

A matrix approach to the statistics of longevity in the gamma-Gompertz and related mortality models^{*†}

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1 Abstract

2 **BACKGROUND:** The gamma-Gompertz model is a fixed frailty model in which baseline mor-
3 tality increases exponentially with age, frailty has a proportional effect on mortality, and frailty
4 at birth follows a gamma distribution. Mortality selects against the more frail, so the marginal
5 mortality rate decelerates, eventually reaching an asymptote. The gamma-Gompertz is one of a
6 wider class of frailty models, characterized by the choice of baseline mortality, effects of frailty,
7 distributions of frailty, and assumptions about the dynamics of frailty.

8 **OBJECTIVES:** To develop a matrix model to compute all the statistical properties of longevity
9 from the gamma-Gompertz and related models.

10 **METHODS:** I develop a matrix version of the gamma-Gompertz model using the vec-permutation
11 matrix formulation for age-stage models.

12 **RESULTS:** The model permits calculation of the mean, variance, coefficient of variation, skew-
13 ness and all moments of longevity, the marginal mortality and survivorship functions, the dynamics
14 of the frailty distribution, and other quantities. The matrix formulation extends naturally to other
15 frailty models. I apply the analysis to the gamma-Gompertz model (for humans and laboratory
16 animals), the gamma-Makeham model, and the gamma-Siler model, and to a hypothetical dynamic
17 frailty model characterized by diffusion of frailty with reflecting boundaries.

18 The matrix model permits partitioning the variance in longevity into components due to het-
19 erogeneity and to individual stochasticity. In several published human data sets, heterogeneity
20 accounts for less than 10% of the variance in longevity. In laboratory populations of five inverte-
21 brate animal species, heterogeneity accounts for 46% to 83% of the total variance in longevity.

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1 Introduction

The gamma-Gompertz (G-G) model is one of a class of models that investigate the effects on survival and longevity of cryptic heterogeneity — heterogeneity that is either unobservable or unobserved — in mortality (e.g. ??????). The goal of this paper is to present a matrix formulation that permits easy computation of all the properties of the G-G model, and of other related models for heterogeneity in mortality.

The G-G model specifies a baseline age-specific mortality rate and modifies this baseline by a factor, called frailty, that reflects the heterogeneity among individuals. The baseline mortality function is the Gompertz model, in which mortality increases exponentially with age t ,

$$\mu(t) = ae^{bt}. \tag{1}$$

Frailty is introduced as a proportional hazard multiplier z ; the mortality of an individual with frailty z at age t is

$$\mu(z, t) = z\mu_0(t) \tag{2}$$

where $\mu_0(t)$ is the baseline mortality schedule.

The dynamics at any age of a cohort subject to such a mortality model depend on the distribution of frailty. Let that distribution at age t be $\pi(z, t)$. Because frailty is cryptic, observations on the cohort reveal not the individual mortality schedules, but rather the marginal mortality rate

$$\mu^*(t) = \int \pi(z, t)\mu(z, t)dz \tag{3}$$

The survivorship function of an individual of frailty z is

$$S(z, t) = \exp\left(-\int_0^t \mu(z, x)dx\right) \tag{4}$$

$$= S_0(t)^z \tag{5}$$

where $S_0(t)$ is the survivorship resulting from the baseline mortality schedule $\mu_0(t)$. The marginal survivorship function is

$$S^*(t) = \int \pi(z, 0)S(z, t)dz. \tag{6}$$

The dynamics of the cohort differ from the dynamics of any frailty class because the distribution of frailty changes as the cohort ages (?). The more frail individuals tend to die sooner, and the cohort is progressively dominated by individuals of lower frailty. The distribution of frailty is

$$\pi(z, t) = \frac{\pi(z, 0)S(z, t)}{\int \pi(z, 0)S(z, t)dz}. \tag{7}$$

In the G-G model, frailty is a fixed property of an individual, and the cohort begins life with frailty distributed according to a gamma distribution

$$z \sim \text{gamma}(k, \lambda) \tag{8}$$

with shape parameter k and scale parameter¹ λ . The mean and variance of z are $E(z) = k/\lambda$ and $V(z) = k/\lambda^2$. Thus, when $E(z) = 1$, the distribution is given by $\text{gamma}(1/\sigma^2, 1/\sigma^2)$.

¹The probability density function is

$$\text{gamma}(k, \lambda) = \frac{1}{\Gamma(k)\lambda^k} z^{k-1}e^{-z/\lambda}. \tag{9}$$

77 The marginal mortality rate (3) for the G-G model is a sigmoid function of age,

$$\mu^*(t) = \frac{ae^{bt}}{1 + \frac{a\sigma^2}{b}(e^{bt} - 1)} \quad (10)$$

78 (?), converging to an asymptote at b/σ^2 as t gets large. Thus, the G-G model is an attractive
 79 explanation for the widely observed pattern of decelerating increase in mortality with age, in both
 80 humans and other species (e.g., ????).

81 Although it is simple to state, and widely used, deriving the consequences of the G-G model
 82 is mathematically challenging. ? has recently obtained an expression for life expectancy at birth,
 83 by integrating the survivorship function (24). The result, a function of the Gompertz parameters
 84 a and b and the gamma distribution parameters k and λ , is written in terms of the Gaussian
 85 hypergeometric² function ${}_2F_1$:

$$e_0(a, b, k, \lambda) = \frac{1}{bk} {}_2F_1\left(k, 1; k + 1; 1 - \frac{1}{b\lambda}\right) \quad (11)$$

86 Life expectancy, however, is only one of many demographic properties implied by a mortality
 87 model. My goal here is to present a matrix formulation that provides all the moments of longevity,
 88 various measures of life disparity, and the full dynamics of the joint distribution of age and frailty.
 89 It will become apparent from the construction of this model that it applies equally to a much
 90 broader class of frailty models, and I will present examples.

91 Section 2 derives the matrix model, using methods developed for cases in which individuals
 92 are jointly classified by age and stage. Section 3 derives the fundamental matrix, the moments
 93 of longevity, the distribution of age at death, and other indices from the matrix model including
 94 (Section 3.4) the dynamics of the frailty distribution over the life of the cohort. Section 4 analyzes
 95 an example from ?. Section 5 discusses some interesting generalizations and explores several other
 96 examples.

97 2 The matrix formulation of the gamma-Gompertz model

98 **Notation.** In what follows, matrices are denoted by upper-case boldface letters, and vectors by
 99 lower-case boldface letters. Where necessary, the dimensions of matrices and vectors are denoted
 100 by subscripts; thus \mathbf{I}_n is an identity matrix of order n and $\mathbf{1}_n$ is a $n \times 1$ vector of ones. The vector
 101 \mathbf{e}_i is the i th unit vector. The symbol \circ denotes the Hadamard, or element-by-element product. The
 102 symbol $\|\mathbf{x}\|$ denotes the 1-norm of the vector \mathbf{x} . The number of age classes is ω and the number of
 103 frailty groups is g .

104 The matrix G-G model is an age-stage classified model in which stages correspond to frailty
 105 classes. Age-stage classified matrix models have been analyzed in other contexts by ??? and ?. The
 106 model is created using the vec-permutation formalism (?) and analyzed using absorbing Markov
 107 chain theory (???)

Note that two parameterizations of the gamma distribution are widely used. Equation (9) is common in demography. MATLAB uses the parameterization

$$\text{gamma}(k, \theta)$$

where k is a shape parameter and θ is a rate parameter. In this parameterization, $E(z) = k\theta$ and $V(z) = k\theta^2$. Thus in MATLAB the distribution with mean equal to 1 is gamma $(1/\sigma^2, \sigma^2)$.

²For more on the Gaussian hypergeometric distribution, see ? Abramowitz and Stegun (1965, 15.1.1), or the online version in the NIST Digital Library of Mathematical Functions, <http://dlmf.nist.gov/15>.

108 To construct the matrix G-G model, let us introduce some notation. Age is described by a
 109 set of discrete age classes $1, \dots, \omega$. The baseline mortality rates are contained in a vector $\boldsymbol{\mu}_0$ of
 110 dimension $\omega \times 1$. For the Gompertz mortality model, the baseline mortality rate vector is

$$\boldsymbol{\mu}_0 = a \begin{pmatrix} e^{0b} \\ e^{1b} \\ e^{2b} \\ \vdots \\ e^{(\omega-1)b} \end{pmatrix} \quad (12)$$

111 Frailty is described by a set of g discrete frailty classes; the frailty values of these classes are
 112 given by a $g \times 1$ vector \mathbf{z} . To create the frailty classes, first specify a maximum frailty, where the
 113 cumulative gamma distribution reaches some high value; say, 0.9999. Since very high values of
 114 frailty are rapidly eliminated, this end of the distribution is, in practice, not very important. Then
 115 specify a minimum value of frailty as some very small number; for the applications reported here,
 116 a value on the order of 10^{-7} was adequate. Since individuals with very low frailty will persist for
 117 a long time in the population, it is important that z_{\min} be small.

118 Experience suggests that logarithmically spaced values between z_{\min} and z_{\max} work well, because
 119 they provide more detail in the frailty distribution at the low end, precisely where individuals will
 120 persist the longest. An alternative is to evenly divide the inverse of the cumulative distribution
 121 function, so that values are most closely spaced where the concentration of initial probability is
 122 greatest. Given the vector \mathbf{z} of frailty classes, the vector of mortality rates by age, for frailty class
 123 i , is

$$\boldsymbol{\mu}_i = z_i \boldsymbol{\mu}_0 \quad (13)$$

124 The distribution of individuals among frailty classes at age t is given by the vector $\boldsymbol{\pi}(t)$; the
 125 initial frailty distribution of the cohort is given by $\boldsymbol{\pi}(0)$. In the matrix G-G model, $\boldsymbol{\pi}(0)$ is a discrete
 126 gamma distribution with mean of 1 and a specified variance.

127 2.1 Constructing transition matrices and the vec-permutation model

128 To construct the age-stage model, define a survival matrix for each frailty class, and a matrix of
 129 frailty transitions for each age class, as follows.

- 130 1. Create a survival matrix \mathbf{U}_i for each frailty class i . It contains survival probabilities on the
 131 first subdiagonal and zeros elsewhere, and is of dimension $\omega \times \omega$.

$$\mathbf{U}_i = \begin{pmatrix} 0 & 0 & \dots & 0 \\ e^{-\mu(z_i,0)} & 0 & \dots & 0 \\ \vdots & \ddots & & \vdots \\ 0 & \dots & e^{-\mu(z_i,\omega-1)} & 0 \end{pmatrix} \quad (14)$$

- 132 2. Create a matrix \mathbf{D}_j describing transitions among frailty classes for each age class j . In the
 133 gamma-Gompertz model, frailty does not change, so $\mathbf{D}_j = \mathbf{I}_g$ for all j .
- 134 3. Create block-diagonal matrices \mathbf{U} and \mathbb{D} by placing the \mathbf{U}_i (respectively, \mathbf{D}_j) on the diagonal
 135 with zeros elsewhere. Both matrices are of dimension $\omega g \times \omega g$.

$$\mathbf{U} = \begin{pmatrix} \mathbf{U}_1 & \dots & 0 \\ \vdots & \ddots & \vdots \\ 0 & \dots & \mathbf{U}_g \end{pmatrix} \quad \mathbb{D} = \begin{pmatrix} \mathbf{D}_1 & \dots & 0 \\ \vdots & \ddots & \vdots \\ 0 & \dots & \mathbf{D}_g \end{pmatrix} \quad (15)$$

136 In the gamma-Gompertz model, $\mathbb{D} = \mathbf{I}_{\omega g}$.

137 The state of the cohort at age t is given by a vector $\tilde{\mathbf{n}}(t)$, which is derived from the array

$$\mathcal{N}(t) = \begin{pmatrix} n_{11} & \cdots & n_{1g} \\ \vdots & & \vdots \\ n_{\omega 1} & \cdots & n_{\omega g} \end{pmatrix} \quad (16)$$

138 that describes the abundance of all age-frailty categories. The population vector is

$$\tilde{\mathbf{n}} = \text{vec } \mathcal{N}^\top; \quad (17)$$

139 that is,

$$\tilde{\mathbf{n}} = \begin{pmatrix} n_{11} \\ \vdots \\ n_{1g} \\ \vdots \\ n_{\omega 1} \\ \vdots \\ n_{\omega g} \end{pmatrix} \quad (18)$$

140 The i th block of entries in $\tilde{\mathbf{n}}$ contains a sub-vector giving the abundance of the frailty classes within
141 age class i .

142 The joint age-frailty composition of the cohort is projected as

$$\tilde{\mathbf{n}}(t+1) = \tilde{\mathbf{U}}\tilde{\mathbf{n}}(t) \quad (19)$$

143 where the projection matrix is

$$\tilde{\mathbf{U}} = \mathbb{D}\mathbf{K}\mathbf{U}\mathbf{K}^\top \quad (20)$$

144 In equation (20), \mathbf{K} (to be more precise, $\mathbf{K}_{\omega, g}$) is the vec-permutation matrix, or commutation
145 matrix (??); see ? for a demographic description.

146 Because frailty is fixed in the gamma-Gompertz model, $\tilde{\mathbf{U}}$ reduces to $\tilde{\mathbf{U}} = \mathbf{K}\mathbf{U}\mathbf{K}^\top$. However, it
147 is good practice to retain the matrix \mathbb{D} as a reminder of its potential use when frailty is dynamic
148 rather than static.

149 2.2 The absorbing Markov chain

150 The matrix $\tilde{\mathbf{U}}$ is the transient matrix of an absorbing Markov chain ??????. The transition matrix
151 of this chain is

$$\mathbf{P} = \left(\begin{array}{c|c} \tilde{\mathbf{U}} & \mathbf{0} \\ \hline \mathbf{M} & \mathbf{I} \end{array} \right) \quad (21)$$

152 where \mathbf{M} is a mortality matrix describing the transitions from transient (i.e., living) states to
153 absorbing (i.e., dead) states. The fundamental matrix of the chain defined by $\tilde{\mathbf{U}}$ is

$$\tilde{\mathbf{N}} = \left(\mathbf{I}_{\omega g} - \tilde{\mathbf{U}} \right)^{-1} \quad (22)$$

154 with dimension $\omega g \times \omega g$. The (i, j) entry of $\tilde{\mathbf{N}}$ is the expected number of visits to state i , conditional
155 on starting in a state j , where states describe the full joint age \times frailty distribution. From the
156 fundamental matrix we can compute all the statistics of the cohort survival properties. We turn
157 now to these analyses.

158 **3 Analysis of the model**

159 The fundamental matrix $\tilde{\mathbf{N}}$ of the joint chain contains all the information necessary to derive
 160 the marginal survival function \mathbf{s}^* (a vector of dimension $\omega \times 1$) and the corresponding marginal
 161 fundamental matrix \mathbf{N}^* (of dimension $\omega \times \omega$).

162 **3.1 The marginal survival function**

163 To obtain the marginal dynamics of the cohort age distribution, first average the columns of $\tilde{\mathbf{N}}$ over
 164 the initial frailty distribution. If, as in the gamma-Gompertz case, the initial frailty distribution
 165 has positive support only in the first age class, the result is

$$\tilde{\mathbf{s}} = \tilde{\mathbf{N}} [\mathbf{e}_1 \otimes \boldsymbol{\pi}(0)].$$

166 This column vector gives the average, over $\boldsymbol{\pi}(0)$, of the number of visits to each age-frailty state
 167 by an individual in the first age class.

168 Next, sum the rows of the vector $\tilde{\mathbf{s}}$ within each frailty class to obtain the marginal mean number
 169 of visits to each age class for an individual in the initial cohort. Because the underlying demographic
 170 model is age-classified, a transient state (i.e., an age class) *can be visited at most once; hence the*
 171 *mean number of visits is the probability of visiting.* Thus the vector of mean number of visits is in
 172 fact the marginal survivorship function \mathbf{s}^* :

$$\mathbf{s}^* = (\mathbf{I}_\omega \otimes \mathbf{1}_g^\top) \tilde{\mathbf{s}} \tag{23}$$

$$= (\mathbf{I}_\omega \otimes \mathbf{1}_g^\top) \tilde{\mathbf{N}} [\mathbf{e}_1 \otimes \boldsymbol{\pi}(0)] \tag{24}$$

173 **3.2 The marginal fundamental matrix**

174 The fundamental matrix $\tilde{\mathbf{N}}$ gives the number of visits to each age-frailty class. We need the
 175 marginal fundamental matrix \mathbf{N}^* , which gives the expected number of visits to each age class. All
 176 the statistics of longevity can be obtained from this matrix (e.g., ?????). To obtain \mathbf{N}^* , note that
 177 the vector \mathbf{s}^* is the first column of \mathbf{N}^* , and that the full matrix is

$$\mathbf{N}^* = \begin{pmatrix} s_1^* & 0 & 0 & \cdots & 0 \\ s_2^* & \frac{s_2^*}{s_2^*} & 0 & \cdots & 0 \\ s_3^* & \frac{s_3^*}{s_2^*} & \frac{s_3^*}{s_3^*} & \cdots & 0 \\ s_4^* & \frac{s_4^*}{s_2^*} & \frac{s_4^*}{s_3^*} & \cdots & 0 \\ \vdots & \vdots & \vdots & \cdots & 1 \end{pmatrix} \tag{25}$$

178 (? , Eq. 10.5.4)³ This can be written

$$\mathbf{N}^* = [\mathbf{s}^* \mathbf{1}_s^\top \text{diag} (\mathbf{s}^*)^{-1}] \circ \mathbf{Y} \tag{26}$$

179 where \mathbf{Y} is a lower triangular matrix with ones on and below the diagonal and zeros elsewhere.

³It seems appropriate to note that Keyfitz presented this result in the first edition of the book in 1977.

180 3.3 Longevity, variation, and disparity

181 When longevity statistics are calculated from \mathbf{N}^* , they give the marginal results for the cohort
 182 starting with the initial frailty distribution $\boldsymbol{\pi}(0)$. These include the following.

- 183 1. The moments of the number of visits to each of the transient states. Because transient states
 184 (age classes) in an age-classified model can be visited no more than once, these moments may
 185 be less interesting in the G-G model than in models with more complicated stage structure.
 186 Letting \mathbf{N}_i^* be the matrix of the i th moments of the number of visits, \mathbf{N}_1^* is given by (25),
 187 and the higher moments include

$$\mathbf{N}_2^* = (2\mathbf{N}_{\text{dg}}^* - \mathbf{I}) \mathbf{N}_1^* \quad (27)$$

$$\mathbf{N}_3^* = \left[6 (\mathbf{N}_{\text{dg}}^*)^2 - 6\mathbf{N}_{\text{dg}}^* + \mathbf{I} \right] \mathbf{N}_1^* \quad (28)$$

$$\mathbf{N}_4^* = \left[24 (\mathbf{N}_{\text{dg}}^*)^3 - 36 (\mathbf{N}_{\text{dg}}^*)^2 + 14\mathbf{N}_{\text{dg}}^* - \mathbf{I} \right] \mathbf{N}_1^*. \quad (29)$$

188 where \mathbf{N}_{dg}^* is a diagonal matrix with the diagonal elements of \mathbf{N}_1^* on the diagonal and zeros
 189 elsewhere (e.g., ???); for a mathematical source see ?.

- 190 2. The moments and statistics of longevity. Longevity is equivalent to the time until absorbtion
 191 in one of the absorbing states. The vector $\boldsymbol{\eta}_1$ of mean longevities (i.e., life expectancies) of
 192 each age class is obtained from the column sums of \mathbf{N}^* , and subsequent moments as follows,
 193 where $\boldsymbol{\eta}_i$ is the vector of i th moments of longevity:

$$\boldsymbol{\eta}_1^\top = \mathbf{1}_\omega^\top \mathbf{N}^* \quad (30)$$

$$\boldsymbol{\eta}_2^\top = \boldsymbol{\eta}_1^\top (2\mathbf{N}^* - \mathbf{I}) \quad (31)$$

$$\boldsymbol{\eta}_3^\top = \boldsymbol{\eta}_1^\top \left[6 (\mathbf{N}^*)^2 - 6\mathbf{N}^* + \mathbf{I} \right] \quad (32)$$

$$\boldsymbol{\eta}_4^\top = \boldsymbol{\eta}_1^\top \left[24 (\mathbf{N}^*)^3 - 36 (\mathbf{N}^*)^2 + 14\mathbf{N}^* - \mathbf{I} \right]. \quad (33)$$

194 (???). These moments provide a complete set of longevity statistics, including the variance,
 195 standard deviation, coefficient of variation, and skewness of longevity:

$$V(\boldsymbol{\eta})^\top = \boldsymbol{\eta}_2 - \boldsymbol{\eta}_1 \circ \boldsymbol{\eta}_1 \quad (34)$$

$$SD(\boldsymbol{\eta}) = \sqrt{V(\boldsymbol{\eta})} \quad (35)$$

$$CV(\boldsymbol{\eta}) = \text{diag}(\boldsymbol{\eta}_1) SD(\boldsymbol{\eta}) \quad (36)$$

$$Sk(\boldsymbol{\eta}) = \text{diag}(V(\boldsymbol{\eta}))^{-3/2} [\boldsymbol{\eta}_3 - 3\boldsymbol{\eta}_1 \circ \boldsymbol{\eta}_2 + 2\boldsymbol{\eta}_1 \circ \boldsymbol{\eta}_1 \circ \boldsymbol{\eta}_1] \quad (37)$$

- 196 3. The joint and marginal distributions of age and stage at death. Becasue the matrix G-G model
 197 is an age-stage structured model, the joint and marginal distributions of age and frailty class
 198 at death are obtained using the mortality matrix $\tilde{\mathbf{M}}$ in (21).

199 If $\tilde{\mathbf{M}}$ is created by defining absorbing states corresponding to the age and frailty class at
 200 death, then $\tilde{\mathbf{M}} = \text{diag}(\mathbf{1}_{\omega g}^\top - \mathbf{1}_{\omega g}^\top \tilde{\mathbf{U}})$. Then column j of the matrix

$$\tilde{\mathbf{B}} = \tilde{\mathbf{M}} \tilde{\mathbf{N}} \quad (38)$$

201 gives the joint distribution of age and frailty at death, conditional on reaching the age-frailty
 202 combination in column j (?).

203 Averaging the first g columns of $\tilde{\mathbf{B}}$ over the initial frailty distribution gives a vector $\tilde{\phi}$ con-
 204 taining the distribution of age and frailty at death of a cohort with initial frailty distribution
 205 $\boldsymbol{\pi}(0)$:

$$\tilde{\phi} = \tilde{\mathbf{B}} \left[\mathbf{e}_1 \otimes \boldsymbol{\pi}(0) \right] \quad (39)$$

206 The marginal distributions of age and of frailty at death can be shown to be

$$\phi_{\text{age}}^* = (\mathbf{I}_\omega \otimes \mathbf{1}_g^\top) \tilde{\phi} \quad (40)$$

$$\phi_{\text{frailty}}^* = (\mathbf{1}_\omega^\top \otimes \mathbf{I}_g) \tilde{\phi} \quad (41)$$

207 3.4 Projecting the distributions of frailty and age

208 The population vector giving the abundance by age and frailty class is projected by the matrix $\tilde{\mathbf{U}}$
 209 in (20):

$$\tilde{\mathbf{n}}(t+1) = \tilde{\mathbf{U}}\tilde{\mathbf{n}}(t) \quad (42)$$

210 If, as in the G-G model, the initial cohort has support only in the first age class, with distribution
 211 $\boldsymbol{\pi}(0)$, then $\tilde{\mathbf{n}}(0) = (\mathbf{e}_1^\top \otimes \mathbf{I}_g) \boldsymbol{\pi}(0)$.

212 Let $\tilde{\mathbf{p}}(t)$ be the vector giving the *proportional* age-frailty distribution at time t . It is projected
 213 by

$$\tilde{\mathbf{p}}(t+1) = \frac{\tilde{\mathbf{U}}\tilde{\mathbf{p}}(t)}{\|\tilde{\mathbf{U}}\tilde{\mathbf{p}}(t)\|}, \quad (43)$$

214 with $\tilde{\mathbf{p}}(0) = \tilde{\mathbf{n}}(0)/\|\tilde{\mathbf{n}}(0)\|$.

215 The marginal age vector and the marginal proportional age distribution vector are obtained by
 216 summing over frailty classes within age classes, as in (40) for the death distribution. They are
 217 given by

$$\mathbf{n}^*(t) = (\mathbf{I}_\omega \otimes \mathbf{1}_g^\top) \tilde{\mathbf{n}}(t) \quad (44)$$

$$\mathbf{p}^*(t) = (\mathbf{I}_\omega \otimes \mathbf{1}_g^\top) \tilde{\mathbf{p}}(t) \quad (45)$$

218 The marginal frailty vector and the marginal proportional frailty distribution are obtained by
 219 summing $\tilde{\mathbf{n}}$ and $\tilde{\mathbf{p}}$ over age within each frailty class, as in (41) for the death distribution:

$$\mathbf{m}^*(t) = (\mathbf{1}_\omega^\top \otimes \mathbf{I}_g) \tilde{\mathbf{n}}(t) \quad (46)$$

$$\boldsymbol{\pi}(t) = (\mathbf{1}_\omega^\top \otimes \mathbf{I}_g) \tilde{\mathbf{p}}(t) \quad (47)$$

220 From the marginal distribution $\boldsymbol{\pi}(t)$ of frailty, all the statistics, particularly the mean and variance,
 221 of frailty can be calculated, to quantify the effects of selection as a function of age.

222 4 An example: Swedish females

223 An example of the calculations is provided by using the G-G parameters a , b , and k estimated by ?
 224 from period mortality data on Swedish females from 1891 to 2010. I used these parameters, for the
 225 arbitrarily selected year 1950, to create the matrices $\hat{\mathbb{A}}$ and $\hat{\mathbb{D}}$ ($\hat{a} = 0.0340$, $\hat{b} = 0.1200$, $\hat{k} = 8.2300$).
 226 Calculations were carried out with $\omega = 150$ age classes and $g = 100$ logarithmically spaced frailty
 227 classes. The results are shown in a series of figures. The marginal mortality rate $\mu^*(t)$ is shown
 228 in Figure 1. That rate increases nearly exponentially until about age 75, at which point the effects
 229 of selection become apparent and the increase decelerates; mortality converges to an asymptote at
 230 about age 100.

231 4.1 Statistics of longevity

232 The mean, standard deviation, coefficient of variation, and skewness of longevity, computed from \mathbf{N}^*
233 using (34)–(37), are shown in Figure 2. The remaining life expectancy and its standard deviation
234 both decrease with age, but the relative variation, as measured by the coefficient of variation,
235 increases up to about age 90 and then decreases slightly. The distribution of longevity goes from
236 negative to positive skewness with increasing age.

237 4.2 Dynamics of frailty

238 The selection against more frail individuals is seen in the dynamics of the distribution of frailty $\pi(t)$
239 (Figure 3). Both the mean and the standard deviation of frailty decrease with age, with the decline
240 becoming visually evident after about age 50. The CV of frailty is known to remain constant with
241 age in the G-G model. In the matrix calculation, it is very nearly constant, increasing slightly at
242 about age 75.

243 4.3 Effects of the G-G parameters

244 The matrix formulation makes it easy to explore the effects of the G-G parameters on the statistics
245 of longevity. Figures 5–7 show the effects of varying a , b , and the variance, $1/k$, of the frailty
246 distribution over several orders of magnitude around the values for Swedish females in 1950.

247 Life expectancy declines with increases in a ; roughly speaking, a 10-fold increase in a reduces
248 life expectancy at birth by about 10 years (Figure 5). The standard deviation of longevity also
249 declines with increasing a , more dramatically at older ages. Together, these changes lead to a
250 coefficient of variation that increases with a . The skewness of the distribution of longevity also
251 increases with a . When measured at birth, it is negative, but by age 60 skewness changes from
252 negative to positive as a increases.

253 Increases in b also reduce life expectancy (Figure 6). The standard deviation of longevity peaks
254 at an intermediate value of b , and declines sharply at higher or lower values. The coefficient of
255 variation of longevity increases with b until at older ages it eventually declines. At very high
256 values of b the expectation and standard deviation of longevity at older ages become zero and the
257 coefficient of variation is undefined.

258 The effects of changes in the variance $\sigma^2 = 1/k$ of the initial frailty distribution $\pi(0)$ are shown
259 in Figure 7. Expected longevity is relatively insensitive to the σ^2 until it becomes much higher than
260 that observed for Swedish females, at which point the mean, variance, and coefficient of variation
261 of longevity all begin to increase with σ^2 . The skewness of longevity increases to a peak (at values
262 much higher than those observed), and then declines again at extremely high values of σ^2 .

263 4.4 Numerical reliability

264 Based on his results using the Gaussian hypergeometric function (11), ? reports a life expectancy of
265 77.29 years for Swedish females in 1950. Evaluating his formula with the Gaussian hypergeometric
266 function as implemented in MATLAB or in Wolfram Alpha gives a result of 76.22 years. The
267 matrix calculation yields 77.23 years or, when adjusted by 0.5 years to correspond to a trapezoidal
268 integration of the survival function, 76.73 years. The differences among the various implementations
269 of the calculation are small (Table 1).

270 Because the matrix calculation is a discrete model, the results are influenced by the number
271 of frailty classes g and the number of age classes ω included in the model. The number of frailty
272 classes determines how closely $\pi(0)$ can approximate a gamma distribution, and the ability of $\pi(t)$

273 to capture the distribution of frailty at late ages when selection has been operating for a long time.
 274 The number of age classes determines the extent to which the longevity statistics are influenced
 275 by the death of all remaining individuals at age ω , which is not part of the G-G model, but must
 276 appear in any finite state approximation.

277 Figure 4 shows the effect of choices of ω and of g on the expectation and the standard deviation
 278 of longevity. For these parameters, choosing $\omega > 100$ and $g > 40$ provides reliable estimates of
 279 both the mean and the variation of longevity for this data set. Too small a value of ω reduces both
 280 the mean and the standard deviation, because the survival curve is truncated at age ω .

281 5 Generalizations and extensions of the model

282 The calculations of \mathbf{N}^* , \mathbf{s}^* , and all the quantities derived from them depend only on the vec-
 283 permutation model structure (20). As a result, the analysis can be extended from the G-G model to
 284 other models for baseline mortality, other models for the effects of frailty, other frailty distributions,
 285 and other models for the dynamics of frailty. I describe some of these extensions here.

286 5.1 Models for baseline mortality

287 The baseline mortality schedule $\boldsymbol{\mu}_0$ is used to create the frailty-specific mortality schedules $\boldsymbol{\mu}_i$ in
 288 equation (13). These schedules are used to create the matrices \mathbf{U}_i that appear in (15). The Gom-
 289 pertz model is only one possible choice of a baseline schedule. Here, I examine some alternatives;
 290 results, in the same format as Figures 2, 3, and 5–7 are collected in Appendix A.

291 For example, the Gompertz-Makeham model

$$\mu(x) = ae^{bx} + c, \quad (48)$$

292 is obtained by adding an age-independent mortality hazard c to the Gompertz model. The gamma-
 293 Makeham model results from incorporating a proportional frailty effect

$$\boldsymbol{\mu}_i = z_i \left(ae^{bi} + c \right) \quad (49)$$

294 where the z_i have a gamma distribution at age 0.

295 Simply modifying $\boldsymbol{\mu}_0$ in equation (13) transforms the G-G model to the gamma-Makeham
 296 model, with

$$\boldsymbol{\mu}_0 = \begin{pmatrix} e^{0b} \\ \vdots \\ e^{(\omega-1)b} \end{pmatrix} + c \begin{pmatrix} 1 \\ \vdots \\ 1 \end{pmatrix} \quad (50)$$

297 All analyses of the gamma-Makeham model then follow from $\tilde{\mathbf{N}}$, computed from $\boldsymbol{\mu}(z)$, just as with
 298 the G-G model.

299 ? estimated the parameters in the gamma-Makeham model as part of an analysis of the effects
 300 of education on the mortality of male and female cohorts in Turin, Italy, from 1971 to 2007. I
 301 analyzed the data for the baseline cohort of women, for which AIC calculations indicated that
 302 the gamma-Makeham model was much more well-supported by the data than the G-G model (?).
 303 Figure A.1 shows the expectation, standard deviation, CV, and skewness of remaining longevity
 304 as a function of age. The patterns are qualitatively similar to the gamma-Gompertz results for
 305 Swedish females (Figure 2). Selection reduces the mean and the standard deviation of frailty as age
 306 approaches 100, and the log of the marginal mortality rate increases with age in a sigmoid fashion
 307 (Figure A.2).

308 There is no reason to stop at the gamma-Makeham model. ? and ? have added gamma-
 309 distributed frailty to the Siler model for mortality

$$\mu_0(x) = e^{a_1 - b_1 x} + e^{a_2 + b_2 x} + e^{a_3} \quad (51)$$

310 where $\exp(a_1 - b_1 x)$ is a declining force of infant mortality, $\exp(a_2 + b_2 x)$ is an increasing force of old
 311 age mortality, and $\exp(a_3)$ is a constant force of background mortality. An analysis of the gamma-
 312 Siler model requires only substituting this expression for μ_0 for the gamma-Gompertz mortality
 313 function in (12).

314 ? estimated parameters for a gamma-Siler model for cohorts of Swedish females born from
 315 1875–1916. Here I show results for the year 1900. The expectation, standard deviation, CV, and
 316 skewness of remaining longevity are shown as functions of age in Figure A.3. The patterns differ
 317 from those of the G-G and gamma-Makeham models mainly in that they show the effects of the
 318 infant mortality term. This effect is also apparent in the marginal mortality function (Figure A.4c)
 319 which declines sharply after birth, remains low, and then increases, eventually reaching a plateau
 320 at older ages.

321 These examples use parametric functions for the baseline mortality schedule, but they can easily
 322 be extended to semiparametric or nonparametric estimates. The estimated mortality function
 323 simply needs to be incorporated into the matrices \mathbf{U}_i .

324 5.2 Models for the effects of frailty

325 In the G-G model, frailty affects mortality as a proportional hazard. Other models for the effects of
 326 frailty can be incorporated into the construction of the matrices \mathbf{U}_i , by replacing the proportional
 327 hazard formulation in (13) with an expression appropriate to the frailty effects.

328 For example, ? briefly considered a model with accelerated aging, in which

$$\mu(z, x) = \mu_0(zx) \quad (52)$$

329 and point out that, if the baseline mortality schedule is Gompertz, then small changes in z can
 330 have large effects on the mortality, especially at later ages. Figure 3 of ? shows an example with
 331 two frailty classes.

332 Accelerated failure time (AFT) models typically specify frailty in terms of its effect on the
 333 survival function, so that

$$s(z, x) = s(zx) \quad (53)$$

334 which implies that

$$\mu(z, x) = z\mu(zx). \quad (54)$$

335 To incorporate such a model in the matrix calculations require only an appropriate change in the
 336 the expression (13) for the mortality rate of each frailty class.

337 5.3 Models for the distribution of frailty

338 The gamma distribution is attractive as a distribution of frailty for its mathematical properties,
 339 and theoretical results suggest that it is likely to underlie mortality trajectories that reach a plateau
 340 at old ages (?). However, any initial distribution $\pi(0)$ can be incorporated in the calculation of the
 341 marginal survival \mathbf{s}^* in (24), and the dynamics of the frailty distribution generated by (47). This
 342 includes other parametric distributions as well as specification of discrete frailty classes (e.g., ?).

343 **5.4 Models for the dynamics of frailty**

344 In the G-G model, frailty is a fixed property of an individual. However, individual heterogeneity
 345 may be dynamic, increasing (debilitation) or decreasing (recuperation) over time due to stress,
 346 disease, etc. The matrix model readily incorporates any finite-state Markov chain as a model for
 347 dynamic heterogeneity, by properly specifying the matrices \mathbf{D}_i , for $i = 1, \dots, \omega$.

348 For example, ? considered a model with two frailty states, z_1 and z_2 . Individuals begin life with
 349 frailty z_1 with mortality schedule $\mu_1(x)$, and change from state one to state two at a rate $\lambda(x)$. The
 350 second frailty state might represent a morbid event such as a heart attack. This model generalizes
 351 to a model considered by ? and ? with a countably infinite number of frailty classes. The mortality
 352 rate is $\mu_i = \mu_0 + z_i\mu$, and frailty increases at the rate $\lambda_0 + z_i\lambda$. The debilitation process leads to
 353 a stochastic increase in individual frailty over time. The resulting sigmoid trajectory of marginal
 354 mortality cannot be distinguished for that produced by the G-G model with an additive Makeham
 355 term (??).

356 Every individual need not begin with the same frailty. ? considered a model in which the
 357 cohort starts with some initial frailty distribution, and then frailty of each individual proceeds in
 358 accordance with the LeBras model. ? modelled the dynamics of frailty as a diffusion process, in
 359 which individuals may, with equal probability, become more frail or recuperate to a lower frailty
 360 level.

361 To incorporate dynamic heterogeneity into the matrix model, consider a hypothetical model
 362 where individual frailty changes as a diffusion process with reflecting boundaries. Frailty is as
 363 likely to increase as to decrease, but it cannot decline below 0 or increase above some maximum
 364 limit. If the changes in frailty follow a diffusion process, then the discrete time transition matrix
 365 \mathbf{D} can be written

$$\mathbf{D} = e^{k\mathbf{Q}} \tag{55}$$

366 where \mathbf{Q} is the intensity matrix of a continuous-time, nearest-neighbor random walk with

$$q_{ij} = \begin{cases} 1 & j = i - 1 \\ -2 & j = i \\ 1 & j = i + 1 \\ 0 & \text{otherwise} \end{cases} \tag{56}$$

367 except that $q_{1,1} = q_{g,g} = -1$. The coefficient k adjusts the speed of diffusion (e.g., ?). Unlike the
 368 LeBras model, this diffusion model does not change the rate of indisposition or recuperation as the
 369 frailty changes, but such dynamics can be easily incorporated.

370 Adding diffusion to the G-G model for Swedish females gives the results shown in Figure 10.
 371 Both life expectancy and the standard deviation of longevity are maximized at intermediate values
 372 of diffusion. There is a balance between creation of diversity by diffusion, and removal of diversity
 373 by selection (a balance familiar from mutation-selection calculations in population genetics). At
 374 sufficiently high rates of diffusion, individual move among frailty levels so rapidly that they cannot
 375 avoid exposure to high levels of frailty (this reduces life expectancy), and because all individuals
 376 experience this random movement the variance in frailty is also reduced.

377 The interaction between the creation of heterogeneity by diffusion and its elimination by se-
 378 lection is shown in Figure 11. The standard deviation of frailty increases from its value at birth,
 379 under the impact of diffusion. Eventually, mortality increases enough to reduce the variation by
 380 selection. At very high levels of diffusion, the heterogeneity is almost totally determined by the
 381 diffusion. At low levels of diffusion, the increase in heterogeneity is smaller, and its reduction due
 382 to selection more prominent.

383 5.5 Some animal mortality patterns

384 In an exploration of the effects of heterogeneity on the distribution of age at death, ? estimated
385 G-G parameters from data on laboratory populations of five species of invertebrate animals: a
386 bean beetle (*Callosobruchus maculatus*), the medfly (*Ceratitis capitata*), the fruit fly *Drosophila*
387 *melanogaster*, the nematode *Caenorhabditis elegans*, and a parasitoid wasp (*Diachasimimorpha*
388 *longicaudata*.) Because the estimated G-G parameters for these species do not appear in the
389 original paper, they are listed here in Table B.1.

390 These species exhibit considerably greater variance in frailty than do the human examples
391 considered so far, ranging from 0.90 to 2.18. In contrast, estimates of the initial variance for
392 Swedish females from 1891 to 2010 range from 0.10 to 0.14 (?). Figure 9 shows the marginal
393 mortality rate for each species as a function of age, with age and mortality rate both standardized
394 relative to life expectancy at birth. The marginal mortality rates reaches a plateau at an age of
395 about 1 life expectancy, at a standardized mortality rate of 0.1 to 0.5. Further interspecific analyses
396 would be interesting.

397 6 Heterogeneous frailty vs. individual stochasticity

398 Inter-individual variance in longevity is often interpreted as evidence of heterogeneity among in-
399 dividuals in their mortality risks. This interpretation is incorrect, because variance in longevity
400 also arises from *individual stochasticity*; the random variation in the fates of individuals subject to
401 the same risks as they move through the life cycle (???). The only way to partition variance into
402 components due to heterogeneity and to individual stochasticity is with a model that contains both
403 sources; the matrix formulation here does so.

404 Thus, the variance in longevity shown in Figure 8 contains both components. As the variance
405 in the initial frailty distribution $\pi(0)$ approaches zero, the remaining variance in longevity is due
406 to individual stochasticity. It is apparent from Figure 8 that the observed variance in longevity is
407 only slightly greater than that accounted for by individual stochasticity.

408 This is not always the case: Table 2 compares the decomposition of the variance for Swedish
409 females (the G-G, gamma-Makeham, and gamma-Siler models) with that for the animal species
410 from ?. In the human populations, heterogeneity accounts for only 2–7% of the variance in longevity.
411 The greater variance in frailty in the experimental animal data makes a much higher contribution
412 to the variance in longevity, from 46% to 83%.

413 It is important to remember that the relative contributions of individual stochasticity and
414 individual heterogeneity depend not only on the variance in frailty, but also on the mortality
415 schedule, and deserve further empirical investigation.

416 7 Conclusions

417 The gamma-Gompertz and related frailty models provide a powerful way to analyze the mortality
418 of heterogeneous cohorts (?). They do so by capturing the interacting effects of changing mortal-
419 ity with age and selection among individuals with different frailty states. Frailty models can be
420 characterized by their components:

- 421 1. a baseline mortality rate,
- 422 2. a mode of action by which frailty affects the baseline mortality rate,
- 423 3. the dynamics of individual frailty over time, and

425 In the matrix G-G model, the baseline mortality rate follows the Gompertz model (1) and frailty
 426 affects the baseline as a proportional hazard, as in (13). The frailty dynamics are fixed, so that
 427 \mathbb{D} in (15) is an identity matrix, and the initial distribution $\pi(0)$ is a gamma distribution with a
 428 mean of 1. Many other models can be created by changing one or more of these components. The
 429 result, however, is always a model classifying individuals by two criteria: age and frailty. The vec-
 430 permutation matrix model (20) methodologically keeps track of both criteria, and makes it easy
 431 to calculate the properties of the joint age \times frailty distribution, the marginal age-specific mortality
 432 and survival functions, and a complete set of statistics of longevity.

433 Table 3 gives a step-by-step protocol for the analysis of the G-G model. Other choices of
 434 baseline mortality rate (e.g., the Makeham model or the Siler model considered in Section 5.1),
 435 the action of frailty (e.g., accelerated failure time models), the dynamics of frailty (e.g., the frailty
 436 diffusion models discussed in Section 5.4), or the initial distribution of frailty require only simple
 437 modifications of the appropriate steps in Table 3.

438 The effects of model parameters on the statistics of longevity, shown in Figures 5, 6, 7, and
 439 8, reveal interesting patterns. In the G-G model, life expectancy declines with increasing values
 440 of a and b , which is not unexpected. It increases with increasing variance in initial frailty, which
 441 is less intuitively easy to explain. The effects on variance, CV, and skewness of longevity are
 442 more diverse. The standard deviation of longevity declines with increases in a , is maximized at
 443 intermediate values of b , and increases with initial variance in frailty. The skewness of longevity
 444 increases with increases in a and b , and increases and eventually declines again with increases in
 445 initial variance in frailty.

446 Variance in longevity is sometimes interpreted as evidence of inequality among individuals
 447 within the population, but this is not necessarily true. The variance calculated from a specified
 448 mortality schedule assumes that all individuals experience the rates defined by that schedule; hence
 449 differences among individuals reflects the stochastic outcome of those rates. Frailty models, how-
 450 ever, include heterogeneity among individuals, so manipulation of the variance of the initial frailty
 451 distribution $\pi(0)$ makes it possible to decompose variation in longevity into contributions from
 452 individual stochasticity and heterogeneous frailty (Figure 8. In several human mortality studies
 453 based on the gamma-Gompertz, gamma-Makeham, and gamma-Siler models, the contribution of
 454 heterogeneous frailty to the variance in longevity is small (2% – 7%). In a set of laboratory studies
 455 of invertebrate animals, it is much higher (50%–80%; see Table 2). More analyses of such patterns
 456 will be presented elsewhere.

457 Finally, I note that the results here provide connections to several other problems related to
 458 frailty. First, there is no need to limit the analysis to age-classified models. The effects of frailty
 459 on stage-classified (e.g., educational status, health status) or multi-stage models can be analyzed
 460 simply by using the appropriate formulation in \mathbb{U} . The formulation as an absorbing Markov chain
 461 can be used to compute likelihood functions from data on individuals. This is used by animal
 462 ecologists working with mark-recapture data (e.g., ??) and implemented in some software packages
 463 (??). Because frailty is inherently unobserved, issues of identifiability arise, which can also be
 464 addressed using the Markov chain formulation (?).

465 Finally, it is a significant advantage that the matrix model is directly amenable to sensitivity
 466 analysis using matrix calculus methods (e.g., ????????) These methods provide the sensitivity of
 467 the moments of longevity, the joint distribution of age and stage at death, and the survivorship
 468 and mortality functions to changes in any of the parameters of the model.

469 **8 Acknowledgments**

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474 am grateful for the hospitality of the Max Planck Institute for Demographic Research.

Table 1: Comparison of life expectancy results from ?, from this paper, and from Missov’s theorem implemented in MATLAB and calculated using Wolfram Alpha. The adjusted value from this paper has had 0.5 years subtracted to make the result directly comparable to a trapezoidal computation of the integral of $S^*(t)$. For the matrix calculations, $\omega = 150$ and $g = 100$.

Source	Expected longevity
Missov (2013)	77.29
Missov via Matlab	76.22
Missov via Wolfram	76.22
Matrix method	77.23
Matrix method (adjusted)	76.73

Table 2: Decomposition of the variance in longevity for human populations and laboratory populations of invertebrate species. The variance σ^2 in initial frailty, the variance $V(\eta)$ in longevity and the components of $V(\eta)$ due to individual stochasticity and to heterogeneous frailty, and the proportion of the variance due to heterogeneity. Species are listed in order of increasing initial variance in frailty. Data from ?.

Species	σ^2	$V(\eta)$	Stochasticity	Heterogeneity	Proportion
Sweden 1950 G-G	0.122	122.9	114.1	8.72	0.071
Turin G-M	0.096	351.6	347.3	104.3	0.012
Sweden 1900 G-S	0.120	1091.7	1074.1	17.60	0.016
nematode	0.90	18.0	9.7	8.3	0.46
fruit fly	0.94	88.1	46.1	42.0	0.48
beetle	1.31	12.7	5.2	7.5	0.59
medfly	1.34	81.8	29.5	52.3	0.64
wasp	2.18	30.3	5.1	25.2	0.83

Table 3: A protocol for analysis of the gamma-Gompertz model.

1. Specify the Gompertz parameters a and b , and the gamma distribution parameter k . Choose values for the numbers of age classes ω and the number of frailty classes g .
2. Generate the baseline mortality vector $\boldsymbol{\mu}_0$ from (12)
3. Specify the frailty classes z_i , $i = 1, \dots, g$, and the discrete approximation to the gamma distribution $\boldsymbol{\pi}$. Logarithmically-spaced frailty classes are recommended.
4. Create the matrices \mathbf{U}_i , for $i = 1, \dots, g$, as in equation (14).
5. Create the block diagonal matrices \mathbb{U} and \mathbb{D} according to (15).
6. Create the joint transition matrix $\tilde{\mathbf{U}}$ according to (20).
7. Analyze the model
 - (a) Compute the fundamental matrix $\tilde{\mathbf{N}}$ from (22).
 - (b) Compute the marginal survival function \mathbf{s}^* from (24).
 - (c) Generate the marginal fundamental matrix \mathbf{N}^* from (25).
 - (d) Generate life expectancy and other indices of longevity from \mathbf{N}^* using (26)–(37).
8. Project the dynamics of the age-railty distribution $\tilde{\mathbf{n}}(t)$ with (42). Obtain the marginal age abundance vector \mathbf{n}^* using (44) and the marginal age distribution vector \mathbf{p}^* using (45).
9. Obtain the marginal frailty abundance vector \mathbf{m}^* using (46) and the frailty distribution $\boldsymbol{\pi}(t)$ from equation (47).
10. If desired, create the mortality matrix $\tilde{\mathbf{M}}$ and generate the distributions of age and of frailty at death from equations (39)–(41).

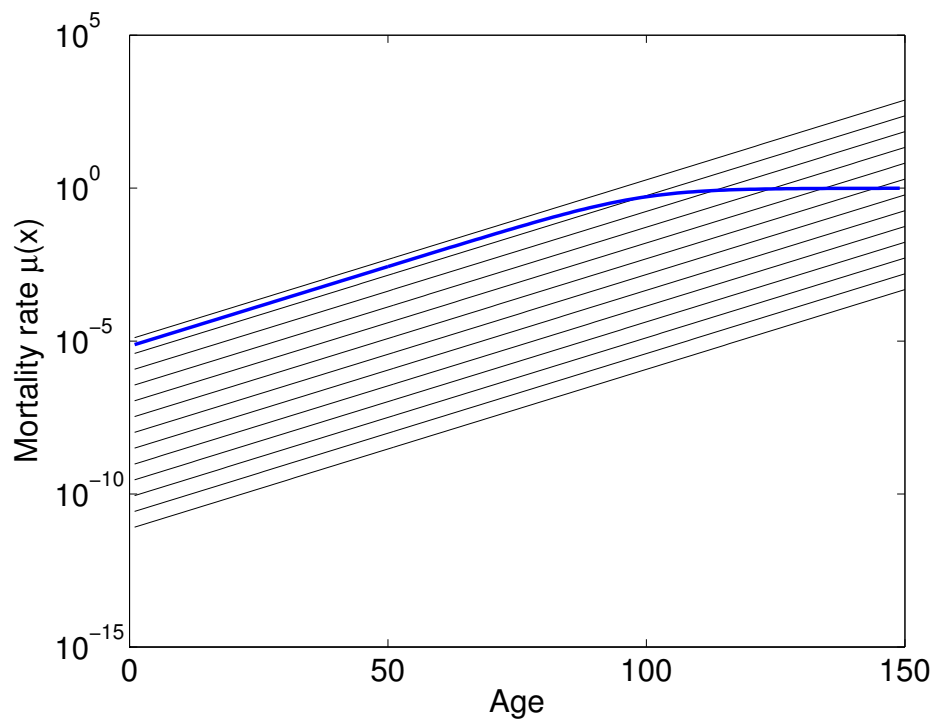


Figure 1: Gamma-Gompertz mortality rate $\mu(t)$ as a function of age, t . Black lines show the age-specific mortality rates for a few of the frailty classes in the model. The blue line shows the marginal hazard $\mu^*(t)$. Parameters for Swedish females as reported in ? for the year 1950.

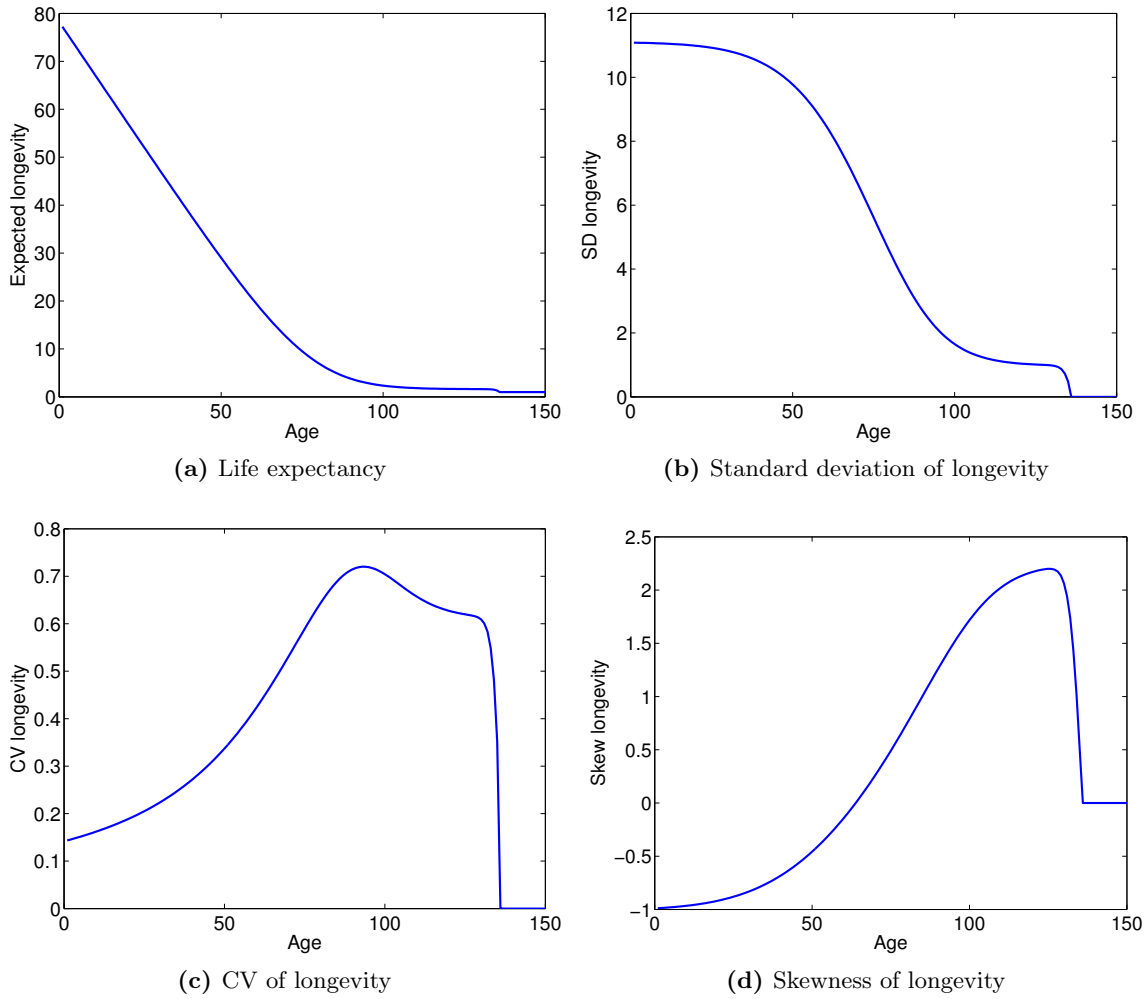


Figure 2: Statistics of longevity for the gamma-Gompertz model, as a function of age, using parameters reported in ? for the year 1950. (a) Life expectancy. (b) Standard deviation of longevity. (c) Coefficient of variation of longevity. (d) Skewness of longevity. [add vertical line at age 110]

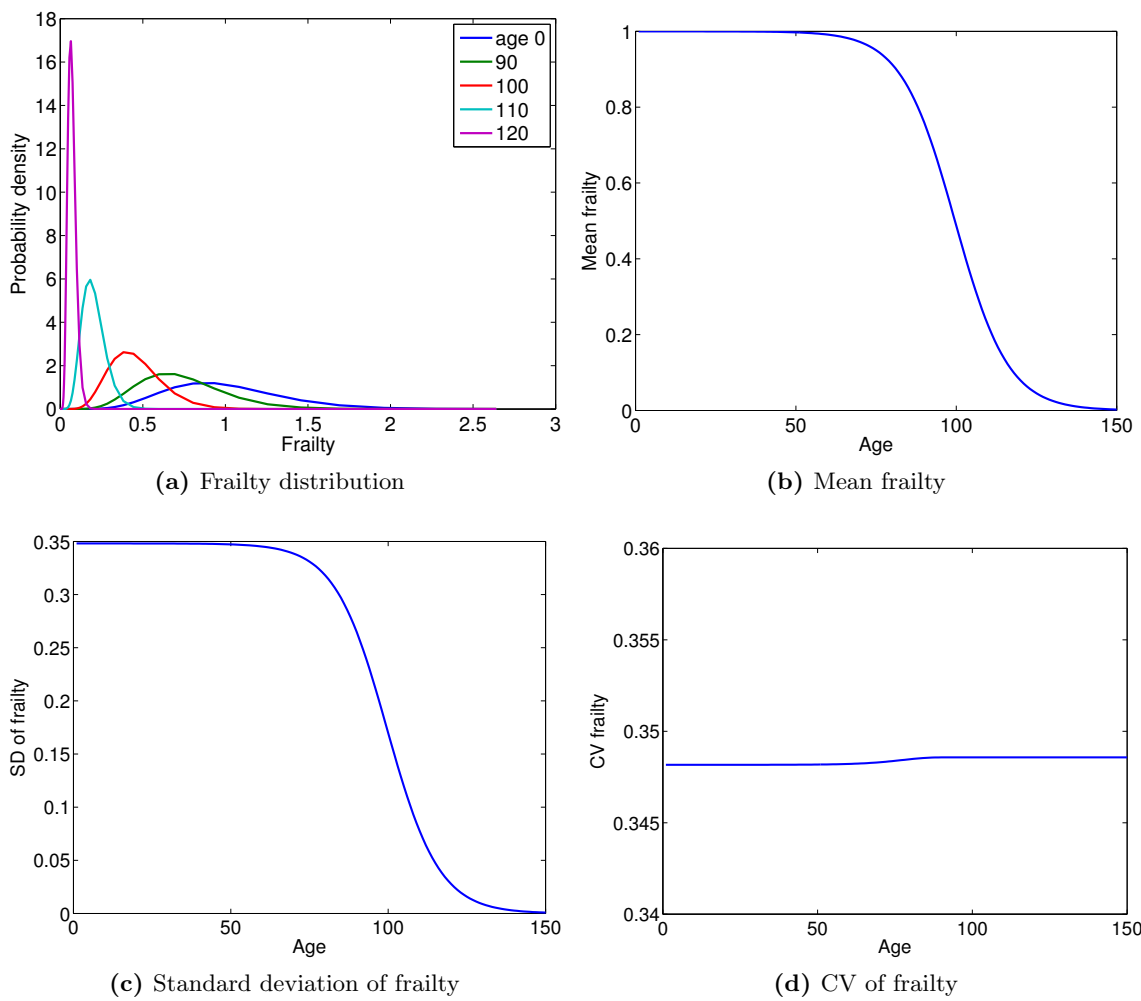


Figure 3: Changes due to selection in the distribution of frailty, and in the mean, CV, and skewness of that distribution, over the life of a cohort. Parameters for Swedish females, as reported in ? for the year 1950.

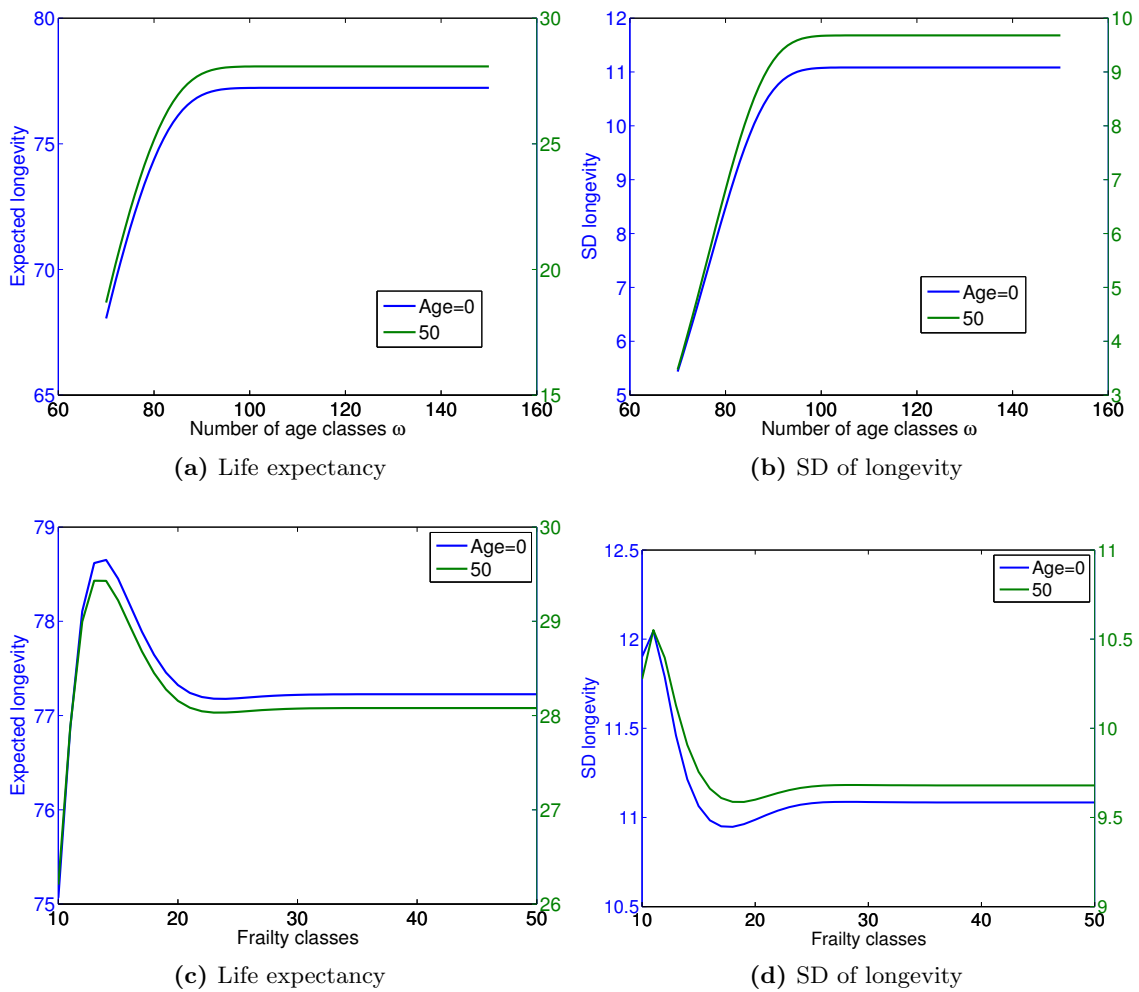
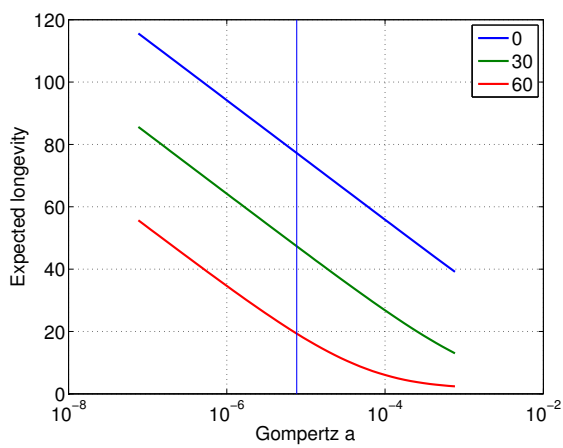
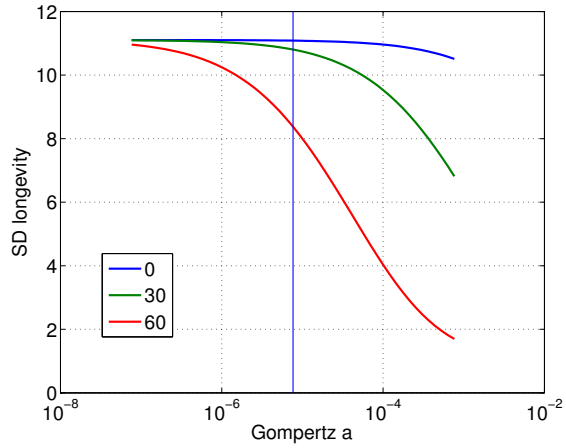


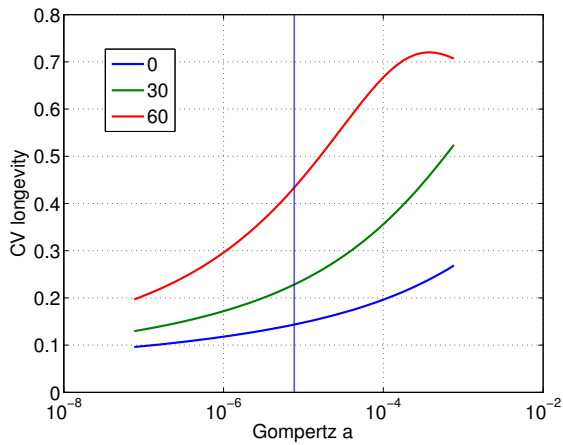
Figure 4: Effect of the number of age classes (ω) and the number of frailty classes (g) on the estimates of life expectancy and the standard deviation of longevity, at ages 0 and 50. (a) and (b) show the effects of age classes, with $g = 100$. (c) and (d) show the effect of the number of frailty classes, with $\omega = 150$.



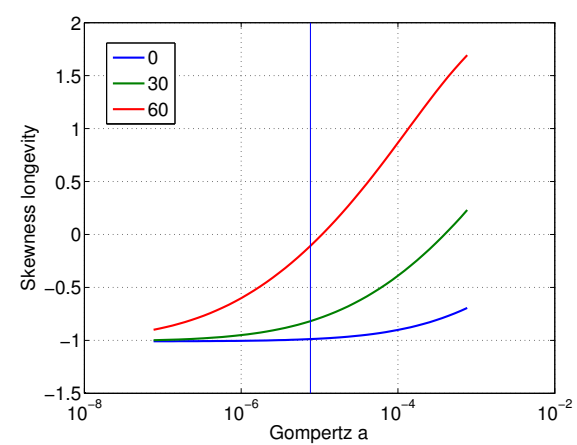
(a) Life expectancy



(b) Standard deviation of longevity



(c) CV of longevity



(d) Skewness of longevity

Figure 5: Statistics of longevity for the gamma-Gompertz model, using parameters reported in ? for Swedish females in 1950, as a function of the Gompertz parameter a .

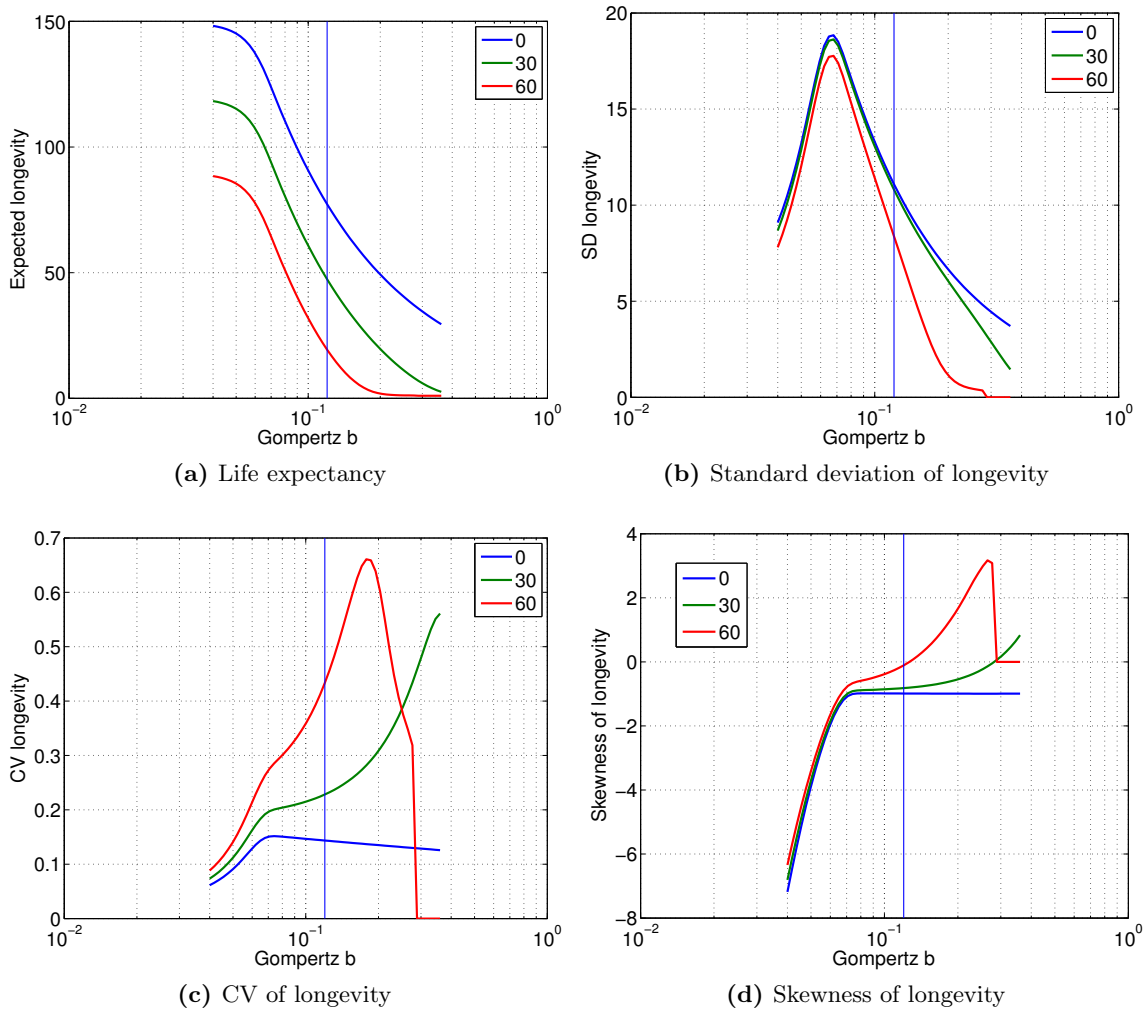


Figure 6: Statistics of longevity for the gamma-Gompertz model, using parameters reported in ? for Swedish females in 1950, as a function of the Gompertz parameter b , for ages 0, 30, and 60 years.

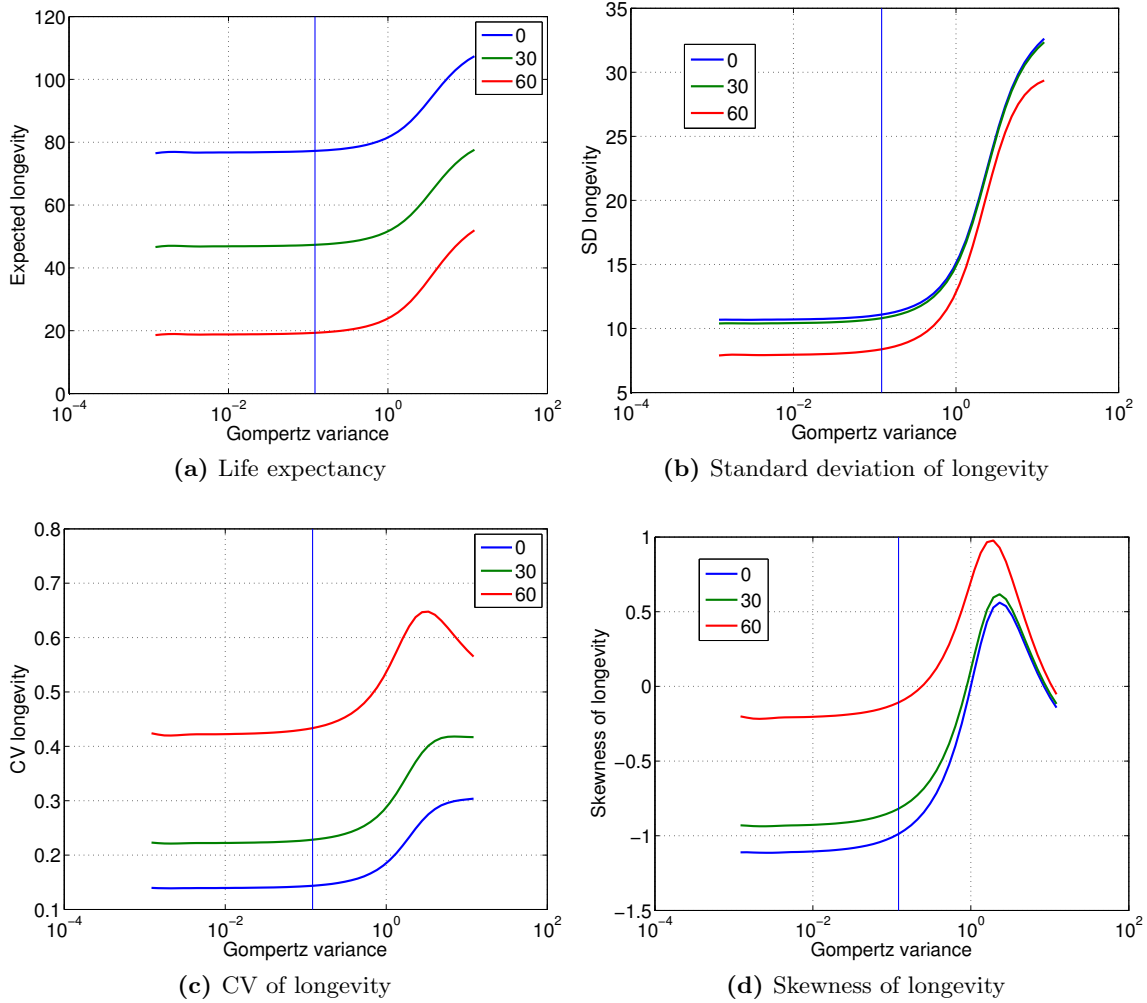
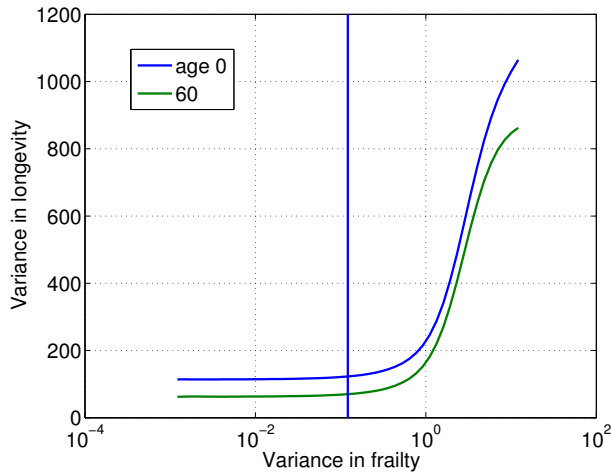
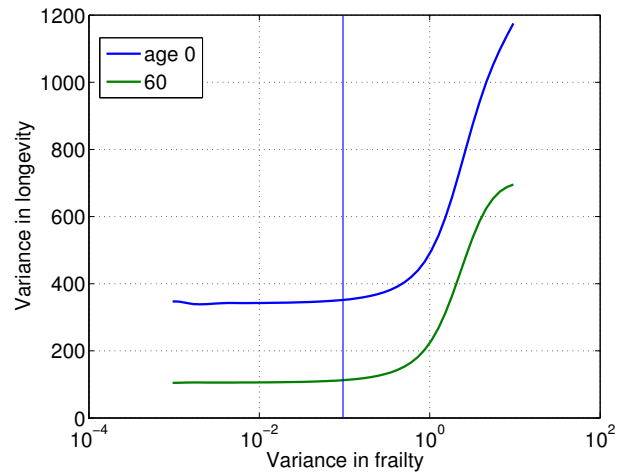


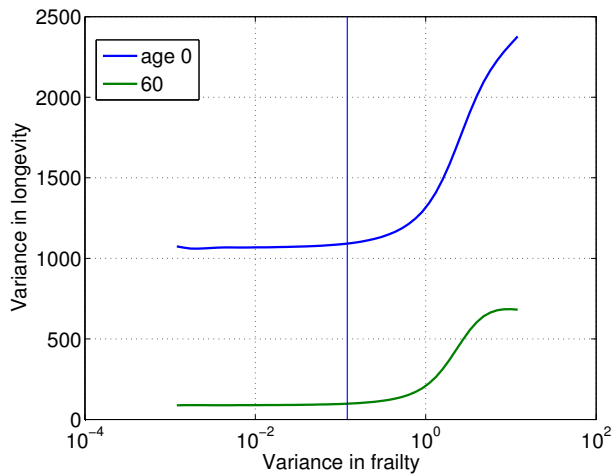
Figure 7: Statistics of longevity at ages 0, 30, and 60 years for the gamma-Gompertz model as a function of the variance in the gamma distribution of frailty (equal to $1/k$). Parameters as reported in ? for Swedish females in 1950. The vertical lines indicate the observed value of variance.



(a) Gamma-Gompertz model



(b) Gamma-Makeham model



(c) Gamma-Siler model

Figure 8: The variance of longevity, at ages 0 and 60, as a function of the variance of the initial frailty distribution. (a) The gamma-Gompertz model, calculated from parameters reported by ? for Swedish females in 1950. (b) The gamma-Makeham model, calculated from parameters reported by ? for a cohort model of the female population of Turin. (b) The gamma-Siler model, calculated from parameters reported by ? for Swedish females born in 1900. The vertical lines indicates the observed values of initial variance in frailty.

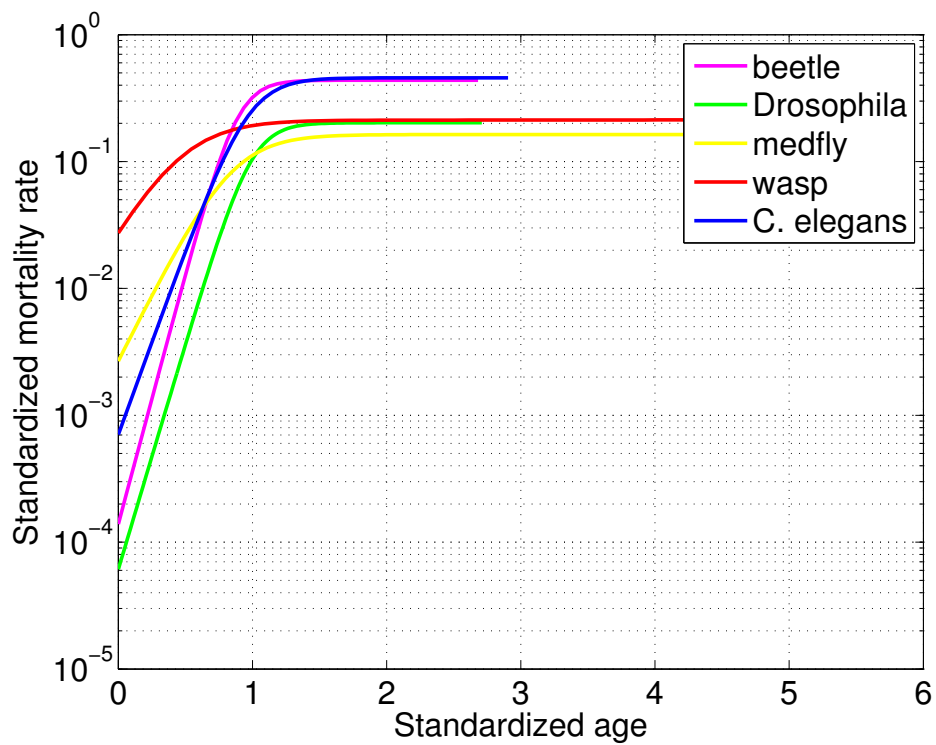
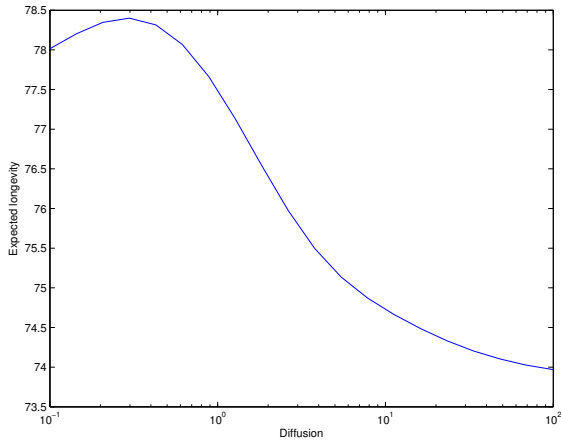
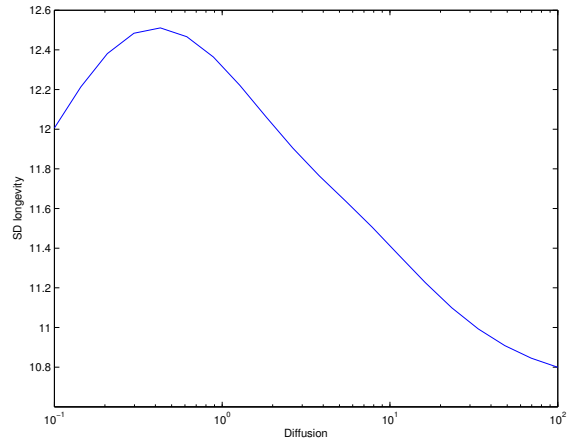


Figure 9: The marginal mortality rate $\mu^*(t)$ as a function of standardized age t , for five species of invertebrate animals, based on G-G parameters estimated by ?. The age abscissa is scaled by dividing age by the life expectancy at birth. The mortality rate is standardized by multiplying by the same life expectancy.



(a) Life expectancy



(b) Standard deviation of longevity

Figure 10: The expectation and the standard deviation of longevity at birth for the gamma-Gompertz model with added diffusion of frailty. Parameters as reported in ? for Swedish females in 1950.

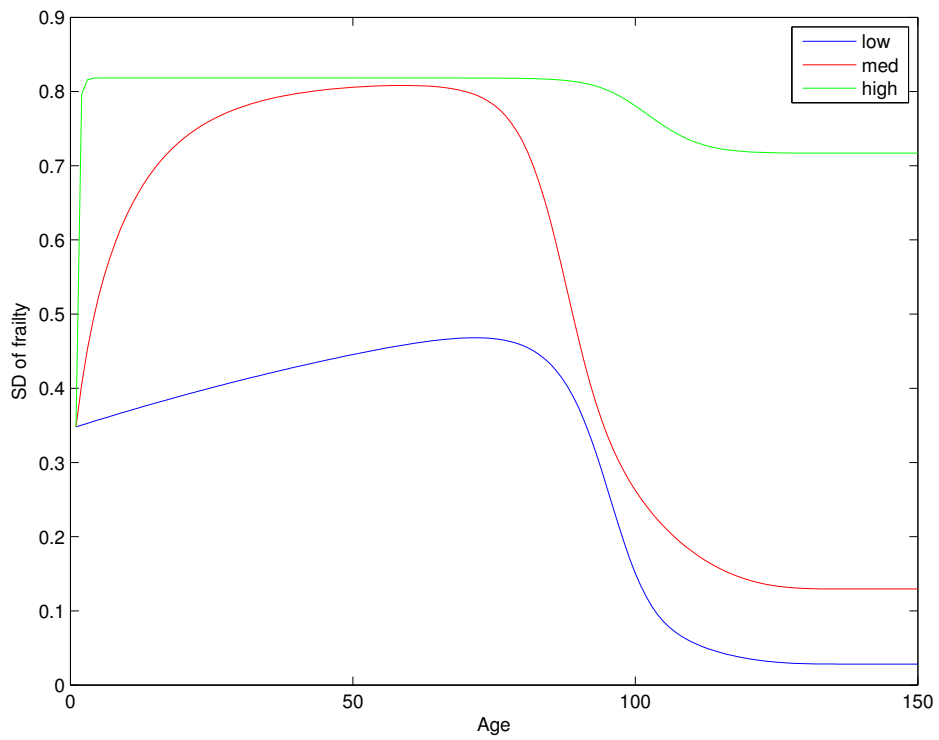


Figure 11: The standard deviation of frailty in a gamma-Gompertz model with diffusion of frailty, at low, medium, and high values of diffusion. Parameters as reported in ? for Swedish females in 1950.

477 **A Gamma-Makeham and gamma-Siler models**

478 This appendix collects results on the statistics of longevity and the dynamics of frailty for the
479 gamma-Makeham model and the gamma-Siler model, in the same format used for results from the
480 G-G model in Figures 2 and 3.

481 **A.1 Gamma-Makeham**

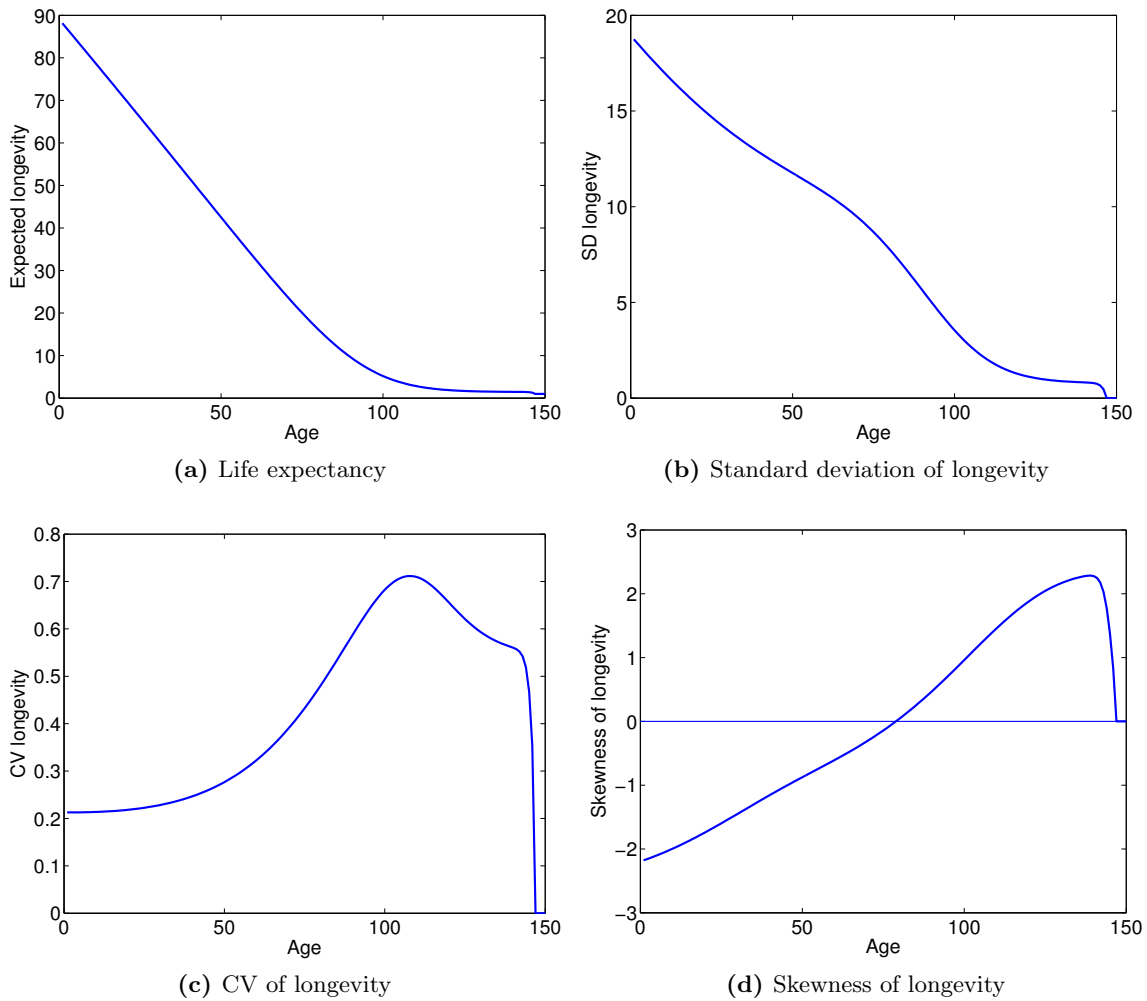
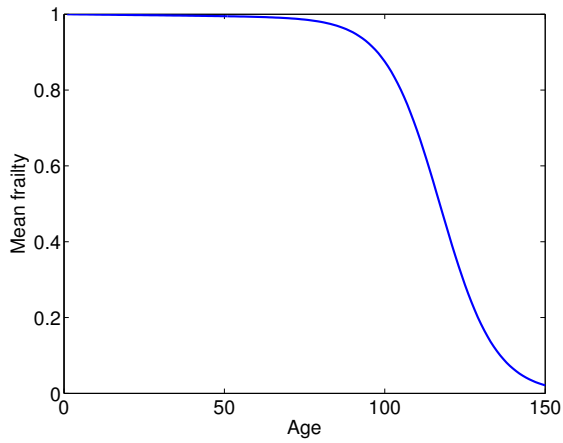
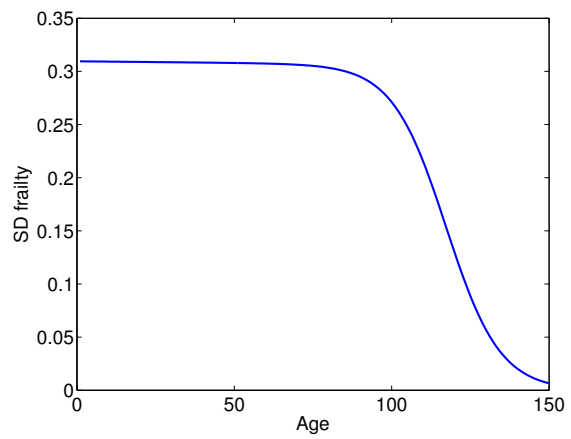


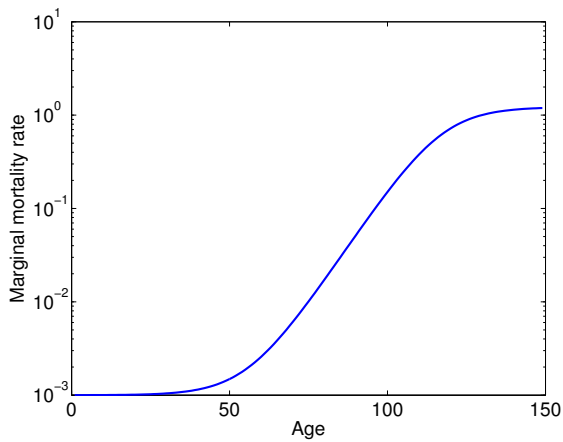
Figure A.1: Statistics of longevity for the gamma-Makeham model, as a function of age. Calculated from parameters reported by ? for a cohort model for the female population of Turin.



(a) Mean frailty



(b) Standard deviation of frailty



(c) Marginal mortality

Figure A.2: Statistics of frailty, and marginal mortality rate μ^* , for the gamma-Makeham model, as a function of age. Calculated from parameters reported by ? for a cohort model for the female population of Turin.

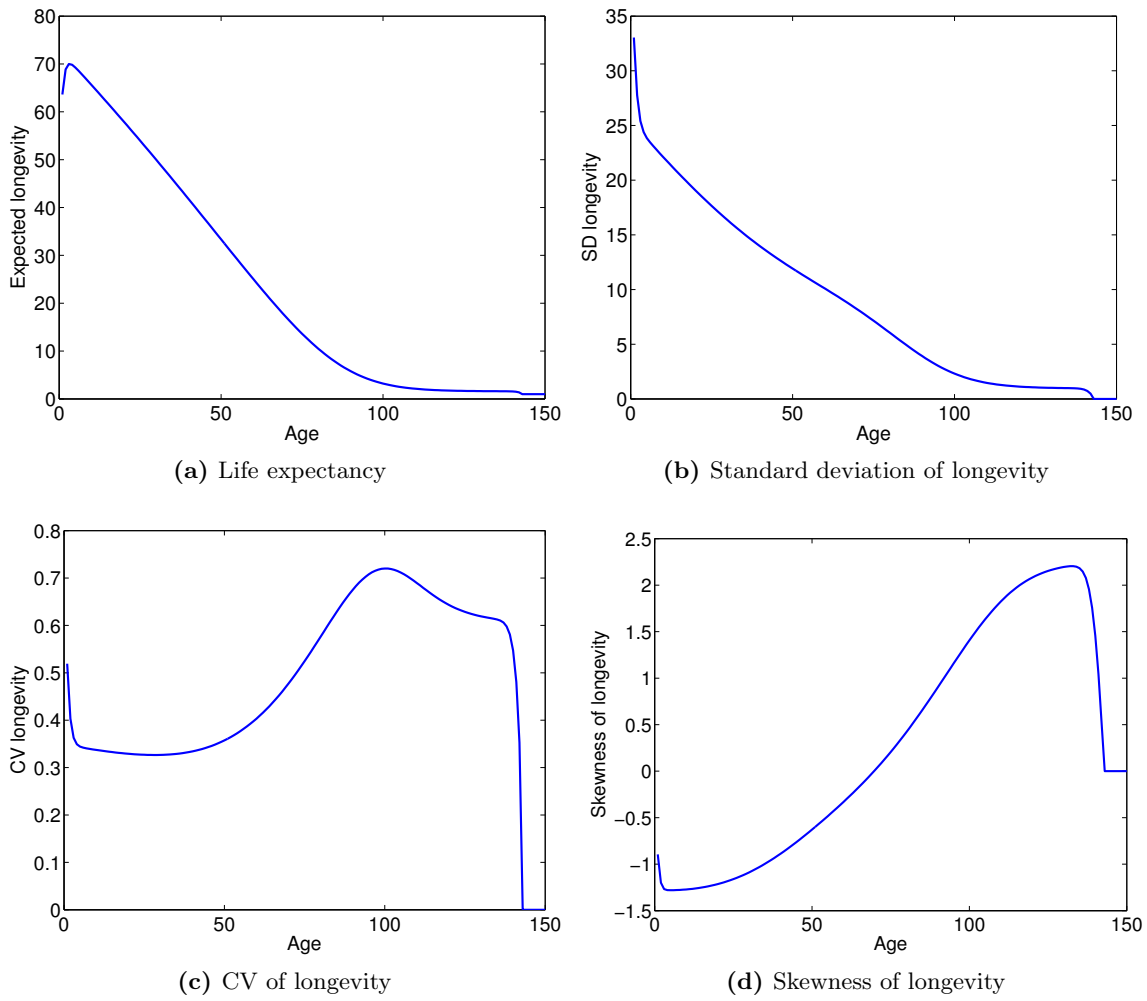
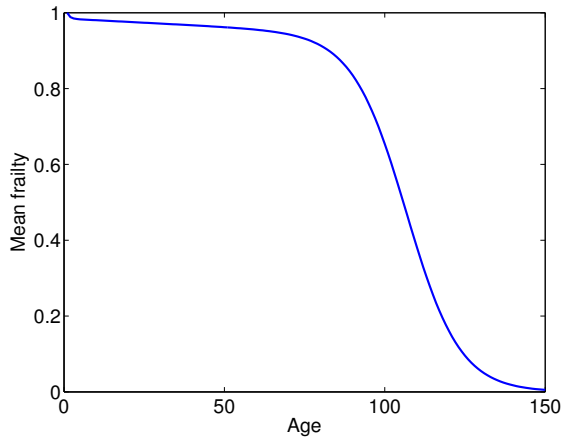
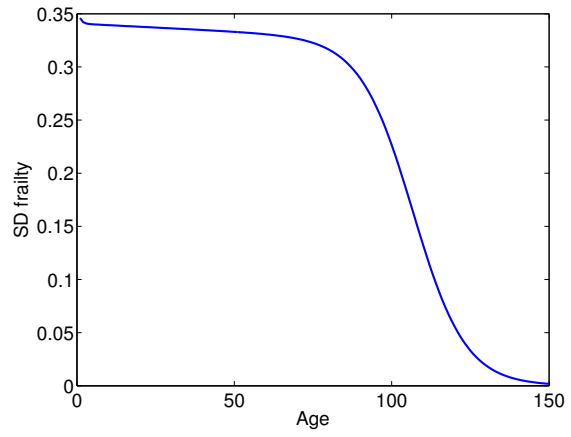


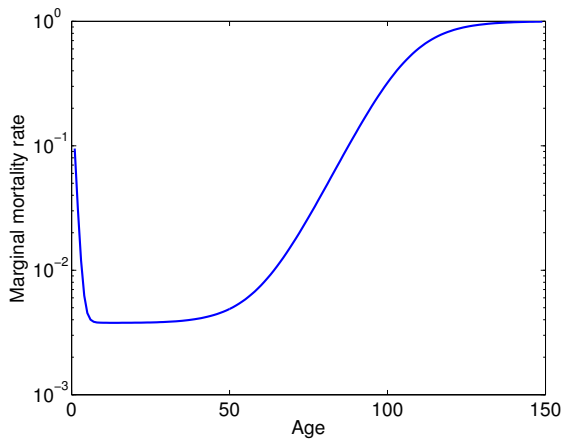
Figure A.3: Statistics of longevity for the gamma-Siler model, as a function of age. Calculated from parameters reported by ? for Swedish females born in 1900.



(a) Mean frailty



(b) Standard deviation of frailty



(c) Marginal mortality

Figure A.4: Statistics of frailty, and marginal mortality rate μ^* , for the gamma-Siler model, as a function of age. Calculated from parameters reported by ? for Swedish females born in 1900.

483 **B Parameters for animal species**

Table B.1: Gamma-Gompertz parameters for the invertebrate animal species analyzed by ?; data provided by Horiuchi (personal communication).

Species	a	b	k
Medfly	0.0027	0.2168	0.7530
Nematode	6.9970×10^{-4}	0.4059	1.1264
Wasp	0.0278	0.4575	0.4640
<i>Drosophila</i>	6.0558×10^{-5}	0.1878	1.0796
Beetle	1.3760×10^{-4}	0.5671	0.7721