COMMENT OPEN

ESHG

Public attitudes challenge clinical practice on genetic risk disclosure in favour of healthcare-provided direct dissemination to relatives

Anna Rosén ¹^M, Mateja Krajc², Hans Ehrencrona ^{3,4} and Svetlana Bajalica-Lagercrantz^{5,6}

© The Author(s) 2023

European Journal of Human Genetics (2024) 32:6-7; https://doi.org/10.1038/s41431-023-01428-3

ENABLING PREVENTIVE MEASURES IN THE ERA OF PRECISION MEDICINE

The increased usage of genetic testing for treatment stratification within the era of precision medicine entails the potential to detect germline genetic risk variants. Germline genetic testing often has implications not only for the individual patient but also for their genetic relatives. This is especially true for high-penetrance pathogenic variants associated with conditions such as familial hypercholesterolemia and hereditary cancer risk syndromes like Lynch syndrome and the hereditary breast and ovarian cancer syndrome. For these conditions, targeted prevention programs are available, and cascade screening is cost-effective [1, 2]. It is therefore highly relevant to find effective strategies to disclose information from the genetic investigation to healthy relatives at risk. Informing relatives at risk enables equitable access to pre-test genetic counselling and a possibility for them to make an informed decision about predictive genetic testing as well as prevention.

GENETIC RISK DISCLOSURE—TIME TO CHANGE TO A HEALTHCARE-MEDIATED DIRECT APPROACH?

Current practice in most countries is to encourage index patients to inform their relatives about the potential impact of a genetic risk assessment. Several studies have explored the barriers and facilitators of family communication, and some have also tested interventions to improve efficacy. It seems that tailored genetic counselling with additional follow-up can increase both the proportion of informed relatives and relatives who contact the genetics clinic, but the data are not conclusive [3, 4]. However, another large meta-analysis on hereditary cancer risk disclosure shows that with family-mediated disclosure, the uptake of genetic counselling in relatives is about 35%, whereas the uptake almost doubles (63%) when using a healthcare-mediated direct contact approach [5].

WHAT DO PEOPLE THINK OF DIRECT CONTACT (FROM A HYPOTHETICAL POINT OF VIEW)?

In this issue of *European Journal of Human Genetics*, Tiller et al. [6] present interesting data on questionnaires directed to the general public. Hence, most participants lacked the experience of belonging to a family with a disease pattern often seen in a hereditary condition. The respondents were briefly introduced to the concept of medically actionable genetic conditions, the importance of sharing information with (genetic) relatives, and current standard practice with family-mediated risk disclosure. On a hypothetical question, most respondents (85%) expressed a preference for being informed about potential genetic risks for future health problems that can be prevented or treated early. However, it remains unclear to what extent respondents with real-life experience of familial disease aggregation would be in favour of such risk awareness.

When provided with two different types of information letters, 67% of respondents preferred a letter with more specific information about the variant in the family, health risks, and preventive measures, whereas 21% preferred a letter containing more general information. Notably, when asked **from whom** they preferred to receive the letter, less than a tenth (8.4%) preferred to receive the letter from a family member. Thus, only a minority preferred to receive information in the way commonly used in current clinical practice in most countries, where the index often is provided with a 'family letter' to further distribution to relatives.

Tiller et al. also show that the majority (68%) would prefer that healthcare providers disseminate the letter directly to them. Interestingly, one-third of them would also like to be contacted by a family member for an explanation. The preference that healthcare providers are directly involved in genetic information disclosure to at-risk relatives is in agreement with reports of public opinion data from Belgium [7], Sweden [8], and Denmark [9]. The Australian data contribute to an increasing body of evidence showing that—in a hypothetical situation—the public envisions a practice that is not implemented today.

Received: 29 June 2023 Accepted: 3 July 2023 Published online: 20 July 2023

¹Department of Radiation Sciences, Oncology, Umeå University, Umeå, Sweden. ²Department of Clinical Cancer Genetics, Institute of Oncology Ljubljana, Ljubljana, Slovenia. ³Division of Clinical Genetics, Department of Laboratory Medicine, Lund University, Lund, Sweden. ⁴Department of Genetics, Pathology and Molecular Diagnostics, Office for Medical Services, Region Skåne, Lund, Sweden. ⁵Department of Oncology-Pathology, Karolinska Institutet, Stockholm, Sweden. ⁶Hereditary Cancer Unit, Karolinska University Hospital, Stockholm, Sweden. ^{Se}email: anna.rosen@umu.se

FUTURE PERSPECTIVES

Even though family-mediated disclosure is well established, it entails complex ethical questions. For example, the dissemination of information to at-risk relatives depends on the willingness of the index case. An approach with direct healthcare-provided risk disclosure could also be challenging. There are only a limited number of studies investigating the real experience of receiving risk information through healthcare-provided direct contact. Even though these studies indicate that the direct approach may be accepted [9] and safe concerning anxiety levels [10, 11], further analyses of the impact on at-risk individuals are needed. There is also a lack of studies evaluating the effectiveness of the direct contact approach concerning the uptake of cascade testing among family members and the degree of enrolment in surveillance. This could preferably be approached by large randomised controlled trials. Of note, the implementation of the direct approach challenges issues concerning patient's autonomy and confidentiality, and relatives right not to know. On the other hand, the direct approach has the potential to safeguard the relatives' right to receive information with potential relevance for their health, i.e., their right to know.

The healthcare system faces a growing need to accommodate an ever-increasing number of (healthy) at-risk individuals. Supplementing family-mediated disclosure with healthcareprovided direct information could constitute an improvement. Hereditary aspects of germline genetic findings are challenging healthcare to a paradigm shift from the patient to the family as the unit of care. This involves a multitude of aspects including not only how to practically identify and define at-risk individuals but also how to administer and store family-level data. Countryspecific legislation and privacy rules govern the possibilities and limitations of these endeavours.

REFERENCES

- Sessa C, Balmana J, Bober SL, Cardoso MJ, Colombo N, Curigliano G, et al. Risk reduction and screening of cancer in hereditary breast-ovarian cancer syndromes: ESMO Clinical Practice Guideline. Ann Oncol. 2023;34:33–47.
- Nherera L, Marks D, Minhas R, Thorogood M, Humphries SE. Probabilistic costeffectiveness analysis of cascade screening for familial hypercholesterolaemia using alternative diagnostic and identification strategies. Heart. 2011;97:1175–81.
- Young AL, Imran A, Spoelma MJ, Williams R, Tucker KM, Halliday J, et al. Probandmediated interventions to increase disclosure of genetic risk in families with a BRCA or Lynch syndrome condition: a systematic review. Eur J Hum Genet. 2023;31:18–34.
- Ballard LM, Band R, Lucassen AM. Interventions to support patients with sharing genetic test results with at-risk relatives: a synthesis without meta-analysis (SWiM). Eur J Hum Genet. 2023. https://www.ncbi.nlm.nih.gov/pubmed/37344572.
- Frey MK, Ahsan MD, Bergeron H, Lin J, Li X, Fowlkes RK, et al. Cascade testing for hereditary cancer syndromes: should we move toward direct relative contact? A systematic review and meta-analysis. J Clin Oncol. 2022;40:4129–43.
- Tiller JM, Stott A, Finlay K, Boughtwood T, Madelli EO, Horton A, et al. Direct notification by health professionals of relatives at-risk of genetic conditions (with patient consent): views of the Australian public. Eur J Hum Genet. 2023. https:// www.ncbi.nlm.nih.gov/pubmed/37280361.

 Phillips A, Dewitte I, Debruyne B, Vears DF, Borry P. Disclosure of genetic risk in the family: a survey of the Flemish general population. Eur J Med Genet. 2023;66:104800. 7

- Andersson A, Hawranek C, Ofverholm A, Ehrencrona H, Grill K, Hajdarevic S, et al. Public support for healthcare-mediated disclosure of hereditary cancer risk information: results from a population-based survey in Sweden. Hered Cancer Clin Pract. 2020;18:18.
- Petersen HV, Frederiksen BL, Lautrup CK, Lindberg LJ, Ladelund S, Nilbert M. Unsolicited information letters to increase awareness of Lynch syndrome and familial colorectal cancer: reactions and attitudes. Fam Cancer. 2019;18:43–51.
- Sermijn E, Delesie L, Deschepper E, Pauwels I, Bonduelle M, Teugels E, et al. The impact of an interventional counselling procedure in families with a BRCA1/2 gene mutation: efficacy and safety. Fam Cancer. 2016;15:155–62.
- Aktan-Collan K, Haukkala A, Pylvanainen K, Jarvinen HJ, Aaltonen LA, Peltomaki P, et al. Direct contact in inviting high-risk members of hereditary colon cancer families to genetic counselling and DNA testing. J Med Genet. 2007;44:732–8.

AUTHOR CONTRIBUTIONS

AR drafted the comment. MK, HE and SB-L provided feedback and helped edit the manuscript. All authors have read and approved the final draft.

FUNDING

No funding was received for writing this comment. Open access funding provided by Umea University.

COMPETING INTERESTS

The authors declare no competing interests.

ADDITIONAL INFORMATION

Correspondence and requests for materials should be addressed to Anna Rosén.

Reprints and permission information is available at http://www.nature.com/ reprints

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http:// creativecommons.org/licenses/by/4.0/.

© The Author(s) 2023