

Pricing Longevity Bonds Using Affine-Jump Diffusion Models

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Abstract

Historically, actuaries have been calculating premiums and mathematical reserves using a deterministic approach, by considering a deterministic mortality intensity, which is a function of the age only, extracted from available (static) life tables and by setting a flat ("best estimate") interest rate to discount cash flows over time. Since neither the mortality intensity nor interest rates are actually deterministic, life insurance companies and pension funds are exposed to both financial and mortality (systematic and unsystematic) risks when pricing and reserving for any kind of long-term living benefits, particularly on annuities and pensions. In this paper, we assume that an appropriate description of the demographic risks requires the use of stochastic models. In particular, we assume that the random evolution of the stochastic force of mortality of an individual can be modelled by using doubly stochastic processes. The model is then embedded into the well know affine-jump framework, widely used in the term structure literature, in order to derive closed-form solutions for the survival probability. We show that stochastic mortality models provide an adequate framework for the development of longevity risk hedging tools, namely mortality-linked contracts such as longevity bonds or mortality derivatives.

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1 Introduction

Longevity risk, i.e., the risk that members of some reference population might live longer, on average, than anticipated, has recently emerged as one of the largest sources of risk faced by life insurance companies, pension funds, annuity providers, life settlement investors and a number of other potential players in the marketplace for this risk. For example, given the uncertainty about future developments in mortality and life expectancy, pension funds and annuity providers run the risk that the net present value of their pension promises and annuity payments will turn out higher than expected, as they will have to pay out a periodic sum of income that will last for an uncertain life span.

This risk is amplified by the current problems in state-run social security systems. Given the long-term demographic trends observed in most developed countries (with low fertility rates and an ageing population), salaries and wages earned by active workers will have to finance the pensions of a growing number of retiree, making traditional pay-as-you-go social security systems unsustainable. This will most likely force public pensions systems to moderate benefit promises in the future, reducing state-provided pension income. Additionally, the market trend away from defined-benefit corporate pension schemes towards defined-contribution plans and the move towards funded pension systems means that “Second Pillar” employer-related pension benefits will inevitably become more uncertain too. Moreover, traditional family networks in which the younger members of a family were encouraged to take care of the older ones, are being broken down by the extended mobility of the workforce.

In a scenario of unknown longevity, retirees can reduce the risk of exhausting assets before passing away by consuming less per year, but such a tactic then increases the chance that they might die with too much wealth left unconsumed. In other words, dying with too little wealth is undesirable, but having too much wealth is also undesirable, since it represents foregone consumption opportunities. In this scenario, individuals will have to become more self-reliant and will wish to diversify their sources of income in retirement, assigning in particular a greater weight to private solutions, namely annuities. As a consequence, annuity providers will face an increasing longevity risk.

These trends in mortality lead to the use of projected survival models when pricing and reserving for life annuities and other long-term living benefits. A number of different projection models have been proposed and are actually used in actuarial practice. In spite of this, the future mortality trend is actually random

and hence, whatever kind of model is adopted, systematic deviations from the forecasted mortality may take place.

The risk of systematic deviations is different in nature from that of random fluctuations around the trend, a well-known type of risk in the insurance business, in both the life and the non-life insurance areas. Effectively, the risk of systematic deviations arises from either a “model” or a “parameter” risk, which are unquestionably non-pooling risks. Long-term trends observed in mortality affect both young and old ages. Although the terms “longevity risk” and “mortality risk” are very often used indistinctly, by longevity risk we mean the risk that members of some reference population might live longer, on average, than anticipated.

Longevity risk is a critical problem both because of the uncertainty of longevity projections, on one hand, and because of the huge amounts of liabilities at risk, on the other. Life annuities are probably the most important insurance product concerned by the longevity risk, but this risk should also be carefully considered when dealing with other long-term living benefits, insurance covers (particularly within the area of health insurance) and other retail products such as reverse mortgages. Additionally, many life insurers and reinsurers globally have become nervous about their exposure to catastrophic mortality events. This has resulted in the issuance of several bonds transferring mortality catastrophe risk to investors.

Despite its significance, traditionally life companies and pension sponsors were short of solutions to manage longevity risk. Until recently, hedging strategies were limited to product redesign (e.g., participating annuities), to the adoption of conservative pricing policies, to the actuarial management of an insurer’s surplus (internal capital) and, in some cases, to the use of prospective lifetables.

In order for a market for hedging of longevity risk to develop there are several prerequisites. These include the development of generally accepted technology and models to quantify the risk and the successful design and implementation of financial products and markets to hedge the risk. There has been a significant increase in research addressing these issues in recent years, namely those involving the securitization of risks, new capital market solutions and new reinsurance treaties.

In this paper, we assume that an appropriate description of the demographic risks requires the use of stochastic models. In particular, we assume that the random evolution of the stochastic force of mortality of an individual can be modelled by using doubly stochastic processes. The model is then embedded into the well know affine-jump framework, widely used in the term structure literature,

in order to derive closed-form solutions for the survival probability. We show that stochastic mortality models provide an adequate framework for the development of longevity risk hedging tools, namely mortality-linked contracts such as longevity bonds or mortality derivatives. The paper is organized as follows. In Section 2 we briefly review both the traditional “dynamic approach” and the new “stochastic mortality approach”. In Section 3 we use doubly stochastic processes in order to model the random evolution of the stochastic force of mortality in a manner that is common in the credit risk literature. The model is then embedded into the well known affine-jump term structure framework, widely used in the term structure or credit risk literature, in order to derive closed-form solutions for the survival probability. In Section 4 we calibrate the model to the Portuguese projected lifetables. Results indicate that the model is flexible enough to accommodate the rectangularization phenomena and that jumps seem to be an appropriate way to describe the random variations observed in mortality. Section 5 concludes.

2 Modeling mortality and longevity risk

One of the key conditions for the development of longevity-linked products and markets and for the hedging of longevity risk is the development of generally agreed market models for risk measurement. Whereas traditional market risks such as equity market, interest rate, exchange rate, credit and commodity risks have well consolidated methodologies for quantifying risk-based capital and for establishing market prices, longevity and mortality risk has historically been a very opaque risk. For a long time, only demographers, actuaries and insurance companies showed any interest in measuring and managing this risk, mainly for pricing purposes. A number of explanations can be given for this, particularly the fact that it is a non-financial risk that has been measured and analyzed in a different way from financial risks, generally adopting deterministic or scenario based approaches.

Historically, actuaries have been calculating premiums and mathematical reserves using a deterministic approach, by considering a deterministic mortality intensity, which is a function of the age only, extracted from available (static) lifetables and by setting a flat (“best estimate”) interest rate to discount cash flows over time. Since neither the mortality intensity nor interest rates are actually deterministic, life insurance companies are exposed to both financial and mortality (systematic and unsystematic) risks when pricing and reserving for any kind of long-term living benefits, particularly on annuities. In particular, the cal-

culuation of expected present values requires an appropriate mortality projection in order to avoid significant underestimation of future costs.

In order to protect the company from mortality improvements, actuaries have different solutions, among them to resort to projected (dynamic or prospective) lifetables, i.e., lifetables including a forecast of future trends of mortality instead of static lifetables. Static lifetables are obtained using data collected during a specific period (1 to 4 years) whereas dynamic lifetables incorporate mortality projections. To illustrate the problems with this approach, consider a female individual born in 2006. Her mother is 30-year-old and her grand-mother 60. To estimate the life expectancy of the newborn, the death probability at age 30 will be her mother's one and at age 60 her grand-mother's one, observed in 2006. This means that in a situation where longevity is increasing, static lifetables underestimate lifelengths and thus premiums relating to life insurance contracts. Conversely, dynamic lifetables will project mortality into the future accounting for longevity improvements.

Since the future mortality is actually unknown, there is enormous likelihood that future death rates will turn out to be different from the projected ones, and so a better assessment of longevity risk would be one that consists of both a mean estimate and a measure of uncertainty. Such assessment can only be performed using stochastic models to describe both demographic and financial risks. In this section, we briefly review both the traditional "dynamic approach" and the new "stochastic mortality approach".

2.1 Discrete-time dynamic approach

Although the subject of estimating future levels of mortality has received enormous attention lately, actuarial models of mortality and life tables for pricing and projecting pension and related life product cash flows have been developed over centuries. Tuljapurkar and Boe (1998), Tabeau (2001), GAD (2001), Pitacco (2004), Wong-Fupuy and Haberman (2004), Booth (2006) and Bravo (2007) provide a detailed review of historical patterns in mortality and longevity forecasting models.

In such a sensitive issue, there are a number of different opinions regarding how long people will live in the future. Some argue that lifespans will continue to increase at least as rapidly as experienced over the last decades due to, e.g., new medical breakthroughs or healthier life styles. Others disagree and project that increases in lifetime will decelerate, and potentially even decline (at least for certain risk groups), since any increase in longevity would have to occur by virtue

of declines in mortality for older age groups, or for other underlying causes of death. Other controversial points in this debate refer to the possible existence of a biological limit to human life and whether we are actually approaching it (see, e.g., Vaupel (1997), Olshansky and Carnes (1997) and Watson Wyatt (2005)).

The competing views on the evolution of lifespans, medical advancement or the existence of a biologic limit to human life translate into the question of how to appropriately model future longevity. Mortality forecasting methods currently in use can be categorized in many different ways. They can be clustered into epidemiological methods, projection by cause of death, extrapolative models, expert-opinion models and relational models.

Epidemiological models analyze the relationship between specific risk factors (e.g., smoking, obesity, socio-economic status, marital status and specific diseases) and their effects on mortality. The idea is to estimate the impact of specified risk factors on mortality rates (not on causes of death), from which mortality forecasts can be generated by projecting these risk factors into the future, given certain distributional assumptions. The practical usefulness of this approach lacks an accurate knowledge of the relationship between risk factors and mortality, something that science may achieve in the future. In relational models, future mortality rates are assumed to follow the dynamics of observed mortality for a more “advanced” population. The assumption is that the mortality profile of the forecasted population (e.g., of a developing country) will converge to the “target” population over some future time horizon.

Extrapolative models assume that future mortality can be estimated by projecting into the future the same trends observed in the recent to medium-term past. While the models can be either deterministic or stochastic, the basic idea is that future mortality will continue to improve at the same rate as in the past (observation window). This approach is the most popular among official bureaus all over the world but should be used with caution since it depends on the reliability of base mortality data and neglects, to a certain extent, the uncertainty related to the evolution of causes of death, future medical advances or to environmental risk factors. A variant of this approach are the methods that involve projection by cause of death, i.e., methods that disaggregate total mortality and forecast mortality rates for each cause of death separately by extrapolating past trends. Finally, expert-opinion models are also based on an extrapolative model, but explicitly include assumptions by the forecaster in respect to the future evolution of mortality. The idea is to incorporate additional information not captured by the statistical model ensuring that forecasted values are not pushed beyond

reasonable limits.

The classical approach to incorporating improvement in longevity for forecasting future mortality within extrapolative models is to fit an appropriate parametric function (e.g., Makeham model) to each calendar year data. Then, each of the parameter estimates is treated as independent time series, extrapolating their behaviour to the future in order to provide the actuary with projected lifetables (see, e.g., CMIB (1976) and Heligman e Pollard (1980)). Despite simple, this approach has serious limitations. In the first place, the approach strongly relies on the appropriateness of the parametric function adopted. Secondly, parameter estimates are very unstable, a feature that undermines the reliability of univariate extrapolations. Thirdly, the time series for parameter estimates are not independent and often robustly correlated. Although applying multivariate time series methods for the parameter estimates is theoretically possible, this will complicate the approach and introduce new problems.

Lee and Carter (1992) developed a simple model for describing the long term trends in mortality as a function of a simple time index. The method models the logarithm of a time series of age-specific death rates $\mu_x(t)$ as the sum of an age-specific component α_x , that is independent of time, and a second component, expressed as a product of a time-varying parameter denoting the general level of mortality κ_t , and an age-specific component β_x that signals the sensitiveness of mortality rates at each age when the general level of mortality changes. Formally, we have

$$\ln \mu_x(t) = \alpha_x + \beta_x \kappa_t + \epsilon_{x,t}, \quad (1)$$

where $\epsilon_{x,t} \sim \mathcal{N}(0, \sigma_\epsilon^2)$ is a white-noise, representing transitory shocks and the parameters α_x , β_x and κ_t have to be constrained by

$$\sum_{t=t_{\min}}^{t_{\max}} \kappa_t = 0 \quad \text{and} \quad \sum_{x=x_{\min}}^{x_{\max}} \beta_x = 1, \quad (2)$$

in order to ensure model identification.

Parameter estimates are obtained by ordinary least squares, i.e., by solving the following minimization program

$$(\hat{\alpha}_x, \hat{\beta}_x, \hat{\kappa}_t) = \arg \min_{\alpha_x, \beta_x, \kappa_t} \left\{ \sum_{x=x_{\min}}^{x_{\max}} \sum_{t=t_{\min}}^{t_{\max}} (\ln \mu_x(t) - \alpha_x - \beta_x \kappa_t)^2 \right\}. \quad (3)$$

Lee and Carter (1992) solve (3) by resorting to *Singular Value Decomposition* techniques but alternative estimation procedures can be implemented consider-

ing iterative methods (see, e.g., Bravo (2007)) or Weighted Least-Squares (see, e.g., Wilmoth (1993)). The resulting time-varying parameter estimates are then modeled and forecasted using standard Box-Jenkins time series methods. Finally, from this forecast of the general level of mortality, projected age-specific death rates are derived using the estimated age-specific parameters.

There have been several extensions to the Lee-Carter model including different error assumptions and estimation procedures. Lee (2000), Lee and Miller (2001), Tuljapurkar and Boe (1998), Brouhns *et al.* (2002a), Wong-Fillipoy and Haberman (2004), Bravo (2007) and Cairns *et al.* (2007) discuss the model and extensions. Brouhns *et al.* (2002a) and Renshaw and Haberman (2003c) develop an extension of the Lee-Carter model allowing for Poisson error assumptions and apply the model to Belgian data. This Poisson log-bilinear approach can be stated as

$$D_{x,t} \sim \text{Poisson}(\mu_x(t) E_{x,t}), \quad (4)$$

where $D_{x,t}$ denotes the number of deaths recorded at age x during year t , from an exposure-to-risk (i.e., the number of person-years from which $D_{x,t}$ arise), $E_{x,t}$, and $\mu_x(t)$ is given by (1).

One of the main advantages of the Poisson log-bilinear model over the Lee-Carter model is that specification (4) allows us to use maximum-likelihood methods to estimate the parameters instead of resorting to least squares (SVD) methods. Formally, we estimate the parameters α_x , β_x and κ_t by maximizing the log-likelihood derived from model (1)-(4)

$$\ln \mathcal{V}(\boldsymbol{\alpha}, \boldsymbol{\beta}, \boldsymbol{\kappa}) = \sum_{t=t_{\min}}^{t_{\max}} \sum_{x=x_{\min}}^{x_{\max}} \{D_{x,t}(\alpha_x + \beta_x \kappa_t) - E_{x,t} \exp(\alpha_x + \beta_x \kappa_t)\} + c, \quad (5)$$

where $\boldsymbol{\alpha} = (\alpha_{x_{\min}}, \dots, \alpha_{x_{\max}})'$, $\boldsymbol{\beta} = (\beta_{x_{\min}}, \dots, \beta_{x_{\max}})'$, $\boldsymbol{\kappa} = (\kappa_{t_{\min}}, \dots, \kappa_{t_{\max}})'$ and c is a constant.

The presence of the bilinear term $\beta_x \kappa_t$ makes it impossible to estimate the model using standard statistical packages that include Poisson regression. Because of this, we resort to an iterative method for estimating log-linear models with bilinear terms proposed by Goodman (1979). Even with a Poisson error assumption, heterogeneity by age-group in mortality indicates over-dispersion of errors.

Empirical studies to date suggest the need for more than a single factor to model longevity improvement, something that the Lee-Carter approach with a single factor and varying improvement impacts by age does not appear to capture.

This is important because if pricing models can often perform reasonably well with only a single factor, hedging requires a more complete picture of the dynamics of longevity improvements. In this sense, Bell (1997), Booth *et al.* (2002) and Renshaw and Haberman (2003c,d) include a second log-bilinear term in (1) and estimate parameters by considering the first two term in a SVD. Additionally, they adopt a multivariate setting in order to project the evolution of the time indices $\kappa_{t,i}$ ($i = 1, 2, \dots$).

Carter and Prskawetz (2001) consider the possibility of time varying parameters α_x and β_x . Renshaw and Haberman (2003a) include additional non-linear age factors when modeling the so-called “mortality reduction factors” within a Generalized Linear Models (GLM’s) approach. Renshaw and Haberman (2006) and Currie *et al.* (2004) include a cohort factor including year of birth as a factor impacting the rate of longevity improvement. This cohort factor is found to be significant in UK mortality data. Renshaw e Haberman (2005) and Bravo (2007) develop a version of the Lee-Carter model considering positive asymptotic mortality. This result is, for most age groups, more consistent with observed mortality patterns when compared with that of the original model. Wilmoth and Valkonen (2002) develop an extension of the Lee-Carter model aimed to investigate differential mortality by considering a number of alternative covariates other than age and calendar time.

Cairns, Blake and Dowd (2006b) develop and apply a two-factor model similar to the Lee-Carter model with a smoothing of age effects using a logit transformation of mortality rates. Cairns *et al.* (2007) analyze England and Wales and US mortality data showing that models that allow for an age effect, a quadratic age effect and a cohort effect fit the data best although the analysis of error distributions in these models revealed disappointing. De Jong and Tickle (2006) formulate the Lee-Carter model in a *state space framework*.

2.2 Stochastic mortality modeling

Models following the approach of Lee and Carter typically adapt discrete-time time series models to capture the random element in the stochastic development of mortality rates. Given the unknown nature of future mortality, some authors have recently developed models in a continuous-time framework by modeling mortality intensity as a stochastic process (see, e.g., Milevsky and Promislow (2001), Dahl (2004), Biffis and Millosovich (2004, 2006), Biffis (2005), Dahl and Møller (2005), Miltersen and Persson (2005), Cairns *et al.* (2006a), Schrager (2006), Bravo (2007) and references therein).

Modeling the mortality intensity as a stochastic process allows us to capture two of its more significant features: time dependency and uncertainty of the future development. Additionally, this framework provides a more accurate description of both premiums and liabilities of life insurance companies and contributes to a proper quantification of systematic mortality risk (also called longevity risk) faced by them. This framework and model application provides the theoretical foundation for financial pricing of longevity dependent financial claims and for the development of longevity risk hedging tools, namely mortality-linked contracts such as longevity bonds or other longevity-linked derivatives.

Up to now, a number of different stochastic mortality models have been proposed - for a detailed classification see Cairns *et al.* (2006a) and Bravo (2007). Most of these stochastic mortality models are *short rate mortality models*, i.e., they model the spot mortality rate $q_x(t)$, or the spot force of mortality $\mu_x(t)$. We can also find *forward mortality models*, i.e., approaches that model the dynamic of the forward mortality intensity $f_x^\mu(t, T)$, a *positive-mortality modeling framework* for the spot survival probability $p_x(t)$, in line with the term structure approach developed by Flesaker e Hughson (1996), Rogers (1997) and Rutkowski (1997), or market-models for the forward survival probability.

Milevsky and Promislow (2001) were the first to propose a stochastic “hazard rate” or force of mortality. With the intention of pricing guaranteed annuitization options in variable annuities, the authors demonstrate, first in a discrete-time framework, how to price and hedge a plain vanilla mortality option using a portfolio composed by zero coupon bonds, insurance contracts and endowment contracts. Moreover, they price the same option in a continuous-time risk-neutral framework assuming that the dynamics of the short interest rate and of the mortality intensity evolve independently over time according to a Cox-Ingersoll-Ross-process and a stochastic *mean reverting Brownian Gompertz-type* model, respectively.

Dahl (2004) develops a general stochastic model for the mortality intensity. The author derives partial differential equations for both the price at which some insurance contracts should be sold on the financial market and for the general mortality derivatives in the presence of stochastic mortality. In addition, he envisages solutions by which systematic mortality risk can be transferred to the financial market. Dahl and Moller (2005) derive risk-minimizing strategies for insurance liabilities in a market without derivative securities. Biffis and Millosovich (2004) expand this framework to a bidimensional setting in order to deal effectively with several sources of risk that simultaneously affect insurance contracts.

In Biffis (2005), affine jump-diffusion processes are used to model both finan-

cial and demographic factors. Specifications of the model with an affine term structure are employed and closed form mathematical expressions (up to the solutions of standard Riccati ordinary differential equations) are derived for some classic life insurance contracts. In Section ?? we illustrate this approach with a particular example.

Schrager (2006) presents an affine stochastic mortality model, that simultaneously describe the evolution of mortality for different age groups as opposed to the previous formulation in which a single cohort is considered. The author fits the model to Dutch mortality data using Kalman filters and presents alternative valuation approaches for a number of mortality-contingent contracts.

Biffis and Denuit (2005) and Biffis *et al.* (2006) generalize the model proposed by Lee and Carter (1992) to a stochastic setting. The authors assume that the dynamics of the time-varying parameter κ_t can be described by stochastic differential equations.

While most of the models presented so far assume independence between financial and demographic risk factors, Miltersen and Persson (2005) allow for correlations. The authors adopt the well know Heath-Jarrow-Morton no-arbitrage approach and model the forward mortality intensity (instead of the spot mortality intensity), taking the whole forward-mortality curve as an infinite-dimensional state variable. Similar to standard term structure literature, they derive no-arbitrage conditions for the drift term.

These models have generally been implemented for single age cohorts. To allow for multiple ages in the modeling, dependence across ages must be modeled in a proper way. Although these arbitrage-free models currently provide the most potential as a standard modeling framework for pricing and hedging longevity risk based products, there are a number of modeling issues that need to be addressed and that are yet to be fully explored. Important issues include the modeling of morbidity and ill-health, the use of multiple state models to capture the dependence between competing risk factors or incorporating cause of death as risk factors.

3 Affine-Jump diffusion processes for mortality

In this section we draw a parallel between insurance contracts and certain credit-sensitive securities and exploit some results of the intensity-based approach to credit risk modeling. Specifically, we use doubly stochastic processes (also known as *Cox processes*) in order to model the random evolution of the stochastic force

of mortality of an individual aged x in a manner that is common in the credit risk literature. The model is then embedded into the well know affine-jump term structure framework, widely used in the term structure literature, in order to derive closed-form solutions for the survival probability.

3.1 Mathematical framework

We are given a filtered probability space $(\Omega, \mathcal{F}, \mathbb{F}, \mathbb{P})$ and concentrate on an individual aged x at time 0. Following the pioneering work of Artzner and Delbaen (1995) in the credit risk literature and the proposals by Dahl (2004) and Biffis (2005) among others in the mortality area, we model his/her random lifetime as an \mathbb{F} -stopping time τ_x admitting a random intensity μ_x . Specifically, we consider τ_x as the first jump-time of a nonexplosive \mathbb{F} -counting process N recording at each time $t \geq 0$ whether the individual has died ($N_t \neq 0$) or survived ($N_t = 0$). The stopping time τ_x is said to admit an intensity μ_x if the compensator of N does, i.e., if μ_x is a nonnegative predictable process such that $\int_0^t \mu_x(s) ds < \infty$ for all $t \geq 0$ and such that the compensated process $M_t = \left\{ N_t - \int_0^t \mu_x(s) ds : t \geq 0 \right\}$ is a local \mathbb{F} -martingale. If the stronger condition $\mathbb{E} \left(\int_0^t \mu_x(s) ds \right) < \infty$ is satisfied, then M_t is an \mathbb{F} -martingale.

From this, we derive

$$\mathbb{E} (N_{t+\Delta t} - N_t | \mathcal{F}_t) = \mathbb{E} \left(\int_t^{t+\Delta t} \mu_x(s) ds \middle| \mathcal{F}_t \right), \quad (6)$$

based on which we can write

$$E (N_{t+\Delta t} - N_t | \mathcal{F}_t) = \mu_x(t) \Delta t + o(\Delta t), \quad (7)$$

an expression comparable with that of the instantaneous probability of death $\Delta t q_{x+t}$ derived in the traditional deterministic context.

By further assuming that N is a Cox (or doubly stochastic) process driven by a subfiltration \mathbb{G} of \mathbb{F} , with \mathbb{F} -predictable intensity μ it can be shown, by using the law of iterated expectations, that the probability of an individual aged $x+t$ at time t surviving up to time $T \geq t$, on the set $\{\tau > t\}$, is given by

$$\mathbb{P}(\tau > T | \mathcal{F}_t) = \mathbb{E} \left[e^{-\int_t^T \mu_{x+s}(s) ds} \middle| \mathcal{F}_t \right]. \quad (8)$$

Readers who are familiar with mathematical finance and, in particular, with the interest rate literature, can without difficulty observe that the right-hand-side of equation (8) represents the price at time t of a unitary default-free zero coupon

bond with maturity at time $T > t$, if the intensity μ is to represent the short-term interest rate.

One of the main advantages of this mathematical framework is that we can approach the survival probability (8) by using well known affine-jump diffusion processes. In particular, an \mathbb{R}^n -valued affine-jump diffusion process X is an \mathbb{F} -Markov process whose dynamics is given by

$$dX_t = \delta(t, X_t)dt + \sigma(t, X_t)dW_t + \sum_{h=1}^m dJ_t^h, \quad (9)$$

where W is a \mathbb{F} -standard Brownian motion in \mathbb{R}^n and each component J^h is a pure-jump process in \mathbb{R}^n with jump-arrival intensity $\{\eta^h(t, X_t) : t \geq 0\}$ and time-dependent jump distribution ν_t^h on \mathbb{R}^n . An important requirement of affine processes is that the drift $\delta : D \rightarrow \mathbb{R}^n$, the instantaneous covariance matrix $\sigma\sigma^T : D \rightarrow \mathbb{R}^{n \times n}$ and the jump-arrival intensity $\eta^h : D \rightarrow \mathbb{R}_+$ must all have an affine dependency on X . The jump-size distribution is determined by its Laplace transform.

The convenience of adopting affine processes in modeling the mortality intensity comes from the fact that, for any $a \in \mathbb{C}^n$, for given $T \geq t$ and an affine function R defined by $R(t, X) = \rho_0(t) + \rho_1(t) \cdot X$, under certain technical conditions we have

$$\phi^X(a, X_t, t, T) \doteq \mathbb{E} \left[e^{-\int_t^T R(s, X_s)ds} e^{a \cdot X_T} \middle| \mathcal{F}_t \right] = e^{\alpha(t) + \beta(t) \cdot X_t}, \quad (10)$$

where $\alpha(\cdot) \doteq \alpha(\cdot; a, T)$, $\beta(\cdot) \doteq \beta(\cdot; a, T)$ satisfy generalized Ricatti ordinary differential equations, that can be solved at least numerically and, in some cases, as we will see below, analytically.

3.2 Mortality intensity as a stochastic process

Turning now to the problem of modeling adequately the dynamics of mortality, we illustrate the approach developed in the previous section by developing a new model for the mortality intensity that considers the classic Feller equation together with a jump component. Formally, we assume that the mortality intensity $\mu_{x+t}(t)$ solves the following stochastic differential equation

$$\begin{aligned} d\mu_{x+t}(t) &= a\mu_{x+t}(t)dt + \sigma\sqrt{\mu_{x+t}(t)}dW(t) + dJ(t) \\ \mu_{x+t}(0) &= \bar{\mu}_x, \end{aligned} \quad (11)$$

with

$$J(t) = \sum_{i=1}^{N_t} \varepsilon_i. \quad (12)$$

where $\bar{\mu}_x > 0$, $a > 0$, $\sigma \geq 0$ and $W(t)$ is a standard Brownian motion.

We assume that $J(t)$ is a compound Poisson process, independent of W , with constant jump-arrival intensity $\eta \geq 0$, where $\{\varepsilon_i : i = 1, \dots, \infty\}$ are i.i.d. variables. Following the results by Kou (2002), among others, we consider jump sizes that are random variables double asymmetric exponentially distributed with density

$$f(z) = \pi_1 \left(\frac{1}{v_1} \right) e^{-\frac{z}{v_1}} \mathbb{I}_{\{z \geq 0\}} + \pi_2 \left(\frac{1}{v_2} \right) e^{\frac{z}{v_2}} \mathbb{I}_{\{z < 0\}} \quad (13)$$

where $\pi_1, \pi_2 \geq 0$, $\pi_1 + \pi_2 = 1$, represent, respectively, the probabilities of a positive (with average size $v_1 > 0$) and negative (with average size $v_2 > 0$) jump. By setting $\pi_1 = 0$ we are interested only on the importance of longevity risk (see, e.g., Biffis (2005)). By setting $\eta = 0$ the model becomes deterministic. When $v_1 = v_2$ and $\pi_1 = \pi_2 = \frac{1}{2}$ we get the so-called “first Laplace law”. By adopting equation (13) we consider the significance of both positive mortality shocks (e.g., new medical breakthroughs) and negative mortality shifts (e.g., bird flu).

In the spirit of (10), let us now assume that the survival probability $_{T-t}p_{x+t}(t)$ is represented by an exponentially affine function. By applying the framework described above, we have that

$$_{T-t}p_{x+t}(t) \equiv e^{\mathcal{A}(\tau) + \mathcal{B}(\tau) \cdot \mu_{x+t}(t)} \quad (14)$$

where $\tau = T - t$.

It can be shown that the solution to this problem admits the following Feynman-Kac representation

$$\begin{aligned} v(t, \mu_{x+t}(t)) \left\{ -\dot{\mathcal{A}}(\tau) - \dot{\mathcal{B}}(\tau) \mu_x(t) + a \mu_{x+t}(t) \mathcal{B}(\tau) + \frac{\sigma^2}{2} \mu_{x+t}(t) \mathcal{B}^2(\tau) \right. \\ \left. + \eta \left(\frac{\pi_1}{1 - v_1 \mathcal{B}(\tau)} + \frac{\pi_2}{1 + v_2 \mathcal{B}(\tau)} - 1 \right) - \mu_{x+t}(t) \right\} = 0, \end{aligned} \quad (15)$$

where $v(t, \mu_{x+t}(t)) = _{T-t}p_{x+t}(t)$.

Dividing both sides of this equation by $v(t, \mu_{x+t}(t))$ we get

$$\begin{aligned} \left[-\dot{\mathcal{B}}(\tau) + a \mathcal{B}(\tau) + \frac{\sigma^2}{2} \mathcal{B}^2(\tau) - 1 \right] \mu_{x+t}(t) \\ + \left[-\dot{\mathcal{A}}(\tau) + \eta \left(\frac{\pi_1}{1 - v_1 \mathcal{B}(\tau)} + \frac{\pi_2}{1 + v_2 \mathcal{B}(\tau)} - 1 \right) \right] = 0, \end{aligned} \quad (16)$$

where $A(\tau)$ and $B(\tau)$ are solutions to the following system of ODEs'

$$\dot{\mathcal{B}}(\tau) = a\mathcal{B}(\tau) + \frac{1}{2}\sigma^2\mathcal{B}^2(\tau) - 1 \quad (17)$$

$$\dot{\mathcal{A}}(\tau) = \eta \left(\frac{\pi_1}{1 - v_1\mathcal{B}(\tau)} + \frac{\pi_2}{1 + v_2\mathcal{B}(\tau)} - 1 \right) \quad (18)$$

with boundary conditions

$$\mathcal{B}(0) = 0, \mathcal{A}(0) = 0. \quad (19)$$

where $\dot{\mathcal{B}}(\tau) = \frac{\partial}{\partial \tau}\mathcal{B}(\tau)$, $\dot{\mathcal{A}}(\tau) = \frac{\partial}{\partial \tau}\mathcal{A}(\tau)$.

By solving the system (17)-(18)-(19), we get the following closed-form solutions for $\mathcal{A}(\tau)$ and $\mathcal{B}(\tau)$

$$\begin{aligned} \mathcal{A}(\tau) = & \eta\pi_1 \left\{ \frac{\alpha_0\tau}{(\alpha_0 - v_1)} + \frac{v_1(\alpha_0 + \alpha_1) [\ln(\alpha_0 + \alpha_1) - \ln(\alpha_0 - v_1 + (\alpha_1 + v_1)e^{\kappa\tau})]}{\kappa(\alpha_0 - v_1)(\alpha_1 + v_1)} \right\} \\ & + \eta\pi_2 \left\{ \frac{\alpha_0\tau}{(\alpha_0 + v_2)} + \frac{v_2(\alpha_0 + \alpha_1)}{\kappa(\alpha_1 - v_2)(\alpha_0 + v_2)} [-\ln(\alpha_0 + \alpha_1) \right. \\ & \left. + \ln(\alpha_0 + v_2 + (\alpha_1 - v_2)e^{\kappa\tau})] \right\} - \eta\tau \end{aligned} \quad (20)$$

$$\mathcal{B}(\tau) = \frac{1 - e^{\kappa\tau}}{\alpha_0 + \alpha_1 e^{\kappa\tau}} \quad (21)$$

with $\kappa = \sqrt{a^2 + 2\sigma^2}$, $\alpha_0 = \frac{(a+\kappa)}{2}$ and $\alpha_1 = \frac{(\kappa-a)}{2}$, defined for

$$-\frac{1}{v_2} < \mathcal{B}(\tau) < \frac{1}{v_1}. \quad (22)$$

We observe that the model stipulates an increasing (deterministic) trend for the mortality intensity, around which random fluctuations occur due to the stochastic component and due to the jump component. Additionally, the model offers a realistic process for the stochastic mortality rate since it ensures that the variable cannot take negative values. The model assumes that both negative and positive jumps can be registered in mortality, a feature that contrasts with similar models that are interested in sudden improvements in mortality (e.g., due to medical advances) only. We think this gives a more appropriate description of mortality, in which unexpected increases in mortality can occur (e.g., caused by natural catastrophes or epidemics). The model offers a nice analytical solution, easy to use in pricing and reserving applications within the life insurance industry.

4 Calibration to the Portuguese projected lifetables

As a first application of the above models, we have calibrated model (11) to the Portuguese projected lifetables. Portuguese projected lifetables were obtained by fitting model (4)-(1)-(2) to a matrix of crude Portuguese death rates, from year 1970 to 2004 and for ages 0 to 84. The data, discriminated by age and sex, refers to the entire Portuguese population and has been supplied by the Portuguese National Institute of Statistics (INE - Instituto Nacional de Estatística). The database used comprised two elements: the observed number of death $d_{x,t}$ given by age and year of death, and the observed population size $l_{x,t}$ at December 31 of each year. We follow the INE definition of population at risk using the population counts at the beginning and at the end of a year and take migration into account. The Poisson parameters α_x , β_x and κ_t implicated in (1) are estimated by maximum-likelihood methods using the iterative procedure described in Section 2.1.¹

The closing of lifetables was performed using the method proposed by De-nuit and Goderniaux (2005) to extrapolate mortality rates at very old ages. The method is a two step method. First, a quadratic function is fitted to age-specific estimated mortality rates in a given age-band. Second, the estimated function is used to extrapolate mortality rates up to a pre-determined maximum age. Formally, the following log-quadratic model is fitted by weighted least-squares

$$\ln \hat{q}_x(t) = a(t) + b(t)x + c(t)x^2 + \epsilon_x(t), \quad x \in [65, 84] \quad (23)$$

to age-specific mortality rates observed at older ages (in our case $x \in [65, 84]$), where $\epsilon_x(t) \sim \mathcal{N}(0, \sigma^2(t))$, with additional constraints

$$q_{x_{\max}} = 1 \quad (24)$$

$$q'_{x_{\max}} = 0 \quad (25)$$

where q'_x denotes the first derivative of q_x with respect to age x_{\max} . Constraints (24) and (25) impose a concave configuration to the curve of mortality rates at old ages and the existence of a horizontal tangent at $x = x_{\max} = 120$. We then use this function to extrapolated mortality rates up to age x_{\max} .

In fitting the model, we have adopted the ordinary least squares method, i.e., we minimize the quadratic deviations between the model survival probabilities, ${}_{T-t}p_{65}^{\text{model}}(t)$, and the prospective lifetable ones, ${}_{T-t}p_{65}^{\text{TP}}(t)$ for an individual aged

¹For more details see Bravo (2007).

65. Formally, parameter estimates Θ solve the following optimization problem

$$\hat{\Theta} = \arg \min_{\Theta} \left\{ Q^2 = \sum_{T=t+1}^{t+(x_{\max}-65)+1} \left({}_{T-t}p_{65}^{\text{model}}(t) - {}_{T-t}p_{65}^{\text{TP}}(t) \right)^2 \right\} \quad (26)$$

where $x_{\max} = 120$ and $t \in \{1970, 1980, 1990, 2004\}$.

Table 1 reports the optimal values of the parameters, the calibration error and the initial value of $\mu_{x+t}(t)$, $\mu_{65}(t)$, chosen to be equal to $-\ln(p_{65}(t))$, for both male and female populations. Figure 1 report, for the generations aged 65 in $t \in [1970, 2004]$, the survival function of the stochastic process analysed and of the prospective lifetable one.

Male				
	$t = 1970$	$t = 1980$	$t = 1990$	$t = 2004$
$\mu_{65}(t)$	0.02765901	0.02774125	0.02558451	0.01689187
a	0.09516212	0.09033169	0.08739382	0.09949474
σ	0.00001013	0.00001131	0.00000981	0.00000978
η	0.0117887	0.03936915	0.06544481	0.05226689
v_1	0.02654017	0.02876439	0.02726195	0.02757463
v_2	0.001128449	0.0001023	0.0001102	0.00009724921
Q^2	0.000483312	0.001135141	0.00423265	0.007431117
Female				
	$t = 1970$	$t = 1980$	$t = 1990$	$t = 2004$
$\mu_{65}(t)$	0.01472793	0.01375416	0.01163745	0.007780187
a	0.1119171	0.1096041	0.1101916	0.1199389
σ	0.00001044	0.00001033	0.00001082	0.00001049
η	0.01180289	0.03190069	0.05536174	0.05693019
v_1	0.0284391	0.02890383	0.02727089	0.02644525
v_2	0.0001189	0.0001098	0.0001072	0.0001066
Q^2	0.0003536312	0.0007131984	0.003482145	0.006155311

Table 1: Parameter estimates

The calibration error is quite small and the parameter estimates show that the value of σ is very low, particularly when compared with that of both positive and negative average size jumps. We can observe that the fit is very good, even when we consider the importance of the rectangularization phenomena, highly significant in the 2004 generation. The results also suggest that jumps seem to be an appropriate way to describe the random variations observed in mortality.

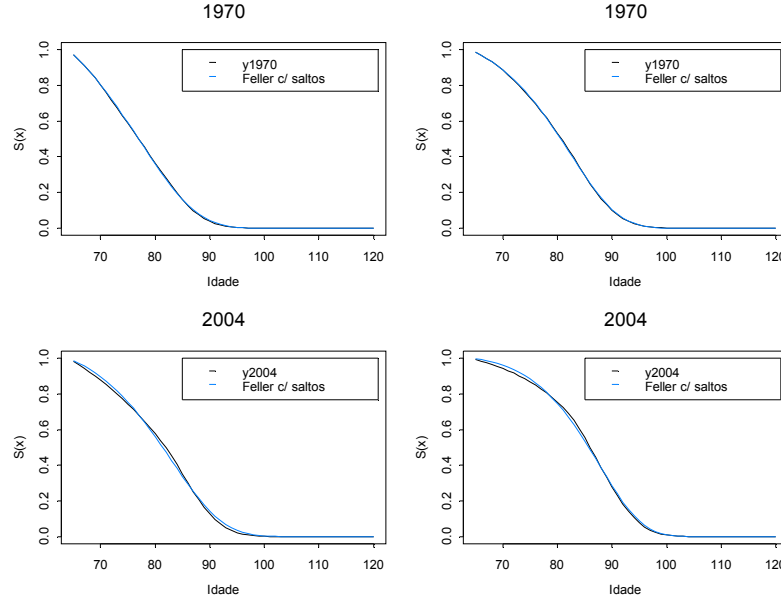


Figure 1: Survival probability $_{T-t}p_{65}(t)$ as a function of age $x + T - t$ for $t = 1970$ and $t = 2004$ (the left panel corresponds to the male population)

5 Conclusion

In this paper, we have reviewed both the traditional discrete-time dynamic approach to mortality projection and the new “stochastic mortality approach” to mortality and longevity risk modeling. We have describe the random evolution of mortality by using doubly stochastic processes. The intensity is then described as an affine-jump diffusion process, considering jump sizes that are random variables double asymmetric exponentially distributed. The model is compatible with both both negative and positive jumps in mortality, a feature that contrasts with similar models that are interested in sudden improvements in mortality (e.g., due to medical advances) only. Survival probabilities have been provided in closed-form. The intensity process has been calibrated to the Portuguese population using projected lifetables built using the Poisson Lee-Carter method. The results show that fit is very good and that the model is flexible enough to accommodate some of the traditional demographic phenomena.

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