of terms used synonymously with cascade testing (e.g., "cascade screening" and "familial genetic testing"). Complete search strategies are provided in Supplemental Material, Appendix B. Hand-searches were also performed by manually examining the references of relevant literature reviews to identify any additional studies that may have been missed due to incomplete or inaccurate indexing in the electronic search databases. All references were uploaded to Covidence Systematic Review software (https://www. covidence.org) [9], a systematic review management system for study selection.

Study selection

Two of four reviewers (MCR, NYW, SS, and WDD) independently reviewed each title and abstract for eligibility, and disagreements were resolved through discussion. The same procedure was repeated for full-text review. Articles that focused on the disclosure of genetic information to family members and the actual uptake of genetic testing by relatives were both included to comprehensively capture barriers and facilitators to cascade testing both from probands' and the family members' perspectives. Conference abstracts, meeting reports, literature reviews, guidelines, and simulation modeling studies were excluded. Articles relating to other types of genetic testing and disclosure (whole-population or universal genetic testing, parental disclosure of genetic testing to children, newborn/ neonatal/pre-natal testing, or proband testing), those that lacked a methods section or relevant outcomes (no barriers/ facilitators described, study focused on clinical outcomes for cascade testing only, or study did not explicitly study cascade testing) and those that only reported prevalence of genetic testing were also excluded.

Data extraction and quality assessment

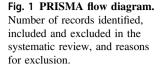
Data extraction forms were developed in Covidence using the PICOS framework for the following information: population (sample size and percentage of women), intervention (characteristics of the cascade testing intervention including disease area(s), whether counseling and resources related to cascade testing were provided, primary mediator of cascade testing), comparator (if applicable), outcomes (barriers and facilitators) and setting (country, scale, clinical, academic) [10]. Barriers and facilitators experienced either by proband or relatives in cascade testing were qualitatively described in some studies and, in others, were examined for their ability to predict relatives' uptake of genetic testing or proband's disclosure to relatives through methods like regression. Our coding used an inductive approach and reflected the language used by study authors. The forms were developed iteratively and piloted on a subset of five articles after which two reviewers independently extracted data from each study. Disagreements were resolved through discussion. Barriers and facilitators were organized according to the Social Ecological Model, a theoretical framework that allows for the examination of the interactions between personal and environmental factors on health behaviors [11].

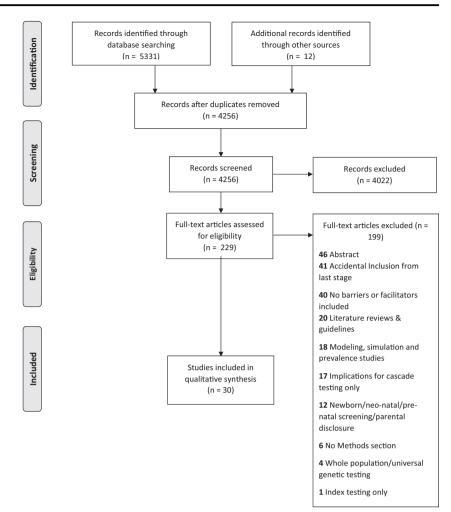
The methodological quality of each study was assessed using the Mixed Method Appraisal Tool (MMAT), version 2018 [12]. Both reviewers independently assessed whether the study met the corresponding MMAT criteria for each study type (RCT, descriptive, observation, qualitative, or mixed-methods). Meta-analysis was not conducted given the significant heterogeneity in study design, populations, setting, and outcomes.

Results

Of the 4256 unique studies that were identified through database searching and hand-searching, 229 articles were assessed for full-text eligibility. Thirty articles [13–42] were included in our analysis after excluding articles as per the inclusion and exclusion criteria (see Fig. 1 for PRISMA flow diagram).

Study characteristics for the included articles are available in Table 1. Twelve studies [13, 17, 22, 24–26, 28, 29, 35, 36, 38, 40] were conducted in the US while the remaining were conducted in Australia (n = 4) [14, 15, 19, 20], Europe (n = 9; 4 UK or parts of UK [23, 27, 31, 42], 4 Netherlands [21, 33, 34, 43], 1 France





[39]), Asia (n = 4; 2 Israel [18, 41], 1 Vietnam [30], 1Japan [16]), and South America (n = 1 from Brazil)[37]). Most studies were conducted within a single-center (n = 12 [16, 17, 20, 22-24, 30, 34, 35, 37, 40, 41]), followed by national- (n = 10) [13, 14, 18, 21, 26, 29, 31, 33, 38, 43], regional- (n = 4) [19, 39, 40, 42], multicenter (n = 3) [27, 28, 36] and state- (n = 1) [25] level studies. The majority of studies (n = 18) [13, 14, 16-19, 25, 26, 28, 30, 33, 35-41] used a descriptive study design using survey/interview/case series/observational few data. with a qualitative studies (n = 9)[15, 21, 23, 24, 27, 29, 31, 42, 43] and mixed-method studies (n = 2) [20, 34] and one randomized controlled trial [22]. Studies spanned disease areas including familial hypercholesterolemia (n = 10) [13, 15, 19, 25, 30, 33, 35, 37, 42, 43], Lynch syndrome (n = 6)[16, 17, 21, 24, 28, 38], hereditary breast and ovarian cancer (n = 6) [18, 22, 29, 36, 40, 41], cystic fibrosis (n = 3)[20, 31, 39], inherited cardiac conditions (n = 2) [23, 27], Fragile X syndrome (n = 2) [26, 34], and long QT syndrome (n = 1) [14]. Only one study explicitly reported that counseling was included as a component of cascade testing [42], while two studies noted that resources to assist with cascade testing (letter, written material and information sheet) were provided to patients [23, 42]. In most studies, the stakeholder who contacted relatives was not explicitly defined (n = 16) [13, 14, 16, 18–21, 26–29, 31, 35, 36, 38, 39], with patients (n = 5)[22, 23, 34, 41, 44], study team members (n = 4)[24, 30, 40, 45], providers (n = 2) [17, 37] and members of a screening program (n = 3) [15, 33, 43] contacting relatives in the remaining studies. Three studies were conducted solely among women [13, 22, 36], one study was conducted solely among men [29], and six studies did not report the proportion of females in the study sample [20, 27, 30, 31, 34, 43]. Sixteen studies did not contain information about race and ethnicity of the study sample [14-16, 19-21, 23, 27, 30, 33, 34, 37, 39, 41-43, 45].

Overall, both reviewers indicated that studies met three or more criteria, with few studies falling below this threshold. Barriers and facilitators at the individual, interpersonal and environmental levels are summarized in Tables 2 and 3, respectively. Notably, no studies in our analysis investigated any environmental facilitators. Only

Benson ¹³ 2 Burns ¹⁴ 2 Campbell ²⁵ 2 Cheune ²⁰ 2	Year of study	Countrav													
s	publication	Country	Setting	Year(s) of data collection	Scale	Design	Data Source	N (Study sample)	% White	% Female	Disease Area	Counseling related to cascade testing	Cascade testing primarily mediated by	Provision of resources related to cascade testing	MMAT Score
\$	2016	USA	Members of the WomenHeart and FH Foundation databases	2014	National	Descriptive 3	Survey data	761	86	100 1	Familial hypercholesterolemia	NR	NR	NR	4
0	2015	Australia	Patients enrolled in the Australian Genetic Heart Disease Registry	NR	National	Descriptive	Survey data	75	NR	73 1	Long QT syndrome	NR	NR	NR	4
	2017	USA	2014 Minnesota State Fair	2014	State	Descriptive	Survey data	971	NR	59	Familial hypercholesterolemia	NR	Study team	NR	3
	2010	USA	Tertiary referral cancer center and public 1996–2008 county hospital in California	1996–2008	Multi- center	Descriptive	Survey data	1135	60	100	Hereditary breast and ovarian cancer	NR	NR	NR	5
de Souza 2 Silva ²¹	2018	Brazil	Academic medical center	NR	Single- center	Descriptive	Survey data	183	NR	54	Familial hypercholesterolemia	NR	Provider (trained specialized health professional)	NR	3
Dilzell ²² 2	2014	USA	 Academic medical center, (2) patients affiliated with Lynch Sydnrome International, an advocacy group 	2012-2013	National	Descriptive	Survey data	50	16	1 62	Lynch syndrome	NR	NR	NR	3
Dugueperoux ²³ 2	2016	France	District in western Brittany where CF incidence is high	1980–2004	Regional	Descriptive (Observational data	128	NR	58	Cystic Fibrosis	NR	NR	NR	3
Finlay ²⁴ 2	2008	USA	Academic medical center	2004	Single- center	Descriptive	Survey data	132	93	73	Hereditary breast and ovarian cancer	NR	Study team	NR	4
Hagoel ²⁵ 2	2000	Israel	National cancer control center	NR	Single- center	Descriptive (Observational data	438	NR	70	Hereditary breast and ovarian cancer	NR	Patient	NR	4
Hallowell ²⁶ 2	2011	Scotland	Regional cascade testing service	2010	Regional	Qualitative]	Interview data	38	NR	55	Familial hypercholesterolemia	Yes	Patient	Yes (letters and written material)	3
Hardcastle ¹⁵ 2	2014	Australia	Lipid disorders clinic in Royal Hospital	NR	Regional	Qualitative]	Interview data	18	NR	44	Familial hypercholesterolemia	NR	Screening program	NR	4
Ishii ¹⁶ 2	2011	Japan	Cancer Institute	2005-2009	Single- center	Descriptive	Survey data	40	NR	63	Lynch syndrome	NR	NR	NR	5
Lerman ¹⁷ 1	1999	USA	Academic medical center	1996–1998	Single- center	Descriptive	Survey data	139	66	55	Lynch syndrome	NR	Provider	NR	4
Lieberman ¹⁸ 2	2018	Israel	Medical center	NR	National	Descriptive	Survey data	1771	NR	79	Hereditary breast and ovarian cancer	NR	NR	NR	4
Maxwell ¹⁹ 2	2009	Australia	Western Australia	2008	Regional	Descriptive 1	Interview and survey data	430	NR	59	Familial hypercholesterolemia	NR	NR	NR	4
8	2013	Australia	Children's Hospital	NR	Single- center	Mixed-	Interview data	249	NR	NR	Cystic fibrosis	NR	NR	NR	3
Mesters ²¹ 2	2005	Netherlands	Individuals enrolled in the Netherlands Foundation for the Detection of Hereditary Tumors	NR	National	Qualitative]	Interview data	30	NR	73 1	Lynch syndrome	NR	NR	NR	5
Montgomery ²² 2	2013	USA	Fox Chase Cancer Center	2000-2003	Single- center	RCT	RCT data	422	92	100	Hereditary breast and ovarian cancer	NR	Patient	NR	4
Ormondroyd ²³ 2	2014	UK	Academic medical center	2010-2011	Single- center	Qualitative]	Interview data	22	NR	09	Inherited cardiac conditions	NR	Patient	Yes (invitation letter and information sheet)	5
Pentz ²⁴ 2	2005	USA	Academic medical center	NR	Single- center	Qualitative]	Interview data	80	85	65	Lynch syndrome	NR	Study team	NR	5
	2006	USA	National survey research registry	2012	National	Descriptive	Survey data	679 families	92	06	Fragile X syndrome	NR	NR	NR	3
Smart ²⁷ 2	2010	UK	Genetics Knowledge Park program	2005-2006	Multi- center	Qualitative]	Interview data	27	NR	NR	Inherited cardiac conditions	NR	NR	NR	5
	2008	USA	Cancer genetics clinics in academic medical centers	NR	Multi- center	Descriptive	Survey data	174	16	70	Lynch syndrome	NR	NR	NR	4
Suttman ²⁹ 2	2018	USA	Academic medical center	NR	National	Qualitative]	Interview data	21	67	0	Hereditary breast and ovarian cancer	NR	NR	NR	5

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lable 1 (continued)	(panuc)														
Study Se	Setting					Methods		Population	Ē		Intervention				
τ <mark>.</mark> Υ Ι	Year of Country study publication		Setting	Year(s) of Scale data collection	1	Design	Data Source	N (Study sample)	% White	% Female	% White % Female Disease Area	Counseling related to cascade testing	Cascade testing primarily mediated by	Provision of resources related to cascade testing	MMAT Score
Truong ³⁰ 20	2018	Vietnam	Vietnam Academic medical center	NR	Single- center	Descriptive	Single- Descriptive Case series data 112 center		NR	NR	Familial hypercholesterolemia	NR	Study team	NR	3
Ulph ³¹ 20	2015 1	England	Research university	2008	National	Qualitative	National Qualitative Interview data	67	68	NR	Cystic Fibrosis	NR	NR	NR	5
van El ⁴³ 20	2018 1	Netherlands	Netherlands Research university	NR	National	Qualitative	National Qualitative Interview data	6	NR	NR	Familial hypercholesterolemia	NR	Screening program	NR	5
van Maarle ³³ 2001		Netherlands	Netherlands Academic medical center	1998	National	National Descriptive Survey data	Survey data	677	NR	54	Familial hypercholesterolemia	NR	Screening program	NR	5
Van Rijn ³⁴ 19	1 1997	Netherlands	Netherlands University hospital	1997	Single- center	Mixed- methods	Interview data	504	NR	NR	Fragile X Syndrome NR	NR	Patient	NR	3
Wurtmann ³⁵ 20	2018 1	USA	Academic medical center	2016	Single- center	Descriptive Survey data	Survey data	38	89	87	Familial hypercholesterolemia	NR	NR	NR	3

significant and descriptive results from our analysis are described below. In addition, nonsignificant factors are presented in Table 3.

Individual barriers

Demographics

One study reported that having low income was negatively associated with uptake of genetic testing among relatives [26], while other studies reported positive associations with other demographic factors (presented under Facilitators).

Knowledge

Studies described low or perceived lack of knowledge among probands and/or relatives as a barrier for both disclosure [25] and uptake of testing [23, 27, 31]. Probands in two studies cited not knowing who was at risk for the disease as a barrier to disclosure [14, 40].

Attitudes, beliefs and emotional responses of the individual

A number of barriers for probands and/or relatives to disclosure were reported, including: actionability of results [25], religion [25], discomfort with topic [25], sadness [25], surprise [25], lack of trust in the genetic information presented [25], laziness [25], no dissemination plan [20], time needed to communicate [25], emotional or general difficulty in sharing information [14, 25, 35, 40], and a preference for doctors to explain [25]. Paternalism [43], psychological burden [43], anticipation of regret [33], location of relatives [27], low perceived susceptibility [23], attitudes toward genetic testing [23], and lack of motivation [15] were described by probands and/or relatives as barriers to uptake of genetic testing. High depression symptoms had a negative effect on disclosure by probands [22], while the presence of depression symptoms among probands had a negative association with uptake. A belief that costs outweighed benefits was described as a barrier to uptake by probands [17]. General privacy concerns [25, 28, 43], deferment or rejection of responsibility [15, 25, 35], anxiety or guilt [14, 20, 25, 27, 28], logistical concerns [16, 35, 44, 45], fear of discrimination in the context of marriage or employment [18, 35], and limited recall of diagnosis and competing pressures [20, 25, 27] were all described as barriers to disclosure and uptake by probands and/or relatives. Finally, lack of readiness for discussion among probands was described as barrier to uptake [18].

Table 2 Barriers to disclosure of genetic information and uptake of cascade testing at the individual level.

Barriers	Disclosure of C	Genetic Informati	ion to Relatives	Uptake of Genetic	Testing by Relatives	
	Probands only	Relatives only	Probands/ Relatives	Probands only	Relatives only	Probands/ Relatives
Individual barriers						
Demographics						
Income					Neg (Low) ²⁶	
Knowledge						
Knowledge/Perceived knowledge			Desc ²⁵			Qual ^{23,27,31}
Not knowing who was at risk in the family						
Reluctance to cause fear, stress, and negative emotions	Desc ²⁸			Desc ¹⁸		
Perceived susceptibility						Desc(Low) ²
Attitudes, beliefs and emotional n	responses of the	individual				
Attitudes toward genetic testing						Desc ^{23,27}
Belief that costs outweigh benefits				Desc ¹⁸		
Lack of motivation						Qual ¹⁵
Depression Symptoms	Neg (High) ²²			Neg (Present) ¹⁷		
Distress	NS ²²					
General privacy concerns	Desc ²⁸		Desc ²⁵			Desc ⁴⁷
Deferment/rejection of responsibility		Desc ³⁵	Desc ²⁵			Qual ¹⁵
Actionability of results			Desc ²⁵			
Religion			Desc ²⁵			
Discomfort with topic			Desc ²⁵			
Embarrassment/Shame			Desc ^{25,20}			Desc ³¹
Sadness			Desc ²⁵			
Surprise			Desc ²⁵			
Lack of trust			Desc ²⁵			
Lazy			Desc ²⁵			
Anxiety/Guilt	Desc ^{14,28}		Desc ^{25,20}			Desc ²⁷
Emotional or general difficulty in sharing information	Desc ^{14,24}	Desc ³⁵	Desc ²⁵			
Logistical concerns		Desc ³⁵	Desc ²⁵	Qual ²⁶		Desc ¹⁶
Fear of discrimination in the context of marriage or employment		Desc ³⁵		Desc ¹⁸		
Readiness for discussion				Desc ¹⁸		
Limited recall of diagnosis and competing pressures			Desc ^{25,20}			Qual ²⁷
No dissemination plan			Desc ²⁰			
Paternalism						Desc ⁴⁷
Psychological burden						Desc ⁴⁷
Anticipation of regret						Desc ³³
Location of relatives			Desc ²⁵			Qual ²⁷
Time needed to communicate			Desc ²⁵			
Prefer doctors to explain			Desc ²⁵			

Table 2 (continued)

Barriers	Disclosure of C	Genetic Informati	on to Relatives	Uptake of Genetic Te	esting by Relatives	
	Probands only	Relatives only	Probands/ Relatives	Probands only	Relatives only	Probands/ Relatives
Perceptions of relatives, relative	es' reported res	ponses, and att	itudes toward	relatives		
Not wanting to upset relatives	Desc ¹⁴	Desc ³⁵				
No at-risk family members			Desc ²⁵			
Relative lack of concern or interest			Desc ^{25,20}	Desc ¹⁸		
Relative not willing to listen /does not care	Qual ²⁴	Desc ³⁵	Desc ²⁵			Qual ²⁷
Relative hostility toward advice or rejection		Desc ³⁵	Desc ²⁵	Desc ¹⁸		
Relative did not believe participant			Desc ²⁵			
Concern regarding family reaction			Desc ²⁵			
Family lack of understanding	Qual ²⁴ , Desc ²⁸	Desc ³⁵	Desc ^{25,20}			
Family may not agree to testing			Desc ²⁵			
Avoidance/Right not to know/ Ignorance						Qual ¹⁵ , Desc ^{20,33,47}
Relatives' stage of life			Desc ²⁰			Desc ¹⁶ , Qual ²⁷
Relative appears healthy						Desc ¹⁶
Interpersonal barriers						
Family communication, support	rt and dynamics					
Impact of disease on family				Desc ³⁴	Neg (Mostly or somewhat negative) ²⁶	
Disappointing experience with disclosure early in process			Desc ²⁰			Desc ³¹
Emotional distance or estrangement or conflict or resentment	Desc ²⁸		Qual ²¹	Qual ²⁶ , Desc ¹⁸		Qual ^{23,27}
General communication concerns			Desc ²⁵	Neg (No communication) ²⁶ , Desc ³⁴		Qual ^{24,27}
Not in contact/close with family	Desc ²⁴	Desc ³⁵	Desc ²⁵ , Qual ²¹			Desc ¹⁶ , Desc ²⁷
Language barrier			Desc ²⁵			
Provider factors						
Provider awareness				Qual ²⁹		Qual ⁴³
Provider engagement						Qual ³¹
Environmental barriers						
Accessibility of testing						
Finances/Cost		Desc ³⁵	Desc ²⁵			Qual ⁴³
Insurance		Desc ³⁵	Desc ²⁵			Qual ⁴³
Access to genetic testing			Desc ²⁰			Qual ³¹
Extra clinical referrals						Qual ⁴³

Neg – negative, Desc- barrier or facilitator described in descriptive studies, Qual- barrier or facilitator described in qualitative studies.

Table 3 Facilitators for disclosure of genetic information and uptake of cascade testing at the individual level.	c information and uptake of	cascade testing at the	individual level.		
Facilitators	Disclosure of genetic information to relatives Probands only Relatives only	ormation to relatives Relatives only	Probands/Relatives	Uptake of genetic testing by relatives Probands only Relatives	atives Probands/Relatives
Individual facilitators					
Demographics					
Age			Pos (High) ²⁰	NS ^{13,21}	NS ^{20,17}
Gender	NS^{28}	Pos (Female) ²²			NS ¹⁷ , Pos (Female) ²³ , Pos (Asian) ²⁰
Race/Ethnicity			Pos (Asian) ²⁰ , NS (Latina) ²⁰ , NS (African American) ²⁰	NS ¹³	NS (Latina) ²⁰ , NS (African American) ²⁰
Income				Pos (High) ¹³	NS ¹⁷
Education				NS^{21}	Pos (College Degree) ¹⁷
Employment				NS^{21}	
Marital Status				Pos (Married) ¹³	Pos (Married) ¹⁷
Socioeconomic status			NS^{20} , Pos (High) ²⁰		Pos (High) ²⁰
Clinical factors					
Personal history of disease	$NS^{22,28}$			NS ^{13,21}	$NS^{20,17}$, Pos^{25}
Family history of disease				NS^{21}	
Smoking				NS^{13}	
Prior history of risk factors				Desc ¹⁹	
Genetic test results	Pos (unambiguous) ²² , NS (inconclusive) ²²				
Knowledge					
Knowledge/Perceived knowledge			Pos (High) ²⁰		
Perceived susceptibility			Desc (High) ²¹		
Attitudes, beliefs and emotional responses of the individual					
Distress	NS^{22}				
Wanting to know				Desc^{20}	Desc ³³
Intrinsic motivation			Desc^{21}		
Perceived control	Pos (High) ²²				
Forced by circumstances					Desc ³³
Need for emotional support	Qual ²⁴ , Desc ²⁸	Desc^{35}	Desc^{20}	Desc ¹⁸	
Satisfaction with decision to undertake genetic testing			Pos (High) ²⁰		Pos (High) ²⁰
Perceptions of relatives, relatives' reported responses, and attitudes toward relatives	ed responses, and attitudes	toward relatives			
Relatives' right to know			Qual ²⁹	Desc ³⁴	Qual ^{23, 24}

Duty in //e p from Desc ²⁸ p from Desc ²⁸ d dynamics genetic Pos (Extremely or somewhat in favor) ²² , NS Opposed/Neutral) ²² , NS NS ²² Desc ²⁸ Desc ²⁸ Desc ²⁸ d dual ²⁴ , Desc ²⁸ at risk telatives	Table 3 (continued)					
ormation will help in letticions or ficticy, letticions or ficticy, letticions or ficticy, letticions or ficticy, letticions or ficticy, letticions or ficticy, letticions of the discardelle from Daca ³ Dec	Moral obligation toward relatives/Duty to inform		Desc^{35}	Desc ^{14,21}	Desc ^{18,26,34}	Qual ³¹
eteitves in the family Day at harm in the family Day and the disease/help from Des ²⁴ that in the family Day at harm in the family Day (Day Day at harm in the family Day (Day Day Day at harm in the family Day (Day Day Day Day at harm in the family Day Day at harm in the family Day Day at harm in the family Day Day Day Day Day Day Day Day Day Da	Feeling that information will help in making medical decisions or lifestyle		Desc ³⁵	Desc ¹⁴		Qual ^{27,31}
at ham in the family/Duy and the discuss/help from Bee ¹ Bee ³ that are some with a same same same same same same same sa	Concern about relatives			Desc^{29}	Desc^{20}	Qual ²³
and the discase/help from Dec ³⁸ g the same sorrow all action, support and dynamics cation, support and dynamics as the factor of the second and information for dynamics as the factor of the second and information for dynamics as the factor of the second and information for dynamics as cating are and the second and information for dynamics as the factor of the second and information for dynamics as the factor of the second and and the second as the consoln for dynamics as	Desire to prevent harm in the family/Duty to protect		Desc ³⁵		Desc ¹⁸	Qual ^{15,27}
g the same sorrow illators ication, support and dynamics leadive's opinion on genetic Pas (Extremely or anowhatin li favor) ²² , NS coposed/Neutral) ²² , NS answhatin li favor) ²² , NS answhatin li favor) ²² , NS posed/Neutral) ²² , NS posed/NS posed/Neutral) ²² , NS posed/NS p		Desc ²⁸				Desc ¹⁶
illators eation, support and dynamics eation, support and dynamics conversion or genetic Pos (Extremely or <u>Opposed/Neutral</u>) ²² , NS Opposed/Neutral) ²² , NS Opposed/Neutral) ²² , NS intervet oping status NS ²² intervet coping status NS ²² is with information Desc ²³ Desc ¹⁴ besc ¹⁴ Desc ¹⁴ Desc ²⁴ is not information at the transport of transport of the transport of transport of the transport of the transport of transport of the transport of transp	To avoid feeling the same sorrow					Desc ¹⁶
tetion, support and dynamics elative's opinion or genetic Pos (Externely or (Opposed/Neural) ²² , NS enses/Relationship to the iter's coping status NS ²² so with information besc ²⁸ NS ²² so with information besc ²⁸ Desc ¹⁴ besc ²⁴ Desc ²⁴ a communication family and information/sport from area besc ³⁵ Desc ³⁵ Oul ²⁶ family letter, Personal and Linformation/sport from area besc ³⁵ Desc ³⁵ Oul ²⁶ a family letter, Personal and Linformation/sport from area support group action dual family networks at risk Desc ³⁵ Desc ³⁵ Desc ³⁵ a gual ²⁶ a family network at risk Desc ³⁵ Desc ³⁵ Desc ³⁶ a gual ³⁶ a predic connector a genetic connector	Interpersonal facilitators					
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Desc ³⁵	Assistance in identifying relatives at risk Assistance in making contact with relatives (physician or hospital) and dissemination plan		Desc ³⁵ Desc ³⁵	Desc ²¹	Desc ²⁶	Qual ^{15, 23,24}
	Provider follow-up					Desc^{47}
	Speaking with a genetic counselor		Desc^{35}			

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Perceptions of relatives, relatives' reported responses, and attitudes toward relatives

Lack of concern or interest by relatives [18, 20, 25], the possibility of relatives not agreeing to testing [25], relatives not believing participants [25] and concern regarding family reaction [25] were described as barriers to disclosure among probands and/or relatives. Not wanting to upset relatives and family [14, 35], and lack of understanding [20, 25, 28, 35, 40] were described as barriers to disclosure among probands and/or relatives. Unwilling and uncaring attitudes of relatives [25, 27, 35, 40], relevance for relative's stage of life [16, 20], and hostility toward advice or rejection [18, 25, 35] were described as barriers to disclosure and uptake among probands and/or (or right not relatives. Avoidance to know) [15, 20, 33, 43] and a perception that the relative appeared to be healthy [16] were barriers to uptake among probands and/or relatives.

Interpersonal barriers

Family communication, support, and dynamics

Emotional distance, estrangement, conflict, or resentment was described as a barrier among probands and/or relatives for both disclosure [21, 28] and uptake [18, 23, 42]. Similarly, not being in contact or not being emotionally close with family was described as a barrier among probands and/or relatives for both disclosure [21, 25, 35, 40] and uptake [16, 27]. General communication concerns was described as a barrier among probands and/or relatives for disclosure [25] and uptake [24, 27, 34]. In another study, no communication had a negative effect among probands on uptake of cascade testing [26]. A disappointing experience with disclosure early in the process was described as a barrier among probands and/or relatives for both disclosure [20] and uptake [31]. The impact on family relationship was described as a barrier among probands for disclosure [28]. The negative impact of disease on the family itself was described as a barrier among probands for uptake [34], and also reported to have a negative association among relatives for uptake [26]. Finally, language barriers among probands and/or relatives were barriers for disclosure [25].

Provider factors

Low provider awareness [29, 43] and lack of provider engagement [31] were described as barriers for probands and/or relatives for uptake of cascade testing.

Environmental barriers

Accessibility of genetic testing was the main environmental barrier studied in the literature. In two studies, probands and/or relatives described finances/cost [25, 43] or insurance coverage [25, 43] as barriers to both disclosure and genetic testing. In another study, relatives described finances, cost or insurance coverage as barriers to disclosure [35]. Access to genetic testing services was described as a barrier for probands and/or relatives for disclosure [20] and uptake [31]. Extra clinical referrals required to pursue genetic testing was also described as a barrier for probands and/or relatives for disclosure for probands and/or relatives for uptake [43].

Individual facilitators

Demographics

Older age was positively associated with disclosure among probands and/or relatives in one study [36]. Females had a positive association with disclosure (relatives) [22] and uptake of testing (proband and or relatives) [39]. Asian race of probands and/or relatives was positively associated with disclosure [36] and uptake [36]. Married individuals were more likely to engage both in disclosure (probands) [13] and uptake of testing (probands and/or relatives) [17]. High income of probands had a positive association with uptake of genetic testing by relatives in one study [13]. Last, high socioeconomic status of probands and/or relatives was associated with both disclosure and uptake of genetic testing in one study [36].

Clinical factors

Personal history of probands and/or relatives was positively associated with uptake [36] for probands and/or relatives in one study. Prior history of risk factors for relatives was described to be a facilitator for genetic testing [19]. Finally, receipt of unambiguous genetic test results by probands was positively associated with disclosure.

Attitudes, beliefs, and emotional responses of the individual

A need for emotional support from relatives was described as a facilitator in multiple studies for disclosure [20, 35, 40] and uptake [18] among both probands and/or relatives. A high satisfaction with the decision to undertake genetic testing was positively associated with both disclosure [36] and uptake [36] among both probands and/or relatives. Similarly, knowledge of screening and risk reduction recommendations was positively associated with disclosure among both probands and/or relatives [36]. Intrinsic motivation [21] and high levels of perceived susceptibility [21] were described as facilitators for disclosure among proband and/or relatives. In one study, high levels of perceived control [22] was positively associated with disclosure among probands. Lastly, probands and/or relatives indicated that feeling forced by circumstances [33] also enabled uptake of genetic testing.

Perceptions of relatives, relatives' reported responses, and attitudes toward relatives

Prominently, a moral obligation toward relatives or a duty to inform was described in multiple studies as a facilitator for probands and/or relatives for both disclosure [14, 21, 35] and uptake [18, 31, 34, 42]. A desire to prevent harm in the family or duty to protect was described as a facilitator for probands and/or relatives for both disclosure [18, 35] and uptake [15, 27]. Relatives' right to know (separate from duty to inform) [23, 24, 29, 34] and a belief that information would help in making medical or lifestyle decisions [14, 27, 31, 35] were also facilitators for probands and/or relatives for both disclosure and uptake. Concern for relatives was described as a facilitator for disclosure [29] and uptake [20, 23]. A perception of relatives having a positive opinion on genetic testing was positively associated with disclosure among probands. A need for help in understanding the disease was a facilitator for disclosure (probands) [28] and uptake (probands and/or relatives) [16]. A desire to prevent relatives from feeling the same sorrow [16] was also a facilitator for uptake among probands and/or relatives.

Interpersonal facilitators

Family communication, support, and dynamics

The degree of closeness was investigated in several studies. When the relatives were children, there was a positive effect on disclosure [22]. In one study, siblings, aunts and uncles were more likely to pursue genetic testing [39]. When the relatives were first-degree relatives of probands, there was a positive association with genetic testing [41]. Solidarity [33] and support [43] from family members were described as facilitators for uptake of genetic testing by probands and/or relatives. Encouraging relatives to get testing was described as a facilitator for disclosure (probands and/or relatives) [28] and uptake [34] (probands) [34]. Providing relatives with information about risk was described as a facilitator for disclosure among probands and/or relatives [14, 28]. Active and open communication was described as a facilitator for disclosure among probands and/or relatives [29].

Provider factors

Materials to pass to relatives (e.g., genetic counseling note, family letter) were described as facilitators for both disclosure (relatives) [35] and uptake (probands and/or relatives) [38, 42]. Assistance in identifying relatives at risk was described among probands as a facilitator for uptake [42], and among relatives for disclosure [35]. Assistance in making contact with relatives and a dissemination plan were described as facilitators for disclosure [21, 35] and uptake [15, 23, 24] among probands and/or relatives. A physician's recommendation was described as a facilitator for both disclosure [21, 28, 40] and uptake [16] by probands and/or relatives. A referral to a genetics clinic [38] and provider follow-up [43] were described as facilitators for uptake among probands and/or relatives. Finally, speaking with a genetic counselor was described as a facilitator for disclosure by relatives [35].

Discussion

Several individual and interpersonal factors, and a few environmental factors, were described as barriers and facilitators to cascade testing for genetic conditions in studies included in our review. In particular, attitudes, beliefs and emotional responses both relating to the individual and their relatives were identified, with a large number of these factors reported in one study that used a survey design. Our findings suggest that there is a need to verify the role of these factors in the uptake of cascade testing using rigorous methods.

Factors relating to provider awareness and engagement were described as facilitators in included studies; conversely, lack of provider awareness and engagement were described as barriers in three studies. Previous studies evaluating cascade testing programs or interventions have shown that direct methods, where trained providers directly contact at-risk relatives of probands, are effective [46, 47]. Two studies [42, 43] included in this review examined the acceptability of direct vs. indirect approaches (where cascade testing is primarily patient-mediated) through qualitative methods, and findings from these studies indicate that even though direct methods may be more effective, patients expressed a preference for patient-mediated approaches as this emphasized autonomy and privacy, and was less threatening to relatives. Further, the regulatory landscape for provider-directed communication is not described in our included studies, and cannot be directly inferred (except in certain settings, such as the United States). Thus, the extracted facilitators from these two studies and others suggest several approaches for improving cascade testing in settings where a patient-mediated method is the norm,

necessary (due to regulations) or preferred. In these settings, assistance in identifying at-risk relatives, creating a dissemination plan, receiving materials to pass on to relatives and follow-up were potential facilitators for cascade testing. These approaches could alleviate reported barriers regarding low or perceived lack of knowledge, as well as barriers such as not knowing who was at risk in the family.

Provider recommendation to share results with family was also described as a facilitator in several studies. This finding matches that of another study that examined nondirective approaches to counseling in families with BRCA1/ 2 gene variants, and suggested that more directive approaches are warranted in hereditary cancers, even if direct provider contact with relatives is not possible [48]. Overall, our results indicate that the role of the provider is critical in the process of cascade testing, and that the nature of provider engagement with patients and their relatives need to be optimized based on the environment and patientpreferences.

Family support, communication and dynamics play a key role in cascade testing, and several factors played a role either as barriers or facilitators. Indeed, it is well-known that genetic testing affects family relationships, with effects ranging from health benefits for relatives to strained relationships with family members. Interventions that strengthen family communication and increase family support, in addition to provider engagement, could optimize cascade testing. The process of genetic counseling typically incorporates psychosocial support for patients and their relatives, and interventions focused on enhancing family support and communication could be integrated in this process [49].

Our results also indicate that few studies have assessed factors outside the individual and interpersonal levels. In particular, contextual characteristics of the study setting and environmental barriers such as access to insurance, costs, etc. may be less relevant in countries with single payersystems, but may play a more important role in countries such as the United States. Future work should examine the relative changeability and importance of multilevel barriers and facilitators for cascade testing. In addition it will be important not only to identify individual and interpersonal level determinants of cascade testing but also those on the environmental level (encompassing the organizational or institutional, community and public policy levels within the Social Ecological Model) to enable a comprehensive understanding of barriers and facilitators specific to various settings and contextual factors, as well as to build effective interventions.

Finally, studies across autosomal dominant and recessive disorders were included to capture barriers and facilitators to the process of cascade testing, whenever family communication and testing were explicitly studied. However, the motivation for cascade testing for variants that confer risk of developing disease (e.g., BRCA1/2 variants) versus those that confer risk for children inheriting a disease (e.g., cystic fibrosis) may drive the differences in the nature of individual and interpersonal barriers and facilitators described in our results.

Limitations

First, we performed a narrative synthesis of the literature, as a meta-analysis was not feasible given that studies in this area did not assess effect sizes of barriers and facilitators on cascade testing. Second, inconsistent terminology was used across studies for the factors investigated, so we were able to merge the extracted barriers and facilitators in only a few instances. Third, there is a potential for bias as we integrated findings from both quantitative and qualitative studies, a majority of studies used a descriptive study design, a majority of the barriers to disclosure were extracted from one study [25], and most of the non-demographic factors were described as barriers or facilitators, instead of being reported as effect-sizes. Fourth, as with any systematic review, it is possible that we may have missed relevant literature. Finally, from an ethical perspective, receiving all information necessary to make an informed decision about testing and being offered testing may be the most appropriate measures for evaluating interventions for cascade testing. However, we used family communication and receiving genetic testing as approximate outcomes, because many studies did not sufficiently discriminate between these distinct processes in outcome measurement. Thus, some barriers from our results (e.g., fear of negative impact on family relationships affecting uptake) may arise from receiving information about testing, being offered testing, but ultimately choosing not to engage in testing.

Conclusions

Findings from this systematic review can inform additional formative work. Future research should examine the role of identified barriers and facilitators for cascade testing using rigorous, theory-informed methods. Taken together this work can inform future development of interventions to improve cascade testing outcomes.

Funding This work was supported by the National Center for Advancing Translational Sciences, National Institutes of Health through Grant KL2TR002490 to MCR. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript. The content is solely the responsibility of the authors and does not necessarily represent the official position of the National Institutes of Health or the Centers for Disease Control and Prevention.

Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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References

- Gupta S, Ahnen DJ, Chen L-M, Chung DC, Cooper G, Early DS, et al. NCCN guidelines Version 3. 2019 genetic/familial high-risk assessment: Colorectal [Internet]. [cited 2019 Dec 18]. 2019. https://www.nccn.org/professionals/physician_gls/pdf/genetics_colon. pdf.
- Cascade Testing: Testing Women for Known Hereditary Genetic Mutations Associated With Cancer - ACOG [Internet]. [cited 2019 Dec 18]. https://www.acog.org/Clinical-Guidance-and-Publica tions/Committee-Opinions/Committee-on-Gynecologic-Practice/ Cascade-Testing-Testing-Women-for-Known-Hereditary-Genetic-Mutations-Associated-With-Cancer.
- Goldberg AC, Hopkins PN, Toth PP, Ballantyne CM, Rader DJ, Robinson JG, et al. Executive summary familial hypercholesterolemia: screening, diagnosis and management of pediatric and adult patients clinical guidance from the national lipid association expert panel on familial hypercholesterolemia background and rationale. J Clin Lipidol. 2011;5:1–8.
- Beitsch PD, Whitworth PW, Hughes K, Patel R, Rosen B, Compagnoni G, et al. Underdiagnosis of hereditary breast cancer: are genetic testing guidelines a tool or an obstacle? J Clin Oncol. 2019;37:453–60.
- Abul-Husn NS, Manickam K, Jones LK, Wright EA, Hartzel DN, Gonzaga-Jauregui C, et al. Genetic identification of familial hypercholesterolemia within a single U.S. health care system. Science. 2016;354:aaf7000.
- Hampel H, De La Chapelle A. The search for unaffected individuals with lynch syndrome: do the ends justify the means? Cancer Prev Res. 2011;4:1–5.
- Roberts MC, Dotson WD, DeVore CS, Bednar EM, Bowen DJ, Ganiats TG, et al. Delivery of cascade screening for hereditary conditions: a scoping review of the literature. Health Aff. 2018;37:801–8.
- Moher D, Liberati A, Tetzlaff J, Altman DG. Preferred reporting items for systematic reviews and meta-analyses: the PRISMA statement. J Clin Epidemiol. 2009;62:1006–12.
- 9. Covidence Knowledge Base [Internet]. [cited 2019 Oct 21]. https://support.covidence.org/help.
- Schardt C, Adams MB, Owens T, Keitz S, Fontelo P. Utilization of the PICO framework to improve searching PubMed for clinical questions. BMC Med Inf Decis Mak. 2007;7:16.
- Golden SD, Earp JAL. Social ecological approaches to individuals and their contexts: twenty years of health education & behavior health promotion interventions. Health Educ Behav. 2012;39:364–72.
- Nha HONG Q, Pluye P, Fàbregues S, Bartlett G, Boardman F, Cargo M, et al. MIXED METHODS APPRAISAL TOOL (MMAT) VERSION 2018 User guide.
- Benson G, Witt DR, VanWormer JJ, Campbell SM, Sillah A, Hayes SN, et al. Medication adherence, cascade screening, and lifestyle patterns among women with hypercholesterolemia: Results from the WomenHeart survey. J Clin Lipidol. 2016;10:937–43.
- Burns C, McGaughran J, Davis A, Semsarian C, Ingles J. Factors influencing uptake of familial long QT syndrome genetic testing. Am J Med Genet A. 2016;170a:418–25.

- Hardcastle S, Legge E, Laundy C, Egan S, French R, Watts G, et al. Patients' Perceptions and Experiences of Familial Hypercholesterolemia, Cascade Genetic Screening and Treatment. Int J Behav Med. 2015;22:92–100.
- Ishii N, Arai M, Koyama Y, Ueno M, Yamaguchi T, Kazuma K, et al. Factors affecting encouragement of relatives among families with Lynch syndrome to seek medical assessment. Fam Cancer. 2011;10:649–54.
- Lerman C, Hughes C, Trock BJ, Myers RE, Main D, Bonney A, et al. Genetic testing in families with hereditary nonpolyposis colon cancer. JAMA. 1999;281:1618–22.
- Lieberman S, Lahad A, Tomer A, Koka S, BenUziyahu M, Raz A, et al. Familial communication and cascade testing among relatives of BRCA population screening participants. Genet Med. 2018;20:1446–54.
- Maxwell SJ, Molster CM, Poke SJ, O'Leary P. Communicating familial hypercholesterolemia genetic information within families. Genet Test Mol Biomark. 2009;13:301–6.
- McClaren BJ, Aitken M, Massie J, Amor D, Ukoumunne OC, Metcalfe SA. Cascade carrier testing after a child is diagnosed with cystic fibrosis through newborn screening: investigating why most relatives do not have testing. Genet Med. 2013;15:533–40.
- Mesters I, Ausems M, Eichhorn S, Vasen H. Informing one's family about genetic testing for hereditary non-polyposis colorectal cancer (HNPCC): a retrospective exploratory study. Fam Cancer. 2005;4:163–7.
- Montgomery SV, Barsevick AM, Egleston BL, Bingler R, Ruth K, Miller SM, et al. Preparing individuals to communicate genetic test results to their relatives: report of a randomized control trial. Fam Cancer. 2013;12:537–46.
- Ormondroyd E, Oates S, Parker M, Blair E, Watkins H. Presymptomatic genetic testing for inherited cardiac conditions: a qualitative exploration of psychosocial and ethical implications. Eur J Hum Genet. 2014;22:88–93.
- Pentz RD, Peterson SK, Watts B, Vernon SW, Lynch PM, Koehly LM, et al. Hereditary nonpolyposis colorectal cancer family members' perceptions about the duty to inform and health professionals' role in disseminating genetic information. Genet Test. 2005;9:261–8.
- Campbell M, Humanki J, Zierhut H. A novel approach to screening for familial hypercholesterolemia in a large public venue. J Community Genet. 2017;8:35–44.
- Raspa M, Edwards A, Wheeler A, Bishop E, Bailey D. Family communication and cascade testing for fragile X syndrome. J Genet Couns. 2016;25:1075–84.
- Smart A. Impediments to DNA testing and cascade screening for hypertrophic cardiomyopathy and Long QT syndrome: a qualitative study of patient experiences. J Genet Couns 2010;19:630–9.
- Stoffel EM, Ford B, Mercado RC, Punglia D, Kohlmann W, Conrad P, et al. Sharing genetic test results in Lynch syndrome: communication with close and distant relatives. Clin Gastroenterol Hepatol. 2008;6:333–8.
- Suttman A, Pilarski R, Agnese DM, Senter L. "Second-class status?" insight into communication patterns and common concerns among men with hereditary breast and ovarian cancer syndrome. J Genet Couns. 2018;27:885–93.
- Truong TH, Kim NT, Nguyen MNT, Pang J, Hooper AJ, Watts GF, et al. Homozygous familial hypercholesterolaemia in Vietnam: case series, genetics and cascade testing of families. Atherosclerosis. 2018;277:392–8.
- Ulph F, Cullinan T, Qureshi N, Kai J. Parents' responses to receiving sickle cell or cystic fibrosis carrier results for their child following newborn screening. Eur J Hum Genet. 2015;23:459–65.
- Louter L, Defesche J, Roeters van Lennep J. Cascade screening for familial hypercholesterolemia: Practical consequences. Atheroscler Suppl. 2017;30:77–85.

- 33. van Maarle MC, Stouthard ME, Marang-van de Mheen PJ, Klazinga NS, Bonsel GJ. How disturbing is it to be approached for a genetic cascade screening programme for familial hypercholesterolaemia? Psychological impact and screenees' views. Community Genet. 2001;4:244–52.
- 34. Van Rijn MA, De Vries BBA, Tibben A, Van Den Ouweland AMW, Halley DJJ, Niermeijer MF. DNA testing for fragile X syndrome: Implications for parents and family. J Med Genet. 1997;34:907–11.
- 35. Wurtmann E, Steinberger J, Veach PM, Khan M, Zierhut H. Risk communication in families of children with familial hypercholesterolemia: identifying motivators and barriers to cascade screening to improve diagnosis at a Single Medical Center. J Genet Couns. 2018;28:50–8.
- Cheung EL, Olson AD, Yu TM, Han PZ, Beattie MS. Communication of BRCA results and family testing in 1,103 high-risk women. Cancer Epidemiol Biomark Prev. 2010;19: 2211–9.
- 37. de Souza Silva PR, Jannes CE, Oliveira TGM, Gómez LMG, Krieger JE, Santos RD, et al. Predictors of family enrollment in a genetic cascade screening program for familial hypercholesterolemia. Arq Bras Cardiol. 2018;111:578–84.
- Dilzell K, Kingham K, Ormond K, Ladabaum U. Evaluating the utilization of educational materials in communicating about Lynch syndrome to at-risk relatives. Fam Cancer. 2014;13:381–9.
- Dugueperoux I, L'Hostis C, Audrezet MP, Rault G, Frachon I, Bernard R, et al. Highlighting the impact of cascade carrier testing in cystic fibrosis families. J Cyst Fibros. 2016;15:452–9.
- 40. Finlay E, Stopfer JE, Burlingame E, Evans KG, Nathanson KL, Weber BL, et al. Factors determining dissemination of results and uptake of genetic testing in families with known BRCA1/2 mutations. Genet Test. 2008;12:81–91.

- Hagoel L, Dishon S, Almog R, Silman Z, Bisland-Becktell S, Rennert G. Proband family uptake of familial-genetic counselling. Psychooncology. 2000;9:522–7.
- Hallowell N, Jenkins N, Douglas M, Walker S, Finnie R, Porteous M, et al. Patients' experiences and views of cascade screening for familial hypercholesterolemia (FH): a qualitative study. J Community Genet. 2011;2:249–57.
- van El CG, Baccolini V, Piko P, Cornel MC. Stakeholder views on active cascade screening for familial hypercholesterolemia. Healthcare. 2018;6:108.
- 44. Hallowell N, Jenkins N, Douglas M, Walker S, Finnie R, Porteous M, et al. A qualitative study of patients' perceptions of the value of molecular diagnosis for familial hypercholesterolemia (FH). J Community Genet. 2017;8:45–52.
- Tong A, Sainsbury P, Craig J. Consolidated criteria for reporting qualitative research (COREQ): A 32-item checklist for interviews and focus groups. Int J Qual Heal Care. 2007;19:349–57.
- 46. Suthers GK, Armstrong J, McCormack J, Trott D. Letting the family know: balancing ethics and effectiveness when notifying relatives about genetic testing for a familial disorder. J Med Genet. 2006;43:665–70.
- 47. Marks D, Thorogood M, Neil SM, Humphries SE, Neil HA. Cascade screening for familial hypercholesterolaemia: implications of a pilot study for national screening programmes. J Med Screen. 2006;13:156–9.
- 48. Sermijn E, Goelen G, Teugels E, Kaufman L, Bonduelle M, Neyns B, et al. The impact of proband mediated information dissemination in families with a BRCA1/2 gene mutation. J Med Genet. 2004;41:e23.
- 49. Sturm AC. The role of genetic counselors for patients with familial hypercholesterolemia. Curr Genet Med Rep. 2014; 2:68–74.