Minimum reversion in multivariate time series

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Abstract

We propose a new multivariate time series model in which we assume that each component has a tendency to revert to the minimum of all components. Such a specification is useful to describe phenomena where each member in a population which is subjected to random noise mimicks the behaviour of the best performing member .

We show that the proposed dynamics generate co-integrated processes. We characterize the model's asymptotic properties for the case of two populations and show a stabilizing effect on long term dynamics in simulation studies. An empirical study involving human survival data in different countries provides an example which confirms the occurrence of the phenomenon of reversion to the minimum in real data.

Keywords: multivariate time series, co-integrated time series, human mortality modelling, rank-dependent drift

Introduction 1

When multivariate time series are used to describe the joint dynamics of stochastic processes, often an a priori assumption of a stabilizing mechanism is made. We expect, for example, that many economic variables will fluctuate randomly over time, but we do not find it plausible that they will diverge while doing so. This is because certain long-term equilibrium relationships are assumed to be present between such variables, despite the volatility we see in them over short time horizons. Many economic examples are now known (see for example Engle and Granger [1987]) and other fields in which such co-integration relationships are found include neuroscience (Østergaard et al. [2017]) and gene differentiation in populations (Hössjer and Ryman [2014]).

A possible model feature that implements such a stabilizing mechanism in multivariate time series is mean reversion: the tendency of all components to drift towards a constant, which has the interpretation of the long term average of the series. Adding a mean reversion term to all increments of a multivariate discrete random walk makes the series second-order stationary. This also guarantees that the distance between any two components will be stationary.

In this paper we propose a different mechanism: instead of making all components tend to a priori chosen constants, we impose that at every time step they move, in expectation, towards the minimal value among all components. When different components in the time series signify the

value of a certain common variable among different groups, such an assumption can be interpreted as all groups mimicking the behaviour of the group which currently has the 'best' value¹.

If knowledge about what is beneficial or detrimental for a group to minimize a certain quantitative indicator is communicated between different groups, and each group is capable of using this knowledge to their benefit, the best practices are implemented in all groups over time. The overall effect would be that the group which has achieved the best (i.e. the minimal) value would be 'followed' over time by those with worse achievements, as they 'learn' or 'mimic' what is beneficial. We will show that this effect will lead to a co-integration relationship between groups but also to an additional downward drift that no group would achieve on its own or if the minimum reversion effect would not be present. In that sense, learning from the best performer generates improvements for all that would not occur if individual groups were left to their own devices.

This hypothesis of 'reversion to the minimum' in multivariate time series can be tested by specifying models with and without such an effect and applying the Bayesian Information Criterion to compare the goodness of fit. In this paper we provide such specifications, analyse the properties of models where it is present, and give an empirical example of a specific time series where clear evidence for this effect is found.

There are a number of models in the literature that are related to the one we propose in this paper. Systems of diffusion processes have been proposed in which the drift and diffusion coefficients of an individual process depend on the rank of that process within the system. For example, Fernholz [2002] and Banner et al. [2005] introduce the Atlas model to describe the market capitalization of firms in equity markets, and Ichiba et al. [2013] apply their model to define optimal investment strategies. The Atlas model is applied by Sartoretti and Hongler [2013] to describe the dynamics of a swarm, and Ruzmaikina and Aizenman [2005] and Shkolnikov [2009] consider the evolution of competing particle systems in discrete time and study the distribution of the gaps between the particles. Balázsa et al. [2014] introduced a continuous time Markov jump model for interacting particles with a jump rate which depends linearly on the distance of the particle from the center of mass of the whole group.

In contrast to the models proposed in the literature, we consider a system of discrete time processes under the assumption that the distribution of each component's stochastic increments does not depend on the rank of the component but on its distance to the minimum component. This feature leads to multivariate time series with short term volatility and long term equilibrium relationships, while it gives a clear interpretation of the presence of the stabilizing mechanism in the long run.

The remainder of the paper is organised as follows. In section 2 we introduce our model and analyse some of its theoretical properties. Section 3 provides a study in which we provide empirical evidence that a 'minimum reversion' effect can be found in human mortality data. Finally, we provide some conclusions and suggestions for further research in Section 4.

2 Minimum Reversion Model

2.1 Specification

Let $\mathcal{T} = \{t_0, t_0 + 1, ..., t_1\}$ for certain $t_0, t_1 \in \mathbb{N}$ with $t_1 > t_0$ and let $\{\kappa_t\}_{t \in \mathcal{T}}$ be a multivariate time series in \mathbb{R}^C with components $\kappa_{t,c}$ with $c \in \mathcal{C} = \{1, 2, ..., C\}$ for a given $C \in \mathbb{N} \setminus \{0, 1\}$. We denote

 $^{^{1}}$ We consider reversion to the minimum in this paper and only consider biometric variables for which low values are deemed to be best. One could of course also apply our analysis in cases where there is a drift towards the maximum. Our choice here is motivated by the particular empirical example we give in the last section, where we consider multivariate time series for human mortality.

the minimum value among the different components at time $t \in \mathcal{T}$ by

$$m_t := \min_{c \in \mathcal{C}} \kappa_{t,c},\tag{1}$$

and specify the following model for the dynamics of the components $\kappa_{t,c}$ with $c \in \mathcal{C}$:

$$\kappa_{t+1,c} - \kappa_{t,c} = \mu_c + \zeta_c(\kappa_{t,c} - \kappa_{t-1,c}) + \sigma_c Z_{t+1,c} + \lambda_c(m_t - \kappa_{t,c}).$$
(2)

The $\mu_c \in \mathbb{R}, \zeta_c \in [0, 1), \lambda_c \in [0, 1)$ and $\sigma_c \in \mathbb{R}^+$ are given constants. We define $Z_t = (Z_{t,1}, \ldots, Z_{t,C})'$ and assume that the $\{Z_t\}_{t\in\mathcal{T}}$ are independent and identically distributed C-dimensional random variables with a multivariate Gaussian distribution. More precisely, we assume that

$$Z_{t,c} = \rho_c W_{t,0} + \sqrt{1 - \rho_c^2 W_{t,c}}, \qquad (3)$$

for independent standard Gaussian variables $\{W_{t,c}\}_{c \in \mathcal{C} \cup \{0\}, t \in \mathcal{T}}$, which creates a correlation structure parameterized by the C constants $\rho_c \in (-1, 1)$.

The constant drift μ_c , the first order autoregression coefficient ζ_c and Gaussian increment $\sigma_c Z_{t+1,c}$ are standard features in time series modelling. Our innovation concerns the λ_c parameters which quantify the 'learning effect'. To discuss the properties of the newly proposed term in (2) we first consider the model with $\mu_c = \zeta_c = 0$ and $\lambda_c > 0$. This introduces a downward drift $\lambda_c(m_t - \kappa_{t,c})$ for κ_c when c is not the component with the lowest current value.

The specification in (2) with $\mu_c = \zeta_c = 0$ and $\lambda_c > 0$ also creates a downward trend in the minimum process m_t defined in (1). To form some intuition for this result we consider the case where $\rho_c = 0$ for all c, that is, the $Z_{t,c}$ are independent for any fixed t. In that case we have

$$\mathbb{P}(m_{t+1} \le a \mid \kappa_t) = 1 - \prod_{c \in \mathcal{C}} \Phi\left(\frac{(1-\lambda_c)\kappa_{t,c} - a + \lambda_c m_t}{\sigma_c}\right).$$

Substituting $a = m_t$ gives an expression for $\mathbb{P}(m_{t+1} \leq m_t \mid \kappa_t)$ and since $\kappa_{t,c} \geq m_t$ for all c and $\kappa_{t,c} = m_t$ for at least one c, this probability must be strictly greater than $\frac{1}{2}$ and smaller than or equal to $1 - 2^{-C}$. This establishes that the probability of a downward movement is always greater than $\frac{1}{2}$ and that it is largest when all components attain the same value, that is, $\kappa_{t,c} = m_t$ for all $c \in \mathcal{C}$. The probability is increasing with the number of components C.

Individual components are thus not stationary but they turn out to be co-integrated, as the following result shows.

Proposition 1. If all processes in (2) have a common minimum reversion parameter and a common drift and there is no autoregressive term, so $\mu_c = \mu$, $\lambda_c = \lambda > 0$ and $\zeta_c = 0$ for all $c \in C$, then the processes $\{\kappa_{\cdot,c}\}_{c\in\mathcal{C}}$ are co-integrated.

Proof. Fix a $c^* \in \mathcal{C}$ and define $\tilde{\kappa}_{t,c} := \kappa_{t,c} - \kappa_{t,c^*}$ for any $c \in \mathcal{C}$. We then find for any $c \neq c^*$

$$\begin{aligned} \tilde{\kappa}_{t,c} &= (1-\lambda)(\kappa_{t-1,c} - \kappa_{t-1,c^*}) + \tilde{Z}_t \\ &= (1-\lambda)\tilde{\kappa}_{t-1,c} + \tilde{Z}_t, \qquad \qquad \tilde{Z}_t = \sigma_c Z_{t,c} - \sigma_{c^*} Z_{t,c^*}. \end{aligned}$$

Since $0 < \lambda \leq 1$ we obtain that $\tilde{\kappa}_{t,c}$ is a stationary AR(1) process for all $c \neq c^*$. Furthermore, we find that $m_t = \min_{a \in C} (\kappa_{t,a^*} + \tilde{\kappa}_{t,a}) = \kappa_{t,a^*} + \min_{a \in C} \tilde{\kappa}_{t,a^*}$ and the

Furthermore, we find that
$$m_t = \min_{c \in \mathcal{C}} (\kappa_{t,c^*} + \kappa_{t,c}) = \kappa_{t,c^*} + \min_{c \in \mathcal{C}} \kappa_{t,c}$$
 and therefore

$$\begin{aligned} \Delta \kappa_{t+1,c^*} &:= \kappa_{t+1,c^*} - \kappa_{t,c^*} &= \mu_{c^*} + \lambda(m_t - \kappa_{t,c^*}) + \sigma_{c^*} Z_{t+1,c^*} \\ &= \lambda \min_{c \in \mathcal{C}} \tilde{\kappa}_{t,c} + \mu_{c^*} + \sigma_{c^*} Z_{t+1,c^*}. \end{aligned}$$

The first term in the last expression is a minimum over stationary processes and the other terms are stationary too, hence $\Delta \kappa_{t,c^*}$ is stationary.

2.2 A analysis of the two dimensional case

To characterize the dynamics of minimum reversion, and to show that it induces a downward drift even for the case where the drift parameters μ_c are taken to be zero, we look at the simplest nontrivial case in two dimensions (C = 2). This means the dynamics of the time series become:

$$\kappa_{t+1,1} - \kappa_{t,1} = \lambda_1(\min_{c \in \{1,2\}} \kappa_{t,c} - \kappa_{t,1}) + \sigma_1 Z_{t+1,1} + \mu, \tag{4}$$

$$\kappa_{t+1,2} - \kappa_{t,2} = \lambda_2 (\min_{c \in \{1,2\}} \kappa_{t,c} - \kappa_{t,2}) + \sigma_2 Z_{t+1,2} + \mu.$$
(5)

We take $\lambda_1 = \lambda_2 := \lambda$ for the analysis in this section and assume that the correlation between the i.i.d. Gaussian variables $Z_{t+1,1}$ and $Z_{t+1,2}$ equals ρ for all t.

We define²

$$m_t = \min_{c \in \{1,2\}} \kappa_{t,c}$$
 and $M_t = \max_{c \in \{1,2\}} \kappa_{t,c}$

Proposition 2. For the model (4)-(5) with $\lambda_1 = \lambda_2 := \lambda$ we have that

$$\lim_{t \to \infty} \mathbb{E}[M_{t+1} - M_t] = \lim_{t \to \infty} \mathbb{E}[m_{t+1} - m_t] = \mu - s\sqrt{\frac{\lambda}{2\pi(2-\lambda)}}$$
(6)

and

$$\lim_{t \to \infty} \mathbb{E}[M_t - m_t] = s \sqrt{\frac{2}{\lambda \pi (2 - \lambda)}}$$
(7)

with $s = \sqrt{\sigma_1^2 + \sigma_2^2 - 2\rho\sigma_1\sigma_2}$.

Proof. Let the indices of the minimum and maximum be denoted by $i_t = \arg \min_{c \in \{1,2\}} \kappa_{t,c}$ and $I_t = \arg \max_{c \in \{1,2\}} \kappa_{t,c}$. We first determine the distribution of the minimum and maximum³. Substituting in (4)-(5) gives, for

$$E_t = M_t + \lambda (m_t - M_t) + \sigma_{I_t} Z_{t+1}^{I_t} - (m_t + \sigma_{i_t} Z_{t+1}^{i_t}),$$

that

$$\begin{aligned} m_{t+1} &= m_t + \sigma_{i_t} Z_{t+1}^{i_t} + \mu + E_t \mathbf{1}_{E_t < 0} \\ M_{t+1} &= M_t + \sigma_{I_t} Z_{t+1}^{I_t} + \mu + \lambda (m_t - M_t) - E_t \mathbf{1}_{E_t \ge 0}. \end{aligned}$$

We find

$$\begin{split} \mathbb{E}[m_{t+1}|(m_t, M_t)] &= m_t + \mu + (1-\lambda)(M_t - m_t)\mathbb{P}(E_t < 0) \\ &+ \mathbb{E}[(\sigma_{I_t} Z_{t+1}^{I_t} - \sigma_{i_t} Z_{t+1}^{i_t}) \mathbf{1}_{E_t < 0} \mid (m_t, M_t)] \\ \mathbb{E}[M_{t+1}|(m_t, M_t)] &= M_t + \mu + \lambda(m_t - M_t) - (1-\lambda)(M_t - m_t)\mathbb{P}(E_t > 0) \\ &- \mathbb{E}[(\sigma_{I_t} Z_{t+1}^{I_t} - \sigma_{i_t} Z_{t+1}^{i_t}) \mathbf{1}_{E_t > 0} \mid (m_t, M_t)]. \end{split}$$

The term $H := \sigma_{I_t} Z_{t+1}^{I_t} - \sigma_{i_t} Z_{t+1}^{i_t}$ is independent of (m_t, M_t) and has a Gaussian distribution with mean zero and variance $s^2 := \sigma_{i_t}^2 + \sigma_{I_t}^2 - 2\rho\sigma_{i_t}\sigma_{I_t} = \sigma_1^2 + \sigma_2^2 - 2\rho\sigma_1\sigma_2$ so $\mathbb{E}[H\mathbf{1}_{H\leq a}] = -se^{-(a/s)^2/2}/\sqrt{2\pi}$ for all $a \in \mathbb{R}$. We thus find that $\mathbb{E}[m_{t+1}|(m_t, M_t)]$ equals⁴

$$m_t + \mu + (1 - \lambda)(M_t - m_t)\Phi(-\frac{(1 - \lambda)(M_t - m_t)}{s}) - se^{-\frac{1}{2}((1 - \lambda)(M_t - m_t)/s)^2}/\sqrt{2\pi}$$

= $m_t + \mu + s(\tilde{D}_t\Phi(-\tilde{D}_t) - e^{-\frac{1}{2}\tilde{D}_t^2}/\sqrt{2\pi}) = m_t + \mu + sf(\tilde{D}_t),$ (8)

²If $\kappa_{t,1} = \kappa_{t,2}$ we let $m_t = \kappa_{t,1}$ and $M_t = \kappa_{t,2}$, but this event has zero probability for all $t \in \mathcal{T}$.

³Since the variables are Gaussian this distribution can be obtained directly as well, see Roberts [1966]. ⁴Here and in the sequel we use Φ to denote the cumulative probability function for a standard Gaussian random variable. if we define the shorthand notation $D_t = M_t - m_t$ and $\tilde{D}_t = (1 - \lambda)D_t/s$. The function $x \to f(x) = x\Phi(-x) - \exp(-x^2/2)/\sqrt{2\pi}$ increases monotonically from its minimum $f(0) = -1/\sqrt{2\pi} = -0.3989$ for x = 0 towards zero for $x = \infty$. Likewise,

$$\mathbb{E}[M_{t+1}|(m_t, M_t)] = M_t + \mu + \lambda(m_t - M_t) - s f(D_t).$$

Since $D_{t+1} = M_{t+1} - m_{t+1} = |(1-\lambda)D_t + s\tilde{H}|$ for a \tilde{H} with the standard normal distribution, we must have that $\tilde{D}_{t+1} = |(1-\lambda)^2 D_t/s + (1-\lambda)s\tilde{H}/s| = (1-\lambda) |\tilde{D}_t + \tilde{H}|$.

Let V(x) = |x| + 1, and define $\Delta V(x) = \mathbb{E}[V(\tilde{D}_{t+1}) - V(\tilde{D}_t) | \tilde{D}_t = x]$, then we have for $\beta = \lambda$ and $b = (1 - \lambda)\mathbb{E}|H|$ that $\Delta V(x) \leq -\beta V(x) + b$ which shows that the process \tilde{D} satisfies the geometric ergodicity conditions in Theorem 15.0.1 of Meyn et al. [2009] on the domain \mathbb{R}^+ . This proves that for any choice of $\tilde{D}_0 \geq 0$ the distribution function G_t of \tilde{D}_t converges to a stationary distribution G. This implies that for $x \geq 0$ the function G should satisfy

$$G(x) = \mathbb{P}(\tilde{D}_t \le x) = \mathbb{P}((1-\lambda) | \tilde{D}_t + \tilde{H} | \le x) = \int_0^\infty [N(\frac{x}{1-\lambda} - y) - N(-\frac{x}{1-\lambda} - y)] dG(y).$$

The solution to this equation is $G(x) = \mathbf{1}_{x\geq 0} \int_0^x 2\exp(-\frac{1}{2}(u/\tilde{\sigma})^2)/(\tilde{\sigma}\sqrt{2\pi})du$ with the parameter $\tilde{\sigma} = (1-\lambda)/\sqrt{\lambda(2-\lambda)}$, i.e. the stationary distribution of \tilde{D} is the distribution of |X| if $X \sim N(0, \tilde{\sigma}^2)$. The stationary distribution of $M_t - m_t = D_t = s\tilde{D}_t/(1-\lambda)$ is therefore the distribution of $s|X|/(1-\lambda)$, which gives (7).

Due to (8), the expected change in m_t under the stationary distribution for \tilde{D}_t equals μ plus the effect of the reversion to the minimum. This effect was shown above to be $s\mathbb{E}[f(\tilde{D}_t)] = s\int_0^{\infty} f(x)dG(x) = s(\tilde{\sigma} - \sqrt{1 + \tilde{\sigma}^2})/\sqrt{2\pi}$ which gives the second equality in (6) after some rewriting. The first one follows from the fact that \tilde{D} converges to a stationary distribution.

If we choose certain parameter values, the asymptotic drifts in proposition 2 can also be estimated by Monte Carlo simulations. We generated 10^5 paths to approximate the stationary distributions and used these to estimate the limit of the expectation of the extra drift (Table 1) and the limit of the expectation of the difference between the minimum and maximum (Table 2) for s = 1, $\mu = 0$ and in the absence of correlation between the increments for the two time series. Simulations also allow us to estimate the extra drift generated by the minimum reversion term in more than two dimensions, i.e. for $|\mathcal{C}| > 2$ and we show some examples Tables 1 and 2. As expected, we find that both the drift generated by the minimum reversion and the difference between the minimum and maximum increases with the population size C. The former increases and the latter decreases when the strength of the reversion to the minimum, which is determined by the parameter λ , increases.

3 Evidence for a Learning Effect in Mortality Rates

In this section we use the time series model in (2) to model mortality rates in several countries. The implicit assumption is that a population with the high mortality rate "copies" the behaviour of individuals in the population with low mortality. Such a model is consistent with the spread of medical advances and changes in behaviour like a reduction in smoking prevalence.

3.1 Modelling Mortality - The Common Age Effect Model

To obtain the time series, we first fit a stochastic mortality model to observed death counts. We assume that the number of deaths, D_{xtc} , in population $c \in C$ at age $x \in \mathcal{X}$ in calendar year $t \in \mathcal{T}$ is

Table 1: Generated drift for different minimum reversion parameters λ according to (6) (second column) and using 10⁵ simulations (third to last column).

	Exact	Simulation					
λ	$ \mathcal{C} = 2$	$ \mathcal{C} = 2$	$ \mathcal{C} = 3$	$ \mathcal{C} = 4$	$ \mathcal{C} = 8$	$ \mathcal{C} = 16$	$ \mathcal{C} = 32$
0.0125	-0.0447	-0.0448	-0.0671	-0.0817	-0.1129	-0.1401	-0.1641
0.025	-0.0635	-0.0635	-0.0952	-0.1158	-0.1602	-0.1987	-0.2329
0.05	-0.0903	-0.0903	-0.1355	-0.1649	-0.2280	-0.2828	-0.3314
0.1	-0.1294	-0.1294	-0.1942	-0.2362	-0.3266	-0.4052	-0.4748
0.2	-0.1881	-0.1881	-0.2821	-0.3432	-0.4745	-0.5887	-0.6899
0.4	-0.2821	-0.2821	-0.4231	-0.5147	-0.7118	-0.8830	-1.0348

Table 2: Expectation of the stationary distribution for $M_t - m_t$ for different minimum reversion parameters λ according to (7) (second column) and using 10⁵ simulations (third to last column).

	Exact	Simulation					
λ	$ \mathcal{C} = 2$	$ \mathcal{C} = 2$	$ \mathcal{C} = 3$	$ \mathcal{C} = 4$	$ \mathcal{C} = 8$	$ \mathcal{C} = 16$	$ \mathcal{C} = 32$
0.0125	7.1589	7.1593	10.7386	13.0627	18.0644	22.4106	26.2618
0.025	5.0781	5.0774	7.6185	9.2656	12.8138	15.8962	18.6282
0.05	3.6137	3.6132	5.4202	6.5931	9.1178	11.3112	13.2565
0.1	2.5887	2.5886	3.8831	4.7229	6.5316	8.1031	9.4965
0.2	1.8806	1.8806	2.8209	3.4313	4.7454	5.8866	6.8988
0.4	1.4105	1.4104	2.1156	2.5735	3.5591	4.415	5.1742

a random variable with a Poisson distribution, that is,

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$$D_{xtc} \sim \operatorname{Pois}\left(\mu_{xtc} E_{xtc}\right) \tag{9}$$

where μ_{xtc} is the hazard rate (also known as the "force of mortality" in the literature) and E_{xtc} refers to the central exposure to risk.

We then use a stochastic model for the force of mortality μ_{xtc} that incorporates the populationspecific time series $\{\kappa_t\}_{t\in\mathcal{T}}$ defined in (2). As we are modelling the mortality in multiple populations simultaneously and wish to make our model suitable for a wide age range, we use a modification of the Lee-Carter model (Lee and Carter [1992]) with common age effects, as suggested by Kleinow [2015]:

$$\log \mu_{xtc} = \alpha_x + \beta_x \kappa_{t,c} \tag{10}$$

where the age effects α_x and β_x do not depend on the population c. Having age effects that are common to all populations ensures that the individual components $\kappa_{t,c}$ (known as "period effects") are comparable across populations as they are rescaled by a common vector β and shifted by a common vector α . The parameters in (10) are not identifiable since

$$\alpha_x + \beta_x \kappa_{t,c} = \tilde{\alpha}_x + \tilde{\beta}_x \tilde{\kappa}_{t,c} \quad \text{when} \quad \tilde{\alpha}_x = \alpha_x - \frac{K_1}{K_2} \beta_x, \quad \tilde{\beta}_x = \frac{\beta_x}{K_2}, \quad \tilde{\kappa}_{t,c} = K_1 + K_2 \kappa_{t,c}$$

for any real numbers K_1 and $K_2 \neq 0$. To identify a unique set of parameters, we impose the following constraints on the parameter vectors α and β :

$$\alpha_{x_r} = 0$$
 and $\beta_{x_r} = 1$



Figure 1: Estimated values of α_x (left) and β_x (right) for the mortality rates of females based on data for the period 1921–2011. We can clearly see the phenomenon known as "age heaping": when people report the death of a person without knowing their exact age, they often report the age to be a multiple of 10.

for a fixed reference age $x_r \in \mathcal{X}$. Applying those two constraints means that $\log \mu_{xtc} = \kappa_{t,c}$ for $x = x_r$ in every population $c \in \mathcal{C}$. In other words, we can interpret the period effect $\kappa_{t,c}$ as the fitted log mortality rate at the reference age x_r in population c. For our empirical analysis we choose the reference age $x_r = 70$.

We estimate the age effects α and β and the period effects κ using the maximum likelihood method based on the Poisson model in (9) for observed deaths counts D_{xtc} and exposure values E_{xtc} .

3.2 Data

The empirical death counts and exposure data have been obtained from the Human Mortality Database (HMD), see HMD [2018]. We consider data for ages $\mathcal{X} = \{0, 1, \ldots, 95\}$ and for two ranges of calendar years: $\mathcal{T} = \{1951, \ldots, 2011\}$ or $\mathcal{T} = \{1921, \ldots, 2011\}$. Table 3 shows a list of countries included in our study together with the HMD country codes. As indicated in the table, there are some countries for which data from 1921 are not available and those countries have been excluded from our empirical study for that range or years.

3.3 Empirical Results

Figure 1 shows the estimated values of the age effects α_x and β_x obtained from data for females based on the observation period 1921–2011. The age effects for the other data sets are not shown since they all have a very similar shape. To illustrate our data and the effect of the minimum reversion effect, Figure 2 shows the estimated values of $\kappa_{t,c}$ for females for the years 1921–2011 in the dataset, together with two projected scenarios based on the model in (2). On the lefthand side the projection includes the minimum reversion effect based on the estimated parameter λ and on the righthand side we set $\lambda = 0$.

Table 3: List of countries included in our study. For all countries mortality data for males and females are considered for calendar years 1951–2011. The table also indicates which countries are included in our study for the longer observation period 1921–2011. Note that for France and England & Wales the total populations are considered rather than the civilian population.

HMD Code	1921 - 2011	Country	HMD Code	1921 - 2011	
NLD	\checkmark	Sweden	SWE	\checkmark	
DNK	\checkmark	Belgium	BEL	\checkmark	
FIN	\checkmark	England & Wales	GBRTENW	\checkmark	
FRATNP	\checkmark	Switzerland	CHE	\checkmark	
AUS	\checkmark	Italy	ITA	\checkmark	
AUT		Ireland	IRL		
NOR	\checkmark	Japan	JAP		
CAN	\checkmark	New Zealand	NZL_NP		
\mathbf{PRT}		Spain	ESP	\checkmark	
USA		Iceland	ISL	\checkmark	
	HMD Code NLD DNK FIN FRATNP AUS AUT NOR CAN PRT USA	HMD Code 1921–2011 NLD ✓ DNK ✓ FIN ✓ FRATNP ✓ AUS ✓ AUT ✓ NOR ✓ CAN ✓ PRT USA	HMD Code1921–2011CountryNLD \checkmark SwedenDNK \checkmark BelgiumFIN \checkmark England & WalesFRATNP \checkmark SwitzerlandAUS \checkmark ItalyAUTIrelandNOR \checkmark JapanCAN \checkmark New ZealandPRTSpainUSA	$\begin{array}{ccccc} \mathrm{HMD}\ \mathrm{Code} & 1921–2011 & \mathrm{Country} & \mathrm{HMD}\ \mathrm{Code} \\ & & & & & \\ \mathrm{NLD} & \checkmark & \mathrm{Sweden} & \mathrm{SWE} \\ \mathrm{DNK} & \checkmark & \mathrm{Belgium} & \mathrm{BEL} \\ \mathrm{FIN} & \checkmark & \mathrm{England}\ \& \ \mathrm{Wales} & \mathrm{GBRTENW} \\ \mathrm{FRATNP} & \checkmark & \mathrm{Switzerland} & \mathrm{CHE} \\ \mathrm{AUS} & \checkmark & \mathrm{Italy} & \mathrm{ITA} \\ \mathrm{AUT} & & \mathrm{Ireland} & \mathrm{IRL} \\ \mathrm{NOR} & \checkmark & \mathrm{Japan} & \mathrm{JAP} \\ \mathrm{CAN} & \checkmark & \mathrm{New}\ \mathrm{Zealand} & \mathrm{NZL}\ \mathrm{NP} \\ \mathrm{PRT} & & \mathrm{Spain} & \mathrm{ESP} \\ \mathrm{USA} & \mathrm{Iceland} & \mathrm{ISL} \end{array}$	

To investigate the significance of the learning effect, we compare the Bayesian Information Criterion (BIC) for models without minimum reversion ($\lambda = 0$) with the more general model in (2). If K denotes the number of parameters and $n = |\mathcal{C}||\mathcal{T}|$ is the total number of observations across all populations and all years, the BIC value equals $K \log n - 2 \log L$, where L is the maximum value of the likelihood function L for the model in (2). This means

$$2\log L = -n\log|\Sigma| - \sum_{t=t_0+2}^{t_1} (\Delta \kappa_t - \eta_{t-1})' \Sigma^{-1} (\Delta \kappa_t - \eta_{t-1}) + \Gamma$$

where Γ is a constant which does not depend on the parameters that must be estimated, $\Delta \kappa_t$ and η_t are vectors with components

$$\Delta \kappa_{t,c} = \kappa_{t,c} - \kappa_{t-1,c}$$
 and $\eta_{t,c} = \mu_c + \zeta_c \Delta \kappa_{t,c} + \lambda_c (m_t - \kappa_{t,c}),$

and Σ is the covariance matrix of $\Delta \kappa_t$ so $\Sigma_{ij} = \sigma_i \sigma_j \rho_i \rho_j$ for $i \neq j$ and $\Sigma_{ii} = \sigma_i^2$.

We determined the BIC values for different specifications in which parameter values may or may not be constrained to be the same for all different groups in C. Based on this analysis, the parameters σ_c and ρ_c are taken population-specific while the parameters $\mu_c = \mu$, $\lambda_c = \lambda$ and $\zeta_c = \zeta$ are common to all populations. The obtained BIC values and parameter estimates⁵ are shown in Table 4 for four data sets.

We notice that the BIC values always improve when λ is not restricted to be zero which shows that the learning effect adds to the goodness of fit of the model. We also observe that the estimated drift parameter $\hat{\mu}$ is reduced when minimum reversion is included in the model, from which we conclude that some of the mortality improvements in the populations are driven by learning effects from others who have lower mortality rates.

⁵We do not show the values ρ_c and σ_c for every group c in C but only report $\bar{\rho}$, the average of ρ_c over all groups c, and $\bar{\sigma}$, which is defined by the requirement that $\bar{\sigma}^2$ is the average of σ_c^2 over all groups c. Estimated parameter values for all countries, genders and data periods are available upon request.

Table 4: Goodness of fit and parameter values of the model in (2) without and with minimum reversion, estimated for different data sets.

Model	$\log L$	K	BIC	$\hat{\mu}$	$\hat{\lambda}$	$\hat{\zeta}$	$\bar{\sigma}$	$\bar{ ho}$
Females	in calenda	ar yea	ars 1921–20	11				
$\lambda = 0$	3886.16	30	-7558.49	-0.0224		-0.2998	0.0338	0.4996
$\lambda \neq 0$	3890.61	31	-7560.27	-0.0194	0.0191	-0.2895	0.0336	0.4977
Males in	ı calendar	years	3 1921-2011					
$\lambda = 0$	4151.49	30	-8089.15	-0.0150		-0.1890	0.0297	0.5207
$\lambda \neq 0$	4155.65	31	-8090.35	-0.0131	0.0185	-0.1804	0.0295	0.5224
Females	in calenda	ar yea	ars 1951–20	11				
$\lambda = 0$	3854.23	42	-7411.38	-0.0210		-0.3414	0.0288	0.4717
$\lambda \neq 0$	3861.62	43	-7419.08	-0.0161	0.0177	-0.3394	0.0285	0.4641
Males in	ı calendar	years	1951 - 2011					
$\lambda = 0$	4100.51	42	-7903.94	-0.0246		-0.3320	0.0252	0.5845
$\lambda \neq 0$	4105.98	43	-7907.81	-0.0192	0.0167	-0.3219	0.0250	0.5832

4 Conclusions and Further Research

We have shown that model specifications that include a minimum reversion term resulted in better BIC values than specifications without such a term when they were fitted to several datasets from the human mortality database. Visual inspection of the projections generated by the time series with minimum reversion in 2 also shows a clear improvement. This testifies to the usefulness of the incorporation of a "learning effect" in the time series.

Several extensions of the proposed model are possible. In particular, the minimum in (2) could be replaced by other rank statistics. Another possible direction of research is the study of a continuous time version of minimum reversion, where the drift term of a diffusion process in multiple dimensions is a function of the distance of the process to the minimum of its components. The properties of these processes could then be compared to the rank-based diffusion processes for particle systems mentioned in the Introduction of this paper.

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Figure 2: Estimated values of $\kappa_{t,c}$ for the mortality rates of females based on data for the period 1921–2011. The simulated scenarios are based on the full model in (2) (left) and the nested model with $\lambda = 0$ (right).