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SYSTEMATIC REVIEW



## Recommendations for economic evaluations of cell and gene therapies: a systematic literature review with critical appraisal

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### ABSTRACT

**Objective:** No consensus exists on the ideal methodology to evaluate the economic impact and value of new, potentially curative gene therapies. We aimed to identify and describe published methodologic recommendations for the economic evaluation of gene therapies and assess whether these recommendations have been applied in published evaluations.

**Methods:** This study was conducted in three stages: a systematic literature review of methodologic recommendations for economic evaluation of gene therapies; an assessment of the appropriateness of recommendations; and a review to assess the degree to which the recommendations were applied in published evaluations.

**Results:** A total of 2,888 references were screened, 83 articles were reviewed to assess eligibility, and 20 papers were included. Fifty recommendations were identified, and 21 reached consensus thresholds. Most evaluations were based on naive treatment comparisons and did not apply consensus recommendations. Innovative payment mechanisms for gene therapies were rarely considered. The only widely applied recommendations related to modeling choices and methods.

**Conclusions:** Methodological recommendations for economic evaluations of gene therapies are generally not being followed. Assessing the applicability and impact of the recommendations from this study may facilitate the implementation of consensus recommendations in future evaluations.

### ARTICLE HISTORY

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### KEYWORDS

Cost-effectiveness analysis; economic evaluation; gene therapy; health technology assessments; health-related quality of life; onasemnogene abeparvovec; spinal muscular atrophy; systematic literature review

## 1. Introduction

Gene therapies function via several mechanisms, such as replacing a disease-causing gene with a healthy copy, inactivating a disease-causing gene, or introducing a new or modified gene to treat a disease [1]. Gene therapies are potentially life-changing for a diverse range of diseases, such as neuromuscular diseases, inherited blindness, metabolic disorders, and hematologic malignancies [2]. Because of technical limitations, medical ethics, and regulatory hurdles, very few approved gene therapies are available for treatment. However, more gene therapies are expected to be approved as technology advances and clinical trials progress [3,4]. An estimated >1 million patients will be treated with gene therapies by the year 2034, leading to an estimated global cost of > \$300 billion [5]. The innovative treatment paradigm and clinical benefits associated with the expected launch of additional gene therapies in the coming years may be met with reimbursement and funding challenges because of the need for health care payment structures to balance greater upfront costs with undetermined long-term clinical safety and effectiveness [6–9].

The use of randomized controlled clinical trials is often unfeasible for gene therapies [5]. Therefore, most clinical studies supporting the market authorization of gene therapies are small, open-label, and single-arm trials [1,5,10]. In health technology assessments (HTAs), limited clinical evidence and greater upfront treatment costs for gene therapies have challenged reimbursement, and specific decision-making considerations have been necessary [10–12]. Substantial challenges remain in how HTAs will appraise the relative degrees of effectiveness, safety, and value-for-money for gene therapies vs. non-gene therapies, based on less comprehensive evidence [10,11].

Several national health institutions and HTA bodies in Europe and North America, including the National Institute for Health and Care Excellence (NICE), the National Institute for Health Research, and the Institute for Clinical and Economic Review in the United States, have assessed the methodologic questions related to the economic evaluation of advanced therapeutic medical products (ATMPs) – including gene therapies – and have provided varying potential

### Article highlights

- Currently, no consensus exists on the ideal methodology to evaluate the economic impact and benefits of new, potentially curative gene therapies, so novel approaches for economic evaluations of gene therapies are needed.
- Fifty methodologic recommendations for the economic evaluation of gene therapies were identified in the published literature, summarized, and critically appraised, with 21 recommendations reaching consensus thresholds and then assessed for applicability and impact in the published economic evaluations.
- Our study found that most evaluations were based on naive treatment comparisons and did not apply consensus recommendations, innovative payment mechanisms for gene therapies were rarely considered, and the only widely applied recommendations related to modeling choices and methods.
- Although analysts conducting economic evaluations of gene therapies have access to many publications that provide methodologic recommendations and guidelines, these published recommendations are generally not being followed.
- Assessing the applicability and impact of the recommendations from this study may facilitate the implementation of consensus recommendations in future economic evaluations of gene therapies to assist with reimbursement and funding challenges for existing health-care payment structures.

recommendations. These included additional scenario analyses to explore long-term benefits, threshold analyses to identify treatment effectiveness, cure proportion modeling in case a percentage of patients is likely to be cured, and reporting of net health benefits in addition to incremental cost-effectiveness (CE) ratios (ICERs) with a measurement of uncertainty [13–17]. Experts from academia and industry have also addressed methodologic questions related to the evaluation of ATMPs and specifically gene therapies. These experts suggest that a completely new reference case is not needed for gene therapies, but some aspects of economic evaluation

should be considered further, because of the unique aspects of gene therapies [18–21].

### 1.1. Aim

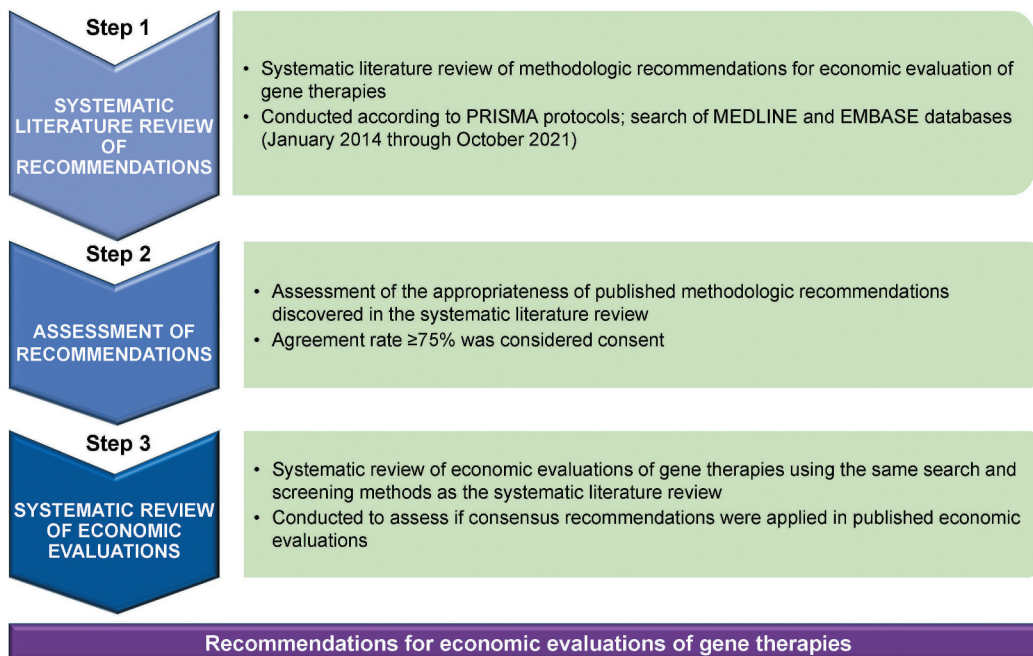
The CE of gene therapies is important to assess because of greater upfront costs, but this assessment is challenging because of the limited relative effectiveness data and the uncertainty around long-term outcomes [10,11]. No consensus currently exists on the ideal methodology to evaluate the economic impact and benefits of new, potentially curative gene therapies, along with reimbursement and funding challenges for existing health care payment structures. Novel approaches for economic evaluations of gene therapies are therefore needed. We aimed to identify and describe the most widely accepted methodologic recommendations for the economic evaluation of gene therapies and assess whether these recommendations were applied in published evaluations.

## 2. Methods

We conducted this study in three stages (Figure 1): a systematic literature review of methodologic recommendations for the economic evaluation of gene therapies; an assessment of the appropriateness of the recommendations; and a review to assess if the consensus recommendations were applied in published evaluations.

### 2.1. Systematic literature review of recommendations

For stage 1, we completed a systematic literature review of recommendations for economic evaluations of gene therapies according to a Preferred Reporting Items for Systematic



**Figure 1.** Study design PRISMA, Preferred Reporting Items for Systematic Reviews and Meta-Analyses. We conducted a systematic literature review of methodologic recommendations for economic evaluation of gene therapies, an assessment of the appropriateness of the recommendations, and a review to assess if consensus recommendations were applied in published economic evaluations.

**Table 1.** Search strategy: key words.

Key word	Key word search
Gene therapy	Gene therapy or innovative medicine or replacement therapy or regenerative medicine or advanced therapeutic medicinal product or ATMP or curative therapy or life-extending or potential cure or curative treatment or curative medicine Gene adjunctive therapy Genetic adjunctive therapy DNA adjunctive therapy Somatic adjunctive gene therapy Somatic adjunctive genetic therapy Gametic adjunctive gene therapy 4D-125 IVT injection or 4D-310 or AAV-CNGB3 or AAV-CNGA3 or AAV directed hLDLR gene therapy or AAVRPE65 or AAV1 or AAV2 or 5-hPDE6B or AAV2 or 5-RPGR or AAV2/8.TBG.hARSB or AAV2-hCHM or AAV2-hRPE65v2, voretigene neparovec-rzyl or AAV2-REP1 or AAV5-hFIX or AAV5-hFIXco-Padua or AAV8-RPGR or AAV9-GLB1 or AAV-mediated REP1 gene replacement or AAVrh.10UCUCLN2 or ABO-102 or Ad2/HIF-1 $\uparrow$ $\pm$ /VP16 or Ad5 or Ad5.hAC6 or Ad5CMV-p53 gene or Ad5-yCD or mutTKSR39rep-ADP or ADA gene transfer or AdCD40L or adeno-associated viral with human factor IX or Ad-p53 or ADV or ADV-tk or ADV/HSV-tk or ADVEGFXC1 or ADVM-022 or ADVM-043 or ADV-Tk or AdvVEGE-D or AGTC-402 or Aldesleukin or Alipogene Tiparovec or Allovectin-7 or ALVAC-MART-1 vaccine or Antigen-specific T Cells CART or Anti-MAGE-A3-DP4 TCR PBL or Anti-NY ESO-1 mTCR PBL or Anti-NY-ESO-1 T-cell receptor PBL or AProArt or ARU-1801 or ASKBio009 or AT132 or AT342 or AT845 or AT-GTX-501 or AT-GTX-502 or autologous anti-MART-1-F5 T-cell receptor or Autologous CD34 positive cells transduced with a lentiviral vector containing human WAS gene or autologous CD34+ cell transduced with G2SCID vector or autologous CD34+ cells genetically modified or autologous CD34+ cells transduced with a lentiviral vector containing the human SGSB gene or Autologous CD34+ cells transduced with WASP lentiviral vector or autologous hematopoietic stem cell transplantation or autologous hematopoietic stem cells genetically modified with GLOBE lentiviral
Economic evaluations	Cost-effectiveness analyses or cost-utility analyses or cost-benefit analyses or economic evaluations or economic evaluation or pharmacoconomics or health economic or cost-effectiveness or cost effectiveness or cost-utility or cost utility or cost-benefit or cost benefit or health technology assessment

Note: AAV, adeno-associated virus; ATMP, advanced therapeutic medical product; hLDLR, human low-density lipoprotein receptor.

Reviews and Meta-Analyses (PRISMA) protocol. The review was based on a literature search performed in MEDLINE and EMBASE databases, covering the period from January 2014 through October 2021. Search key words were related to economic evaluations of gene therapies (Table 1).

Publications were selected if the primary objective was to review, list, discuss, or provide recommendations or solutions for challenges related to economic evaluations of ATMPs (gene therapies and regenerative medicines). Evaluations had to consider both costs and health outcomes to be included, and one of the evaluated treatment strategies had to be a gene therapy or chimeric antigen receptor T-cell (CAR-T) therapy. The following were excluded: animal studies, articles addressing genetic tests, genotyping and whole-genome sequencing interventions, clinical trials, cost-minimization analyses, cost-of-illness or disease burden studies, conference abstracts, publications not offering economic evaluation methodologic recommendations, and articles not available in English. No restriction was placed on geographic scope.

The review of recommendations was completed by a search for relevant white papers published by HTA agencies (NICE, Scottish Medicines Consortium, Haute Autorité de Santé, Canadian Agency for Drugs and Technologies in Health, US Institute for Clinical and Economic Review, and Institute for Quality and Efficiency in Health Care) and a manual cross-referencing search through published literature reviews on the topic.

Abstracts and full texts were independently screened by two reviewers. After a first selection of references based on title and abstract, full texts were screened again by two different analysts. Detailed information, including product class, intervention of investigation, target diseases, population, region, sponsor, use of surrogate endpoints, and time frame, was extracted from selected articles and summarized. Disagreements in the screening, extraction, and summary process were resolved by discussions between analysts and a senior researcher. We listed recommendations in a

comprehensive manner, without judging relevance, and then classified the recommendations by theme.

## 2.2. Assessment of recommendations

For stage 2, we evaluated the appropriateness of published methodologic recommendations discovered in the systematic literature review and selected which recommendations that we, in our collective judgment, believed to be worthwhile and valuable. Our assessment was based on our relevant health economic experience from academia or HTA-related organization membership(s) in Europe and the United States along with our experience related to economic evaluations of gene therapies.

In reviewing the recommendations, we (the eight authors of this paper) indicated if each proposed recommendation was relevant for the economic evaluation of gene therapy ('Agree,' 'Neutral,' or 'Disagree') and whether publications should explicitly report how corresponding issues were addressed. An agreement rate of  $\geq 75\%$  ( $n = 6/8$ ) was determined to identify a consensus recommendation. We grouped the final list of consensus recommendations into five categories: input data (recommendations addressing limitations of clinical data, such as small patient numbers and single-arm trials); modeling choices (methods, considering in particular the uncertainty around long-term effects of gene therapies); health-related quality of life (HRQOL; measurement and evaluation of outcomes, including the challenges related to the pediatric population); estimation of costs; and evaluation framework.

## 2.3. Systematic review of economic evaluations of gene and cell therapies

For stage 3, we conducted a separate review of economic evaluations of ATMPs in order to assess the degree of concordance by analysts to our consensus methodologic



recommendations. The systematic review of economic evaluations was conducted using the same key words and methods used in the prior review.

Detailed information was extracted from selected articles, including product class (e.g. gene therapy or CAR-T); intervention of investigation (generic name); target disease (classified based on the International Classification of Diseases, 11th edition); population (e.g. children or adults); region of study; study sponsor (e.g. private or public); utilization of surrogate endpoints; clinical trial design; use of observational data; indirect treatment comparison (ITC) methods; time frame; discount rates; perspective; utility elicitation methods; types of scenario analyses; types of sensitivity analyses; details of innovative payment mechanisms; and use of real-world evidence post-launch.

Data extraction was conducted by one analyst and checked for completeness and accuracy by a second analyst. A senior analyst was consulted in any case of discrepancy. The quality of reporting was assessed using the Drummond checklist for assessing economic evaluations for gene therapies [22]. This checklist was developed to clarify the extent to which various factors, including clinical effectiveness, elements of value (value to caregivers and insurance carriers, and improvement in life expectancy), and other influences (such as discounts/alternative payment methods), are identified and considered in an economic evaluation [22].

### 3. Results

#### 3.1. Systematic literature review of recommendations

An initial search retrieved 4,302 records. Of these, 2,888 records remained for title and abstract screening after duplicates were removed (Figure 2). Based on the inclusion/exclusion criteria, 2,805 articles were excluded by the reviewers, and after the first selection of references based on title and abstract, 85 full-text articles were excluded. We reviewed the full texts of the remaining 83 articles to assess eligibility, and 20 papers (18 methodologic publications and two white papers from U.S. and Canadian HTA

agencies) were included in the descriptive synthesis (Table 2) [14,18,19,21,23–38].

#### 3.2. Assessment of recommendations

We identified 50 recommendations from the systematic literature review and grouped these recommendations into five categories for review and assessment (Table 3). After the review, 21 consensus recommendations were identified per the agreement rate of  $\geq 75\%$  ( $n = 6/8$ ).

##### 3.2.1. Input data

Seventeen of the identified input data recommendations were associated with limitations to clinical data for gene therapies, including surrogate endpoint validations, use of nonrandomized trial data, methods for estimating relative efficacy and safety, and utilization of expert opinion to obtain information and make a probabilistic representation in the absence of data.

There is scarcity of long-term observed data for gene therapies, particularly at the time of initial regulatory approval or reimbursement consideration [14,21,24]. There was agreement ( $n = 7/8$ ) with using surrogate endpoint data to assess clinical efficacy; however, only half ( $n = 4/8$ ) reported that evidence of a correlation between treatment effects on surrogate endpoints and final endpoints should be presented.

Several publications documented that the rationale for conducting noncomparative studies (single-arm trials) should be clearly elucidated [14,21–24,32] and others recommended the use of other nonrandomized data to provide complementary information to a single-arm trial to allow for an estimation of relative effectiveness [19,24,25,31]. We unanimously agreed ( $n = 8/8$ ) that the rationale behind conducting noncomparative studies should be clearly provided and with the use of observational data to serve as a control arm.

Some publications recommended an assessment of the feasibility of conducting indirect comparisons using network meta-analysis or other statistical approaches when direct comparison is not possible [14,24]. We supported the use of

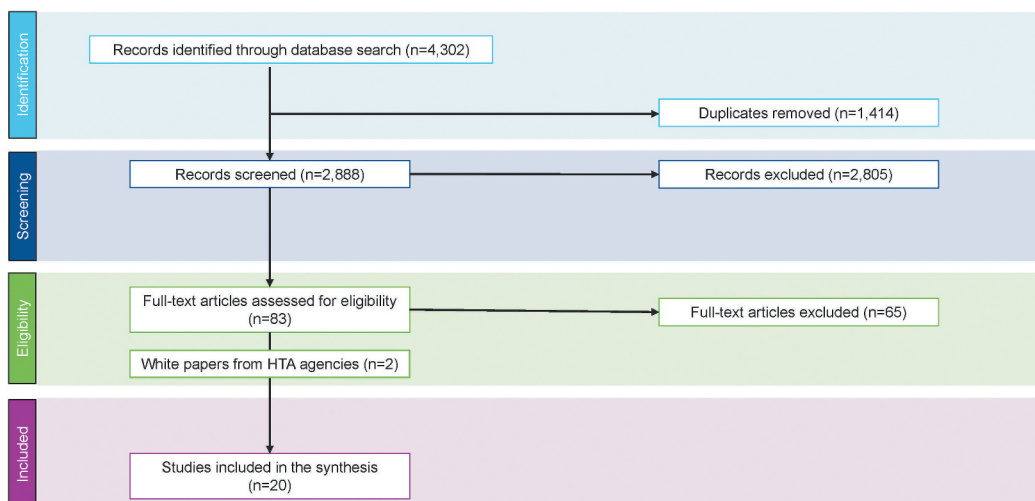


Figure 2. PRISMA Diagram for the Search on Methodologic Recommendations HTA, health technology assessment; PRISMA, Preferred Reporting Items for Systematic Reviews and Meta-Analyses.

**Table 2.** Publications identified in the systematic literature review.

Publication	Funding	Region	Type of Considered Gene Therapy	Recommendation
Ten Ham, 2020 [23]	None	Unknown	<i>In-vivo</i> and <i>ex-vivo</i> gene therapy	Require new reference case
Coyle, 2020 [24]	Private: Novartis/AveXis	Germany	Cell and gene therapy	Require new reference case
Angelis, 2020 [25]	None	United Kingdom	Advanced therapy medicinal products, for cell and gene therapies	Require new reference case
Annemans, 2020 [26]	Public: INAMI/RIZIV	Belgium	Specialized treatments (orphan medical product)	Other information
Gonçalves, 2020 [27]	Private: CTI Clinical Trial & Consulting	Unknown	Advanced therapy medicinal products	Other information
Garrison, 2019 [28]	Private: Novartis/AveXis	United States	One-time gene replacement therapies	Require adaptation/considerations
Petrou, 2019 [29]	None	Cyprus	CAR-T cell therapy	Other information
Gavan, 2019 [30]	None	United Kingdom, United States	Genomic-based diagnostic and gene therapies	Require adaptation/considerations
Raymakers, 2019 [31]	None	Canada	CAR-T therapy	Require adaptation/considerations
Jönsson, 2019 [32]	Private: Gilead/Kite Pharma	Europe	Advanced therapy medicinal products	Require adaptation/considerations
Drummond, 2019 [21]	Private: Novartis/AveXis	United Kingdom, United States, France	Gene therapy	Require adaptation/considerations
Hampson, 2018 [33]	Public: ICER	United States	Gene therapy	Require adaptation/considerations
Hettle, 2017 [14]	Public: NIHR	United Kingdom	Regenerative medicines and cell therapy products	Require adaptation/considerations
Aballéa, 2020 [19]	Private: Novartis/AveXis	France	Gene replacement therapy	Require new reference case
Chapman, 2021 [34]	NA	United States	Single or short-term therapies with potential cures	NA
Gonçalves, 2022 [35]	NA	Portugal	Advanced therapy medicinal products	NA
Jørgensen, 2018 [36]	NA	United Kingdom	Gene therapy	NA
Pochopień, 2021 [18]	NA	France	Gene therapy	NA
CADTH, 2018 [37]	Public: CADTH	Canada	Gene therapy	Require adaptation/considerations
Marsden, 2017 [38]	Public: ICER	United States	Regenerative medicine	Require adaptation/considerations

Note: CADTH, Canadian Agency for Drugs and Technologies in Health; CAR-T, chimeric antigen receptor T-cell; ICER, Institute for Clinical and Economic Review; INAMI-RIZIV, Belgian National Institute for Health and Disability Insurance; NA, not available; NIHR, National Institute for Health and Care Research.

network meta-analysis when feasible in these cases ( $n = 6/8$ ). In addition, we agreed that matching-adjusted indirect comparisons (MAIC) ( $n = 6/8$ ) and propensity score matching (PSM) ( $n = 7/8$ ) may be considered when multivariate and network meta-analysis are not feasible.

### 3.2.2. Modeling choices

We identified 12 recommendations around modeling choices and methods, covering time horizon and extrapolation, scenario analyses, parametric sensitivity analysis, value of information, and discount rate.

Five publications recommended reporting analyses over different time frames and/or to consider different curative time frames or variance in treatment waning or to use a threshold analysis to determine the duration of beneficial effect that would be needed to achieve standard CE thresholds [19,23,25,32,38]. Extrapolation approaches, such as cure proportion modeling, were suggested as the standard reference case for gene therapies whenever relevant, whereas survival analysis was suggested to address uncertainty based on other modeling approaches [19,25]. We agreed ( $n = 7/8$ ) that scenario analyses with different time frames or efficacy waning parameters should be performed, but we did not reach agreement consensus regarding reporting a threshold analysis on the minimum duration of effect

required to reach CE thresholds or the use of cure proportion modeling.

Several publications highlighted the importance of conducting and reporting both deterministic and probabilistic sensitivity analyses in the case of gene therapies [14,19,21,23,31,32,38]. We agreed with the importance of these analyses ( $n = 6/8$ ). We also agreed ( $n = 6/8$ ) with the Institute for Clinical and Economic Review's recommendation to conduct optimistic and conservative scenarios on treatment benefits (e.g. duration of benefit, magnitude/quality of benefit, proportion that achieve a specific benefit, different types of survival models, and relative treatment benefit under alternative assumptions), with the selection of assumptions and inputs used in the optimistic and conservative scenarios being described and justified [38].

Three publications supported lesser discount rates for health outcomes than for costs [19,24,32], five recommended retaining standard reference case discount rates for health outcomes and costs (usually equal) in base-case analysis and to conduct sensitivity analyses with different discount rates for benefits and costs [14,21,23,25,38], and one stated that variable discount rates over time would be more appropriate than a uniform and constant discount rate [9]. We agreed ( $n = 7/8$ ) with applying the reference

Table 3. Assessment of identified recommendations for the economic evaluations of gene therapies.

Subject	Description	Number of Respondents			Number of Missing responses
		Agree	Neutral	Disagree	
<b>INPUT DATA</b>		7	0	1	0
<b>Surrogate endpoints validation</b>	<p>Because of the lack of long-term observed clinical benefits for gene therapies, surrogate endpoint data could be used to assess clinical efficacy</p> <p>If agreed with previous recommendation: evidence on surrogate outcomes validation should be presented based on an explicit hierarchy approach, including:</p> <ul style="list-style-type: none"> <li>Evidence of biological credibility of relationship between surrogate and final outcomes</li> <li>Evidence of the correlation between surrogate endpoint and final endpoint across individuals</li> <li>Evidence of correlation between treatment effects on surrogate endpoint and on final endpoint, based on other clinical studies in the same area</li> </ul> <p>The rationale behind conducting noncomparative studies (single-arm trials) should be clearly provided</p> <p>The inclusion of other nonrandomized data to provide complementary information to a single-arm trial should be considered</p> <p>If agreed with previous recommendation: evidence to justify the inclusion of potential other nonrandomized data into the analysis should include discussions on the following points:</p> <ul style="list-style-type: none"> <li>Preliminary data suggest that the magnitude of treatment benefits vs. historical cohort data is dramatic</li> <li>Primary endpoint is objective, durable, and reproducible</li> <li>Heterogeneity in patient population and study outcomes impacts are explored</li> <li>Confounding factors are well-known and controlled with suitable statistical adjustment methods</li> <li>Generalizability and transferability of the clinical data toward the historical cohort are proactively assessed</li> <li>Other(s)</li> </ul>	6	1	0	1
<b>Use of nonrandomized data</b>	<p>The rationale behind conducting noncomparative studies (single-arm trials) should be clearly provided</p> <p>The inclusion of other nonrandomized data to provide complementary information to a single-arm trial should be considered</p> <p>If agreed with previous recommendation: evidence to justify the inclusion of potential other nonrandomized data into the analysis should include discussions on the following points:</p> <ul style="list-style-type: none"> <li>Preliminary data suggest that the magnitude of treatment benefits vs. historical cohort data is dramatic</li> <li>Primary endpoint is objective, durable, and reproducible</li> <li>Heterogeneity in patient population and study outcomes impacts are explored</li> <li>Confounding factors are well-known and controlled with suitable statistical adjustment methods</li> <li>Generalizability and transferability of the clinical data toward the historical cohort are proactively assessed</li> <li>Other(s)</li> </ul>	3	4	1	0
<b>Relative effectiveness estimation</b>	<p>When direct comparison is not possible, the feasibility of conducting an indirect comparison using multivariate and network meta-analyses should be assessed</p> <p>When multivariate and network meta-analyses are not feasible, other statistical approaches can be considered, such as:</p> <ul style="list-style-type: none"> <li>Matching-adjusted indirect comparisons</li> <li>Simulated treatment comparison</li> <li>Meta-regression</li> <li>Propensity score matching</li> <li>Other(s)</li> </ul>	4	1	0	0
<b>Expert opinion elicitation</b>	<p>Because of limited data available in the context of gene therapies, structured expert opinion elicitation to obtain information and make a probabilistic representation of their knowledge should be used in absence of empirical data for some inputs</p> <p>Analyses over different time frames should be reported, considering different curative time frames or variation in treatment waning</p>	7	1	0	0
<b>MODELING</b>		2	6	0	0
<b>Time horizon and extrapolation</b>	<p>Cure proportion modeling (including mixture and nonmixture cure models) should be considered as the standard reference case for 'single and short-term therapies' whenever relevant, while survival analysis based on other modeling approaches should be used to address uncertainty when feasible</p> <p>Threshold analyses may be run to determine the duration of beneficial effect that would be needed to achieve standard cost-effectiveness thresholds</p>	3	4	0	1
<b>Scenario analyses</b>	<p>Capping economic surplus at the time threshold for loss of exclusivity could be used (e.g. using a 12-year cutoff scenario)</p> <p>Scenario analyses to measure the impact of model assumptions, methodologic choices, and data sources should be conducted, as well as more optimistic and conservative scenarios on treatment benefits</p> <p>Both deterministic and probabilistic sensitivity analyses should be routinely conducted and reported</p>	0	7	1	0
<b>Parametric sensitivity analysis</b>	<p>VOI analysis could be employed to assess uncertainty and inform further evidence collection</p> <p>No change to reference case discounting should be applied to both health outcomes and costs. Sensitivity analysis including the use of varying discount rates for benefits and costs (e.g. 0%–5%) to explore the magnitude of impacts of discount rates should be conducted</p>	6	2	0	0
<b>VOI</b>		6	2	0	0
<b>Discount rate</b>	<p>If disagreed with previous recommendation:</p> <p>Variable discount rates over time are more appropriate than applying a uniform and constant discount rate for both benefits and costs in the reference case</p> <p>Discount rates on health effects should generally be lower than the discount rate applied to costs in the reference case</p>	1	5	2	0
		7	0	1	0
		1	0	0	7
		1	0	0	7

(Continued)

Table 3. (Continued).

Subject	Description	Number of Respondents			Number of Missing responses	
		Agree	Neutral	Disagree		
HRQOL	<b>HRQOL in children</b> In absence of a validated instrument to value HRQOL in very young children, alternative approaches, such as vignette studies (using direct elicitation from proxies), must be used (if treatment is targeting very young children)	6	2	0	0	
ESTIMATION OF COSTS	<b>HRQOL in caregivers Perspective</b> HRQOL in caregivers and families in both health care payer and societal perspectives should be considered	7	0	1	0	
	<b>Other costs</b> Cost-effectiveness analysis should be conducted in two reference case analyses, from both health care payer and societal perspectives Other costs and resources considered should include: Infrastructure cost on the health care system Patient expenses for traveling to specialized medical centers	8	0	0	0	
EVALUATION FRAMEWORK	<b>Innovative payment mechanism</b> Innovative payment mechanisms, to facilitate access to gene therapies while safeguarding the sustainability of health care budgets, should be considered in the analysis	4	4	0	0	
	<b>Attributing economic surplus</b> A 'shared surplus' scenario could be considered with sharing the economic surplus with the health sector by apportioning only a part (e.g. 50%) of cost offsets or a part of cost offsets and QALY gains to the value of the therapy If the economic surplus is perceived to be unreasonably high primarily because of existing extremely expensive care, a scenario capping cost offsets at a defined threshold (e.g. \$150,000 per year in the United States) could be considered	0	5	3	0	
EVALUATION FRAMEWORK	<b>Additional elements of value</b> Specific elements of value to consider for gene therapies may include: Severity of disease (e.g. weighting QALYs according to disease severity) Scientific spillovers Value of caregivers Insurance value/peace of mind/reduction in uncertainty Value of hope Value of cure Fear of contagion Reduction in inequity	0	8	0	0	
	To include broader elements of value into decision analysis, alternative approaches to traditional cost-utility analysis could be considered, such as: Multiple criteria decision analysis Cost-benefit analysis	4	0	0	4	
	Because of the challenges related to the valuation of benefits associated with life-changing treatment for very young children, alternative approaches to standard cost-utility analyses may be considered, including: Cost-benefit analysis Valuation of benefits in terms of saved young life equivalents Given a small target population, and specific elements of value that gene therapy could provide, greater cost-effectiveness thresholds could be considered	2	2	1	3	
	In the case of economic evaluations conducted after launch, real-world evidence should be used to confirm the treatment benefits and bridge the evidence gaps in the initial evaluation	5	0	1	2	
		3	1	2	2	
		2	2	2	2	
		3	2	1	2	
		1	3	1	3	
		2	3	0	3	
	<b>Valuation of health benefits for very young children</b>		0	6	2	0
			2	2	1	3
	<b>WTP threshold</b>		5	2	1	0
<b>Reevaluation</b>		2	6	0	0	
		1	7	0	0	
		8	0	0	0	

Note: HRQOL, health-related quality of life; QALYs, quality-adjusted life-years; VOI, value of information; WTP, willingness-to-pay.



case discount rates in the base-case analysis for gene therapies and conducting sensitivity analyses with different rates.

### 3.2.3. Health-related quality of life

Thirteen recommendations identified concerned the measurement and valuation of outcomes. These included recommendations related to HRQOL for children and caregivers.

In the absence of validated instruments to assess and value HRQOL for children, alternative approaches, such as vignette studies, are recommended [19]. We supported this recommendation ( $n = 6/8$ ).

Two publications highlighted the importance of considering the impact of gene therapies on the HRQOL of caregivers and families, irrespective of whether costs are assessed from a health care payer or societal perspective [25,27]. We agreed ( $n = 7/8$ ) that HRQOL of caregivers and patient families should be considered in economic evaluations of gene therapies.

### 3.2.4. Estimation of costs

Publications addressed several aspects of the estimation of costs: perspective, additional costs not usually considered in economic evaluations, innovative payment mechanisms for gene therapies, and cost offsets.

Many publications recommended conducting economic evaluations from both health care payer and societal perspectives [19,21,23–25,27,30,33]. We unanimously agreed ( $n = 8/8$ ) with this recommendation. Some costs not usually considered in economic evaluations could be substantial for gene therapies, particularly those related to infrastructure changes and travel to distant, specialized facilities where gene therapies may be delivered [32]. Although half ( $n = 4/8$ ) agreed with the suggestion to account for other costs, consensus was not reached.

Several publications recommended exploring the impact of innovative payment mechanisms in CE analyses [14,21,23–25,35–37]. We all agreed ( $n = 8/8$ ) with the recommendation to consider innovative payment mechanisms to facilitate access to gene therapies while recognizing the sustainability of health care budgets.

### 3.2.5. Evaluation framework

Publications assessed general recommendations related to the framework of evaluation and decision-making, including elements of value to consider beyond QALYs, analytical frameworks in which those elements may be considered, whether a greater CE threshold is relevant for gene therapies, and collection of real-world evidence after launch.

Elements of importance not normally captured in QALYs for gene therapies according to reviewed publications were severity of disease, scientific spillovers [28], insurance value, value of hope, value of cure, fear of contagion, and reduction in inequity [27,28,32]. We unanimously agreed ( $n = 8/8$ ) that severity of disease should be considered, and we also agreed ( $n = 7/8$ ) that value of caregivers should be considered. None of the other elements reached consensus. Four publications discussed whether the cost per QALY model could be adapted to account for the elements of value cited above or if the cost-utility analysis (CUA) framework needed to be changed more fundamentally (i.

e. using SAVEs instead of QALYs or multiple criteria decision analysis [MCDA] or cost-benefit analysis [CBA] instead of CUA) [20,24,28,32]. There was very little support for such approaches.

Four publications proposed the use of a greater CE ratio for gene therapies [18,21,24,35], with one proposing the establishment of explicit budget impact thresholds to highlight access challenges and to trigger negotiation with manufacturers [25]. Only one out of eight supported a greater threshold for gene therapies, while the others ( $n = 7/8$ ) were neutral.

A few publications posited that post-launch real-world evidence collection is critical to confirm the treatment benefits and fill the evidence gaps from the initial regulatory submission [19,21,24,27]. We all agreed ( $n = 8/8$ ) that, after the launch of a gene therapy, real-world evidence is important to confirm the benefits of treatment and to provide further evidence on other elements of value.

## 3.3. Systematic review of economic evaluations of gene and cell therapies

One hundred and sixty references were selected for full-text review, and 126 articles were excluded by the reviewers based on the exclusionary criteria. Three articles were added based on a manual cross-referencing search through published literature reviews on the topic. A total of 37 publications were included after we screened the titles and abstracts (Table 4) [36,39–74]. These 37 economic evaluations investigated 10 different marketed gene therapies. Other publications covered adeno-associated virus (AAV)-mediated gene therapies (in three studies) and hypothetical cell or gene therapies (in two studies). CAR-T cell therapies were assessed in 16 studies, and gene therapies other than CAR-T cell therapies were assessed in 21 economic evaluations. These interventions were assessed in 12 different pathologies. Fifteen studies included adult patients only, 11 studies included pediatric patients only, and 11 studies included both children and adults. Most studies were conducted in the United States ( $n = 22$ ) and the United Kingdom ( $n = 8$ ). Half of the economic evaluations ( $n = 19$ ) were funded by private companies and 15 were funded by public and private organizations.

Criteria of the Drummond checklist [22] were applied in reviewed economic evaluations. However, there were methodologic weaknesses related to the identification of relevant costs and lack of or limited sensitivity analysis.

### 3.3.1. Review of input data

Four consensus recommendations regarding limitations of clinical data were retained: justifying the validity of surrogate endpoints, at least with a biologic argument; justifying the utilization of single-arm trial designs when applicable; using observational data for patients not receiving gene therapy as a control arm in the absence of comparative clinical trials; and methods for ITCs. Most published economic evaluations reviewed did not adhere to the consensus recommendations, with only one study providing information on validation of surrogate endpoints [56].

Of the publications reporting the use of single-arm clinical trials, only five justified their use [47,49,63,65,74]. Justifications covered practical and ethical reasons, low incidence/rarity of



Table 4. Characteristics of the studies in the publications included for the review of economic evaluations.

Publication	Class (CAR-T/Other Gene Therapy)	Intervention	Pathology	Population	Region	Sponsor
Bolous, 2021 [39]	Other gene therapy	AAV-mediated gene therapy	Hemophilia	Adults	United States	Public
Machin, 2018 [40]	Other gene therapy	AAV-mediated gene therapy	Hemophilia	Adults	United States	Public
Halioua-Haubold, 2019 [41]	Other gene therapy	AAV-mediated gene therapy	Retinal dystrophy	Adults	United States	Public
Salcedo, 2021 [42]	Other gene therapy	Hypothetical cell or gene therapy	Sickle cell disease	Children + adults	United States	Public
Fleeman, 2017 [43]	Other gene therapy	Talimogene laherparepvec	Melanoma and nonvisceral disease	Adults	United States	Public
Almutairi, 2019 [44]	Other gene therapy	Talimogene laherparepvec + ipilimumab	Melanoma	Adults	United States	Private
Cher, 2020 [45]	CAR-T	Tisagenlecleucel	Large B-cell lymphoma	Adults	Singapore	None
Thielen, 2020 [46]	CAR-T	Tisagenlecleucel	Acute lymphoblastic leukemia	Children	Netherlands	Private
Moradi-Lakeh, 2021 [47]	CAR-T	Tisagenlecleucel	Acute lymphoblastic leukemia + Diffuse large B-cell lymphoma	Children + young adults	Switzerland	Private
Wakase, 2021 [48]	CAR-T	Tisagenlecleucel	Large B-cell lymphoma	Adults	Japan	Private
Qi, 2021 [49]	CAR-T	Tisagenlecleucel	Large B-cell lymphoma	Adults	United States	Private
Lin, 2018 [50]	CAR-T	Tisagenlecleucel	Acute lymphoblastic leukemia	Children	United States	Private
Furzer, 2020 [51]	CAR-T	Tisagenlecleucel	Acute lymphoblastic leukemia	Children	Canada	Public
Sarkar, 2019 [52]	CAR-T	Tisagenlecleucel	Acute lymphoblastic leukemia	Children	United Kingdom	Public
Walton, 2019 [53]	CAR-T	Tisagenlecleucel	Acute lymphoblastic leukemia	Children + young adults	United Kingdom	Public
Whittington, 2018 [54]	CAR-T	Tisagenlecleucel	Acute lymphoblastic leukemia	Children + young adults	United States	Public
Ribera Santasusana, 2020 [55]	CAR-T	Tisagenlecleucel	Acute lymphoblastic leukemia	Children + young adults	Spain	Private
Farmer, 2020 [56]	Other gene therapy	Voretigene neparovvec	Retinal dystrophy	Children + adults	United Kingdom	Public
Uhrmann, 2020 [57]	Other gene therapy	Voretigene neparovvec	Retinal dystrophy	Children + adults	Germany	None
Vinoto, 2020 [58]	Other gene therapy	Voretigene neparovvec	Retinal dystrophy	Children + adults	United Kingdom	Private
Johnson, 2019 [59]	Other gene therapy	Voretigene neparovvec	Retinal dystrophy	Children + adults	United States	Private
Zimmermann, 2019 [60]	Other gene therapy	Voretigene neparovvec	Retinal dystrophy	Children + adults	United States	Public
Jørgensen, 2018 [36]	Other gene therapy	Hypothetical novel GT	Parkinson's disease	Adults	United States/United Kingdom	Private
Cook, 2020 [61]	Other gene therapy	Valoctogene roxaparovvec	Hemophilia	Adults	United States	Private
Zuluaga-Sanchez, 2019 [62]	Other gene therapy	Nusinersen	Spinal muscular atrophy	Children	Sweden	Private
South, 2019 [63]	Other gene therapy	Retrovirally transduced autologous CD34+ cells	Adenosine deaminase deficiency – severe combined immunodeficiency	Children	United Kingdom	Public
Simons, 2021 [64]	CAR-T	KTE-X19	Mantle cell lymphoma	Adults	United States	Private
Liu, 2021 [65]	CAR-T	Axicabtagene ciloleucel	Large B-cell lymphoma	Adults	United States	Private
Whittington, 2019 [66]	CAR-T	Axicabtagene ciloleucel	Non-Hodgkin's Lymphoma	Adults	United States	Public
Roth, 2018 [67]	CAR-T	Axicabtagene ciloleucel	Melanoma	Adults	United States	Private
Lin, 2019 [68]	CAR-T	Axicabtagene ciloleucel/tisagenlecleucel	Large B-cell lymphoma	Adults	United States	Private
Dean, 2021 [69]	Other gene therapy	Onasemnogene abeparovvec	Spinal muscular atrophy	Children	United States	Private
Malone, 2019 [70]	Other gene therapy	Onasemnogene abeparovvec	Spinal muscular atrophy	Children	United States	Private
Malone, 2019 [70]	Other gene therapy	Onasemnogene abeparovvec	Spinal muscular atrophy	Children	United States/United Kingdom	Private
Broekhoff, 2021 [71]	Other gene therapy	Onasemnogene abeparovvec	Spinal muscular atrophy	Children	Netherlands	Public
Connock, 2020 [72]	Other gene therapy	Onasemnogene abeparovvec	Spinal muscular atrophy	Children	United Kingdom	None
Shih, 2021 [73]	Other gene therapy	Onasemnogene abeparovvec/nusinersen + newborn screening	Spinal muscular atrophy	Children	Australia	Public
Kansal, 2021 [74]	Other gene therapy	Betibeglogene autotemcel	Transfusion-dependent β-thalassemia	Children + adults	United States	Private

Note: AAV, adeno-associated virus.

the disease, and lack of confounding variables without details provided. Approximately 70% of studies ( $n = 26/37$ ) used other nonrandomized data, such as natural history studies or registries. Some studies provided precise comparison information between populations, and others provided complementary information on the rationale and robustness of the data without providing additional details.

Where comparators were not included in pivotal clinical trials, most of the studies ( $n = 26/32$ ) employed naive comparisons. Two studies used MAIC [47,65] and one used a scenario analysis [48]. Two studies used published network meta-analyses [67,72].

### 3.3.2. Review of modeling choices

Consensus recommendations related to modeling choices and methods included conducting analyses over different time frames or with different durations of treatment effect; using the reference case discount rates and conducting sensitivity analyses with different rates for costs and outcomes; conducting deterministic and probabilistic sensitivity analyses on model parameters; and reporting optimistic and pessimistic scenarios related to treatment benefits. Our systematic review of economic evaluations found that the degree of adherence to these recommendations was generally positive, except for reporting optimistic and pessimistic scenarios related to treatment benefits.

Most studies (86% [ $n = 32/37$ ]) used a lifetime time horizon in the base-case scenario. Different time frames and/or assumptions on treatment effects over time were reported in 18 economic evaluations as sensitivity analyses. The time frame or duration of effect often had a greater impact on the results.

A majority of the economic evaluations (78% [ $n = 29/37$ ]) used standard discount rates, identical for both health outcomes and costs, and ran sensitivity analyses to vary discount rates. The choice of the discount rate was identified as significantly affecting the results [48,71].

Twenty-six of 37 publications included both deterministic and probabilistic sensitivity analyses. For the 11 remaining publications, four provided a deterministic sensitivity analysis only [41,55,57,63], two provided a probabilistic sensitivity analysis only [53,54], and the others did not conduct sensitivity analyses [36,56,61,66,72].

Only five of the 37 publications (14%) reported pessimistic/optimistic scenarios on treatment benefits [50,52,67,69,70]. These two extreme scenarios were reported in addition to the standard sensitivity analyses. ICER values varied widely between optimistic and pessimistic scenarios.

### 3.3.3. Review of health-related quality of life

Consensus recommendations related to the measurement and valuation of outcomes included the use of alternative approaches, such as vignette studies to obtain utility values when no valid generic instruments (such as EuroQol-5D [EQ-5D], a standard measure of clinical and economic HRQOL via surveys) exist for the targeted population and the need to account for caregivers' and families' HRQOL when impacted by the patient's disease. Our systematic review of economic

evaluations determined that the degree of adherence to these consensus recommendations was poor.

Of the 14 evaluations including children aged 5 years or younger, five evaluations used utility values elicited from vignette studies [58,59,62,73,74]. Indications covered retinal dystrophy, SMA, and TDT. Other evaluations including very young children considered utility values based on EQ-5D (youth version).

Six studies considered HRQOL for caregivers and families, including only four of 14 studies of pediatric populations [46,50,58,62,69,73]. The impact of HRQOL inclusion on the results of economic evaluations for caregivers depends on the gene therapy and assessed indications, and differences also can be seen within the same indication.

### 3.3.4. Review of estimation of costs

Two recommendations for estimating costs reached consensus: conducting analyses from both health care payer and societal perspectives and exploring the impact of innovative payment mechanisms on incremental costs. Twenty-four studies considered a payer perspective [36,39,40,42,44,45,47–51,54,55,58,61,64–70,72,74], and three studies considered the societal perspective [57,71,73]. Four studies conducted both health care payer and societal analyses [46,52,60,62].

Discussions and analyses considering innovative/alternative payment mechanisms were reported in approximately 20% of the economic evaluations ( $n = 7/37$ ) [36,50,52,54,55,69,74]. Several forms of performance-based payments were considered, such as assuming payment for treatment acquisition for responders at 1 month, payment triggered by a remission duration reaching a given threshold, and payment only for initial complete response. Thus, payment mechanisms varied according to the nature and duration of treatment effects. The impact of innovative payment mechanisms on ICERs varied substantially between studies.

Many published evaluations were conducted pre-launch, and no real-world evidence was available. Therefore, only three economic evaluations considered real-world evidence (two included sensitivity analyses), using real-world data on adverse events and health care resource utilization [48,49,51]. The use of estimates of treatment effects based on real-world evidence often led to a reduction of the ICER. We reviewed several economic evaluations that discussed the importance of collecting real-world evidence.

## 4. Discussion

New gene therapies will be reaching the market in upcoming years, and the cumulative budget impact of these therapies is expected to be substantial [5]. Health economic evaluations will play an important role in pricing and reimbursement decisions related to gene therapies. However, economic evaluations of gene therapies raise many methodologic challenges, which have been discussed in the literature [1–12,14]. We found many recent publications that provided methodologic recommendations for economic evaluations of gene therapies or, more broadly, ATMPs [14,18,19,21,23–38]. In the current study, we summarized these recommendations,

critically appraised the recommendations, and then assessed their applicability and impact in published economic evaluations of gene therapies. We found that, although analysts conducting economic evaluations of gene therapies have access to many publications that provide methodologic recommendations and guidelines, most of these recommendations were generally not followed.

The recommendations originated from academia, HTA agencies, and industry. Some consensus for these recommendations was observed between these publications, including conducting analyses from both health care payer and societal perspectives; considering the impact of innovative payment mechanisms on CE; collecting real-world evidence and updating evaluations after launch; conducting analyses over different time frames; and reviewing the evidence supporting the validation of surrogate endpoints. Several other recommendations appeared in one or two publications only. However, areas of disagreement were not observed between publications, except for divergent views on discount rates.

Some recommendations aimed to resolve important issues associated with economic evaluations of gene therapies, but would require paradigmatic changes in evaluation methodology, and are therefore unlikely to be implemented (e.g. the use of CBA or MCDA instead of CUA or the use of SAVEs instead of QALYs).

Recommendations that would provide more information to decision-makers and improve transparency without changing results of the reference case (e.g. presenting evidence on validation of surrogate endpoints; justifying the use of single-arm studies; providing scenario analyses; discussing elements of value that are not captured in QALYs, such as scientific spillovers, insurance value, or reduction in inequities) achieved consensus, and some of these recommendations were frequently implemented in reviewed economic evaluations.

The importance of providing some justification of the validity of surrogate endpoints reached consensus, but not on the exact criteria of validation. With one exception, reviewed studies did not provide any evidence of validation of surrogate endpoints. This is not surprising because a published review [75] reported that the pivotal trial evidence supporting marketing approvals for products going through expedited approval pathways were often based on non-validated surrogate endpoints.

Published economic evaluations often involved comparisons between a gene therapy and standard of care, using a single-arm trial to inform health outcomes with the gene therapy and an observational study to inform outcomes of standard of care. There was relative consensus that investigators using such comparisons should be able to justify the objective and reproducible nature of the endpoints, assess the consequences of heterogeneity in patient population and study outcomes, and control for confounding factors. However, justifications for using such comparisons in reviewed economic evaluations were missing in a majority of publications or limited to comments about the comparability between populations.

Indirect treatment comparisons are an important area of possible improvement for future economic evaluations of gene therapies [76]. In the absence of head-to-head studies, it is generally recommended to perform ITCs using network meta-analyses when feasible, which requires randomized

controlled trials. When only single-arm studies are available, ITCs may be performed using MAIC or PSM, if individual patient data are available for the comparator [76]. There was consensus about these recommendations.

Another recurring challenge in economic evaluations of gene therapies is the measure and valuation of HRQOL. A recommendation reaching consensus was to conduct vignette studies to obtain health state utility values for this population. Several publications used vignette studies, but most of those studies were also flawed, and in experimental vignette studies, flaws in study design or conduct may limit data integrity or introduce biased results [77]. The full potential of vignette studies has not yet been realized [77]. Only one evaluation actually used the approach from the consensus recommendation, having vignettes valued by a sample of the general public using a direct utility method [56].

Only four of 14 economic evaluations for pediatric populations considered HRQOL for caregivers, even though severe pediatric conditions may be expected to have a substantial impact on caregiver HRQOL [78]. According to studies that incorporated disutility values for caregivers, the impact on the ICER appeared to be small to moderate.

Several recommendation papers argued that costs should be valued from two perspectives – health care payer and societal [19,21,23–25]. We fully agreed with this, but only 11% of studies reported analyses from both perspectives, possibly because economic evaluations may be related to a specific country's HTA guidelines [17]. Another explanation may be that authors of economic evaluations simply considered that the fraction of costs not paid by the health care payer was modest. Thus, the differences in ICERs between societal and health care payer perspectives, in articles reporting both, never exceeded 25%. While we all agreed with the recommendation to explore the influence of alternative payment mechanisms in terms of CE, <20% of reviewed economic evaluations reported such analyses.

Consensus recommendations related to modeling methods and choices were more frequently followed than other recommendations, specifically deterministic and probabilistic sensitivity analyses and sensitivity analyses around discount rates. The time frame and discount rates often had a large impact on results, which confirms the importance of conducting these sensitivity or scenario analyses. The greater variability in results according to discount rates raises the question of which discount rates are the most appropriate. We generally recommended following standard methodologic guidelines for the base-case analysis, which will ensure comparability between studies. However, debate exists on whether the discount rates recommended by some HTA agencies are truly appropriate [79].

There were several limitations to our study. Reviewed studies were in the English language only. In addition, economic evaluations were identified through MEDLINE and EMBASE, and we did not search for reports published on the websites of HTA agencies. A substantial number of articles were not included because of the exclusion criteria implemented during the systematic literature review (e.g. articles addressing genetic tests, genotyping and whole-genome sequencing interventions, clinical trials, cost-minimization analyses, and



cost-of-illness analyses). Finally, we reviewed recommendations without providing additional specific insight into our assessment of the recommendations. Having specific insight for recommendations that did not reach the consensus threshold may have provided additional understanding.

## 5. Expert opinion

Despite the potential clinical benefits associated with some gene therapies, obstacles to efficient market access, including reimbursement and funding challenges, prevail. Although it is important to assess the CE of gene therapies because there are greater upfront costs associated with these treatments, it is challenging to perform this assessment because of the limited relative effectiveness data available and the uncertainty around the long-term effectiveness and safety of these treatments. No consensus exists on the ideal methodology to evaluate the economic impact and benefits of new, potentially curative gene therapies or the associated reimbursement and funding challenges for existing health care payments structures. This is concerning because a growing number of gene therapies are expected to be approved in the coming years. Therefore, novel approaches for economic evaluation of gene therapies are urgently needed to address these issues.

We aimed to identify and describe the most widely accepted published methodologic recommendations for the economic evaluation of gene therapies and to assess whether these recommendations were applied in published evaluations. Economic evaluations of gene therapies in the current medical literature highlight several issues that have been accepted as limitations of economic evaluations in other therapeutic areas. The studies we reviewed in this systematic literature review largely presented ways to evaluate gene therapies as appropriately as possible within the standard CE analysis framework.

We reviewed the recommendations in the current medical literature associated with the economic evaluation of gene therapies and identified several consensus recommendations. Because of a lack of long-term observed clinical benefits for gene therapies, surrogate endpoint data should be used to assess clinical efficacy. The rationale for conducting noncomparative studies should be clearly elucidated, and the inclusion of other nonrandomized data should be considered to provide complementary information to a single-arm trial. For access to gene therapies, consideration of innovative payment mechanisms was supported, which would facilitate patient access while maintaining sustainable health care budgets. The HRQOL of caregivers and patients' families must be considered in economic evaluations of gene therapies. After the launch of a gene therapy, gathering real-world evidence is important to confirm the benefits of treatment and to provide further evidence following initial evaluations.

Fully addressing the limitations of economic evaluation in the context of gene therapies may require methodologic changes beyond those that health economists currently appear willing to accept. However, with the clinical progress made over recent years, gene therapies are now considered a potentially paradigm-shifting treatment, making it possible to

treat incurable diseases with unmet needs. The results of the current study highlight the considerable HTA challenges that remain. It is important to understand what additional data are needed to convince decision-makers to pay for the potential long-term treatment benefits associated with gene therapies. These guidelines were summarized to critically appraise and assess the applicability and impact of the recommendations in published evaluations, which may facilitate the implementation of more important recommendations in future evaluations.

## 6. Conclusions

Health economists conducting evaluations of gene therapies have access to a relatively large number of publications that provide methodologic recommendations. Economic evaluations of gene therapies highlight several issues that have been accepted as limitations of economic evaluations in other therapeutic areas. Studies from the reviewed literature generally presented ways to evaluate gene therapies as appropriately as possible within the standard CE analysis framework. Fully addressing the limitations of economic evaluation in the context of gene therapies may require methodologic changes beyond those that health economists may readily accept. Most of these recommendations are currently generally not followed by published economic evaluations of gene therapies. Assessing the applicability and impact of the recommendations from this study may facilitate the implementation of important recommendations in future evaluations.

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## Declaration of interest

M Toumi and S Aballea are former employees of Creativ-Ceutical and consultants for Novartis Gene Therapies, Inc. (NGT). O Dabbous is an employee of NGT and owns stock/other equities. S Sullivan is a consultant for AbbVie, Bayer, Incyte, Nanoscope, Neurocrine, NGT, Novo Nordisk, and Spark Therapeutics. P Neumann served on advisory boards/as a consultant for AbbVie, Amgen, Bayer, the Congressional Budget Office, Janssen, Merck, NGT, Novartis, Novo Nordisk, Vertex, and Curta Inc., and received research grants from the Alzheimer's Association, Amgen, the Bill and Melinda Gates Foundation, Lundbeck, the National Institutes of Health, and the National Pharmaceutical Council. J-M Graf von der Schulenburg and M Drummond are consultants for NGT. S Tunis has received honoraria from NGT and served on advisory boards for BioMarin and UCB Pharma. D Malone has served as a consultant for Novartis, NGT, Sarepta, Pharmacyclics, Currax, and Seres, and served on advisory boards for BioMarin and Novartis. The authors have no other relevant affiliations or financial involvement with any organization or entity with a financial interest in or financial conflict with the subject matter or materials discussed in the manuscript apart from those disclosed.



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