

Stone Age Distributed Computing

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Abstract

The traditional models of distributed computing focus mainly on networks of computer-like devices that can exchange large messages with their neighbors and perform arbitrary local computations. Recently, there is a trend to apply distributed computing methods to networks of sub-microprocessor devices, e.g., biological cellular networks or networks of nano-devices. However, the suitability of the traditional distributed computing models to these types of networks is questionable: do tiny bio/nano nodes “compute” and/or “communicate” essentially the same as a computer? In this paper, we introduce a new model that depicts a network of randomized finite state machines operating in an asynchronous environment. Although the computation and communication capabilities of each individual device in the new model are, by design, much weaker than those of a computer, we show that some of the most important and extensively studied distributed computing problems can still be solved efficiently.

1 Introduction

Networks are at the core of many scientific areas, be it social sciences (where networks for instance model human relations), logistics (e.g. traffic), or electrical engineering (e.g. circuits). *Distributed computing* is the area that studies the power and limitations of distributed algorithms and computation in networks. Due to the major role that the Internet plays today, models targeted at understanding the fundamental properties of networks focus mainly on “Internet-capable” devices. The standard model in distributed computing is the so called *message passing* model, where nodes may exchange large messages with their neighbors, and perform arbitrary local computations.

Some networks though, are not truthfully represented by the classical message passing model. For example, *wireless* networks such as ad hoc or sensor networks, whose research has blossomed in the last decade, require some adaptations of the message passing model so that it meets the limited capabilities of the underlying wireless devices more precisely. More recently, there is a trend to apply distributed computing methods, and in particular, the message passing model, to networks of sub-microprocessor devices, for instance networks of biological cells or nano-scale mechanical devices. However, the suitability of the message passing model to these types of networks is far from being certain: do tiny bio/nano nodes “compute” and/or “communicate” essentially the same as a computer? Since such nodes will be fundamentally more limited than silicon-based devices, we believe that there is a need for a network model, where nodes are by design below the computation and communication capabilities of Turing machines.

Networked Finite State Machines. In this paper, we take a radically different approach: Instead of imposing additional restrictions on the existing models for networks of computer-like devices, we introduce an entirely new model, referred to as *networked finite state machines (nFSM)*, that depicts a network of randomized finite state machines progressing in asynchronous steps (refer to Section 2 for a formal description). Under the nFSM model, nodes communicate by transmitting messages belonging to some finite communication alphabet Σ such that a message $\sigma \in \Sigma$ transmitted by node u is delivered to its neighbors (the same σ to all neighbors) in an asynchronous fashion; each neighbor v of u has a port corresponding to u in which the last message delivered from u is stored.

The access of node v to its ports is limited: each state q in the state set Q of the FSM is associated with some *query letter* $\sigma = \sigma(q) \in \Sigma$; if node v resides in state q at some step of the execution, then the next state and the message transmitted by v at this step are determined by q and by the number $\sharp(\sigma)$ of occurrences of σ in v ’s ports. The crux of the model is that $\sharp(\sigma)$ is calculated according to the *one-two-many*¹ principle: the node can only count up to some predetermined

¹ The one-two-many theory states that some small isolated cultures (e.g., the Piraha tribe of the Amazon [20]) did not develop a counting system that goes beyond 2. This is reflected in their languages that include words for “1”, “2”, and “many” that stands for any number larger than 2.

bounding parameter $b \in \mathbb{Z}_{>0}$ and any value of $\sharp(\sigma)$ larger than b cannot be distinguished from b .

In particular, the nFSM model satisfies the following *model requirements*, that we believe, make it more applicable to the study of networks consisting of weaker devices such as those mentioned above.

(M1) The model is applicable to arbitrary network topologies.

(M2) All nodes run the same protocol executed by a (randomized) FSM.

(M3) The network operates within an asynchronous environment, with node activation patterns independent of message delivery patterns.

(M4) All features of the FSM (specifically, the state set Q , message alphabet Σ , and bounding parameter b) are of constant size independent of any parameter of the network (including the degree of the node executing the FSM).

The last requirement is perhaps the most interesting one as it implies that a node cannot perform any calculation that involves numbers beyond some predetermined constant. This comes in contrast to many distributed algorithms operating under the message passing model that strongly rely on the ability of a node to perform such calculations (e.g., count up to some parameter of the network or a function thereof).

Results. Our investigation of the new model begins by implementing an nFSM synchronizer that practically allows the algorithm designer to assume a synchronous environment (Section 3). Then, we show that the computational power of a network operating under the nFSM model is essentially equivalent to that of a randomized Turing machine with linear space bound (cf. linear bounded automaton). In comparison, the computational power of a network operating under the message passing model is trivially equivalent to that of a (general) Turing machine, therefore there exist distributed problems that can be solved under the message passing model in constant time but cannot be solved under the nFSM model at all (Section 6).

Nevertheless, we show that arguably the most important and extensively studied problems in distributed computing admit efficient — namely, with run-time polylogarithmic in the number of nodes — algorithms operating under the nFSM model. Specifically, we develop such algorithms for computing a maximal independent set (MIS) in arbitrary graphs (Section 4) and for 3-coloring of (undirected) trees (Section 5). We also develop an efficient algorithm that computes a maximal matching in arbitrary graphs, but this requires a small unavoidable modification of the nFSM model that goes beyond the scope of the current version of the paper.

Related Work. As mentioned above, the message passing model is the gold standard when it comes to understanding distributed algorithms. Several variants exist for this model, differing mainly in the bounds imposed on the message size and the level of synchronization. Perhaps the most popular message passing variants are the fully synchronous *local* and *congest* models [26, 31, 36], assuming that in each round, a node can send messages to its neighbors (different

messages to different neighbors), receive and interpret the messages sent to it from its neighbors, and perform an arbitrary local computation² determining, in particular, the messages sent in the next round. The difference between the two variants is cast in the size of the communicated messages: the local model does not impose any restrictions on the message size, hence it can be used for the purpose of establishing general lower bounds, whereas the congest model is more information-theoretic, with a (typically logarithmic) bound on the message size. Indeed, most theoretical literature dealing with distributed algorithms relies on one of these two models.

As the congest model still allows for sending different messages to different neighbors in each round, it was too powerful for many settings. Instead, with the proliferation of wireless networks, new more restrictive message passing models appeared such as the *radio network* model [13]. In radio networks, nodes still operate in synchronous rounds, where in each round a node may choose to transmit a message or stay silent. A transmitted message is received by all neighbors in the network if the neighbors do not experience interference by concurrently transmitting nodes in their own neighborhood. There are several variants, e.g. whether nodes have collision detection, or not.

Since the radio network model is still too powerful for some wireless settings, more restrictive models were suggested. One such example is the *beeping* model [17, 16], where in each round a node can either beep or stay silent, and a silent node can only distinguish between the case in which no node in its neighborhood beeps and the case in which at least one node beeps. Efficient algorithms and lower bounds for the MIS problem under the beeping model were developed by Afek et al. [2, 1]. Note that the beeping model resembles our nFSM model in the sense that the “beeping rule” can be viewed as counting under the one-two-many principle with bounding parameter $b = 1$. However, it is much stronger in other perspectives: (i) the beeping model assumes synchronous communication and does not seem to have a natural asynchronous variant, thus it does not satisfy requirement (M3); and (ii) the local computation is performed by a Turing machine whose memory is allowed to grow with the network (this is crucial for the algorithms of Afek et al. [2, 1]), thus it does not satisfy requirements (M2) and (M4).

Our nFSM model is a generalization of the extensively studied *cellular automaton* model [30, 18, 38] that captures a network of FSMs, arranged in a grid topology (some other highly regular topologies were also considered), where the transition of each node depends on its current state and the states of its neighbors. Still, the nFSM model differs from the cellular automaton model in many aspects; in particular, the latter model is not applicable for non-regular network topologies, in contrast to requirement (M1), and to the most part, it also does not support asynchronous environments (at least not as asynchrony is grasped in the current paper), in contrast to requirement (M3).

Another model that resembles the nFSM model is that of *communicating automata* [12]. This

² It is important to point out that even though the local and congest models allow for arbitrary local computations, the existing literature hardly ever assumes anything that cannot be computed in time polynomial in the size of the information received thus far; the rare exceptions are typically clearly mentioned in the text.

model also assumes that each node in the network operates a FSM in an asynchronous manner, however the steps of the FSMs are message driven: for each state q of node v and for each message m that node v may receive from an adjacent node u while residing in state q , the transition function of v should have an entry characterized by the 3-tuple (q, u, m) that determines its next move. As such, different nodes would typically operate different FSMs, hence the model does not satisfy requirement (M2), and more importantly, the size of the FSM operated by node v inherently depends on the degree of v , hence it does not satisfy requirement (M4). Moreover, the node activation pattern is driven by the incoming messages, so it also does not satisfy requirement (M3).

Applicability to Biological Cellular Networks. Regardless of the theoretical interest in implementing efficient algorithms using weaker assumptions, we believe that our new model and results should be appealing to anyone interested in understanding the computational aspects of biological cellular networks. A basic dogma in biology (see, e.g., [33]) states that all cells communicate and that they do so by emitting special kinds of proteins (e.g., cytokines and chemokines in the immune system) that can be recognized by designated receptors, thus enabling neighboring cells to distinguish between different concentration levels of these proteins, which, after a signaling cascade, leads to different gene expression.

Translated to the language of the nFSM model, the emitted proteins correspond to the letters of the communication alphabet, where the actual emission corresponds to transmitting a letter, and the ability of a cell to distinguish between different concentration levels of these proteins corresponds to the manner in which the nodes in our model interpret the content of their ports. Using an FSM as the underlying computational model of each node seems to be the right choice especially in the biological setting as demonstrated by Benenson et al. [11] who showed that essentially any FSM can be implemented by enzymes found in cells' nuclei. One may wonder if the specific problems studied in the current paper have any relevance to biological cellular networks. Indeed, Afek et al. [2] discovered that a biological process that occurs during the development of the nervous system of a fly is in fact equivalent to solving the MIS problem.

2 Model

Throughout, we assume a network represented by a finite undirected graph $G = (V, E)$. Under the *networked finite state machines (nFSM)* model, each node $v \in V$ runs a protocol depicted by the 8-tuple

$$\Pi = \langle Q, Q_I, Q_O, \Sigma, \sigma_0, b, \lambda, \delta \rangle,$$

where

- Q is a finite set of *states*;

- $Q_I \subseteq Q$ is the subset of *input states*;
- $Q_O \subseteq Q$ is the subset of *output states*;
- Σ is a finite *communication alphabet*;
- $\sigma_0 \in \Sigma$ is the *initial letter*;
- $b \in \mathbb{Z}_{>0}$ is a *bounding parameter*; let $B = \{0, 1, \dots, b-1, \geq b\}$ be a set of $b+1$ distinguishable symbols;
- $\lambda : Q \rightarrow \Sigma$ assigns a *query letter* $\sigma \in \Sigma$ to every state $q \in Q$; and
- $\delta : Q \times B \rightarrow 2^{Q \times (\Sigma \cup \{\varepsilon\})}$ is the *transition function*.

It is important to point out that protocol Π is oblivious to the graph G . In fact, the number of states in Q , the size of the alphabet Σ , and the bounding parameter b are all assumed to be universal constants, independent of any parameter of the graph G . In particular, the protocol executed by node $v \in V$ does not depend on the degree of v in G . We now turn to describe the semantics of the nFSM model.

Communication. Node v communicates with its adjacent nodes in G by *transmitting* messages. A transmitted message consists of a single letter $\sigma \in \Sigma$ and it is assumed that this letter is delivered to all neighbors u of v . Each neighbor u has a *port* $\psi_u(v)$ (a different port for every adjacent node v) in which the last message σ received from v is stored. At the beginning of the execution, all ports store the initial letter σ_0 . It will be convenient to consider the case in which v does not transmit any message (and hence does not affect the corresponding ports of the adjacent nodes) as a transmission of the special *empty symbol* ε .

Execution. The execution of node v progresses in discrete *steps* indexed by the positive integers. At each step $t \in \mathbb{Z}_{>0}$, v resides in some state $q \in Q$. Let $\lambda(q) = \sigma \in \Sigma$ be the query letter that λ assigns to state q and let $\sharp(\sigma)$ be the number of occurrences of σ in v 's ports in step t . Then, the pair (q', σ') of state $q' \in Q$ in which v resides in step $t+1$ and message $\sigma' \in \Sigma \cup \{\varepsilon\}$ transmitted by v in step t (recall that ε indicates that no message is transmitted) is chosen *uniformly at random* (and independently of all other random choices) among the pairs in

$$\delta(q, f_b(\sharp(\sigma))) \subseteq Q \times (\Sigma \cup \{\varepsilon\}) ,$$

where $f_b : \mathbb{Z}_{\geq 0} \rightarrow B$ is defined as

$$f_b(x) = \begin{cases} x & \text{if } 0 \leq x \leq b-1 ; \\ \geq b & \text{otherwise .} \end{cases}$$

Informally, this can be thought of as if v queries its ports for occurrences of σ and “observes” the exact value of $\sharp(\sigma)$ as long as it is smaller than the bounding parameter b ; otherwise, v merely “observes” that $\sharp(\sigma) \geq b$ which is indicated by the symbol $\geq b$.

Input and Output. Initially (in step 1), each node resides in some of the input states in Q_I . The choice of the initial state of node $v \in V$ reflects the input passed to v at the beginning of the execution. This allows our model to cope with distributed problems in which different nodes get different input symbols. When dealing with problems in which the nodes do not get any initial input (such as the graph theoretic problems addressed in this paper), we shall assume that Q_I contains a single *initial* state.

We say that the (global) execution of the protocol is in an *output configuration* if all nodes reside in output states of Q_O . If this is the case, then the output of node $v \in V$ is determined by the output state $q \in Q_O$ in which v resides.

Asynchrony. The nodes are assumed to operate in an *asynchronous* environment. This asynchrony has two facets: First, for the sake of convenience, we assume that the actual application of the transition function in each step $t \in \mathbb{Z}_{>0}$ of node $v \in V$ is instantaneous (namely, lasts zero time) and occurs at the end of the step;³ the length of step t of node v , denoted $L_{v,t}$, is defined as the time difference between the application of the transition function in step $t - 1$ and that of step t . It is assumed that $L_{v,t}$ is finite, but apart from that, we do not make any further assumptions on this length, that is, the step length $L_{v,t}$ is determined by the adversary independently of all other step lengths $L_{v',t'}$. In particular, we do not assume any synchronization between the steps of different nodes whatsoever.

Another facet of the asynchronous environment is that a message transmitted by node v in step t (if such a message is transmitted) is assumed to reach the port $\psi_u(v)$ of an adjacent node u after a finite time delay, denoted $D_{v,t,u}$. We assume that if v transmits message $\sigma_1 \in \Sigma$ in step t_1 and message $\sigma_2 \in \Sigma$ in step $t_2 > t_1$, then σ_1 reaches u before σ_2 does. Apart from this “FIFO” assumption, we do not make any other assumptions on the delays $D_{v,t,u}$. In particular, this means that under certain circumstances, the adversary may overwrite message σ_1 with message σ_2 in port $\psi_u(v)$ of u so that u will never “know” that message σ_1 was transmitted.⁴

Consequently, a *policy* of the adversary is captured by: (1) the length $L_{v,t}$ of step t of node v for every $v \in V$ and $t \in \mathbb{Z}_{>0}$; and (2) the delay $D_{v,t,u}$ of the delivery of the transmission of node v in step t to an adjacent node u for every $v \in V$, $t \in \mathbb{Z}_{>0}$, and $u \in N(v)$.⁵ Assuming that the adversary is oblivious to the random coin tosses of the nodes, an adversarial policy is depicted by infinite sequences of $L_{v,t}$ and $D_{v,t,u}$ parameters.

³ This assumption can be lifted at the cost of a more complicated definition of the adversarial policy described soon.

⁴ Often, much stronger assumptions are made in the literature. For example, a common assumption for asynchronous environments is that the port of node u corresponding to the adjacent node v is implemented by a buffer so that messages cannot be “lost”. We do not make any such assumption for our nFSM model.

⁵ We use the standard notation $N(v)$ for the *neighborhood* of node v in G , namely, the subset of nodes adjacent to v .

For further information on asynchronous environments, we point the reader to one of the standard textbooks [31, 28].

Correctness and Run-Time Measures. A protocol Π for problem P is said to be *correct* under the nFSM model if for every instance of P and for every adversarial policy, Π reaches an output configuration within finite time with probability 1, and for every output configuration reached by Π with positive probability, the output of the nodes is a valid solution to P . Given a correct protocol Π , the complexity measure that interests us in the current paper is the *run-time* of Π defined as follows.

Consider some instance \mathcal{I} of problem P . Given an adversarial policy \mathcal{A} and a sequence (actually an n -tuple of sequences) \mathcal{R} of random coin tosses that lead to an output configuration within finite time, the run-time $T_{\Pi}(\mathcal{I}, \mathcal{A}, \mathcal{R})$ of Π on \mathcal{I} with respect to \mathcal{A} and \mathcal{R} is defined as the (possibly fractional) number of *time units*⁶ that pass from the beginning of the execution until the first time the protocol reaches an output configuration, where a time unit is defined to be the maximum among all step length parameters $L_{v,t}$ and delivery delay parameters $D_{v,t,u}$ appearing in \mathcal{A} before the output configuration is reached. Let $T_{\Pi}(\mathcal{I}, \mathcal{A})$ denote the random variable that depicts the run-time of Π on \mathcal{I} with respect to \mathcal{A} . Following the standard procedure in this regard, we say that the run-time of a correct protocol Π for problem P is $f(n)$ if for every n -node instance \mathcal{I} of P and for every adversarial policy \mathcal{A} , it holds that $T_{\Pi}(\mathcal{I}, \mathcal{A})$ is at most $f(n)$ in expectation and with high probability. The protocol is said to be *efficient* if its run-time is polylogarithmic in the size of the network (cf. [26]).

3 Convenient Transformations

In this section, we show that the nFSM protocol designer may, in fact, assume a slightly more “user-friendly” environment than the one described in Section 2. This is based on the design of black-box compilers transforming a protocol that makes strong assumptions on the environment into one that does not make any such assumptions. Specifically, the assumptions that can be lifted that way are synchrony (Section 3.1), and multiple-letter queries (Section 3.2).

3.1 Implementing a Synchronizer

As described in Section 2, the nFSM model assumes an asynchronous environment. Nevertheless, it will be convenient to extend the nFSM model to *synchronous* environments. One natural such extension augments the model described in Section 2 with the following two *synchronization prop-*

⁶ Note that time units are defined solely for the purpose of the analysis. Under an asynchronous environment, the nodes have no notion of time and in particular, they cannot measure a single time unit.

erties for every two adjacent nodes $u, v \in V$ and for every $t \in \mathbb{Z}_{>0}$:

(S1) when node u is in step t , node v is in step $t - 1$, t , or $t + 1$; and

(S2) at the end of step $t + 1$ of u , port $\psi_u(v)$ stores the message transmitted by v in step t of v 's execution (or the last message transmitted by v prior to step t if v does not transmit any message in step t).

An environment in which properties (S1) and (S2) are guaranteed to hold is called a *locally synchronous* environment. Local-only communication can never achieve global synchrony, however, research in the message passing model has shown that local synchrony is often sufficient to provide efficient algorithms [4, 6, 5]. To distinguish a protocol assumed to operate in a locally synchronous environment from those making no such assumptions, we shall often refer to the execution steps of the former as *rounds* (cf. fully synchronized protocols). Our goal in this section is to establish the following theorem.

Theorem 3.1. *Every nFSM protocol $\Pi = \langle Q, Q_I, Q_O, \Sigma, \sigma_0, b, \lambda, \delta \rangle$ designed to operate in a locally synchronous environment can be simulated in an asynchronous environment by a protocol $\hat{\Pi}$ at the cost of a constant multiplicative run-time overhead.*

The procedure in charge of the simulation promised in Theorem 3.1 is referred to as a *synchronizer* [4]. The remainder of Section 3.1 is dedicated to the design (and analysis) of a synchronizer for the nFSM model.

Overview. Round $t \in \mathbb{Z}_{>0}$ of node $v \in V$ under Π is simulated by $O(1)$ contiguous steps under $\hat{\Pi}$; the collection of these steps is referred to as v 's *simulation phase* of round t . Protocol $\hat{\Pi}$ is designed so that v maintains the value of $t \bmod 3$, referred to as the *trit* (ternary digit) of round t , which is also encoded in the message transmitted by v at the end of round t .⁷ The main principle behind our synchronizer is that node v will not move to the simulation phase of round $t + 1$ while its ports still contain messages sent in a round whose trit is $t - 1 \bmod 3$.

Under Π , the decisions made by node v at round t should be based on the messages transmitted by all neighbors u of v at round $t - 1$. However, during v 's simulation phase of round t , port $\psi_v(u)$ may contain messages transmitted at round $t - 1$ or at round t under Π . The latter case is problematic since the message transmitted by u in the simulation phase of round $t - 1$ is overwritten by that transmitted in the simulation phase of round t . To avoid this obstacle, a message transmitted by node u under $\hat{\Pi}$ at the end of the simulation phase of round t also encodes the message that u transmitted under Π at round $t - 1$.

So, if v resides in a state whose query letter is $\sigma \in \Sigma$ in round t under Π , then under $\hat{\Pi}$, v should query for all $\hat{\Sigma}$ -letters encoding a transmission of σ at round $t - 1$. Since there are several such letters, a carefully designed feature should be used so that $\hat{\Pi}$ accounts for their combined number.

⁷ Note that maintaining the value of $t \bmod 2$ is insufficient for the sake of reaching synchronization.

Protocol $\widehat{\Pi}$. Let

$$\widehat{\Pi} = \langle \widehat{Q}, \widehat{Q}_I, \widehat{Q}_O, \widehat{\Sigma}, \widehat{\sigma}_0, b, \widehat{\lambda}, \widehat{\delta} \rangle .$$

Consider node $v \in V$ and round $t \in \mathbb{Z}_{>0}$. As the name implies, node v 's *simulation phase* of round t under $\widehat{\Pi}$, denoted $\phi_v(t)$, corresponds to round t of Π . Protocol $\widehat{\Pi}$ is designed so that at every step in $\phi_v(t)$ other than the last one, v does not transmit any message (indicated by transmitting ε), and at the last step of the simulation phase, v always transmits some message $\widehat{\sigma} \in \widehat{\Sigma}$, denoted $M_v(t)$.

The alphabet $\widehat{\Sigma}$ is defined to be

$$\Sigma' = (\Sigma \cup \{\varepsilon\}) \times (\Sigma \cup \{\varepsilon\}) \times \{0, 1, 2\} .$$

The semantics of the message $M_v(t) = (\sigma, \sigma', j)$ sent by node v at the last step of the simulation phase $\phi_v(t)$ is that: v transmits $\sigma \in \Sigma \cup \{\varepsilon\}$ at round $t - 1$ under Π ; v transmits $\sigma' \in \Sigma \cup \{\varepsilon\}$ at round t under Π ; and $j = t \bmod 3$. Following that logic, we set $\widehat{\sigma}_0 = (\varepsilon, \sigma_0, 0)$.

The state set \widehat{Q} of $\widehat{\Pi}$ is defined to be

$$\widehat{Q} = \left(\bigcup_{q \in Q} (P_q \cup S_q) \right) \times \{0, 1, 2\} ,$$

where $P_q \times \{j\}$ and $S_q \times \{j\}$, $q \in Q$, $j \in \{0, 1, 2\}$, are referred to as the *pausing* and *simulating* features, respectively, whose role will be clarified soon. Suppose that v resides in state $q \in Q$ in step t under Π and that $j = t \bmod 3$. Then, throughout $\phi_v(t)$, node v resides in some state in $(P_q \cup S_q) \times \{j\}$. In particular, in the first steps of the simulation phase, v resides in states of the pausing feature $P_q \times \{j\}$, and then at some stage it switches to the simulating feature $S_q \times \{j\}$ and remains in its states until the end of the simulation phase.

The Pausing Feature. For the simulation phase of round t , we denote the letters in $(\Sigma \cup \{\varepsilon\}) \times (\Sigma \cup \{\varepsilon\}) \times \{j - 2\}$ as *dirty* and the letters in $(\Sigma \cup \{\varepsilon\}) \times (\Sigma \cup \{\varepsilon\}) \times \{j - 1, j\}$ as *clean*.⁸ The purpose of the pausing feature $P_q \times \{j\}$ is to pause the execution of v until its ports do not contain any dirty letter. This is carried out by including in $P_q \times \{j\}$ a state $p_{\sigma, \sigma'}$ for every $\sigma, \sigma' \in \Sigma \cup \{\varepsilon\}$; the query letter of $p_{\sigma, \sigma'}$ is (the dirty letter) $\widehat{\lambda}(p_{\sigma, \sigma'}) = (\sigma, \sigma', j - 2)$ and the transition function $\widehat{\delta}$ is designed so that v moves to the next (according to some fixed order) state in the feature $P_q \times \{j\}$ if and only if there are no ports storing the query letter.

We argue that the pausing feature guarantees synchronization property (S1). For the sake of the analysis, it is convenient to assume the existence of a fully synchronous simulation phase of a virtual round 0; upon completion of this simulation phase (at the beginning of the execution), every node $v \in V$ transmits the message $M_v(0) = \widehat{\sigma}_0$. We are now ready to establish the following lemma.

⁸ Throughout this section, arithmetic involving the parameter j is done modulo 3.

Lemma 3.2. *For every $t \in \mathbb{Z}_{>0}$, $v \in V$, and $u \in N(v)$, when v completes the pausing feature of $\phi_v(t)$, port $\psi_v(u)$ stores either $M_u(t-1)$ or $M_u(t)$.*

Proof. By induction on t . The base case of round $t = 0$ holds by our assumption that $\phi_v(0)$ and $\phi_u(0)$ are fully synchronous. Assume by induction that the assertion holds for round $t-1$. Applying the inductive hypothesis to both u and v , we conclude that (1) when v completes the pausing feature of $\phi_v(t-1)$, port $\psi_v(u)$ stores either $M_u(t-2)$ or $M_u(t-1)$; and (2) when u completes the pausing feature of $\phi_u(t-1)$, port $\psi_u(v)$ stores either $M_v(t-2)$ or $M_v(t-1)$.

Let τ_u and τ_v denote the times at which u and v complete the pausing feature of $\phi_u(t)$ and $\phi_v(t)$, respectively. Since v cannot complete the pausing feature of $\phi_v(t)$ while $M_u(t-2)$ is still stored in $\psi_v(u)$, it follows that at time τ_v , port $\psi_v(u)$ stores the message $M_u(t')$ for some $t' \geq t-1$. Our goal in the remainder of this proof is to show that $t' \leq t$. If $\tau_v < \tau_u$, then t' must be exactly $t-1$, which concludes the inductive step for that case.

So, assume that $\tau_v > \tau_u$ and suppose by contradiction that $t' \geq t+1$. Using the same line of arguments as in the previous paragraph, we conclude that at time τ_u , port $\psi_u(v)$ stores the message $M_v(t-1)$. Node u cannot complete the pausing feature of $\phi_u(t+1)$ while $M_v(t-1)$ is still stored in $\psi_u(v)$, hence v must have transmitted $M_v(t)$ before u completed the pausing feature of $\phi_u(t+1)$. But this means that v completed the pausing feature of $\phi_v(t)$ before u could have transmitted $M_u(t+1)$, in contradiction to the assumption that $\psi_v(u)$ stores $M_u(t')$ for some $t' \geq t+1$ at time τ_v . The assertion follows. \square

Consider two adjacent nodes $u, v \in V$. If node u is at round $t-1$ when an adjacent node v is at round $t+1$, then v completed the pausing feature of $\phi_v(t)$ before u transmitted $M_u(t-1)$, in contradiction to Lemma 3.2. Therefore, our synchronizer satisfies synchronization property (S1). Furthermore, a similar argument shows that between the time v completed the pausing feature of $\phi_v(t)$ and the time v completed the simulation phase $\phi_v(t)$ itself, the content of $\psi_v(u)$ may change from $M_u(t-1)$ to $M_u(t)$ (if it was not already $M_u(t)$), but it will not store $M_u(t')$ for any $t' > t$. This fact is crucial for the implementation of the simulation feature.

The Simulation Feature. Upon completion of the pausing feature $P_q \times \{j\}$, v moves on to the simulation feature $S_q \times \{j\}$. The purpose of this feature is to perform the actual simulation of round t in v , namely, to determine the state (of Q) dominating the simulation phase of the next round and the message transmitted when moving from the simulation phase of the current round to that of the next round.

To see how this works out, suppose that $\lambda(q) = \sigma \in \Sigma$. We would have wanted node v to count (up to the bounding parameter b) the number of occurrences of $\hat{\Sigma}$ -letters in its ports that correspond to the transmission of σ at round $t-1$ under Π , that is, the number of occurrences of

letters in $\Gamma_{t-1} \cup \Gamma_t$, where

$$\Gamma_{t-1} = \{(\sigma', \sigma, j-1) \mid \sigma' \in \Sigma \cup \{\varepsilon\}\} \quad \text{and} \quad \Gamma_t = \{(\sigma, \sigma', j) \mid \sigma' \in \Sigma \cup \{\varepsilon\}\}.$$

More formally, the application of the transition function $\widehat{\delta}$ at the end of the simulation phase $\phi_v(t)$ should be based on $f_b(\sum_{\gamma \in \Gamma_{t-1} \cup \Gamma_t} \#(\gamma))$, where $\#(\gamma)$ stands for the number of occurrences of the letter γ in the ports of v at the end of $\phi_v(t)$.

Identifying the integer b with the symbol $\geq b$, we observe that the function $f_b : \mathbb{Z}_{\geq 0} \rightarrow B$ satisfies

$$f_b(x + y) = \min \{f_b(x) + f_b(y), b\}$$

for every $x, y \in \mathbb{Z}_{\geq 0}$. A natural attempt to compute $f_b(\sum_{\gamma \in \Gamma_{t-1} \cup \Gamma_t} \#(\gamma))$ would include in the feature $S_q \times \{j\}$ a state $s_{\gamma,i}$ for every letter $\gamma \in \Gamma_{t-1} \cup \Gamma_t$ and integer $i \in \{0, \dots, b\}$; the query letter of $s_{\gamma,i}$ would be $\widehat{\lambda}(s_{\gamma,i}) = \gamma$ and the transition function $\widehat{\delta}$ would be designed so that v moves from $s_{\gamma,i}$ to $s_{\gamma',i'}$, where γ' follows γ in some fixed order of the letters in $\Gamma_{t-1} \cup \Gamma_t$ and $i' = \min\{i + f_b(\#(\gamma)), b\}$.

However, care must be taken with this approach since $\#(\gamma)$ may decrease (respectively, increase) during $\phi_v(t)$ for $\gamma \in \Gamma_{t-1}$ (resp., for $\gamma \in \Gamma_t$) due to new incoming messages. To avoid this obstacle, we design the feature $S_q \times \{j\}$ so that first, it computes $\varphi_1 \leftarrow f_b(\sum_{\gamma \in \Gamma_{t-1}} \#(\gamma))$; next, it computes $\varphi_2 \leftarrow f_b(\sum_{\gamma \in \Gamma_t} \#(\gamma))$; and finally, it computes “again” $\varphi_3 \leftarrow f_b(\sum_{\gamma \in \Gamma_{t-1}} \#(\gamma))$. If $\varphi_1 = \varphi_3$, then the current simulation phase is over and $\widehat{\delta}$ is applied, simulating $\delta(q, f_b(\varphi_1 + \varphi_2))$; otherwise, the feature $S_q \times \{j\}$ is invoked from scratch. Since the value of $f_b(\sum_{\gamma \in \Gamma_{t-1}} \#(\gamma))$ cannot increase during the simulation phase, and since $\varphi_1 \leq b$, the feature $S_q \times \{j\}$ is invoked at most b times throughout the execution of the simulation phase. By induction on t , we conclude that our synchronizer satisfies synchronization property (S2), which concludes the correctness proof of the simulation.

Accounting. It remains to show that all ingredients of protocol $\widehat{\Pi}$ are of constant size and that the run-time of protocol $\widehat{\Pi}$ incurs at most a constant multiplicative overhead on top of that of protocol Π . The former claim is established by following our synchronizer construction, observing that $|\widehat{\Sigma}| = O(|\Sigma|^2)$ and $|\widehat{Q}| = O(|Q| \cdot (|\Sigma|^2 + |\Sigma| \cdot b))$ (recall that the bounding parameter b remains unchanged). For the latter claim, we need the following definition: given some node subset $U \in V$ and round $t \in \mathbb{Z}_{>0}$, let $\tau(U, t)$ denote the first time at which u completed simulation phase $\phi_u(t)$ for all nodes $u \in U$. The following proposition can now be established.

Proposition 3.3. *For every node $v \in V$ and round $t \in \mathbb{Z}_{>0}$, the time difference $\tau(\{v\}, t+1) - \tau(N(v) \cup \{v\}, t)$ is (up)bounded by a constant.*

Proof. Since each transmitted message has a delay of at most 1 unit of time, it follows that by time $\tau(N(v) \cup \{v\}, t) + 1$, message $M_u(t)$ must reach $\psi_v(u)$ for all $u \in N(v)$. The pausing and simulation features of $\phi_v(t+1)$ are then completed within $O(|\Sigma|^2)$ and $O(|\Sigma| \cdot b)$ steps, respectively. The assertion follows as each step lasts for at most 1 unit of time. \square

Employing Proposition 3.3, we conclude by induction on t that $\tau(V, t) = O(t)$ for every $t \in \mathbb{Z}_{>0}$, hence if the execution of protocol Π requires T rounds, then the execution of protocol $\hat{\Pi}$ is completed within $O(T)$ time units. Theorem 3.1 follows.

3.2 Multiple-Letter Queries

Recall that according to the model presented in Section 2, each state $q \in Q$ is associated with a query letter $\lambda(q)$ and the application of the transition function when node v resides in state q is determined by $f_b(\sharp(\sigma))$, where $\sharp(\sigma)$ is the number of occurrences of the letter σ in the ports of v . From the perspective of the protocol designer, it is often more convenient to assume that the node queries on all letters simultaneously, namely, that the application of the transition function is determined by the vector $\langle f_b(\sharp(\sigma)) \rangle_{\sigma \in \Sigma}$.

Now that we may assume a synchronous environment, this stronger multiple-letter queries assumption can easily be supported. Indeed, at the cost of increasing the number of states and the run-time by constant factors, one can subdivide each round into $|\Sigma|$ subrounds, dedicating each subround to a different letter in Σ , so that at the end of the round, the state of v reflects $f_b(\sharp(\sigma))$ for every $\sigma \in \Sigma$.

Theorem 3.4. *Every nFSM protocol with multiple-letter queries can be simulated by an nFSM protocol with single-letter queries at the cost of a constant multiplicative run-time overhead.*

4 Maximal Independent Set

Given a graph $G = (V, E)$, the *maximal independent set (MIS)* problem asks for a node subset $U \subseteq V$ which is independent in the sense that $(U \times U) \cap E = \emptyset$, and maximal in the sense that $U' \subseteq V$ is not independent for every $U' \supset U$. Distributed MIS algorithms with logarithmic run-time operating in the message passing model were presented by Luby [27] and independently, by Alon et al. [3];⁹ Luby's algorithm has since become a specimen of distributed algorithms; in the last 25 years, researchers have tried to improve it, if only e.g., with an improved bit complexity [29], on special graph classes [34, 25], or in a weaker communication model [1]. An $\Omega(\sqrt{\log n})$ -lower bound on the run-time of any distributed MIS algorithm operating in the message passing model was established by Kuhn et al. [23]. Our goal in this section is to design an nFSM protocol for the MIS problem with run-time $O(\log^2 n)$.

Outline of the Key Technical Ideas. Our protocol is inspired by the existing message passing MIS algorithms. Common to all these algorithms is that they are based on the concept of grouping

⁹ The focus of [27] and [3] was actually on the PRAM model, but their algorithms can be adapted to the message passing model.

consecutive rounds into *phases*, where in each phase, nodes compete against their neighbors over the right to join the MIS. Existing implementations of such competitions require at least one of the following three capabilities: (1) performing calculations that involve super-constant numbers; (2) communicating with each neighbor independently; or (3) sending messages of super-constant size, specifically, of size $c \log n$ for some constant $c > 0$. The first two capabilities are clearly out of the question for an nFSM protocol. The third one is also not supported by the nFSM model, but perhaps one can divide a message with a logarithmic number of bits over logarithmic many rounds, sending 1 (or $O(1)$) bits per round (cf. Algorithm B in [29])?

This naive attempt results in phases of length $c \log n$. However, no FSM can count the rounds in a $c \log n$ long phase — a task essential for deciding if the current phase is over and the next one should begin. Furthermore, to guarantee fair competition, the phases must be aligned across the network, thus ruling out the possibility to start node v 's phase i before phase $i - 1$ of some node $u \in N(v)$ is finished. In fact, an efficient algorithm that requires $\omega(1)$ long aligned phases cannot be implemented under the nFSM model. So, how can we decide if node v joins the MIS using constant size messages without the ability to maintain long aligned phases?

This issue is resolved by relaxing the requirements that the phases are aligned and of a pre-determined length, introducing a feature referred to as a *tournament*. Our tournaments are only “softly” aligned and their lengths are determined probabilistically, in a manner that can be maintained under the nFSM model. Nevertheless, they enable a fair competition between neighboring nodes, as desired.

The Protocol. Employing Theorems 3.1 and 3.4, we assume a locally synchronous environment and use multiple-letter queries. The state set of the protocol is $Q = \{\text{WIN}, \text{LOSE}, \text{DOWN}_1, \text{DOWN}_2, \text{UP}_0, \text{UP}_1, \text{UP}_2\}$, with $Q_I = \{\text{DOWN}_1\}$ (the initial state of all nodes) and $Q_O = \{\text{WIN}, \text{LOSE}\}$, where WIN (respectively, LOSE) indicates membership (resp., non-membership) in the MIS output by the protocol. The states in $Q_A = Q - Q_O$ are called the *active* states and a node in an active state is referred to as an *active* node. We take the communication alphabet Σ to be identical to the state set Q , where the letter transmissions are designed so that node v transmits letter q whenever it moves to state q from some state $q' \neq q$; no letter is transmitted in a round at which v remains in the same state. Letter DOWN_1 is the initial letter stored in all ports at the beginning of the execution. The bounding parameter is set to $b = 1$.

A schematic description of the transition function is provided in Figure 1; its logic is as follows. Each state $q \in Q_A$ has a subset $D(q) \subseteq Q_A$ of *delaying states*: node v remains in the current state q as long as (at least) one of its neighbors is in some state in $D(q)$. This is implemented by querying on the letters (corresponding to the states) in $D(q)$, staying in state q as long as at least one of these letters is found in the ports. Specifically, state DOWN_1 is delayed by state DOWN_2 , which is delayed by all three UP states. State UP_j , $j = 0, 1, 2$, is delayed by state $\text{UP}_{j-1 \bmod 3}$, where state

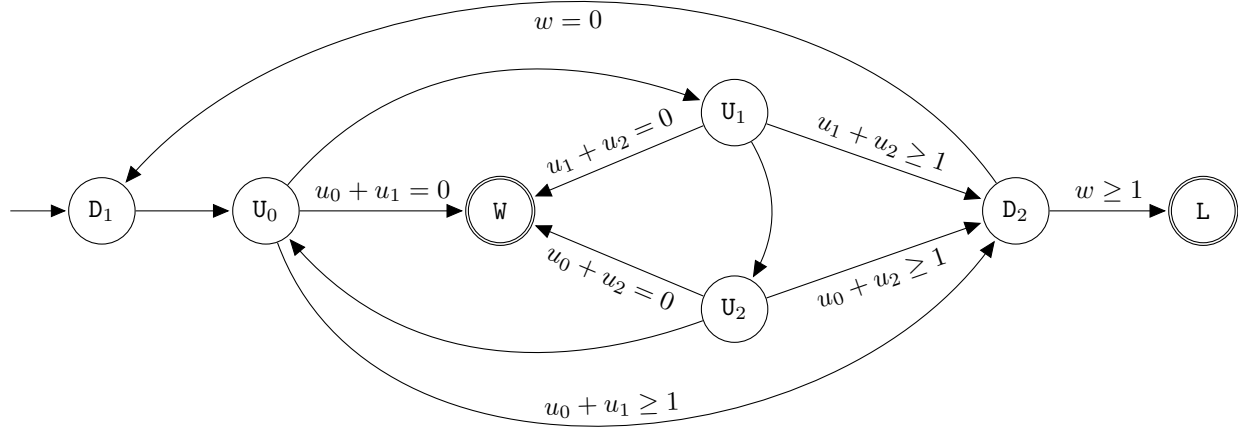


Figure 1: The transition function of the MIS protocol with state names abbreviated by their first (capital) letters. The node stays in state q (a.k.a. *delayed*) as long as $\sharp(q') > 0$ for any state q' such that a $q' \rightarrow q$ transition is defined (for clarity, this is omitted from the figure). Assuming that the node is not delayed, each transition specified in the figure is associated with a condition on the number of appearances of the query letters in the ports (depicted by the corresponding lower-case letter) so that the transition is followed only if the condition is satisfied (an empty condition is satisfied by all port configurations); if some port configuration satisfies several transition conditions, then one of them is chosen uniformly at random.

UP_0 is also delayed by state $DOWN_1$.

States WIN and LOSE are sinks in the sense that a node that moves to one of these states will stay there indefinitely. Assuming that node v does not find any delaying letter in its ports, the logic of the UP and DOWN states is as follows. From state $DOWN_1$, v moves to state UP_0 . From state $DOWN_2$, v moves to state $DOWN_1$ if $\sharp(WIN) = 0$, that is, if it does not find any WIN letter in its ports; otherwise, it moves to state LOSE. When in state UP_j , v tosses a fair coin and proceeds as follows: if the coin turns head, then v moves to state $UP_{j+1 \bmod 3}$; if the coin turns tail, then v moves to state WIN if $\sharp(UP_j) = \sharp(UP_{j+1 \bmod 3}) = 0$; and to state $DOWN_2$ otherwise. This completes the description of our nFSM protocol for the MIS problem.

Turns and Tournaments. Our protocol is designed so that an active node v traverses the DOWN and UP states in a (double-)circular fashion: an inner loop of the UP states (moving from state UP_j to state $UP_{j+1 \bmod 3}$) nested within an outer loop consisting of the DOWN states and the inner loop. Of course, v may spend more than one round at each state $q \in Q_A$ (delayed by adjacent nodes in states $D(q)$); we refer to a maximal contiguous sequence of rounds that v spends in the same state

$q \in Q_A$ as a q -turn, or simply as a *turn* if the actual state q is irrelevant. A maximal contiguous sequence of turns that starts at a DOWN_1 -turn and does not include any other DOWN_1 -turn (i.e., a single iteration of the outer loop) is referred to as a *tournament*. We index the tournaments and the turns within a tournament by the positive integers. Note that by definition, every tournament i of v starts with a DOWN_1 -turn, followed by a non-empty sequence of UP -turns. If tournament $i + 1$ of v exists, then tournament i ends with a DOWN_2 -turn; otherwise, it ends with an UP -turn. The following observation is established by induction on the rounds.

Observation 4.1. *Consider some node $v \in V$ in turn $j \in \mathbb{Z}_{>0}$ of tournament $i \in \mathbb{Z}_{>0}$ and some active node $u \in N(v)$.*

- *If this is a DOWN_1 -turn of v ($j = 1$), then u is in either (A) the last (DOWN_2 -)turn of tournament $i - 1$; (B) turn 1 of tournament i ; or (C) turn 2 of tournament i .*
- *If this is an UP -turn of v ($j \geq 2$), then u is in either (A) turn $j - 1$ of tournament i ; (B) turn j of tournament i ; (C) turn $j + 1$ of tournament i ; or (D) the last (DOWN_2 -)turn $j' \leq j + 1$ of tournament i .*
- *If this is a DOWN_2 -turn of v (the last turn of this tournament), then u is in either (A) an UP -turn $j' \geq j - 1$ of tournament i ; (B) the last (DOWN_2 -)turn of tournament i ; or (C) turn 1 of tournament $i + 1$.*

Given some $U \subseteq V$ and $i, j \in \mathbb{Z}_{>0}$, let $T_U(i, j)$ denote the first time at which every node $v \in U$ satisfies either

- (1) v is inactive;
- (2) v is in tournament $i' > i$;
- (3) v is in the last (DOWN_2 -)turn of tournament i ; or
- (4) v is in turn $j' \geq j$ of tournament i .

Employing Observation 4.1, the delaying states feature guarantees that

$$T_v(i, j + 1) \leq T_{N(v) \cup \{v\}}(i, j) + 1 \quad (1)$$

for every $v \in V$ and $i, j \in \mathbb{Z}_{>0}$. Since $T_U(i, j) \leq T_V(i, j)$ for every $U \subseteq V$, we can apply inequality (1) to each node $v \in V$, concluding that

$$T_V(i, j + 1) \leq T_V(i, j) + 1,$$

which immediately implies that

$$T_V(i, k + 1) \leq T_V(i, 1) + k. \quad (2)$$

Geometric Random Variables. Consider some $v \in V$ and $i \in \mathbb{Z}_{>0}$. Assuming that tournament i of v exists, let $X_v(i)$ denote its *length* in terms of number of turns. For the sake of simplifying the analysis, if tournament i is the last tournament of v , then we actually take $X_v(i)$ to be its length plus

1 (this is done in order to compensate for the missing DOWN_2 -turn in the end of the tournament.) The logic of the UP states implies that $X_v(i)$ is a random variable that obeys distribution $\text{Geom}(1/2)+2$, namely, a fixed term of 2 plus the geometric distribution with parameter $1/2$, independently of $X_{v'}(i')$ for any $v' \neq v$ and/or $i' \neq i$. Since the maximum of n independent $\text{Geom}(1/2)$ -random variables is $O(\log n)$ with high probability, inequality 2 yields the following observation.

Observation 4.2. *For every $i \in \mathbb{Z}_{>0}$, $T_V(i, 1)$ is finite with probability 1 and*

$$T_V(i+1, 1) \leq T_V(i, 1) + O(\log n)$$

with high probability.

Our protocol is designed so that node v moves to an output state (WIN or LOSE) in the end of each tournament with positive probability. Moreover, the logic of state DOWN_2 guarantees that if node v moves to state WIN in the end of tournament i , then all its active neighbors move to state LOSE in the end of their respective tournaments i . By Observation 4.2, we conclude that our protocol reaches an output configuration with probability 1 and that every output configuration reflects an MIS. It remains to bound the run-time of our protocol.

The Virtual Graph G^i . Let V^i be the set of nodes for which tournament i exists and let $G^i = (V^i, E^i)$ be the subgraph induced on G by V^i , where $E^i = E \cap (V^i \times V^i)$.¹⁰ Given some node $v \in V^i$, let $N^i(v) = \{u \in V^i \mid (u, v) \in E\}$ be the neighborhood of node v in G^i and let $d^i(v) = |N^i(v)|$ be its degree. Note that the graph G^i is virtual and defined solely for the sake of the analysis; in particular, we do not assume that there exists some time at which the graph induced by any meaningful subset of the nodes (say, the nodes in tournament i) agrees with G^i . The key observation in this context is that conditioned on G^i , the random variables $X_v(i)$, $v \in V^i$, are (still) independent and obey distribution $\text{Geom}(1/2) + 2$. Moreover, the graph G^{i+1} is fully determined by the random variables $X_v(i)$, $v \in V^i$. Our analysis relies on the following lemma.

Lemma 4.3. *There exist two constants $0 < p, c < 1$ such that $|E^{i+1}| \leq c|E^i|$ with probability at least p .*

We will soon turn to proving Lemma 4.3, but first, let us explain why it suffices for the completion of our analysis. Define the random variable $Y = \min\{i \in \mathbb{Z}_{>0} : |E^i| = 0\}$. Lemma 4.3 implies that Y is stochastically dominated by a random variable that obeys distribution $\text{NB}(O(\log n), 1 - p) + O(\log n)$, namely, a fixed term of $O(\log n)$ plus the negative binomial distribution with parameters $O(\log n)$ and $1 - p$, hence $Y = O(\log n)$ in expectation and with high probability. Since the nodes in $V - V^i$ are all in an output state (and will remain in that state), and since the logic of the UP states implies that a degree-0 node in G^i will move to state WIN in the end of tournament i (with probability 1) and thus, will not be included in V^{i+1} , we can employ Observation 4.2 to conclude that the run-time of our protocol is $O(\log^2 n)$.

¹⁰ The notation G^i used in this section should not be confused with the i^{th} power of G .

The remainder of this section is dedicated to establishing Lemma 4.3. The proof technique we use for that purpose resembles (a hybrid of) the techniques used in [3] and [29] for the analysis of their MIS algorithms. We say that node $v \in V^i$ is *good* in G^i if

$$|\{u \in N^i(v) \mid d^i(u) \leq d^i(v)\}| \geq d^i(v)/3,$$

i.e., if at least third of v 's neighbors in G^i have degrees smaller or equal to that of v . The following lemma is established in [3].

Lemma 4.4 ([3]). *More than half of the edges in E^i are incident on good nodes in G^i .*

Disjoint Winning Events. Consider some good node v in G^i with $d = d^i(v) > 0$ and let $\hat{N}^i(v) = \{u \in N^i(v) \mid d^i(u) \leq d\}$. Recall that the definition of a good node implies that $|\hat{N}^i(v)| \geq d/3$. We say that node $u \in \hat{N}^i(v)$ *wins* v in tournament i if

$$X_u(i) > \max \left\{ X_w(i) \mid w \in N^i(u) \cup \hat{N}^i(v) - \{u\} \right\}$$

and denote this event by $A^i(u, v)$. The main observation now is that if u wins v in tournament i , then in the end of their respective tournaments i , u moves to state WIN and v moves to state LOSE. Moreover, the events $A^i(u, v)$ and $A^i(w, v)$ are disjoint for every $u, w \in \hat{N}^i(v)$, $u \neq w$.

Let u_1, \dots, u_k be the nodes in $N^i(u) \cup \hat{N}^i(v)$, where $0 < k \leq 2d$ by the definition of a good node. Let $B^i(u, v)$ denote the event that the maximum of $\{X_{u_\ell}(i) \mid 1 \leq \ell \leq k\}$ is attained at a single $1 \leq \ell \leq k$. Since $X_{u_1}(i), \dots, X_{u_k}(i)$ are independent random variables that obey distribution $\text{Geom}(1/2) + 2$, it follows that $\mathbb{P}(B^i(u, v)) \geq 2/3$. Therefore,

$$\mathbb{P}(A^i(u, v)) = \mathbb{P}(A^i(u, v) \mid B^i(u, v)) \cdot \mathbb{P}(B^i(u, v)) \geq \frac{1}{k} \cdot \frac{2}{3},$$

which implies that

$$\begin{aligned} \mathbb{P}(v \notin V^{i+1} \mid v \text{ is good in } G^i) &\geq \mathbb{P}\left(\bigvee_{u \in \hat{N}^i(v)} A^i(u, v)\right) \\ &= \sum_{u \in \hat{N}^i(v)} \mathbb{P}(A^i(u, v)) \geq \frac{d}{3} \cdot \frac{1}{2d} \cdot \frac{2}{3} = \frac{1}{9}. \end{aligned}$$

Combined with Lemma 4.4, we conclude that $\mathbb{E}[|E^{i+1}|] < \frac{35}{36} |E^i|$. Lemma 4.3 follows by Markov's bound.

Theorem 4.5. *There exists an nFSM protocol that computes an MIS in any n -node graph with run-time $O(\log^2 n)$.*

5 Coloring a Tree with 3 Colors

Given a graph $G = (V, E)$, the *coloring* problem asks for an assignment of colors to the nodes such that no two neighboring nodes have the same color. A coloring using at most k colors is called a

k-coloring. The smallest number of colors needed to color graph G is called its *chromatic number*, denoted by $\chi(G)$. In general, $\chi(G)$ is difficult to compute even in a centralized model [10]. As such, the distributed computing community is generally satisfied already with a $(\Delta + 1)$ -, $O(\Delta)$ -, or even $\Delta^{O(1)}$ -coloring, where $\Delta = \Delta(G)$ is the largest degree in the graph G , with possibly $\Delta(G) \gg \chi(G)$ [15, 32, 19, 26, 37, 7, 22, 9, 8, 35]. However, even for relatively simple graph classes, Δ may grow with n . As the output of each node under the nFSM model is taken from a constant size set, we must and will tackle a graph class that features a small chromatic number: trees.

Any tree T has a chromatic number $\chi(T) = 2$. Unfortunately, it is easy to show that in general, the task of 2-coloring trees requires run-time proportional to the diameter of the tree even under the message passing model, and hence cannot be achieved by an efficient distributed algorithm. The situation improves dramatically once 3 colors are allowed; indeed, Cole and Vishkin [15] presented a distributed algorithm that 3-colors directed paths, and in fact, any directed tree (directed in the sense that each node knows the port leading to its unique parent), in time $O(\log^* n)$. Linial [26] showed that this is asymptotically optimal.

Since it is not clear how to represent directed trees in the nFSM model, we focus on undirected trees, designing an nFSM protocol that 3-colors any n -node (undirected) tree in run-time $O(\log n)$. A lower bound result of Kothapalli et al. [21] shows that this cannot be improved (asymptotically) even by a message passing algorithm as long as the size of each message is $O(1)$.

Employing Theorems 3.1 and 3.4, we assume a locally synchronous environment and use multiple-letter queries. The description of the protocol will not dwell into the level of defining the states and transition function (as we did in Section 4 for the MIS protocol), but the reader will be easily convinced that this protocol can indeed be implemented under the nFSM model.

The Modes. At all times, each node $v \in V$ is in one of the following three *modes*.

- (1) Mode **COLORED**: the color of v is determined (v is in an output state) and it no longer takes an active part in the protocol.
- (2) Mode **ACTIVE**: the color of v has not been determined yet and v takes an active part in the protocol.
- (3) Mode **WAITING**: the color of v has not been determined yet and v is waiting for one of its neighbors to be colored before it resumes taking an active part in the protocol (going back to mode **ACTIVE**).

Initially, all nodes are in mode **ACTIVE**. When an **ACTIVE** node moves to mode **COLORED**, assigned with color $c \in \{1, 2, 3\}$, it transmits a ‘my color is c ’ message and it does not transmit any more messages; when an **ACTIVE** node moves to mode **WAITING**, it transmits an ‘I am **WAITING**’ message and it does not transmit any more messages until it returns to mode **ACTIVE**, in which case it transmits an ‘I am **ACTIVE**’ message. Therefore, the message stored in the port of node v

corresponding to neighbor u of v always indicates (perhaps among other things) the current mode of u .

The Phases. The execution of the protocol is divided into *phases* indexed by the positive integers, where each phase consists of 4 rounds. Consider some phase $i \in \mathbb{Z}_{>0}$. Let V^i be the set of **ACTIVE** nodes at the beginning of phase i and let F^i be the forest induced on T by V^i (F^i may contain one or more trees), referred to as the **ACTIVE forest**. Given some node $v \in V^i$, let $N^i(v) = \{u \in V^i \mid (u, v) \in E\}$ be the neighborhood of v in F^i and let $d^i(v) = |N^i(v)|$ be its degree.

The structure of the phases is as follows. Consider some node $v \in V^i$. In round 1 of the phase, v transmits an ‘I am **ACTIVE**’ message. Setting the bounding parameter of the protocol to $b = 3$, we conclude that in round 2, v can distinguish between the cases $d^i(v) = 0$, $d^i(v) = 1$, $d^i(v) = 2$, and $d^i(v) \geq 3$ simply by querying its ports for ‘I am **ACTIVE**’ messages; in other words, v “knows” $f_3(d^i(v))$, i.e., its degree calculated with respect to the one-two-many principle with bounding parameter $b = 3$. Employing this “knowledge”, v transmits $f_3(d^i(v))$ in round 2 of phase i , so in round 3, the port of v corresponding to u stores a message indicating $f_3(d^i(u))$ for every node $u \in N^i(v)$.

Rounds 3 and 4 of phase i are dedicated to Procedure **RandColor** that we will describe soon. Whether or not v runs Procedure **RandColor** depends on the degree of v and on the degrees of its **ACTIVE** neighbors. Specifically, v runs Procedure **RandColor** if: (1) $d^i(v) = 0$; (2) $d^i(v) = 1$ with $N^i(v) = \{u\}$ and $d^i(u) = 1$; or (3) $d^i(v) = 2$ with $N^i(v) = \{u_1, u_2\}$ and $d^i(u_1), d^i(u_2) \leq 2$. In contrast, if $d^i(v) = 1$ with $N^i(v) = \{u\}$ and $d^i(u) \geq 2$, then v moves to mode **WAITING** without running Procedure **RandColor**, in which case we say (just for the sake of the analysis) that v *waits on* u . Otherwise ($d^i(v) \geq 3$ or $d^i(v) = 2$ with some neighbor $u \in N^i(v)$ such that $d^i(u) \geq 3$), v remains in mode **ACTIVE** without running Procedure **RandColor**.

As stated beforehand, the **COLORED** nodes do not take an active part in the protocol. A **WAITING** node v moves to mode **ACTIVE** in the end of phase i if some neighbor u of v , $u \in V^i$, moves to mode **COLORED** during phase i (v spots this event by querying on ‘my color is c ’ messages).

Procedure RandColor. Responsible for the actual color assignments, Procedure **RandColor** takes 2 rounds (rounds 3 and 4 of some phase). Only an **ACTIVE** node may run the procedure, and when the procedure is over, the node either stays in mode **ACTIVE** or moves to mode **COLORED**. Consider some node v running the procedure and let $C(v) \subseteq \{1, 2, 3\}$ be the subset of colors which are not yet assigned to the neighbors of v in T . (Our analysis shows that if v is **ACTIVE**, then $C(v) \neq \emptyset$.) As every **COLORED** node transmits a message indicating its color, v can determine $C(v)$ by querying its ports.

In the first round of Procedure **RandColor**, v picks some color $c \in C(v)$ uniformly at random and transmits a ‘proposing color c ’ message. In the second round of the procedure, if v finds a

‘proposing color c ’ (with the same c) in its ports, then it remains in mode **ACTIVE**. Otherwise (no neighbor of v competes with v over color c), it moves to mode **COLORED** and transmits a ‘my color is c ’ message. This completes the description of our protocol.

The Waiting Hierarchy. The ‘waits on’ relation induces a hierarchy referred to as the *waiting hierarchy* which is represented by a (collection of) directed tree(s) defined over a subset of the edges of the tree T . Our protocol is designed so that if v waits on u , moving to mode **WAITING** in phase i , then in phases $1, \dots, i$, u was **ACTIVE**, and in phase $i + 1$, u is either **ACTIVE** or **COLORED**. Moreover, if u is **ACTIVE** and $v \in N(u)$ is **WAITING**, then v must be waiting on u . Note also that if v waits on u and u moves to mode **COLORED** in phase j , then v moves back to mode **ACTIVE** in (the beginning of) phase $j + 1$ and $d^{j+1}(v) = 0$.

Observation. *In the beginning of phase i , $|C(v)| \geq \min\{d^i(v) + 1, 3\}$ for every $i \in \mathbb{Z}_{>0}$ and node $v \in V^i$.*

Proof. As long as $d^i(v) \geq 3$, no neighbor of v can run Procedure **RandColor**, and hence no neighbor of v can move to mode **COLORED**. Therefore, $C(v) = \{1, 2, 3\}$ in the beginning of the first phase $i \in \mathbb{Z}_{>0}$ such that $d^i(v) \leq 2$. From that moment on, every **ACTIVE** neighbor of v that moves to mode **COLORED** decreases both $|C(v)|$ and $d^i(v)$ by 1. The assertion is completed by recalling that non-**ACTIVE** neighbors of v must be waiting on v and hence, cannot move to mode **COLORED** before v does. \square

Corollary 5.1. *Consider some node $v \in V^i$ that runs Procedure **RandColor**. If $d^i(v) = 0$, then v moves to mode **COLORED** with probability 1. Otherwise ($d^i(v)$ is either 1 or 2), v moves to mode **COLORED** with a positive constant probability.*

Let \tilde{V}^i be the restriction of V^i to nodes v that were **ACTIVE** in all phases $1, \dots, i$; this is, \tilde{V}^i does not include **WAITING** nodes that became **ACTIVE** again (recall that these will move to mode **COLORED** in the next phase with probability 1). Let \tilde{F}^i be the forest induced on T by \tilde{V}^i . Given some node $v \in \tilde{V}^i$, let $\tilde{N}^i(v) = \{u \in \tilde{V}^i \mid (u, v) \in E\}$ be the neighborhood of v in \tilde{F}^i and let $\tilde{d}^i(v) = |\tilde{N}^i(v)|$ be its degree. Observe that if $v \in \tilde{V}^i$, then $v \in V^i$ and $\tilde{d}^i(v) = d^i(v)$. Therefore, if $v \in V^i - \tilde{V}^i$, then $d^i(v) = 0$, in which case v runs Procedure **RandColor** in phase i and Corollary 5.1 guarantees that $v \notin V^{i+1}$.

The correctness of the protocol can now be established: The logic of Procedure **RandColor** implies that every output configuration is a legal coloring. Since **ACTIVE** leaves are removed from \tilde{F}^i with probability 1 and since every tree has at least two leaves, it follows that $\hat{V}^{1+k} = \emptyset$ for $k = \lceil n/2 \rceil$. Combining the properties of the waiting hierarchy with Corollary 5.1, we conclude that the execution reaches an output configuration within at most k additional phases. It remains to analyze the run-time of our protocol.

Good nodes. Consider some tree T' . We say that node v of T' is *good* if v is a leaf or if the degree of v is 2 and both neighbors of v are of degree at most 2.

Observation 5.2. *In every tree, at least a $(1/5)$ -fraction of the nodes are good.*

Consider some $i \in \mathbb{Z}_{>0}$ and some node $v \in \tilde{V}^i$. Let T' be the tree to which v belongs in \tilde{F}^i . We argue that if v is good in T' , then $v \notin \tilde{V}^i$ with a positive constant probability. Indeed, if v is a leaf in T' , which means that $\tilde{d}^i(v) = d^i(v) = 1$, then it either moves to mode **WAITING** with probability 1 (if the neighbor of v has a higher degree) or it runs Procedure **RandColor**, in which case Corollary 5.1 guarantees that v moves to mode **COLORED** with a positive constant probability; if $\tilde{d}^i(v) = d^i(v) = 2$ and both neighbors of v in F^i (and in T') are of degree at most 2, then v runs Procedure **RandColor**, in which case Corollary 5.1 again guarantees that v moves to mode **COLORED** with a positive constant probability. Since Corollary 5.1 also guarantees that nodes of degree 0 in \tilde{F}^i move to mode **COLORED** with probability 1, we can employ Observation 5.2 and Markov's bound to establish the following observation.

Observation 5.3. *There exists two constants $0 < p, c < 1$ such that $|\tilde{V}^{i+1}| \leq c|\tilde{V}^i|$ with probability at least p .*

Similarly to the analysis in Section 4, define the random variable $Y = \min\{i \in \mathbb{Z}_{>0} : |\tilde{V}^i| = 0\}$. Observation 5.3 implies that Y is stochastically dominated by a random variable that obeys distribution $\text{NB}(O(\log n), 1 - p) + O(\log n)$, namely, a fixed term of $O(\log n)$ plus the negative binomial distribution with parameters $O(\log n)$ and $1 - p$, hence $Y = O(\log n)$ in expectation and with high probability. Since Y bounds from above the depth of the waiting hierarchy, it follows that the execution reaches an output configuration within $2Y$ phases, which completes the analysis.

Theorem 5.4. *There exists an $n\text{FSM}$ protocol that 3-colors any n -node (undirected) tree with run-time $O(\log n)$.*

6 Computational Power

A *deterministic linear bounded automaton* (dLBA) is a (deterministic) Turing machine whose working tape is restricted to the cells specifying the input (this is equivalent to a $\text{DSPACE}(O(n))$ Turing machine). A *non-deterministic linear bounded automaton*, a.k.a., *linear bounded automaton* (LBA), is the non-deterministic version of a dLBA, and a *randomized linear bounded automaton* (rLBA) is the randomized version. Kuroda [24] proved that the class of languages that can be decided by an LBA is exactly the context-sensitive languages, corresponding to the Type-1 grammars in Chomsky's hierarchy of formal languages [14]. Whether LBAs are equivalent to dLBAs and where exactly do rLBAs lie between the two are major open questions in computational complexity (cf. the first LBA problem). The following two lemmas show that in terms of its computational power (regardless of run-time considerations), an $n\text{FSM}$ protocol is essentially equivalent to an rLBA.

Lemma 6.1. *An $n\text{FSM}$ protocol on a graph G of arbitrary topology can be simulated by an rLBA.*

Proof. The input for the Turing machine is the graph G , given as an adjacency list. In order to simulate the execution of the nFSM protocol, we store some additional information in the entries of the adjacency list as follows: For each node v , we store its current state and the next letter it transmits. For every node u in the list of neighbors $N(v)$ attached to v , we store the entry of u 's port that corresponds to v . In each round of the nFSM protocol, the rLBA performs two sweeps of the list of nodes: The first sweep serves to calculate v 's next state q and transmitted letter σ for all nodes v , based on v 's current state and the messages in its ports, according to the nFSM state machine, which is hard-wired in the rLBA. However, the calculated letter σ is not being “transmitted” yet, so the calculations for subsequent nodes in the list are not messed up, but rather stored in the corresponding place next to v . In the second sweep, for every node v , the letter σ is being “transmitted”, that is, the lists of neighbors are traversed, and at each occurrence of v , the current letter is replaced by σ . This way, we simulate every round of the nFSM protocol. In total, our simulation requires additional $O(1)$ space per node and $O(1)$ space per edge, hence it can be implemented with an rLBA. The assertion follows. \square

Lemma 6.2. *An rLBA can be simulated by an nFSM protocol on a path.*

Proof. Let n be the number of cells in the tape of the rLBA. Then, the path network has n nodes, each corresponding to one cell of the tape, i.e., we identify a node v of the path nFSM with a certain cell on the tape. Let Γ be the working alphabet and P be the state space of the rLBA. The nFSM protocol is designed so that the state of node v indicates: (1) which letter from Γ is written in v ; (2) if the head of the rLBA currently points to v ; (3) the current state of the rLBA, which is allowed to be incorrect if (2) is false; and (4) if the head is currently located to the left or to the right of v . Hence, we fix $Q = \Gamma \times \{0, 1\} \times P \times \{L, R\}$. The alphabet of the nFSM is $\Sigma = \{L, R\} \times P$.

Suppose that the input to the rLBA is $\gamma_1 \dots \gamma_n \in \Gamma^n$. Then, we assume that the initial state of the i th node in the path is (γ_i, h, p_0, L) , where p_0 is the initial state of the Turing machine and

$$h = \begin{cases} 1 & \text{if } i = 1 \\ 0 & \text{if } i > 1. \end{cases}$$

Note that the distinction between the initial state of the first node in the path and the initial states of all other nodes is without loss of generality. Indeed, as the first and last nodes have degree 1 and all interior nodes have degree 2, it is easy for a node to “decide” (under the nFSM model) if it is an interior node. Distinguishing between the first and last nodes is unavoidable if one wants to distinguish between the inputs $\gamma_1 \dots \gamma_n$ and $\gamma_n \dots \gamma_1$.

At all times, we maintain the invariant that exactly one node is in a state in $\Gamma \times \{1\} \times P \times \{L, R\}$ — denote this node as *active* — whereas all other nodes are in a state in $\Gamma \times \{0\} \times P \times \{L, R\}$. Only the active node can transmit messages; all other nodes remain silent and listen. If a non-active node v receives a message indicating that the head should move to the left (respectively, right), and v 's state indicates that the head is currently to its right (resp., left), then v becomes the active

node; otherwise, v does not react to this message. Now, the nodes simulate the behavior of the rLBA by calculating the next state of the rLBA based on the rLBA's transition function (which is hard-wired in the FSM) and updating their own states accordingly. The assertion follows. \square

References

- [1] Y. Afek, N. Alon, Z. Bar-Joseph, A. Cornejo, B. Haeupler, and F. Kuhn. Beeping a maximal independent set. In *Proceedings of the 25th international conference on Distributed computing (DISC)*, pages 32–50, 2011.
- [2] Y. Afek, N. Alon, O. Barad, E. Hornstein, N. Barkai, and Z. Bar-Joseph. A Biological Solution to a Fundamental Distributed Computing Problem. *Science*, 331(6014):183–185, Jan. 2011.
- [3] N. Alon, L. Babai, and A. Itai. A fast and simple randomized parallel algorithm for the maximal independent set problem. *J. Algorithms*, 7:567–583, December 1986.
- [4] B. Awerbuch. Complexity of network synchronization. *J. ACM*, 32(4):804–823, 1985.
- [5] B. Awerbuch, B. Patt-Shamir, D. Peleg, and M. E. Saks. Adapting to asynchronous dynamic networks (extended abstract). In *STOC*, pages 557–570, 1992.
- [6] B. Awerbuch and D. Peleg. Network synchronization with polylogarithmic overhead. In *FOCS*, pages 514–522, 1990.
- [7] L. Barenboim and M. Elkin. Distributed $(\Delta+1)$ -coloring in linear (in Δ) time. In *STOC*, pages 111–120, 2009.
- [8] L. Barenboim and M. Elkin. Combinatorial algorithms for distributed graph coloring. In *DISC*, pages 66–81, 2011.
- [9] L. Barenboim and M. Elkin. Deterministic distributed vertex coloring in polylogarithmic time. *J. ACM*, 58(5):23, 2011.
- [10] M. Bellare, O. Goldreich, and M. Sudan. Free bits, pcps, and nonapproximability-towards tight results. *SIAM J. Comput.*, 27(3):804–915, 1998.
- [11] Y. Benenson, T. Paz-Elizur, R. Adar, E. Keinan, Z. Livneh, and E. Shapiro. Programmable and autonomous computing machine made of biomolecules. *Nature*, 414(6862):430–434, Nov. 2001.
- [12] D. Brand and P. Zafiropulo. On communicating finite-state machines. *J. ACM*, 30:323–342, April 1983.
- [13] I. Chlamtac and S. Kutten. On Broadcasting in Radio Networks—Problem Analysis and Protocol Design. *Communications, IEEE Transactions on [legacy, pre - 1988]*, 33(12):1240–1246, 1985.
- [14] N. Chomsky. Three models for the description of language. *IRE Transactions on Information Theory*, 2:113–124, 1956. <http://www.chomsky.info/articles/195609--.pdf>.

- [15] R. Cole and U. Vishkin. Deterministic coin tossing with applications to optimal parallel list ranking. *Inf. Control*, 70(1):32–53, July 1986.
- [16] A. Cornejo and F. Kuhn. Deploying wireless networks with beeps. In *Proceedings of the 24th international conference on Distributed computing (DISC)*, pages 148–162, 2010.
- [17] R. Flury and R. Wattenhofer. Slotted Programming for Sensor Networks. In *International Conference on Information Processing in Sensor Networks (IPSN)*, Stockholm, Sweden, April 2010.
- [18] M. Gardner. The fantastic combinations of John Conway’s new solitaire game ‘life’. *Scientific American*, 223(4):120–123, 1970.
- [19] A. V. Goldberg, S. A. Plotkin, and G. E. Shannon. Parallel symmetry-breaking in sparse graphs. *SIAM J. Discrete Math.*, 1(4):434–446, 1988.
- [20] P. Gordon. Numerical Cognition Without Words: Evidence from Amazonia. *Science*, 306(5695):496–499, Oct. 2004.
- [21] K. Kothapalli, C. Scheideler, M. Onus, and C. Schindelhauer. Distributed Coloring in $\tilde{O}(\sqrt{\log n})$ Bit Rounds. In *20th International Parallel and Distributed Processing Symposium (IPDPS)*, 2006.
- [22] F. Kuhn. Weak graph colorings: distributed algorithms and applications. In *Proceedings of the twenty-first annual symposium on Parallelism in algorithms and architectures*, SPAA ’09, pages 138–144, New York, NY, USA, 2009. ACM.
- [23] F. Kuhn, T. Moscibroda, and R. Wattenhofer. What cannot be computed locally! In *Proceedings of the twenty-third annual ACM symposium on Principles of distributed computing (PODC)*, pages 300–309, 2004.
- [24] S.-Y. Kuroda. Classes of languages and linear-bounded automata. *Information and Control*, 7(2):207–223, 1964.
- [25] C. Lenzen and R. Wattenhofer. MIS on trees. In *Proceedings of the 30th annual ACM SIGACT-SIGOPS symposium on Principles of distributed computing (PODC)*, pages 41–48, New York, NY, USA, 2011.
- [26] N. Linial. Locality in distributed graph algorithms. *SIAM J. Comput.*, 21:193–201, Feb. 1992.
- [27] M. Luby. A simple parallel algorithm for the maximal independent set problem. *SIAM J. Comput.*, 15:1036–1055, November 1986.
- [28] N. A. Lynch. *Distributed Algorithms*. Morgan Kaufmann, 1st edition, 1996.

- [29] Y. Métivier, J. M. Robson, N. Saheb-Djahromi, and A. Zemmari. An optimal bit complexity randomised distributed MIS algorithm. *Distributed Computing*, 23(5-6):331–340, Jan. 2011.
- [30] J. V. Neumann. *Theory of Self-Reproducing Automata*. University of Illinois Press, Champaign, IL, USA, 1966.
- [31] D. Peleg. *Distributed computing: a locality-sensitive approach*. Society for Industrial and Applied Mathematics, Philadelphia, PA, USA, 2000.
- [32] S. Plotkin. *Graph-theoretic techniques for parallel, distributed, and sequential computation*. MIT/LCS/TR. Laboratory for Computer Science, Massachusetts Institute of Technology, 1988.
- [33] D. Sadava. *Life: The Science of Biology*. Sinauer Associates, 2011.
- [34] J. Schneider and R. Wattenhofer. An Optimal Maximal Independent Set Algorithm for Bounded-Independence Graphs. In *Journal of Distributed Computing*, March 2010.
- [35] J. Schneider and R. Wattenhofer. Distributed Coloring Depending on the Chromatic Number or the Neighborhood Growth. In *18th International Colloquium on Structural Information and Communication Complexity (SIROCCO), Poland*, June 2011.
- [36] J. Suomela. Survey of local algorithms. *To appear in: ACM Computing Surveys*, 2012. <http://www.cs.helsinki.fi/u/josuomel/doc/local-survey.pdf>.
- [37] M. Szegedy and S. Vishwanathan. Locality based graph coloring. In *STOC*, pages 201–207, 1993.
- [38] S. Wolfram. *A new kind of science*. Wolfram Media, Champaign, Illinois, 2002.