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Using Observational Data to Calibrate Simulation Models

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Abstract

Background—Individual-level simulation models are valuable tools for comparing the impact of clinical or public health interventions on population health and cost outcomes over time. However, a key challenge is ensuring that outcome estimates correctly reflect real-world impacts. Calibration to targets obtained from randomized trials may be insufficient if trials do not exist for populations, time periods, or interventions of interest. Observational data can provide a wider range of calibration targets, but requires methods to adjust for treatment-confounder feedback. We propose the use of the parametric g-formula to estimate calibration targets and present a case-study to demonstrate its application.

Methods—We used the parametric g-formula applied to data from the HIV-CAUSAL Collaboration to estimate calibration targets for 7-year risks of CDC-defined AIDS and/or death (AIDS/death) under 3 treatment initiation strategies. We compared these targets to projections from the Cost-Effectiveness of Preventing AIDS Complications (CEPAC) model for treatment-naïve individuals presenting to care in 1996–1999, 2000–2002, or 2003 or onwards.

Results—The parametric g-formula estimated a decreased risk of AIDS/death over time and with earlier treatment. The uncalibrated CEPAC model successfully reproduced targets obtained via the g-formula for baseline 1996–1999, but over-estimated calibration targets in contemporary populations and failed to reproduce time trends in AIDS/death risk. Calibration to g-formula targets improved CEPAC model fit for contemporary populations.

Conclusion—Individual-level simulation models are developed based on best available information about disease processes in one or more populations of interest, but these processes can

change over time or between populations. The parametric g-formula provides a method for using observational data to obtain valid calibration targets and enables updating of simulation model inputs when randomized trials are not available.

Keywords

Agent-based model; calibration; HIV; g-formula

Introduction

Simulation models, such as agent-based or individual-level models, are often used to estimate the population impact of implementing public health interventions or clinical treatment strategies. Individual-level simulation models estimate (counterfactual) outcome distributions under alternative policy or treatment strategies, and therefore answer causal questions of the form: What would the outcome distribution in a population be if a particular strategy had been implemented?

An individual-level simulation model is defined by parameters that describe the relationships between treatments, outcomes, and other variables.⁽¹⁾ A key practical challenge to creating these models is that these parameters cannot be directly estimated from the population of interest. Otherwise, the outcome distributions could be estimated directly from the data using causal inference techniques rather than via simulation. Instead, model parameters typically need to be selected from a variety of sources and thus calibration procedures are required before the model can be applied to the population of interest.^(2–5)

Calibration is the process of comparing the simulation model results to outcome distributions estimated using another analytic method, and has two steps. First, investigators must identify appropriate benchmarks as calibration targets. Second, they must ensure that the simulation model reproduces these targets in the population of interest.⁽⁴⁾ A large literature is devoted to the second step, including methods for searching the input parameter space and determining goodness of fit.^(6–9) Here, we focus on the first step, for which two broad types of calibration targets may be of interest.

One possible target is the observed outcome distribution in the population of interest. This can be estimated via a randomized trial or using observational data.⁽⁵⁾ However, replicating the observed outcomes requires knowing the distribution of treatment strategies in the population so that these can be included in the analysis. This information may be difficult to obtain except from randomized trials, where it is fixed by design. Because of this, randomized trials are often seen as the gold-standard for obtaining calibration targets, but there are several limitations to relying solely on randomized trials. First, a given trial typically compares only a limited number of interventions, and the particular strategies of interest for a model may not be included. Second, trials are often not available in the population for which the simulation model is being calibrated, or may have been conducted in a highly specific subset of the population and thus of limited generalizability.

A second possible target is the counterfactual outcome distribution under a single treatment strategy selected from among the strategies present in the population. If sufficient

observational data are available from the population of interest, this distribution can be validly estimated using causal inference techniques, such as the parametric g-formula, which is a generalization of standardization to settings with time-varying treatment and confounders (10, 11) and similar in many aspects to individual-level simulation models.(12)

Here, we propose the application of the parametric g-formula to obtain a range of calibration targets for individual-level simulation models when longitudinal data (on treatments, outcomes, and confounders) is available in a population of interest. As a case study, we apply the parametric g-formula to observational data to estimate calibration targets for the Cost-Effectiveness of Preventing AIDS Complications (CEPAC) model. We consider the question of when to initiate antiretroviral therapy (ART) for people living with HIV.

The structure of this paper is as follows. We first describe the scenario of interest, the observational dataset, and the parametric g-formula. We then describe the estimation of calibration targets, the CEPAC model, and the calibration process. The calibration process is presented in two stages: (I) estimation of the calibration targets via the parametric g0formula and assessment of original parameterization fit, and (II) update of model parameters to improve fit. We conclude with a discussion of the assumptions required for the parametric g-formula to validly estimate the true counterfactual outcome distributions.

Methods

Scenario

We consider 3 treatment initiation strategies for people living with HIV individuals in Europe and North America: (a) immediate ART initiation at diagnosis of HIV infection (currently recommended in the US); (b) early ART initiation when CD4 count falls below 500 cells/ μ l; and (c) late ART initiation when CD4 count falls below 350 cells/ μ l. ART was defined as a regimen of antiretroviral drugs including at least two nucleoside reverse transcriptase inhibitors (NRTIs) and either one or more protease inhibitors (PI) or boosted PIs, one non-nucleoside reverse transcriptase inhibitor (NNRTI), one integrase inhibitor (INSTI), one entry or fusion inhibitor, or three or more NRTIs.

Until relatively recently, it was debated whether the benefits of early treatment initiation might be offset by an increased risk of side effects and resistance that limits future treatment options.(13–15) As of 2017, these strategies have been assessed in both observational cohorts and randomized trials, and there is consensus that early initiation is preferable. We use these strategies here as simple demonstration targets,(16–18) but the same approach described here could be applied to a wider range of strategies or to more complex strategies, such as scheduled treatment interruption plans or treatment switching strategies, provided data exist.

We considered two types of outcomes: death and a combined outcome of death or AIDS (defined as first diagnosis of an AIDS-defining opportunistic illness (19)). For each outcome, we estimated two calibration targets: the 7-year outcome risk and the 7-year survival curve under each treatment initiation strategy.

Data source

We used data from the HIV-CAUSAL Collaboration (20, 21) to estimate our calibration targets. This collaboration combines data from prospective cohorts of HIV-positive individuals in Brazil, Canada, France, Greece, Netherlands, Spain, Switzerland, UK, and USA. We excluded data from Brazil from our analyses because a separate parameterization of the CEPAC model exists for Brazil. The estimates presented here are based on data pooled in December 2015.

Our analyses included all HIV-positive individuals in the HIV-CAUSAL Collaboration aged 18 years or older, who were ART-naïve, did not have AIDS, were not pregnant, and had a CD4 cell count and HIV RNA measurement within the past three months of baseline. The start of follow-up for each individual was defined as the first month in which all eligibility criteria were met during three baseline time periods: 1996–1999, 2000–2002, and 2003 and onwards.

Each individual was followed until the earliest of death; censoring, defined as 12 months after the last recorded CD4 cell count and HIV RNA measurements or pregnancy (if known); or the study-specific administrative end of follow-up between February 2010 and March 2013. Unlike in a previous application of the parametric g-formula to estimate the impact of treatment initiation on survival in these data,(17) we did not require that baseline occur within 6 months of HIV diagnosis because date of diagnosis is not used in CEPAC.

Parametric g-formula

We used the parametric g-formula to estimate our calibration targets adjusting for time-fixed and time-varying confounders.(10, 11, 22) The implementation of the parametric g-formula involves two steps: (I) parametric estimation of the joint distribution of covariates and outcome over time; (II) simulation of the counterfactual outcome distribution under different treatment strategies.(22) A brief description of each step follows. For more details we refer readers to previous applications of the parametric g-formula. (17, 23–25)

Step I: Parametric estimation of the joint distribution of the data

As in previous analyses (16–18), we considered the following covariates: baseline CD4 count in cells/ μ l (<50, 50–99, 100–199, 200–349, 350–499, ≥500), HIV RNA log copies per mL (<4, 4–5, >5), sex, race, geographical origin (Europe and the USA, sub-Saharan Africa, rest of the world, unknown), transmission group (heterosexual, homosexual or bisexual, injection drug user, and other/unknown), age in years (<35, 35–50, >50), calendar year (<2000, 2000–04, 2005–10, 2011–13), and cohort; and time-varying CD4 cell count, HIV RNA, and AIDS (when not an outcome). Importantly, these time-varying confounders are also affected by prior treatment history leading to treatment-confounder feedback.

We next fit parametric regression models to the HIV-CAUSAL data to estimate the joint distribution of the time-varying covariates, treatment initiation, and mortality over time. We used logistic regression to model the time-varying indicators for death, AIDS, and treatment initiation, and linear regression to model the natural logarithms of HIV RNA and CD4 cell count. All models included the two most recent values of the time-varying covariates, as

well as time since last CD4 count and HIV RNA measurements, and all baseline covariates. Models for CD4 count and HIV RNA also included product terms for the number of months since ART initiation.

Step II: Estimation of the counterfactual outcome distribution under several treatment strategies

For each treatment strategy, we used the parametric models fit in Step I to generate a cohort of the same size as the original data via Monte-Carlo simulation. At each time point, we assigned treatment according to the treatment strategy. We then calculated the outcome distribution in the simulated data.

We used non-parametric bootstraps with 500 samples to obtain 95% confidence intervals. For each bootstrap, we took a random sample of the HIV-CASUAL data with replacement and repeated the full process of Steps I –II. We then calculated the 95% confidence interval based on the 2.5th and 97.5th percentiles of the bootstrap outcome estimates. All analyses were conducted using the GFORMULA macro(26) in SAS 9.4.

As a sensitivity analysis, we estimated the outcome distributions under no intervention. That is, we assigned treatment at each month based on the estimated conditional probability of treatment observed in the HIV-CAUSAL data from Step 2.

CEPAC Model

We calibrated the Cost-Effectiveness of Preventing AIDS Complications (CEPAC) model. The CEPAC model is a large, well-established individual-level simulation model that has been used to assess cost-effectiveness of a range of clinical and public health policies aimed at reducing the burden of disease and improving survival among people living with HIV(27–29). Simulated individuals in CEPAC exist in one of three health states: chronic infection, acute OIs, or death. Time-varying covariates, including CD4 count, HIV RNA level, age, and sex, govern transitions between health states. There are three distinct probabilities that govern mortality each month: the probability of chronic AIDS-related mortality, the probability of acute OI-related mortality, and the probability of non-AIDS-related mortality. Full specification of the model is provided on the CEPAC website (30) and elsewhere (28, 31).

The CEPAC model was originally designed using data from the Multicenter AIDS Cohort Study (MACS) —a cohort study of HIV-positive men which began enrollment in 1984.(32) CEPAC was originally calibrated to data from cohorts of people living with HIV in the US; the prior calibration is described in detail elsewhere.(33) However, changes in available treatments, testing strategies, and demographics of people living with HIV over time require that the model be periodically recalibrated to ensure it continues to provide valid estimates of survival and AIDS progression in contemporary settings. Here, we compared the estimated outcome distributions from CEPAC to calibration targets obtained using the parametric g-formula applied to HIV-CAUSAL data in the three baseline time periods. Because more recent cohort data more accurately reflect the current clinical course of HIV disease and treatment, we aimed for the recalibrated model to better fit the data from 2003 onwards.

Calibration

We simulated each treatment strategy and time period in CEPAC using 10^5 individuals. For each simulated population, we used the observed baseline distributions of CD4 count (mean and standard deviation), HIV RNA strata (% within strata), age (mean and standard deviation), gender (% male), and transmission risk group from individuals in HIV-CAUSAL who met the inclusion criteria within that baseline time period (Table 1, Appendix 1).

We used a previously published set of CEPAC parameter values for modeling AIDS-related mortality, OI incidence, and changes to CD4 count and HIV RNA over time (Table 2).^(28, 32, 34–36) Sex-stratified lifetables for the probability of non-AIDS-related mortality were obtained from the Eurostat website.⁽³⁷⁾ Key parameter values are given in Table 2. For more detail on these parameters, their sources, and how they govern transitions between health states in CEPAC see Appendix 2.

We focused our calibration on two parameters, selected because their initial values had been determined based on discussions with clinicians and other experts in the field, without a substantial empirical evidence base. Specifically, we assessed the on-treatment multiplier for OI incidence and the on-treatment multiplier for chronic AIDS-related mortality. In the original CEPAC calibration, both multipliers were set to 1.0. We performed a grid search by varying the on-treatment multipliers jointly between 1 and 0 by 0.2 units for a total of 36 calibration runs. We chose a simple grid search, rather than more formal search procedures such as Nelder Mead, because our goal was chiefly to demonstrate the method of obtaining calibration targets, rather than performing a full calibration of CEPAC.

We assessed the fit of the 7-year mortality and combined AIDS or mortality risks from the calibration runs in comparison to the values of the calibration targets estimated using the parametric g-formula applied to HIV-CAUSAL. Fit was determined based on the number of treatment strategies and outcomes for which the CEPAC risk estimates were within the 95% confidence intervals for the g-formula risks estimates. We also visually inspected survival and AIDS-free survival curves produced by the calibration runs of CEPAC and those obtained via the g-formula.

As sensitivity analyses, we compared the distributions of mean CD4 count and HIV RNA over time estimated using CEPAC and the parametric g-formula (Appendix 3), and assessed the impact on the CEPAC model results of varying a range of additional input parameters (Appendix 4).

Results

Stage I: Comparison of the g-formula estimates with the original CEPAC model results

The HIV-CAUSAL Collaboration included 28,269 eligible individuals in 1996–1999, 25,433 in 2000–2002, and 78,093 in 2003 and onwards. Baseline characteristics are shown in Table 1. The median duration of follow-up was 65 months (IQR: 27–151) in 1996–1999, 64 months (IQR: 27–126) in 2000–2002, and 36 months (IQR: 18–66) in 2003 and onwards. Treatment initiation was similar in the three time periods—63% of participants initiated

ART during follow-up when baseline was between 1996 and 1999, 61% when baseline was between 2000 and 2002, and 65% when baseline was in 2003 or later.

Table 3 compares the observed risks from HIV-CAUSAL with those estimated via the g-formula under no intervention on treatment, and indicates that the g-formula provides a good fit with the observational data. Additional sensitivity results for covariates are presented in Figures A2–A5. The g-formula estimates for AIDS onset are somewhat further from the observational data in the 1996–1999 baseline. This may be due to differences in confounders early in the HIV epidemic, differences in the subset of individuals infected with HIV presenting to care. A similar process could be used to assess the calibrated CEPAC model, if the distribution of treatment initiation strategies in HIV-CAUSAL were known, by taking a weighted average of CEPAC results over the treatment strategies followed in the data.

The calibration targets obtained from the g-formula under the three interventions of interest are reported in Table 4. Using the g-formula, the estimated 7-year risk of mortality under immediate ART initiation was 8.0% (95% CI: 7.5, 8.5) for baseline in 1996–1999, 6.9% (95% CI: 6.4, 7.4) for baseline in 2000–2002, and 4.1% (95% CI: 3.9, 4.4) for baseline in 2003 and onwards (Table 4). The estimated 7-year risk of AIDS or death under immediate ART initiation was 14.0% (95% CI: 13.4, 14.6) in 1996–1999, 12.6% (95% CI: 12.0, 13.3) in 2000–2002, and 7.9% (95% CI: 7.6, 8.3) in 2003 and onwards (Table 4). Earlier treatment initiation resulted in decreased risks for both outcomes, in all time periods.

The original CEPAC parameterization (using 1.0 for each of the on-treatment multipliers) resulted in 7-year mortality risk estimates that were within the 95% confidence intervals of the g-formula estimates under all treatment strategies for 1996–1999, but not for the 2000–2002 or 2003 and onwards (Table 3, original CEPAC column). The CEPAC estimates of the risk of combined AIDS/death outcome were higher than the g-formula estimates in all time periods. Appendix Figures A1 and A2 display the estimated survival and AIDS-free survival curves obtained via the g-formula and the original CEPAC model.

Stage II: Update of the simulation model parameters via calibration to g-formula targets

Figure 1 displays the survival and AIDS-free survival curves from all 36 CEPAC calibration runs compared to the g-formula estimates from 2003 and onwards (the CEPAC estimates are shown in grey and the observational estimates are shown in black). For the strategies of immediate initiation and initiation at CD4 <500, multiple CEPAC calibration parameter sets returned estimated survival curves that fit well with the observational estimates for both outcomes, although survival and AIDS-free survival curves were somewhat steeper in the first 6012 months of follow-up when using the g-formula (Figure 1). In contrast, for initiation at CD4 <350, all CEPAC calibration parameter sets underestimated both survival and AIDS-free survival compared with the observational data (Figure 1).

Figure 2 shows the 7-year mortality and combined AIDS or death risks in the 36 CEPAC calibration runs. The top left corner of each sub-figure represents the 7-year risk when the on-treatment multipliers for both OI incidence and chronic AIDS-mortality were set to 0, for a given treatment strategy. This is equivalent to a situation in which individuals on ART are not at risk for these outcomes. The bottom right corner of each sub-figure represents the

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scenario where both multipliers are set to 1, which would occur if the probability of OIs or chronic AIDS-related mortality was the same regardless of treatment status. In each sub-figure, the location in the heat map which is closest to the 7-year risk estimated from the observational data is indicated by the black box; the area of the heat map which produces estimates within the 95% confidence intervals estimated from the observational data is enclosed in grey boxes.

The g-formula mortality risk at 7 years was reproducible for immediate initiation and initiation at CD4 <500 when the multiplier for chronic AIDS-related mortality was close to zero and the multiplier for OI incidence was at or below 0.2 (Figure 2). For initiation at CD4 <500 cells/ μ l the best fitting on-treatment multiplier for chronic AIDS-related mortality changed to between 0.4 and 0.6. For initiation at CD4 <350, all calibration runs returned estimates greater than the upper bound of the 95% confidence intervals from the g-formula estimates for both outcomes.

Based on the 36 calibration runs, we fine-tuned the on-treatment parameter values by applying a range of multipliers to the incidence of opportunistic infections and chronic AIDS-related mortality in CEPAC for simulated patients on ART and projected overall mortality and composite outcome of AIDS or death using this grid of possible multipliers. We compared these projections to those generated by parametric g-formula estimates from HIV-CAUSAL. Subsequently, we validated the newly calibrated CEPAC projections to mortality estimates from the COHERE cohort.(38) This fine-tuning resulted in final calibrated parameter values of 0.2 for OI incidence and 0.1 for chronic AIDS-related mortality. We used these values to run CEPAC in all three time periods and present the risk estimates obtained from this calibrated model under the three treatment initiation strategies in Table 3.

The mortality risk and the combined AIDS or mortality risk estimates from the calibrated CEPAC model were very close to the corresponding observational estimates for 2003 and onwards under immediate treatment initiation and under initiation at CD4 counts above 500 cells/ μ l, but over-estimated the observational estimates under initiation below 350 cells/ μ l (Table 3, calibrated column). Further calibration using the approach described here could be performed to improve model performance for delayed treatment initiation strategies. Here, we omit this calibration, since our goal is to demonstrate the process of using the g-formula for calibration rather than present a full calibration of CEPAC.

Discussion

We have described a step-by-step procedure to identify counterfactual calibration targets from observational data for use in calibrating an individual-level simulation model. To do so, we estimated risks under several treatment strategies in an observational study (HIV-CAUSAL Collaboration) using the parametric g-formula, identified model (CEPAC) input parameters for which empirical data did not exist, and modified values of these parameters to match the counterfactual outcome estimates.

The parametric g-formula allows us to estimate the outcome distributions under any treatment strategy of interest followed by at least some individuals in the data. Unlike

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traditional regression approaches, the parametric g-formula is able to provide unbiased outcome estimates even in the presence of treatment-confounder feedback – that is, even when time-varying confounders are themselves affected by prior treatment.(11, 12, 22, 39) A similar approach could be used to estimate counterfactual calibration targets under other types of interventions, such as treatment interruption, monitoring of treatment efficacy, or treatment switching, and is particularly useful when complex interventions which depend on the evolution of time-varying markers of disease progression are of interest. For example, the g-formula could be used to obtain calibration targets for a cardiovascular risk model under a range of statin use strategies. Finally, even when trials with appropriate interventions exist, non-adherence and loss to follow-up may make intention-to-treat estimates insufficient. In these cases, obtaining appropriate calibration targets may require adjustment for time-varying non-adherence or analysis of intermediate outcomes, and can benefit from the use of methods such as the parametric g-formula. (10, 11, 22, 40, 41)

Our case study was based on the HIV-CAUSAL Collaboration, a consortium of observational cohorts that collect longitudinal data on treatments, confounders, and outcomes. The validity of this calibration procedure relies critically on the validity of the parametric g-formula estimates. The parametric g-formula requires the assumptions of no unmeasured treatment-outcome confounders and no misspecification of the parametric models for treatment, confounders, and outcomes. Although unmeasured confounding is always possible in observational studies, we adjusted our models for the main clinical indications for treatment initiation during the study period such as CD4 count, HIV RNA, and AIDS diagnosis, as well as for other potential confounders such as age, sex, geographic origin, and mode of transmission. We also assessed the potential for model misspecification in our parametric g-formula estimates by estimating the outcome distributions under the observed distribution of treatment initiation strategies in HIV-CAUSAL, and found good agreement between the data and our model results.

In our analysis, survival and AIDS-free survival both increased substantially between the three baseline time periods when using the g-formula. This trend for improved survival of people living with HIV has been observed elsewhere. (42) The original CEPAC parameterization resulted in estimates that fit well with the g-formula estimates from 1996-1999, but not with those from most recent periods and CEPAC does not currently model changes in parameters by calendar time. After calibration, the CEPAC model fit for the 2003 onwards baseline period was greatly improved especially under immediate treatment initiation and initiation at CD4 counts below 500 cells/ μ l.

The validity of the calibration also relies on the assumption that the population used for calibration experiences the same treatment effects and outcome risks as the target population for model inference. In our case, a majority of HIV-CAUSAL participants live in Europe, while the CEPAC-US model has generally been used to estimate outcomes in the United States. Differences in access to, and cascades of, care, underlying mortality risks, or infectious disease transmission between these regions may explain some of the differences we observed for the strategy of initiating treatment at CD4 counts below 350 cells/ μ l in recent years.(42) Individuals who delayed treatment initiation in the post-2003 period are likely to have presented to care later in the disease process, and those presenting late may

differ between US and European populations. Therefore, estimates of calibration targets based on HIV-CAUSAL for this strategy may be less appropriate to compare with US-specific CEPAC estimates.

However, comparison of our calibrated CEPAC results with previous studies suggests that, for individuals who are retained in care and on treatment, mortality risk may not differ substantially between American and European populations. A study of a highly adherent cohort of people living with HIV in the USA found a projected mean life expectancy of 69.3 years from the time of seroconversion,(43) whereas we estimate a mean life expectancy of 67.6 years using the newly calibrated CEPAC model. The calibrated CEPAC model also matched previously published AIDS-related death rates from the COHERE collaboration within a margin of 5 deaths per 10,000 people over a 2.7 year period. Five-year survival estimates from the calibrated CEPAC model were within 2% of previously published HIV-CAUSAL estimates for the strategy of immediate treatment initiation.(20)

Here, we used a simplified process for varying and assessing parameter fit – a grid search of parameter space, and a comparison with 95% confidence intervals of the calibration targets. More sophisticated search techniques and methods of assessing goodness of fit between model outputs and calibration targets exist and could result in further improvements in the calibration.(8, 44–48) In addition, in order to focus on the process of obtaining a range of counterfactual calibration targets, we did not perform a full calibration to fit the model results for all outcomes. However, we were able to greatly improve the fit of the CEPAC model to the strategy of immediate treatment initiation for the period of 2003 onwards, which is the strategy currently recommended for people living with HIV in the US and will therefore likely serve as a comparison strategy for future analyses.(49)

In summary, the original CEPAC model performed well when compared to calibration targets estimated from data starting in 1996–1999 – the time period for which the model was developed – but recalibration of CEPAC parameters was required to obtain a fit with targets estimated from 2003 onwards. This highlights the fact that populations can change over time such that periodic recalibration of individual-level simulation models may be needed to ensure that these models remain relevant over time. Similarly, recalibration of simulation models may be needed when the target of inference changes to a new population. Calibration must thus be an ongoing process to keep simulation models up-to-date and relevant for the population, or populations, of interest. However, randomized controlled trials will likely not be available for all populations or time periods of interest. The parametric g-formula can provide a solution by increasing the availability of calibration targets for these models.

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Appendix

1. HIV-CAUSAL Collaboration detailed baseline characteristics

The main text (Table 1) provides a summary of relevant HIV-CAUSAL cohort baseline covariates necessary for parameterizing the CEPAC runs. We provide additional information on the cohort profiles in Appendix Tables A1–A4 to provide further context to our results.

2. CEPAC input parameter sources

The main text (Table 2) provides the value of key CEPAC input parameters. We provide additional information on the source of these parameters here to provide further context to our results (Appendix Table A5).

Life table data were obtained based on average values for France, Greece, Netherlands, Spain, Switzerland, and the UK. For 1996–1999, we used lifetables for the period 1996–2013; for 2000–2002, we used 2000–2013; and for 2003 and onwards, 2003–2013. This parameter is independent of treatment status.

Among individuals not on ART, CD4 cell counts decline each month (34, 35), and HIV RNA remains at an individual-specific set point. The monthly probabilities of opportunistic infections (OIs), chronic AIDS-related mortality, and acute OI-related mortality vary over time based on current CD4 cell count in a given month. When not on ART, these probabilities were obtained from the Multi-center AIDS Cohort Study (MACS). (32)

Use of ART affects simulated individuals through three main mechanisms: reducing in HIV RNA, increasing CD4 cell count, and lowering the probabilities of OIs and of mortality. The overall probability of virologic suppression (HIV RNA reduced to lowest stratum) within 6 months of treatment initiation is 86.0%. (36) To model heterogeneity of response to ART, each individual is assigned a probability of treatment response, a measure of treatment adherence, which is multiplied by the overall probability of virologic suppression to determine the individual's final probability of suppression. CD4 cell count increases by a mean of 88 cells/ μ l in the first two months after treatment initiation, and a mean of 4 cells/ μ l in the following months (36, 50). This CD4 cell count increase reduces the probability of OI incidence, acute OI-related mortality, and chronic AIDS-related mortality, since these probabilities are CD4-dependent. In addition, the probabilities of chronic AIDS-related mortality, and OI incidence can be further adjusted by 'on treatment' multipliers stratified by CD4 cell count and OI type (Pneumocystis pneumonia, Mycobacterium avium complex, Toxoplasmosis, Cytomegalovirus, fungal infections, other OIs).

Individuals who achieve virologic suppression have a monthly probability of treatment failure, meaning that treatment no longer decreases the individual's HIV RNA or increases their CD4 count. Individuals who fail treatment will experience stable, or declining CD4 counts, and rising HIV RNA. At their next modeled clinic visit, these individuals can be detected as failing and changed to a new ART regimen. There are 6 lines of ART available in the model all with the same efficacy; individuals who fail all 6 lines remained on the last failed ART regimen.

3. Survival and AIDS-free survival distributions by antiretroviral therapy initiation strategy and baseline time period

Figures A1 and A2 show the survival and AIDS-free survival estimates over time for each baseline time period and antiretroviral therapy initiation strategy, estimated using the parametric g-formula applied to HIV-CAUSAL data and using the original CEPAC parameterization.

4. Mean of study variables over time observed in HIV-CAUSAL and estimated using the parametric g-formula

Figures A3 – A5 compare the means of treatment and study covariates over time estimated using the parametric g-formula under the strategy “start treatment with the observed conditional probability from the data” with the corresponding means observed in HIV-CAUSAL. Though this comparison can detect model misspecification in the parametric g formula, the lack of differences does not guarantee no model misspecification under other strategies. If the distribution of treatment initiation strategies followed by individuals in HIV-CAUSAL were known, a similar comparison could be made between CEPAC estimates of treatment and covariates over time and the observed means. However, parameterizing treatment to match the observed conditional probability distributions in a simulation model may often not be feasible.

5. Progression of CD4 count and viral load

Some differences in survival and AIDS onset estimates between CEPAC and the g-formula may be related to differences in progression of CD4 count and HIV RNA over time. Figure A6 shows the estimated mean CD4 cell count and mean HIV RNA levels from the g-formula applied to HIV-CAUSAL, and from CEPAC for the 2003 baseline, by treatment initiation strategy, among individuals remaining alive at each month over follow-up. The CEPAC runs shown here were obtained from the original parameterization, where the treatment multipliers for OIs and for chronic AIDS-related mortality were both set to 1. Mean CD4 counts improved over the three time periods and were higher when treatment was initiated earlier for both CEPAC and the parametric g-formula. Mean CD4 counts improved more rapidly and reached much higher levels by the end of follow-up in CEPAC than with the parametric g-formula. Mean HIV RNA level decreased over the three time periods and with earlier treatment initiation. HIV RNA levels were more variable in the parametric g-formula estimates than in CEPAC, especially in the earlier baseline periods.

6. Additional parameters assessed in calibration

We first varied a range of parameters to identify those to which the CEPAC results were most sensitive. Specifically, we varied the 1-month probabilities of OI incidence and chronic AIDS-related mortality when not on-treatment, the on-treatment multipliers for OI incidence and chronic AIDS-related mortality, the highest CD4 count stratum at which OIs and/or chronic AIDS-related mortality could occur while on treatment, the probability of acute OI-related mortality (on and off treatment), the probability of treatment response, and the probability of treatment failure after suppression. To assess the importance of specific baseline characteristics, we also varied the distributions for baseline CD4 cell counts and HIV RNA strata, and assessed using transmission risk group distributions and sex-stratified lifetables from the USA only (28).

Changes to the on-treatment multipliers for OIs and chronic AIDS-related mortality, distribution of probability of response, and probability of treatment failure were the most effective in reducing the discrepancies between the CEPAC and observational estimates. We selected the on-treatment multipliers for our main calibration example since empirical data regarding the appropriate value of these parameters did not exist.

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Veterans Aging Cohort Study-Virtual Cohort

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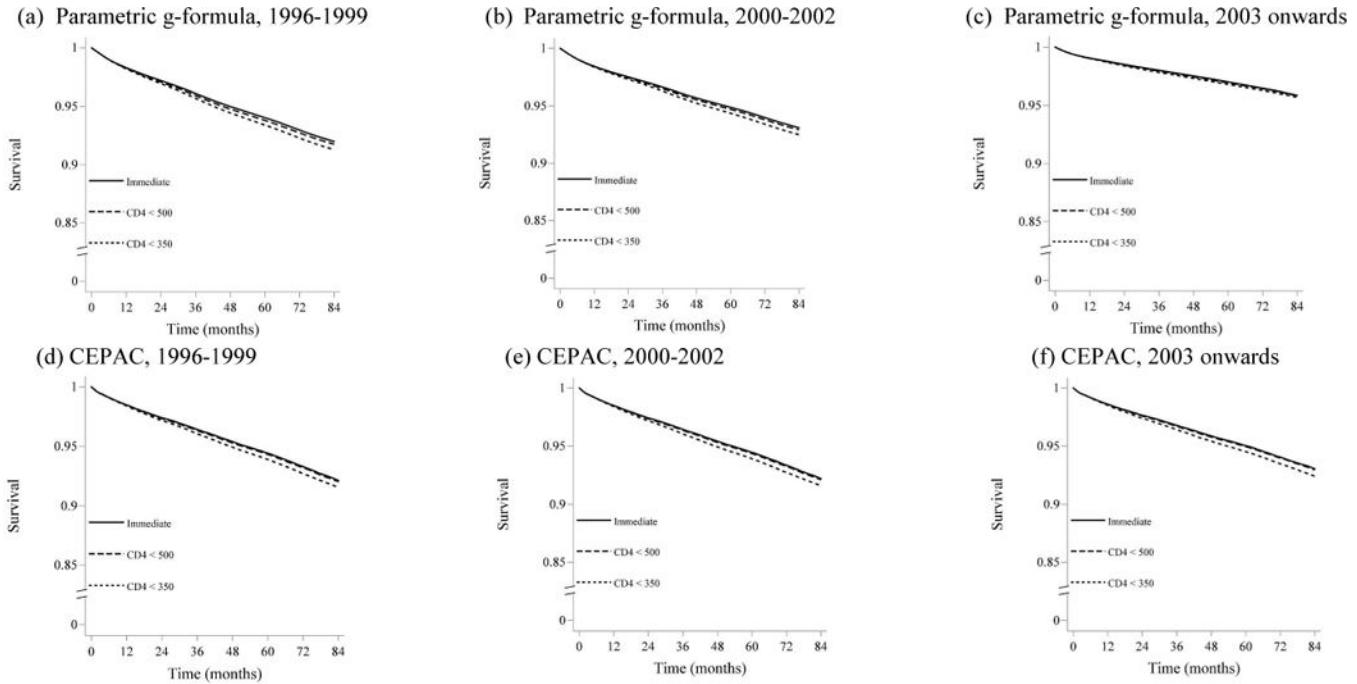
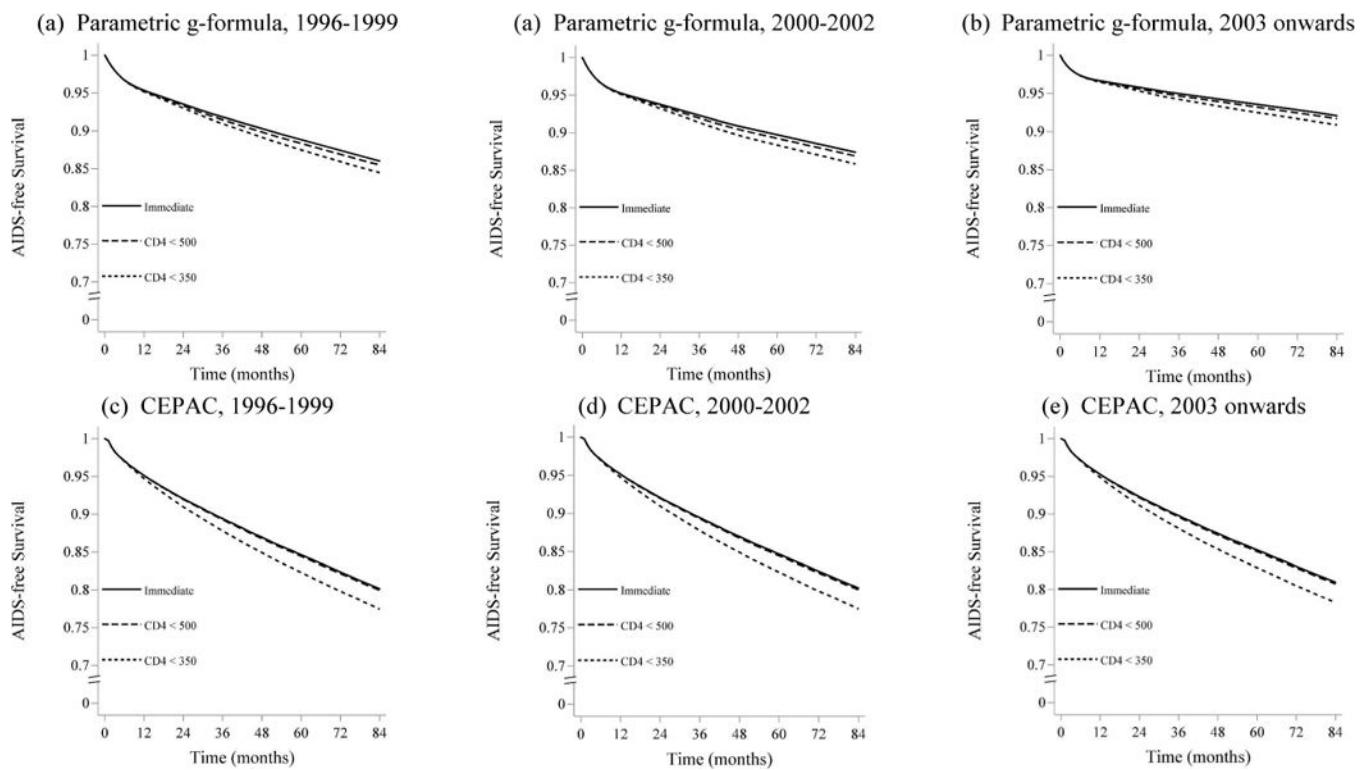


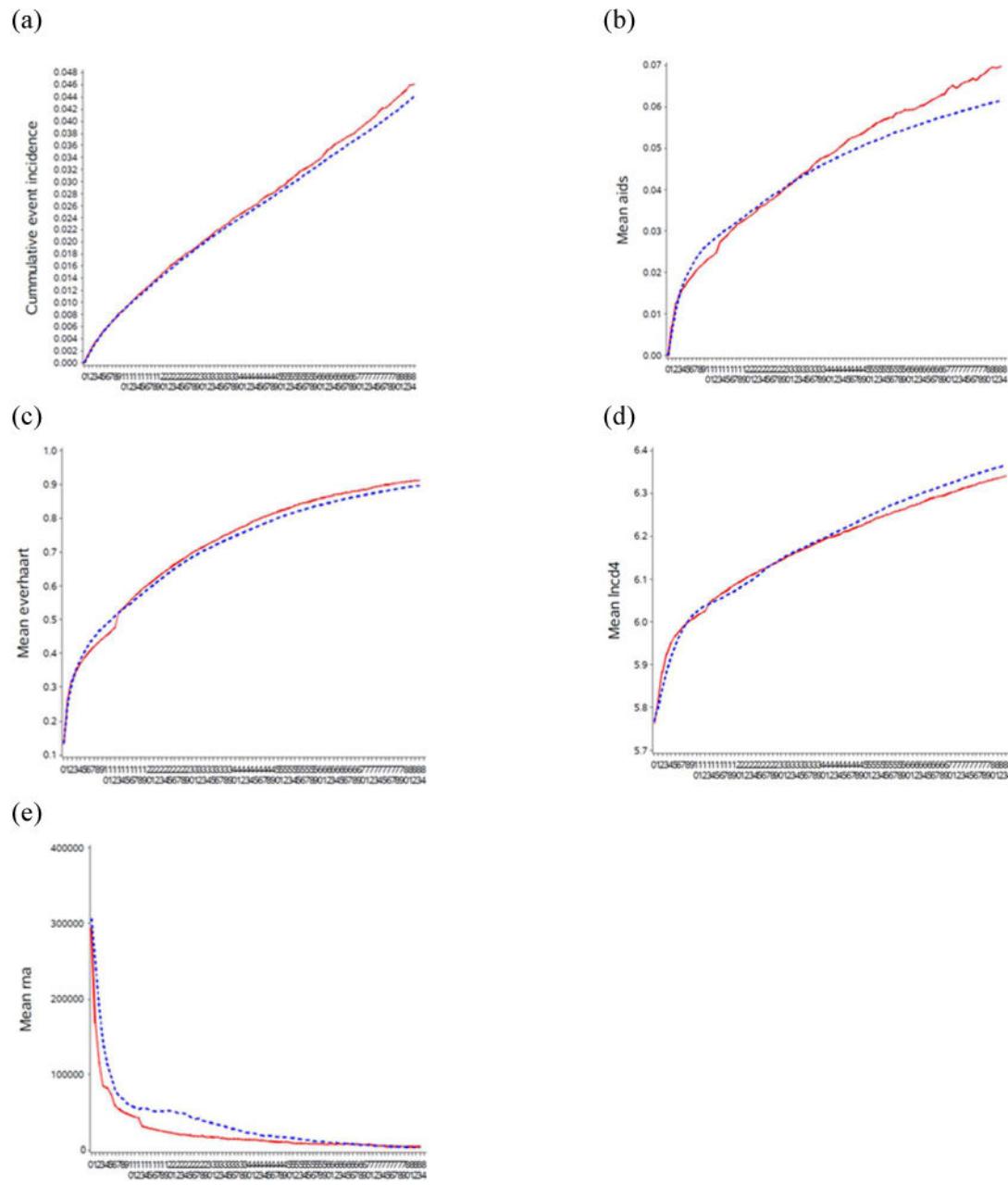
Figure A1.

Survival distribution stratified by baseline time period and antiretroviral therapy initiation strategy estimated using the parametric g-formula applied to HIV-CAUSAL data and using the original CEPAC parameterization. (a) Baseline from Jan 1, 1996 – Dec 31, 1999, parametric g-formula; (b) Baseline from Jan 1, 2000 – Dec 31, 2002, parametric g formula; (c) Baseline on or after Jan 1, 2003, parametric g-formula; (d) Baseline from Jan 1, 1996 – Dec 31, 1999, CEPAC; (e) Baseline from Jan 1, 2000 – Dec 31, 2002, CEPAC; (f) Baseline on or after Jan 1, 2003, CEPAC. All CEPAC estimates use initial parameterization of 1.0 for on-ART multipliers of opportunistic infection incidence and chronic AIDS-related mortality. CEPAC: Cost-Effectiveness of Preventing AIDS Complications model

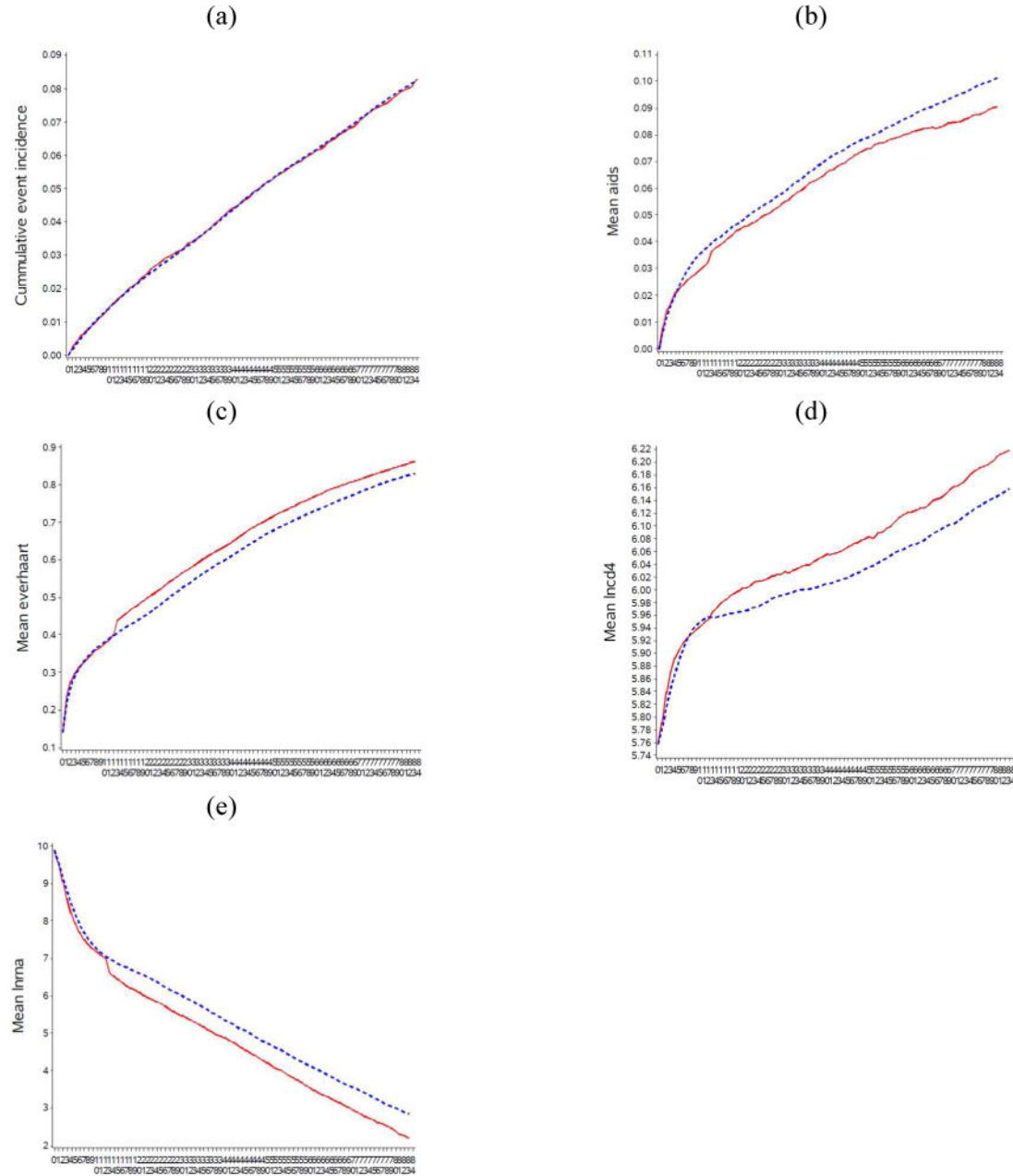
**Figure A2.**

AIDS-free survival baseline time period and antiretroviral therapy initiation strategy estimated using the parametric g-formula applied to HIV-CAUSAL data and using the original CEPAC parameterization. (a) Baseline from Jan 1, 1996 – Dec 31, 1999, parametric g-formula; (b) Baseline from Jan 1, 2000 – Dec 31, 2002, parametric g-formula; (c) Baseline on or after Jan 1, 2003, parametric g-formula; (d) Baseline from Jan 1, 1996 – Dec 31, 1999, CEPAC; (e) Baseline from Jan 1, 2000 – Dec 31, 2002, CEPAC; (f) Baseline on or after Jan 1, 2003, CEPAC. All CEPAC estimates use initial parameterization of 1.0 for on-ART multipliers of opportunistic infection incidence and chronic AIDS-related mortality.

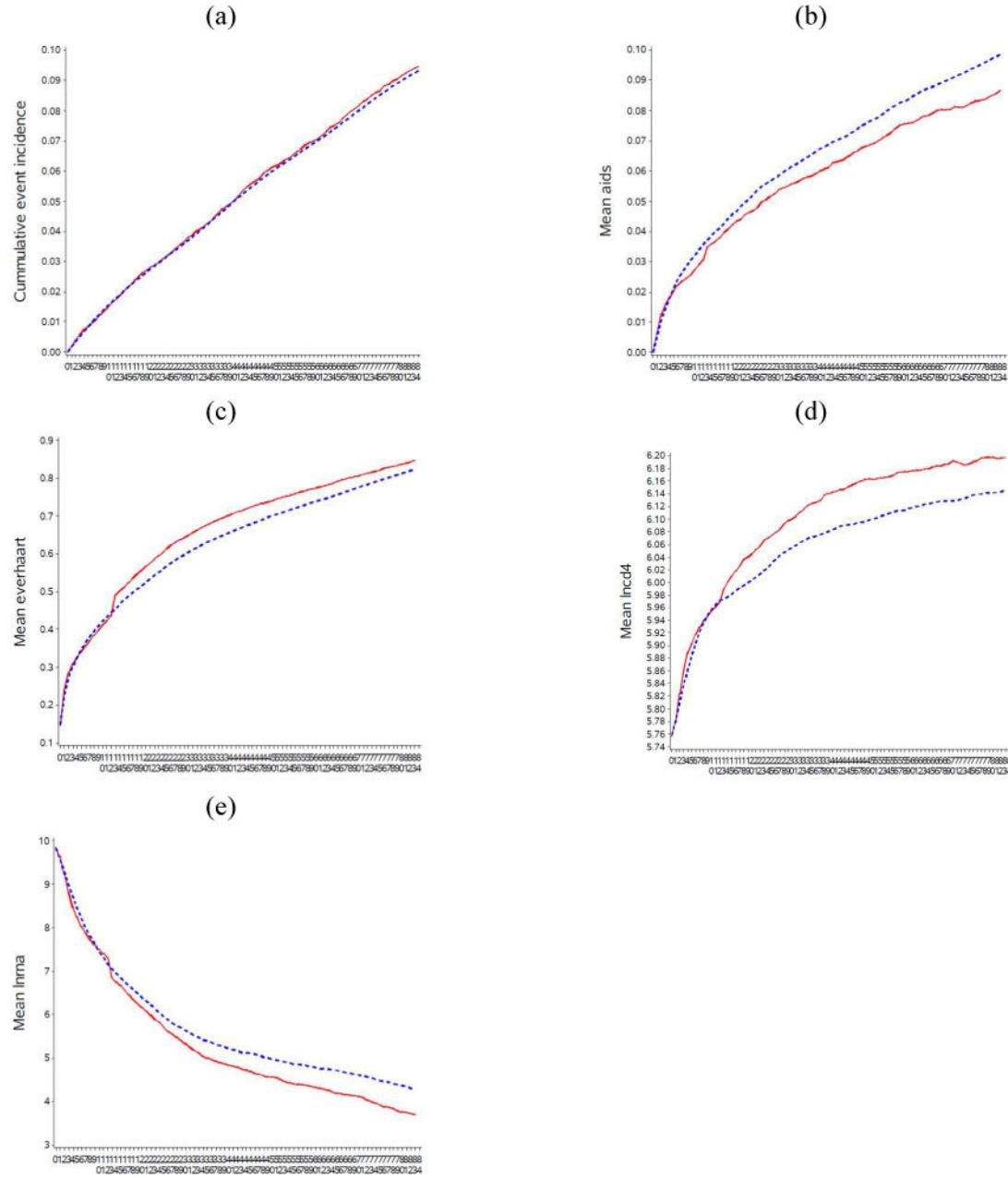
CEPAC: Cost-Effectiveness of Preventing AIDS Complications model

**Figure A3.**

Mean of the main study variables under no intervention when outcome is mortality: observed (solid line) and estimated via the parametric g-formula (dotted line). HIV-CAUSAL Collaboration, on or after Jan 1, 2003. (a) Cumulative incidence of death; (b) Cumulative incidence of AIDS; (c) Mean proportion on treatment; (d) Mean CD4 count, natural log scale; (e) Mean HIV RNA.

**Figure A4.**

Mean of the main study variables under no intervention when outcome is mortality: observed (solid line) and estimated via the parametric g-formula (dotted line). HIV-CAUSAL Collaboration, Jan 1, 2000 – Dec 31, 2002. (a) Cumulative incidence of death; (b) Cumulative incidence of AIDS; (c) Mean proportion on treatment; (d) Mean CD4 count, natural log scale; (e) Mean HIV RNA, natural log scale.

**Figure A5.**

Mean of the main study variables under no intervention when outcome is mortality: observed (solid line) and estimated via the parametric g-formula (dotted line). HIV-CAUSAL Collaboration, Jan 1, 1996 – Dec 31, 1999. (a) Cumulative incidence of death; (b) Cumulative incidence of AIDS; (c) Mean proportion on treatment; (d) Mean CD4 count, natural log scale; (e) Mean HIV RNA, natural log scale.

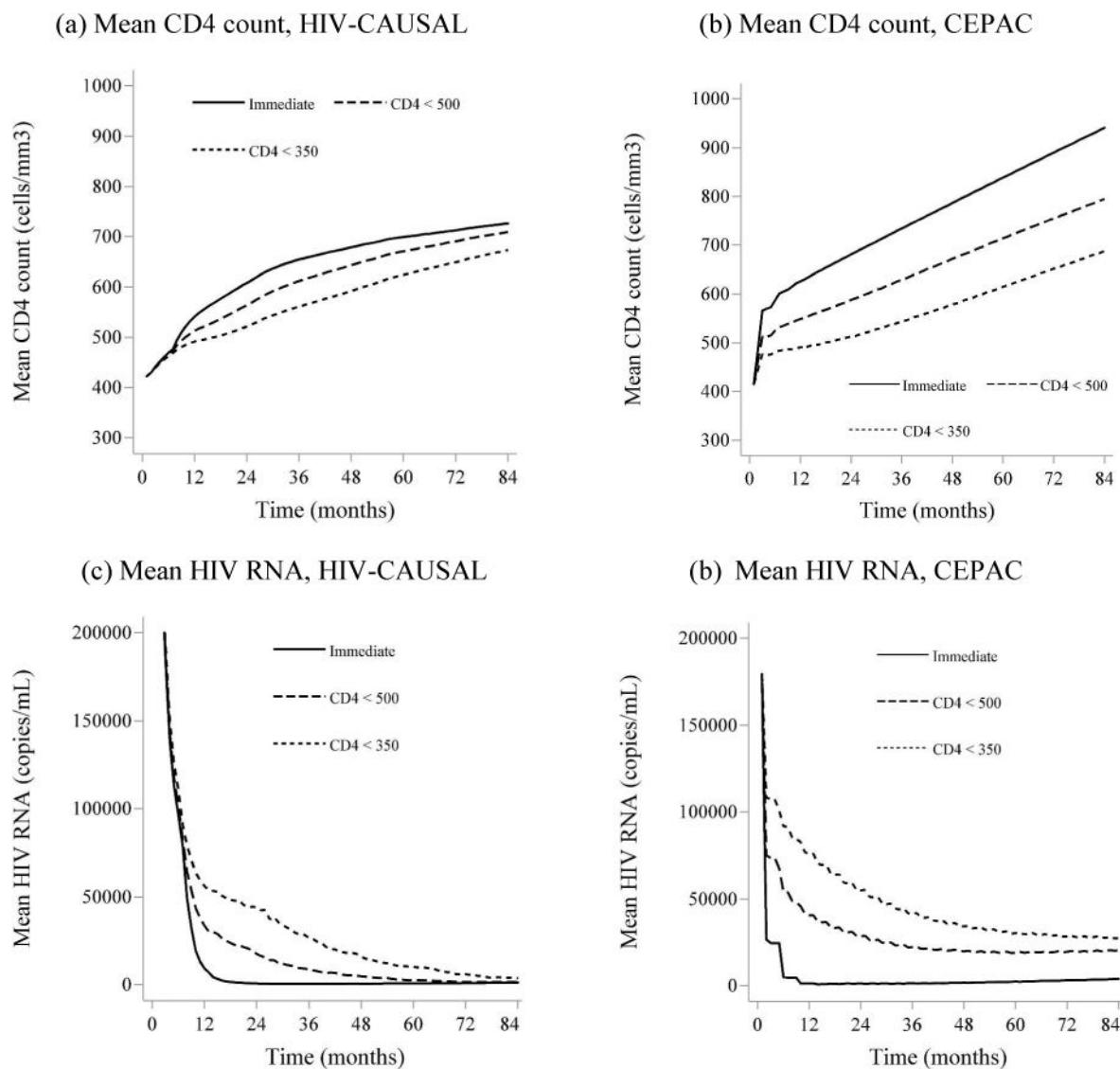


Figure A6.
Mean of CD4 count and HIV RNA under intervention

Estimating using CEPAC and via the parametric g-formula in HIV-CAUSAL Collaboration, using baseline on or after Jan 1, 2003. All CEPAC estimates use initial parameterization of 1.0 for multipliers. (a) Mean CD4 count in HIV-CAUSAL (cells/ μ L); (b) Mean CD4 count in CEPAC (cells/ μ L); (c) Mean HIV RNA in HIV-CAUSAL (copies/mL); (d) Mean HIV RNA in CEPAC (copies/mL).

Table A1

Distribution of HIV-CAUSAL data by country of cohort, %

Cohort	Jan 1 1996 to Dec 31 1999	Jan 1 2000 to Dec 31 2002	On or after Jan 1 2003
Canada	1.0	0.9	1.4
France	50.0	39.5	27.0

Cohort	Jan 1 1996 to Dec 31 1999	Jan 1 2000 to Dec 31 2002	On or after Jan 1 2003
Greece	2.3	2.0	4.8
Netherlands	2.0	6.0	12.6
Spain	7.0	10.6	17.4
Switzerland	8.6	6.2	4.1
UK	16.2	21.3	24.4
USA	12.7	13.5	9.7

Table A2

Detailed baseline characteristics, HIV-CAUSAL Collaboration: Jan 1, 1996 Dec 31, 1999

		Individuals (n)	Person-years	Proportion starting ART during follow-up (%)	Median (IQR) follow-up (months)	Deaths (n)	Incidence of death (per 1000 person-years)	Age (years)
CD4 cell count, cells/ μl	<50	1,519	9,781.25	0.84	53 (15, 145)	354	36.2	20.0
	50–100	1,172	7,693.50	0.84	53 (17, 149)	209	27.2	19.0
	100–200	2,979	19,403.50	0.80	51 (17, 142)	374	19.3	20.0
	200–350	6,295	40,830.67	0.72	50 (17, 141)	485	11.9	30.0
	350–500	6,612	41,859.00	0.62	47 (17, 135)	414	9.9	30.0
	>500	9,692	55,965.08	0.46	40 (15, 117)	525	9.4	40.0
HIV RNA, copies/mL	<10,000	10,187	56,539.58	0.45	36 (14, 110)	571	10.1	40.0
	10,000–100,000	11,498	74,533.67	0.67	50 (17, 141)	900	12.1	60.0
	>100,000	6,584	44,459.75	0.82	58 (19, 146)	890	20.0	50.0
Sex	Men	21,361	136,126.17	0.65	48 (17, 139)	2,059	15.1	10.0
	Women	6,908	39,406.83	0.57	41 (15, 114)	302	7.7	20.0
Age, years	<35	13,835	81,588.00	0.59	42 (16, 119)	572	7.0	40.0
	35–50	11,745	75,364.92	0.65	48 (16, 142)	1,152	15.3	80.0
	>50	2,689	18,580.08	0.73	64 (19, 150)	637	34.3	50.0
Transmission group	Heterosexual	8,401	52,008.58	0.64	47 (17, 130)	402	7.7	30.0
	Homosexual or bisexual	9,197	69,782.58	0.67	70 (22, 162)	545	7.8	30.0
	Injecting drug user	5,132	22,833.33	0.52	29 (11, 72)	429	18.8	30.0
	Other or unknown	5,539	30,908.50	0.65	37 (13, 117)	985	31.9	70.0

Table A3

Detailed baseline characteristics, HIV-CAUSAL Collaboration: baseline, Jan 1, 2000 Dec 31, 2002

		Individuals (n)	Person-years	Proportion starting ART during follow-up (%)	Median (IQR) follow-up (months)	Deaths (n)	Incidence of death (per 1000 person-years)
CD4 cell count, cells/ μ l	<50	1,375	8,071.00	0.88	58 (18, 122)	253	31.3
	50–100	1,109	6,494.00	0.87	57 (19, 121)	166	25.6
	100–200	2,645	14,813.58	0.83	52 (18, 118)	271	18.3
	200–350	5,678	32,327.08	0.72	51 (17, 121)	342	10.6
	350–500	5,922	34,159.50	0.58	49 (18, 122)	326	9.5
	>500	8,704	46,830.08	0.42	43 (17, 112)	350	7.5
HIV RNA, copies/mL	<10,000	8,685	44,931.42	0.43	40 (15, 108)	403	9.0
	10,000–100,000	10,631	60,588.25	0.65	50 (18, 120)	683	11.3
	>100,000	6,117	37,175.58	0.82	61 (21, 124)	622	16.7
Sex	Men	19156	132819.7	0.63	52 (19, 123)	1,471	13.4
	Women	5554	34921.58	0.57	39 (15, 98)	237	7.3
Age, years	<35	18,803	110,041.17	0.57	42 (17, 108)	282	4.9
	35–50	6,630	32,654.08	0.63	52 (18, 123)	784	11.9
	>50	11,014	57,643.25	0.72	63 (19, 124)	642	33.7
Transmission group	Heterosexual	11,270	65,988.17	0.61	44 (17, 108)	310	6.5
	Homosexual or bisexual	3,149	19,063.83	0.64	75 (26, 138)	305	5.3
	Injecting drug user	8,953	47,355.00	0.49	26 (11, 60)	242	20.8
	Other or unknown	8,280	57,125.67	0.65	42 (15, 116)	851	32.0

Table A4

Detailed baseline characteristics, HIV-CAUSAL Collaboration: On or after Jan 1, 2003

		Individuals (n)	Person-years	Proportion starting ART during follow-up (%)	Median (IQR) follow-up (months)	Deaths (n)	Incidence of death (per 1000 person-years)
CD4 cell count, cells/ μ l	<50	3,847	13,935.50	0.87	35 (15, 65)	347	24.9
	50–100	3,279	12,158.25	0.89	35 (15, 69)	212	17.4
	100–200	8,067	29,194.92	0.88	34 (15, 66)	344	11.8
	200–350	18,069	64,787.17	0.80	34 (15, 64)	432	6.7
	350–500	18,661	63,856.75	0.62	31 (14, 61)	297	4.7
	>500	26,170	85,179.33	0.44	29 (13, 57)	374	4.4

		Individuals (n)	Person-years	Proportion starting ART during follow-up (%)	Median (IQR) follow-up (months)	Deaths (n)	Incidence of death (per 1000 person-years)	Average
HIV RNA, copies/mL	<10000	20,793	68,189.42	0.49	29 (13, 58)	374	5.5	3
	10000–100000	34,976	120,595.83	0.66	31 (14, 62)	798	6.6	6
	>100000	22,324	80,326.67	0.79	34 (15, 65)	834	10.4	6
Sex	Men	62,498	219,436.33	0.66	32 (15, 63)	1,769	8.1	1
	Women	15,595	49,675.58	0.64	28 (12, 56)	237	4.8	2
Age, years	<35	34,977	112,735.75	0.59	28 (13, 57)	257	2.3	2
	35–50	32,120	118,048.75	0.69	35 (15, 66)	866	7.3	6
	>50	10,996	38,327.42	0.76	33 (15, 63)	883	23.0	7
Transmission group	Heterosexual	24,673	82,956.25	0.67	30 (13, 60)	462	5.6	3
	Homosexual or bisexual	37,434	136,221.58	0.63	34 (16, 65)	433	3.2	3
	Injecting drug user	3,361	9,018.58	0.58	20 (11, 44)	191	21.2	1
	Other or unknown	12,625	40,915.50	0.71	29 (13, 59)	920	22.5	7

Table A5

CEPAC model input parameters sources, all time periods

Parameter	Source
HIV natural history parameters	
Mean monthly CD4 count decrease when not on ART, stratified by CD4 cell count and HIV RNA	Research in Access to Care for the Homeless (REACH), San Francisco Men's Health Study, and Multi-Center AIDS Cohort Study (MACS) (32, 34, 35)
Monthly risk of opportunistic infections (OIs), stratified by CD4 cell count and OI	MACS (32); Walensky et al, 2010, <i>Clin Infect Dis</i> (28)
Monthly risk for acute OI-related death, among those with OI, stratified by OI	Walensky et al, 2010, <i>Clin Infect Dis</i> (28)
Efficacy of antiretroviral therapy	
Mean monthly CD4 count decrease when on ART, stratified by time since treatment initiation	Sax et al, 2010, <i>Lancet</i> (36)
Viral suppression after 24 weeks, %	Sax et al, 2010, <i>Lancet</i> (36)
Monthly risk of late failure after initial viral suppression	Sax et al, 2010, <i>Lancet</i> (36)
Probability of treatment response	Sax et al, 2010, <i>Lancet</i> (36); VOLTART trial(50)

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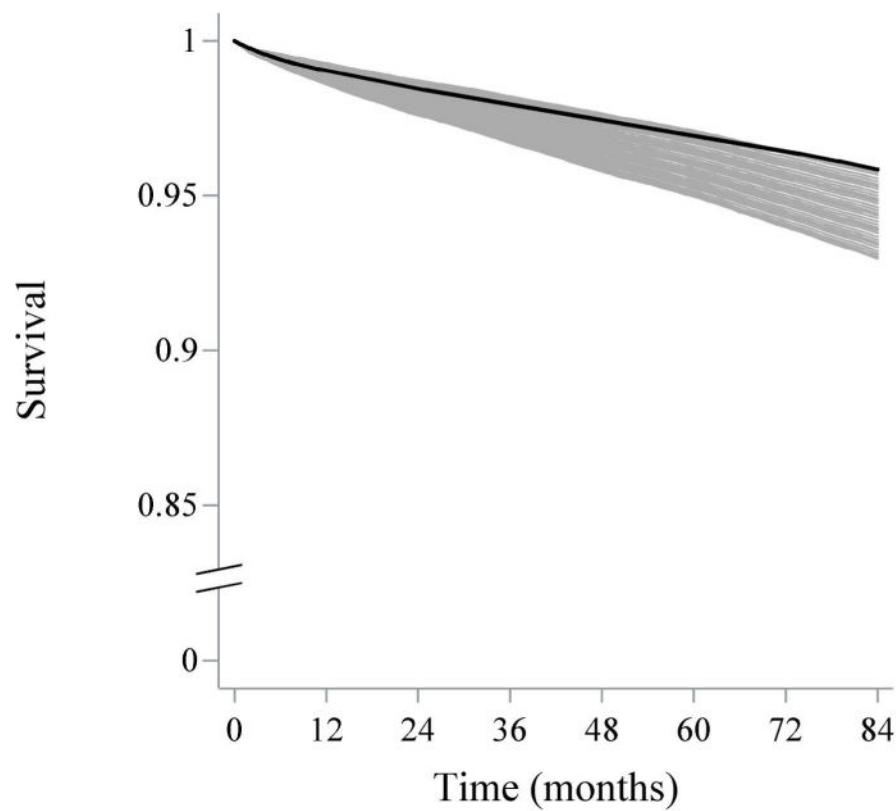
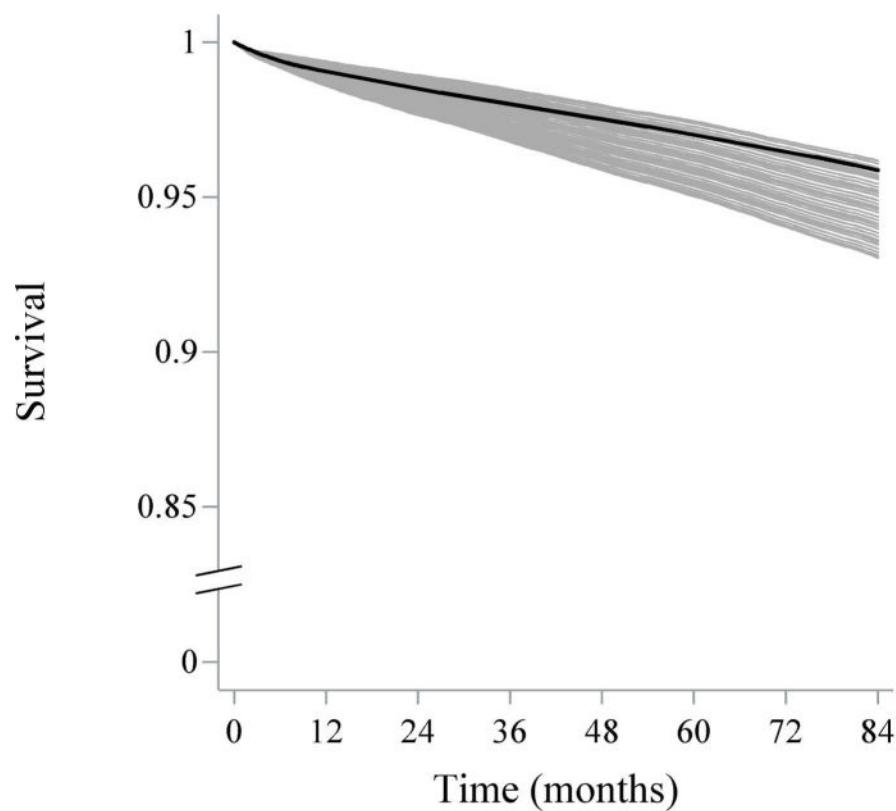
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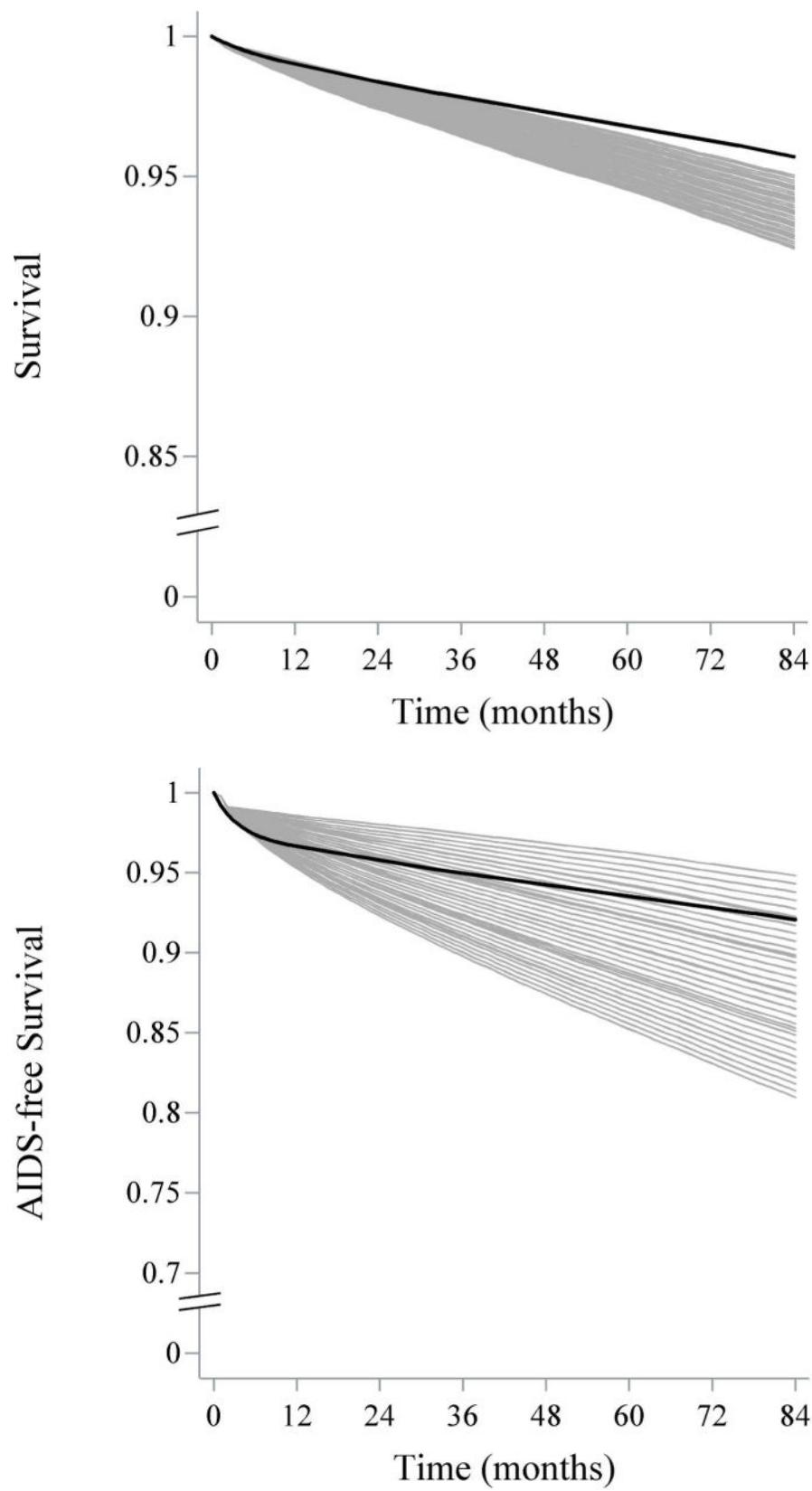


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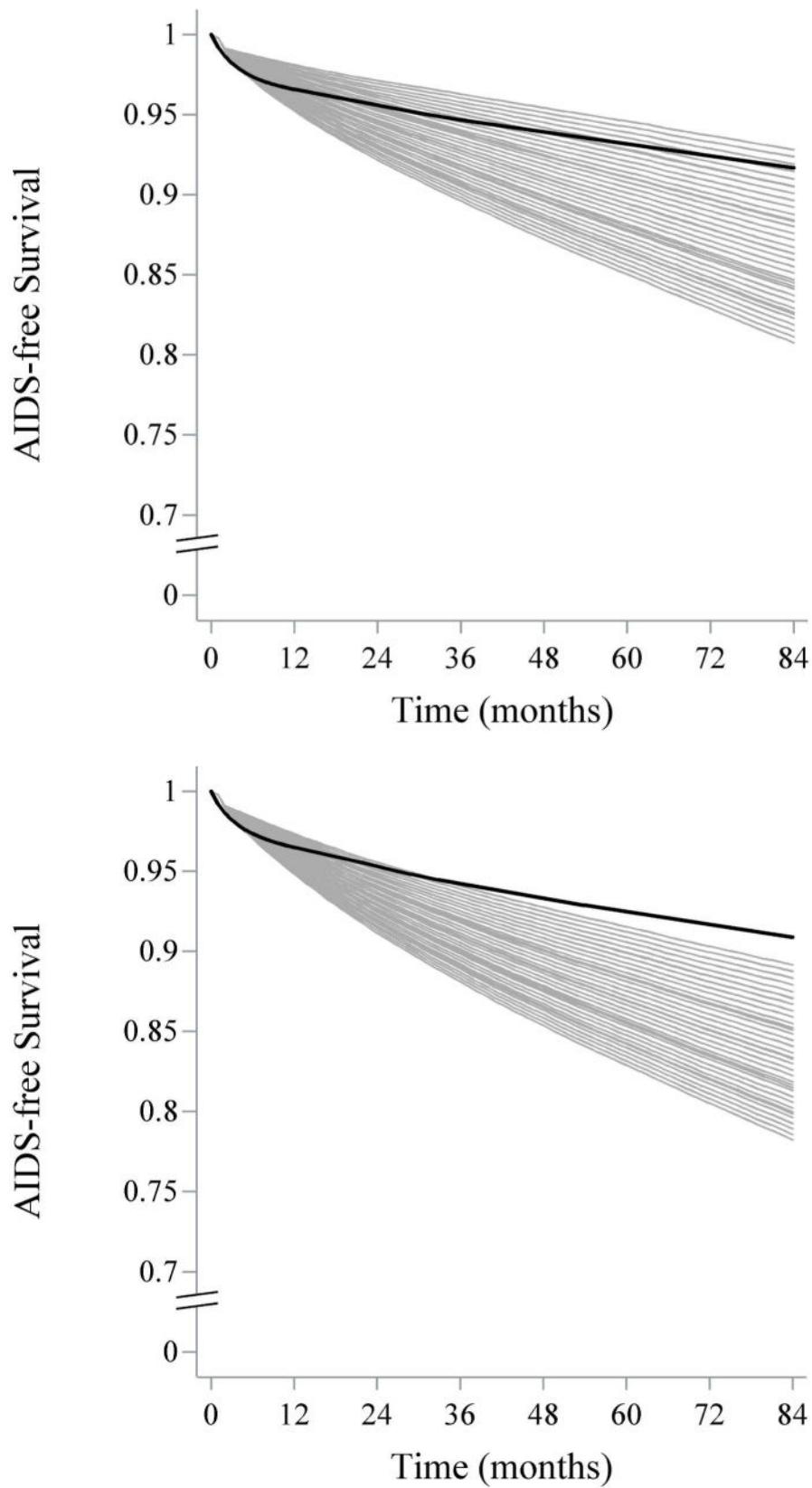
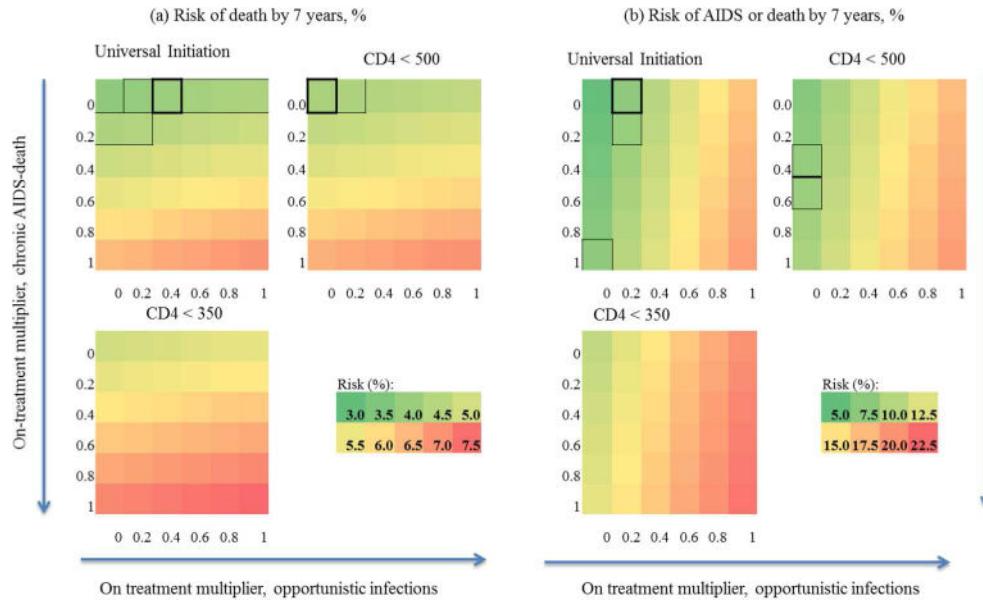


Figure 1.

Survival and AIDS-free survival over follow-up estimated via the parametric g-formula applied to HIV-CAUSAL data and each of 36 CEPAC calibration runs, varying on-treatment multipliers for opportunistic infection incidence and chronic AIDS-related mortality from 0 to 1 by 0.2.

CEPAC calibration runs (grey);, parametric g-formula estimates (black). All runs have baseline on or after Jan 1, 2003. Survival (a–c), AIDS-free survival (d–f). Immediate universal initiation(a,d); Initiation at CD4 <500 cells/ μ l(b,e); and Initiation at CD4 <350 cells/ μ l(c,f). CEPAC: Cost-Effectiveness of Preventing AIDS Complications model.

**Figure 2.**

7-year mortality (a) and combined mortality/AIDS risk (b) from CEPAC calibration runs for baseline on or after Jan 1, 2003, varying on-ART multipliers for chronic AIDS-related mortality and opportunistic infections from 0 to 1 by 0.2, stratified by treatment strategy. Scales for each outcome shown below the strategy ‘CD4 < 500’. On-treatment multiplier for chronic AIDS-related mortality increases from 0 to 1 down y-axis, on-treatment multiplier for opportunistic infections increases from 0 to 1 across x-axis, following direction of arrows. Black boxes indicate closest matches to parametric g-formula estimates when baseline is on or after Jan 1, 2003; grey boxes indicate 95% confidence interval for parametric g-formula using 500 bootstrap samples. For the strategy ‘treat at CD4 < 350’, no CEPAC runs resulted in risk estimates within the g-formula 95% confidence intervals for either outcome. CEPAC: Cost-Effectiveness of Preventing AIDS Complications model.

Table 1

Baseline characteristics of eligible individuals in the HIV-CAUSAL Collaboration by time period for variables used in CEPAC*

		Jan 1 1996 to Dec 31 1999 N = 28,269		Jan 1 2000 to Dec 31 2002 N = 25,433		On or after Jan 1 2003 N = 78,093	
CD4 cell count, cells/ μ L	<50	1,519	5.4	1,375	5.4	3,847	4.9
	50–100	1,172	4.1	1,109	4.4	3,279	4.2
	100–200	2,979	10.5	2,645	10.4	8,067	10.3
	200–350	6,295	22.3	5,678	22.3	18,069	23.1
	350–500	6,612	23.4	5,922	23.3	18,661	23.9
	>500	9,692	34.3	8,704	34.2	26,170	33.5
	Mean (SE)	426.5 (267.9)	—	426.3 (266.9)	—	421.2 (256.8)	—
HIV RNA, copies/mL	>100,000	6,584	23.3	6,117	24.1	22,324	28.6
	30,000–100,000	5,958	21.1	5,775	22.7	19,824	25.4
	10,000–30,000	5,363	19.0	4,774	18.8	14,991	19.2
	3,000–10,000	4,438	15.7	4,075	16.0	10,835	13.9
	500–3,000	3,812	13.5	3,326	13.1	7,777	10.0
	20–500	2,105	7.4	1,337	5.3	2,263	2.9
	0–20	9	0.0	29	0.1	79	0.1
Sex	Men	21,361	75.6	18,803	73.9	62,498	80.0
	Women	6,908	24.4	6,630	26.1	15,595	20.0
Age, years	Mean (SE)	36.9 (9.4)	—	37.9 (10.0)	—	37.9 (10.9)	—
Transmission group	Heterosexual	8,401	29.7	8,953	35.2	24,673	31.6
	Homosexual or bisexual	9,197	32.5	8,280	32.6	37,434	47.9
	Injection drug user	5,132	18.2	3,053	12.0	3,361	4.3
	Other/unknown	5,539	19.6	5,147	20.2	12,625	16.2

* CEPAC: Cost-Effectiveness of Preventing AIDS Complications model; OIs: opportunistic infections

CEPAC* model input parameters[†] all time periods

Mean monthly CD4 count decrease (cells/ μ L), by HIV	CD4 > 500 cells/ μ L	CD4: 350 – 499 cells/ μ L	CD4: 200 – 299 cells/ μ L	CD4: 100 – 199 cells/ μ L	CD4: 50 – 99 cells/ μ L	CD4 < 50 cells/ μ L
RNA stratum, copies/mL						
>100,000	9.54	7.94	6.34	4.66	3.30	1.37
30,000–100,000	9.54	7.94	6.34	4.66	3.30	1.37
10,000–30,000	6.85	5.70	4.55	3.34	2.37	0.98
3,000–10,000	5.87	4.88	3.90	2.87	2.03	0.84
500–3,000	4.77	3.97	3.17	2.33	1.65	0.69
20–500	2.45	2.03	1.63	1.19	0.85	0.35
0–20	2.45	2.03	1.63	1.19	0.85	0.35
Monthly risk of opportunistic infections (OIs), per 10,000	CD4 > 500 cells/ μ L	CD4: 350 – 499 cells/ μ L	CD4: 200 – 299 cells/ μ L	CD4: 100 – 199 cells/ μ L	CD4: 50 – 99 cells/ μ L	CD4 < 50 cells/ μ L
Pneumocystis pneumonia	4.0	8.0	16.0	61.8	110.0	83.6
Mycobacterium avium complex	1.0	3.0	3.0	12.0	26.0	46.9
Toxoplasmosis	0.5	1.0	2.0	3.0	10.0	7.0
Cytomegalovirus	1.0	3.0	7.0	13.0	31.0	81.7
Fungal infections	1.0	3.0	3.0	15.0	25.0	31.9
All other OIs	6.0	12.0	27.0	66.8	114.3	115.0
Monthly risk for acute OI-related death, among those with OI, %	Pneumocystis pneumonia	Mycobacterium avium complex	Toxoplasmosis	Cytomegalovirus	Fungal infections	All other OIs
3.5	4.5	18.2	4.8	3.6	4.3	
Efficacy of antiretroviral therapy	Viral suppression after 24 weeks, %		Monthly risk of late failure after initial viral suppression, %	On-treatment multiplier for OI incidence	On-treatment multiplier for chronic AIDS-related mortality	
		86.0 **	0.5 ***	1.0		

* CEPAC: Cost Effectiveness of Preventing AIDS Complications model; OIs: opportunistic infections

[†] Parameter values for original CEPAC model before calibration

** Values represent mean risk; individual's risk of viral suppression and late failure depends on probability of treatment response

Table 3

Observed and estimated 7 year risk in each baseline cohort under no intervention, HIV-CAUSAL Collaboration

Baseline period	Outcome	Observed risk at 7 years, %	Estimated risk at 7 years, % (95% CI) ^{**}
Jan 1, 1996 – Dec 31, 1999	Mortality	9.5	9.3 (8.8, 9.9)
	AIDS or death	16.2	16.6 (16.0, 17.3)
Jan 1, 2000 – Dec 31, 2002	Mortality	8.3	8.2 (7.8, 8.7)
	AIDS or death	15.4	16.0 (15.4, 16.7)
Jan 1, 2003 onwards	Mortality	4.6	4.4 (4.2, 4.7)
	AIDS or death	9.5	9.5 (9.2, 9.8)

^{**} HIV-CAUSAL estimates based on the parametric g-formula adjusted for measured time-varying confounders (CD4 count, HIV-RNA and AIDS) and baseline characteristics (calendar period and age of HIV diagnosis, risk group, gender, geographical origin, ethnicity and cohort). 95% CI: confidence interval

Table 4

Risk of all-cause mortality and combined AIDS or death outcome in each baseline cohort by ART initiation strategy estimated using the parametric g-formula and the CEPAC model.

Baseline period	ART initiation strategy (CD4/ μ l)	All-cause mortality			AIDS or mortality		
		CEPAC*	Original†	Calibrated‡	Parametric g-formula**	CEPAC*	Original†
Jan 1, 1996–Dec 31, 1999	Immediate universal	7.9	5.1	8.0 (7.5, 8.5)	19.8	8.9	14.0 (13.4, 14.6)
	CD4 >500	8.0	5.5	8.2 (7.8, 8.7)	20.0	10.7	14.5 (13.9, 15.1)
	CD4 >350	8.5	6.2	8.7 (8.3, 9.2)	22.6	14.1	15.5 (15.0, 16.1)
	Immediate universal	7.8	5.0	6.9 (6.4, 7.4)	19.8	8.8	12.6 (12.0, 13.3)
	CD4 >500	7.9	5.4	7.1 (6.6, 7.6)	20.0	10.6	13.1 (12.5, 13.8)
	CD4 >350	8.4	6.1	7.5 (7.1, 8.0)	22.5	14.0	14.2 (13.6, 14.9)
Jan 1, 2000–Dec 31, 2002	Immediate universal	7.0	4.2	4.1 (3.9, 4.4)	19.0	8.0	7.9 (7.6, 8.3)
	CD4 >500	7.0	4.6	4.2 (3.9, 4.4)	19.2	9.7	8.3 (8.0, 8.6)
	CD4 >350	7.6	5.2	4.3 (4.1, 4.5)	21.8	13.1	9.1 (8.8, 9.4)

* CEPAC: Cost-Effectiveness of Preventing AIDS Complications model; OIs: opportunistic infections

** HIV-CAUSAL estimates based on the parametric g-formula adjusted for measured time-varying confounders (CD4 count, HIV-RNA and AIDS) and baseline characteristics (calendar period and age of HIV diagnosis, risk group, gender, geographical origin, ethnicity and cohort). 95% CI: confidence interval

† Original CEPAC settings: opportunistic infection (OI) incidence on-treatment multiplier and chronic AIDS-related mortality on-treatment multiplier both equal to 1.0

‡ Calibrated CEPAC settings: opportunistic infection (OI) incidence on-treatment multiplier equal to 0.2 and chronic AIDS-related mortality on-treatment multiplier equal to 0.1.