Variance bounds for a class of biochemical reactions with bursts using a discrete expansion

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Abstract—We consider the problem of quantifying the variance in the number of molecules of a species, in biochemical reactions with nonlinear reaction rates. We address this problem for a particular configuration where a species is formed with bursts, with a nonlinear rate that depends on another spontaneously formed species. By making use of an appropriately formulated expansion based on the Newton series, in conjunction with spectral properties of the master equation, we derive an analytical expression that provides a hard bound for the variance. We also show that this bound is exact when the propensities are linear. Furthermore, numerical simulations demonstrate that this is very close to the actual variance.

I. INTRODUCTION

Chemical Master Equations (CMEs) are frequently used to model biochemical systems. Nevertheless, when the reaction rates are nonlinear analytical expressions for the moments are in general not feasible. Approximate analysis can be carried out via simulations, such as trajectories generated via the Gillespie Algorithm, [1]. Using, however those for evaluating moments requires considerable computational effort. Alternative approaches that lead to an approximate analysis include the Finite State Projection algorithm [2] and the Linear Noise Approximations (LNAs), or Van Kampen’s Ω-Expansion where the last two involve a linearisation of the reaction rates.

Recent studies have developed techniques for computing bounds for moments. In [3] bounds are computed by means of semidefinite programming. The idea of moment semidefinite programs was used in [4] and in [5], [6] optimization problems are formulated with positive semidefinite constraints associated with moment matrices for chemical reactions. Approximate moment dynamics with closure properties are derived in [7], [8] and [9] by selecting an appropriate nonlinear system to approximate the infinite set of moment Ordinary Differential Equations (ODEs) emerging from the CME.

In this paper we present an alternative approach whereby an appropriately formulated expansion based on the Newton series, is used to derive a hard bound for the variance for a class of biochemical reactions. In particular, the variance bound is derived for a species that is formed with bursts with a nonlinear rate that is a function of another spontaneously formed species. We also show that the bound derived is exact when the reaction rates are linear. Finally, we use numerical simulations to investigate how close the bound is to the actual variance, and we find that, unlike LNAs, this is very close to the true variance even in regimes where the reaction rate is highly nonlinear. A side result associated with the covariance of the species under consideration is also presented.

It should be noted that the methodology used to derive the bound presented is of independent interest as it exploits appropriate spectral properties of the master equation. Generalizations of this approach to larger classes of biochemical reactions is part of ongoing work.

The paper is structured as follows. In Section II we introduce the notation and provide the problem formulation. The main results are given in Section III. A numerical evaluation of the results is provided in Section IV. Finally, conclusions are drawn in Section V. The proofs of the results presented in the text are provided in the appendix.

II. PROBLEM FORMULATION

A. Notation

The following list introduces various symbols used within the paper

\((A)_k\) Falling factorial of \(A \in \mathbb{Z}^\geq\) with \(k \in \mathbb{Z}^\geq\): if \(k \neq 0\) then \((A)_k = \prod_{n=0}^{k-1} A - n\), if \(k = 0\) then \((A)_0 = 1\)

\(\Delta_k[\cdot]\) \(k\) step finite difference of the form \(\Delta_k[f](A) \equiv f(A + k) - f(A)\)

\(\Delta^P_k[\cdot]\) \(k\) step finite difference operator applied \(p\) times

\(\delta_{a,b}\) \(\delta_{a,b} = 1\), if \(a = b\), 0 otherwise

\(E[X]\) Expectation of random variable \(X\)

\(E_P[\cdot]\) Expectation operator with respect to the Poisson distribution

\(\mathbb{R}[A]\) Set of polynomials in \(A\) with real valued coefficients

\(\mathbb{R}^>\) Set of positive real numbers \(\{x \in \mathbb{R} : x > 0\}\)

\(\mathbb{R}^\geq\) Set of non negative real numbers \(\{x \in \mathbb{R} : x \geq 0\}\)

\(\mathbb{Z}^>\) Set of non negative integers \(\{0, 1, 2, 3, \ldots\}\)

\(U\) Discrete uniform probability distribution with support \(\{a, b\}\) with \(a, b \in \mathbb{Z}^>\)

\(\mathbb{I}\) Unity operator: \(\mathbb{I}[X] = X\)

\(A \sim \mathcal{P}(x)\) Random variable \(A\) has Poisson probability distribution with mean \(x\)

\(\text{cov}(X, Y)\) Covariance of random variables \(X\) and \(Y\)

\(\text{var}(X)\) Variance of random variable \(X\)

\(\mathcal{B}(p, n)\) Binomial probability distribution, with parameters \(0 < p \leq 1\) and \(n \in \mathbb{Z}^\geq\)

\(\mathcal{G}(p)\) Geometric probability distribution, with parameter \(0 < p \leq 1\)

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B. Problem Formulation

We consider the following system

\[
\begin{align*}
A & \xrightarrow{F} A + 1 \quad B \xrightarrow{R(A)} B + Q \\
A & \xrightarrow{\gamma_A} A - 1 \quad B \xrightarrow{\gamma_B} B - 1
\end{align*}
\]

(1)

where \(A\) and \(B\) are two biochemical species. Random variable \(A(t)\) denotes the number of molecules of species \(A\) at time \(t\) and similarly random variable \(B(t)\) denotes the number of molecules of species \(B\) at time \(t\). The random variable \(Q\) denotes the increase in the number of molecules of species \(B\) when a birth takes place. The parameter \(F \in \mathbb{R}^2\) is a constant that denotes the rate at which the species \(A\) is produced, \(R : \mathbb{Z}^2 \rightarrow \mathbb{R}^2\) is the rate of production of the species \(B\) and constants \(\gamma_A \in \mathbb{R}^2\) and \(\gamma_B \in \mathbb{R}^2\) represent the death rate of each molecule of species \(A\) and \(B\) respectively.

For any \(a, b \in \mathbb{Z}^2\), we denote by \(P(a, b, t)\) the probability \(A(t) = a, B(t) = b\). We assume that the random variable \(Q\) is independent from the random variables \(A(t), B(t)\) and denote by \(P_Q(q)\) the probability \(Q = q\). The CME, a version of the Chapman-Kolmogorov equation for Markov processes, for system (1) is

\[
\frac{\partial P(a, b, t)}{\partial t} = \gamma_A ((a + 1)P(a + 1, b, t) - aP(a, b, t)) + F[P(a - 1, b, t) - P(a, b, t)] + \gamma_B [(b + 1)P(a, b + 1, t) - bP(a, b, t)] + R(a) \sum_{q=1}^{b} P_Q(q)P(a, b - q) - P(a, b, t)
\]

(2)

The problem we are trying to address is to find an expression of the mean and variance of \(B\). This is a non-trivial problem due to the nonlinear rate \(R(A)\).

In this paper we present a bound on the variance of \(B\) obtained by exploiting a discrete expansion of \(R(A)\) based on the Newton series. The bound becomes exact when \(R(A)\) is linear. Furthermore, we show via numerical examples that it is very close to the true variance even in regimes where \(R(A)\) is highly nonlinear, in contrast to a LNA approximation for the variance.

It should be noted that filtering problems associated with (1) have also been studied in the literature. The optimal causal filter for estimating \(A\) when \(B\) is observed was derived in [10] and a close to optimal version was deployed in vitro in [11]. The average squared difference between \(A\) and \(B\) was quantified in [12].

The results that will be presented are associated with the equilibrium distribution of species \(A\) and \(B\) and we therefore make the following assumption.

Assumption 1: System (2) reaches a stationary distribution denoted by \(P(a, b)\) i.e. \(\lim_{t \to +\infty} P(a, b, t) = P(a, b)\).

Note that from the CME (2) \(P(a, b)\) satisfies the following

\[
\gamma_A [(a + 1)P(a + 1, b, a) - aP(a, b, a)] + F[P(a - 1, b, a) - P(a, b, a)] + \gamma_B [(b + 1)P(a, b + 1, a) - bP(a, b, a)] + R(a) \sum_{q=1}^{b} P_Q(q)P(a, b - q) - P(a, b, a) = 0
\]

(3)

For convenience in the notation we denote by \(A\) the random variable representing the number of molecules of species \(A\) at equilibrium, and its probability distribution is denoted by \(P\). Similarly, we denote by \(B\) the random variable representing the number of molecules of species \(B\) at equilibrium. Random variable \(Q\) can have any discrete distribution as long as it takes values in the set of positive integers excluding 0, and is independent of \(A\) and \(B\). It should also be noted that \(A \sim P(F/\gamma_A)\), i.e. random variable \(A\) has a Poisson distribution with mean \(F/\gamma_A\) which is a known result that follows from the fact that \(F\) and \(\gamma_A\) are constant (see e.g. [13]).

III. MAIN RESULTS

Before we present our main results we state a condition upon \(R(A)\).

Assumption 2: \(R : \mathbb{Z}^2 \rightarrow \mathbb{R}^2\) is a ratio of polynomials, i.e. \(R(A) = \{ \frac{f}{g} \mid f, g \in \mathbb{R}[A], g \neq 0 \ \forall A \in \mathbb{Z}^2 \}\).

Note that this assumption is mild within a biological perspective as reaction rates are typically rational functions of molecule numbers. In order to give our first result we define the quantities in (4) which are needed for the variance bound derived.

\[
\sigma_0 = \mathbb{E}_P[R(A)], \quad \sigma_1 = -\mathbb{E}_P[R(A)] + \frac{\mathbb{E}_P[AR(A)]}{\mathbb{E}_P[R(A)]}
\]

(4)

Note that in (4) \(\mathbb{E}_P[\cdot]\) denotes the expectation with respect to the Poisson distribution of the random variable \(A\).

Lemma 1: Let \(A \sim P(F/\gamma_A)\). Then the quantities \(\sigma_0\) and \(\sigma_1\) from (4) exist and are finite.

We now give our main result which is an expression that provides a lower bound on the variance of \(B\).

Proposition 1: Consider the system in (3) with \(R(A)\) satisfying Assumption 2. The following inequality holds

\[
\begin{align*}
\text{var}(B) > & (\gamma_A + \gamma_B)\sigma_0 (\mathbb{E}[Q^2] + \mathbb{E}[Q]) + 2(\mathbb{E}[Q])^2 \sigma_1^2 \mathbb{E}_P[A] \\
& 2\gamma_B (\gamma_B + \gamma_A)
\end{align*}
\]

(5)

where \(\sigma_0\) and \(\sigma_1\) are given in (4).

Remark 1: If \(R(A)\) is a linear function then, equation (5) holds with equality. This is stated in Proposition 2.

Remark 2: Parameters \(\sigma_0\) and \(\sigma_1\) exist and are bounded as stated in Lemma 1. Note that \(\mathbb{E}_P[\cdot]\) is the expectation with respect to a Poisson distribution with a known mean and may be computed to an arbitrary high precision by taking a sufficient amount of terms in the evaluation of the expectation.

Below we show that the bound is exact if all transition rates are linear.
Proposition 2: Consider the system in (3). If $R(A) = R_c A$ with $R_c \in \mathbb{R}^>$ the bound for the variance in Proposition 1 holds with equality and is given by (6)

$$\text{var}(B) = R_c \mathbb{E}_P[A] \left[ R_c \mathbb{E}(Q)^2 + (\gamma_A + \gamma_B)(\mathbb{E}[Q^2] + \mathbb{E}(Q)) \right]$$

Remark 3: The variance in the case of Proposition 2 can be computed analytically using LNA approaches, see e.g. [13] and [14].

Finally we give a side result on the covariance of $A$ and $B$.

Proposition 3: Consider the systems described in (3). Then $\text{cov}(A, B) = \text{cov}(B, A)$ satisfies

$$\text{cov}(A, B) = \mathbb{E}[Q] \frac{\sigma_1 \mathbb{E}_P[A]}{\gamma_A + \gamma_B}$$

where $\sigma_1$ is given in (4).

Remark 4: Note that Proposition 3 holds with equality also when $R(A)$ is nonlinear.

The results stated above have a number of useful properties relative to other more conventional approaches for quantifying the variance. As mentioned in Section I, Proposition 1 provides a bound for the variance when the transition rate $R(a)$ is nonlinear whereas existing methods like LNA and numerical simulations give only approximate values. As it will be seen in Section IV the variance calculation through the LNA is not as close to the variance of $B$ as is the bound provided in Proposition 1. This is especially important in the case of highly nonlinear propensity functions where LNA becomes less accurate. Furthermore, the numerical investigation in Section IV demonstrates that the bound maintains its accuracy despite the presence of bursts.

IV. EXAMPLES

In order to evaluate the conservativeness of the bound for the variance given in (5) we numerically investigate how close this is to the actual variance. The simulations have been performed using the software package GillesPy2 [15]. We select as function $R(A)$ a nonlinear function commonly found in biological systems, i.e. a Hill function $R(a) = \frac{(a/A_0)^{n_h}}{1+(a/A_0)^{n_h}}$ where $n_h \in \mathbb{R}^+$ and $A_0 \in \mathbb{R}^>$. We set as parameters $n_h = 9$ and $A_0 = 100$ and $F = 1$. This function clearly fulfills the conditions in Assumption 2.

Fig. 1 displays four cases corresponding to different burst distributions [16]. The figure highlights that the bound is not conservative regardless of the distribution of $Q$ and it is very close to $\text{var}(B)$ for different values of $\mathbb{E}_P[A]$ obtained by varying $\gamma_A$. The figures also show a comparison of the bound with the variance of $B$ computed through the LNA of system (3). Note that the computation carried out via LNA performs poorly for certain regimes. Specifically, the performance is particularly poor when $A$ is with high probability in the nonlinear regime of the Hill function, while it is more accurate when $A$ is predominantly in the linear regime.

The left hand side of Fig. 2 shows the relative error, expressed as a percentage, between the variance of $B$ and the bound corresponding to different values of $\mathbb{E}_P[A]$ (obtained by varying $\gamma_A$) and $\gamma_B$ when $Q$ has a geometric distribution, i.e. $Q \sim \mathcal{G}(0.5)$. The right hand side of Fig. 2 displays the relative error between the variance of $B$ and the variance obtained trough LNA. We find that the bound is still not conservative and performs consistently better than variance obtained trough LNA for different values of $\gamma_A$ and $\gamma_B$.

V. CONCLUSIONS

We have derived a hard bound for the variance in the molecule numbers of a species formed with bursts, with a nonlinear rate that depends on another spontaneously formed species. The bound follows from a discrete expansion based on the Newton series, and exploits spectral properties of the master equation. We have also shown that the bound holds with equality if the propensity functions are linear. The accuracy of the bound has been investigated with numerical simulations which demonstrate that this is very close to the actual variance also when the rate of formation of the species is highly nonlinear, and despite the presence of bursts. Future work will focus on extending the results to larger classes of CMEs.
VI. APPENDIX

In the appendix we provide the proofs of the results presented in the main text.

A. Proof of Lemma 1

Proof: The proof follows by noting that \( \sigma_0 \) and \( \sigma_1 \) are expectations of rational functions with respect to a Poisson distribution.

B. Proof of Proposition 1

Proof:

We divide our proof in three parts. In the first part we compute \( T_a'(1) \), where \( T_a(z) = \sum_{b=0}^{\infty} z^b P(a, b) \) is a probability generating function that is used for the calculation of the moments of \( B \). In particular, we show that \( T_a'(1) \) satisfies an equation that involves a difference operator that depends on the CME.

In Part 2 we derive an expression for \( T_a'(1) \) in terms of the eigenfunctions of this operator. This expression depends on the Newton series expansion of \( R(A) \).

In Part 3 we make use the results in Part 1, 2 to derive an expression for the variance of \( B \), and hence deduce a bound in terms of \( \sigma_0, \sigma_1 \), as stated in Proposition 1.

Part I: Consider (3) and let us make use of the generating function \( T_a(z) = \sum_{b=0}^{\infty} z^b P(a, b) \) and \( M(z) = \sum_{b=0}^{\infty} z^b Q(q) \). Under these transformations we compute the following equation.

\[
\gamma_A [(a + 1)Ta_{a+1}(z) - aTa_{a}(z)] + F[Ta_{a-1}(z) - Ta_{a}(z)] + \gamma_B (1 - z)Ta_{a}(z) + R(a)(M(a) - 1)Ta_{a}(z) = 0 \quad (8)
\]

Note that (8) provides an alternative way to express (3). By setting \( z = 1 \) we have \( T_a(1) = P_A(a) \), where \( A \) in the considered case has a Poisson distribution, see [13]. Secondly, we can compute moments of \( B \) by differentiating (8) with respect to \( z \) and setting \( z = 1 \). In particular, we have

\[
T_a'(1) = \sum_{b=0}^{\infty} bP(a, b), \quad T_a''(1) = \sum_{b=0}^{\infty} b(b-1)P(a, b) \quad (9)
\]

Hence by differentiating (8) with respect to \( z \) and then setting \( z = 1 \) we obtain the following expression involving the first moment

\[
\gamma_A [(a + 1)Ta_{a+1}(1) - aTa_{a}(1)] + F[Ta_{a-1}(1) - Ta_{a}(1)] - \gamma_B Ta_{a}(1) + R(a)(M'(1)Ta_{a}(1)) = 0 \quad (10)
\]

where \( M'(1) = E[Q] \).

For the second moment we have

\[
\gamma_A [(a + 1)Ta_{a+1}(1) - aTa_{a}(1)] + F[Ta_{a-1}(1) - Ta_{a}(1)] - 2\gamma_B Ta_{a}(1) + R(a)(M''(1)Ta_{a}(1) + 2M'(1)Ta_{a}(1)) = 0 \quad (11)
\]

where \( M''(1) = E[Q^2] - E[Q] \). By summing (10) over all \( a \) yields

\[
\sum_{a=0}^{\infty} T_a'(1) = \frac{M'(1)}{\gamma_B} \sum_{a=0}^{\infty} R(a)Ta_{a}(1) \quad (12)
\]

Note that the left hand side of (12) is the expectation of \( B \) with respect to the known distribution of \( A \). From the definition of \( T_a'(1) \) and by noting that \( T_a(1) = P_A(a) \), (12) can be written as

\[
E[B] = \frac{E[Q]}{\gamma_B} E_P[R(A)] \quad (13)
\]

For the second moment of \( B \) we proceed analogously as for the first moment by summing (11) over all \( a \) and obtain

\[
\sum_{a=0}^{\infty} T_a''(1) = \frac{1}{2\gamma_B} \left( M''(1) \sum_{a=0}^{\infty} R(a)Ta_{a}(1) + 2M'(1) \sum_{a=0}^{\infty} R(a)Ta_{a}(1) \right) \quad (14)
\]

From (9) and (14) we obtain

\[
E[B^2] - E[B] = \frac{1}{2\gamma_B} (M''(1)E_P[R(A)] + 2M'(1)E[BR(A)]) \quad (15)
\]

where for the last term on the right hand side of (14) we have applied the definition of \( T_a'(1) \).

We now seek in the remainder of the proof to derive an expression for the last term in (15).

We now consider the following representation of \( T_a''(1) \), where function \( G(a) \) is to be determined:

\[
T_a'(1) = \frac{T_a(1)G(a)}{\gamma_B} \quad (16)
\]

We substitute (16) into (10) and obtain

\[
\frac{\gamma_A}{\gamma_B} [(a + 1)Ta_{a+1}(1)G(a + 1) - aTa_{a}(1)G(a)] + \frac{F}{\gamma_B} [Ta_{a-1}(1)G(a - 1) - Ta_{a}(1)G(a)] - G(a)Ta_{a}(1) + M'(1)R(a)Ta_{a}(1) = 0 \quad (17)
\]
Note that from the definition of the Poisson distribution we have:

$$T_{a+1}(1) = \frac{F}{\gamma_A(a+1)} T_a(1), \quad T_{a-1}(1) = \frac{a\gamma_A}{F} T_a(1)$$  \hspace{1cm} (18)

By rearranging (17) and using (18) leads to

$$T_a(1)[(S - 1)G(a) + M'(1)R(a)] = 0$$  \hspace{1cm} (19)

where $S$, defined in (20), is an operator acting upon $G(a)$

$$S = \gamma_B^{-1}(\gamma_A a\Delta_1 + F\Delta_1) = \frac{\gamma_A}{\gamma_B} (a\Delta_1 + EP[A]\Delta_1)$$  \hspace{1cm} (20)

By making use of (19) we find that $G(a)$ satisfies

$$(1 - S)G(a) = M'(1)R(a)$$  \hspace{1cm} (21)

We now seek to exploit spectral properties of $S$ in order to derive an expression for $G(a)$ and hence of $T'_a(1)$.

Part 2: In this part of the proof we derive an expression for $T'_a(1)$.

We first focus on the operator $S$ and its corresponding eigenfunctions. First let us note that $\Delta_1[(a)k] = k(a)_{k-1}$ and $a\Delta_1[(a)k] = -k(a)_{k}$ and hence we can write $S[(a)k]$ as

$$S[(a)k] = -\frac{k\gamma_A}{\gamma_B} [(a)_{k} - EP[A](a)_{k-1}]$$  \hspace{1cm} (22)

We will derive an expression for $G(a)$ in terms of the eigenfunctions and eigenvalues of operator $S$. We derive those below, following the approach in [17]. To facilitate the analysis of (22) we introduce operator $F$, which for a monomial $a^k$, $a, k \in \mathbb{Z}^+$ it satisfies the property $F[a^k] = (a)k$. It is also defined to be linear, i.e. $F[a_{1}^k + a_{2}^k] = (a_{1})^k + (a_{2})^k$, $F[\lambda a^k] = \lambda(a)_{k}$, $\lambda \in \mathbb{R}$.

We can hence rewrite (22) as

$$S[F[a^k]] = -\frac{k\gamma_A}{\gamma_B} F[a^{k-1}(a - EP[A])]$$  \hspace{1cm} (23)

which can also be written as

$$S[F[a^k]] = -\frac{\gamma_A}{\gamma_B} \left[ \frac{d}{da} (a - EP[A]) \right]$$  \hspace{1cm} (24)

We now apply operator $S$, in combination with operator $F$, to $(a - EP[A])^k$ which yields

$$S[F[(a - EP[A])^k]] = -\frac{\gamma_A}{\gamma_B} \left[ \frac{d}{da} (a - EP[A])^k \right]$$  \hspace{1cm} (25)

By observing that $\frac{d}{da} [(a - EP[A])^k] = k(a - EP[A])^{k-1}$ we have

$$S[F[(a - EP[A])^k]] = -k\frac{\gamma_A}{\gamma_B} F[(a - EP[A])^k]$$  \hspace{1cm} (26)

We now use the binomial expansion to rewrite the eigenfunctions and corresponding eigenvalues, which take the form

$$\lambda_n = -n\frac{\gamma_A}{\gamma_B}$$  \hspace{1cm} (27)

$$\psi_n(a) = \sum_{k=0}^{n} \binom{n}{k} (-EP[A])^k (a)_{n-k}$$  \hspace{1cm} (28)

Using the expressions derived for the eigenfunctions and eigenvalues of $S$ we now proceed to derive an expression for $G(a)$. In particular, by noting the decomposition of $R(a)$ in terms of the eigenfunctions $\psi_n$ given in Proposition 4 we make the following ansatz on $G(a)$

$$G(a) = M'(1) \sum_{n=0}^{\infty} \sigma_n \zeta(n) \psi_n(a)$$  \hspace{1cm} (29)

where $\zeta(n)$ is a function to be determined and $\sigma_n$ are as defined in Proposition 4. By substituting (29) in (21) we get

$$M'(1) \sum_{n=0}^{\infty} \sigma_n \left(1 + n\frac{\gamma_A}{\gamma_B}\right) \zeta(n) \psi_n(a) = M'(1)R(a)$$  \hspace{1cm} (30)

Noting the series expansion of $R(a)$ in Proposition 4 with $r = EP[A]$ we deduce that

$$\zeta(n) = \frac{\gamma_B}{n\gamma_A + \gamma_B}$$  \hspace{1cm} (31)

Therefore from (16), (29), (31) we deduce the following expression for $T'_a(1)$

$$T'_a(1) = \frac{T_a(1)M'(1)}{\gamma_B} \sum_{n=0}^{\infty} \sigma_n \frac{\gamma_B}{n\gamma_A + \gamma_B} \psi_n(a)$$  \hspace{1cm} (32)

Part 3: We now make use of the series expansion of $T'_a(1)$ in (32) to compute the moments of $B$.

We evaluate first the unknown term in the right hand side of (15). In particular, by noting that $\mathbb{E}[BR(A)] = \sum_{n=0}^{\infty} R(a)T'_a(1)$ and from the expression of $T'_a(1)$ in (32) we have

$$\mathbb{E}[BR(A)] =$$

$$= \frac{M'(1)}{\gamma_B} \sum_{a=0}^{\infty} \sum_{n=0}^{\infty} \sigma_n \frac{\gamma_B}{n\gamma_A + \gamma_B} \psi_n(a) T'_a(1)$$

$$= \mathbb{E}[Q] \frac{\gamma_B}{EP} \left[ \sum_{\lambda=0}^{\infty} \sum_{n=0}^{\infty} \sigma_n \sigma_n' \frac{\gamma_B}{n\gamma_A + \gamma_B} \psi_n(A) \psi_n'(A) \right]$$  \hspace{1cm} (33)

where $M'(1) = \mathbb{E}[Q]$ and in the second step we have used the results of Proposition 4. We now evaluate the expectation in (33) as follows

$$\mathbb{E}[P] \left[ \sum_{n=0}^{\infty} \sum_{\lambda=0}^{\infty} \sigma_n' \sigma_n \frac{\gamma_B}{n\gamma_A + \gamma_B} \psi_n(A) \psi_n'(A) \right] =$$

$$= \sum_{n=0}^{\infty} \sigma_n^2 \frac{\gamma_B n! (EP[A])^n}{n\gamma_A + \gamma_B}$$  \hspace{1cm} (34)

where the equality holds given the results of Proposition 5. Hence we can now write the following expression for $\mathbb{E}[BR(A)]$

$$\mathbb{E}[BR(A)] = \frac{M'(1)}{\gamma_B} \sum_{n=0}^{\infty} \sigma_n^2 \frac{\gamma_B n! (EP[A])^n}{n\gamma_A + \gamma_B}$$  \hspace{1cm} (35)
We finally express $\text{var}(B)$ as in (36) by making use of (13), (15), and (35)

\begin{equation}
\text{var}(B) = \\
\frac{1}{2\gamma_B} \left[ (E[Q^2] - E[Q])E_P[R(A)] \\
+ \frac{2E(Q)^2}{\gamma_B} \sum_{n=0}^{\infty} \gamma_B n! (E_P[A])^n \frac{n\gamma_A + \gamma_B}{n!} \right] \\
+ \frac{E_P[R(A)]E[Q]}{\gamma_B} \left( E_P[R(A)]E[Q] \right)^2
\end{equation}

(36)

It should be noted that from Proposition 5 we have $E_P[\psi_n(A)] = \delta_{n,0}$ and hence by computing the expectation with respect to the Poisson distribution on both sides of (40) we have that $E_P[R(A)] = \sigma_0$. By substituting $\sigma_0$ in (36) and through some manipulation we obtain

\begin{equation}
\text{var}(B) = \\
\frac{1}{2\gamma_B} \left[ (E[Q^2] - E[Q])\sigma_0 \\
+ \frac{2E(Q)^2}{\gamma_B} \sum_{n=1}^{\infty} \gamma_B n\sigma_0 (E_P[A])^n \frac{n\gamma_A + \gamma_B}{n!} \right] + \frac{E[Q]}{\gamma_B} \sigma_0
\end{equation}

(37)

Since all terms in (37) are positive by using only the first term in the summation appearing in (37) we obtain

\begin{equation}
\text{var}(B) > \\
\frac{(\gamma_A + \gamma_B)\sigma_0 (E[Q^2] + E[Q]) + 2E(Q)^2 \sigma_0 E_P[A]}{2\gamma_B (\gamma_B + \gamma_A)}
\end{equation}

(38)

C. Expansion of $R(A)$

The proposition below is based on ideas in [12]. The proof is omitted due to page constraints.

**Proposition 4:** Consider $R(A) : \mathbb{Z}^+ \to \mathbb{R}^+$, that satisfies Assumption 2. Let $A \in \mathbb{Z}^+$ and consider the function

\begin{equation}
\psi_n(A) = \sum_{k=0}^{n} \binom{n}{k} (-r)^k (A)_{n-k}
\end{equation}

(39)

where $r \in \mathbb{R}^+$. Then the following equality holds:

\begin{equation}
R(A) = \sum_{n=0}^{\infty} \sigma_n \psi_n(A)
\end{equation}

(40)

where

\begin{equation}
\sigma_n = \sum_{k=n}^{\infty} \binom{k}{n} \frac{\Delta^k[R(A)]}{k!} \bigg|_{A=0} r^{k-n}
\end{equation}

(41)

D. Proof of Proposition 2

**Proof:** The proof follows by substituting the values of $\sigma_0$ and $\sigma_1$ in (5) when $R(\alpha) = R_c \alpha$ and by comparing with that obtained using LNA (see [13] and [14]).

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**REFERENCES**


