Loosely-Stabilizing Leader Election on Arbitrary Graphs in Population Protocols

Yuichi Sudo^{1,2}, Fukuhito Ooshita², Hirotsugu Kakugawa², and Toshimitsu Masuzawa²

 ¹ NTT Secure Platform Laboratories 3-9-11 Midori-cho, Musashino, Tokyo, 180-8585, Japan sudo.yuichi@lab.ntt.co.jp
 ² Graduate School of Information Science and Technology, Osaka University 1-5 Yamadaoka, Suita, Osaka, 565-0871, Japan {y-sudou,f-oosita,kakugawa,masuzawa}@ist.osaka-u.ac.jp

Abstract. In the population protocol model Angluin et al. proposed in 2004, there exists no self-stabilizing protocol that solves leader election on complete graphs without knowing the exact number of nodes. To circumvent the impossibility, we previously introduced the concept of loosestabilization, which relaxes the closure requirement of self-stabilization. A loosely-stabilizing protocol guarantees that starting from any initial configuration a system reaches a loosely-safe configuration, and after that, the system keeps its specification (e.g. the unique leader) not forever, but for a sufficiently long time. Our previous work presented a loosely-stabilizing protocol that solves the leader election on complete graphs using only the upper bound N of n, not the exact value of n. We take this work one step further in this paper: We propose two looselystabilizing protocols that solve leader election for *arbitrary graphs*. One is a deterministic protocol that uses the identifiers of nodes while the other is a probabilistic protocol that works on anonymous networks. Given the upper bounds N and Δ of the number of nodes and the maximum degree of nodes respectively, both protocols keep a unique leader for $\Omega(Ne^N)$ expected steps after entering a loosely-safe configuration. The former enters a loosely-safe configuration within $O(m\Delta N \log n)$ expected steps while the latter does within $O(m\Delta^2 N^3 \log N)$ expected steps where m is the number of edges of the graph.

Keywords: Loose-stabilization, Population protocols, Leader election.

1 Introduction

The *population protocol* (PP) model, which was presented by Angluin et al.[1], represents wireless sensor networks of mobile sensing devices that cannot control their movement. Two devices (say *agents*) communicate with each other only when they come sufficiently close to each other (we call this event an *interaction*). One example represented by this model is a flock of birds where each bird is equipped with a sensing device with a small transmission range; each device

© Springer International Publishing Switzerland 2014

can communicate with another device only when the corresponding birds come sufficiently close to each other. This unique but meaningful model has attracted broad attention, and there have been numerous studies involving it.

Self-stabilizing leader election (SS-LE) requires that starting from any configuration, a system (say *population*) reaches a safe-configuration in which a unique leader is elected, and after that, the population has the unique leader forever. Self-stabilizing leader election is important in the PP model because (i) many population protocols in the literature work on the assumption that a unique leader exists [1,2,3], and (ii) self-stabilization tolerates any finite number of transient faults and this property suits systems consisting of numerous cheap and unreliable nodes. (Such systems are the original motivation of the PP model.) However, there exists strict impossibility of SS-LE in the PP model: no protocol can solve SS-LE for complete graphs, arbitrary graphs, trees, lines, degree-bounded graphs and so on unless the exact size of the graph (the number of agents n is available [3]. This impossibility holds even if we strengthen the PP model by assigning unique identifies to agents, allowing agents to use random numbers, introducing memory of communication links (mediated population protocols [10]), or allowing more than two agents (k agents) to interact at the same time (the PP_k model [5]).

Accordingly, many studies of SS-LE took either one of the following two approaches. One approach is to accept the assumption that the exact value of n is available and focus on the space complexity of the protocol. Cai et al. [6] proved that n states of each agent is necessary and sufficient to solve SS-LE for a complete graph of n agents. Mizoguchi et al.[12] and Xu et al.[14] improved the space-complexity by adopting the mediated population protocol model and the PP_k model respectively. The other approach is to use *oracles*, a kind of failure detectors. Fischer and Jiang [8] took this approach for the first time. They introduced oracle Ω ? that informs all agents whether at least one leader exists or not and proposed two protocols that solve SS-LE for rings and complete graphs by using Ω ?. Beauquier et al.[4] presented an SS-LE protocol for arbitrary graphs that uses two copies of Ω ?. Canepa et al.[7] proposed two SS-LE protocols that use Ω ? and consume only 1 bit of each agent: one is a deterministic protocol for trees and the other is a probabilistic protocol for arbitrary graphs although the position of the leader is not static and moves among the agents.

Our previous work [13] took another approach to solve SS-LE. We introduced the concept of loose-stabilization, which relaxes the closure requirement of selfstabilization: we allow protocols to deviate from the specification after following it for a sufficiently long time. Concretely, starting from any initial configuration, the population must reach a loosely-safe configuration within a relatively short time; after that, the specification of the problem (the unique leader) must be kept for a sufficiently long time, though not forever. We then proposed a loosely-stabilizing protocol that solves leader election on complete graphs using only an upper bound N of n, not using the exact value of n. Starting from any configuration, the protocol enters a loosely-safe configuration within $O(nN \log n)$ expected steps. After that, the unique leader is kept for $\Omega(Ne^N)$ expected steps. Since the specification is kept for an exponentially long time, we can say this loosely-stabilizing protocol is practically equivalent to a self-stabilizing leader election protocol. Furthermore, this protocol works on any complete graph whose size is no more than N while protocols using the exact value of n work only on the complete graph of size n.

Some works on population protocols assume the probabilistic distribution regarding the interactions of agents: any interaction occurs uniformly at random [1,2,13]. This assumption have been used partly for evaluating the time complexity of protocols. We also adopt this assumption because the measure of time is crucial in the concept of loose-stabilization.

1.1 Our Contribution

In this paper, we consider loosely-stabilizing leader election for arbitrary undirected graphs. We adopt two settings: the population with agent-identifiers as in [9] ¹ and the population in which agents can use random numbers for statetransition as in [7]. As mentioned above, no self-stabilizing protocol can solve SS-LE for arbitrary graphs, even in these settings, unless the exact value of n is available. For each setting, we propose two protocols $P_{\rm ID}$ and $P_{\rm RD}$ respectively. To elect the unique leader, we take "the minimum ID selection" approach for $P_{\rm ID}$ utilizing the identifiers of agents while we take a novel approach we call "virus war mechanism" for $P_{\rm RD}$ utilizing random numbers.

Given upper bounds N of n and Δ of the maximum degree of nodes, both protocols keep the unique leader for $\Omega(Ne^N)$ expected steps after entering a loosely-safe configuration. Protocol P_{ID} enters a loosely-safe configuration within $O(mN\Delta \log n)$ expected steps while P_{RD} does within $O(mN^3\Delta^2 \log N)$ expected steps where m is the number of edges of the graph. Both protocols consume only $O(\log N)$ bits of each agent's memory. We can say this space complexity is small because even space optimal self-stabilizing protocols that use exact value of n consume $O(\log n)$ bits of each agent[6,12]. For simplicity, our protocols are presented for undirected graphs. However, they work on *directed* graphs with slight modification which is discussed in the conclusion.

Angluin et al.[1] proves that for any population protocol P working on complete graphs, there exists a protocol that simulates P on any arbitrary graph. However, this simulation can be achieved assuming that all the agents have the common initial states at the start of the execution. Since we cannot assume the specific initial states (This is the essence of self-stabilization), we cannot translate our previous loosely-stabilizing algorithm[13] for complete graphs to a loosely-stabilizing algorithm that works for arbitrary graphs.

¹ Strictly speaking, our model with identifiers is stronger than the model in [9]. We use identifiers to compare their values while Guerraoui et al.[9] only allow equality-test of identifiers and prohibited any other calculation of identifiers such as value-comparing.

2 Preliminaries

This section defines the model we consider for this paper. The model includes both agent-identifiers and random numbers while protocols P_{ID} and P_{RD} use only one of them. In what follows, we denote set $\{z \in \mathbb{N} \mid x \leq z \leq y\}$ by [x, y].

A population is a simple and weakly-connected directed graph $G(V, E, \mathrm{id})$ where V ($|V| \ge 2$) is a set of agents, $E \subseteq V \times V$ is a set of directed edges and *id* defines unique identifiers of agents. Each edge represents a possible interactions (or communication between two agents): If $(u, v) \in E$, agents u and v can interact with each other where u serves as an *initiator* and vserves as a *responder*. Each agent v has the unique identifier $\mathrm{id}(v) \in I$ ($I = [0, \mathrm{id}_{\max}]$, $\mathrm{id}_{\max} \in O(n^c)$ for constant c). We say that G is undirected if it satisfies $(u, v) \in E \Leftrightarrow (v, u) \in E$. We define n = |V| and m = |E|.

A protocol P(Q, Y, I, R, T, O) consists of a finite set Q of states, a finite set Y of output symbols, a set of possible identifiers I, a range of random numbers $R \subset \mathbb{N}$, transition function $T: (Q \times I) \times (Q \times I) \times R \to Q \times Q$, and output function $O: (Q \times I) \to Y$. When an interaction between two agents occurs, T determines the next states of the two agents based on the current states of the agents, identifiers of the two agents, and a random number $r \in R$ generated at each interaction. The *output of an agent* is determined by O: the output of agent v with state $q \in Q$ is O(q, id(v)). We assume that the set of possible identifiers I is a given parameter and not subject to protocol design.

A configuration is a mapping $C: V \to Q$ that specifies the states of all the agents. We denote the set of all configurations of protocol P by $\mathcal{C}_{all}(P)$. We say that configuration C changes to C' by interaction e = (u, v) and integer $r \in R$, denoted by $C \stackrel{e,r}{\to} C'$, if we have (C'(u), C'(v)) = T(C(u), id(u), C(v), id(v), r) and C'(w) = C(w) for all $w \in V \setminus \{u, v\}$. A scheduler determines which interaction occurs at each time. In this paper, we consider a uniformly random scheduler $\Gamma = \Gamma_0, \Gamma_1, \ldots$: each $\Gamma_t \in E$ is a random variable such that $\Pr(\Gamma_t = (u, v)) = 1/m$ for any $t \geq 0$ and any $(u, v) \in E$. We also define the random number sequence as $A = R_1, R_2, \ldots$: each number $R_t \in R$ is a random variable such that $\Pr(R_t = r) = 1/|R|$ for any $t \geq 0$ and $r \in R$. Given an initial configuration C_0, Γ , and Λ , the execution of protocol P is defined as $\Xi_P(C_0, \Gamma, \Lambda) = C_0, C_1, \ldots$ such that $C_t \stackrel{\Gamma_t, R_t}{\to} C_{t+1}$ for all $t \geq 0$. We denote $\Xi_P(C_0, \Gamma, \Lambda)$ simply by $\Xi_P(C_0)$ when no misunderstanding can arise.

The leader election problem requires that every agent should output L or Fwhich means "leader" or "follower" respectively. We say that a finite or infinite sequence of configurations $\xi = C_0, C_1, \ldots$ preserves a unique leader, denoted by $\xi \in LE$, if there exists $v \in V$ such that $O(C_t(v), \mathrm{id}(v)) = L$ and $O(C_t(u), \mathrm{id}(u)) = F$ for any $t \geq 0$ and $u \in V \setminus \{v\}$. For $\xi = C_0, C_1, \ldots$, the holding time of the leader $\mathrm{HT}(\xi, LE)$ is defined as the maximum $t \in \mathbb{N}$ that satisfies $(C_0, C_1, \ldots, C_{t-1}) \in LE$. We define $\mathrm{HT}(\xi, LE) = 0$ if $C_0 \notin LE$. We denote $\mathbf{E}[\mathrm{HT}(\Xi_P(C), LE)]$ by $\mathrm{EHT}_P(C, LE)$. Intuitively, $\mathrm{EHT}_P(C, LE)$ is the expected number of interactions for which the population keeps the unique leader after protocol P starts from configuration C. For configuration sequence $\xi = C_0, C_1, \ldots$ and a set of configurations \mathcal{C} , we define convergence time $\operatorname{CT}(\xi, \mathcal{C})$ as the minimum $t \in \mathbb{N}$ that satisfies $C_t \in \mathcal{C}$. We define $\operatorname{CT}(\xi, \mathcal{C}) = |\xi|$ if $C_t \notin \mathcal{C}$ for any $t \geq 0$, where $|\xi|$ is the length of ξ . We denote $\operatorname{E}[\operatorname{CT}(\Xi_P(C), \mathcal{C})]$ by $\operatorname{ECT}_P(C, \mathcal{C})$. Intuitively, $\operatorname{ECT}_P(C, \mathcal{C})$ is the expected number of interactions by which the population enters a configuration in \mathcal{C} after P starts from C.

Definition Protocol P(Q, Y, I, R, T, O) is an (α, β) -loosely-stabilizing leader election protocol if there exists set S of configurations satisfying two inequalities $\max_{C \in \mathcal{C}_{au}(P)} \text{ECT}_{P}(C, S) \leq \alpha$ and $\min_{C \in S} \text{EHT}_{P}(C, LE) \geq \beta$.

2.1 Chernoff Bounds

In this section, we quote the three variants of Chernoff bounds [11] used in several proofs of this paper.

Lemma 1 (from Eq. (4.2) in [11]). The following inequality holds for any binomial random variable X:

$$\Pr(X \ge 2\mathbf{E}[X]) \le e^{-\mathbf{E}[X]/3}.$$

Lemma 2 (from Eq. (4.5) in [11]). The following inequality holds for any binomial random variable X:

$$\Pr(X \le \mathbf{E}[X]/2) \le e^{-\mathbf{E}[X]/8}.$$

Lemma 3 (from Eq. (4.5) in [11]). The following inequality holds for any binomial random variable X:

$$\Pr(X \le \mathbf{E}[X]/4) \le e^{-9\mathbf{E}[X]/32}.$$

3 Leader Election with Identifiers

This section presents loosely-stabilizing leader election protocol P_{ID} , which works on arbitrary undirected graphs with unique identifiers of agents (Protocol 1). In the protocol description, we regard a state of agents as a collection of *variables* (e.g. timer), and denote a transition function as pseudo code that updates variables of initiator x and responder y. We denote the value of variable var of agent $v \in V$ by v.var. We also denote the value of var in state $q \in Q$ by q.var.

This protocol elects the agent with the minimum identifier, denoted by v_{\min} , as the leader. Each agent v tries to find the minimum identifier and stores it on v.lid. At interaction, two agents x and y compare their lid and store the smaller value on their lid (Lines 3 and 6), by which the smallest identifier $id(v_{\min})$ eventually spreads to all the agents. Then, after some point, v_{\min} is the unique leader because output function O makes only agents v satisfying id(v) = v.lid output L and other agents output F.

Protocol 1. Leader Election with Identifiers $P_{\rm ID}$

Variables of each agent: lid $\in I$, timer $\in [0, t_{\max}]$ **Output function** *O*: if v.lid = id(v) holds, then the output of agent v is L; Otherwise, F. **Interaction** between initiator x and responder y: 1: if x.lid > id(x) then $x.lid \leftarrow id(x)$ endif 2: if x.lid < y.lid then 3: $y.\texttt{lid} \leftarrow x.\texttt{lid}$ $x.\texttt{timer} \leftarrow y.\texttt{timer} \leftarrow \max(x.\texttt{timer} - 1, 0)$ 4: 5: else if x.lid > y.lid then $x.\texttt{lid} \leftarrow y.\texttt{lid}$ 6: 7: $x.\texttt{timer} \leftarrow y.\texttt{timer} \leftarrow \max(y.\texttt{timer} - 1, 0)$ //x.lid = y.lid at this time 8: else $x.\texttt{timer} \leftarrow y.\texttt{timer} \leftarrow \max(x.\texttt{timer} - 1, y.\texttt{timer} - 1, 0)$ 9: 10: end if 11: if id(x) = x.lid or id(y) = y.lid then // a leader resets timers 12: $x.\texttt{timer} \leftarrow y.\texttt{timer} \leftarrow t_{\max}$ 13: else if x.timer = 0 then // a new leader is created at timeout $x.\mathtt{lid} \leftarrow y.\mathtt{lid} \leftarrow \min(\mathrm{id}(x), \mathrm{id}(y))$ 14:15: $x.\texttt{timer} \leftarrow y.\texttt{timer} \leftarrow t_{\max}$ 16: end if

However, in the initial configuration, some agents may have false identifiers (or the integers that are not identifiers of any agent in the population) on lid. A false identifier may spread to the population instead of $id(v_{\min})$ if it is smaller than $id(v_{\min})$. We define $ID = \{id(v) \mid v \in V\}$, which is the correct identifiers set (Note that ID \subseteq I). Protocol P_{ID} removes false identifiers $i \notin$ ID from lid of all the agents by the *timeout mechanism*. Specifically, if $x.lid \neq y.lid$, we take the timer value of the agent with smaller lid, decrease it by one, and substitute the decreased value into x.lid and y.lid (Lines 4 and 7). If x.lid = y.lid, we take the larger value of x.timer and y.timer, decrease it by one, and substitute the decreased value into x.lid and y.lid (Line 9). We call this event larger value propagation. If x or y is a leader, both timers are reset to $t_{\rm max}$ (Line 12). We call this event timer reset. When a timer becomes zero, agents x and y suspect that there exists no leader in the population. In this case, they elect the one with a smaller identifier as a leader by substituting $\min(id(x), id(y))$ into x.lid and y.lid (Line 14). We call this event *timeout*. Agents with false identifiers never experience timer reset; thus, their timers keep on decreasing. Hence, timeout eventually occurs and their lids satisfy $lid \in ID$. This mechanism rarely ruins the stability of the unique leader because agents with $lid \in ID$ keep high value timers because of timer reset and lager value propagation.

Complexity Analysis The upper bound t_{\max} of variable timer is the only parameter of P_{ID} , which affects the correctness and complexities of the protocol. We assume $t_{\max} \geq 8\delta \max(d, 2 + \log n)$ where δ is the maximum degree of the agents and

d is the diameter of population G. (Note that δ is an even number because G is undirected.) We prove the following equations under this assumption:

$$\max_{C \in \mathcal{C}_{all}} \operatorname{ECT}_{P_{\mathrm{ID}}}(C, \mathcal{S}_{\mathrm{id}}) = O(m\delta\tau \log n), \tag{1}$$

$$\min_{C \in \mathcal{S}_{id}} \operatorname{EHT}_{P_{ID}}(C, LE) = \Omega(\tau e^{\tau}), \qquad (2)$$

where $\tau = t_{\max}/(8\delta)$ and S_{id} is the set of configurations in which $v.lid = id(v_{\min})$ and $v.timer > t_{\max}/2$ hold for all $v \in V$ and $v_{\min}.timer = t_{\max}$ holds. When upper bounds N of n and Δ of δ are available and we assign $t_{\max} = 8N\Delta$, protocol P_{ID} is an $(O(m\Delta N \log n), \Omega(Ne^N))$ -loosely-stabilizing leader election protocol.

First, we analyze the expected holding time. Let $C_0 \in S_{\text{id}}$ and $\Xi_{P_{\text{ID}}}(C_0) = C_0, C_1, \ldots$. To prove (2), it suffices to show that both $C_0, \ldots, C_{2m\tau} \in LE$ and $C_{2m\tau} \in S_{\text{id}}$ hold with probability at least $p_{\text{suc}} = 1 - O(ne^{-\tau})$. Then, we have $\min_{C_0 \in S_{\text{id}}} \text{EHT}_{P_{\text{ID}}}(C_0, LE) \geq 2m\tau/(1 - p_{\text{suc}}) = \Omega(\tau e^{\tau})$.

Lemma 4. The probability that every $v \in V$ joins only less than $t_{\max}/2$ interactions among $\Gamma_0, \ldots, \Gamma_{2m\tau-1}$ is at least $1 - ne^{-\tau}$.

Proof. For any $v \in V$ and $t \geq 0$, v joins interaction Γ_t with probability at most δ/m . Thus, the number of interactions v joins during the $2m\tau$ interactions is bounded by binomial random variable $X \sim B(2m\tau, \delta/m)$. Applying a variant of Chernoff bound (Lemma 1), we have

$$\Pr(X \ge t_{\max}/2) = \Pr(X \ge 2\mathbf{E}[X]) \qquad \because t_{\max} = 8\delta\tau$$
$$\le e^{-\mathbf{E}[X]/3}$$
$$= e^{-2\delta\tau/3} \qquad \text{(By Chernoff Bound of Lemma 1)}$$
$$\le e^{-\tau}. \qquad \because \delta \ge 2$$

Summing up the probabilities for all $v \in V$ gives the lemma.

Lemma 5. Let $C_0 \in \mathcal{L}_{\text{lid}}$ and $\Xi_{P_{\text{ID}}}(C_0) = C_0, C_1, \ldots$ Then, we have the following inequality:

$$\Pr(\forall v \in V, \ C_{2m\tau}(v).\texttt{timer} > t_{\max}/2) \ge 1 - 2ne^{-\tau}.$$

Proof. It suffices to show $\Pr(C_{2m\tau}(v).\texttt{timer} > t_{\max}/2) \geq 1 - 2e^{-\tau}$ for any agent $v \in V$. We denote the shortest path from v_{\min} to v by (v_0, v_1, \ldots, v_k) where $v_0 = v_{\min}, v_k = v, 0 \leq k \leq d$ and $(v_{i-1}, v_i) \in E$ for all $i = 1, \ldots, k$. For any $t \in [0, 2m\tau]$, we define $v_{\text{head}}(t)$ as v_l with maximum $l \in [1, k]$ such that there exist t_1, t_2, \ldots, t_l satisfying $0 \leq t_1 < t_2 < \cdots < t_l < t$ and $\Gamma_{t_i} \in$ $\{(v_{i-1}, v_i), (v_i, v_{i-1})\}$ for $i = 1, 2, \ldots, l$. We define $v_{\text{head}}(t) = v_0$ if such l does not exist. Intuitively, $v_{\text{head}}(t)$ is the head of the agents in path (v_0, v_1, \ldots, v_k) to which a large timer value is propagated from v_{\min} . (Remember that v_{\min} resets the timers to t_{\max} .) We define J(t) as the number of integers $i \in [0, t]$ such that $v_{\text{head}}(i)$ joins interaction Γ_i . Intuitively, J(t) is the number of interactions that the head agent joins among $\Gamma_0, \ldots, \Gamma_t$. Obviously, we have $C_t(v_{\text{head}}(t))$.timer $\geq t_{\text{max}} - J(t)$ for any $t \geq 0$.

In what follows, we prove $\Pr(v_{\text{head}}(2m\tau) = v) \ge 1 - e^{-\tau}$ and $\Pr(J(2m\tau) < t_{\max}/2) \ge 1 - e^{-\tau}$, which give $\Pr(C_{2m\tau}(v).\texttt{timer} > t_{\max}/2) \ge 1 - 2e^{-\tau}$. For any $i \in [1, k]$, a pair v_{i-1} and v_i interacts with probability 2/m at each interaction. Hence, we can say each interaction makes v_{head} forward with probability 2/m. Therefore, by letting Z be a binomial random variable such that $Z \sim B(2m\tau, 2/m)$, we have

$$\begin{aligned} \Pr(v_{\text{head}}(t) = v) &= 1 - \Pr(Z < k) \\ &\geq 1 - \Pr(Z < d) \\ &\geq 1 - \Pr\left(Z < \frac{1}{4} \cdot \mathbf{E}[Z]\right) \qquad \because d \le \tau = \frac{1}{4} \cdot \mathbf{E}[Z] \\ &\geq 1 - e^{-9\mathbf{E}[Z]/32} \qquad \text{(By Chernoff bound of Lemma 3)} \\ &> 1 - e^{-\tau}. \end{aligned}$$

The probability that $v_{\text{head}}(t)$ joins interaction Γ_t is at most δ/m regardless of t. Hence, by letting Z' be a binomial random variable such that $Z' \sim B(2m\tau, \delta/m)$, we have

$$\Pr(J(2m\tau) < t_{\max}/2) > 1 - \Pr(Z' \ge t_{\max}/2)$$

= 1 - $\Pr(Z' \ge 2\mathbf{E}[Z'])$
> 1 - $e^{-\mathbf{E}[Z']/3}$ (By Chernoff bound of Lemma 1)
= 1 - $e^{-2\delta\tau/3}$
> 1 - $e^{-\tau}$. $\because \delta \ge 2$

Thus, we have shown $\Pr(C_{2m\tau}(v).\texttt{timer} > t_{\max}/2) \ge 1 - 2e^{-\tau}.$

Lemma 6. $\min_{C \in S_{id}} \text{EHT}_{P_{ID}}(C, LE) = \Omega(\tau e^{\tau}).$

Proof. We have $C_0, \ldots, C_{2m\tau} \in LE$ and $C_{2m\tau} \in S_{id}$ if $C_0 \in S_{id}$ holds, no timeout happens, and any agent interacts at most $t_{\max}/2$ times during $2m\tau$ interactions. Hence, probability p_{suc} discussed in the beginning of this section is at least $1 - 3ne^{-\tau}$ by Lemmas 4 and 5, which leads to the lemma.

Next, we analyze the expected convergence time. To prove (1), we define two sets of configurations: $C_{\text{lid}} = \{C \in C_{\text{all}}(P_{\text{ID}}) \mid \forall v \in V, C(v). \text{lid} \in \text{ID}\}$ and $\mathcal{L}_{\text{lid}} = C_{\text{lid}} \cap \{C \in C_{\text{all}}(P_{\text{ID}}) \mid C(v_{\min}). \text{lid} = \text{id}(v_{\min}) \land C(v_{\min}). \text{timer} = t_{\max}\}.$

Lemma 7. $\max_{C \in \mathcal{C}_{all}(P_{ID})} \text{ECT}_{P_{ID}}(C, \mathcal{C}_{lid}) = O(m\delta\tau \log n).$

Proof. Let z be the maximum value of v.timer such that v.lid \notin ID. This z decreases by one every time all interactions of E occur. Thus, it takes at most $\frac{m}{m} + \frac{m}{m-1} + \ldots \frac{m}{1} \leq m(1 + \log m)$ expected steps to decrease z by one. Hence, $\max_{C \in \mathcal{C}_{all}(P_{\text{ID}})} \text{ECT}_{P_{\text{ID}}}(C, \mathcal{C}_{\text{Id}}) \leq t_{\max}m(1 + \log m) = O(m\delta\tau \log n).$

Lemma 8. $\max_{C \in \mathcal{C}_{\text{lid}}} \text{ECT}_{P_{\text{ID}}}(C, \mathcal{L}_{\text{lid}}) = O(m).$

Proof. We have v_{\min} .lid = id (v_{\min}) and v_{\min} .timer = t_{\max} just after v_{\min} interacts in any configuration of C_{lid} . This takes O(m) expected interactions.

Lemma 9. $\max_{C \in \mathcal{L}_{\text{lid}}} \text{ECT}_{P_{\text{ID}}}(C, \mathcal{S}_{\text{id}}) = O(m\tau).$

Proof Sketch. Let $C_0 \in \mathcal{L}_{\text{lid}}$ and $\mathcal{Z}_{P_{\text{ID}}}(C_0) = C_0, C_1, \dots$ By similar argument to Lemmas 4 and 5, we can prove $\Pr(C_{2m\tau} \in \mathcal{S}_{\text{id}}) > 1 - 2ne^{-\tau}$. Since $C \in \mathcal{L}_{\text{lid}}$ cannot change to $D \notin \mathcal{L}_{\text{lid}}$, we have $\max_{C \in \mathcal{L}_{\text{lid}}} \operatorname{ECT}_{P_{\text{ID}}}(C, \mathcal{S}_{\text{id}}) \leq 2m\tau + 3ne^{-\tau} \cdot \max_{C \in \mathcal{L}_{\text{lid}}} \operatorname{ECT}_{P_{\text{ID}}}(C, \mathcal{S}_{\text{id}})$. Solving this inequality gives the lemma.

The following lemma immediately follows from Lemmas 7, 8, and 9.

Lemma 10. $\max_{C \in \mathcal{C}_{all}(P_{ID})} \text{ECT}_{P_{ID}}(C, \mathcal{S}_{id}) = O(m\delta\tau \log n).$

Lemmas 6 and 10 gives the following theorem.

Theorem 1 Protocol P_{ID} is a $(O(m\delta\tau \log n), \Omega(\tau e^{\tau}))$ loosely-stabilizing leader election protocol for arbitrary graphs when $t_{\max} \ge 8\delta \max(d, 2 + \log n)$.

Therefore, given upper bound N and Δ of n and δ respectively, we get a $(O(m\Delta N \log n), \Omega(Ne^N))$ loosely-stabilizing leader election protocol for arbitrary graphs by assigning $t_{\max} = 8N\Delta$.

4 Leader Election with Random Numbers

This section presents loosely-stabilizing leader election protocol $P_{\rm RD}$. It works on arbitrary undirected anonymous graphs with a random number generated at each interaction (Protocol 2). Random numbers are used in Line 11: When the protocol enters Line 11, the code is executed with probability p = 1/|R|. This is implemented as the code is executed only when a specific number is generated. For example, p = 0.01 if we assign R = [0, 99] and treat 0 as a specific number.

Each agent has binary variable $DoA \in \{DEAD, ALIVE\}$ and three timers $timer_L$, $timer_V$ and $timer_S$. The output function defines leaders based on DoA: agent v is a leader if v is alive (or v.DoA = ALIVE), and a follower if v is dead (or v.DoA = DEAD). Protocol P_{RD} consists of a timeout mechanism (Lines 1-7) and a virus-war mechanism (Lines 8-14). By using $timer_L$, the timeout mechanism creates a leader when it is suspected that no leader exists. By using $timer_V$ and $timer_S$, the virus-war mechanism reduces the number of leaders.

The timeout mechanism is almost the same as P_{ID} . By the timer reset and the larger value propagation, timeout eventually occurs when no leader exists, and all agents keep high timer values with high probability when one ore more leaders exist. At timeout, a dead agent becomes a leader (Line 5).

In the virus-war mechanism, each leader tries to kill other leaders by viruses and become the unique leader. We say that agent v has a virus if $v.timer_V > 0$, and v wears a (head) shield if $v.timer_S > 0$. A leader creates a new virus with probability p when it interacts as an initiator (Line 11). When creating a virus,

Protocol 2. Leader Election with Random Numbers $P_{\rm RD}$

Variables of each agent: $DoA \in \{DEAD, ALIVE\}, timer_{L} \in [0, t_{max}], timer_{V} \in [0, t_{virus}], timer_{S} \in [0, t_{shld}]$ **Output function** *O*: if v.DoA = ALIVE holds, then the output of agent v is L, otherwise F. **Interaction** between initiator x and responder y: 1: $x.\texttt{timer}_{L} \leftarrow y.\texttt{timer}_{L} \leftarrow \max(x.\texttt{timer}_{L} - 1, y.\texttt{timer}_{L} - 1, 0)$ 2: if x.DoA = ALIVE or y.DoA = ALIVE then 3: $x.\texttt{timer}_\texttt{L} \leftarrow y.\texttt{timer}_\texttt{L} \leftarrow t_{\max}$ // a leader resets timer 4: else if $x.timer_L = 0$ then // a new leader is created at timeout 5: $x.DoA \leftarrow ALIVE$ 6: $x.\texttt{timer}_{\texttt{L}} \leftarrow y.\texttt{timer}_{\texttt{L}} \leftarrow t_{\max}$ 7: end if 8: $x.timer_{V} \leftarrow y.timer_{V} \leftarrow \max(x.timer_{V} - 1, y.timer_{V} - 1, 0)$ 9: $x.\texttt{timer}_{s} \leftarrow \max(0, x.\texttt{timer}_{s} - 1)$ 10: if x.DoA = ALIVE then Execute $(x.timer_v \leftarrow t_{virus}, x.timer_s \leftarrow t_{shld})$ with probability p 11:// An alive initiator creates a new virus and a new shield with probability p. 12: end if

13: if $x.timer_V > 0$ and $x.timer_S = 0$ then $x.DoA \leftarrow DEAD$ endif 14: if $y.timer_V > 0$ and $y.timer_S = 0$ then $y.DoA \leftarrow DEAD$ endif

the agent wears a shield so as not to be killed by the new virus (Line 11). A virus spreads among agents by interactions (Line 8), and an agent is killed when it has a virus without a shield (Lines 13-14). A virus has TTL (time to live), which is memorized on timer_V and decreased by one at each interaction of its owner (line 8). When timer_V becomes zero, the virus vanishes and looses the ability to kill agents. A shield also has TTL, which is memorized on timer_S and decreased by one at each interaction of its owner (Line 9). When timer_S becomes zero, the shield vanishes and looses the ability to protect its owner from viruses.

The virus-war mechanism correctly works if p is sufficiently small and $t_{\rm shld}$ is sufficiently greater than $t_{\rm virus}$. Consider the case multiple leaders exist. Since p is small, all viruses and shields eventually vanishes. After that, some agent eventually creates a new virus and shield. The created virus kills all other agents unless some of them also create a new virus and shield before the virus reaches them. Since p is sufficiently small, the probability of the latter is small. Thus, the unique leader is elected within a relatively short time. Even after that, the unique leader keeps on creating new viruses, each of which may kill the leader. However, the leader is not killed for an extremely long time: since $t_{\rm shld} \gg t_{\rm virus}$, the leader's shield rarely vanishes before all viruses vanish from the population.

Complexity Analysis We have four parameters in $P_{\rm RD}$: three upper bounds $t_{\rm max}$, $t_{\rm virus}$, and $t_{\rm shld}$ of the timers, and probability p. We assume $t_{\rm virus} = t_{\rm max}/2$, $t_{\rm max} \ge 8\delta \max(d, 2 + \log(13n\delta \lceil \log n \rceil))$, $t_{\rm shld} \ge 2\delta t_{\rm max} \lceil \log n \rceil$ and $p \le 1000$

 $(4mt_{\rm shld})^{-1}$. We prove the following equations under this assumption:

$$\max_{C \in \mathcal{C}_{all}} \text{ECT}_{P_{\text{RD}}}(C, \mathcal{S}_{\text{RD}}) = O(mp^{-1}), \tag{3}$$

$$\min_{C \in \mathcal{S}_{\mathrm{RD}}} \operatorname{EHT}_{P_{\mathrm{RD}}}(C, LE) = \Omega(\tau e^{\tau}), \tag{4}$$

where $\tau = t_{\text{max}}/(8\delta)$ and S_{RD} is the set of configurations we define later. When upper bounds N and Δ are available and we assign $t_{\text{max}} = 8N\Delta$, $t_{\text{shld}} = 2\Delta t_{\text{max}} \lceil \log N \rceil$ and $p = (4N^2 t_{\text{shld}})^{-1}$ (*i.e.*, $R = [0, 4N^2 t_{\text{shld}} - 1]$), then P_{RD} is an $(O(m\Delta^2 N^3 \log N), \Omega(Ne^N))$ -loosely-stabilizing leader election protocol.

Before proving equations (3) and (4), we define five sets of configurations:

$$\begin{split} \mathcal{G}_{\text{half}} &= \{ C \in \mathcal{C}_{\text{all}}(P_{\text{RD}}) \mid \exists v \in V, \ C(v).\texttt{DoA} = \text{ALIVE} \land C(v).\texttt{timer}_{\texttt{s}} > t_{\text{shld}}/2 \}, \\ \mathcal{V}_{\text{clean}} &= \{ C \in \mathcal{C}_{\text{all}}(P_{\text{RD}}) \mid \forall v \in V, \ C(v).\texttt{timer}_{\texttt{v}} = 0 \}, \\ \mathcal{L}_{\text{half}} &= \{ C \in \mathcal{C}_{\text{all}}(P_{\text{RD}}) \mid \#_L(C) \geq 1 \land \forall v \in V, \ C(v).\texttt{timer}_{\texttt{L}} > t_{\text{max}}/2 \}, \\ \mathcal{L}_{\text{one}} &= \{ C \in \mathcal{C}_{\text{all}}(P_{\text{RD}}) \mid \#_L(C) = 1 \}, \\ \mathcal{S}_{\text{RD}} &= (\mathcal{G}_{\text{half}} \cup \mathcal{V}_{\text{clean}}) \cap \mathcal{L}_{\text{half}} \cap \mathcal{L}_{\text{one}}, \end{split}$$

where $\#_L(C)$ denotes the number of leaders in configuration C. Note that \mathcal{G}_{half} requires that not all agents but at least one leader has timers more than $t_{shld}/2$.

First, we analyze the expected holding time. Let $C_0 \in \mathcal{S}_{\text{RD}}$ and $\mathcal{Z}_{P_{\text{RD}}}(C_0) = C_0, C_1, \ldots$ To prove (4), it suffices to show that both $C_0, \ldots, C_{8m\delta\tau \lceil \log n \rceil} \in LE$ and $C_{8m\delta\tau \lceil \log n \rceil} \in \mathcal{S}_{\text{RD}}$ hold with probability no less than $p_{\text{suc}} = 1 - O(n\delta \log n \cdot e^{-\tau})$. Then, $\min_{C_0 \in \mathcal{S}_{\text{RD}}} \text{EHT}_{P_{\text{RD}}}(C_0, LE) \geq 8m\delta\tau \lceil \log n \rceil \tau/(1 - p_{\text{suc}}) = \Omega(\tau e^{\tau})$.

We define two predicates PROP_i and HALF_i for any $i \geq 0$: $\operatorname{PROP}_i = 1$ if $C_{2m\tau(i+1)}(v)$.timer_L > $t_i - t_{\max}/2$ for all $v \in V$, otherwise $\operatorname{PROP}_i = 0$, where $t_i = \max_{v \in V} C_{2m\tau i}(v)$; $\operatorname{HALF}_i = 1$ if every agent joins only less than $t_{\max}/2$ interactions among $\Gamma_{2m\tau i}, \ldots, \Gamma_{2m\tau(i+1)-1}$, otherwise $\operatorname{HALF}_i = 0$. Intuitively, $\operatorname{PROP}_i = 1$ means the maximum value of timer_L propagates to all the agents well during the $2m\tau$ interactions, and $\operatorname{HALF}_i = 1$ means every agent does not interact so much during the $2m\tau$ interactions.

Lemma 11. Let $C_0 \in S_{\text{RD}}$ and $\Xi_{P_{\text{RD}}}(C_0) = C_0, C_1, \ldots$ Then, we have both $C_0, \ldots, C_{8m\delta\tau \lceil \log n \rceil} \in LE$ and $C_{8m\delta\tau \lceil \log n \rceil} \in S_{\text{RD}}$ if the following conditions hold:

(A) $\#_L(C_t) \ge 1$ for all $t = 0, \dots, 8m\delta\tau \lceil \log n \rceil$, (B) $C_{8m\delta\tau \lceil \log n \rceil} \in \mathcal{G}_{half} \cup \mathcal{V}_{clean}$, (C) $\operatorname{PROP}_i = 1$ for all $i = 0, \dots, 4\delta \lceil \log n \rceil - 1$, and (D) $\operatorname{HALF}_i = 1$ for all $i = 0, \dots, 4\delta \lceil \log n \rceil - 1$.

Proof. We have $C_{2m\tau i}(v)$.timer_L > $t_{\max}/2$ for any $i \in [0, 4\delta \lceil \log n \rceil]$ from (A) and (C). Since no agent interacts more than $t_{\max}/2$ times among each $2m\tau$ interactions (i.e. (D)), timeout does not occur at any interaction $\Gamma_0, \ldots, \Gamma_{8m\delta\tau \lceil \log n \rceil - 1}$, by which we obtain $C_0, \ldots, C_{8m\delta\tau \lceil \log n \rceil} \in LE$. We also obtain $C_{8m\delta\tau \lceil \log n \rceil} \in \mathcal{L}_{half} \cap \mathcal{L}_{one} \cap (\mathcal{G}_{half} \cup \mathcal{V}_{clean}) = \mathcal{S}_{RD}$ from above discussion and (B).

Lemma 12. The probability that every agent joins only less than $t_{\text{shld}}/2$ interactions as an initiator among $\Gamma_0, \ldots, \Gamma_{8m\delta\tau \lceil \log n \rceil - 1}$ is at least $1 - ne^{-\delta\tau}$.

Proof. For any $v \in V$ and $t \geq 0$, v joins interaction Γ_t as an initiator with probability at most $\delta/(2m)$ since v has at most $\delta/2$ outgoing edges. Thus, the number of interactions v joins as an initiator during the $8m\delta\tau\lceil\log n\rceil$ interactions is bounded by binomial random variable $X \sim B(8m\delta\tau\lceil\log n\rceil, \delta/(2m))$. We have

$$\begin{aligned} \Pr(X \ge t_{\text{shld}}/2) &\leq \Pr(X \ge 8\delta^2 \tau \lceil \log n \rceil) & \because t_{\text{shld}} \ge 16\delta^2 \tau \lceil \log n \rceil \\ &= \Pr(X \ge 2\mathbf{E}[X]) \\ &\leq e^{-\mathbf{E}[X]/3} \quad \text{(By Chernoff Bound of Lemma 2)} \\ &= e^{-4\delta^2 \tau \lceil \log n \rceil/3} \\ &= e^{-\delta\tau}. \end{aligned}$$

Summing up these probabilities gives the lemma.

Lemma 13. Let $C_0 \in \mathcal{S}_{\text{RD}}$ and $\Xi_{P_{\text{RD}}}(C_0) = C_0, C_1, \ldots$. Then, we have $\Pr(\forall t \in [0, 8m\delta\tau \lceil \log n \rceil - 1], \ \#_L(C_t) \ge 1) \ge 1 - ne^{-\delta\tau}$.

Proof. By Lemma 12, it suffices to show that $\#_L(C_t) \geq 1$ holds for all $t \in [0, 8m\delta\tau \lceil \log n \rceil]$ when we assume every agent joins only less than $t_{\text{shld}}/2$ interactions as an initiator among $\Gamma_0, \ldots, \Gamma_{8m\delta\tau \lceil \log n \rceil - 1}$. Since $C_0 \in \mathcal{S}_{\text{RD}}$, we have $C_0 \in \mathcal{G}_{\text{half}} \cup \mathcal{V}_{\text{clean}}$. If $C_0 \in \mathcal{G}_{\text{half}}$, there exists a leader v such that $C_0(v)$.timers $> t_{\text{shld}}/2$. By the assumption, v decrease its timers by at most $t_{\text{shld}}/2$; thus, v is never killed and remains a leader in $C_0, \ldots, C_{8m\delta\tau \lceil \log n \rceil}$. If $C_0 \in \mathcal{V}_{\text{clean}}$, no leader is killed before a new virus is created. Even if some leader u creates a new virus at interaction Γ_t $(0 \leq t < 8m\delta\tau \lceil \log n \rceil)$, u wears a new shield at the same time. Hence, u remains a leader in $C_t, \ldots, C_{8m\delta\tau \lceil \log n \rceil}$ by the assumption.

We define the first round time $\operatorname{RT}_{\Gamma}(1)$ as the minimum t satisfying $\forall e \in E, 0 \leq \exists t' \leq t, \Gamma_{t'} = e$. For any $i \geq 2$, we define the *i*-th round time $\operatorname{RT}_{\Gamma}(i)$ as the minimum t satisfying $\forall e \in E, \operatorname{RT}_{\Gamma}(i-1) < \exists t' \leq t, \Gamma_{t'} = e$. Lemma 15 bounds $\operatorname{RT}_{\Gamma}(i)$ from above with high probability. To prove the lemma, we firstly prove Lemma 14.

Lemma 14. Let v_1, v_2, \ldots, v_l be any l (l < n) agents in V. There exists at least 2l edges of E that are incident to at least one of the l agents.

Proof. Since l < n, there exists agent $r \in V$ that differs from any v_1, v_2, \ldots, v_l . Since G is strongly connected, there exists a rooted spanning tree T on G where r is the root agent of T. Then, every v_i $(i \in [1, k])$ has two edges between v_i and the parent agent of v_i in T. (Remind that G is undirected, that is, $(u, v) \in E \Leftrightarrow (v, u) \in E$ for any $u, v \in V$.) These edges are mutually exclusive. Thus, we have 2l edges of E that are incident to at least one of the l agents.

Lemma 15. $\Pr(\operatorname{RT}_{\Gamma}(i) < 2im \lceil \log n \rceil) \ge 1 - ne^{-i/4}$ holds for any $i \ge 1$.

Proof. Each round j ($j \ge 1$) finishes when every agent $v \in V$ interacts in round j. Consider the case that k ($k \ge 1$) agents have not yet interacted in round j and

only n-k agents have interacted in round j. We call the former uninvolved agents and the latter involved agents. If k < n, one of the k uninvolved agents joins the next interaction and becomes an involved agent with probability more than 2k/mby Lemma 14. If k = n, some uninvolved agent joins the next interaction with probability 1. Let $X_{j,k}$ $(j \ge 1, k \ge 1)$ be the random variable that corresponds to the number of trials to the first success in which the success probability of each trial is 2k/m. From the above discussion, we obtain

$$\Pr(\operatorname{RT}_{\Gamma}(i) \ge 2im\lceil \log n \rceil) \le \Pr\left(\sum_{j=1}^{i} \left(1 + \sum_{k=1}^{n-1} X_{j,k}\right) \ge 2im\lceil \log n \rceil\right)$$

$$\le \Pr\left(\sum_{k=1}^{n-1} \sum_{j=1}^{i} X_{j,k} \ge 2im\lceil \log n \rceil - i\right).$$
(5)

For binomial random variable $Y_k \sim B(\lceil \frac{im}{k} \rceil, \frac{2k}{m})$, we have $\Pr(\sum_{j=1}^i X_{j,k} > \frac{im}{k}) \leq \Pr(\sum_{j=1}^i X_{j,k} \geq \lceil \frac{im}{k} \rceil) \leq \Pr(Y_k \leq i)$. Hence, we have

$$\Pr\left(\sum_{j=1}^{i} X_{j,k} > \frac{im}{k}\right) \leq \Pr(Y_k \leq i)$$

$$\leq \Pr\left(Y_k \leq \frac{1}{2} \cdot \mathbf{E}[Y_K]\right)$$

$$\leq e^{-\mathbf{E}[Y_k]/8} \quad (By \text{ Chernoff Bound of Lemma 2})$$

$$\leq e^{-i/4}.$$
(6)

From Inequalities (5) and (6), we have

$$\Pr(\mathrm{RT}_{\Gamma}(i) \ge 2im\lceil \log n \rceil) \le \Pr\left(\sum_{k=1}^{n-1} \sum_{j=1}^{i} X_{j,k} \ge 2im\lceil \log n \rceil - i\right)$$
$$\le \Pr\left(\sum_{k=1}^{n-1} \sum_{j=1}^{i} X_{j,k} > \sum_{k=1}^{n-1} \frac{im}{k}\right)$$
$$\le \sum_{k=1}^{n-1} \Pr\left(\sum_{j=1}^{i} X_{j,k} > \frac{im}{k}\right)$$
$$\le ne^{-i/4},$$

where $\sum_{k=1}^{n-1} \frac{im}{k} \leq im(1 + \log n) - i < 2im\lceil \log n \rceil - i$ is used for the second inequality. Thus, $\Pr(\operatorname{RT}_{\Gamma}(i) < 2im\lceil \log n \rceil) \geq 1 - ne^{-i/4}$ holds. \Box

Lemma 16. Let $C_0 \in \mathcal{S}_{\text{RD}}$ and $\Xi_{P_{\text{RD}}}(C_0) = C_0, C_1, \ldots$. Then, we have $\Pr(C_{8m\delta\tau \lceil \log n \rceil} \in \mathcal{G}_{\text{half}} \cup \mathcal{V}_{\text{clean}}) \ge 1 - 2ne^{-\delta\tau}$. **Proof.** Assume that $\operatorname{RT}_{\Gamma}(t_{\operatorname{virus}}) < 8m\delta\tau \lceil \log n \rceil$ holds and every agent joins only less than $t_{\operatorname{shld}}/2$ interactions as an initiator among $\Gamma_0, \ldots, \Gamma_{8m\delta\tau \lceil \log n \rceil - 1}$. These assumptions lead to $C_{8m\delta\tau \lceil \log n \rceil} \in \mathcal{G}_{\operatorname{half}} \cup \mathcal{V}_{\operatorname{clean}}$ as follows. If a new virus is not created among $\Gamma_0, \ldots, \Gamma_{8m\delta\tau \lceil \log n \rceil - 1}$, then all viruses in the initial configuration vanish during the period since each round decreases the maximum value of timerv by at least one. Thus, $C_{8m\delta\tau \lceil \log n \rceil} \in \mathcal{V}_{\operatorname{clean}}$ holds. If some agent v creates a new virus at Γ_t , then v wears a new shield at the same time. Thus, $C_{t+1}(v)$.timers = t_{shld} . Since v interacts as an initiator only less than $t_{\operatorname{shld}}/2$ times among $\Gamma_{t+1}, \ldots, \Gamma_{8m\delta\tau \lceil \log n \rceil - 1}$, we have $C_{8m\delta\tau \lceil \log n \rceil}(v)$.timers > $t_{\operatorname{shld}}/2$, which means $C_{8m\delta\tau \lceil \log n \rceil} \in \mathcal{G}_{\operatorname{half}}$. By $t_{\operatorname{virus}} = 4\delta\tau$ and Lemmas 12 and 15, the probability that the two assumptions hold is at least $1 - 2ne^{-\delta\tau}$.

Lemma 17. $Pr(PROP_i = 1) \ge 1 - 2ne^{-\tau}$ for any $i \ge 0$.

Proof. The same argument as the proof of Lemma 5 gives the lemma. \Box

Lemma 18. $\Pr(\text{HALF}_i = 1) \ge 1 - ne^{-\tau}$ for any $i \ge 0$.

Proof. Each interaction is independent. Thus, Lemma 4 gives the lemma. \Box

Lemma 19. $\min_{C \in \mathcal{S}_{RD}} EHT_{P_{RD}}(C, LE) = \Omega(\tau e^{\tau}).$

Proof. Probability p_{suc} , discussed in the beginning of this section, is at least $1 - 3ne^{-\delta\tau} - 4\delta \lceil \log n \rceil \cdot 3ne^{-\tau} \ge 1 - 13n\delta \lceil \log n \rceil e^{-\tau}$ by Lemmas 11, 13, 16, 17 and 18, which leads to the lemma.

Next, we analyze the expected convergence time. We define two sets of configurations: $\mathcal{N}_{oVG} = \{C \in \mathcal{C}_{all}(P_{RD}) \mid \forall v \in V, \ C(v).\texttt{timer}_{V} = C(v).\texttt{timer}_{S} = 0\}$ and $\mathcal{L} = \{C \in \mathcal{C}_{all}(P_{RD}) \mid \#_{L}(C) \geq 1\}.$

Lemma 20. $\max_{C \in \mathcal{C}_{all}(P_{RD})} ECT_{P_{RD}}(C, \mathcal{S}_{RD}) = O(mp^{-1}).$

Proof Sketch. Probability *p*, with which a leader creates a virus at each interaction, is sufficiently small $(p < 1/(4mt_{shld}))$. Thus, the probability that all viruses and shields vanish (i.e. the population enters a configuration of \mathcal{N}_{oVG}) within $2mt_{\text{shld}}$ interactions is at least $1 - (2mt_{\text{shld}} \cdot p + O(ne^{-\tau})) > 1/2 - O(ne^{-\tau})$. Even if the reached configuration of \mathcal{N}_{oVG} does not have any leader, the timeout mechanism creates a leader, and the population enters a configuration of $\mathcal{N}_{oVG} \cap$ \mathcal{L} . This takes less than $16m\delta\tau \lceil \log n \rceil$ interactions with probability $1 - O(ne^{-\tau})$. After the population enters into $\mathcal{N}_{oVG} \cap \mathcal{L}$, additional [m/p] interactions create a new virus with probability $1 - e^{-2}$. Let v be a leader that creates the virus. Since v wears a new shield at the same time, v is not killed and remains a leader during the next $2m\tau$ interactions with probability $1 - O(e^{-\tau})$. On the other hand, the virus spreads to all the agents within these $2m\tau$ interactions with probability $1 - O(ne^{-\tau})$, killing all the agents other than v. A leader other than v may create a new virus during the $2m\tau$ interactions, and survives with a shield. However, this probability is at most $2m\tau \cdot p < 1/4$. Hence, v becomes the unique leader within the $2m\tau$ interactions with probability $3/4 - O(ne^{-\tau})$. After the $2m\tau$ interactions,

all the agents have $\mathtt{timer}_{L} > t_{\max}/2$ with probability $1 - O(ne^{-\tau})$ by the larger value propagation, and $v.\mathtt{timer}_{S} > t_{\mathrm{shld}}/2$ holds with probability $1 - O(ne^{-\tau})$. Hence, the population enters a configuration of $\mathcal{L}_{\mathrm{one}} \cap \mathcal{L}_{\mathrm{half}} \cap \mathcal{G}_{\mathrm{half}} \subset \mathcal{S}_{\mathrm{RD}}$ within the $2m\tau$ interactions with probability $3/4 - O(ne^{-\tau})$. As a result, starting from any configuration, the population enters into $\mathcal{S}_{\mathrm{RD}}$ within $O(mp^{-1})$ interactions with probability $1/4 - e^{-2} - O(ne^{-\tau}) > 0.11 - o(1)$, which gives the lemma. \Box

Lemmas 19 and 20 gives the following theorem.

Theorem 2 Protocol P_{RD} is a $(O(mp^{-1}), \Omega(\tau e^{\tau}))$ loosely-stabilizing leader election protocol for arbitrary graphs when $t_{\text{max}} \geq 8\delta \max(d, 2 + \log(13n\delta \lceil \log n \rceil))$, $t_{\text{virus}} = t_{\text{max}}/2$, $t_{\text{shld}} \geq 2\delta t_{\text{max}} \lceil \log n \rceil$ and $p \leq (4mt_{\text{shld}})^{-1}$.

Therefore, given upper bound N and Δ of n and δ respectively, we get a $(O(m\Delta^2 N^3 \log N), \Omega(Ne^N))$ loosely-stabilizing leader election protocol for arbitrary graphs by assigning $t_{\text{max}} = 8N\Delta$, $t_{\text{virus}} = t_{\text{max}}/2$, $t_{\text{shld}} = 2\Delta t_{\text{max}} \lceil \log N \rceil$ and $p = (4N^2 t_{\text{shld}})^{-1}$.

5 Conclusion

We have presented two loosely-stabilizing leader election protocols for arbitrary undirected graphs in the PP model: one works with agent-identifiers and the other works with random numbers. Both protocols keep a unique leader for an exponentially long expected time after entering a loosely-safe configuration. The protocols use only upper bounds N of n and Δ of δ while any self-stabilizing leader election protocol needs the exact knowledge of n. The restriction of the protocols to *undirected* graph is only for simplicity of protocol description and complexity analysis. The proposed protocols also work on arbitrary *directed* graphs with slight modification: it is only necessary that a responder also executes some actions of an initiator (Line 1 of Protocol 1 and Lines 10-12 of Protocol 2). Our future work is to develop a loosely-stabilizing leader election protocol without agent-identifiers or random numbers for arbitrary graphs. We will also tackle with loosely-stabilizing leader election for some classes of graphs (e.g. rings and trees). We are also interested in the empirical evaluation of the holding time of loosely-stabilizing protocols. Since our probabilistic evaluation of the holding time in this paper is not tight, the actual holding time of the protocols should be much longer. By simulation experiments, we will empirically evaluate the actual holding time (and convergence time) for various network sizes and graph topologies.

Acknowledgements. This work was supported by JSPS KAKENHI Grant Numbers 24500039, 24650012, 25104516, 26280022, and 26330084.

References

 Angluin, D., Aspnes, J., Diamadi, Z., Fischer, M.J., Peralta, R.: Computation in networks of passively mobile finite-state sensors. Distributed Computing 18(4), 235–253 (2006)

- Angluin, D., Aspnes, J., Eisenstat, D.: Fast computation by population protocols with a leader. In: Dolev, S. (ed.) DISC 2006. LNCS, vol. 4167, pp. 61–75. Springer, Heidelberg (2006)
- Angluin, D., Aspnes, J., Fischer, M.J., Jiang, H.: Self-stabilizing population protocols. ACM Transactions on Autonomous and Adaptive Systems 3(4), 13 (2008)
- Beauquier, J., Blanchard, P., Burman, J.: Self-stabilizing leader election in population protocols over arbitrary communication graphs. In: Baldoni, R., Nisse, N., van Steen, M. (eds.) OPODIS 2013. LNCS, vol. 8304, pp. 38–52. Springer, Heidelberg (2013)
- Beauquier, J., Burman, J., Rosaz, L., Rozoy, B.: Non-deterministic population protocols. In: Baldoni, R., Flocchini, P., Binoy, R. (eds.) OPODIS 2012. LNCS, vol. 7702, pp. 61–75. Springer, Heidelberg (2012)
- Cai, S., Izumi, T., Wada, K.: How to prove impossibility under global fairness: On space complexity of self-stabilizing leader election on a population protocol model. Theory of Computing Systems 50(3), 433–445 (2012)
- 7. Canepa, D., Potop-Butucaru, M.G.: Stabilizing leader election in population protocols (2007), http://hal.inria.fr/inria-00166632
- Fischer, M., Jiang, H.: Self-stabilizing leader election in networks of finite-state anonymous agents. In: Shvartsman, A. (ed.) OPODIS 2006. LNCS, vol. 4305, pp. 395–409. Springer, Heidelberg (2006)
- 9. Guerraoui, R., Ruppert, E.: Even small birds are unique: Population protocols with identifiers. Rapport de Recherche CSE-2007-04, Department of Computer Science and Engineering, York University, York, ON, Canada (2007)
- Michail, O., Chatzigiannakis, I., Spirakis, P.G.: Mediated population protocols. Theoretical Computer Science 412(22), 2434–2450 (2011)
- 11. Mitzenmacher, M., Upfal, E.: Probability and Computing: Randomized Algorithms and Probabilistic Analysis. Cambridge University Press (2005)
- Mizoguchi, R., Ono, H., Kijima, S., Yamashita, M.: On space complexity of selfstabilizing leader election in mediated population protocol. Distributed Computing 25(6), 451–460 (2012)
- Sudo, Y., Nakamura, J., Yamauchi, Y., Ooshita, F., Kakugawa, H., Masuzawa, T.: Loosely-stabilizing leader election in a population protocol model. Theoretical Computer Science 444, 100–112 (2012)
- Xu, X., Yamauchi, Y., Kijima, S., Yamashita, M.: Space complexity of selfstabilizing leader election in population protocol based on k-interaction. In: Higashino, T., Katayama, Y., Masuzawa, T., Potop-Butucaru, M., Yamashita, M. (eds.) SSS 2013. LNCS, vol. 8255, pp. 86–97. Springer, Heidelberg (2013)