

Direct-to-consumer misleading information on cancer risks calls for an urgent clarification of health genetic testing performed by commercial companies

Antoine de Pauw, Mathias Schwartz, Chrystelle Colas, Lisa Golmard,

Dominique Stoppa-Lyonnet

▶ To cite this version:

Antoine de Pauw, Mathias Schwartz, Chrystelle Colas, Lisa Golmard, Dominique Stoppa-Lyonnet. Direct-to-consumer misleading information on cancer risks calls for an urgent clarification of health genetic testing performed by commercial companies. European Journal of Cancer, 2020, 132, pp.100 - 103. 10.1016/j.ejca.2020.03.007. hal-03490846

HAL Id: hal-03490846 https://hal.science/hal-03490846

Submitted on 22 Aug2022

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution - NonCommercial 4.0 International License

Version of Record: https://www.sciencedirect.com/science/article/pii/S0959804920301489 Manuscript_0d64eec756734ad3accbac05390fb36c

Title:

Direct-to-consumer misleading information on cancer risks calls for an urgent clarification of health genetic testing performed by commercial companies.

Authors:

Antoine de Pauw*(1,2), Mathias Schwartz*(1), Chrystelle Colas (1,2), Lisa Golmard (1,2),

Dominique Stoppa-Lyonnet (1,3, 4).

* These authors contributed equally to this work

Affiliations:

- (1) Department of Genetics, Institut Curie, Paris, France
- (2) Paris Sciences Lettres Research University, Paris, France
- (3) INSERM U830, Institut Curie Paris, Paris, France
- (4) Paris-University, Paris, France

Authors contributions:

AP and MS wrote the first draft of the manuscript; CC, LG and DS revised the manuscript for major intellectual content; AP, CC, and DS performed specialized medical consultations; MS and LG analyzed the genetic data.

Corresponding author:

Pr. Dominique Stoppa-Lyonnet

Department of Genetics, Institut Curie, 26 rue d'Ulm 75005 Paris, France

dominique.stoppalyonnet@curie.fr

+33 (0) 1 44 32 46 97

Abstract

DTC commercial companies offer genetic tests that are presented as allowing individuals the opportunity to increase their capacities to be in charge of their own healthcare managements. DTC companies deny performing medical tests, yet they provide data based on sequencing multigene panel or whole exome. This contradiction allows these companies to escape the requirements of a regulated medical practice that guarantees the quality of the tests, as well as the information and support for tested individuals. Herein, we illustrate the lack of such requirements by analyzing the bad experience of a young man who dealt with DTC health genetic testing companies. There is an emergency for DTC testing to be either deprived of any medically relevant information, or carried out in a legally regulated medical framework.

A striking example of bad practices

Mr. C., a 33-year-old French man with no relevant personal or family history of cancer or genetic disease, had an experience with a DTC commercial company which illustrates how "health" genetic tests are *de facto* medical information. This young man decided in December 2018 "by curiosity" to benefit from a Christmas discount on the Whole Genome Sequencing (WGS) offered by a DTC company. Mr. C. chose the *Premium Whole Genome for Health* test that proposed the analysis of genes involved in 70 genetic diseases or syndromes, and thus to "get actionable insights to drive his personalized prevention". By ordering the test on the company's website, Mr. C. was supposed to accept a 10,000-word "Terms of service" agreement, consisting of no less than 29 clauses, with only one of them referring to WGS, or rather, its limitations. In fact, Mr. C. did not give any specific consent. He sent a salivary sample in January 2019. Three months later, he received an e-mail to download an eighteen-page PDF file of genetic results. This "Wellness and Lifestyle Report" listed the results of the 70 explored diseases, each of them annotated by a green or red light. Among them, eight

cancer predisposition syndromes were listed, with notable exception of both Hereditary Breast and Ovarian Cancer syndrome (HBOC caused by BRCA1 and BRCA2 genes) and Hereditary Non-Polyposis Colorectal Cancer syndrome (HNPCC caused by mismatch repair genes). Mr. C.'s report exhibited two "red lights" in the "cancer" section, and therefore two supposed monoallelic pathogenic variants: one in the APC gene associated with familial adenomatous polyposis (FAP), and one in the TP53 gene associated with Li-Fraumeni Syndrome (LFS). FAP confers a very high risk of colorectal cancer before the age of 40, and requires preventive removal of the colon and rectum when polyposis occurs. LFS is associated with a major risk of multiple cancers including sarcoma and brain tumors, often diagnosed within the pediatric age range with a recommended annual whole-body MRI monitoring. In addition, the risk of transmission to the child for both these high-risk cancer predispositions, following an autosomal dominant inheritance, is 50%. The PDF file sent to Mr C. included a simple cut and paste from text published by National Institutes of Health (NIH) on exposed cancer risks and prevention modalities regarding these two predispositions. However, the file did not state that Mr. C. should consult a healthcare specialist provider, nor that these results need to be confirmed by an independent test. Terrified to learn that he was at very high risk for multiple and early-onset cancers, Mr. C. contacted the Institut Curie Genetics Department by email and sent the report. Even though the nomenclature used to name the pathogenic variants did not follow the Human Genome Variation Society (HGVS) guidelines,[1] we confirmed that both variants have indeed been reported as pathogenic for a period of time: NM_000038.5: c.423-1G>A, p.?, and NM_000546.5: c.734G>A, p.(Gly245Asp) (HGVS nomenclature), at the APC and *TP53* loci respectively.[2,3]

We then quickly arranged a meeting with Mr. C. at our genetic clinics. We inquired on his current state of health, as well as his personal and family medical histories. We proposed to carry out genetic testing targeted on the *APC* and *TP53* reported variants on two independent

tissue samples (whole blood and buccal swab), as is routine. Mr. C. agreed to the analysis and signed a consent form. During the five-week wait for the results, and seeing as Mr. C. mentioned some occasional rectal bleeding, a gastro-intestinal endoscopy was performed. This examination turned out to be normal and no histopathological abnormalities were detected in systematic gastric and colonic biopsies.

Sanger DNA sequencing failed to detect any of the variants reported by the DTC laboratory. We then asked Mr. C. to get back in contact with the company in order to obtain the raw data of his WGS analysis. He sent us the FASTQ (raw sequencing data) and VCF (*variant call format*, listing all detected genetic variants) files. The low sequencing depth at these positions, the poor quality of the variant calling, as well as the low allelic ratios of altered-to-wildtype alleles (data available on request), indicated that both variants were indeed obvious false positives.

In brief, the lack of both pre and post-genetic testing information, the lack of consent, the lack of post-result accompaniment, the generated anxiety and the prescription of an at-risk medical examination, induced by two false positive results, summarize the bad experience of Mr. C. in the brand new world of DTC health genetic testing by commercial companies.

A critical need for clarification

While this report might seem anecdotal, it is unfortunately not an isolated case and it illustrates how DTC genetic testing often blurs the lines between a medical device and a consumer product.[4,5] Most DTC companies have aggressive commercial marketing plans based on the enhanced autonomy of consumers and their allegedly increased capacity to be in charge of their own healthcare, without the intermediary of physicians and long waiting lists

for specialist appointments.[6] However, there is often a strong contradiction between, on the one hand DTC company results they report that *de facto* have medical value, and on the other hand pre-test information – when the consumer reads it carefully – which declines all responsibility for the results. This deliberate ambiguity allows DTC commercial companies to escape both the legal requirements of medical practice, and the regimented recommendations made to health providers in the field of genetics.[7,8] In addition, because people who have received such positive DTC test results are often afraid and seek advice from public genetic clinics, undue costs to public services are induced.[9]

The two false positive results of this report also illustrate how loose sequence interpretation, left to algorithms and automatic report writing devices, with no one officially responsible for the result, can lead to a disaster. Indeed, any professional with skills in medical genetics would have been cautious facing such a result. A young adult without any symptoms or family history, carrying two unlinked pathogenic variants, each associated with a high-risk cancer predisposing syndrome, whose prevalence in the general population is individually less than 1/10,000, the likelihood of observing both in a single individual being less than $1/10^8$. This result seems virtually impossible in an asymptomatic adult. A recent study has indicated that 40% of variants in a variety of genes reported in DTC raw data were false positives, underlying the importance of confirming any positive DTC results in a clinical grade laboratory with a legally recognized competence in medical genetics.[10] In addition, genetic tests with medical purposes performed by DTC companies could be falsely reassuring for customers. Indeed, "whole genome sequencing" appellation can be misleading if some genes (causing HBOC or HNPCC for instance) are not analyzed, as in the case of Mr C.. Another source of falsely reassuring information is the use of technologies that do not cover the whole coding sequence of analyzed genes. For example, the Food and Drug Administration allowed in 2017 the company 23andMe to test three frequent *BRCA1* and *BRCA2* pathogenic variants. It is clear that testing only three variants does not provide much medical information. On the contrary, a negative result can provide false reassurance. Actually, alleged "health" information sold to customers might be a pretext to get their genetic data on which they lose any control. [11]

Besides wrongly interpreting sequencing results, another cause of false positives is the overinterpretation of variants of unknown significance (VUS, class 3) - or even benign variants-, some of them being often reported as disease causing. In a survey of 3,672 women who had BRCA1/BRCA2 testing, Kurian et al. reported that in the USA 51% of women carrying a BRCA1/BRCA2 VUS underwent bilateral prophylactic mastectomy even in the absence of a significant family history.[12] Although not specific to them, these misinterpretations of VUS are at higher risks in DTC companies than in accredited laboratories. Here again, the contradiction between a market-driven genetic testing proposal, and the will to escape any possible medical responsibility, explains many reported cases of variant over-interpretation. Another essential criterion of the quality of a genetic test is the demonstration of its predictive value, that is indeed its clinical utility, a criterion little known by DTC companies, especially when they recommend specific diets or physical exercises according to non-validated genetic data.[7,8,13] At last difficult question raised here is that, by definition, DTC tests are aimed at the general population, ignoring the possible modifying factors of the penetrance of a genetic trait. This is of particular concern for cancer predisposing genes, such as the BRCA1 and BRCA2 genes. Yet still a subject of debate, the United States Preventive Services Task Force (USPSTF) and the American College of Medical Genetics (ACMG) do not encourage general population BRCA1/2 testing and recently made cautious recommendations toward broadening criteria.[14-16] At present, the knowledge of medical genetics, and in particular cancer genetics, is still in progress. It is the time of worldwide epidemiological genetic studies conducted by international consortia such as ENIGMA, IBCCS and InSiGHT in the field of cancer.[17–19]

Predictive genetic tests leading to irreversible cancer-preventive surgical decisions, to specific cancer treatments or to the identification of recessive disease carrier for reproductive purposes, are indeed already present in the field of medical practice. DTC practices challenge the current legislation. There is an emergency for DTC testing to be either really deprived of any medically relevant information or carried out in a legally regulated medical framework guaranteeing the quality of tests, as well as the information and support of individuals. The fact that individuals have the free will to access information on and from their own genome, does not imply that laboratories can freely deliver false and dangerous information, on the contrary. With the risk of losing credibility due to such improper practices, driven by the market rather than the clinical utility, the future of medical genetics is at stake.

Acknowledgments:

The authors would like to thank Mr. C. for accepting to share his experience, Prof. Stanislas Lyonnet for peer reviewing the manuscript, and Ms. Sarah Colautti for reviewing the English.

References

- [1] den Dunnen JT, Dalgleish R, Maglott DR, Hart RK, Greenblatt MS, McGowan-Jordan J, et al. HGVS Recommendations for the Description of Sequence Variants: 2016 Update. Hum Mutat 2016;37:564–9. https://doi.org/10.1002/humu.22981.
- [2] Spirio L, Green J, Robertson J, Robertson M, Otterud B, Sheldon J, et al. The identical 5' splice-site acceptor mutation in five attenuated APC families from Newfoundland demonstrates a founder effect. Hum Genet 1999;105:388–98. https://doi.org/10.1007/s004399900153.
- [3] Srivastava S, Zou Z, Pirollo K, Blattner W, Chang EH. Germ-line transmission of a mutated p53 gene in a cancer-prone family with Li–Fraumeni syndrome. Nature 1990;348:747–9. https://doi.org/10.1038/348747a0.
- [4] Kalokairinou L, Howard HC, Slokenberga S, Fisher E, Flatscher-Thöni M, Hartlev M, et al. Legislation of direct-to-consumer genetic testing in Europe: a fragmented regulatory landscape. J Community Genet 2018;9:117–32. https://doi.org/10.1007/s12687-017-0344-2.
- [5] Horton R, Crawford G, Freeman L, Fenwick A, Wright CF, Lucassen A. Direct-toconsumer genetic testing. BMJ 2019:15688. https://doi.org/10.1136/bmj.15688.
- [6] Marietta C, McGuire AL. Currents in Contemporary Ethics Direct-to-Consumer Genetic Testing: Is it the Practice of Medicine? J Law Med Ethics 2009;37:369–74. https://doi.org/10.1111/j.1748-720X.2009.00380.x.
- [7] European Society of Human Genetics. Statement of the ESHG on direct-to-consumer genetic testing for health-related purposes. Eur J Hum Genet 2010;18:1271–3. https://doi.org/10.1038/ejhg.2010.129.
- [8] ACMG Board of Directors. Direct-to-consumer genetic testing: a revised position statement of the American College of Medical Genetics and Genomics. Genet Med 2016;18:207–8. https://doi.org/10.1038/gim.2015.190.
- [9] Royal College of General Practitioners, The British Society for Genetic Medicine. Position Statement on Direct to Consumer Genomic Testing 2019.
- [10] Tandy-Connor S, Guiltinan J, Krempely K, LaDuca H, Reineke P, Gutierrez S, et al. False-positive results released by direct-to-consumer genetic tests highlight the importance of clinical confirmation testing for appropriate patient care. Genet Med 2018;20:1515–21. https://doi.org/10.1038/gim.2018.38.
- [11] Roberts JL, Hawkins J. When health tech companies change their terms of service. Science 2020;367:745–6. https://doi.org/10.1126/science.aaz6732.
- [12] Kurian AW, Li Y, Hamilton AS, Ward KC, Hawley ST, Morrow M, et al. Gaps in Incorporating Germline Genetic Testing Into Treatment Decision-Making for Early-Stage Breast Cancer. J Clin Oncol 2017;35:2232–9. https://doi.org/10.1200/JCO.2016.71.6480.
- [13] Colas C, Golmard L, de Pauw A, Caputo SM, Stoppa-Lyonnet D. "Decoding hereditary breast cancer" benefits and questions from multigene panel testing. The Breast 2019;45:29–35. https://doi.org/10.1016/j.breast.2019.01.002.
- [14] US Preventive Services Task Force, Owens DK, Davidson KW, Krist AH, Barry MJ, Cabana M, et al. Risk Assessment, Genetic Counseling, and Genetic Testing for *BRCA* -Related Cancer: US Preventive Services Task Force Recommendation Statement. JAMA 2019;322:652. https://doi.org/10.1001/jama.2019.10987.

- [15] Rutgers E, Balmana J, Beishon M, Benn K, Evans DG, Mansel R, et al. European Breast Cancer Council manifesto 2018: Genetic risk prediction testing in breast cancer. Eur J Cancer 2019;106:45–53. https://doi.org/10.1016/j.ejca.2018.09.019.
- [16] Green RF, Ari M, Kolor K, Dotson WD, Bowen S, Habarta N, et al. Evaluating the role of public health in implementation of genomics-related recommendations: a case study of hereditary cancers using the CDC Science Impact Framework. Genet Med 2019;21:28–37. https://doi.org/10.1038/s41436-018-0028-2.
- [17] Spurdle AB, Healey S, Devereau A, Hogervorst FBL, Monteiro ANA, Nathanson KL, et al. ENIGMA-Evidence-based network for the interpretation of germline mutant alleles: An international initiative to evaluate risk and clinical significance associated with sequence variation in BRCA1 and BRCA2 genes. Hum Mutat 2012;33:2–7. https://doi.org/10.1002/humu.21628.
- [18] Antoniou AC, Goldgar DE, Andrieu N, Chang-Claude J, Brohet R, Rookus MA, et al. A weighted cohort approach for analysing factors modifying disease risks in carriers of high-risk susceptibility genes. Genet Epidemiol 2005;29:1–11. https://doi.org/10.1002/gepi.20074.
- [19] Peltomaki P, Vasen H. Mutations predisposing to hereditary nonpolyposis colorectal cancer: Database and results of a collaborative study. The International Collaborative Group on Hereditary Nonpolyposis Colorectal Cancer. Gastroenterology 1997;113:1146–58. https://doi.org/10.1053/gast.1997.v113.pm9322509.