

Journal Pre-proof

A Systematic Review of Value Criteria for Next-Generation Sequencing/
Comprehensive Genomic Profiling to Inform Value Framework Development

Federico Augustovski, PhD, Carla Colaci, MSc, Mackenzie Mills, PhD, Danitza Chavez, MSc, Fernando Argento, MSc, Verónica Alfie, MSc, Andrés Pichon Riviere, PhD, Panos Kanavos, PhD, Andrea Alcaraz, MSc

PII: S1098-3015(24)00071-8

DOI: <https://doi.org/10.1016/j.jval.2024.02.002>

Reference: JVAL 3964

To appear in: *Value in Health*

Received Date: 12 September 2023

Revised Date: 6 February 2024

Accepted Date: 13 February 2024

Please cite this article as: Augustovski F, Colaci C, Mills M, Chavez D, Argento F, Alfie V, Riviere AP, Kanavos P, Alcaraz A, A Systematic Review of Value Criteria for Next-Generation Sequencing/ Comprehensive Genomic Profiling to Inform Value Framework Development, *Value in Health* (2024), doi: <https://doi.org/10.1016/j.jval.2024.02.002>.

This is a PDF file of an article that has undergone enhancements after acceptance, such as the addition of a cover page and metadata, and formatting for readability, but it is not yet the definitive version of record. This version will undergo additional copyediting, typesetting and review before it is published in its final form, but we are providing this version to give early visibility of the article. Please note that, during the production process, errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

Copyright © 2024, International Society for Pharmacoeconomics and Outcomes Research, Inc.
Published by Elsevier Inc.



Target Journal: [VIH]

TITLE

A Systematic Review of Value Criteria for Next-Generation Sequencing/Comprehensive Genomic Profiling to Inform Value Framework Development

Authors

Federico Augustovski, PhD¹, Carla Colaci, MSc¹, Mackenzie Mills, PhD², Danitza Chavez, MSc², Fernando Argento, MSc¹, Verónica Alfie, MSc¹, Andrés Pichon Riviere, PhD¹, Panos Kanavos, PhD²; Andrea Alcaraz, MSc¹.

1- Health Technology Assessment and Health Economics Department, Institute for Clinical Effectiveness and Health Policy (IECS)

2- Medical Technology Research Group, London School of Economics (LSE)

Corresponding author:

Federico Augustovski, MD, MSc, PhD

Health Technology Assessment and Health Economics Department

Institute for Clinical Effectiveness and Health Policy (IECS)

Doctor Emilio Ravignani 2024

Buenos Aires - Argentina

Email: Federico Augustovskigustovsi@iecs.org.ar

Phone number: +54 9 221 523 0188

Precis: A systematic review of value dimensions that could be considered for Next Generation Sequencing - Comprehensive Genomic Profiling as well as other healthcare technologies

Word Count: 4317

Number of Pages: 21

Number of Figures: 3

Number of Tables: 3

Appendix: Pages: 27 Figures: 1 Tables: 2

Authors contribution

Concept and design: Federico Augustovski, Andrés Pichon Riviere, Andrea Alcaraz.

Acquisition of data: Carla Colaci, Fernando Argento, Verónica Alfie, Andrea Alcaraz.

Analysis and interpretation Of Data: Carla Colaci, Fernando Argento, Verónica Alfie, Andrea Alcaraz, Federico Augustovski

Drafting of the manuscript: Fernando Argento, Andrés Pichon Riviere, Mackenzie Mills, Panos Kanavos, Andrea Alcaraz

Critical revision of the paper for important intellectual content: Federico Augustovski, Carla Colaci, Verónica Alfie, Fernando Argento, Andrés Pichon Riviere, Mackenzie Mills, Danitza Chavez, Panos Kanavos, Andrea Alcaraz.

Statistical analysis: Fernando Argento, Carla Colaci

Obtaining funding: Andrés Pichon Riviere, Mackenzie Mills, Danitza Chavez, Panos Kanavos, Andrea Alcaraz, Federico Augustovski

Supervision: Federico Augustovski, Andrés Pichon Riviere, Panos Kanavos, Mackenzie Mills

Funding/Support: This project was funded by Precision Cancer Consortium (PCC), a pharma/biotech consortium consisting of AstraZeneca, Bayer, GSK, Janssen, Lilly, Novartis, and Roche. Funding source had no role in the design, conduction, analysis and reporting of this

study. PCC members reviewed a previous version of the manuscript. The final version was independently produced and approved by all authors.

Acknowledgements:

Members of the Project Steering Committee:

Jean Mossman – Patient advocacy (London School of Economics, UK)

Mark Lawler –Chair in Translational Cancer Genomics (Queen’s University Belfast, Northern Ireland)

Michael Drummond - Professor Emeritus (University of York, UK)

Nicola Normanno - Past President of the Italian Cancer Society (Italy); President International Quality Network for Pathology (IQN Path, Luxembourg)

Laura Sampietro-Colom - Head of Innovation Assessment Unit (Health Technology Assessment, Clinic Barcelona University Hospital, Spain)

Albrecht Stenzinger - Professor at Institute of Pathology (University Hospital Heidelberg, Germany)

Benjamin Horbach - Global Policy Leader, Market Access (Roche, representing PCC)

Highlights section:

- Value frameworks play a crucial role in enhancing transparency and facilitating informed decision-making within the healthcare system. They serve as valuable guides, highlighting key dimensions that need evaluation when assessing various healthcare technologies.
- IECS (Institute for Clinical Effectiveness and Health Policy) recently developed a value framework for diagnostic technologies, based on a targeted systematic review and a participatory process with HTA stakeholders in Latin America.

- In this paper we report a systematic review to update the previous framework and develop an comprehensive evidence based list of criteria and subcriteria that could be considered for creation of a NGS/CGP targeted value framework as well as a starting point for value frammeworks targeted to other health technologies

Journal Pre-proof

TITLE**A Systematic Review of Value Criteria for Next-Generation Sequencing/Comprehensive Genomic Profiling to Inform Value Framework Development****Abstract**

Objective: To comprehensively identify and map an exhaustive list of value criteria for the assessment of Next-Generation Sequencing/Comprehensive Genomic Profiling (NGS/CGP), to be used as an aid in decision making.

Methods: We conducted a systematic review to identify existing Value Frameworks (VFs) applicable to any type of healthcare technology. VFs and criteria were mapped to a previously published Latin American -LA- VF (Augustovski et al., VIH 2021) to harmonize definitions and identify additional criteria and or subcriteria. Based on this analysis, we extracted a comprehensive, evidence-based list of criteria to be considered in the design of an NGS/CGP VF.

Results: A total of 42 additional VFs were compared to the LA VF, 88% were developed in high-income countries, 30% targeted genomic testing and 16% specifically targeted oncology. A total of 242 criteria and subcriteria were extracted; 227 (94%) were fully/partially included in the LA VF; 15 (6%) were new. Clinical benefit and economic aspects were the most common criteria. VFs oriented to genomic testing showed greater overlap with the previous VFs. Considering all criteria and subcriteria, a total of 18 criteria and 36 individual sub-criteria were identified.

Conclusion: Our study provides an evidence-based set of criteria and subcriteria for healthcare decision-making useful for NGS/CGP. The resulting list can be beneficial to inform decision making and will serve as a foundation to co-create a multi-stakeholder NGS/CGP VF that is aligned with the needs and values of health systems and could help to improve patient access to high value technologies.

Keywords: Value framework, HTA, Systematic Review, Diagnostic, NGS, CGP

Introduction

Value frameworks, or value assessment frameworks (VF) transparently and explicitly define the individual dimensions and criteria that are important in a decision-making process and often reflect the preferences or values of the different actors involved in their construction and use. In the context of healthcare, they provide an indication of which factors are most important, and by extension less important, in assessing the value of a health intervention. The scope, role, and application of VFs in healthcare has been highly variable. While drugs and therapeutic technologies have been the primary target of VFs, there are numerous examples of "generic" VFs used for the evaluation of a range of different health technologies (such as the Genetic testing Evidence Track Tool [GETT]¹, UK Genetic Testing Network [UKGTN]², ACCE model³ framework among others). Due to the complexity and specificities of different contexts, VFs have been increasingly developed in the last one or two decades. Some VFs have been developed on a worldwide scale, but there are also regional or country-specific VFs, or even frameworks aimed at decision-making in specific health conditions (e.g. oncology) or in specific settings (e.g. patient-centred decisions).^{4,5}

Collaborative and multi-stakeholder frameworks (encompassing regulatory bodies, academia, patient organizations, and pharmaceutical companies) could expedite and standardize data collection for the purpose of evaluating potential benefits, risks, economic value, among other criteria, in the context of health.⁶

Next-Generation Sequencing/Comprehensive Genomic Profiling (NGS/CGP) are high-throughput sequencing technologies that enable rapid analysis of DNA and RNA. These tests allow for broad genomic profiling evaluations, whole exome sequencing (WES) or whole genome sequencing (WGS). NGS/CGP technologies have transformed the way we use genetic data in healthcare by allowing the rapid generation of large amounts of sequence data. This has spurred developments in various domains, including genetics, genomics, transcriptomics, epigenetics, metagenomics, and personalized medicine (PM).⁷ In the latter domain, PM has enabled the detection of somatic driver mutations, allowed resistance mechanisms to be defined, facilitated quantification of mutational burden, and detected germline mutations. In the context of PM, there is a need to build or adapt a value framework for the specificities of NGS/CGP to inform resource allocation decisions using evidence-based principles and values.

Previous studies have already emphasized the difficulty of evaluating diagnostic technologies, such as how to view them in the context of the patient's treatment pathway, or to be able to assess direct benefits at such an early stage within this context. NGS/CGP not only does not escape these difficulties, but new ones are also encountered, such as the real relationship between genetic alterations and diseases, simultaneous diagnosis of multiple diseases or the incidental finding of diseases and their possible impact. This is important, as these technologies are becoming more numerous and increasingly integrated into medical care (e.g. in oncology care pathways). Thus, constructing or adapting a precise value framework for implementing NGS/CGP becomes crucial. To accomplish this objective, resource allocation decisions concerning NGS/CGP must be strongly evidence-based, ensuring they align with the principles of evidence-based medicine. Due to the dynamic nature of the field and the continuing emergence of new diagnostic tests and medicines, together with the infrastructure investment decisions that these technologies often require, there is growing uncertainty associated with the decision-financing point for NGS/CGP technologies. In this context, technology assessments can play an important role in facilitating informed coverage decisions. Taking into account that many generic VFs have been developed for specific diseases/conditions, they may not necessarily address the proper use of NGS/CGP.

In 2020, the Institute for Clinical Effectiveness and Health Policy (IECS) developed a Diagnostic Technology Value Framework aimed broadly at diagnostic test technologies for Latin American (Latam) decision-makers.⁴ The IECS value framework consists of 15 criteria or domains (defined as a set of criteria or attributes that assist in defining the global value of particular health technologies) and a total of 21 sub-criteria or sub-domains (defined as different aspect or components “nested” within a criterion or domain, in cases where a dimension includes several different aspects) for assessing the value of a diagnostic technology. The previous targeted literature review and the specificity of this framework focused on diagnostic tests, as well as the wide variety of criteria and subcriteria- led us to consider this framework as a starting or reference VF. However, the framework addresses all types of diagnostic technologies, so it may not address the particular needs of genetic testing, and particularly NGS/CGP. Appendix Figure 1 shows Latam VF (IECS) criteria and subcriteria. Given the growing use of genomic testing worldwide, this project was undertaken to review and describe an extensive list of criteria and subcriteria for the further development of a new value framework for NGS/CGP technologies for decision-makers. In this diagnostic value framework, the value was mainly focussed on assessing the changes that

can be made to an existing treatment following a more accurate diagnosis. However, a VF for NGS/CGP demands a broader approach. For example, dealing the value of changes in a range of treatments, and also additional value, arising from the treatment possibilities for other potential patients in the future, by identifying the patients with the gene alterations that future treatments might target. Therefore, since NGS/CGP may be associated with more aspects of value it was necessary to approach the generation of a VF in a more comprehensive way. Using this broad approach it may also be relevant to a wider range of health technologies. So, we using the recent IECS framework as a starting VF, we performed a systematic review to define an updated exhaustive list of potential criteria or subcriteria.

The current project was co-led by IECS and the London School of Economics (LSE). In this paper, we present the results of the first stage of the project, which includes a systematic review of value frameworks used for the evaluation of NGS/CGP, a mapping of their criteria and subcriteria, and the development of a comprehensive list of non-overlapping value criteria and subcriteria (non-overlapping in this case is understood in a non-restrictive way, as criteria or subcriteria that could be evaluated individually, even though some of them could be related to a lesser or greater extent.). This proposed list can not only serve as the starting point for the further development of a collaborative NGS/CGP value framework but also provides an important reference for future endeavors to develop value frameworks for these or other health technologies.⁸ While this work provides a list of criteria that could be further contextualized to any jurisdiction, the next step of this project will include, using this exhaustive list as a starting point, to develop a NGS/CGP VF for Europe through a collaborative process that will include several consensus building rounds (i.e Delphi panel) with key stakeholders.

Methods

The methodology employed in this study consisted of three distinct stages. The initial stage encompassed a systematic literature review, during which various VFs were identified. Subsequently, in the second stage, a mapping exercise was conducted to ascertain the alignment of criteria and subcriteria between the identified VFs through the systematic literature review (SLR) and the original Latam VF.

In the last stage, the entire author team collaboratively addressed the discrepancies that arose during the mapping, leading to the final integration and inclusion of the new criteria/sub-criteria in the original IECS VF. A detailed explanation of each step is presented below. The outcome of this study was the development of a comprehensive list of potentially relevant and mutually exhaustive criteria and sub-criteria.

First stage: Systematic review

Contemplating that many domains of interest can be addressed not only by value frameworks targeting genetic testing but also by generic frameworks, our systematic review employed a threefold approach to comprehensively identify available value frameworks, without any language restriction. Firstly, we updated the previous SLR search strategy applied to the original IECS's VF⁴, extending the search from 2018 to October 2022. This step aimed to capture more recent value frameworks associated with diagnostic technologies, encompassing both genetic and non-genetic diagnostics. Secondly, we added an extensive literature search across MEDLINE and EMBASE, also until October 2022. This was a tailored-sensitive search strategy, focusing on value frameworks and genomic tests, specifically Next-Generation Sequencing (NGS) and Comprehensive Genomic Profiling (CGP) technologies. In this component, we did not include an initial publication date. Thirdly, in order to identify any value frameworks possibly overlooked by the previous two components, we performed a specific search using the term value framework without date restriction. More detail of the search strategy can be seen in the Supplementary Material (S1).

To augment our search, we supplemented it by exploring grey literature, with the aim of detecting VFs from key health technology assessment agencies from diverse countries, and scientific societies, and embraced the utilization of general search engines.

Study screening and selection were performed independently by two reviewers from the research team. Disagreements were resolved by consensus of the entire review team. All phases of study selection were performed using COVIDENCE^{®9,10}, a web-based platform designed for the systematic review process. The eligibility criteria were defined to capture both generic and targeted frameworks that could be used to assess NGS/CGP technologies. This included: a) generic value frameworks aimed at health technologies; b) value frameworks aimed more specifically at

genomic tests; c) value frameworks aimed more specifically at diagnostic tests (genomic or otherwise); and d) value frameworks specifically targeting NGS or CGP technologies. Studies were considered if they included at least two value criteria.

Data extraction was performed independently by reviewers from the research team and discrepancies were resolved by consensus of the entire team. In the case of studies with several publications, the main study was considered as the main reference and the evidence was supplemented with data from the grey literature for completeness. For each of the frameworks surveyed, the criteria and sub-criteria that make up the framework, the reference framework, the region, the area of use, the scope, and the perspective (the recommended context of use) were extracted.

Second stage: Mapping

Once data from all included VFs had been extracted, the similarities and differences between these frameworks and the IECS framework were assessed. In order to perform this mapping, we assessed the degree of inclusion of each criterion and subcriteria of all included value frameworks, using a four-category scale: fully included (criteria or subcriteria that were considered to be fully captured by the IECS framework), partially included, not included, and methodological criteria (those that are part of the methodology for conducting an evaluation, but not as a particular final criterion or subcriterion that can be used to assess value - one example can be how to determine the population to be included in the assessment). The mapping was performed by two researchers and any discrepancies were discussed with the research team. Criteria and sub-criteria were considered as separate units and were matched against the IECS value framework independently. In turn, a comparative analysis of the criteria-subcriteria included according to the scope of application of VF, the perspective of VF, and the therapeutic area of VF was performed.

Third stage: Integration of the criteria/subcriteria

Upon completion of the search for value frameworks and the execution of criteria/sub-criteria mapping, the selection process encompassed those elements that exhibited partial overlap or no overlap at all with the criteria/sub-criteria of the IECS VF. This set of criteria/sub-criteria underwent independent evaluation by two distinct working groups—the IECS researchers and the LSE researchers' team—with the aim of discerning which among them should be included in the

value framework designed for the assessment of NGS/CGP. Any discrepancies were collectively resolved through a series of successive meetings involving the entire working team. Finally, those criteria/sub-criteria judged to be potentially relevant were incorporated into the final mutually exclusive list of criteria and subcriteria to be further assessed in a following step of the project by a panel of experts.

Journal Pre-proof

Results

The review of peer-review literature yielded a total of 2067 non-duplicate records, of which 31 articles were identified for full-text review. Following full-text review, 18 articles were excluded: nine for not being value frameworks, three for the wrong outcome, three due to inadequate study design (e.g., systematic reviews, case studies), three for being a value framework out of scope (e.g., frameworks with only one criterion, non-health-oriented framework). A total of 13 studies were included from the grey literature, a further 16 studies were identified, of which, after evaluation, 13 met the inclusion criteria. Two of the grey literature studies surveyed were used to supplement the evidence extracted from two studies of VF in the literature from indexed journals, leading to a final 11 VF from the grey literature. An additional 19 studies were included from the previous literature search of the Augustovski *et al.* study, in which a total of 20 studies were identified.⁴ One of the studies (a methodological manual) was excluded as the team considered that it did not meet the inclusion criteria for the present review. See Figure 1 for more information.

Figure 1. Study flow diagram. PRISMA 2020 flow diagram¹¹

First stage: Systematic review

43 VFs were extracted (including the IECS VF), all of which are listed in Table 1 below (more detailed information on the criteria/sub-criteria can be found in Supplementary Material (S3 Table 1)).

Fifteen were developed in the USA, followed by five developed for all of Europe, four developed worldwide, four in Canada, four in the United Kingdom, three in Australia, two in Germany, and two in Latin America. One framework was developed in Spain, Brazil, and Iran. One framework did not refer to its country/region.

In terms of scope, fourteen (32.5%) of the frameworks were designed to evaluate general diagnostic tests, thirteen (30.2%) frameworks were developed to evaluate genomic tests, twelve (27.9%) were general frameworks, and only four (9.3%) frameworks were designed for the evaluation of medicines.

In terms of therapeutic area, twenty-nine (67.4%) of the frameworks were developed for any health condition, seven (16.3%) for oncology, and two (4.7%) for rare diseases. Only one framework was developed strictly for genetic diseases, independent of any disease area. Two frameworks (4.7%) did not specify the target therapeutic area, while the other two focused on other areas of use.

Thirty-five VF (81.4%) were *de novo* developed frames where no other frameworks were considered as a reference. Only eight VF (18.6%) remarked that they used other frameworks as references. The ACCE, EGAPP, and USPSTF frameworks were referenced most frequently.^{3,12,13}

The perspective of the use of the frameworks was mainly for use in technology assessment agencies, with twenty-four VF developed in this context (55.8%), followed by seven (16.3%) for use in professional societies, five (11.6%) for use in ministries of health, two (4.7%) for hospitals and only one framework was designed for use from a patient perspective. Table 1 shows the total number of studies per category extracted.

Second stage: Mapping

Mapping of criteria and subcriteria was undertaken following completion of data extraction. We used the IECS VF as the reference framework and we mapped the rest of the VFs to find commonalities and differences. When including all criteria and subcriteria from each of the 42 frameworks (excluding the IECS VF), a total of 242 individual criteria and subcriteria were retrieved. Of these, 200 criteria and subcriteria were fully captured within the IECS framework, 8 were partially captured, 15 were new, and 19 were non-final criteria or values and referred to methodological aspects (See Figure 2).

Figure 2. Scheme of the methodology and general results of the mapping (non-overlapping criteria and subcriteria).

When we mapped the forty-two VFs with the IECS VF, they had at a minimum 2 shared criteria and a maximum of 10 shared criteria. The most frequently included criteria within the 43 VF were clinical benefit (n=40) and economic aspects (n=34). Other commonly occurring criteria included non-clinical benefit (n=22), safety (n=19) and organizational impact (n=17). Less frequently included criteria were absence of alternative diagnostic technologies (n=3), environmental impact (n= 2) and priority in the health system (n=1). Table 2 displays the 42 FVs categorized by scope,

denoting their respective application areas and the intended perspectives they were designed for. Each criterion aligning with the IECS FV is marked with a checkmark, while those criteria not aligning are marked with a hyphen.

Sub-analysis on the criteria included within identified VFs was conducted based on VF scope, perspective, and therapeutic area (See Figure 3). In terms of scope, both similarities and differences in the criteria included are present between generic frameworks, diagnostic frameworks, genomic frameworks, and medicines frameworks. Most of the frameworks assessed included clinical benefit and economic aspects criteria, regardless of VF scope. Medicines frameworks had a high frequency of including safety (100%) and disease burden ($\approx 75\%$) relative to other framework scopes (25-30%). However, nonclinical benefits were included less frequently in drug frameworks. Value frameworks with a genomic scope had a higher frequency of including legal and ethical criteria ($\approx 50\%$ compared to $\approx 25\%$ for other scopes) and organizational impact ($\approx 50\%$ compared to $\approx 0\text{-}12\%$ for other scopes).

When examining the perspectives of the VFs, the findings revealed that they were developed for application in Health Technology Assessments (n=23), Ministry of Health (MoH) standpoint (n=4), professional societies (n=7), while a subset (n=8) did not specify their perspective.

There were no major differences between the criteria included across VF perspectives. Organizational impact was the most frequently included criteria within HTA frameworks ($\approx 50\%$ vs $\approx 12.5\text{-}25\%$). Non-clinical benefits were less frequently included in scientific society frameworks compared to the rest $\approx 12\%$ vs. $\approx 50\text{-}60\%$).

In terms of therapeutic area, categories for sub-analysis included frameworks for all health conditions (n= 28), frameworks for oncology (n= 7) and frameworks for others (n= 7). Frameworks designed for a specific therapeutic area that is not oncology were grouped into a single “other” category given small sample size. Inclusion of criteria was generally similar across the three therapeutic area groups. However, the oncology-oriented frameworks generally evaluated fewer criteria (mainly clinical benefit, safety, economic evaluation, and to a lesser degree, disease burden). Frameworks for all health conditions and frameworks for non-oncology therapeutic areas (“others”) included other criteria at a higher frequency, such as organizational impact ($\approx 15\%$ vs $\approx 40\text{-}60\%$) and non-clinical benefits (0% vs $\approx 40\text{-}70\%$).

Figure 3. Analysis of the criteria considered according to different VF categories (scope, perspective, and area)

Third stage: Integration of the criteria/subcriteria

A total of 15 criteria/sub-criteria were identified that did not overlap with any of the criteria/sub-criteria in the original IECS framework (See Supplementary Material S4. Table 2 contains more information about these criteria/subcriteria). Out of these, 4 were deemed irrelevant for this study after reaching consensus with the entire team. Finally, 2 were included as additional criteria ('Public health/population benefit', 'Quality assurance/quality improvement program'), and 7 as sub-criteria ('Technical aspects', 'Fear of contagion', 'Real option value', 'Research priorities', 'Risk of overutilization', 'External pressures', 'Degree of investment in research and development'). On the other hand, out of the 9 criteria/sub-criteria that partially matched the original IECS value framework criteria, after team consensus, it was decided to include 8 as sub-criteria ('Test for disadvantaged or underserved communities', 'Test for rare diseases', 'Equity in health financing', 'Impact on education', 'Impact on career', 'Impact on stigma', 'Effective access to the test and subsequent treatment').

It's important to highlight that after team discussion, a consensus was reached to add the sub-criterion 'Safety and Data Governance', given the significance of patients being owners of their own information. Additionally, a criterion labeled 'Other' was introduced to encompass certain sub-criteria that weren't well-defined under other main criteria but hold importance within this context.

Table 3 shows the final list of non-overlapping and potentially relevant criteria and subcriteria, proposed for further contextualization exercises such as the upcoming next phase of the project consisting of a VF co-creation for Europe with consensus building activities with key stakeholders, or other VF design initiatives. This final list contains 18 criteria and 36 sub-criteria.

Discussion

The systematic review unveiled a total of 42 value frameworks, encompassing a collective sum of 242 individual criteria and subcriteria (when including overlap). Among these, 15 criteria were not encompassed in the original IECS framework. Most of these frameworks were conducted from a health technology assessment (HTA) perspective and targeted all health conditions and technologies. The vast majority of the frameworks adequately addressed both clinical benefit and economic endpoints. Safety, another very common criterion, was not universally addressed in the frameworks evaluated. This may be in part because the search was oriented to value frameworks for diagnostic/genomic tests, technologies in which it may be less important compared to frameworks oriented towards medicines evaluation.

There are inherent tensions and trade-offs among the usefulness, applicability, and precision of different value frameworks. For rapid and general decision-making, generic VF are probably preferred by decision-makers, but they have the downside of being less fit for purpose when targeting more specific technologies such as diagnostic tests, genetic/genomic tests, or particular disease populations such as oncology patients.

As previously stated, the evaluation of diagnostic tests presents inherent complexities compared to other medical technologies. Amongst diagnostic health technologies genomic panel tests, such as Next-Generation Sequencing, introduce additional intricacies. Frequently, there is a significant gap between the utilization of these tests and the subsequent emergence of their derived benefits. This situation makes it intricate to ascertain their genuine advantages.

Despite our review having identified thirteen frameworks specifically designed for the evaluation of genomic profiling, none of them were explicitly tailored to technologies like NGS/CGP, which facilitate the simultaneous assessment of numerous genes.

Within the criteria and subcriteria of genomic testing value frameworks, we found specific criteria that warrant further consideration, such as analytical validity or penetrance. While these criteria could conceivably be taken into account within the patient's clinical pathway, it is pertinent to explicitly contemplate the potential importance of incorporating these types of criteria and subcriteria. However, frameworks like ACCE and EGAPP, some of the earliest frameworks and foundations for subsequent frameworks, fail to address the unique challenges posed by broad

genomic panels such as Whole Exome Sequencing (WES) or Whole Genome Sequencing (WGS).^{12,54} This raises the question of whether specific genomic testing criteria suffice in this context. Another interesting aspect to highlight is that these frameworks pay more attention to ethical and legal aspects relative to other frameworks. One issue that necessitates consideration in these cases pertains to incidental findings. Such findings hold relevance in the context of NGS/CPG, as large-scale genomic testing may reveal genomic alterations associated with diseases unrelated to the specific pathology for which the tests were conducted, thereby having substantial clinical ramifications.^{55,56} In many instances, it becomes necessary to determine what information should be communicated to patients and how it ought to be conveyed. While the incorporation of these specific considerations may not suffice in conferring complete value to these technologies, it nevertheless represents a step in the right direction.

Although in this study we do not consider how each of the criteria of the framework is technically evaluated, and if and how are they weighted in a deliberative or quantitative way, it is important to understand that NGS/CPG may present particular difficulties.⁵⁷ For example, for economic evaluation, the choice of comparator, the time and type of analysis, and the structure of the model may be particularly challenging. Additional challenges are present for the evaluation of clinical benefit in the clinical (PICO) question. Normally, in the evaluation of a diagnostic technology, the relationship of a test to its treatment can be established, while in this case there may be multiple associated therapeutic interventions.⁵⁸

This review serves as an initial step in the development of a value framework specifically adapted to CGP/NGS. Additionally, our study, which provides a unique evidence base set of non overlapping value criteria and subcriteria, can also help to inform future value framework initiatives. Despite the identification of numerous value frameworks, none are expressly designed for wide-range gene panels, and only thirteen of these are designed for general genomic testing. Consequently, our understanding of the specific areas requiring evaluation in the context of NGS/CGP technology remains limited. Nevertheless, we strongly believe that this study serves as a starting point for establishing criteria values for these novel and innovative technologies.

This paper has strengths and limitations. Some limitations that warrant consideration. Notably, the initial process of mapping criteria and sub-criteria of all included VF was undertaken by the research team, drawing from the LATAM value framework. Such a process involved qualitative

research methods and it inherently involves an element of interpretation. Although we followed a rigorous qualitative process and the final decision regarding the inclusion, definition and grouping of criteria and subcriteria involved all the researchers from both institutions, there exists the potential for the omission of criteria that could hold substantial significance for others. Moreover, the act of mapping, inherently tied to a specific framework, can limit the distinctions among the appraised frameworks. For instance, while two distinct frameworks assessing different facets of clinical benefit, they were subsumed under the overarching criteria of clinical benefit, and their individual disparities were not delineated and carried forward. Regarding our study strengths, foremost among these is the comprehensive incorporation of diverse scopes, perspectives, and domains for which the frameworks under study were formulated. This inclusive approach affords a more comprehensive and holistic understanding of the intrinsic values attributed to healthcare technologies. Furthermore, while the principal focus of this study pertains to NGS/CGP technologies, the search strategy extended beyond this domain, thereby encompassing criteria germane to alternative technological realms, including general diagnostic technologies. The resultant synthesis of these broader criteria lends added relevance to the assessment of NGS/CGP technologies.

Conclusion

Our study establishes a robust set of evidence-based criteria and subcriteria to inform healthcare decision-making in the realm of NGS/CGP genomic testing. This set of criteria will form the cornerstone for the multi/stakeholder collaborative development of NGS/CGP Value Framework in Europe, and can also help to inform future value framework initiatives. This holds the potential to drive meaningful advances in patient health outcomes and access and help to address some of the current healthcare system challenges.

Journal Pre-proof

References

1. Rousseau F, Lindsay C, Charland M, et al. Development and description of GETT: a genetic testing evidence tracking tool. *Clin Chem Lab Med.* 2010;48(10):1397-1407. doi:10.1515/CCLM.2010.291
2. UK Genetic Testing Network (UKGTN) new test recommendations. National Health Service (NHS). https://www.engage.england.nhs.uk/consultation/specialised-services-consultation/user_uploads/ukgtncpag-cover-paper.pdf. Published 2002. Accessed January 30, 2023.
3. ACCE Model Process for Evaluating Genetic Tests. Centers for Disease Control and Prevention (CDC). <https://www.cdc.gov/genomics/gtesting/acce/index.htm>. Published December 28, 2010. Accessed December 13, 2023.
4. Augustovski F, Alfie V, Alcaraz A, García Martí S, Drummond MF, Pichon-Riviere A. A Value Framework for the Assessment of Diagnostic Technologies: A Proposal Based on a Targeted Systematic Review and a Multistakeholder Deliberative Process in Latin America. *Value Health.* 2021;24(4):486-496. doi:10.1016/j.jval.2020.11.008
5. Mandelblatt JS, Ramsey SD, Lieu TA, Phelps CE. Evaluating Frameworks That Provide Value Measures for Health Care Interventions. *Value Health.* 2017;20(2):185-192. doi:10.1016/j.jval.2016.11.013
6. Mateo J, Steuten L, Aftimos P, et al. Delivering precision oncology to patients with cancer. *Nat Med.* 2022;28(4):658-665. doi:10.1038/s41591-022-01717-2
7. Kumar KR, Cowley MJ, Davis RL. Next-Generation Sequencing and Emerging Technologies. *Semin Thromb Hemost.* 2019;45(7):661-673. doi:10.1055/s-0039-1688446
8. Europe's Beating Cancer Plan - European Union. https://www.google.com/search?q=Europe%27s+Beating+Cancer+Plan&rlz=1C5CHFA_enES836ES837&oq=Europe%27s+Beating+Cancer+Plan&aqs=chrome..69i57j0i22i3019.225j0j4&sourceid=chrome&ie=UTF-8#:~:text=Europe%27s%20Beating%20Cancer%20Plan%20%2D%20European,europa.eu%20%E2%80%BA%20eu+cancer%2Dplan+en+0.
9. Babineau J. Product Review: Covidence (Systematic Review Software). *J Can Health Libr Assoc.* 2014;35(2):68-71. doi:10.5596/c14-016
10. Covidence systematic review software. Veritas Health Innovation. www.covidence.org.
11. Page MJ, McKenzie JE, Bossuyt PM, et al. The PRISMA 2020 statement: an updated guideline for reporting systematic reviews. *BMJ.* 2021;372. doi:10.1136/bmj.n71
12. Teutsch SM, Bradley LA, Palomaki GE, et al. The Evaluation of Genomic Applications in Practice and Prevention (EGAPP) Initiative: methods of the EGAPP Working Group. *Genet Med.* 2009;11(1):3-14. doi:10.1097/GIM.0b013e318184137c
13. U.S. Preventive Services Task Force. *Guide to Clinical Preventive Services: Methods.*; 2003. <https://play.google.com/store/books/details?id=PtcFxFxKeFcC>.

14. Angelis A, Kanavos P. Multiple Criteria Decision Analysis (MCDA) for evaluating new medicines in Health Technology Assessment and beyond: The Advance Value Framework. *Soc Sci Med.* 2017;188:137-156. doi:10.1016/j.socscimed.2017.06.024
15. Pearson SD. The ICER Value Framework: Integrating Cost Effectiveness and Affordability in the Assessment of Health Care Value. *Value Health.* 2018;21(3):258-265. doi:10.1016/j.jval.2017.12.017
16. Institute for clinical and economic review (ICER) 2020-2023 Value Assessment Framework. https://icer.org/wp-content/uploads/2020/10/ICER_2020_2023_VAF_102220.pdf. Published January 31, 2020. Accessed December 20, 2022.
17. Schnipper LE, Davidson NE, Wollins DS, et al. Updating the American Society of Clinical Oncology Value Framework: Revisions and Reflections in Response to Comments Received. *J Clin Orthod.* 2016;34(24):2925-2934. doi:10.1200/JCO.2016.68.2518
18. Campolina AG, Estevez-Diz MDP, Abe JM, de Soárez PC. Multiple Criteria Decision Analysis (MCDA) for evaluating cancer treatments in hospital-based health technology assessment: The Paraconsistent Value Framework. *PLoS One.* 2022;17(5):e0268584. doi:10.1371/journal.pone.0268584
19. dosReis S, Butler B, Caicedo J, et al. Stakeholder-Engaged Derivation of Patient-Informed Value Elements. *The Patient - Patient-Centered Outcomes Research.* 2020;13(5):611-621. doi:10.1007/s40271-020-00433-8
20. Harris RP, Wilt TJ, Qaseem A, High Value Care Task Force of the American College of Physicians. A value framework for cancer screening: advice for high-value care from the American College of Physicians. *Ann Intern Med.* 2015;162(10):712-717. doi:10.7326/M14-2327
21. Merlin T, Farah C, Schubert C, Mitchell A, Hiller JE, Ryan P. Assessing personalized medicines in Australia: a national framework for reviewing codependent technologies. *Med Decis Making.* 2013;33(3):333-342. doi:10.1177/0272989X12452341
22. Shams L, Yazdani S, Nasiri T. The Value Framework Governing Iran's Health System Policy: A Practical Gap. *Int J Prev Med.* 2022;13:96. doi:10.4103/ijpvm.IJPVM_446_20
23. Rogowski WH, Schleidgen S. Using needs-based frameworks for evaluating new technologies: an application to genetic tests. *Health Policy.* 2015;119(2):147-155. doi:10.1016/j.healthpol.2014.11.006
24. Garrison LP Jr, Neumann PJ, Willke RJ, et al. A Health Economics Approach to US Value Assessment Frameworks-Summary and Recommendations of the ISPOR Special Task Force Report [7]. *Value Health.* 2018;21(2):161-165. doi:10.1016/j.jval.2017.12.009
25. Lakdawalla DN, Doshi JA, Garrison LP Jr, Phelps CE, Basu A, Danzon PM. Defining Elements of Value in Health Care-A Health Economics Approach: An ISPOR Special Task Force Report [3]. *Value Health.* 2018;21(2):131-139. doi:10.1016/j.jval.2017.12.007
26. Pichon-Riviere A, Garcia-Marti S, Oortwijn W, Augustovski F, Sampietro-Colom L. Defining the Value of Health Technologies in Latin America: Developments in Value Frameworks to Inform the Allocation of Healthcare Resources. *Int J Technol Assess Health Care.* 2019;35(1):64-68. doi:10.1017/S0266462319000072

27. Giacomini M, Miller F, Browman G. Confronting the “gray zones” of technology assessment: evaluating genetic testing services for public insurance coverage in Canada. *Int J Technol Assess Health Care*. 2003;19(2):301-316. doi:10.1017/s0266462303000278
28. Guidelines for preparing assessments for the Medical Services Advisory Committee. Medical Services Advisory Committee. Australia Government. Department of Health and Age Care. [http://www.msac.gov.au/internet/msac/publishing.nsf/Content/E0D4E4EDDE91EAC8CA2586E0007AF C75/\\$File/MSAC%20Guidelines-complete-16-FINAL\(18May21\).pdf](http://www.msac.gov.au/internet/msac/publishing.nsf/Content/E0D4E4EDDE91EAC8CA2586E0007AF C75/$File/MSAC%20Guidelines-complete-16-FINAL(18May21).pdf). Published May 2021. Accessed January 3, 2023.
29. Anonychuk A, Beastall G, Shorter S, Kloss-Wolf R, Neumann P. A framework for assessing the value of laboratory diagnostics. *Healthc Manage Forum*. 2012;25(3, Supplement):S4-S11. doi:10.1016/j.hcmf.2012.07.015
30. NICE health technology evaluations: the manual. National Institute for Health Care Excellence (NICE). <https://www.nice.org.uk/process/pmg36/resources/nice-health-technology-evaluations-the-manual-pdf-72286779244741>. Published January 31, 2022.
31. Institut national d'excellence en santé et services sociaux. Assessment mechanism for medical biology tests at INESSS. <https://www.inesss.qc.ca/en/themes/sante/medical-biology-procedures/assessment-mechanism-for-medical-biology-tests-at-inesss.html#c1192>. Accessed January 24, 2023.
32. General Methods. Version 5.0. Institute for Quality and Efficiency in Health Care (IQWiG). https://www.iqwig.de/methoden/general-methods_version-5-0.pdf. Published July 10, 2017. Accessed January 10, 2017.
33. A Framework for Comprehensive Assessment of the Value of Diagnostic Tests. Advanced Medical Technology Association (AdaMed). https://www.advamed.org/wp-content/uploads/2017/05/advameddiagnosticframeworkreport_09.pdf. Published May 2017. Accessed January 12, 2023.
34. Wurcel V, Cicchetti A, Garrison L, et al. The Value of Diagnostic Information in Personalised Healthcare: A Comprehensive Concept to Facilitate Bringing This Technology into Healthcare Systems. *Public Health Genomics*. 2019;22(1-2):8-15. doi:10.1159/000501832
35. Frueh FW, Quinn B. Molecular diagnostics clinical utility strategy: a six-part framework. *Expert Rev Mol Diagn*. 2014;14(7):777-786. doi:10.1586/14737159.2014.933075
36. MolDX® Program (Administered by Palmetto GBA). Palmetto GBA. [https://www.palmettogba.com/palmetto/providers.nsf/Files/Technical_Assessment_Summary_Form_GEN-PF-001.pdf/\\$FILE/Technical_Assessment_Summary_Form_GEN-PF-001.pdf](https://www.palmettogba.com/palmetto/providers.nsf/Files/Technical_Assessment_Summary_Form_GEN-PF-001.pdf/$FILE/Technical_Assessment_Summary_Form_GEN-PF-001.pdf). Accessed January 20, 2023.
37. Mann G, Squire SB, Bissell K, et al. Beyond accuracy: creating a comprehensive evidence base for TB diagnostic tools. *Int J Tuberc Lung Dis*. 2010;14(12):1518-1524. <https://www.ncbi.nlm.nih.gov/pubmed/21144235>.
38. Canestaro WJ, Pritchard DE, Garrison LP, Dubois R, Veenstra DL. Improving the Efficiency and Quality of the Value Assessment Process for Companion Diagnostic Tests: The Companion test Assessment Tool (CAT). *J Manag Care Spec Pharm*. 2015;21(8):700-712. doi:10.18553/jmcp.2015.21.8.700

39. Lee DW, Neumann PJ, Rizzo JA. Understanding the medical and nonmedical value of diagnostic testing. *Value Health*. 2010;13(2):310-314. doi:10.1111/j.1524-4733.2009.00597.x
40. EUnetHTA. HTA Core Model for Diagnostic Technologies v 1.0r Work Package 4. EUnetHTA. <https://www.eunethta.eu/hta-core-model-for-diagnostic-technologies-1-0r/>. Published December 2008. Accessed January 15, 2023.
41. Blancquaert I. Testing for BRCA: the Canadian experience. The Evaluation of Clinical Validity and Clinical Utility of Genetic Tests, Summary of an expert workshop. *Cambridge, UK: PHG Foundation; 2007*. 2007:23-25.
42. Calonge N, Green NS, Rinaldo P, et al. Committee report: Method for evaluating conditions nominated for population-based screening of newborns and children. *Genet Med*. 2010;12(3):153-159. doi:10.1097/GIM.0b013e3181d2af04
43. Severin F, Borry P, Cornel MC, et al. Points to consider for prioritizing clinical genetic testing services: a European consensus process oriented at accountability for reasonableness. *Eur J Hum Genet*. 2015;23(6):729-735. doi:10.1038/ejhg.2014.190
44. Fryback DG, Thornbury JR. The efficacy of diagnostic imaging. *Med Decis Making*. 1991;11(2):88-94. doi:10.1177/0272989X9101100203
45. Annemans L, Aymé S, Le Cam Y, et al. Recommendations from the European Working Group for Value Assessment and Funding Processes in Rare Diseases (ORPH-VAL). *Orphanet J Rare Dis*. 2017;12(1):50. doi:10.1186/s13023-017-0601-9
46. Garrison ZBJMF. *Improving the Health Technology Assessment of Complementary Diagnostics.pdf*. Office of Health Economics; 2016.
47. A Novel Approach to Evaluating the Utility of Genetic Tests: A Coverage Decision Framework. InformedDNA. Genetics, Decoded. https://informeddna.com/wp-content/uploads/2022/03/InformedDNA_Coverage_Decision_Framework_wp.pdf. Published November 2019. Accessed January 13, 2023.
48. Krahn M, Miller F, Bayoumi A, et al. DEVELOPMENT OF THE ONTARIO DECISION FRAMEWORK: A VALUES BASED FRAMEWORK FOR HEALTH TECHNOLOGY ASSESSMENT. *Int J Technol Assess Health Care*. 2018;34(3):290-299. doi:10.1017/S0266462318000235
49. Harris RP, Helfand M, Woolf SH, et al. Current methods of the US Preventive Services Task Force: a review of the process. *Am J Prev Med*. 2001;20(3 Suppl):21-35. doi:10.1016/s0749-3797(01)00261-6
50. Shah-Manek B, Galanto JS, Nguyen H, Ignoffo R. Value Frameworks for the Patient-Provider Interaction: A Comparison of the ASCO Value Framework Versus NCCN Evidence Blocks in Determining Value in Oncology. *J Manag Care Spec Pharm*. 2017;23(6-a Suppl):S13-S20. doi:10.18553/jmcp.2017.23.6-a.s13
51. Wendy Bruening, Eileen Erinoff, Karen M. Schoelles, Fang Sun. *Addressing Challenges in Genetic Test Evaluation Evaluation Frameworks and Assessment of Analytic Validity*. Agency for Healthcare Research and Quality (AHRQ); 2011.

52. de Guzmán. José Antonio Castilla Alcalá Eduardo Briones Pérez Soledad Márquez Calderón de la BACP. *Guide for Decision-Making on the Introduction of New Genetic Tests in the National Health System (GEN Guide)*. Ministry of Health and Consumer Affairs.; 2007.
53. Drug Abacus. Memorial Sloan Katherine Hospital. <https://www.drugpricinglab.org/tools/drug-abacus/>. Accessed January 23, 2023.
54. ACCE Model Process for Evaluating Genetic Tests. <https://www.cdc.gov/genomics/gtesting/acce/index.htm>. Published September 22, 2022. Accessed June 29, 2023.
55. Blackburn HL, Schroeder B, Turner C, Shriver CD, Ellsworth DL, Ellsworth RE. Management of Incidental Findings in the Era of Next-generation Sequencing. *Curr Genomics*. 2015;16(3):159. doi:10.2174/1389202916666150317232930
56. Martinez-Martin N, Magnus D. Privacy and ethical challenges in next-generation sequencing. *Expert review of precision medicine and drug development*. 2019;4(2):95. doi:10.1080/23808993.2019.1599685
57. Phillips KA, Deverka PA, Marshall DA, et al. Methodological Issues in Assessing the Economic Value of Next-Generation Sequencing Tests: Many Challenges and Not Enough Solutions. *Value Health*. 2018;21(9):1033. doi:10.1016/j.jval.2018.06.017
58. Tarride JE, Gould T, Thomas DM. Challenges of Conducting Value Assessment for Comprehensive Genomic Profiling. *Int J Technol Assess Health Care*. 2022;38(1). doi:10.1017/S026646232200040X

Journal Pre-proof

Table 1. Value frameworks identified

Author	Year	Region	Scope	Target area of use	Reference framework	Perspective
Augustovski et al ⁴	2021	Latin America	Diagnostic technology	All health conditions	No	HTA
CDC ³	2000-2004	USA	Diagnostic technology	All health conditions	No	Professionals Society
Angelis et al ¹⁴	2017	UK	General framework	All health conditions	No	HTA
Pearson et al /ICER ^{15,16}	2018/ 2020-2023	USA	General framework	All health conditions	No	HTA
ASCO ¹⁷	2016	USA	Drugs framework	Oncology	No	Professionals Society
Campolinal et al ¹⁸	2022	Brasil	Drugs framework	Oncology	No	Hospital
DosReis et al ¹⁹	2020	NM	General framework	All health conditions	No	Patients
Harris et al ²⁰	2015	USA	Diagnostic technology	Oncology	No	Professionals Society
Merlin et al ²¹	2013	Australia	Genetic test	Oncology	No	MoH
Shams et al ²²	2022	Iran	General framework	All health conditions	No	MoH
Rogowski et al ²³	2015	Germany	Genetic test	Unspecified	No	Other
Garrison et al Lakdawalla et al ^{24,25}	2018	International	General framework	All health conditions	No	HTA
Pichon-Riviere et al ²⁶	2019	Latin America	General framework	All health conditions	No	HTA
Giacomini et al ²⁷	2003	Canada	Genetic test	All health conditions	No	MoH
MSAC. Medical Services Advisory Committee ²⁸	2021	Australia	General framework	All health conditions	USPSTF	HTA
Anonychuck et al ²⁹	2012	Australia	Diagnostic technology	All health conditions	No	Hospital
NICE ³⁰	2022	UK	General framework	All health conditions	No	HTA
Teutsch et al ¹²	2009	USA	Genetic test	All health conditions	No	MoH
INESS ³¹	2022	Canada (Quebec)	Diagnostic technology	All health conditions	No	HTA
IQWIG ³²	2017	Germany	General framework	All health conditions	No	HTA
AdvaMed ³³	2017	USA	Diagnostic technology	All health conditions	No	HTA
Medtech ³⁴	2019	Europe	Diagnostic technology	Other	No	Other
Frueh et al ³⁵	2014	USA	Diagnostic technology	All health conditions	No	HTA
Palmetto GBA ³⁶	2011	USA	Genetic test	All health conditions	No	Other
Mann et al ³⁷	2010	UK	Diagnostic technology	Other	No	Professionals Society
Canestaro et al ³⁸	2015	International	Diagnostic technology	Oncology	No	HTA
Lee et al ³⁹	2010	International	Diagnostic technology	All health conditions	No	HTA
EUnetTHA ⁴⁰	2015	Europe	Diagnostic technology	All health conditions	No	HTA

Blancquaert et al ⁴¹	2007	Canada	Genetic test	All health conditions	ACCE	HTA
Calonge et al ⁴²	2003	USA	Genetic test	Rare diseases	ACCE y EGAPP	Professionals Society
Rousseau et al, GETT (Genetic testing Evidence Tracking Tool) ¹	2010	International	Genetic test	All health conditions	ACCE y UKGTN	HTA
Severin et al (Eurogentest Evaluation Model) ⁴³	2015	Europe	Genetic test	All health conditions	No	Other
UKGTN (The United Kingdom Genetic Testing Network) ²	2002	United Kingdom	Genetic test	All health conditions	ACCE	HTA
Fryback and Thornbury ⁴⁴	1991	USA	Diagnostic technology	All health conditions	No	Professionals Society
Annemans et al ⁴⁵	2017	Europe	Drugs framework	Rare diseases	No	HTA
Garrison et al ⁴⁶	2016	Europe	Diagnostic technology	All health conditions	No	HTA
InformedDNA ⁴⁷	2019	USA	Genetic test	Genetic conditions	EGAPP/USPSTF	HTA
Krahn et al ⁴⁸	2007	Canada	General framework	Unspecified	No	HTA
Harris et al ⁴⁹	2001	USA	Diagnostic technology	All health conditions	No	Professionals Society
Shah-Manek ⁵⁰	2017	USA	General framework	Oncology	No	HTA
AHRQ (Agency for Healthcare Research and Quality) ⁵¹	2011	USA	Genetic test	All health conditions	EGAPP	HTA
Calderón et al ⁵²	2006	Spain	Genetic test	All health conditions	UK Genetic Testing Network	HTA
Memorial Sloan Katherine Hospital ⁵³	NM	USA	Drugs framework	Oncology	No	Other

Abbreviations: *HTA*: Health Technology Agency, *EGAPP*: Evaluation of Genomic Applications in Practice and Prevention, *USPSTF*: U.S Preventive Services Task Force.

Table 2. Characteristics of the frameworks reviewed and their overlap with the IECS framework.

Source	Target area of use	Perspective	Clin	Safe	Qual	Econ	Org	Prior	Burd	Equi	Eth L	Seve	Alter	Non-C	Envi	Soc	Inno	Other
Diagnostic technology																		
Fryback et al ⁴⁴	All conditions*	PS	✓	✓	-	✓	-	-	-	-	-	-	-	✓	-	-	-	-
ACCE ⁵⁴	All conditions	PS	✓	✓	✓	✓	-	-	✓	-	✓	-	-	-	-	✓	-	-
Canestaro et al ³⁸	Oncology	HTA	✓	-	-	✓	-	-	✓	-	-	-	-	-	-	-	-	-
EUnefTHA ⁴⁰	All conditions	HTA	✓	✓	-	✓	✓	-	✓	✓	-	✓	-	✓	✓	-	-	-
Frueh et al ³⁵	All conditions	HTA	✓	-	-	✓	-	-	-	-	-	-	-	-	-	-	-	-
Garrison et al ²⁴	All conditions	HTA	✓	-	-	✓	-	-	-	-	-	-	-	✓	-	-	✓	-
Harris et al ⁴⁹	All conditions	PS	✓	✓	-	-	-	-	-	-	-	-	-	-	-	-	-	-
Harris et al ²⁰	Oncology	PS	✓	-	-	✓	-	-	-	-	-	-	-	-	-	-	-	-
Lee et al ³⁹	All conditions	HTA	✓	-	-	-	-	-	-	-	-	-	-	✓	-	-	-	-
Mann et al ³⁷	Other	PS	✓	-	-	✓	✓	-	-	✓	-	-	-	-	-	-	-	-
Anonychuck et al ²⁹	All conditions	Hospital	✓	✓	-	✓	✓	-	-	-	✓	-	-	✓	✓	✓	-	-
Medtech ³⁴	Other	Other	-	-	-	✓	✓	-	-	-	-	-	-	✓	-	-	-	-
Drugs Framework																		
Annemans et al ⁴⁵	Rare diseases	HTA	✓	✓	✓	✓	✓	-	✓	-	-	✓	-	✓	-	✓	✓	-
ASCO ¹⁷	Oncology	PS	✓	✓	-	✓	-	-	-	-	-	-	-	-	-	-	-	-
Campolinal et al ¹⁸	Oncology	Hospital	✓	✓	-	✓	-	-	✓	-	-	-	-	-	-	-	-	-
MSK ⁵³	Oncology	Other	✓	✓	-	✓	-	-	✓	✓	-	-	✓	-	-	✓	✓	✓
General framework (drugs and other)																		
NICE ³⁰	All conditions	HTA	✓	-	-	✓	-	-	-	-	-	-	-	✓	-	-	-	-
IQWIG ³²	All conditions	HTA	✓	-	-	✓	✓	-	-	-	✓	-	-	✓	-	✓	-	-
AdvaMed ³³	All conditions	HTA	✓	✓	-	✓	✓	-	-	-	-	-	-	✓	-	-	-	-
Krahn et al ⁴⁸	Unspecified	HTA	✓	✓	-	✓	-	-	-	✓	✓	-	-	-	-	-	-	-
Angelis et al ¹⁴	All conditions	HTA	✓	✓	-	✓	-	-	✓	✓	-	-	✓	✓	-	✓	✓	-
MSAC ²⁸	All conditions	HTA	-	-	-	✓	✓	-	-	-	-	-	-	-	-	-	-	-

DosReis et al ¹⁹	All conditions	Patients	✓	✓	-	✓	-	-	✓	-	-	✓	-	✓	-	✓	✓	-
ICER ^{15,16}	All conditions	HTA	✓	-	-	✓	-	-	-	-	-	-	-	✓	-	✓	✓	-
Lakdawalla et al ²⁵	All conditions	HTA	✓	-	-	✓	-	-	-	✓	-	✓	-	✓	-	✓	✓	✓
NCCN ⁵⁰	Oncology	HTA	✓	✓	✓	-	✓	-	-	-	-	-	-	-	-	-	-	-
Pichon-Riviere et al ²⁶	All conditions	HTA	✓	✓	✓	✓	✓	✓	✓	✓	-	✓	-	-	-	-	✓	-
Shams et al ²²	All conditions	MoH	✓	-	-	✓	-	-	-	✓	-	-	-	✓	-	-	-	✓
Teutsch et al ¹²	All conditions	MoH	✓	-	✓	-	-	-	-	-	-	-	-	-	-	-	-	-
Genetic test																		
AHRQ ⁴⁹	All conditions	HTA	✓	✓	-	-	✓	-	-	-	✓	-	-	-	-	-	-	-
Blancquaert ⁴¹	All conditions	HTA	✓	-	-	-	✓	-	-	-	-	-	-	✓	-	-	-	-
INESS ³¹	All conditions	HTA	✓	-	-	✓	✓	-	-	-	✓	-	-	✓	-	-	-	-
Calderón et al ⁵²	All conditions	HTA	✓	✓	-	✓	-	-	-	✓	✓	-	-	-	-	-	-	-
Calonge et al ⁴²	Rare diseases	PS	✓	-	-	-	-	-	-	-	-	-	-	-	-	✓	-	-
Eurogentest [...] ⁴³	All conditions	Other	✓	-	-	✓	-	-	✓	-	-	-	-	✓	-	-	-	-
GETT ¹	All conditions	HTA	✓	-	-	✓	✓	-	✓	-	✓	-	-	✓	-	-	-	✓
Giacomini et al ²⁷	All conditions	MoH	✓	-	-	✓	✓	-	-	-	-	-	-	✓	-	✓	-	-
InformedDNA ⁴⁷	Genetic	HTA	✓	✓	-	✓	✓	-	✓	✓	✓	✓	-	✓	-	-	-	✓
Merlin et al ²¹	Oncology	MoH	✓	-	-	✓	-	-	-	-	-	-	-	-	-	-	-	-
Palmetto GBA ³⁶	All conditions	Other	✓	✓	-	✓	-	-	-	-	✓	-	-	✓	-	-	-	✓
Rogowski et al ²³	Unspecified	Other	✓	-	-	✓	-	-	✓	-	-	✓	-	-	-	-	-	-
UKGTN ²	All conditions	HTA	✓	-	-	-	✓	-	✓	-	✓	✓	✓	-	-	-	-	-

References: **PS:** Profesional Society, **HTA:** Health Technology Agency, **Clin:** Clinical benefits and test performance; **Safe:** Safety and unwanted consequences; **Qual:** Quality of scientific evidence; **Econ:** Economical aspects; **Org:** Organizational aspects and feasibility within the clinical path; **Prior:** Health priority of the health system; **Burd:** Disease burden; **Equi:** Equity; **EthL:** Ethical and legal aspect; **Seve:** Severity of the disease; **Alter:** Absence of alternative diagnostic technologies; **Non-C:** non clinical benefits; **Envi:** Environmental impact; **Soc:** broader social impact; **Inno:** Innovation.

Note: *All conditions refer to "All health conditions"

Journal Pre-proof

Table 3. Final list of value criteria/sub-criteria and their definitions.

Criteria	Definition	Sub-criteria	Definition
Clinical Benefit and Test Performance	Clinical benefits for people undergoing the test (better health or improvement in clinical outcomes derived from the use of this technology considering the perspective of the individual taking the test) and test performance	Clinical consequences of test use	Clinical consequences during the diagnostic or therapeutic process of carrying out the test (e.g., a change in the therapeutic approach that is associated to an improvement in the state of health becomes apparent from the test result)
		Test performance	Evaluates the diagnostic yield of the test through indicators such as sensitivity, specificity, precision, and reproducibility
		Technical aspects	Evaluates the stability and storage form of the reagents used to perform the tests.
Safety and Unintended consequences	Related to the adverse or unintended effects described for the test being assessed	Procedure safety	Unintended consequences in whom the test is being performed secondary to the performance of the test (e.g., injury to neighboring organs during biopsy)
		Consequences of wrong diagnosis	Unintended consequences in whom the test is being performed secondary to the misdiagnosis (false positives and false negatives)
		Safety of test preparation	Unintended consequences in whom the test is being performed secondary to the preparation for it (e.g., adverse events during bowel preparation for biopsy colonoscopy)
		Safety for test operators	Unintended consequences in test operators (e.g. radiation exposure from taking tomography guided biopsies)
		Risk of overutilization	Refers to the risk of overuse, or abuse, of genomic testing. (e.g: using the test in an individual, or a population, for reasons not specific to the patient's indication).
Quality of scientific evidence	How reliable the scientific evidence to be evaluated, as well as its results is; also the potential that different biases or systematic errors would not allow drawing valid conclusions.		
Economic aspects	Assess how good the health investment in this diagnostic test (cost-effectiveness) is; what its financial or budget impact are, or other aspects such as impact on the out-of-pocket expenditure of the patient and his family	Economic evaluation (Clinical effectiveness and/or Budget Impact Analysis)	Comparison between costs and health results of two or more diagnostic options. Budget impact for the funder when incorporating the test (e.g., includes acquisition costs, maintenance, supplies, training)
		Other costs	Patient and family out-of-pocket expenses, costs related to productivity loss, etc

Organizational aspects and Feasibility within the clinical path	Related to the necessary actions taken by the organization of the system to implement the technology being evaluated so that it can reach the target population, also the feasibility of its implementation. It takes into account which place the test being evaluated occupies in the clinical path.	Impact on the health service provision system	Implementing the test requires modifications of buildings, processes, logistics, etc. within the organization providing health services
		Impact on the patient care path	Implementing the test would be associated with less time in access to the benefit, additional studies, associated practices would be avoided, or the availability of resources would be increased
Health system priorities	Priority of this health problem (for the country or health system, defined by those who design health policies)	Health priority within the health system	Priority of this health problem (for the country or health system, defined by those who design health policies).
		Research priorities	Priority for this technology and this disease in the current or future research agenda.
Disease burden	How important the loss of health, both in mortality and in quality of life is. This includes taking into account the pattern of inheritance, genomic heterogeneity, mutation prevalence, mutation penetrance and neomutation rate.		
Equity	What consequences the implementation of the technology being evaluated would have on equity or inequalities in the system and health	Test for neglected diseases	Test oriented to the diagnosis of neglected diseases
		Test for communicable diseases and high prevalence	Test oriented to the diagnosis of communicable diseases and/or of high prevalence in the region (e.g. HPV/cervical cancer)
		Test in populations with little access to health services	Test considered situations where there is poor access to health services
		Test for disadvantaged or underserved communities	Test oriented to the diagnosis of disadvantaged or underserved communities defined as those relevant communities that have been historically disadvantaged through discrimination, neglect, reduced research funding, or other factors.
		Test for rare diseases	Test oriented to the diagnosis of rare neoplastic diseases. Defined as a disease that affects a small number of people compared to the general population. In Europe, the European Medicines Agency (EMA) considers a disease with a prevalence of less than five in 10,000 people (equivalent to less than one in 2,000) to be rare.
		Equity in health financing	Equity in health financing by promoting health, especially in disadvantaged areas, and helping to meet catastrophic health expenditures

Ethical and legal aspects It considers relevant the social and moral norms and values that derive from the technology in question. It implies an understanding of the consequences of implementing or not implementing a sanitary technology in two aspects: with respect to the values that prevail in the society and with respect to the norms and values that the same technology constructs when it is put into use

Severity of the disease It takes into account the risk of mortality, the risk of disability (and its severity), the quality of life, and the duration of type of cancer.

Absence of alternative diagnostic technologies There is no diagnostic technology available for that type of cancer, or stage, etc.

Non-clinical benefits	Non-clinical benefits (other benefits) for the people who take the test and other participants, as caregivers, family members, etc. It considers other benefits related to the use of the technology, which improve the experience of the test for those who undergo it and also of other participants, such as family members, caregivers, etc. comfort during preparation implementation practice	Experience of who takes the test	Experience of who takes the test / caregivers (comfort, invasiveness, preparation)
		Value of the information	Value of the information provided by the test in special situations (e.g., end-of-life diseases, diseases with poor prognosis, diseases that affect offspring)
		Load on caregivers or family	The test is associated with a lower burden on caregivers or the family of whom the test is performed (e.g., the result of the test results in a lower number of subsequent controls or avoids other unnecessary tests or associated procedures)
		Preparation and/or care	Pre-preparation and / or care after the test (characteristics, need for completion)
		Number of results associated with the test	Number of results associated with the sample (amount of information provided by the test with the sample obtained)
		Test processing time	Sample processing time (suitable for the disease / target population)
		Self test	Self-test (whoever takes the test can do it himself or another person without the need for more training)
		Fear of contagion	The early and correct diagnosis of an infectious diseases limits the spread the disease to others (e.g. HPV/cervical cancer).

	Impact on education	The impact of the diagnosis and treatment on one's education/schooling
	Impact on career	The impact of the diagnosis and treatment on one's career
	Impact on stigma	The impact on the person or the family or society of the diagnosis or treatment, generating embarrassment, self-consciousness, rejection by family or rejection by society.
Environmental Impact	It is a measure that the production, use or implementation of technology would cause in the environment. Eg, technology is associated with a greater generation of toxic waste	
Broader Social Impact	Impact on other sectors beyond health, such as job creation, industrial promotion, technology transfer, and society as a whole.	
Innovation	The diagnostic test being evaluated uses new mechanisms or technologies that were not previously available or is a new test not known so far.	
Quality assurance/ quality improvement program	<p>Internal: What are the controls (positive or negative, normal or abnormal) and what are their origin (samples banking, cell lines)? Specify internal controls related to the analysis carried out and their limits. Are internal quality control procedures applied to the laboratory as a whole or specific to the test? Provide the rates of errors reported. Provide the type of standards (e.g., molecular weight markers) included in the analysis.</p> <p>External: Identify the external quality control programs for this test and specify the type of programs to which you participate. What does this program cover (e.g., analytical aspects of the test or interpretation and reporting)? If no external QC program exists, which type of control do you make (e.g., blind tests, exchanges of various samples between laboratories).</p>	

Public health/ population benefit The public health/ population that this technology will bring when used for this health problem (e.g: detection of susceptible populations to improve their follow-up, screening uses)

Others	Bridge to other future treatments ("Real option value")	When a health technology extends life, creates opportunities for the patient to benefit from other future advances in medicine.
	External pressures	This driver assesses external pressures for test coverage, imposed by providers, members of patient societies, society at large, laboratories and politicians for accelerating unwarranted adoption of tests before solid evidence exists.
	Degree of investment in research and development	The number of human subjects enrolled in the approval trials for the first indication, was used as a proxy for the research and development costs necessary to develop the drug.
	Safety and data governance	This indicator refers to the extent to which test results and their associated patient data are protected from unauthorized access, use, loss, or corruption.
	Effective access to the test and subsequent treatment	Real life effective coverage/access to the test and eventual treatment

Note: In black colour, the original criteria/sub-criteria of the IECS value framework are mentioned. In yellow are highlighted the criteria/sub-criteria that emerged from the systematic review, mapping; and integration as described in the methods and results section.

Figure 1. Study flow diagram. PRISMA 2020 flow diagram

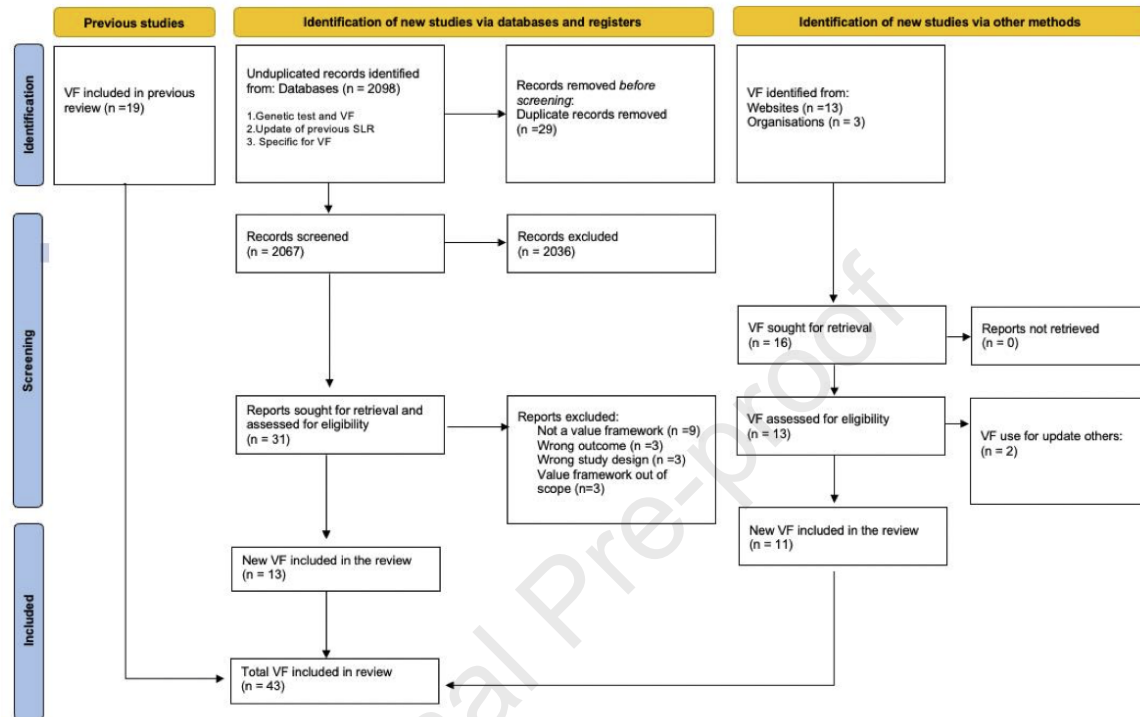


Figure 2. Scheme of the methodology and general results of the mapping (non-overlapping criteria and subcriteria).

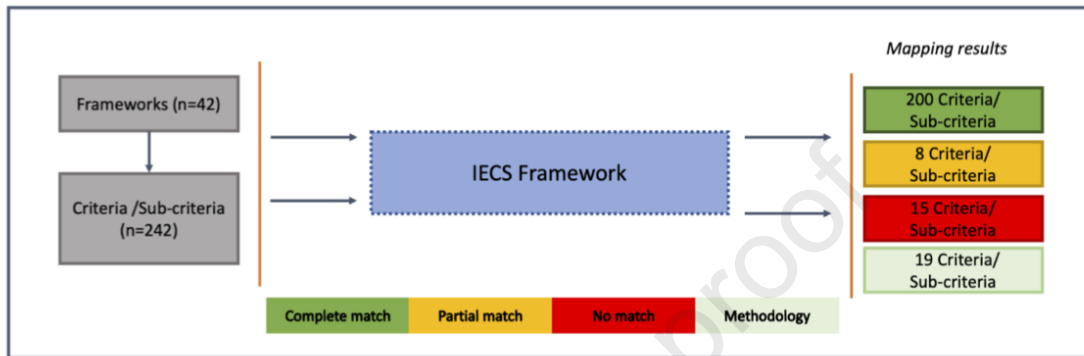
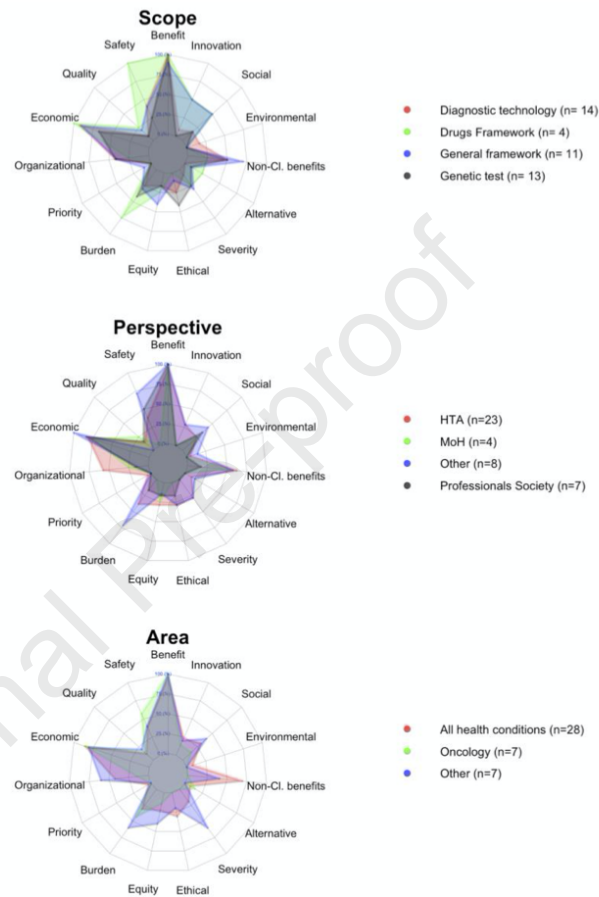


Figure 3. Analysis of the criteria considered according to different VF categories (scope, perspective, and area)



*Abbreviations: **Benefit:** clinical benefits and test performance; **Safety:** Safety and unwanted consequences; **Quality:** Quality of scientific evidence; **Economics:** Economical aspects; **Organizational:** Organizational aspects and feasibility within the clinical path; **Priority:** Health priority of the health system; **Burden:** Disease burden; **Ethical:** Ethical and legal aspect; **Severity:** Severity of the disease; **Alternative:** Absence of alternative diagnostic technologies; **Non-CL benefits:** non clinical benefits; **Environmental:** Environmental impact; **Social:** broader social impact; **HTA:** Health technology agency; **MoH:** Ministry of Health.*