Application of GTH-like algorithm to Markov modulated Brownian motion with jumps

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Abstract

The Markov modulated Brownian motion is a substantial generalization of the classical Brownian Motion. On the other hand, the Markovian arrival process (MAP) is a point process whose family is dense for any stochastic point process and is used to approximate complex stochastic counting processes. In this paper, we consider a superposition of the Markov modulated Brownian motion (MMBM) and the Markovian arrival process of jumps which are distributed as the bilateral ph-type distribution, the class of which is also dense in the space of distribution functions defined on the whole real line. In the model, we assume that the inter-arrival times of the MAP depend on the underlying Markov process of the MMBM. One of the subjects of this paper is introducing how to obtain the first passage probabilities of the superposed process using a stochastic doubling algorithm designed for getting the minimal solution of a nonsymmetric algebraic Riccatti equation. The other is to provide eigenvalue and eigenvector results on the superposed process to make it possible to apply the GTH-like algorithm, which improves the accuracy of the doubling algorithm.

Keywords: Markov modulated Brownian motion, Markovian arrival process, first passage time, doubling algorithm, nonsymmetric algebraic Riccati equation, GTH-like algorithm

1. Introduction

This paper is concerned with the superposed process of the Markov modulated Brownian motion (MMBM) and Markovian arrival process (MAP), and a technique to enhance the performance of an algorithm for computation of the first passage probabilities of the superposed process.

The MMBM, a generalization of the Brownian motion, is a bivariate Markov process of which the drift and diffusion parameters vary according to the states of a underlying Markov process. On the other hand, the MAP is a point process whose jump epochs are also governed by an underlying Markov process. Since the family of MAP is dense for any stochastic point process, it is used to approximate complex stochastic counting processes. Concerning the sizes of jumps of the MAP, we assume that they are mutually independent and distributed as the bilateral ph-type (BPH) distributions with their parameters depending on the states of the underlying Markov processes of both the MMBM and MAP. We note that the bilateral ph-type distribution, introduced by Ahn and Ramaswami (2005), can be represented as a mixture of two independent ph-type distributions, and the class of bilateral ph-type distributions is also dense in the space of distribution functions defined on the whole real line.

Then, the superposition of the MMBM and MAP becomes a generalized process that can be used for modelling complex systems in numerous academic fields including queues, finance, and insurance

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risk theories. For example, of applications, see Belhaj (2010), Jiang and Pistorius (2008), Asmussen (2003), and the references therein. We call this superposed process the generalized Markov modulated Brownian motion and use G-MMBM for abbreviation. In relation to G-MMBM, we deal with two main subjects in this paper. The first one is to provide Markovian representation of the G-MMBM so that we can obtain the first passage probabilities of the G-MMBM using the approaches of Ahn (2016), in which they show that the first passage probabilities can be achieved by using the minimal non-negative solution of a nonsymmetric algebraic Riccati equation (NARE) of the form

$$AZ + ZB + ZCZ + D = \mathbf{0}.$$
 (1.1)

The other is to show eigenvalue and eigenvector results so that we can use the GTH-like algorithm introduced by Alfa *et al.* (2001), which enhances the accuracy of the algorithms for computing the minimal nonnegative solution of the NARE in relation to the first passage probabilities of the G-MMBM.

The remainder of this paper is organized as follows. In Section 2, we introduce the G-MMBM and also the relation between the first passage probabilities of the process and the minimal nonnegative solution of a nonsymmetric algebraic Riccati equation. In Section 3, we provide parameter representation of the G-MMBM which can be obtained using embedding technique. The doubling algorithm and eigenvalue and eigenvector results on the G-MMBM to be used for GTH-like algorithm are introduced in Section 3. Finally, we illustrate the effects of the GTH-like algorithm by using numerical examples.

2. Description of the G-MMBM and related NARE

The G-MMBM is the superposed process of an MMBM and an MAP. We note that the MMBM consists of a modulating irreducible Markov process J_B with a state space $S^{(B)} = S_b^{(B)} \cup S_u^{(B)} \cup S_d^{(B)}$, a drift row-vector $\boldsymbol{\mu}^{(B)} = (\boldsymbol{\mu}_b^{(B)} \, \boldsymbol{\mu}_u^{(B)} \, \boldsymbol{\mu}_d^{(B)})$, and a diffusion row-vector $\boldsymbol{\sigma}^{(B)} = (\boldsymbol{\sigma}_b^{(B)} \, \boldsymbol{0} \, \boldsymbol{0})$; the MAP is modulated by an irreducible Markov process J_M with a state space $S^{(M)} = \{1, \ldots, m\}$ and its transition and jump rates depend on J_B . The resulting G-MMBM is a two-dimensional Markov process and use (X, J) to denote the process, where we call X and J the level and the phase process of the G-MMBM, respectively. Here, the phase process J is the superposed process of J_B and J_M of which the state space S is represented as $S = S_b \cup S_u \cup S_d$ with $S_b = S_b^{(B)} \otimes S^{(M)}$, $S_u = S_u^{(B)} \otimes S^{(M)}$, and $S_d = S_d^{(B)} \otimes S^{(M)}$. Then the level process X can be represented as

$$X(t) = a + \int_0^t \mu_{J(u)} du + \int_0^t \sigma_{J(u)} dB(u) + \int_0^t dM(u),$$
(2.1)

where (i) when $J(t) = (i, j) \in S$, $\mu_{(i,j)} = [\mu^{(B)}]_i$ and $\sigma_{(i,j)} = [\sigma^{(B)}]_i$; (ii) *M* is the jump process with the jump epochs being either the transition epochs of *J*, or the arrival epochs of the compound Poisson processes whose intensity depends on the state of *J*. Moreover, the sizes of the incurred jumps are mutually independent and distributed as the BPH distribution of the form

$$G(dx) = \beta^{+} e^{\mathbf{B}^{+} x} (-\mathbf{B}^{+} \mathbf{1}) dx \, \chi(x > 0) + \beta^{-} e^{-\mathbf{B}^{-} x} (-\mathbf{B}^{-} \mathbf{1}) dx \, \chi(x < 0),$$
(2.2)

in which nonnegative probability vectors β^+ and β^- satisfy $\beta^+ \mathbf{1} + \beta^+ \mathbf{1} = 1$, and \mathbf{B}^+ and \mathbf{B}^- are substochastic matrices which have negative diagonal elements, nonnegative off-diagonal elements, and non-positive row-sums; (iii) and $B = \{B(t), t \ge 0\}$ is a standard Brownian motion that is independent of *J* and *M*.

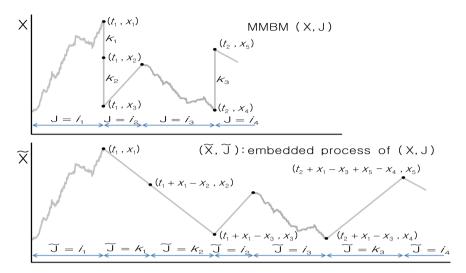


Figure 1: Illustration of how a sample path of (F, J) is transformed to a sample path of its embedded process (\tilde{F}, \tilde{J}) . Note that $i_1, i_3 \in S_b, i_2 \in S_u, i_4 \in S_d, k_1, k_2 \in S_-$, and $k_3 \in S_+$.

In connection with the state spaces *S*, we denote the drift and diffusion row vectors by $\boldsymbol{\mu} = (\mu_i, i \in S)$ and $\boldsymbol{\sigma} = (\sigma_i, i \in S)$, respectively, and we define the sub-vectors $\boldsymbol{\sigma}_k$ and $\boldsymbol{\mu}_k$ as $\boldsymbol{\sigma}_k = (\sigma_i, i \in S_k)$ and $\boldsymbol{\mu}_k = (\mu_i, i \in S_k)$ for k = b, u, d. In addition to the state space *S*, it is necessary to adopt jump-state spaces S_+ and S_- modulating bilateral ph-type jumps, details of which are to be given in Section 3. For these state spaces, we define the drift and diffusion row vectors as $\boldsymbol{\mu}_+ = \mathbf{1}'_+, \boldsymbol{\mu}_- = \mathbf{1}'_-, \boldsymbol{\sigma}_{j^+} = \mathbf{0}_+$, and $\boldsymbol{\sigma}_- = \mathbf{0}_-$ with the prime denoting the transpose operator. Note that, for $k \in \{b, u, d, +, -\}$, we use $\mathbf{0}_k$, $\mathbf{1}_k$, I_k , and $\mathbf{0}_{k,l}$ to denote the s_k (the dimension of S_k)-dimensional row vector of 0s, column vector of 1s, s_k -dimensional identity matrix, and $s_k \times s_l$ -dimensional zero matrix, respectively. We also use *I* and **0** to denote the identity matrix and the zero matrix of appropriate dimension which can be identified from the context. Throughout the paper, for a given vector \mathbf{v} , we use the notation $\Delta_{\mathbf{v}}$ to denote the diagonal matrix with the elements of \mathbf{v} on its diagonal.

2.1. Representation of the G-MMBM through the embedding technique and completed graph

Let (\tilde{X}, \tilde{J}) denote the embedded process of (X, J) to be obtained by using an embedding technique, which replaces the positive and negative jumps in (X, J) with linear stretches having slopes +1 and -1 respectively in \tilde{X} , and lets \tilde{J} track the states (phases) in BPH jumps in addition to the states in J. In Figure 1, we illustrate how a sample path of (X, J) is transformed to its corresponding sample path of (\tilde{X}, \tilde{J}) . Then, the embedded process (\tilde{X}, \tilde{J}) becomes an MMBM without a jump, in which its state space $S^{\mathbf{e}}$, diffusion vector $\sigma^{\mathbf{e}}$, and drift vector $\mu^{\mathbf{e}}$ are respectively given as $S^{\mathbf{e}} = S_b \cup S_u \cup S_+ \cup S_d \cup S_-$, $\sigma^{\mathbf{e}} = (\sigma_b \mathbf{0}_u \mathbf{0}_+ \mathbf{0}_d \mathbf{0}_-)$, and $\mu^{\mathbf{e}} = (\mu_b \mu_u \mu_+ \mu_d \mu_-)$. We denote by T the infinitesimal generator of \tilde{J} and represent it in a partitioned form as

$$T = (T_{i,j}, i, j = b, u, +, d, -) \text{ with } T_{+,-} = \mathbf{0} \text{ and } T_{-,+} = \mathbf{0},$$
(2.3)

where $T_{i,j}$ is a sub-matrix of T containing $[T]_{lm}$, the $(l, m)^{th}$ element of T with $l \in S_i$ and $m \in S_j$. We note that the restriction of T on S should be Q, and that $(-[T]_{ii})^{-1}$, $i \in S$, represents the mean dwelling

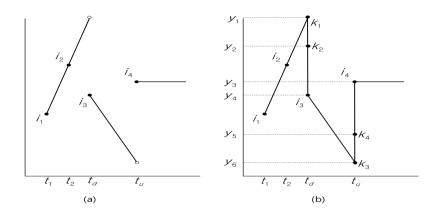


Figure 2: (a) Level process X and phases of J. (b) Completed graph \mathscr{C} and phases of \mathscr{J} .

time of *J* in *i* until a jump or a transition to other states in *S* occurs, whereas $(-[Q]_{ii})^{-1}$ is the mean dwelling time of *J* in *i* until a transition of *J* to other states in *S*. We let $\pi^{\mathbf{e}} = (\pi_b \ \pi_u \ \pi_+ \ \pi_d \ \pi_-)$ be the stationary probability vector of *T* satisfying $\pi^{\mathbf{e}}T = \mathbf{0}$ and $\pi^{\mathbf{e}}\mathbf{1} = 1$. Defining the average drift $\tilde{\mu}$ of the embedded process as $\tilde{\mu} = \sum_{i \in S^{\mathbf{e}}} [\pi^{\mathbf{e}}]_i [\mu^{\mathbf{e}}]_i$, we say that the G-MMBM is positively recurrent, transient, and null recurrent when $\tilde{\mu} < 0, > 0$, and = 0, respectively. We describe the detailed constructions of *S*^e and *T* in Section 3.

All the information for analyzing the G-MMBM is contained in the parameter triple (T, μ^e, σ^e) . So, when we call the parameter of the G-MMBM, it indicates the parameter triple. With regard to the level process \tilde{X} , it is represented as

$$\tilde{X}(t) = a + \int_0^t [\boldsymbol{\mu}^{\mathbf{e}}]_{\tilde{J}(u)} du + \int_0^t [\boldsymbol{\sigma}^{\mathbf{e}}]_{\tilde{J}(u)} dB(u).$$

For description of G-MMBM, it is useful to introduce a terminology called the completed graph \mathscr{C} of the process and its phase process $\mathscr{J}(a, t)$ to be defined for $(a, t) \in \mathscr{C}$. We note that the completed graph \mathscr{C} is a subset of the plane \mathbb{R}^2 containing (X(t), t) for all t and the line segment joining (X(t), t) and (X(t-), t) for all discontinuity points t. The completed graph, introduced in Whitt (1980), is useful for representing quantities in relation to a process with discontinuous sample paths. For more details, refer to Whitt (1980) and Ahn (2016, 2017). We illustrate a G-MMBM, its completed graph, and the phase process defined on the completed graph in Figure 2.

2.2. First passage time of the G-MMBM and a nonsymmetric algebraic Riccati equation

We define the first passage times of X to 0 as $\tau = \inf\{t > 0 : X(t) < 0\}$. In relation to τ , we define an probability matrix $\hat{\mathbf{f}}(a)$ such that its $(i, j)^{th}$ element, $i, j \in S^{\mathbf{e}}$, satisfies

$$\left[\hat{\mathbf{f}}(a)\right]_{ij} = P\left[\mathcal{J}(0,\tau) = j|\mathcal{J}(a,0) = i\right], \ a \ge 0.$$

Then, the $(s_b + s_d + s_-)$ -dimensional squared sub-matrix $\hat{\mathbf{f}}_{bd-,bd-}(a) = ([\hat{\mathbf{f}}(a)]_{ij}; i, j \in S_b \cup S_d \cup S_-)$ has the matrix exponential form $e^{\mathbf{H}a}$, where **H** is a sub-stochastic matrix.

Concerning H, Ahn (2016) showed that it can be obtained by using the minimal solution of a

non-symmetric algebraic Riccati equation of the form

$$\mathbf{A}Z + Z\mathbf{B} + Z\mathbf{C}Z + \mathbf{D} = \mathbf{0},\tag{2.4}$$

where, with definition of the following relevant matrices

$$\mathbf{\Lambda} = \text{diag}\{-[T]_{ii}, i \in S_b\}, \mathbf{\Sigma} = \mathbf{\Delta}_{\boldsymbol{\sigma}_b}, D_1 = \mathbf{\Sigma}^{-2} \mathbf{\Delta}_{\boldsymbol{\mu}_b}, \text{ and } D_2 = \mathbf{\Sigma}^{-1} \left(2\mathbf{\Lambda} + \mathbf{\Sigma}^{-2} \mathbf{\Delta}_{\boldsymbol{\mu}_b}^2 \right)^{\frac{1}{2}},$$

the coefficient matrices satisfy

$$\mathbf{A} = \begin{pmatrix} D_{1}-D_{2} & 2\Sigma^{-1}T_{b,u} & 2\Sigma^{-1}T_{b,+} \\ \mathbf{0}_{u,b} & \Delta_{\mu}^{-1}T_{u,u} & \Delta_{\mu}^{-1}T_{u,+} \\ \mathbf{0}_{+,b} & T_{+,u} & T_{+,+} \end{pmatrix}, \qquad \mathbf{C} = \begin{pmatrix} \Sigma^{-1} & \mathbf{0}_{b,u} & \mathbf{0}_{b,+} \\ \mathbf{0}_{d,b} & \Delta_{-\mu_{d}}^{-1}T_{d,u} & \Delta_{-\mu_{d}}^{-1}T_{d,+} \\ \mathbf{0}_{-,b} & T_{-,u} & \mathbf{0}_{-,+} \end{pmatrix},$$

$$\mathbf{B} = \begin{pmatrix} -D_{1}-D_{2} & \mathbf{0}_{b,d} & \mathbf{0}_{b,-} \\ \Delta_{-\mu_{d}}^{-1}T_{d,b} & \Delta_{-\mu_{d}}^{-1}T_{d,d} & \Delta_{-\mu_{d}}^{-1}T_{d,-} \\ T_{-,b} & T_{-,d} & T_{-,-} \end{pmatrix}, \qquad \mathbf{D} = \begin{pmatrix} 2\Sigma^{-1}(T_{b,b}+\mathbf{A}) & 2\Sigma^{-1}T_{b,d} & 2\Sigma^{-1}T_{b,-} \\ \Delta_{\mu_{u}}^{-1}T_{u,b} & \Delta_{\mu_{u}}^{-1}T_{u,d} & \Delta_{\mu_{u}}^{-1}T_{u,-} \\ T_{+,b} & T_{+,d} & \mathbf{0}_{+,-} \end{pmatrix}.$$
(2.5)

Furthermore, from the results of Section 2.1.2 of Bini *et al.* (2012), Liu and Xue (2012), and Theorem 3.1.1 and 3.1.2 of Ahn (2016), we can derive the following result:

Theorem 1. Let $\sigma(\cdot)$ denote the set of eigenvalues of a given squared matrix \cdot . (a) The NARE (2.4) has a minimal nonnegative solution Z. Moreover, Z is the unique solution such that $\sigma(\mathbf{B} + \mathbf{CZ}) \subset \mathbb{C}_{-} \cup \{0\}$ and $\sigma(\mathbf{A} + Z\mathbf{C}) \subset \mathbb{C}_{-} \cup \{0\}$.

(b) Let Z be the minimal non-negative solution of the NARE (2.4). Then \mathbf{H} satisfies the relation

$$\mathbf{H} = \mathbf{B} + \mathbf{C}Z. \tag{2.6}$$

3. Representation of the superposed process as a G-MMBM

We introduce how to represent the G-MMBM which is the superposed process of an MMBM and Markovian arrival processes (MAP) with BPH-distributed jumps. Since the transition rates and jump rates of the MAP are assumed to be dependent on the process J_B modulating the MMBM, we denote the MAP parameters by $(\Gamma_0^{J_B(t)}, \Gamma_1^{J_B(t)})$. We also assume that MAP jumps are mutually independent and distributed as the BPH distribution of which the parameters depend on the state of the superposed Markov process J. We note that the representation form of the G-MMBM depend on dependence structures of the jumps on the Markov process J.

3.1. BPH jumps depending on only the Markov states before their corresponding jump epochs

In this section, we assume that the jump sizes are mutually independent and distributed as the BPH distribution, of which the parameters depend on the state of J just before their jump epochs. We begin with simple examples for illustration.

Example 1. We consider the superposition of a two-state MMBM and two-state MAP, where

- (i) $S^{(B)} = \{1, 2\}, Q^{(B)} = (q_{ij}), \mu^{(B)} = (\mu_1 \ \mu_2) \text{ and } \sigma^{(B)} = (\sigma_1 \ \sigma_2),$
- (ii) two-state MAP($\Gamma_0^{J_B(t)}, \Gamma_1^{J_B(t)}$) with $S^{(M)} = \{1, 2\}, \Gamma_0^i = (\gamma_{0,kl}^i; k, l \in S^{(M)})$ and $\Gamma_1^i = (\gamma_{1,kl}^i; k, l \in S^{(M)})$ for $i \in S^{(B)}$,

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(iii) an MAP jump occurring at the transition epoch of J_M from k to l while J_B stays in $i \in S^{(B)}$ is assumed to be distributed as ph-type distribution of the form

$$\boldsymbol{\beta}_{k}^{i}e^{B_{k}^{i}x}\left(-B_{k}^{i}\mathbf{1}\right)\chi(x>0).$$

Then, the infinitesimal generator of the embedded process of the superposition can be represented as

($q_{11} + \gamma^1_{0;11}$	$\gamma^1_{0;12}$	q_{12}	0	$(\gamma_{1;11}^1 + \gamma_{1;12}^1)\beta_1^1$	0	0	0)
	$\gamma^1_{0;21}$	$q_{11} + \gamma_{0;22}^1$	0	q_{12}	0	$(\gamma_{1;21}^1 + \gamma_{1;22}^1)\beta_2^1$	0	0
	q_{21}	0	$q_{22} + \gamma_{0;11}^2$	$\gamma^{2}_{0;12}$	0	0	$(\gamma_{1;11}^2{+}\gamma_{1;12}^2)\beta_1^2$	0
	0	q_{21}	$\gamma^{2}_{0;21}$	$q_{22} + \gamma_{0;22}^2$	0	0	0	$(\gamma_{1;21}^2+\gamma_{1;22}^2)\beta_2^2$
	$(-B_1^11)\bar{\gamma}_{1;11}^1$	$(-B_1^11)\bar{\gamma}_{1;12}^1$	0	0	B_1^1	0	0	0
	$(-B_2^11)\bar{\gamma}_{1:21}^1$	$(-B_2^11)\bar{\gamma}_{1;22}^1$	0	0	0	B_2^1	0	0
	0	0	$(-B_1^2 {\bf 1}) \bar{\gamma}_{1;11}^2$	$(-B_1^2 1) \bar{\gamma}_{1;12}^2$	0	0	B_{1}^{2}	0
l	0	0	$(-B_2^2 {\bf 1}) \bar{\gamma}_{1;21}^2$	$(-B_2^2 {\bf 1}) \bar{\gamma}_{1;22}^2$	0	0	0	B_2^2)

where $\bar{\gamma}_{1;kl}^i = \gamma_{1;kl}^i / (\gamma_{1;k1}^i + \gamma_{1;k2}^i)$ for $i \in S^{(B)}$ and $k, l \in S^{(M)}$. The drift and diffusion vectors are given as $(\mu_1 \ \mu_1 \ \mu_2 \ \mu_2 \ \mathbf{1}' \ \mathbf{1}' \ \mathbf{1}')$ and $(\sigma_1 \ \sigma_1 \ \sigma_2 \ \sigma_2 \ \mathbf{0} \ \mathbf{0} \ \mathbf{0} \ \mathbf{0})$, respectively. \Box

Example 2. We consider the special case of Example 3.1. In addition to the assumption (i) and (ii) of Example 3.1, we assume that (ii)' $\gamma_{1:k\ell}^i / \sum_{\ell \in S^{(M)}} \gamma_{1:k\ell}^i = \bar{\gamma}_{1:\ell}^i$ for all $k \in S^{(M)}$; (iii) an MAP jump occurring at the transition epoch of J_M from k to l while J_B stays in $i \in S^{(B)}$ is assumed to be distributed as BPH distribution of the form

$$\boldsymbol{\beta}_{k}^{i-}e^{B^{i-}|x|}\left(-B^{i-}\mathbf{1}\right)\chi(x<0)+\boldsymbol{\beta}_{k}^{i+}e^{B^{i+}x}\left(-B^{i+}\mathbf{1}\right)\chi(x>0).$$

Then, the drift and diffusion vectors are given as $(\mu_1 \mu_1 \mu_2 \mu_2 \mathbf{1'} \mathbf{1'} - \mathbf{1'} - \mathbf{1'})$ and $(\sigma_1 \sigma_1 \sigma_2 \sigma_2 \mathbf{0} \mathbf{0} \mathbf{0} \mathbf{0})$, respectively, and the infinitesimal generator of the embedded process can be represented as

($q_{11} + \gamma_{0;11}^1$	$\gamma^1_{0;12}$	q_{12}	0	$(\gamma_{1;11}^1 + \gamma_{1;12}^1)\beta_1^{1+}$	0	$(\gamma_{1;21}^1 + \gamma_{1;22}^1)\beta_1^{1-}$	0)
	$\gamma^1_{0;21}$	$q_{11} + \gamma_{0;22}^1$	0	q_{12}	$(\gamma^1_{1;21}{+}\gamma^1_{1;22})\beta_2^{1+}$	0	$(\gamma^1_{1;21}{+}\gamma^1_{1;22})\beta_2^{1-}$	0
	q_{21}	0	$q_{22} + \gamma_{0;11}^2$	$\gamma^{2}_{0;12}$	0	$(\gamma_{1;11}^2{+}\gamma_{1;12}^2)\beta_1^{2+}$	0	$(\gamma_{1;11}^2 + \gamma_{1;12}^2)\beta_1^{2-1}$
	0	q_{21}	$\gamma^2_{0;21}$	$q_{22} + \gamma_{0;22}^2$	0	$(\gamma_{1;11}^2{+}\gamma_{1;12}^2)\beta_2^{2+}$	0	$(\gamma_{1;21}^2 + \gamma_{1;22}^2)\beta_2^{2-}$
	$(-B^{1+}1)\bar{\gamma}_{1;1}^1$	$(-B^{1+}{\bf 1})\bar{\gamma}^1_{1;2}$	0	0	B^{1+}	0	0	0
	0	0	$(-B^{2+}1)\bar{\gamma}_{1;1}^2$	$(-B^{2+}1)\bar{\gamma}_{1;2}^2$	0	B^{2+}	0	0
	$(-B^{1-1})\bar{\gamma}_{1;1}^1$	$(-B^{1-1})\bar{\gamma}_{1:2}^{1}$	0	0	0	0	B^{1-}	0
l	0	0	$(-B^{2-1})\bar{\gamma}_{1;1}^2$	$(-B^{2-1})\bar{\gamma}_{1;2}^2$	0	0	0	B^{2-})
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Example 3. We consider the superposition of a two-state MMBM and inhomogeneous Poisson process, where,

- (i) $S^{(B)} = \{1, 2\}, Q^{(B)} = (q_{ij}), \mu^{(B)} = (\mu_1 \ \mu_2) \text{ and } \sigma^{(B)} = (\sigma_1 \ \sigma_2),$
- (ii) Poisson process with intensity $\gamma^{(J_B(t))}$,
- (iii) the size of a jump which occurs while J_B stays in $i \in S^{(B)}$ is assumed to be distributed as BPH distribution of the form

$$\boldsymbol{\beta}^{i+}e^{B^{i+}x}\left(-B^{i+}\mathbf{1}\right)\chi(x>0)+\boldsymbol{\beta}^{i-}e^{B^{i-}x}\left(-B^{i-}\mathbf{1}\right)\chi(x<0).$$

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Then, the infinitesimal generator of the embedded process of the superposition can be represented as

1	$q_{11} - \gamma^{(1)}$	q_{12}	$\gamma^{(1)}\beta^{1+}$	0	$\gamma^{(1)}\beta^{1-}$	0)
I	q_{21}	$q_{22} - \gamma^{(2)}$	0	$\gamma^{(2)}\beta^{2+}$	0		
	$(-B^{1+}1)$	0	B^{1+}	0	0	0	
	0	$(-B^{2+}1)$	0	B^{2+}	0	0	I.
	$(-B^{1-1})$	0	0	0	B^{1-}	0	
1	0	$(-B^{2-1})$	0	0	0	B^{2-})

The drift and diffusion vectors of the embedded process are given as $(\mu_1 \ \mu_2 \ \mathbf{1'} - \mathbf{1'})$ and $(\sigma_1 \ \sigma_2 \ \mathbf{0} \ \mathbf{0})$, respectively. \Box

Now, we consider the general case of the superposition. For the MAP jump occurring when $J = (i, \ell)$, we let $(\beta_{\ell}^{i+}, B_{\ell}^{i-}, B_{\ell}^{i-}, B_{\ell}^{i-})$ denote the parameters of the BPH distribution. If we denote by S_{ℓ}^{i+} and S_{ℓ}^{i-} the state spaces corresponding to B_{ℓ}^{i+} and B_{ℓ}^{i-} , respectively, then $S_{+} = \bigcup_{i \in S^{(B)}, \ell \in S^{(M)}} S_{\ell}^{i+}$ and $S_{-} = \bigcup_{i \in S^{(B)}, \ell \in S^{(M)}} S_{\ell}^{i-}$. The drift and diffusion vectors $\mu^{\mathbf{e}} = (\mu_b \, \mu_u \, \mu_d \, \mathbf{1}'_{j^+} \, \mathbf{1}'_{j^-})$ and $\sigma^{\mathbf{e}} = (\sigma_b \, \mathbf{0} \, \mathbf{0} \, \mathbf{0} \, \mathbf{0})$ satisfy $\mu_b = \mu_b^{(B)} \otimes \mathbf{1}'_m, \mu_u = \mu_u^{(B)} \otimes \mathbf{1}'_m, \mu_d = \mu_d^{(B)} \otimes \mathbf{1}'_m$, and $\sigma_b = \sigma_b^{(B)} \otimes \mathbf{1}'_m$. Moreover, the infinitesimal generator *T* of the embedded process given in the following partitioned form

$$T = \begin{pmatrix} T_{bb} & T_{bu} & T_{bd} & T_{b+} & T_{b-} \\ T_{ub} & T_{uu} & T_{ud} & T_{u+} & T_{u-} \\ T_{db} & T_{du} & T_{dd} & T_{d+} & T_{d-} \\ T_{+b} & T_{+u} & T_{+d} & T_{++} & \mathbf{0} \\ T_{-b} & T_{-u} & T_{-d} & \mathbf{0} & T_{--} \end{pmatrix},$$
(3.1)

in which, with $\chi(\cdot)$ denoting the indicator function and matrices $\mathbf{a}_{ij}^{k\ell}$ for $k, \ell \in \{b, u, d, +, -\}$ and $i \in S_k, j \in S_\ell$ being defined as

$$\mathbf{a}_{ij}^{kk} = q_{ij}I_m + \chi(i=j)\Gamma_0^i \text{ and } \mathbf{a}_{ij}^{k\ell} = q_{ij}I_m \text{ for } k \neq \ell,$$

the submatrices satisfy that

$$\begin{aligned} T_{bb} &= \left(\mathbf{a}_{ij}^{bb}, \ i, j \in S_{b}\right), \ T_{bu} &= \left(\mathbf{a}_{ij}^{bu}, \ i \in S_{b}, j \in S_{u}\right), \ T_{bd} &= \left(\mathbf{a}_{ij}^{bd}, \ i \in S_{b}, j \in S_{d}\right), \\ T_{uu} &= \left(\mathbf{a}_{ij}^{uu}, \ i, j \in S_{u}\right), \ T_{ub} &= \left(\mathbf{a}_{ij}^{ub}, \ i \in S_{u}, j \in S_{b}\right), \ T_{ud} &= \left(\mathbf{a}_{ij}^{uu}, \ i \in S_{u}, j \in S_{d}\right), \\ T_{dd} &= \left(\mathbf{a}_{ij}^{dd}, \ i, j \in S_{d}\right), \ T_{db} &= \left(\mathbf{a}_{ij}^{db}, \ i \in S_{d}, j \in S_{b}\right), \ T_{du} &= \left(\mathbf{a}_{ij}^{du}, \ i \in S_{d}, j \in S_{u}\right), \end{aligned}$$

with $\gamma_{1:k}^i = \sum_{l \in S_M} \gamma_{1:kl}^i$ for $i \in S$ and matrices \mathbf{b}_i^{k+} and \mathbf{b}_i^{k-} for $k \in \{b, u, d\}, i \in S_k$ being defined as

$$\mathbf{b}_i^{k+} = \operatorname{diag}\left\{\gamma_{1;j}^i \beta_j^{i+}, j \in S^{(M)}\right\} \text{ and } \mathbf{b}_i^{k-} = \operatorname{diag}\left\{\gamma_{1;j}^i \beta_j^{i-}, j \in S^{(M)}\right\},$$

$$T_{b+} = \left(\operatorname{diag} \left\{ \mathbf{b}_{i}^{b+}, i \in S_{b} \right\} \mathbf{0} \mathbf{0} \right), T_{u+} = \left(\mathbf{0} \operatorname{diag} \left\{ \mathbf{b}_{i}^{u+}, i \in S_{u} \right\} \mathbf{0} \right), T_{d+} = \left(\mathbf{0} \operatorname{diag} \left\{ \mathbf{b}_{i}^{d+}, i \in S_{d} \right\} \right), \quad (3.2)$$
$$T_{b-} = \left(\operatorname{diag} \left\{ \mathbf{b}_{i}^{b-}, i \in S_{b} \right\} \mathbf{0} \mathbf{0} \right), T_{u-} = \left(\mathbf{0} \operatorname{diag} \left\{ \mathbf{b}_{i}^{u-}, i \in S_{u} \right\} \mathbf{0} \right), T_{d-} = \left(\mathbf{0} \operatorname{diag} \left\{ \mathbf{b}_{i}^{d-}, i \in S_{d} \right\} \right),$$

with matrices $\mathbf{c}_i^{+\ell}$ and $\mathbf{c}_i^{-\ell}$ for $\ell \in \{b, u, d\}$, $i \in S_{\ell}$ being defined as

$$\mathbf{c}_{i}^{+\ell} = \operatorname{diag}\left\{ \left(-B_{j}^{i+1} \mathbf{1} \right) \left(\gamma_{1;j}^{i} \right)^{-1}, j \in S^{(M)} \right\} \Gamma_{1}^{i} \text{ and } \mathbf{c}_{i}^{-\ell} = \operatorname{diag}\left\{ \left(-B_{j}^{i-1} \mathbf{1} \right) \left(\gamma_{1;j}^{i} \right)^{-1}, j \in S^{(M)} \right\} \Gamma_{1}^{i},$$

$$T_{+b} = \begin{pmatrix} \operatorname{diag} \left\{ \mathbf{c}_{i}^{+b}, i \in S_{b} \right\} \\ \mathbf{0} \\ \mathbf{0} \end{pmatrix}, T_{+u} = \begin{pmatrix} \mathbf{0} \\ \operatorname{diag} \left\{ \mathbf{c}_{i}^{+u}, i \in S_{u} \right\} \\ \mathbf{0} \end{pmatrix}, T_{+d} = \begin{pmatrix} \mathbf{0} \\ \operatorname{diag} \left\{ \mathbf{c}_{i}^{+d}, i \in S_{d} \right\} \end{pmatrix}, \quad (3.3)$$
$$T_{-b} = \begin{pmatrix} \operatorname{diag} \left\{ \mathbf{c}_{i}^{-b}, i \in S_{b} \right\} \\ \mathbf{0} \\ \mathbf{0} \end{pmatrix}, T_{-u} = \begin{pmatrix} \mathbf{0} \\ \operatorname{diag} \left\{ \mathbf{c}_{i}^{-u}, i \in S_{u} \right\} \\ \mathbf{0} \end{pmatrix}, T_{-d} = \begin{pmatrix} \mathbf{0} \\ \mathbf{0} \\ \operatorname{diag} \left\{ \mathbf{c}_{i}^{-d}, i \in S_{d} \right\} \end{pmatrix},$$

and with $\mathbf{d}_i^{++} = \text{diag}\{B_k^{i+}, k \in S^{(M)}\}$ and $\mathbf{d}_i^{--} = \text{diag}\{B_k^{i-}, k \in S^{(M)}\}$ for $i \in S$,

$$T_{++} = \begin{pmatrix} \operatorname{diag}\{\mathbf{d}_{i}^{++}, i \in S_{b}\} & \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \operatorname{diag}\{\mathbf{d}_{i}^{++}, i \in S_{u}\} & \mathbf{0} \\ \mathbf{0} & \mathbf{0} & \operatorname{diag}\{\mathbf{d}_{i}^{++}, i \in S_{d}\} \end{pmatrix},$$
(3.4)
$$T_{--} = \begin{pmatrix} \operatorname{diag}\{\mathbf{d}_{i}^{--}, i \in S_{b}\} & \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \operatorname{diag}\{\mathbf{d}_{i}^{--}, i \in S_{u}\} & \mathbf{0} \\ \mathbf{0} & \mathbf{0} & \operatorname{diag}\{\mathbf{d}_{i}^{--}, i \in S_{d}\} \end{pmatrix}.$$

3.2. BPH jumps depending on both the Markov states before and after their corresponding jump epochs

In this section, we consider the superposition in which the MAP jumps depend on the states of J both before and after the jump epoch of the process. We also take a simple example for illustration of the structure of the resulting embedded process.

Example 4. Consider the superposition of a two-state MMBM and two-state MAP, where,

(i)
$$S^{(B)} = \{1, 2\}, Q^{(B)} = (q_{ij}), \mu^{(B)} = (\mu_1 \ \mu_2) \text{ and } \sigma^{(B)} = (\sigma_1 \ \sigma_2),$$

- (ii) two-state MAP($\Gamma_0^{J_B(t)}, \Gamma_1^{J_B(t)}$) with $S^{(M)} = \{1, 2\}, \Gamma_0^i = (\gamma_{0,kl}^i; k, l \in S^{(M)})$ and $\Gamma_0^i = (\gamma_{0,kl}^i; k, l \in S^{(M)})$ for $i \in S^{(B)}$,
- (iii) an MAP jump occurring at the transition epoch of J from (i, k) to (i, ℓ) is assumed to be distributed as ph-type distribution of the form

$$\boldsymbol{\beta}_{k\ell}^{i} e^{B_{k\ell}^{i} x} \left(-B_{k\ell}^{i} \mathbf{1}\right) \chi(x>0).$$

Then, the infinitesimal generator of the embedded process of the superposition can be represented as

($q_{11} + \gamma_{0;11}^1$	$\gamma_{0;12}^{1}$	q_{12}	0	$\gamma^1_{1;11}\beta^1_{11}$	$\gamma^1_{1;12}\beta^1_{12}$	0	0	0	0	0	0)
	$\gamma^1_{0;21}$	$q_{11} + \gamma^1_{0;22}$	0		0	0	$\gamma^1_{1;21}\beta^1_{21}$	$\gamma^1_{1;22}\beta^1_{22}$	0	0	0	0
I	q_{21}	0	$q_{22} + \gamma_{0;11}^2$	$\gamma^{2}_{0;12}$	0	0	0	0	$\gamma^2_{1;11}\beta^2_{11}$	$\gamma_{1;12}^2 \beta_{12}^2$	0	0
	0	q_{21}	$\gamma^{2}_{0;21}$	$q_{22} + \gamma_{0;22}^2$	0	0	0	0	0	0	$\gamma_{1;21}^2 \beta_{21}^2$	$\gamma_{1;22}^2 \beta_{22}^2$
I	$-B_{11}^1$ 1	0	0	0	B_{11}^{1}	0	0	0	0	0	0	0
	0	$-B_{12}^11$	0	0	0	B_{12}^{1}	0	0	0	0	0	0
I	$-B_{21}^1$ 1	0	0	00	0		B_{21}^{1}	0	0	0	0	0
I	0	$-B_{22}^11$	0	0	0	0		B_{22}^{1}	0	0	0	0
	0	0	$-B_{11}^2$ 1	0	0	0	0	0	B_{11}^2	0	0	0
	0	0	0	$-B_{12}^2$ 1	0	0	0	0	0	B_{12}^2	0	0
l	0	0	$-B_{21}^2$ 1	0	0	0	0	0	0	0	B_{21}^2	0
l	0	0	0	$-B_{22}^2$ 1	0	0	0	0	0	0	0	B_{22}^2

Application of GTH-like algorithm to Markov modulated Brownian motion with jumps

Now, we consider the general case of the superposition. For the MAP jump which occurs at the transition epoch of *J* from (i, j) to (i, k), we let $(\beta_{jk}^{i+}, B_{jk}^{i+}, \beta_{jk}^{i-}, B_{jk}^{i-})$ be the parameters of the BPH distribution. If we denote by S_{jk}^{i+} and S_{jk}^{i-} the state space corresponding to B_{jk}^{i+} and B_{jk}^{i-} , respectively, then $S_+ = \bigcup_{i \in S^{(B)}} \bigcup_{j \in S^{(M)}} \bigcup_{k \in S^{(M)}} S_{jk}^{i+}$, and $S_- = \bigcup_{i \in S^{(B)}} \bigcup_{j \in S^{(M)}} \bigcup_{k \in S^{(M)}} S_{jk}^{i-}$. The drift and diffusion vectors $\mu^{\mathbf{e}} = (\mu_b \ \mu_u \ \mu_d \ \mathbf{1}'_+ \ \mathbf{1}'_-)$ and $\sigma^{\mathbf{e}} = (\sigma_b \ \mathbf{0} \ \mathbf{0} \ \mathbf{0} \ \mathbf{0}$ satisfy $\mu_b = \mu_b^{(B)} \otimes \mathbf{1}'_m$, $\mu_u = \mu_u^{(B)} \otimes \mathbf{1}'_m$, $\mu_d = \mu_d^{(B)} \otimes \mathbf{1}'_m$, and $\sigma_b = \sigma_b^{(B)} \otimes \mathbf{1}'_m$. Moreover, the infinitesimal generator *T* of the embedded process given in the following partitioned form

$$\begin{pmatrix} T_{bb} & T_{bu} & T_{bd} & T_{b+} & T_{b-} \\ T_{ub} & T_{uu} & T_{ud} & T_{u+} & T_{u-} \\ T_{db} & T_{du} & T_{dd} & T_{d+} & T_{d-} \\ T_{+b} & T_{+u} & T_{+d} & T_{++} & \mathbf{0} \\ T_{-b} & T_{-u} & T_{-d} & \mathbf{0} & T_{--} \end{pmatrix},$$
(3.5)

in which, with matrices $\mathbf{a}_{ii}^{k\ell}$ for $k, \ell \in \{b, u, d, +, -\}$ and $i \in S_k, j \in S_\ell$ being defined as

$$\mathbf{a}_{ij}^{kk} = q_{ij}I_m + \chi(i=j)\Gamma_0^i \text{ and } \mathbf{a}_{ij}^{k\ell} = q_{ij}I_m \text{ for } k \neq \ell$$

the submatrices satisfy that,

$$T_{bb} = \left(\mathbf{a}_{ij}^{bb}, \ i, j \in S_{b}\right), \quad T_{bu} = \left(\mathbf{a}_{ij}^{bu}, \ i \in S_{b}, j \in S_{u}\right), \quad T_{bd} = \left(\mathbf{a}_{ij}^{bd}, \ i \in S_{b}, j \in S_{d}\right), \quad (3.6)$$

$$T_{uu} = \left(\mathbf{a}_{ij}^{uu}, \ i, j \in S_{u}\right), \quad T_{ub} = \left(\mathbf{a}_{ij}^{ub}, \ i \in S_{u}, j \in S_{b}\right), \quad T_{ud} = \left(\mathbf{a}_{ij}^{ud}, \ i \in S_{u}, j \in S_{d}\right), \quad (3.6)$$

$$T_{dd} = \left(\mathbf{a}_{ij}^{dd}, \ i, j \in S_{d}\right), \quad T_{db} = \left(\mathbf{a}_{ij}^{db}, \ i \in S_{d}, j \in S_{b}\right), \quad T_{du} = \left(\mathbf{a}_{ij}^{du}, \ i \in S_{d}, j \in S_{u}\right),$$

with matrices $\mathbf{b}_i^{\ell+}$ and $\mathbf{b}_i^{\ell-}$ for $\ell \in \{b, u, d\}, i \in S_\ell$ being defined as

$$\mathbf{b}_{i}^{\ell+} = \mathbf{b}_{i}^{\ell+} = \operatorname{diag}\left\{\left(\gamma_{1;k1}^{i}\beta_{k1}^{i+}, \dots, \gamma_{1;km}^{i}\beta_{km}^{i+}\right), k \in S_{M}\right\} \text{ and } \mathbf{b}_{i}^{\ell-} = \operatorname{diag}\left\{\left(\gamma_{1;k1}^{i}\beta_{k1}^{i-}, \dots, \gamma_{1;km}^{i}\beta_{km}^{i-}\right), k \in S_{M}\right\},\$$
$$T_{b+} = \left(\operatorname{diag}\left\{\mathbf{b}_{i}^{b+}, i \in S_{b}\right\} \mathbf{0} \mathbf{0}\right), T_{u+} = \left(\mathbf{0} \operatorname{diag}\left\{\mathbf{b}_{i}^{u+}, i \in S_{u}\right\} \mathbf{0}\right), T_{d+} = \left(\mathbf{0} \operatorname{0} \operatorname{diag}\left\{\mathbf{b}_{i}^{d+}, i \in S_{d}\right\}\right), \quad (3.7)$$

 $T_{b-} = \left(\operatorname{diag} \left\{ \mathbf{b}_{i}^{b-}, i \in S_{b} \right\} \mathbf{0} \mathbf{0} \right), T_{u-} = \left(\mathbf{0} \operatorname{diag} \left\{ \mathbf{b}_{i}^{u-}, i \in S_{u} \right\} \mathbf{0} \right), T_{d-} = \left(\mathbf{0} \operatorname{0} \operatorname{diag} \left\{ \mathbf{b}_{i}^{d-}, i \in S_{d} \right\} \right),$

with matrices $\mathbf{c}_i^{+\ell}$ and $\mathbf{c}_i^{+\ell}$ for $\ell \in \{b, u, d\}$, $i \in S_\ell$ being defined as

$$\mathbf{c}_{i}^{+\ell} = \begin{pmatrix} \operatorname{diag} \left\{ -B_{1,k}^{i+} \mathbf{1}, k \in S^{(M)} \right\} \\ \vdots \\ \operatorname{diag} \left\{ -B_{m,k}^{i+} \mathbf{1}, k \in S^{(M)} \right\} \end{pmatrix} \text{ and } \mathbf{c}_{i}^{-\ell} = \begin{pmatrix} \operatorname{diag} \left\{ -B_{1,k}^{i-} \mathbf{1}, k \in S^{(M)} \right\} \\ \vdots \\ \operatorname{diag} \left\{ -B_{m,k}^{i-} \mathbf{1}, k \in S^{(M)} \right\} \end{pmatrix},$$

$$T_{+b} = \begin{pmatrix} \operatorname{diag} \left\{ \mathbf{c}_{i}^{+b}, i \in S_{b} \right\} \\ \mathbf{0} \end{pmatrix}, T_{+u} = \begin{pmatrix} \mathbf{0} \\ \operatorname{diag} \left\{ \mathbf{c}_{i}^{+u}, i \in S_{u} \right\} \\ \mathbf{0} \end{pmatrix}, T_{+d} = \begin{pmatrix} \mathbf{0} \\ \operatorname{diag} \left\{ \mathbf{c}_{i,i}^{-d}, i \in S_{d} \right\} \end{pmatrix}, \quad (3.8)$$

$$T_{-b} = \begin{pmatrix} \operatorname{diag} \left\{ \mathbf{c}_{i}^{-b}, i \in S_{b} \right\} \\ \mathbf{0} \end{pmatrix}, T_{-u} = \begin{pmatrix} \mathbf{0} \\ \operatorname{diag} \left\{ \mathbf{c}_{i}^{-u}, i \in S_{u} \right\} \\ \mathbf{0} \end{pmatrix}, T_{-d} = \begin{pmatrix} \mathbf{0} \\ \operatorname{diag} \left\{ \mathbf{c}_{i}^{-d}, i \in S_{d} \right\} \end{pmatrix},$$

and with $\mathbf{d}_{i}^{++} = \text{diag}\{\text{diag}\{B_{k,l}^{i+}, l \in S^{(M)}\}, k \in S^{(M)}\}$ and $\mathbf{d}_{i}^{--} = \text{diag}\{\text{diag}\{B_{k,l}^{i-}, l \in S^{(M)}\}, k \in S^{(M)}\}$ for $i \in S_{b} \cup S_{u} \cup S_{d}$,

$$T_{++} = \begin{pmatrix} \operatorname{diag}\{\mathbf{d}_{i}^{++}, i \in S_{b}\} & \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \operatorname{diag}\{\mathbf{d}_{i}^{++}, i \in S_{u}\} & \mathbf{0} \\ \mathbf{0} & \mathbf{0} & \operatorname{diag}\{\mathbf{d}_{i}^{++}, i \in S_{d}\} \end{pmatrix}, \quad (3.9)$$
$$T_{--} = \begin{pmatrix} \operatorname{diag}\{\mathbf{d}_{i}^{--}, i \in S_{b}\} & \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \operatorname{diag}\{\mathbf{d}_{i}^{--}, i \in S_{u}\} & \mathbf{0} \\ \mathbf{0} & \mathbf{0} & \operatorname{diag}\{\mathbf{d}_{i}^{--}, i \in S_{d}\} \end{pmatrix}.$$

4. The structure-preserving doubling algorithm and GTH-like algorithm

We first introduce some relevant definitions and notations. For any matrices $A, B \in \mathbb{R}^{m \times n}$, we write $A \ge B(A > B)$ if $[A]_{ij} \ge [A]_{ij}([A]_{ij} > [B]_{ij}$ for all i, j, where $[A]_{ij}$ denotes the $(i, j)^{th}$ element of A. A real square matrix A is called a Z-matrix if all its off-diagonal elements are non-positive. Any Z-matrix A can be written as sI - B with $B \ge 0$. A Z-matrix A is called an M-matrix if $s \ge \rho(B)$, where $\rho(\cdot)$ is the spectral radius; it is called a singular M-matrix if $s = \rho(B)$ and a non-singular M-matrix if $s > \rho(B)$.

4.1. Structure-preserving doubling algorithm

We recall the parameter triple (T, μ^{e}, σ^{e}) of the G-MMBM given in the previous section and the stationary probability vector π^{e} of the infinitesimal generator. Then the first passage probability of the G-MMBM can be obtained using the minimal solution of the NARE AZ + ZB + ZCZ + D = 0 given in (2.4) and (2.5) as mentioned in Section 2.2. In connection with the NARE (2.4), we define an *M*-matrix *M* as

$$M = \begin{pmatrix} -\mathbf{B} & -\mathbf{C} \\ -\mathbf{D} & -\mathbf{A} \end{pmatrix},\tag{4.1}$$

which is irreducible by the irreducibility of J_B and J_M . Moreover, M is a singular matrix.

To get the nonnegative minimal solution of the NARE, we can use the structure-preserving doubling algorithm (SDA) presented in Guo *et al.* (2006), which is as follow,

Structure-preserving doubling (SDA) algorithm for an NARE AZ + ZB + ZCZ + D = 0

Step 1. Choose
$$\gamma \ge \max\{-[\mathbf{A}]_{ii}, -[\mathbf{B}]_{ii}, i \in S\}$$
 and set

$$\begin{pmatrix} E_0 & G_0 \\ H_0 & F_0 \end{pmatrix} = \begin{pmatrix} \gamma I - \mathbf{B} & -\mathbf{C} \\ -\mathbf{D} & \gamma I - \mathbf{A} \end{pmatrix}^{-1} \begin{pmatrix} \gamma I + \mathbf{B} & \mathbf{C} \\ \mathbf{D} & \gamma I + \mathbf{A} \end{pmatrix}$$

Step 2. $E_{k+1} = E_k (I - G_k H_k)^{-1} E_k; F_{k+1} = F_k (I - H_k G_k)^{-1} F_k,$ $G_{k+1} = G_k + E_k (I - G_k H_k)^{-1} G_k F_k; H_{k+1} = H_k + F_k (I - H_k G_k)^{-1} H_k E_k,$

Step 3. $Z = H_{\infty}$,

The convergence results of the SDA when *M* is an irreducible *M*-matrix is given in the following theorem, which can be found in Theorem 3.1 and 3.2 of Wang *et al.* (2012) and Theorem 3.1 of Xue and Li (2017). For the theorem, we define the Cayley transform of R = -B - CX and S = -A - DY such that

$$R_{\gamma} = (R + \gamma I)^{-1} (R - \gamma I) \text{ and } S_{\gamma} = (S + \gamma I)^{-1} (S - \gamma I).$$
 (4.2)

Theorem 2. Note that $\{E_k\}$, $\{G_k\}$, $\{H_k\}$, and $\{F_k\}$ denote the matrices in the k^{th} iteration of the SDA. (a) For all $k \ge 0$, the matrices E_k , G_k , H_k , and F_k are non-negative. Moreover, for all $k \ge 0$, $I - G_k H_k$ and $I - H_k G_k$ are non-singular M-matrices.

(b) In the positive recurrent case, then $\rho(R_{\gamma}) = 1$ and $\rho(S_{\gamma}) < 1$. Furthermore, $\{H_k\}$ converges to X quadratically with,

$$\limsup_{k\to\infty} \sqrt[2^k]{\|H_k - X\|} \le \rho(S_{\gamma}),$$

 $\{F_k\}$ converges to **0** quadratically with $\limsup_{k\to\infty} \sqrt[2^k]{\|F_k\|} \le \rho(S_{\gamma})$, and $\{E_k\}$ is bounded. The notation ||A|| denotes the maximum of the absolute values of the elements in a Matrix A.

(c) In the transient case, then $\rho(R_{\gamma}) < 1$ and $\rho(S_{\gamma}) = 1$. Furthermore, $\{H_k\}$ converges to X quadratically with

$$\limsup_{k\to\infty} \sqrt[2^k]{\|H_k-X\|} \le \rho(R_{\gamma}),$$

 $\{E_k\}$ converges to **0** quadratically with $\limsup_{k\to\infty} \sqrt[2^k]{||E_k||} \le \rho(R_\gamma)$, and $\{F_k\}$ is bounded. (d) In the null recurrent case, then $\rho(R_\gamma) = 1$ and $\rho(S_\gamma) = 1$. In this case, $\{H_k\}$ converges to X and $\{E_k\}$, $\{F_k\}$ are bounded.

As we can see in SDA algorithm, the initialization phase and the iteration kernel of the SDA involve inverting non-singular M-matrices. In particular, in the null recurrent case, the M-matrices, although non-singular, move towards singular M-matrices at convergence. These inversions are causes of concerns on entry-wise relative accuracy of the computed minimal nonnegative solution. Fortunately, a non-singular M-matrices can be inverted by the GTH-like algorithm of Alfa *et al.* (2001) to almost full entry-wise relative accuracy, which enhances the entry-wise relative accuracy of the computed minimal nonnegative solution.

4.2. GTH-like algorithm

The GTH-like algorithm, proposed by Alfa *et al.* (2001), is used for solving linear systems $A\mathbf{x} = \mathbf{b}$ when *A* is an *M*-matrix satisfying $A\mathbf{v} = \mathbf{c}$ with $\mathbf{v} > 0$ and $\mathbf{c} \ge 0$. It is a subtraction-free algorithm and compute the inverse of *A* with relative errors in the magnitude of the machine precision. For the details on the algorithm, refer to (2001). When it holds that $A\mathbf{v} = \mathbf{c}$ with $\mathbf{v} > 0$ and $\mathbf{c} \ge 0$, we say that the triplet representation of *A* is available and represent as $A = (N_A, \mathbf{v}, \mathbf{c})$ with $N_A = \text{diag}(A) - A$.

Proposition 1. Recall that $\Lambda = diag\{-[T_{b,b}]_{i,i}, i \in S_b\}$. Then, the left and right eigenvectors **u** and **v** corresponding to the eigenvalue 0 of *M* are given as

$$\mathbf{u} := \begin{pmatrix} \mathbf{u}_1 \\ \mathbf{u}_2 \end{pmatrix} = \begin{pmatrix} \begin{bmatrix} (D_1 + D_2)^{-1} \mathbf{\Lambda} \pi'_b \\ \Delta_{-\mu_d} \pi'_d \\ \pi'_- \end{bmatrix} \\ \begin{bmatrix} 0.5 \mathbf{\Sigma} \pi'_b \\ \Delta_{\mu_u} \pi'_u \\ \pi'_+ \end{bmatrix} \end{pmatrix} \text{ and } \mathbf{v} := \begin{pmatrix} \mathbf{v}_1 \\ \mathbf{v}_2 \end{pmatrix} = \begin{pmatrix} \begin{bmatrix} \mathbf{1}_b \\ \mathbf{1}_d \\ \mathbf{1}_- \end{bmatrix} \\ \begin{bmatrix} (D_1 + D_2) \mathbf{\Sigma} \mathbf{1}_b \\ \mathbf{1}_u \\ \mathbf{1}_+ \end{bmatrix} \end{pmatrix}, \quad (4.3)$$

which are unique up to a scalar multiple.

Proof: We can prove by showing that $\mathbf{u}'M = \mathbf{0}$ and $M\mathbf{v} = \mathbf{0}$. Recall that

$$\mathbf{A} = \begin{pmatrix} D_1 - D_2 & 2\Sigma^{-1}T_{b,u} & 2\Sigma^{-1}T_{b,+} \\ \mathbf{0}_{u,b} & \Delta_{\mu u}^{-1}T_{u,u} & \Delta_{\mu u}^{-1}T_{u,+} \\ \mathbf{0}_{+,b} & T_{+,u} & T_{+,+} \end{pmatrix}, \quad \mathbf{C} = \begin{pmatrix} \Sigma^{-1} & \mathbf{0}_{b,u} & \mathbf{0}_{b,+} \\ \mathbf{0}_{d,b} & \Delta_{-\mu_d}^{-1}T_{d,u} & \Delta_{-\mu_d}^{-1}T_{d,-} \\ \mathbf{0}_{-,b} & T_{-,u} & \mathbf{0}_{-,+} \end{pmatrix},$$
$$\mathbf{B} = \begin{pmatrix} -D_1 - D_2 & \mathbf{0}_{b,d} & \mathbf{0}_{b,-} \\ \Delta_{-\mu_d}^{-1}T_{d,b} & \Delta_{-\mu_d}^{-1}T_{d,d} & \Delta_{-\mu_d}^{-1}T_{d,-} \\ T_{-,b} & T_{-,d} & T_{-,-} \end{pmatrix}, \quad \mathbf{D} = \begin{pmatrix} 2\Sigma^{-1}(T_{b,b} + \mathbf{A}) & 2\Sigma^{-1}T_{b,d} & 2\Sigma^{-1}T_{b,-} \\ \Delta_{\mu_d}^{-1}T_{u,b} & \Delta_{\mu_d}^{-1}T_{u,d} & \Delta_{\mu_d}^{-1}T_{u,-} \\ T_{+,b} & T_{+,d} & \mathbf{0}_{+,-} \end{pmatrix}.$$

Note that $\mathbf{u}'M = (-\mathbf{u}'_1\mathbf{B} - \mathbf{u}'_2\mathbf{D} - \mathbf{u}'_1\mathbf{C} - \mathbf{u}'_2\mathbf{A})$. If we temporarily let $-\mathbf{u}'_1\mathbf{B} - \mathbf{u}'_2\mathbf{D} = (\mathbf{z}_b \ \mathbf{z}_d \ \mathbf{z}_-)$, then

$$\mathbf{z}_{b} = \pi_{b} \mathbf{\Lambda} (D_{1} + D_{2})^{-1} (D_{1} + D_{2}) - \pi_{d} T_{d,b} - \pi_{-} T_{-,b} - \pi_{b} (0.5\Sigma) \left(2\Sigma^{-1} \right) (T_{b,b} + \mathbf{\Lambda}) - \pi_{u} T_{u,b} - \pi_{+} T_{+,b} \\ = -\pi_{b} T_{b,b} - \pi_{u} T_{u,b} - \pi_{+} T_{+,b} - \pi_{d} T_{d,b} - \pi_{-} T_{-,b} = \mathbf{0}.$$

Similarly, we can show that

$$\mathbf{z}_{d} = -\pi_{b}T_{b,d} - \pi_{u}T_{u,d} - \pi_{+}T_{+,d} - \pi_{d}T_{d,d} - \pi_{-}T_{-,d} = \mathbf{0},$$

$$\mathbf{z}_{-} = -\pi_{b}T_{b,-} - \pi_{u}T_{u,-} - \pi_{+}T_{+,-} - \pi_{d}T_{d,-} - \pi_{-}T_{-,-} = \mathbf{0}.$$

If we also temporarily let $-\mathbf{u}_1'\mathbf{C} - \mathbf{u}_2'\mathbf{A} = (\mathbf{z}_b \ \mathbf{z}_u \ \mathbf{z}_+)$, then

$$\begin{aligned} \mathbf{z}_{b} &= -\pi_{b} \mathbf{\Lambda} (D_{1} + D_{2})^{-1} \mathbf{\Sigma}^{-1} - \pi_{b} (0.5 \mathbf{\Sigma}) (D_{1} - D_{2}) \\ &= -\pi_{b} (D_{1} + D_{2})^{-1} \left[\mathbf{\Lambda} \mathbf{\Sigma}^{-1} + 0.5 \mathbf{\Sigma} \left(D_{1}^{2} - D_{2}^{2} \right) \right] \\ &= -\pi_{b} (D_{1} + D_{2})^{-1} \left[\mathbf{\Lambda} \mathbf{\Sigma}^{-1} - \mathbf{\Sigma}^{-1} \mathbf{\Lambda} \right] = \mathbf{0}. \end{aligned}$$

Note that the matrices D_1 , D_2 , Σ , and Λ are all diagonal. We can also show that

$$\mathbf{z}_{u} = -\boldsymbol{\pi}_{b}T_{b,u} - \boldsymbol{\pi}_{u}T_{u,u} - \boldsymbol{\pi}_{+}T_{+,u} - \boldsymbol{\pi}_{d}T_{d,u} - \boldsymbol{\pi}_{-}T_{-,u} = \mathbf{0},$$

$$\mathbf{z}_{+} = -\boldsymbol{\pi}_{b}T_{b,+} - \boldsymbol{\pi}_{u}T_{u,+} - \boldsymbol{\pi}_{+}T_{+,+} - \boldsymbol{\pi}_{d}T_{d,+} - \boldsymbol{\pi}_{-}T_{-,+} = \mathbf{0}.$$

We left the proof of $M\mathbf{v} = \mathbf{0}$ to the reader, which is arithmetic.

By examining the iteration kernel of the SDA algorithm, we see the inverse matrices $I - H_kG_k$ and $I - G_kH_k$, which we can apply the GTH-like algorithm provided that their triplet representation are available. In general, the triplet representation are not available. However, the spectral result in Proposition 1 make it possible, which is shown in the following proposition.

Proposition 2. Recall the matrices E_k , G_k , H_k , F_k in the k^{th} iteration of SDA algorithm. Then, it satisfies that, for all $k \ge 0$,

$$(I - H_k G_k)\mathbf{v}_2 = F_k \mathbf{v}_2 + H_k E_k \mathbf{v}_1$$
 and $(I - G_k H_k)\mathbf{v}_1 = E_k \mathbf{v}_1 + G_k F_k \mathbf{v}_2$,

where $\mathbf{v} = (\mathbf{v}'_1 \ \mathbf{v}'_2)'$ is the right eigenvector corresponding to 0 of the matrix M of which the form is given in (4.3).

Proof: Since $M\mathbf{v} = \begin{pmatrix} -B & -C \\ -D & -A \end{pmatrix}\mathbf{v} = \mathbf{0}$ from Proposition 1 and *M* is an M-matrix, the matrix $\begin{pmatrix} \gamma I - B & -C \\ -D & \gamma I - A \end{pmatrix}$ is non-singular M-matrix. Moreover, since

$$\begin{pmatrix} \gamma I + B & C \\ D & \gamma I + A \end{pmatrix} \mathbf{v} = \gamma \mathbf{v} + \begin{pmatrix} B & C \\ D & A \end{pmatrix} \mathbf{v} = \gamma \mathbf{v} - \begin{pmatrix} B & C \\ D & A \end{pmatrix} \mathbf{v} = \begin{pmatrix} \gamma I - B & -C \\ -D & \gamma I - A \end{pmatrix} \mathbf{v},$$

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it holds that

$$\begin{pmatrix} E_0 & G_0 \\ H_0 & F_0 \end{pmatrix} \mathbf{v} = \begin{pmatrix} \gamma I - B & -C \\ -D & \gamma I - A \end{pmatrix}^{-1} \begin{pmatrix} \gamma I + B & C \\ D & \gamma I + A \end{pmatrix} \mathbf{v} = \mathbf{v}.$$

We can also show that

$$\left(\begin{array}{cc} E_k & G_k \\ H_k & F_k \end{array}\right) \mathbf{v} = \mathbf{v}$$

for all k > 0 by induction using the same way in Theorem 3.2 of Xue and Li (2017), which yields that for all $k \ge 0$

$$\left(\begin{array}{cc}I & -G_k\\ -H_k & I\end{array}\right)\mathbf{v} = \left(\begin{array}{cc}E_k & \mathbf{0}\\ \mathbf{0} & F_k\end{array}\right)\mathbf{v}.$$

Multiplying $(I G_k)$ to the right of both sides, we can obtain the result of $(I - G_k H_k)$. Similarly, we can get the result of $(I - H_k G_k)$ by using the equation

$$\begin{pmatrix} F_k & H_k \\ G_k & E_k \end{pmatrix} \begin{pmatrix} \mathbf{v}_2 \\ \mathbf{v}_1 \end{pmatrix} = \begin{pmatrix} \mathbf{v}_2 \\ \mathbf{v}_1 \end{pmatrix}.$$

Since $I - H_k G_k$ and $I - G_k H_k$ are non-singular *M*-matrices for each $k \ge 0$, Proposition 2 makes it possible to compute their inverses using the GTH-like algorithm.

5. Numerical study

For the numerical study, we used MATLAB (R2019a) running on a Windows 10 64 bit in DELL Server with dual processor Intel (R) Xeon (R) Gold 6126 CPU @ 2.60GHz and 256GB of main memory.

To see the effect of GTH-like algorithm, we compare the original SDA algorithm(A_1) with the subtraction-free SDA(A_2) which uses the GTH-like algorithm. For the study, we take one of the examples in Example 3.1, of which $S^{(B)} = \{1\}$, $S^{(M)} = \{1, ..., n\}$, $\Gamma_0^1 = -(15/8)I_n$, $\Gamma_1^1 = -(15/8n)J_n$, $\beta_k^{1-} = \beta_k^{1-} = (1/2n)\mathbf{1}'_n$ for all $k \in S^{(M)}$, $B_k^{1-} = B_k^{1+} = -2I_n$ for all $k \in S^{(M)}$, $\mu_b = (3/2)\mathbf{1}'_n$, and $\sigma_b = \sqrt{7/4}\mathbf{1}'_n$. Then, the infinitesimal generator of the embedded process of the superposition is represented as

$$T = \begin{pmatrix} T_{b,b} & T_{b,+} & T_{b,-} \\ T_{+,b} & T_{+,+} & T_{+,-} \\ T_{-,b} & T_{-,+} & T_{-,-} \end{pmatrix} = \begin{pmatrix} -\frac{15}{8}I_n & \frac{15}{16n}\mathbf{J}_n & \frac{15}{16n}\mathbf{J}_n \\ \frac{2}{n}\mathbf{J}_n & -2I_n & \mathbf{0}_{n\times n} \\ \frac{2}{n}\mathbf{J}_n & \mathbf{0}_{n\times n} & -2I_n \end{pmatrix},$$

where \mathbf{J}_n denotes the *n*-dimensional square matrix of 1's.

We note that the average drift value of (F, J), average value of aggregated amount of fluid per unit time, is (3/2) for all *n*. To investigate the effects of the average drift value, we consider another parameter $m = (2/3) \times \overline{\mu}_m$, which makes the average drift value of $(d_m \times F, J)$ be $\overline{\mu}_m$. For the MMBM $(d_m \times F, J)$ with τ_m and H_m being its first passage time to 0 and the *H*-matrix, it holds that for all *m* and *x*

$$P[\tau < \infty | J(0) = 1, F(0) = x] = P[\tau_m < \infty | J(0) = 1, d_m F(0) = d_m x]$$
$$= \mathbf{e}_1 e^{(d_m x)H_m} \mathbf{1} = \frac{3}{4} e^{-x} + \frac{1}{4} e^{-3x},$$
(5.1)

$(\bar{\mu}, n)$	(0, 10)	(0,100)	(0, 500)	$(10^{-5}, 10)$	$(10^{-5}, 100)$	$(10^{-5}, 500)$
A1	NaN	4.98E-08	NaN	2.84E-16	1.70E-15	5.07E-16
A2	1.55E-15	1.71E–14	1.78E-14	1.94E-16	1.43E-15	9.71E-17
$(\bar{\mu}, n)$	$(10^{-4}, 10)$	$(10^{-4}, 100)$	$(10^{-4}, 500)$	(5, 10)	(5,100)	(5, 500)
A1	2.84E-16	1.69E-15	5.27E-16	2.08E-16	1.71E-15	5.69E-16
A2	2.08E-16	1.51E-15	4.84E-17	1.32E-16	1.54E-15	6.94E-17

Table 1: Comparison of error values

where \mathbf{e}_1 is a unit vector with appropriate dimension, in which its first element is 1 and the others are all 0. For the exact function of *x*, we refer to Example 6.1 of Asmussen (1995).

We choose the values of d_m so that the average drift value $\bar{\mu}_m$ of the MMBM's $(d_m \times F, J)$ is to be 0.00001, 0.0001, 5, and then compute $P[\tau_m < \infty | J(0) = 1, d_m F(0) = 3d_m]$ to compare with the exact value in (5.1). Note that the diffusion and drift vector of $(d_m \times F, J)$ are given as $(d_m \sigma_b \ \mathbf{0} \ \mathbf{0})$ and $(d_m \mu_b \ \mathbf{1'} - \mathbf{1'})$. To check our computations even more, we consider the MMBM (F, J) for which $S = \{1, \ldots, 3n\}$ with n = 10, 100, 500. For the case of $\bar{\mu} = 0$, we used the MMBM (F_0, J) , of which all the parameters are same as (F, J) except μ_b being given as $\mathbf{0}_{n\times 1}$. We compute $P[\tau_0 < \infty | J(0) = 1, F_0(0) = 3] = \mathbf{e}_1 e^{3H_0} \mathbf{1}$, which is equal to 1. Here, τ_0 and H_0 denotes the first passage time to 0 and the *H*-matrix of (F_0, J) . The numerical results are presented in Tables 1. From the table, we can observe that the GTH-like algorithm makes the doubling algorithm more stable and accurate, especially when the G-MMBM is critical, that is, the average value of drift is 0.

6. Concluding remarks

In this paper, we introduce how to parameterize the superposition of the MMBM and MAP with bilateral ph-type distributed jumps and also the non-symmetric algebraic Riccati equation to be used to compute the first passage probability of the superposed process. We note that the coefficient matrices of the equation can be obtained from the parametrization. Furthermore, we provide the spectral results which makes it possible to apply the GTH-like algorithm to a structure-preserving doubling algorithm for solving the equation. Through a numerical study, we have shown that using the GTH-like algorithm enhances the performance of the structure-preserving algorithm and makes it remarkably more stable and accurate especially when the superposed process is in the critical case.

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References

Ahn S and Ramaswami V (2005). Bilateral phase type distributions. Stochastic Models, 21 239–259.

- Ahn S (2016). Total shift during the first passages of Markov modulated Brownian motion with bilateral ph-type jumps: Formulas driven by the minimal solution matrix of a Riccati equation, *Stochastic Models*, **32**, 433–459.
- Ahn S (2017). Time-dependent and stationary analyses of the two-sided reflected Markov modulated Brownian motion with bilateral ph-type jumps, *Journal of the Korean Statistical Society*, **46**, 45–69.

Alfa AS, Xue J, and Ye Q (2001) Accurate computation of the smallest eigenvalue of a diagonally

dominant M-matrix, Math. Comp., 71, 217-236.

Asmussen S (1995). Stationary distributions for fluid flow models with or without Brownian noise. *Stochastic Models*, **11**, 1–20.

Asmussen S (2003). Applied Probability and Queues(2nd ed.), Springer, Verlag.

- Belhaj M (2010). Optimal dividend payments when risk reserves follow a jump diffusion process. Mathematical Finance, 20, 313–325.
- Bini DA, Iannazzo B, and Meini B (2012). *Numerical Solution of Algebraic Riccati Equations*, Siam, Philadelphia.
- Guo XX, Lin WW, and Xu SF (2006). A structure-preserving doubling algorithm for nonsymmetric algebraic Riccati equation, *Numer. Math.*, **103**, 393–412.
- Jiang Z and Pistorius MR (2008). On perpetual American put valuation and first passage in a regimeswitching model with jumps. *Finance Stoch.* **12**, 331–355.
- Liu C and Xue J (2012). Complex nonsymmetric algebraic Riccati equations arising in Markov modulated fluid flows, *SIAM J. Matrix Anal. Appl.*, **33**, 569–596.
- Wang W, Wang W, and Li R (2012). Alternating-directional doubling algorithm for *M*-matrix algebraic Riccati equations, SIAM. J. Matrix Anal. & Appl., 33, 170–194.
- Whitt W (1980). Some useful functions for functional limit theorems, *Mathematics of Operations Research*, **5**, 67–85.
- Xue J and Li R (2017) Highly accurate doubling algorithm for *M*-matrix algebraic Riccati equations, *Numer. Math.* **135**, 733—767.

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