Analysis of the Propagation Time of a Rumour in Large-scale Distributed Systems

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Abstract—The context of this work is the well studied dissemination of information in large scale distributed networks through pairwise interactions. This problem, originally called rumor mongering, and then rumor spreading, has mainly been investigated in the synchronous model. This model relies on the assumption that all the nodes of the network act in synchrony, that is, at each round of the protocol, each node is allowed to contact a random neighbor. In this paper, we drop this assumption under the argument that it is not realistic in large scale systems. We thus consider the asynchronous variant, where at time unit, the rumor is placed on one of the vertices of a given network, and this rumor is propagated to all the vertices of the network through pairwise interactions between vertices. One of the important questions of these protocols is the spreading time, that is time it needs for the rumor to be known by all the vertices of the network.

Several models have been considered to answer this question. The most studied one is the synchronous push/pull model, also called the synchronous random phone call model. This model assumes that all the vertices of the network act in synchrony, which allows the algorithms designed in this model to divide time in synchronized rounds. During each synchronized round, each vertex of the network selects at random one of its neighbor and either sends to it the rumor if it knows it (push operation) or gets the rumor from it if it knows the rumor (pull operation). In the synchronous model, the spread time of a rumor is defined as the number of synchronous rounds necessary for all the nodes to know the rumor. In one of the first papers dealing with the push operation only, Frieze and Grimmet [16] proved that if the underlying graph is complete, then asymptotically almost surely the number of rounds is \(\frac{\log(n)}{2} + \frac{\log(\log(n))}{\log(n)} + o(\log(n))\) where \(n\) is the number of nodes of the graph. Further results have been established (see for example [21], [7] and the references herein), the most recent ones resulting from the observation that the rumor spreading time is closely related to the conductance of the graph of the network [17], [18]. Investigations have also been done in different topologies of the network [9], [11], [14], [25], in the presence of link or vertices failures (see [13]), and dynamic graphs [10].

All the above studies assume that all vertices of the network act synchronously. In distributed networks, and in particular in large scale distributed systems, such a strict synchronization is unrealistic. Several authors have recently dropped this assumption by considering an asynchronous model. Boyd et al [28] consider that each node has a clock that goes off at the time of a rate 1 Poisson process. Each time the ring of a node goes off, the push or pull operations are triggered according to the knowledge of the node. Acan et al [1] go a step further by studying rumor spreading time for any graph topology. They show that both the average and guaranteed spreading time are \(\Omega(n)\), where \(n\) is the number of nodes in the network. Further investigations have been made for different network topologies [26], [15].

I. INTRODUCTION

Randomized rumor spreading is an important mechanism that allows the dissemination of information in large and complex networks through pairwise interactions. This mechanism initially proposed by Deemers et al [12] for the update of a database replicated at different sites, has then been adopted in many applications ranging from resource discovery [19], data-aggregation [22], complex distributed applications [8], or virus propagation in computer networks [6], to mention just a few.

A lot of attention has been devoted to the design and study of randomized rumor spreading algorithms. Initially, some rumor is placed on one of the vertices of a given network,
a) Our contributions: In this paper we consider the population protocol model, which turns out to resemble to the discrete-time version of the asynchronous spreading model. This model provides minimalist assumptions on the computational power of the nodes: nodes are finite-state automata, identically programmed, they have no identity, they do not know how numerous they are, and they progress in their computation through random pairwise interactions. Their objective is to ultimately converge to a state from which the sought property can be derived from any node [5]. In this model, the spreading time is defined as the number of interactions needed for all the nodes of the network to learn the rumor. Angluin et al [3] analyze the spreading time of a rumor by only considering the push operation (which they call the one-way epidemic operation), and show that with high probability, a rumor injected at some node requires $O(n \log n)$ interactions to be spread to all the nodes of the network.

In the present paper we go a step further by considering a more general problem namely, that is all the nodes of the network initially receive an input value, and the objective for each node is to learn the maximal value initially received by any node. Note that the rumor spreading problem is a particular instance of this problem when two input values 1 and 0 are considered respectively representing the knowledge and the absence of knowledge of the rumor. We present a thorough analysis of the number of interactions needed for all the nodes to converge to the correct response. Specifically, we study the expectation, variance and an exact formulation of the distribution of the number of interactions needed to propagate a rumor.

This formulation being hardly usable in practice once $n$ becomes too large, a tight bound is derived. This bound is all the more interesting as usual probabilistic inequalities fail to provide relevant results in this case. Finally, we study the asymptotic behavior of the spreading time when the size of the network tends to infinity.

b) Road map: The remainder of this paper is organized as follows. Section II presents the population protocol model. Section III specifies the problem addressed in this work. Analysis of the spreading time is proposed in Section IV, while we study in Section V its asymptotic behavior. We have simulated our protocol to illustrate our theoretical analysis. Finally, Section VI concludes.

II. Population protocols model

In this section, we present the population protocol model, introduced by Angluin et al. [2]. This model describes the behavior of a collection of nodes that interact pairwise. The following definition is from Angluin et al [4]. A population protocol is characterized by a 6-tuple $(Q, \Sigma, Y, \iota, \omega, f)$, over a complete interaction graph linking the set of $n$ nodes, where $Q$ is a finite set of states, $\Sigma$ is a finite set of input symbols, $Y$ is a finite set of output symbols, $\iota : \Sigma \to Q$ is the input function that determines the initial state of a node, $\omega : Q \to Y$ is the output function that determines the output symbol of a node, and $f : Q \times Q \to Q \times Q$ is the transition function that describes how two nodes interact and update theirs states. Initially all the nodes start with a initial symbol from $\Sigma$, and upon interactions with nodes update their state according to the transition function $f$. Interactions between nodes are orchestrated by a random scheduler: at each discrete time, any two nodes are randomly chosen to interact with a given distribution. Note that its is assumed that the random scheduler is fair, which means that the interactions distribution is such that any possible interaction cannot be avoided forever. The notion of time in population protocols refers to as the successive steps at which interactions occur, while the parallel time refers to as the successive number of steps each node executes [5]. Nodes do not maintain nor use identifiers (nodes are anonymous and cannot determine whether any two interactions have occurred with the same agents or not). However, for ease of presentation the nodes are numbered $1, 2, \ldots, n$. We denote by $C^{(i)}_t$ the state of node $i$ at time $t$. The stochastic process $C = \{C_t, t \geq 0\}$, where $C_t = (C^{(1)}_t, \ldots, C^{(n)}_t)$, represents the evolution of the population protocol. The state space of $C$ is thus $Q^n$ and a state of this process is also called a protocol configuration.

III. Spreading the maximum

We consider in this section the following problem. Each site has initially an integer value. At each discrete instant of time, two distinct nodes are successively chosen and they change their value with the maximum value of each node. More precisely, for all nodes $a$ and $b$, with $a \neq b$, we consider the function $f$ given by

$$f(a, b) = (\max\{a, b\}, \max\{a, b\}).$$

We want to evaluate the time needed so that all the nodes get the same value.

Let $C = \{C_t, t \geq 0\}$ be a discrete-time stochastic process with state space $S = \mathbb{N}^n$. For every $t \geq 0$, the state at time $t$ of the process is denoted by $C_t = (C^{(1)}_t, \ldots, C^{(n)}_t)$, where $C^{(i)}_t$ is the integer value of node $i$ at time $t$. At each instant $t$, two distinct indexes $i$ and $j$ are successively chosen among the set of nodes $1, \ldots, n$ randomly. We denote by $X_t$ the random variable representing this choice and we suppose that this choice is uniform, i.e we suppose that

$$\mathbb{P}\{X_t = (i, j)\} = \frac{1}{n(n-1)} 1_{i \neq j}. $$

Once the couple $(i, j)$ is chosen at time $t$, the process reaches state $C_{t+1}$, at time $t+1$, given by

$$C^{(i)}_{t+1} = C^{(j)}_{t+1} = \max\{C^{(i)}_t, C^{(j)}_t\}$$

and $C^{(m)}_{t+1} = C^{(m)}_t$ for $i \neq j$.

We denote by $M$ the maximum initial value among all the nodes, i.e. $M = \max\{C^{(1)}_0, \ldots, C^{(n)}_0\}$. It is easily checked that for all $t \geq 0$, we have $M = \max\{C^{(1)}_t, \ldots, C^{(n)}_t\}$.

We consider the random variable $T_n$ defined by

$$T_n = \inf\{t \geq 0 \mid C^{(i)}_t = M, \text{ for every } 1, \ldots, n\}.$$

The random variable $T_n$ represents the number of interactions needed for all the nodes in the network to know the maximal value $M$.
We introduce the discrete-time stochastic process $Y = \{Y_t, t \geq 0\}$ with state space $\{1, \ldots, n\}$ defined, for all $t \geq 0$, by

$$Y_t = \left\{ i \mid C(t)^i = M \right\}. $$

The random variable $Y_t$ represents the number of nodes knowing the maximum value $M$ at time $t$. The stochastic process $Y$ is then a homogeneous Markov chain with transition probability matrix $A$. The non zero transition probabilities are given, for $i, j = 1, \ldots, n$, by

$$
\begin{align*}
A_{i,i} &= 1 - \frac{2i(n-i)}{n(n-1)}, \\
A_{i,i+1} &= \frac{2i(n-i)}{n(n-1)}, \text{ for } i \neq n.
\end{align*}
$$

Indeed, when $Y_t = i$, in order to get $Y_{t+1} = i+1$, either the first node must be chosen among the ones with the maximum value (probability $i/n$) and the second agent must be chosen among the ones with the non maximum value (probability $(n-i)/(n-1)$) or the first node must be chosen among the ones with the non maximum value (probability $(n-i)/n$) and the second node must be chosen among the ones with the non maximum value (probability $i/(n-1)$).

The states $1, \ldots, n-1$ of $Y$ are transient and state $n$ is absorbing. The random variable $T_n$ can then be written as

$$T_n = \inf \{ t \geq 0 \mid Y_t = n \}. $$

It is well-known, see for instance [27], that the distribution of $T_n$ is given, for every $k \geq 0$, by

$$\mathbb{P} \{ T_n > k \} = \alpha Q^k 1, \quad (1)$$

where $\alpha$ is the row vector containing the initial probabilities of states $1, \ldots, n-1$, that is $\alpha_i = \mathbb{P} \{ Y_0 = i \}$, $Q$ is the sub-matrix obtained from $A$ by deleting the row and the column corresponding to absorbing state $n$ and $1$ is the column vector of dimension $n-1$ with all its entries equal to 1.

For $i = 0, \ldots, n$, we introduce the notation

$$p_i = \frac{2i(n-i)}{n(n-1)}$$

and we denote by $H_k$ the harmonic series defined by $H_0 = 0$ and $H_k = \sum_{\ell=1}^{k} 1/\ell$, for $k \geq 1$. Note that, for every $i = 0, \ldots, n$, we have $p_i = p_{n-i}$.

If we denote by $S_i$, for $i = 1, \ldots, n-1$, the total time spent by the Markov chain $Y$ in state $i$, then conditionally on the event $Y_0 = i$, $S_i$ has a geometric distribution with parameter $p_i$, for $i = 1, \ldots, n-1$ and in this case, we have $T_n = S_1 + \cdots + S_{n-1}$. It follows that

$$\mathbb{P} \{ T_n > k \mid Y_0 = i \} = \mathbb{P} \{ S_i + \cdots + S_{n-1} > k \},$$

which means that $\mathbb{P} \{ T_n > k \mid Y_0 = i \}$ is decreasing with $i$ and in particular that

$$\mathbb{P} \{ T_n > k \mid Y_0 = i \} \leq \mathbb{P} \{ T_n > k \mid Y_0 = 1 \}. \quad (2)$$

IV. ANALYSIS OF THE SPREADING TIME

In the following we study the expectation and the variance of $T_n$, the number of interactions needed for all the nodes in the network to know the maximal value $M$. We then provide an explicit expression of the distribution of $T_n$, and then a bound and an equivalent for the explicit distribution of $T_n$.

A. Expectation and variance of $T_n$

The mean time $\mathbb{E}(T_n)$ needed so that all the nodes get the same value is then given by

$$\mathbb{E}(T_n) = \alpha (I - Q)^{-1} 1, \quad (3)$$

where $I$ is the identity matrix. This expectation can also be written as

$$\mathbb{E}(T_n) = \sum_{i=1}^{n-1} \alpha_i \mathbb{E}(T_n \mid Y_0 = i).$$

This conditional expectations are given by the following theorem.

**Theorem 1:** For every $n \geq 1$ and $i = 1, \ldots, n$, we have

$$\mathbb{E}(T_n \mid Y_0 = i) = \frac{(n-1)(H_{n-1} + H_{n-i} - H_{i-1})}{2}. $$

**Proof:** If $Y_0 = n$, which means that all the nodes start with same values, then we have $T_n = 0$ and so $\mathbb{E}(T_n \mid Y_0 = n) = 0$. For $i = 1, \ldots, n-1$ we have

$$\mathbb{E}(T_n \mid Y_0 = i) = \sum_{\ell=i}^{n-1} \mathbb{E}(S_{\ell}) = \sum_{\ell=i}^{n-1} \frac{1}{\ell} = \frac{n(n-1)}{2} \sum_{\ell=i}^{n-1} \frac{1}{\ell(n-\ell)} = \frac{n-1}{2} \sum_{\ell=i}^{n-1} \left( \frac{1}{\ell} + \frac{1}{n-\ell} \right) = \frac{(n-1)(H_{n-1} + H_{n-i} - H_{i-1})}{2},$$

which completes the proof. □

In particular, when the maximum value is initially unique, i.e. when $Y_0 = 1$ with probability 1, we have $\alpha_1 = 1$ and thus

$$\mathbb{E}(T_n) = \mathbb{E}(T_n \mid Y_0 = 1) = (n-1)H_{n-1} \sim n \ln(n).$$

More generally, from Relation (2), we have

$$\mathbb{E}(T_n) \leq \mathbb{E}(T_n \mid Y_0 = 1) = (n-1)H_{n-1} \sim n \ln(n).$$

The variance of $T_n$ is obtained similarly.

**Theorem 2:** For every $n \geq 1$ and $i = 1, \ldots, n$, we have

$$\text{Var}(T_n \mid Y_0 = i) = \frac{(n-1)^2}{4} \left( \sum_{\ell=i}^{n-1} \frac{1}{\ell^2} + \sum_{\ell=1}^{n-i} \frac{1}{\ell^2} \right) = \frac{\mathbb{E}(T_n \mid Y_0 = i)}{n}. $$
Proof: If \( Y_0 = n \), which means that all the nodes start with the same values, then we have \( T_n = 0 \) and thus \( \text{Var}(T_n \mid Y_0 = n) = 0 \). For \( i = 1, \ldots, n - 1 \) we have, using the independence of the \( S_\ell \),

\[
\text{Var}(T_n \mid Y_0 = i) = \sum_{\ell = i}^{n-1} \text{Var}(S_\ell) = \sum_{\ell = i}^{n-1} \frac{1-p_\ell}{p_\ell^2}
\]

\[
= \frac{n-1}{n} \sum_{\ell = i}^{n-1} \frac{1}{\ell} - \sum_{\ell = i}^{n-1} \frac{1}{\ell} = \frac{n^2(n-1)^2}{4} \sum_{\ell = i}^{n-1} \frac{1}{\ell^2(n-\ell)^2} - \frac{n-1}{n} \sum_{\ell = i}^{n-1} \frac{1}{\ell(n-\ell)}
\]

\[
= \frac{(n-1)^2}{4} \sum_{\ell = i}^{n-1} \left( \frac{1}{\ell^2} + \frac{1}{(n-\ell)^2} \right) - \frac{n-1}{n} \sum_{\ell = i}^{n-1} \frac{1}{\ell(n-\ell)}
\]

\[
= \frac{(n-1)^2}{4} \sum_{\ell = i}^{n-1} \left( \frac{1}{\ell^2} + \frac{1}{(n-\ell)^2} \right) - \frac{\mathbb{E}(T_n \mid Y_0 = i)}{n}
\]

which completes the proof.

In particular, when the maximum value is initially unique, i.e. when \( Y_0 = 1 \) with probability 1, we have \( \alpha_1 = 1 \) and thus

\[
\text{Var}(T_n) = \text{Var}(T_n \mid Y_0 = 1) = \frac{(n-1)^2}{2} \sum_{\ell = i}^{n-1} \frac{1}{\ell^2} = \frac{n(n-1)^2}{2n^2} \sim n^{-2} n^2 = \frac{\pi^2}{12}.
\]

More generally, from Theorem 2, we have

\[
\text{Var}(T_n \mid Y_0 = i) \leq \frac{(n-1)^2}{4} \left( \sum_{\ell = i}^{n-1} \frac{1}{\ell^2} + \sum_{\ell = i}^{n-i} \frac{1}{\ell^2} \right) \leq \frac{(n-1)^2}{2} \sum_{\ell = i}^{n} \frac{1}{\ell^2} \leq \frac{\pi^2}{12}.
\]

It follows that

\[
\text{Var}(T_n) = \sum_{i=1}^{n-1} \alpha_i \text{Var}(T_n \mid Y_0 = i) \leq \frac{\pi^2}{12}.
\]

B. Explicit expression of the distribution of \( T_n \)

The distribution of \( T_n \), for \( n \geq 2 \), which is given by Relation (1) can be easily computed as follows. Let \( V(k) = (V_1(k), \ldots, V_{n-1}(k)) \) be the column vector defined by \( V_i(k) = \mathbb{P}\{T_n > k \mid Y_0 = i\} \). According to Relation (1), we have \( V(k) = Q^k \mathbb{1} \). Since \( V(0) = \mathbb{1} \), writing \( V(k) = QV(k-1) \) for \( k \geq 1 \), we get for \( i = 1, \ldots, n-2 \),

\[
\begin{cases}
V_i(k) = (1-p_i)V_i(k-1) + p_iV_{i+1}(k-1), \\
V_{n-1}(k) = (1-p_{n-1})V_{n-1}(k-1).
\end{cases}
\]

Recall that we have \( p_i = 2(i-n)/(n(n-1)) \). This recursion can be easily computed since we have, for \( k \geq 0 \),

\[
V_{n-1}(k) = (1-p_{n-1})^k = \left(1 - \frac{2}{n}\right)^k.
\]
Moreover, we have
\[ \mathbb{P}\{T_n > k\} \leq \mathbb{P}\{T_n > k \mid Y_0 = 1\}. \]

**Proof:** The result is trivial for \( n = 2 \) since in this case we have \( T_2 = 1 \). We thus suppose that \( n \geq 3 \). Note that by definition of \( p_i \) we have \( p_i = p_{n-i} \). Consider the sequence \( b_i \) defined for \( i = 1, \ldots, n-2 \), by
\[ b_1 = 1 \text{ and } b_i = \frac{p_i b_{i-1}}{p_i - p_1}, \quad \text{for } i = 2, \ldots, n-2. \]
Observing that
\[ b_i = \frac{i(n-i)b_{i-1}}{(i-1)(n-i-1)}, \]
it is easily checked by recurrence that for \( i = 1, \ldots, n-2 \), we have
\[ b_i = \frac{i(n-2)}{n-i-1}. \]

We show now by recurrence that for all \( i = 1, \ldots, n-2 \), we have
\[ V_{n-i}(k) \leq b_i (1-p_1)^k, \quad \text{for all } k \geq 0 \]
and \( V_{n-i}(k) \sim b_i (1-p_1)^k \).

Both results are true for \( i = 1 \) since \( V_{n-1}(k) = (1-p_{n-1})^k = (1-p_1)^k \). Suppose now that these results are true for a fixed integer \( i \) with \( 1 \leq i \leq n-3 \). From Relations (4), we have
\[
V_{n-i-1}(k) = (1-p_{n-i-1})V_{n-i-1}(k-1) + p_{n-i-1}V_{n-i}(k-1) = (1-p_{i+1})V_{n-i-1}(k-1) + p_{i+1}V_{n-i}(k-1).
\]
Using the recurrence hypothesis, we obtain, for what concerns the inequality,
\[ V_{n-i-1}(k) \leq (1-p_{i+1})V_{n-i-1}(k-1) + p_{i+1}b_i (1-p_1)^{k-1}. \]
Expanding this inequality and using the fact that \( V_{n-i-1}(0) = 1 \), this leads to
\[
V_{n-i-1}(k) \leq (1-p_{i+1})^k + p_{i+1}b_i (1-p_1)^{k-1} - j.
\]
Since \( b_{i+1} \geq 1 \), we get
\[ V_{n-i-1}(k) \leq b_{i+1}(1-p_1)^k \]
In the same way, using a similar calculus, we obtain
\[ V_{n-i-1}(k) \sim b_{i+1}(1-p_1)^k + b_{i+1}(1-p_1)^k. \]
Since \( p_{i+1} > p_1 \), we also get
\[ V_{n-i-1}(k) \sim b_{i+1}(1-p_1)^k. \]

We thus have shown that for all \( i = 1, \ldots, n-2 \), we have
\[ V_{n-i}(k) \leq b_i (1-p_1)^k, \quad \text{for all } k \geq 0 \]
and \( V_{n-i}(k) \sim b_i (1-p_1)^k \).

In particular, for \( i = n-2 \) we obtain
\[ V_2(k) \leq b_{n-2}(1-p_1)^k, \quad \text{for all } k \geq 0 \]
and \( V_2(k) \sim b_{n-2}(1-p_1)^k \).

Consider now the term \( V_1(k) \). From Relations (4) and using the previous inequality, we have
\[
V_1(k) \leq (1-p_1)V_1(k-1) + p_1V_2(k-1)
\leq (1-p_1)V_1(k-1) + p_1b_{n-2}(1-p_1)^{k-1}.
\]
Expanding this inequality and using the fact that \( V_1(0) = 1 \), this leads to
\[
V_1(k) \leq (1-p_1)^k + p_1b_{n-2} \sum_{j=0}^{k-1} (1-p_1)^j (1-p_1)^{k-1-j}
= (1-p_1)^k + p_1b_{n-2}(1-p_1)^{k-1}
= (1-p_1 + kp_1b_{n-2})(1-p_1)^{k-1}
\leq (1 + kp_1b_{n-2})(1-p_1)^{k-1},
\]
which gives
\[
V_1(k) \leq \left(1 + \frac{2k(n-2)^2}{n}\right) \left(1 - \frac{2}{n}\right)^{k-1}.
\]
In the same way, using a similar calculus, we obtain
\[
V_1(k) \leq \frac{2k(n-2)^2}{n} \left(1 - \frac{2}{n}\right)^{k-1}.
\]
Finally, since \( \mathbb{P}\{T_n > k \mid Y_0 = i\} \) is decreasing with \( i \), we have
\[
\mathbb{P}\{T_n > k\} = \sum_{i=1}^{n-1} \mathbb{P}\{T_n > k \mid Y_0 = i\} \mathbb{P}\{Y_0 = i\}
\leq \mathbb{P}\{T_n > k \mid Y_0 = 1\},
\]
which completes the proof.

The bound established in Theorem 4 is all the more interesting as usual probabilistic inequalities fail to provide relevant results in this particular case. For example, Markov inequality leads for all real number \( c \geq 1 \) to
\[
\mathbb{P}\{T_n \geq c \mathbb{E}(T_n)\} \leq \frac{1}{c},
\]
and Bienaym-Tchebychev inequality leads for all real number \( x > 0 \) to
\[
\mathbb{P}\{|T_n - \mathbb{E}(T_n)| \geq x\} \leq \frac{\pi^2 n^2}{12x^2}.
\]

The author of [20] provides a bound, based on Chernoff inequality, for the tail probabilities of the sum of independent, but not necessarily identically distributed, geometric random variables. In the particular case of our protocol computing the maximum, this leads to the following result.
Theorem 5: For all \( n \geq 3 \) and for all real number \( c \geq 1 \), we have
\[
\mathbb{P}(T_n > c \mathbb{E}(T_n)) \leq \frac{1}{c} \left( 1 - \frac{2}{n} \right) (c - 1 - \ln c)(n - 1) H_{n-1}.
\]
The right-hand side term is equal to 1 when \( c = 1 \).

Proof: We have already shown that
\[
\mathbb{P}(T_n > c \mathbb{E}(T_n)) \leq \mathbb{P}(T_n > c \mathbb{E}(T_n) \mid Y_0 = 1).
\]
The upper bound is then an application of Theorem 2.3 of [20], and it is clearly equal to 1 when \( c = 1 \).

Applying Theorem 4 at point \( k = \lfloor c \mathbb{E}(T_n) \rfloor \), we obtain
\[
\mathbb{P}(T_n > c \mathbb{E}(T_n)) \leq \left( 1 + \frac{2c(n - 1)H_{n-1}(n - 2)^2}{n} \right) \times \left( 1 - \frac{2}{n} \right) \mathbb{E}(T_n) - 1.
\]
\[
\leq \left( 1 + \frac{2c(n - 1)H_{n-1}(n - 2)^2}{n} \right) \times \left( 1 - \frac{2}{n} \right) e^{(c-1)H_{n-1}}.
\]
From now on we denote this bound by \( f(c, n) \) and in the same way, we denote by \( g(c, n) \) the bound of \( \mathbb{P}(T_n > c \mathbb{E}(T_n)) \) derived from Theorem 5. We then have, for \( n \geq 3 \) and \( c \geq 1 \),
\[
f(c, n) = \left( 1 + \frac{2c(n - 1)H_{n-1}(n - 2)^2}{n} \right) \times \left( 1 - \frac{2}{n} \right) e^{(c-1)H_{n-1}}.
\]
\[
g(c, n) = \frac{1}{c} \left( 1 - \frac{2}{n} \right) (c - 1 - \ln c)(n - 1) H_{n-1}.
\]
We also introduce the notation
\[
e(c, n) = \mathbb{P}(T_n > c \mathbb{E}(T_n)).
\]

Theorem 6: For every \( n \geq 3 \), there exists a unique \( c^* \geq 1 \) such that \( f(c^*, n) = e(c^*, n) \) and we have
\[
\begin{align*}
    f(c, n) &> g(c, n) \quad \text{for all } 1 \leq c < c^* \\
    f(c, n) &< g(c, n) \quad \text{for all } c > c^*.
\end{align*}
\]
Furthermore,
\[
\lim_{c \to \infty} \frac{f(c, n)}{g(c, n)} = 0.
\]

Proof: See [24]

The graphs on Figures 1, 2 and 3 illustrate the behavior of the bounds \( f(c, n) \) and \( g(c, n) \), depending on \( c \) and for different values of \( n \), compared to the real distribution of \( T_n \) at point \( c \mathbb{E}(T_n) \), i.e. to \( e(c, n) = \mathbb{P}(T_n > c \mathbb{E}(T_n)) \). The bound \( f(c, n) \) that we provided in Theorem 4 clearly shows better accuracy than the Chernoff bound \( g(c, n) \) provided in [20] above the threshold \( c^* \) introduced in Theorem 6. Furthermore, this threshold seems to decrease to 1 as \( n \) tends to infinity, as can be seen on Figure 4.
infinity, depending on the value of \( c \).

We prove in the following corollary that the bounds \( f(c, n) \) and \( g(c, n) \), obtained from Theorem 4 and Theorem 5 respectively with \( k = cE(T_n) \), both tend to 0 when \( n \) goes to infinity.

**Corollary 7:** For all real number \( c > 1 \), we have
\[
\lim_{n \to \infty} f(c, n) = 0 \quad \text{and} \quad \lim_{n \to \infty} g(c, n) = 0.
\]

**Proof:** For all \( x \in [0, 1] \), we have \( \ln(1-x) \leq -x \). Applying this property to the bound \( f(c, n) \) leads to
\[
f(c, n) \leq \frac{1 + 2c(n-1)H_{n-1}(n-2)^2}{n} e^{-2(c(n-1)H_{n-1}(n-2))/n} \leq \frac{(1 + 2c(n-2)^2 H_{n-1})}{n} e^{-2(c(n-1)H_{n-1}(n-2))/n}.
\]
Since \( \ln(n) \leq H_{n-1} \leq 1 + \ln(n-1) \), we get
\[
f(c, n) \leq \frac{(1 + 2c(n-2)^2(1 + \ln(n-1)))}{n} e^{2c \ln(n)} e^{-2c \ln(n)}/n.
\]
For \( x \geq 0 \), the function \( u(x) = x e^{2c (\ln(x) + 2)/x} \) satisfies \( u(x) \leq \exp(2c/e^{(c-2)/c}) \), so we obtain
\[
f(c, n) \leq \frac{1 + 2c(n-2)^2(1 + \ln(n-1))}{n^{2e}} \exp\left(\frac{2c/e^{(c-2)/c}}{n}\right).
\]
The fact that \( c > 1 \) implies that this last term tends to 0 when \( n \to \infty \). Concerning the bound \( g(c, n) \), we have
\[
g(c, n) = \frac{1}{c} \left(1 - \frac{2}{n}\right)(c-1-\ln(c))(n-1)H_{n-1}.
\]
For \( n \to \infty \), \( g(c, n) \) tends to 0 when \( n \) tends to infinity, since \( c - 1 - \ln(c) \) is positive for \( c > 1 \).

**Theorem 8:** For all real \( c \geq 0 \), we have
\[
\lim_{n \to +\infty} P\{T_n > cE(T_n)\} = \begin{cases} 0 & \text{if } c > 1 \\ 1 & \text{if } c < 1. \end{cases}
\]

**Proof:** From Corollary 7, both bounds \( f(c, n) \) and \( g(c, n) \) of \( P\{T_n > cE(T_n)\} \) tend to 0 when \( n \) tends to infinity, so using either \( f(c, n) \) or \( g(c, n) \) we deduce that
\[
\lim_{n \to \infty} P\{T_n > cE(T_n)\} = 0 \quad \text{for all } c > 1.
\]
In the case where \( c < 1 \), Theorem 3.1 of [20] leads to
\[
P\{T_n > cE(T_n)\} \geq 1 - e^{-2(n-1)H_{n-1}(c-1-\ln(c))/n} \geq 1 - e^{-2(n-1)\ln(n)(c-1-\ln(c))/n}.
\]
Since \( c - 1 - \ln(c) > 0 \) for all \( c \in [0, 1] \), the right-hand side term of this inequality tends to 1 when \( n \to \infty \). Thus,
\[
\lim_{n \to \infty} P\{T_n > cE(T_n)\} = 1 \quad \text{if } c < 1.
\]

The results established previously don’t allow us to figure out neither the existence of \( \lim_{n \to \infty} P\{T_n > cE(T_n)\} \) when \( c = 1 \), nor its value. However, numerical results give us a glimpse of its limiting behavior.

In Figure 5, we show the probability \( P\{T_n > E(T_n)\} \) for different values of \( n \). The oscillations of this probability with \( n \) are due to the fact \( T_n \) is a discrete random variable and \( E(T_n) \) is not an integer. That is why we propose in this figure a smoothing of this probability using the sequence
\[
s_n = (1 - a_n)P\{T_n > E(T_n)\} + a_nP\{T_n > E(T_n) + 1\},
\]
where \( a_n \) is the fractional part of \( E(T_n) \), that is \( a_n = E(T_n) - \lfloor E(T_n) \rfloor \). Since \( a_n \in [0, 1] \), we have
\[
P\{T_n > E(T_n) + 1\} \leq s_n \leq P\{T_n > E(T_n)\},
\]
that is why we also show in this figure the probability \( P\{T_n > E(T_n) + 1\} \). We also checked that the sequence \( (s_n) \) is increasing until \( n = 20000 \). This figure suggests the following result proposed as a conjecture.

**Conjecture:** \( \lim_{n \to \infty} P\{T_n > E(T_n)\} \) exists and \( \approx 0.4484 \).

Figure 6 shows the results obtained by simulation concerning the proportion of nodes informed by rumor as a function of the parallel time. Recall that the parallel time refers to as the successive number of steps each node executes [5]. Initially, a
single node is informed of the rumor. This figure illustrates our analysis. For instance, with probability almost 1 one thousand nodes (resp. one million nodes) learn the rumor after no more than 7 (resp 11) interactions for each of them. The complexity in space (number of memory bits) is in \( O(1) \).

VI. Conclusion

In this paper we have provided a thorough analysis of the rumor spreading time in the population protocol model. Providing such a precise analysis is a step towards the design of more complex functionality achieved by combining simple population protocols [23], [3]. Indeed, an important feature of population protocols is that they do not halt. Nodes can never know whether their computation is completed and thus nodes forever interact with their neighbors while their outputs stabilize to the desired value (e.g. the maximal value of any node of the network). By precisely characterizing, for each protocol of interest, with any high probability, the number of interactions each node must execute to converge to the desired value, each node can on its own, decide the time from which the current protocol has stabilized and start the parallel of sequential executions of the next ones.

REFERENCES


