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Quasi-Birth-Death Processes, Tree-Like QBDs, Probabilistic 1-Counter Automata, and Pushdown Systems

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Abstract

We begin by observing that (discrete-time) Quasi-Birth-Death Processes (QBDs) are equivalent, in a precise sense, to (discrete-time) probabilistic 1-Counter Automata (p1CAs), and both Tree-Like QBDs (TL-QBDs) and Tree-Structured QBDs (TS-QBDs) are equivalent to both probabilistic Pushdown Systems (pPDSs) and Recursive Markov Chains (RMCs).

We then proceed to exploit these connections to obtain a number of new algorithmic upper and lower bounds for central computational problems about these models. Our main result is this: for an arbitrary QBD (even a null-recurrent one), we can approximate its termination probabilities (i.e., its G matrix) to within i bits of precision (i.e., within additive error $1/2^i$), in time polynomial in both the encoding size of the QBD and in i , in the unit-cost rational arithmetic RAM model of computation. Specifically, we show that a decomposed Newton’s method can be used to achieve this.

We emphasize that this bound is very different from the well-known “linear/quadratic convergence” of numerical analysis, known for QBDs and TL-QBDs, which typically gives no constructive bound in terms of the encoding size of the system being solved. In fact, we observe (based on recent results for pPDSs) that for the more general TL-QBDs this bound fails badly. Specifically, in the worst case Newton’s method “converges linearly” to the termination probabilities for TL-QBDs, but requires exponentially many iterations in the encoding size of the TL-QBD to approximate these probabilities within any non-trivial constant error $c < 1$.

Our upper bound proof for QBDs combines several ingredients: a detailed analysis of the structure of 1-counter automata, an iterative application of a classic condition number bound for errors in linear systems, and a very recent constructive bound on the performance of Newton’s method for monotone systems of polynomial equations.

1 Introduction

A variety of important stochastic models are finitely presentable but describe an infinite-state underlying stochastic process. Among the many examples are (multi-type) branching processes, (quasi-)birth-death processes, stochastic petri nets, and stochastic context-free grammars. Computation of basic quantities associated with such stochastic models (both transient analyses and steady-state analyses) are fundamental to many applications. Yet the complexity of computing many such quantities is not adequately understood.

This paper begins by observing that there is a close correspondence between different denumerable-state probabilistic models studied, on the one hand, in the queueing theory and structured Markov chain community, and, on the other hand, more recently, in the literature on analysis and model checking of recursive probabilistic procedural programs. Specifically, we observe that discrete-time Quasi-Birth-Death processes (QBDs) are equivalent, in a precise sense, to probabilistic 1-Counter Automata (p1CAs), which are in turn a strict subclass of probabilistic Pushdown Systems (pPDSs), namely they are pPDSs restricted to a 1-letter stack alphabet. Furthermore, we show that Tree-Structured and Tree-Like QBDs (TL-QBDs), which are extensions of QBDs, are indeed equivalent to pPDSs and Recursive Markov Chains (RMCs).

These results are not at all surprising once one gets over the differences in notation and language used by the two communities. Both types of models are infinite-state structured Markov chains that are finitely presented; in the case of QBDs and their tree extensions the notation and methodology is more algebraic, matrix-based, while in the case of pPDSs it is more automata-theoretic and combinatorial.

We exploit these equivalences to obtain several new algorithmic results about these models. A number of results follow immediately from the equivalences and existing results about the various models. For in-

stance, it follows from results on RMCs that quantitative model-checking of linear-time (ω -regular) temporal properties for QBDs and TL-QBDs can be decided in PSPACE in the size of the model ([14]). On the other hand, obtaining any non-trivial approximation of the “termination probabilities” for TL-QBDs (the analog of the G matrix of QBDs), even to within any constant additive factor $c < 1/2$, is at least as hard as long standing open problems in *exact* numerical computation, such as the *square-root sum* problem, whose complexity (in the standard Turing model of computation) is not even known to be in NP [16].

Our main result is a new upper bound on numerical approximation of central quantities associated with QBDs. Specifically, we show that, given a QBD (even a null-recurrent one), the basic G matrix of “termination probabilities” for the QBD (and various other quantities of interest that can be derived from it) can be approximated to within i bits of precision in time polynomial in *both* the encoding size of the QBD and in i , in the unit-cost rational arithmetic RAM (i.e., rational Blum-Shub-Smale) model of computation. More precisely, in the stated time complexity in the unit-cost model, one can compute a matrix $\tilde{G} \geq 0$ such that $\|G - \tilde{G}\|_\infty \leq 1/2^i$. Specifically, we show that the decomposed Newton’s method (studied for RMCs and for arbitrary monotone systems of polynomial equations in [13]) can be used to achieve this bound.

We emphasize that this analysis is very different from the well-known “linear/quadratic convergence” analyses traditional to numerical analysis, which is known to hold (in null-recurrent/non-null-recurrent cases, respectively) on the equations that arise for QBDs and TL-QBDs, using Newton’s method and several other methods (such as logarithmic reduction and cyclic reduction). “Linear/quadratic convergence” results only bound the number of iterations required as a function of the desired error $\epsilon > 0$ (i.e., the desired number of bits i of precision). They completely ignore how large the number of iterations may need to be as a function of the encoding size of the input QBD.

In fact, we observe using recent results for pPDSs ([19]) that this polynomial upper bound for QBDs fails badly for TL-QBDs. Specifically, there are worst-case examples of TL-QBDs which require exponentially many iterations of Newton’s method, as a function of the size of the TL-QBD, in order to approximate termination probabilities (the analog of the G matrix for TL-QBDs) to within any non-trivial constant additive error, thus even to within 1 bit of precision. This is the case even though Newton’s method is “linearly convergent” on these examples. Our results thus reveal a vast difference in the worst case behavior of Newton’s

method on QBDs and TL-QBDs, not apparent from the usual “linear/quadratic” convergence analysis.

Our proof of the new upper bound for QBDs relies on several ingredients. We first perform a detailed analysis of the structure of 1-counter automata, establishing key properties. Firstly, there is a fixed polynomial, $q(n)$, such that for any QBD whose encoding size is n ,¹ the termination probabilities (i.e., entries of the G matrix), which may of course be irrational, are each either 0 or $\geq 1/2^{q(n)}$. This bound fails badly for TL-QBDs, as there are simple examples (already noted for RMCs [13]) of size $O(n)$ for which positive termination probabilities are $1/2^{2^n}$. As a second crucial property, we show that the dependencies among variables in the non-linear (matrix) equation $X = A_{-1} + A_0X + A_1X^2$, associated with a QBD (whose least non-negative solution is the G matrix) have a very special structure when decomposed into strongly connected components (SCCs). Roughly speaking, the SCCs can have non-linear internal structure, but distinct nonlinear SCCs can not “depend” on each other. This special structure does not hold for the equations associated with termination probabilities of TL-QBDs.

These two structural results allow us to bring in other key ingredients in order to establish the polynomial upper bound for QBDs. Specifically, we use an important constructive upper bound recently established in [9] on the performance of Newton’s method for (strongly connected) monotone systems of polynomial equations, combined with our result that QBD termination probabilities can be “polynomially” bounded away from zero, in order to establish that for the non-linear SCCs in the equations for G , a polynomial number of iterations of Newton’s method (as a function of the encoding size and number of bits of precision), starting from the 0 vector, suffice to obtain a desired number of bits of precision for the variables in a non-linear SCC. Finally, to approximate the entire matrix G , we deal with a possibly nested series of linear SCCs “above” nonlinear ones in the DAG of SCCs, by using an iterative application of a classic, but rather delicate, condition number bound for errors in the solution of linear systems resulting from coefficient errors.

On the other hand, as a “lower bound” for QBDs, we show that deciding whether $G_{i,j} \geq p$, for a given rational p , is at least as hard as the square-root sum problem. Thus, resolving *exact* quantitative decision problems for QBDs in polynomial time or even in NP, in the traditional Turing model, is not possible with-

¹In other words, n is the number of bits needed to describe the QBD, by describing all the rational coefficients (given by numerator and denominator in binary) in all the $m \times m$ matrices that define the QBD.

out a major breakthrough in exact numerical analysis. By contrast, for the more general TL-QBDs, we observe that our recent result in [15] for RMCs implies that even the problem of obtaining any non-trivial *approximation* of termination probabilities for TL-QBDs is square-root-sum hard.

These results lead us to suspect that a similar difference should exist in the worst-case behavior on QBDs and TL-QBDs for other numerical solution methods such as the logarithmic or cyclic reduction type algorithms (see, e.g., [2]). We have however not analysed these other algorithms. Indeed, the equivalences we point out open the door for the extensive methods and algorithms developed in the structured Markov chain community (which after all has a much longer history) to be applied to the analysis of the more recently studied models like pPDSs and RMCs, for analysis and model checking of recursive probabilistic procedural program. In the other direction, we feel that the “automata-theoretic” viewpoint, offered by the work on RMCs and pPDSs, and related literature, can be further exploited in research on QBDs, TL-QBDs, and related models. In any case, we believe a cross-fertilization between these two communities will be a fruitful source of research in the near future. A tool called PReMo [33] which implements optimized versions of the decomposed Newton’s method and other methods for the analysis of Recursive Markov Chains (and their controlled and game extensions) has been augmented with an input format for QBDs.

We have conducted a very preliminary comparison of PReMo’s performance on QBDs with that of an existing tool for QBDs: SMCSolver [3]. SMCSolver’s implementation of algorithms like (shifted) cyclic reduction handily beats PReMo (by an order of magnitude or more) on large “dense” QBDs where the input A_i matrices are dense. This is explained by the following facts: firstly, such dense systems are typically not decomposable; moreover, SMCSolver exploits concise matrix representations of the nonlinear equations associated with QBDs, which require $O(n^2)$ encoding size (where n is the number of control states, and assuming bounded size coefficients), whereas PReMo employs an explicit algebraic formula representation of these equations (which allows handling arbitrary monotone systems of nonlinear equations) which for dense input A_i ’s requires $O(n^3)$ encoding size. Algorithms (like cyclic reduction) employed in SMCSolver operate directly on these matrix equations, and thus have far lower cost per iteration. However, unlike PReMo, SMCSolver does not exploit the potential for decomposing these equations (indeed, decomposition can destroy their simple matrix equation form). Thus on very decomposable

systems, PReMo can do better. In particular, we constructed a family of highly decomposable QBDs, where even the G matrix itself is sparse. On these examples, with 500 control states, SMCSolver’s fastest algorithm (shifted cyclic reduction) took over 40 seconds to achieve 10^{-10} error bounds, and on 5000 control states it crashed. PReMo ran in under 1 second even for 5000 control states, to achieve 10^{-10} error. However, these are very contrived examples explicitly designed to be highly decomposable. On more dense examples SMCSolver is far superior to PReMo. An interesting line of future research would be to combine the benefits of the concise matrix representations employed in, e.g., SMCSolver, and the decomposition methods employed in PReMo. One challenge in this regard is this: Newton’s method can also be carried out directly over $O(n^2)$ sized matrix equations for QBDs, with low cost per iteration ($O(n^3)$ operations), using known efficient methods for solving the concise linear matrix equations that arise in each iteration of Newton’s method over QBDs (certain generalized Sylvester matrix equations, see [2]). However, while TL-QBDs and RMCs also have nonlinear equations with $O(n^2)$ matrix representations, no such efficient solution method is known for the more general linear matrix equations that arise in iterations of Newton’s method on them. Finding such a method would make Newton’s method more practical on large “dense” TL-QBDs, RMCs, and pPDSs. But even with such an efficient method, it remains a challenge to combine it well with decomposition, because in general decomposition destroys the matrix form of the equations.

Due to space constraints, all proofs are omitted from this conference paper. Please see the full version [12].

Related work: Quasi-Birth-Death Processes (QBDs) and more generally M/G/1-type and G/M/1-type Markov chains have been studied for decades in queueing theory, performance evaluation, and related areas, both in discrete and continuous time, and so have numerical solution methods for them (see, e.g., the books [25, 26, 23, 2]). In particular, Latouche in [22], studied the behavior of Newton’s method on these models, and showed (building on [27]) that under certain assumptions (namely when $A = \sum_i A_i$ is irreducible and parameter $\rho \neq 1$) the Newton iterates are well defined and converge monotonically and “quadratically” to the matrix G . Several other “quadratically convergent” methods have also been developed, e.g., logarithmic reduction [24], and cyclic reduction (see [2]). Remke et. al. in [29] have studied numerical algorithms for model checking of continuous-time QBDs against properties expressed in the continuous-time temporal logic CSL. Several other models, in particular, (discrete-time) stochastic Petri Nets restricted to

markings where just one place can be unbounded, are already known to be equivalent to QBDs (see, e.g., [28]).

Tree-Structured QBDs (TS-QBDs) are a generalization of QBDs, first studied in [36, 30, 35]. Tree-Like QBDs (TL-QBDs) are a restriction of TS-QBDs, studied in, e.g., [23, 4, 32]. It was already observed in [31] that TL-QBDs and TS-QBDs are equivalent, under a tight notion of equivalence which amounts to an instance of what we use to show equivalence also to pPDSs and RMCs. Bini et. al. [4] studied the performance of several numerical algorithms for TL-QBDs, including Newton’s method. Building on [22], they show that under a similar set of assumptions, Newton’s iterations are defined and converge monotonically and quadratically for various quantities such as the termination probabilities (the analog of the G matrix).

Pushdown automata are of course classic models that date back to the origins of automata theory (see, e.g., [17]). They have many applications, e.g., in parsing of languages. Pushdown systems (the transition graphs of pushdown automata), and equivalent models such as Recursive State Machines, have been studied extensively in the past decade for the analysis and model checking of procedural programs (see, e.g., [8, 1]). In more recent years, researchers have extended these models with probabilistic behavior, i.e., to probabilistic Pushdown Systems (pPDSs) ([10, 6, 11, 5]) and Recursive Markov Chains ([13, 14, 34]), and developed model checking algorithms for them. In particular, results in [14] yield that linear-time ω -regular quantitative model checking of RMCs and pPDSs can be decided in PSPACE (we note that this is an upper bound for an exact decision procedure, not numerical estimation). A key role was played in all these analyses by the computation of termination probabilities (the analog of the G matrix) for RMCs. A number of “lower bounds” were established in [13, 15], showing that these upper bounds could not be substantially improved without major breakthroughs on long standing open problems in exact numerical computation. In [13], a decomposed Newton’s method was studied for approximation of termination probabilities, and it was shown that, after decomposition, Newton’s method converges monotonically, starting from 0, for arbitrary monotone polynomial systems (MSPs) that do have a non-negative solution. Subsequently, [19, 9] studied in much greater detail the performance of (decomposed) Newton’s method on such monotone systems of polynomial equations. They not only established worst-case linear convergence results (even when the Jacobian at the least fixed point (LFP) is singular), but importantly for our results in this paper, they also provided a strong constructive upper bound on the number of it-

erations required for Newton’s method as a function of the encoding size of the polynomial system. For pPDSs and RMCs their upper bounds require exponentially many iterations in the model’s size in order to converge to within a constant factor of termination probabilities, but we will exploit their results, and other things, to show polynomial bounds for QBDs.

1-Counter Automata, which amount to Pushdown Systems with only one stack symbol, are a standard automata-theoretic model, and their relationship to other infinite-state models in automata theory has been well studied (see, e.g., [21, 20]). Probabilistic 1-Counter Automata have not yet been extensively studied in the literature on model checking and verification. Recently however, Brožek and Kučera have informed us [7], that they have obtained a (as yet unpublished) polynomial-time algorithm for deciding whether termination occurs almost surely (with probability 1) starting from a given control state, and counter value 1, for a probabilistic 1-counter automaton.

The rest of this paper is organized as follows. Section 2 gives basic definitions. In Section 3 we show the equivalence between QBDs and p1CA, and between Tree-structured and Tree-like QBDs and pPDS, state some consequences, and show that the square-root sum problem reduces to the decision problem for QBDs. In Section 4 we prove important structural properties of p1CAs, and in Section 5 we use them to analyze the decomposed Newton method for QBDs and prove a polynomial bound on the number of iterations.

2 Definitions

Efficient embeddings and equivalences. We show various probabilistic models are “essentially equivalent”. To make the notion of “essentially equivalent” precise, we use the following definitions.

Definition 1 *For a (countable-state, discrete-time)*

Markov chain $\mathcal{M}$ with states t and t' , we write $t \xrightarrow{\bar{t}, p} t'$ to denote that there is a sequence of states $\bar{t} = t_0, \dots, t_k$, where $t_0 = t$ and $t_k = t'$, and probabilistic transitions (t_0, p, t_1) and $(t_i, 1, t_{i+1})$ for $1 \leq i < k$. (Note that if $k = 1$, this just says that (t, p, t') is a transition of $\mathcal{M}$.)

We shall say that one (countable state) Markov chain $\mathcal{M}$ embeds efficiently in another Markov chain $\mathcal{M}'$, if there exist two polynomial-time computable mappings, f, g , where f is a one-to-one mapping from states of $\mathcal{M}$ to states of $\mathcal{M}'$, and g is a one-to-one mapping that maps a transition (t, p, t') of $\mathcal{M}$ to a sequence, $\bar{t} = t_0 \dots t_k$ of states in $\mathcal{M}'$, with $t_0 = f(t)$ and $t_k = f(t')$, and such that $f(t) \xrightarrow{\bar{t}, p} f(t')$ holds in $\mathcal{M}'$, and furthermore such that none of the auxiliary states $t_1, \dots, t_{k-1}$ are in the range of the mapping f .

Intuitively, this is essentially a monomorphic embedding of one Markov chain inside another, except that a transition (t, p, t') can be “stretched” into a sequence of transitions, using intermediate auxiliary states, and with probability 1 transitions out of these auxiliary states leading to the target, $f(t')$. All models we consider, even countable-state ones, have a finite description. So, for a family $\mathcal{F}$ of finite presentations of Markov chains, each $\mathcal{A} \in \mathcal{F}$, describes a potentially infinite-state underlying chain $M(\mathcal{A})$.

Definition 2 *If $\mathcal{F}$ and $\mathcal{F}'$ are two classes of finitely-presented Markov chains, we say that $\mathcal{F}$ is efficiently subsumed by $\mathcal{F}'$ iff: there is a polynomial-time computable mapping $h : \mathcal{F} \mapsto \mathcal{F}'$, which maps a model $\mathcal{A} \in \mathcal{F}$ to a $h(\mathcal{A}) \in \mathcal{F}'$, and such that there exists a pair of functions $f_{\mathcal{A}}$ and $g_{\mathcal{A}}$, which can themselves be efficiently computed (as Turing machines) from $\mathcal{A}$, and such that $f_{\mathcal{A}}$ and $g_{\mathcal{A}}$ constitute an efficient embedding of $M(\mathcal{A})$ into $M(h(\mathcal{A}))$. Finally, we say two classes $\mathcal{F}$ and $\mathcal{F}'$ of finitely-presented chains are M-equivalent if both of them are efficiently subsumed by the other.*

It is not hard to see that if one family $\mathcal{F}$ of finitely presented Markov chains is efficiently subsumed by another family $\mathcal{F}'$, via a mapping h , then a variety of computational problems for $M(\mathcal{A})$, where $\mathcal{A} \in \mathcal{F}$, efficiently reduce to basically the same analyses of $M(h(\mathcal{A}))$ where $h(\mathcal{A}) \in \mathcal{F}'$. These include both transient analyses (such as reachability or hitting probability) as well as limit distributions.

In all the probabilistic models we define, we assume that all probability coefficients in the models are rational (for computational purposes), and that they are encoded in the standard way, by providing numerator and denominator in binary.

Probabilistic Pushdown Systems. There are a number of equivalent variations on the definition of (probabilistic) Pushdown Systems. We use a standard definition which is convenient for analysis. A *probabilistic Pushdown System* (pPDS) $\mathcal{P} = (Q_P, \Gamma, \Delta)$ consists of a set of control states Q_P , a stack alphabet Γ , and a probabilistic transition relation $\Delta \subseteq (Q_P \times \Gamma) \times [0, 1] \times Q_P \times \{\text{swap}(\Gamma), \text{swap}\&\text{push}(\Gamma \times \Gamma), \text{pop}\}$. That is, a transition has the form $((s, \gamma), p_{(s, \gamma), (s', \mathcal{C})}, (s', \mathcal{C}))$, where based on the control state s and the symbol on top of the stack, γ , with probability $p_{(s, \gamma), (s', \mathcal{C})}$, the transition updates the control state to s' , and performs action $\mathcal{C}$ on the stack: If $\mathcal{C} = \text{swap}(\gamma')$ then the action swaps the top-of-stack symbol, γ , with symbol γ' . If $\mathcal{C} = \text{swap}\&\text{push}(\gamma', \gamma'')$, then the action both swaps γ with γ' and then pushes γ'' on top of the stack. Lastly, if $\mathcal{C} = \text{pop}$, then the action pops the top-stack-symbol γ off the stack. Each such transi-

tion has an associated probability $p_{(s, \gamma), (s', \mathcal{C})}$, and we assume that for each pair (s, γ) of control state and top of stack symbol, $\sum_{(s', \mathcal{C})} p_{(s, \gamma), (s', \mathcal{C})} = 1$. We assume there is a special stack symbol $\perp \in \Gamma$ that marks the bottom of the stack. Accordingly, $\perp$ is never overwritten with a different stack symbol, nor popped off the stack, and is never pushed onto the stack or overwrites a different stack symbol. A stack with letter γ at the top and remaining content $\omega \in \Gamma^*$ will be written $\omega\gamma$ (note that the leftmost symbol in $\omega\gamma$ is $\perp$). A pPDS $\mathcal{P}$ defines a countable-state Markov chain $M(\mathcal{P}) = (V', \Delta')$ in an obvious way. Namely, the states of $M(\mathcal{P})$ are $V' = \{(w, s) \mid s \in Q_P, w \in \perp\Gamma^*\}$, and the probabilistic transitions of $M(\mathcal{P})$ are $\Delta' = \{((w, s), p, (w', s')) \mid ((s, \gamma), p, (s', \mathcal{C})) \in \Delta \text{ \& applying action } \mathcal{C} \text{ to } w \text{ yields } w'\}$. It was shown in [13] that pPDSs are M-equivalent to *Recursive Markov Chains* (RMCs). Since we do not explicitly use RMCs, we will not recall their formal definition.

Probabilistic 1-Counter Automata. A *probabilistic 1-counter automaton* (p1CA), A , is just a pPDS with only one stack symbol γ (other than the special bottom symbol $\perp$). In other words, it is a pPDS with $\Gamma = \{\perp, \gamma\}$. This is not the usual definition: they are typically defined as having a finite number of control states and an additional non-negative counter which can be incremented or decremented during transitions, and such that transitions can be enabled/disabled depending on whether the counter is equal to 0 or not. However, this can easily be seen to be equivalent to a pPDS with one stack symbol, γ . The stack acts as precisely a (unary) counter, and the counter is equal to 0 precisely when the top stack symbol is $\perp$.

Formally, a p1CA is usually defined in the following form, which we will find convenient. A p1CA, A , is 3-tuple $A = (S, \delta, \delta_0)$ where S is a finite set of *control states* and $\delta \subseteq S \times \mathbb{R}_{>0} \times \{-1, 0, 1\} \times S$ and $\delta_0 \subseteq S \times \mathbb{R}_{>0} \times \{0, 1\} \times S$ are *transition relations*. The transition relation δ is enabled when the counter is non-zero, and the transition relation δ_0 is enabled when it is zero. We use $p_{u,v}^{(c)}$ to denote the unique probability such that there is a transition $(u, p_{u,v}^{(c)}, c, v) \in \delta$, and likewise we use $q_{u,v}^{(c)}$ to denote the unique probability such that there is a transition $(u, q_{u,v}^{(c)}, c, v) \in \delta_0$. If such a transition exists, it is unique, and thus $p_{u,v}^{(c)} > 0$ (or $q_{u,v}^{(c)} > 0$) is uniquely determined. If such a transition doesn't exist, we may sometimes assume for convenience that $p_{u,v}^{(c)} = 0$ (or $q_{u,v}^{(c)} = 0$), even though there are no explicit 0-probability transitions provided in the input which describes A . The transition probabilities out of each control state u define a probability distribution, i.e., $\sum_{c=-1}^1 \sum_v p_{u,v}^{(c)} = 1$, and $\sum_{c=0}^1 \sum_v q_{u,v}^{(c)} = 1$. A

p1CA, A , generates a denumerable-state Markov chain $M(A) = (V', \Delta')$ with state set $V' = \{(s, d) \mid s \in S, d \in \mathbb{N}\}$, and probabilistic transition relation $\Delta' = \{((s, 0), p, (s', j)) \mid (s, p, j, s') \in \delta_0\} \cup \{((s, i), p, (s', j)) \mid i > 0, \& (s, p, c, s') \in \delta, \& j = i + c\}$. Obviously, pPDSs with only one stack symbol γ (other than $\perp$) and p1CAs (with unary counter) are M-equivalent.

A *1-counter automaton* (1CA) is just a p1CA without probabilities, i.e., the transition relation is non-deterministic. To each p1CA, $A = (S, \delta, \delta_0)$, we can associate an *underlying* 1CA, $A' = (S, \delta', \delta'_0)$, which ignores probabilities of transitions and treats them non-deterministically. Specifically, a transition $(u, c, v) \in \delta'$ ($\in \delta'_0$) iff $p_{u,v}^{(c)} > 0$, ($q_{u,v}^{(c)} > 0$, respectively). For a 1CA, $\mathcal{A} = (S, \delta, \delta_0)$, a *path* starting at state (s_1, n_1) is a sequence of states $(s_1, n_1), (s_2, n_2), \dots, (s_r, n_r)$, such that, for all $i \in \{1, \dots, r-1\}$, either $n_i > 0$ and $(s_i, n_{i+1} - n_i, s_{i+1}) \in \delta$ or $n_i = 0$ and $(s_i, n_{i+1} - n_i, s_{i+1}) \in \delta_0$. It is called a *non-zero path* if $n_i > 0$ for all $i \in \{1, \dots, r-1\}$. (Note that we allow $n_r = 0$ in non-zero paths.) Such a (non-zero) path is called a *(non-zero) terminating path* if $n_r = 0$, and if so it is said to terminate in state $(s_r, 0)$. For p1CAs, A , we define paths, non-zero paths, etc., as simply the paths, non-zero paths, etc., in the underlying 1CA. Note that for a p1CA, the probability that a particular non-zero path $(s_1, n_1), (s_2, n_2), \dots, (s_r, n_r)$ occurs, in a random walk starting at state (s_1, n_1) of the Markov chain $M(A)$ is precisely $\prod_{1 \leq i < r} p_{s_i s_{i+1}}^{(n_{i+1} - n_i)}$.

Quasi-Birth-Death Processes (QBDs). We consider discrete-time QBDs only. Of course, many analyses for continuous-time QBDs boil down to analyses of their respective embedded discrete-time chains.

A *Quasi-Birth-Death process* (QBD) is a countable state Markov chain whose transition matrix has the following block structure:²

$$\begin{bmatrix} B_0 & B_1 & 0 & 0 & 0 & \dots \\ A_{-1} & A_0 & A_1 & 0 & 0 & \dots \\ 0 & A_{-1} & A_0 & A_1 & 0 & \dots \\ 0 & 0 & A_{-1} & A_0 & A_1 & \dots \\ \dots & \dots & \dots & \dots & \dots & \dots \end{bmatrix}$$

where $B_0, B_1, A_{-1}, A_0, A_1 \in \mathbb{R}_{\geq 0}^{m \times m}$. Thus, the finite input which describes a QBD consists of the five $m \times m$ matrices: B_0, B_1, A_{-1}, A_0 , and A_1 . We can represent each state of a QBD by a pair (i, j) , where $1 \leq i \leq m$ is the index of the state within its block and $j \in \mathbb{N}$ is

²In fact, various slightly different definitions of QBDs are given in the literature, typically differing slightly on the structure of transition probabilities in the boundary cases, i.e., for the first few blocks. These differences are immaterial and these variants can be efficiently embedded in the transition structure described here, as many authors have already observed.

the index of the block. Central to many analyses for QBDs is the computation of the associated G matrix, which we will call the termination probability matrix. This is a $m \times m$ matrix, whose (i, i') entry $G_{i,i'}$ denotes the probability that, starting in state $(i, 1)$, the Markov chain will eventually visit a state in block 0, and such that the first such state it visits is $(i', 0)$. As is well known (e.g., [25]), G is the least non-negative solution to the matrix equation $X = A_{-1} + A_0 X + A_1 X^2$, i.e., for any non-negative solution matrix G' , we have $G \leq G'$ (entry-wise inequality). Other key matrices, which are also central to computations for QBDs, can be derived from the matrix G . Specifically, the R matrix, has $R_{i,i'}$ equal to the expected number of visits to state $(i', n+1)$, starting from state (i, n) , before returning to a state in a block $\leq n$. The matrix U (the “taboo probability” matrix) has $U_{i,i'}$ equal to the probability that starting from state $(i, 1)$ the chain does not visit a state in block 0 until it eventually revisits a state in block 1, and it does so in state $(i', 1)$. The matrices U and R can be obtained from G : $U = A_0 + A_1 G$, and $R = A_{-1}(I - U)^{-1}$. (Of course, the approximate solution of G will introduce errors in the solutions for U and R .) If the QBD is positive recurrent, these matrices can be used to compute steady state probabilities for being in any given state (i, j) (see, e.g., [23]).

Tree-Like and Tree-Structured QBDs. Several slight variants of TL-QBDs (and TS-QBDs) have appeared in the literature. We used the most restrictive definition of TL-QBDs in order to have the strongest results about the equivalence of all these models. Consider the infinite rooted d -ary tree T_d , label every edge with a symbol in $\Gamma = \{1, \dots, d\}$, and label every node with the string $w \in \Gamma^*$ corresponding to the path from the root; the root is labelled with the empty string ϵ . The states of TS-QBDs and TL-QBDs consist of pairs (w, i) , where $w \in \Gamma^*$ is (the label of) a node of the tree T_d and $i \in \{1, \dots, m\}$ acts as a “control state”. The transitions of a TS-QBD are as follows. From a state (ϵ, i) , $i \in \{1, \dots, m\}$, there is a transition to state:

1. (ϵ, j) with probability $f^{i,j}$, where $j \in \{1, \dots, m\}$.
2. (s, j) with probability $u_s^{i,j}$, where $s \in \Gamma$, and $j \in \{1, \dots, m\}$.

From any state (wk, i) , where $w \in \Gamma^*$ and $k \in \Gamma$, and $i \in \{1, \dots, m\}$, there is a transition to state:

3. (w, j) with probability $d_k^{i,j}$.
4. (ws, j) , where $s \in \Gamma$, with probability $a_{k,s}^{i,j}$.
5. (wks, j) , where $s \in \Gamma$, with probability $u_s^{i,j}$.

A TS-QBD can thus be described by a finite collection of $m \times m$ matrices (specifically, $d^2 + 2d + 1$ such matrices)

with rational entries, namely the matrices $D_k, A_{k,s}, U_s$, and F , where $k, s \in \Gamma$, and where their (i, j) entry is $d_k^{i,j}, a_{k,s}^{i,j}, u_s^{i,j}$, and $f^{i,j}$, respectively.

TL-QBDs are defined by restricting TS-QBDs: TL-QBDs are TS-QBDs with the additional requirement that if $k \neq s$, then $A_{k,s} = 0$ (i.e., the zero matrix), and secondly that $A_{k,k} = A_{s,s}$ for all $k, s \in \Gamma$.

3 Equivalences and basic consequences

Proposition 1 *QBDs and p1CAs are M-equivalent.*

The proof is very simple (see [12]). Briefly, the block index of a state of a QBD corresponds to the counter value of a state of the p1CA, the B matrices of a QBD correspond to the probabilities q of transitions on 0 counter and the A matrices correspond to the probabilities p of transitions on nonzero counter.

Theorem 2 *pPDSs, RMCs, TL-QBDs, and TS-QBDs are all M-equivalent.*

Obviously TL-QBDs are a special case of TS-QBDs. Furthermore, TS-QBDs are themselves a special case of pPDSs (equivalently, RMCs [13]), where transitions are constrained as follows: (1) If $((s, \gamma), p_{(s, \gamma), (s', C)}, (s', C)) \in \Delta$, where $C = \text{swap}\&\text{push}(\gamma', \gamma'')$, we must have $\gamma = \gamma'$, i.e., every “swap and push” operation must be just a “push”, and (2) for all $\gamma, \gamma' \in \Gamma$, we must have $p_{(s, \gamma), (s', \text{swap}\&\text{push}(\gamma, \gamma''))} = p_{(s, \gamma'), (s', \text{swap}\&\text{push}(\gamma', \gamma''))}$, i.e., the probability of the “push” does not depend on the top stack symbol. The argument that pPDS can be efficiently embedded in TL-QBDs is a bit more involved (see [12]). Briefly, we can use auxiliary control states in TL-QBDs in order to mimic, e.g., a pPDS’s *swap&push* operation via a sequence of transitions in a TL-QBD.

Thus all the known results for pPDSs and RMCs apply to TL-QBDs, and vice versa. The following corollary highlights a few results for TL-QBDs (and TS-QBDs) that follow from work on pPDSs and RMCs. The *square-root sum problem* asks, given natural numbers $(d_1, \dots, d_n) \in \mathbb{N}^n$ and $k \in \mathbb{N}$, whether $(\sum_i \sqrt{d_i}) \geq k$. This decision problem is contained in PSPACE, but its containment even in NP is a longstanding open problem first posed in the 1970s ([16]), with many applications. See ([13]) for more background.

Corollary 3 1. ([14]) *The quantitative model checking problem for QBDs and TL-QBDs, against linear-time ω -regular temporal properties, is decidable in PSPACE.*

2. ([13, 15]) *The square-root sum problem is polynomial time reducible to the problem of approximating the termination probability (the analog of the*

G matrix) for TL-QBDs, even to within just one bit of precision (or even to within any constant additive factor $c < 1/2$). Furthermore, even deciding whether a termination probability for a TL-QBD is 1 is sqrt-sum-hard.

3. ([19]) *There are TL-QBDs for which at least exponentially many iterations of the (decomposed) Newton’s method ([13]), applied to the non-linear equations for termination probabilities are needed as a function of the TL-QBD’s encoding size, to even converge to within just one bit of precision of a termination probability.*

The following is not a corollary of earlier results. It is proved in [12].

Theorem 4 *The square-root sum problem is polynomial time reducible to the following problem: given a p1CA (QBD) with control states u and v , and given a rational value p decide whether $G_{u,v} \leq p$.*

4 Structural properties of (p)1CAs

This section develops crucial structural properties of (probabilistic) 1-Counter Automata, used in section 5 to establish strong results on the performance of (decomposed) Newton’s method for QBDs. Let $\text{mp}(s, s')$ ($\text{mp}_{\text{n-z}}(s, s')$) denote the length of the shortest (non-zero, respectively) terminating path starting at state $(s, 1)$ and terminating at state $(s', 0)$. If there is no such (non-zero) terminating path, then by definition $\text{mp}(s, s') = \infty$ ($\text{mp}_{\text{n-z}}(s, s') = \infty$, respectively). By convention, a path with a single state has length 0. The next lemma (proved in [12]) shows that in 1CAs whenever a terminating path exists, a “short” (polynomial length) such path also exists.

Lemma 5 *Suppose $\mathcal{A} = (S, \delta, \delta_0)$ is a 1CA where $|S| = k$. For any pair of control states $s, s' \in S$, either $\text{mp}_{\text{n-z}}(s, s') = \infty$ or else $\text{mp}_{\text{n-z}}(s, s') \leq k^3$. Likewise, either $\text{mp}(s, s') = \infty$, or else $\text{mp}(s, s') \leq k^4$.*

Corollary 6 *Let $A = (S, \delta, \delta_0)$ be a p1CA where $|S| = k$, and let $p_{\min} > 0$ be the smallest positive probability on any transition of A . For any pair of states $s, s' \in S$, either $G_{s,s'} = 0$ or $G_{s,s'} \geq p_{\min}^{k^3}$.*

Indeed, $G_{s,s'} > 0$ iff there is a non-zero terminating path starting at $(s, 1)$ and terminating at $(s', 0)$. By Lemma 5, the length of the shortest such path is $\leq k^3$. Therefore its probability is at least $p_{\min}^{k^3}$.

The termination probabilities $G_{u,v}, u, v \in S$ are the smallest nonnegative solution to the following fixed point equation system

$$x_{uv} = p_{uv}^{(-1)} + \left(\sum_{w \in S} p_{uw}^{(0)} x_{wv} \right) + \sum_{y \in S} p_{uy}^{(1)} \sum_{z \in S} x_{yz} x_{zv} \quad (1)$$

We can clean up this system of equations for G by removing the variables x_{uv} for which $G_{u,v} = 0$. This can be done in polynomial-time, even for more general fixed point equations associated with pPDSs and RMCs (see [13]). (After clean-up, the equations may no longer have the simple matrix form.) Henceforth, we consider only cleaned-up equation systems, where only non-zero variables remain.

Based on this equation system we can build a dependency graph, $D = (\tilde{X}, E)$, whose nodes are all non-zero variables $\tilde{X} = \{x_{uv} : u, v \in S \text{ and } G_{u,v} \neq 0\}$ and there is an edge $(x_{uv}, x_{st}) \in E$ iff x_{st} occurs on the rhs of the equation $x_{uv} = \alpha$ corresponding to x_{uv} . We decompose this graph into strongly connected components (SCCs) and sort them topologically. As a result we obtain a sequence of SCCs $X_1, X_2, \dots, X_m$ such that there can exist a path in graph D from variable $x \in X_i$ to variable $x' \in X_j$ only if $i \geq j$. We will write $x_{st} \equiv x_{uv}$ iff $s = u$ and $t = v$. We say a variable x_{uv} *depends* on the value of a variable x_{st} iff either $x_{st} \equiv x_{uv}$, or there is a path from x_{uv} to x_{st} in the graph D . Of course this relation is transitive. We say that an equation $x_{uv} = \alpha$ is *non-linear in a set X' of variables* if, by removing all variables that are not in X' from monomials in α , we are left with an expression α' that is non-linear. We say that SCC X_i is nonlinear if the equation $x_{uv} = \alpha$ of some variable $x_{uv} \in X_i$ is nonlinear in X_i .

Theorem 7 *If the clean equation $x_{uv} = \alpha$, for a variable $x_{uv} \in X_i$ is non-linear in X_i , and if the clean equation for a variable $x_{st} \in X_j$ is non-linear in X_j , and there is a path from x_{uv} to x_{st} in dependency graph D , then there is a path from x_{st} to x_{uv} in D .*

The proof is long and is given in [12]. With the dependency graph D , we associate in the standard way a directed acyclic graph (DAG), H , whose nodes are SCCs of D , and which has an edge from SCC X_i to X_j iff there is an edge in D from some variable in X_i to some variable in X_j . The following is an easy corollary.

Corollary 8 *In the DAG, H , along any directed path of SCCs there is at most one nonlinear SCC.*

5 New upper bounds on Newton's method for QBDs

We will now exploit the structural results about p1CAs established in section 4, to establish strong new upper bounds on the performance of (decomposed) Newton's method on QBDs. In our analysis in this section, we assume a unit-cost exact rational arithmetic RAM model of computation. In other words, individual arithmetic operations on rationals have unit cost, regardless of the size of the rationals.

Recall that in (multi-variate) Newton's method, we are given a suitably differentiable map $F : \mathbb{R}^n \mapsto \mathbb{R}^n$, and we wish to find a solution to the system of equations $F(\mathbf{x}) = \mathbf{0}$. Starting at some $\mathbf{x}^0 \in \mathbb{R}^n$, the method works by iterating $\mathbf{x}^{k+1} := \mathbf{x}^k - (F'(\mathbf{x}^k))^{-1}F(\mathbf{x}^k)$, where $F'(\mathbf{c})$ is the *Jacobian matrix* of partial derivatives, whose (i, j) entry is $\frac{\partial F_i}{\partial x_j}$ evaluated at $\mathbf{c}$.

In the setting of p1CAs, we have a system of n equations in n variables, $x_i = P_i(\mathbf{x})$, which we can denote by $\mathbf{x} = P(\mathbf{x})$. Thus, we wish to find a solution to $F(\mathbf{x}) \doteq P(\mathbf{x}) - \mathbf{x} = \mathbf{0}$. Note that these are polynomial functions, and thus certainly differentiable.

We shall solve this system of equations using the *decomposed Newton's method* of [13], which applies more generally not just to systems $\mathbf{x} = P(\mathbf{x})$ arising for p1CAs, but to any monotone system $\mathbf{x} = P(\mathbf{x})$ of polynomial equations (i.e., where the coefficients in $P(\mathbf{x})$ are non-negative) which has a non-negative solution. Specifically, for any such system $\mathbf{x} = P(\mathbf{x})$ which has been *cleaned up* (i.e., variables which are necessarily zero in any least solution have been removed, something which can be done easily in polynomial time [13]) we form the dependency graph D for the non-zero variables in the corresponding cleaned system of equations, we decompose D into SCCs, and form the DAG of SCCs, H . We then “solve” for the values of variables in each SCC of H , “bottom up” by applying Newton's method starting at the vector 0 to the equations for each SCC, beginning with bottom SCCs. Once one SCC is “solved” the values computed for the variables in that SCC are plugged into equations in higher SCCs that depend on those values.

Of course, since values may in general be irrational and are only converged to in the limit, we have to specify more carefully what we mean by “solve” an SCC. This is where we make crucial use of the special structure of SCCs in the case p1CAs and QBDs. By Corollary 8, for any non-linear SCC, X_i , it must be the case that any other SCC, X_j , for which there is a path in H from X_i to X_j , is linear, i.e., any variable $x_{uv} \in X_j$ has a corresponding clean equation $x_{uv} = \alpha$ which is linear in the variables of X_j , assuming variables in even lower SCCs have been assigned fixed values. It was shown in [13] (in the more general setting of monotone systems arising from RMCs and pPDSs) that for such linear SCCs, X_j , Newton's method converges in just one iteration, starting at the vector 0, to the exact rational least fixed point (LFP) solution we are after (i.e., to the values $G_{u,v}$ for these variables in $x_{uv} \in X_j$). Thus, in a bottom up fashion we can compute the exact solutions $G_{u,v}$ for those variables x_{uv} which are in linear SCCs below any nonlinear SCC. After computing these values we plug them into equations for variables in higher

SCCs that depend on them, and we eliminate the linear SCC which was already solved. We do this until there are no bottom linear SCCs remaining.

We next have to apply Newton's method to non-linear SCCs, which can have irrational solutions which are only converged to in the limit. How many iterations are "enough"? For this, we will use the following recent result by Esparza et. al. (Theorem 3.2 of [9]) on the behavior of Newton's method on precisely such monotone nonlinear systems. Let $P(X)$ be a cleaned monotone system of polynomials (i.e., $P(X)$ consists of n multi-variate polynomials, P_i , $i = 1, \dots, n$, in the variables $X = x_1, \dots, x_n$), such that $X = P(X)$ has a non-negative solution, and since it is cleaned, only positive solutions, and therefore a least fixed point (LFP) solution, $q^* > 0$. A vector q' is said to have i valid bits of q^* if $|q_j^* - q'_j|/q_j^* \leq 2^{-i}$ for every $1 \leq j \leq n$.

Theorem 9 ([9]) *Let $P(X)$ be a cleaned strongly connected monotone system of quadratic polynomials (i.e., $P(X)$ consists of n quadratic multi-variate polynomials in n variables). Let $c_{\min}$ be the smallest nonzero coefficient of any monomial in $P(X)$, and let $\mu_{\min}$ and $\mu_{\max}$ be the minimal and maximal components of the LFP vector $q^* > 0$, respectively. Let $k_f = n \cdot \log(\frac{\mu_{\max}}{c_{\min} \cdot \mu_{\min} \cdot \min\{\mu_{\min}, 1\}})$. Let $\mathbf{x}^j$ denote the vector of values obtained after j iterations of Newton's method on the system $F(X) = P(X) - X$, starting with the initial all 0 vector, $\mathbf{x}^0 = \mathbf{0}$. Then for every $i \geq 0$, $\mathbf{x}^{(\lceil k_f \rceil + i)}$ has i valid bits of q^* .*

For a given p1CA, we hereafter use m to denote the maximum number of bits required to encode the integer numerators and denominators of transition probabilities of the p1CA. Thus, in particular, the smallest non-zero transition probability is $p_{\min} \geq 1/2^m$. Using Theorem 9, the following can be shown (see [12]).

Theorem 10 *Let $P(X)$ be the cleaned strongly connected monotone system of quadratic polynomials associated with a nonlinear SCC, X_i , of the decomposed system of equations associated with a p1CA, and where the exact rational values $G_{u,v}$ associated with variables x_{uv} in already solved "lower" linear SCCs have been substituted for x_{uv} on the right hand side of equations for variables in X_i . Suppose that the p1CA has n control states, and thus $|X_i| \leq n^2$, and let $G|_{X_i}$ denote those entries $G_{u,v}$ of the matrix G , such that $x_{uv} \in X_i$. Then, starting with $\mathbf{x}^0 := \mathbf{0}$, for every $i \geq 0$, the Newton iteration $\mathbf{x}^{(4mn^5 + mn^2 + i)}$ has i valid bits of $G|_{X_i}$.*

Theorem 10 implies that we can compute i bits of the values $G_{u,v}$ for variables x_{uv} in non-linear SCCs of the system $X = P(X)$ associated with a p1CA (QBD),

using only a number of iterations of Newton's method which is polynomially bounded in the size of the p1CA, and linearly bounded in i .

We now have to confront a major difficulty: there may be other, linear, SCCs, X_r , which are "above" such non-linear SCCs in H . Specifically, there may be a linear SCC X_r , from which there is a path in H to a non-linear SCC, X_i . In order to be able to (approximately!) compute $G_{u,v}$ for variables $x_{uv} \in X_r$, we have to first approximately compute the (possibly irrational) values $G_{u',v'}$, for $x_{u'v'} \in X_i$, and substitute this value in occurrences of $x_{u'v'}$ in equations for higher linear SCCs. The question arises: how many bits of precision i , do we need to compute $G_{u',v'}$ to in order to compute $G_{u,v}$ to within i bits of precision? To answer this, we employ a classic bound, based on condition numbers, on errors in the solution of a linear systems.

Theorem 11 (see, e.g., [18], Chap 2.1.2, Thm 3.³) *Consider a system of linear equations, $Bx = b$, where $B \in \mathbb{R}^{n \times n}$ and $b \in \mathbb{R}^n$. Suppose B is non-singular, and $b \neq 0$. Let $x^* = B^{-1}b$ be the unique solution to this linear system, and suppose $x^* \neq 0$. Let $\|\cdot\|$ denote any vector norm and associated matrix norm (when applied to vectors and matrices, respectively). Let $\text{cond}(B) = \|B\| \cdot \|B^{-1}\|$ denote the condition number of B . Let $\varepsilon, \varepsilon' > 0$, be values such that $\varepsilon' < 1$, and $\varepsilon \cdot \text{cond}(B) \leq \varepsilon'/4$. Let $\mathcal{E} \in \mathbb{R}^{n \times n}$ and $\zeta \in \mathbb{R}^n$, be such that $\frac{\|\mathcal{E}\|}{\|B\|} \leq \varepsilon$, $\frac{\|\zeta\|}{\|b\|} \leq \varepsilon$, and $\|\mathcal{E}\| < 1/\|B^{-1}\|$. Then the system of linear equations $(B + \mathcal{E})x = b + \zeta$ has a unique solution x_ε^* such that:*

$$\frac{\|x_\varepsilon^* - x^*\|}{\|x^*\|} \leq \varepsilon'$$

We will apply this theorem using the l_∞ vector norm and induced matrix norm (maximum absolute row sum): $\|x\|_\infty = \max_i |x_i|$ and $\|A\|_\infty = \max_i \sum_j |a_{ij}|$.

Suppose that the fixed point equation system for a linear SCC of a p1CA, which lives "above" some non-linear SCCs in the DAG H , looks like this: $x = Ax + b$. We know that $A \geq 0$ is an irreducible matrix (precisely because the variables being solved for are in the same SCC), $b \geq 0$, and $b \neq 0$ since otherwise the unique solution for this system would be $q^* = 0$, and zero variables were already eliminated. We can of course rewrite this linear equation as $(I - A)x = b$. It follows from a more general result in [13] about the decomposed systems of equations arising for RMCs (pPDSs) (specifically, see Lemma 17 and Theorem 14 of [13]), that $\rho(A) < 1$, where $\rho(A)$ denotes the spectral radius of A , and that therefore $(I - A)$ is non-singular, and furthermore $(I - A)^{-1} = (\sum_{i=0}^{\infty} A^i)$. Thus the LFP of

³Our statement is weaker, but derivable from that theorem.

this equation system is $q^* = (I - A)^{-1}b = (\sum_{k=0}^{\infty} A^k)b$. To prove bounds on errors in “higher” linear SCCs, when values in nonlinear SCCs are approximated, we will need the following two lemmas (proved in [12]):

Lemma 12 *Let $A \in \mathbb{R}_{\geq 0}^{n \times n}$ and $b \in \mathbb{R}_{\geq 0}^n$, such that: $(I - A)^{-1} = \sum_{k=0}^{\infty} A^k$, and $(\sum_{k=0}^{\infty} A^k)b \leq \mathbf{1}$, and A is an irreducible non-negative matrix whose smallest non-zero entry is $c > 0$, and $b \neq 0$ and $p > 0$ is the largest entry of b . Then: $\|\sum_{k=0}^{\infty} A^k\|_{\infty} \leq \frac{n}{pc^n}$.*

Lemma 13 *Let X_r be a linear SCC of the cleaned equation system for a p1CA, whose corresponding linear equation system is $x = Ax + b$, after variables x_{uv} in lower SCCs have been substituted by their exact (possibly irrational) values $G_{u,v}$. Let $p_{\min}$ denote the smallest positive probability on any transition of the p1CA, and let n be its number of control states. Then the following bounds hold:*

1. $\frac{1}{2^{2mn^3+m}} \leq p_{\min}^{2n^3+1} \leq \|(I - A)\|_{\infty} \leq n + 1$
2. $\|(I - A)^{-1}\|_{\infty} \leq \frac{n^2}{p_{\min}^{5n^5}} \leq n^2 \cdot 2^{5mn^5}$
3. $\text{cond}(I - A) \leq \frac{2n^3}{p_{\min}^{5n^5}} \leq 2n^3 \cdot 2^{5mn^5}$
4. $\|b\|_{\infty} \geq p_{\min}^{2n^3+1} \geq \frac{1}{2^{2mn^3+m}}$

For a “higher” linear SCC, X_r , i.e., one which can reach some non-linear SCC in H , let us define its *height*, $h_r < \infty$, to be the maximum finite distance in H between X_r and some lower non-linear SCC that it can reach. Let $h_{\max} = \max_r h_r$, where the maximum is taken over all linear SCCs that can reach a non-linear SCC. Note that, as a very loose upper bound, certainly $h_{\max} \leq n^2$, where $n = |S|$ is the number of control states of the p1CA, because there are at most n^2 variables in the entire system. Now consider the decomposed Newton’s method applied to the fixed point equations for a p1CA, with the following specification for the number of iterations to be applied to each SCC:

1. Use, one iteration of Newton’s method (starting at vector $x^0 = 0$), or any linear system solving method, to solve a remaining bottom linear SCC exactly. Remove the linear SCC, and plug the corresponding values of variables into equations for higher SCCs. Do this until only non-linear bottom SCCs remain, or all SCCs are solved.
2. For each remaining non-linear SCC, apply Newton’s method (starting with vector $x^0 = 0$) to the non-linear equations for these SCCs, using the following number of iterations:

$$4mn^5 + mn^2 + h_{\max}(9mn^5 + 4) + i$$

Afterwards, plug the resulting (approximate) values for variables in each such non-linear SCC into the equations for higher (linear) SCCs.

3. For each remaining linear SCC, use one iteration of Newton’s method (or any other linear system solution method) to solve for the exact (unique) solution of the corresponding linear system (note that the coefficients of these equations will have errors because of the approximations below, but we still seek their exact solution), then remove the linear SCC, and plug these values into higher (linear) SCCs that remain, until no SCCs remain.

Theorem 14 *Given a p1CA (or, equivalently, a QBD), the above algorithm, based on (a decomposed) Newton’s method, approximates every entry of the matrix G of termination probabilities for the p1CA (QBD) to within i bits of precision (i.e., to within additive error $1/2^i$). In the unit-cost arithmetic RAM model of computation (i.e., rational Blum-Shub-Smale model), the algorithm has a running time which is polynomial in both the encoding size of the p1CA (QBD) and in i .*

The proof is in [12]. We briefly sketch the argument. It proceeds by induction on the height, h , of a given “higher” linear SCC, X_r , above non-linear SCCs, to show that for every variable $x_{uv} \in X_r$ we compute $G_{u,v}$ to within $W_h = (h_{\max} - h)(9mn^5 + 4) + i$ bits of precision. The base case $h = 0$ follows from the fact that, applying Theorem 10, all non-linear SCCs are approximated to within $W_0 = h_{\max}(9mn^5 + 4) + i$ bits of precision, and all “lower” linear SCCs are computed exactly. The inductive step shows, by using Theorem 11 and Lemma 13, that at each height $h \geq 1$, solution of the linear SCCs at that height can “lose for us” at most another $9mn^5 + 4$ bits of precision. We emphasize that these (impractical) upper bounds for the number of iterations are very coarse, and are only intended to facilitate our proof that polynomially many iterations of Newton’s method suffice. A more detailed analysis would likely yield polynomial bounds with much smaller exponents as the required number of iterations.

An important remaining open question is this: can polynomial upper bounds for approximating the G matrix for QBDs be established in the standard Turing model of computation, rather than in the unit-cost rational arithmetic RAM model as we have done? A possible approach for doing this is via a detailed analysis of the effect of round-off errors on iterations of Newton’s method over the nonlinear equations for QBDs.

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