Parallel Local Search

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Contents

1

Introduction

A wide variety of problems in practical planning and design situations is concerned with the choice of the best solution from a finite, possibly large, number of alternatives. Many of these problems can be modelled as combinatorial optimization problems. Combinatorial optimization problems are often computationally intractable and, consequently, larger instances of such problems can typically be solved only to proximity.

Local search is a generally applicable approximation technique for combinatorial optimization problems that is able to find good quality solutions, albeit at the expense of substantial running times. It is often argued that parallelism can be used to reduce these running times, which makes it possible to handle larger instances in a given amount of running time, or to find better solutions for a given instance in an equal amount of running time. In this thesis we investigate the potentials of parallel processing for local search.

1.1 Combinatorial optimization

In this section we give a brief introduction to combinatorial optimization. More elaborate introductions can be found in [Papadimitriou & Steiglitz, 1982; Nemhauser & Wolsey, 1988]. A combinatorial optimization problem is either a minimization problem or a maximization problem consisting of a set of problem instances. Without loss of generality, we consider only minimization problems.

Definition 1.1. An *instance* of a combinatorial optimization problem is a pair $(\mathcal{S}, f)$, where $\mathcal{S}$ is a finite set of solutions, and $f : \mathcal{S} \to \mathbb{Z}$ is a function that gives the cost of a solution. The objective is to find a solution in $\mathcal{S}$ with minimal cost. □

The *size* of an instance I, denoted by $size(I)$, is defined as the number of symbols needed to encode I in a compact way. The finiteness of the set $\mathcal{S}$ suggests that an instance $(\mathcal{S}, f)$ can be solved by examining all solutions and selecting the one with minimal cost. Such an enumeration approach is only practical for very small instances if the size of $\mathcal{S}$, which is denoted by $|\mathcal{S}|$, grows superpolynomially with the instance size. The *time complexity function* $t_A : \mathbb{N} \to \mathbb{N}$ of an algorithm A gives for each instance size the largest running time needed by A for solving an instance of that size, where running time is measured in the number of elementary operations such as assignments, comparisons, etc. Hence, $t_A(size(I))$ is an upper bound on the running time needed to solve an instance I. To compare algorithms, one is often interested in the order of their complexity functions. A function f is $\mathcal{O}(f')$ with $f, f' : \mathbb{N} \to \mathbb{N}$ if there exist constants $c, m \in \mathbb{N}$ such that $f(n) \leq c \cdot f'(n)$ for all $n > m$. f is $\Omega(f')$ if there exist constants $c, m \in \mathbb{N}$ such that $f(n) \geq c \cdot f'(n)$ for all $n > m$, and f is $\Theta(f')$ if f is both $\Omega(f')$ and $\mathcal{O}(f')$. An algorithm A is a polynomial-time algorithm if $t_A = \mathcal{O}(f)$ for some polynomial function f. Otherwise A is a superpolynomial-time algorithm. Problems for which a polynomial-time algorithm is known, which implies that instances I can be solved in $size(I)^{\mathcal{O}(1)}$ time, are often called 'easy'. For many other problems no such algorithms are known despite considerable effort to find them. The difference between these kinds of problems is formalized by the theory of NP-completeness. Garey & Johnson [1979] present an overview of the NP-completeness theory. This theory is based on *decision problems* in which one is asked to determine whether there exists a solution with cost of at most $k \in \mathbb{Z}$. A decision problem has only two possible solutions, either 'yes' or 'no'. With each optimization problem a decision problem can be associated. Next, two classes of problems are introduced, called P and NP, with P $\subseteq$ NP. These classes are used to distinguish between easy and hard problems.

Definition 1.2. P is the class of decision problems for which each instance can be solved by a polynomial-time algorithm. □

A *concise certificate* for an instance I is an amount of data with size polynomial in $size(I)$. The class NP can now be defined as follows.

Definition 1.3. NP is the class of decision problems for which each instance I with answer 'yes' has a concise certificate c such that a 'yes' answer for I can be verified by a polynomial-time algorithm using c. □

The concept of reducibility has been proven useful for relating the computational complexity of two decision problems.

Definition 1.4. Problem π is *polynomially reducible* to problem π' if a polynomial-time algorithm exists that maps each instance I of π onto instance I' of π' such that I is a 'yes' instance of π if and only if I' is a 'yes' instance of π'. □

If π is polynomially reducible to π', then π' is at least as difficult as π. This leads to a class of problems in NP that can be considered as the most difficult ones in NP. These problems are called *NP-complete*.

Definition 1.5. A problem $\pi \in$ NP is *NP-complete* if each problem in NP is polynomially reducible to π. The class of NP-complete problems is called NPC. □

Reducibility can be used to prove that a problem π is NP-complete. From Definitions 1.4 and 1.5 it follows that in order to prove that π is NP-complete it suffices to show that $\pi \in$ NP and that a problem $\pi' \in$ NPC is polynomially reducible to π. Note that, if one NP-complete problem can be solved in polynomial time, then all NP-complete problems can be solved in polynomial time. However, so far no polynomial-time algorithm has been designed for any NP-complete problem and it is widely believed that NP-complete problems are *intractable*—that is, any algorithm that solves each instance of an NP-complete problem requires superpolynomial running time.

Finally, a problem is *NP-hard* if it is as least as difficult as any problem in NP, i.e., any problem in NP is polynomially reducible to it. Hence, each NP-complete problem is NP-hard, and if the decision variant of a combinatorial optimization problem is NP-complete, then the optimization problem is NP-hard.

Handling NP-hard problems. Two approaches can be distinguished to handle NP-hard problems. Either one searches for optimal solutions, at the risk of very large, possibly excessive, running times, or one is satisfied with relatively quickly obtainable solutions at the risk of sub-optimality. The first approach finds optimal solutions and accepts the possibility of superpolynomial running times. It seeks to achieve as much improvement as possible over straightforward enumeration, because often many solutions can be discarded as non-optimal without enumerating them explicitly. Dynamic programming, branch-and-bound, and branch-and-cut algorithms are examples of such implicit enumeration techniques.

For many NP-hard optimization problems one has to resort to approximation algorithms since larger instances cannot be solved optimally in acceptable running times. Approximation algorithms aim to find good-quality suboptimal solutions in a moderate amount of time. Two classes of approximation algorithms can be distinguished, viz. *constructive* algorithms and *local search* algorithms.

Constructive algorithms build a solution by starting with an empty solution and consecutively adding elements to the partial solution until a complete solution is obtained. Constructive algorithms typically run in polynomial time as each step requires polynomial time and the total number of steps is also bounded polynomially, but they may find solutions of mediocre quality. Local search algorithms constitute a class of approximation algorithms that are based on repeatedly replacing a solution by a neighboring solution. They often find good-quality solutions but may require substantial running times. Local search is discussed in more detail in Chapter 2.

Analysis of algorithms. The quality of an approximation algorithm is judged by the quality of the solutions it produces, its *effectiveness*, and by the time it requires to obtain them, its *efficiency*. The quality of solutions is measured by their *excess ratio*, the relative deviation of the costs of solutions from optimal solutions. Both effectiveness and efficiency can be considered from a worst-case and an average-case point of view. Worst-case analysis gives upper bounds on the average-case performance. Two approaches can be used to study the average-case behavior, namely, probabilistic or empirical analysis. A probabilistic average-case analysis aims at determining the expected performance of an algorithm by presupposing a probability distribution over the set of problem instances. An empirical analysis is based on a large number of numerical experiments on a set of benchmark instances either originating from practice or randomly generated according to some probability distribution. If optimal solutions are not known, the excess ratio is usually computed using a lower bound for the optimal solution, which establishes an upper bound for the empirical average-case performance. For many approximation algorithms a worst-case or average-case analysis is quite complicated and, therefore, one often resorts to an empirical average-case analysis. Another option is to compare the performance of an algorithm with that of other algorithms in order to determine a relative ranking of algorithms. Here it is essential that a challenging set of test instances is available.

1.2 Parallel processing

In this section we discuss some preliminaries for parallel processing that are relevant for the work described in this thesis. A more elaborate introduction into parallel processing can be found in [Bertsekas & Tsitsiklis, 1989].

1.2.1 Parallel machines

We first discuss some issues related to parallel machine architectures. Our discussion is mainly based on the classification of parallel architectures presented by Bertsekas & Tsitsiklis [1989].

Over the years many parallel machine models have been proposed. Unfortunately, no theoretical model effectively describes the full spectrum of parallel machines. Some models are based on technological developments, while others are based on theoretical observations. The most widely used classification of parallel computers is based on the distinction between *single instruction, multiple data* (SIMD) and *multiple instruction, multiple data* (MIMD) machines. In SIMD machines one global instruction is performed at a time, possibly on different data elements. In MIMD machines different instructions on different data elements can be performed simultaneously. Most modern parallel computers belong to the class of MIMD machines.

Based on the timing of instructions on different processing units it is possible to distinguish between *synchronous* and *asynchronous* machines. A synchronous machine uses one global clocking scheme to synchronize instructions among processors. If several local clocking schemes exist in a parallel machine, typically one per processing unit, the system is called asynchronous. SIMD machines are synchronous by definition, while MIMD machines are mainly asynchronous.

The above model does not address the way information is exchanged between processors. There are two possibilities for exchanging information. In *shared-memory* systems processors may read and write in a common memory accessible by each processor. In *message-passing* systems processors exchange data through messages. Message-passing systems can be characterized by their interconnection network topology that describes how processors are connected. The most common topologies are the ring and the torus. In a ring each processor is connected with two other processors such that a cycle is formed. In a torus processors are positioned in a two-dimensional grid. Processors are connected to four other processors, and each row and column forms a ring. Other frequently used topologies are the tree, and the hypercube. Some parallel computing environments permit the usage of *virtual topologies* in which topologies are simulated on the machine at hand in a user transparent way. It is of course important for the system's performance that the virtual topology is mapped efficiently onto the physical topology.

Other distinctions made between parallel machines are based on the amount of data each processor can handle and the number of processors. In *fine-grained* systems each processor can handle only a small amount of data, while in *coarse-grained* systems each processor deals with a large amount of data. *Massively parallel* systems consist of a large number of, usually relatively simple, processors. Massively parallel systems are typically fine-grained.

An important class of machine models in parallel complexity theory is the *parallel random access machine* (PRAM) model. A *concurrent read, concurrent* *write* PRAM is a shared memory machine in which all processors can simultaneously read from and write to the same memory locations. A *concurrent read, exclusive write* PRAM is a shared memory machine in which all processors can simultaneously read from the same memory location, but exclusive access to a memory location is required for writing. Although none of the currently available parallel machines are true PRAM's, this model supports the study of the intrinsic parallelism in problems and provides theoretical lower bounds for the time complexity of many parallel algorithms. The PRAM model can be adapted to capture features of more realistic parallel machines by considering memory partitioning, communication latency, and sparse network topologies, which all result in less powerful variants of the above PRAM model.

Features of a parallel machine at hand, besides the computational power of processors, can have a large influence on the performance of algorithms when executed on that machine. Important properties of a message-passing MIMD machine are the latency for starting up communication, and the throughput of communication between processors, which also depends on the interconnection network topology. For shared-memory MIMD machines the access time to various parts of memory plays an important role. Computational experiments are therefore important in the study of parallelism.

1.2.2 Parallel algorithms

Useful criteria to evaluate a parallel algorithm for a given problem are its *speed-up* and *efficiency*. The speed-up is the running time of the best known sequential algorithm for this problem divided by the parallel running time, and the efficiency is the speed-up divided by the number of processors [Barr & Hickman, 1993]. Another important issue is the *scalability* of an algorithm, which refers to its performance for increasing number of processors.

It is usually required that the output of parallel algorithms is independent of the number of employed processors. In our research, however, the number of employed processors may determine the part of the solution space that is explored, which may affect the output. Furthermore, we consider randomized algorithms for which different runs may require different running times. As the quality of the output of the local search algorithms we study is measured using the relative excess of final solutions over global minima, we define speed-up as follows.

Definition 1.6. Let $t_{A,\epsilon}(1)$ be the average time to find a solution with a relative excess ϵ over the global minimum using sequential algorithm A and let $t_{B,\epsilon}(P)$ be the average time for this using parallel algorithm B on machine M with P processors, then the *speed-up* of B over A on M is given by $t_{A,\epsilon}(1)/t_{B,\epsilon}(P)$. □

Other useful criteria, besides speed-up and efficiency, for judging the quality of the parallel algorithms we study in this thesis are the extend in which these parallel algorithms are able to find better quality solutions than sequential algorithms in a given amount of time. Also the instance sizes that can be handled by sequential and parallel algorithms in a given amount of time may be of interest. These criteria are of course closely related to the speed-up of algorithms as algorithms with a large speed-up can typically find better quality solutions in a given amount of time since more solutions can be explored. The reduction of running times for given instance sizes can also be employed to handle larger instances in a given amount of time.

Two issues are important in designing parallel algorithms with a good speed-up, viz., *communication overhead* and *load balancing*. On most parallel machines communication is costly compared to computation operations, so attention has to be paid to the communication behavior of parallel algorithms. Load balancing is important to achieve that no processor ever becomes idle while others are working.

1.2.3 Parallel complexity

It is often attempted to define a class of problems for which parallel processing can be profitable. A traditional example is the complexity class NC that consists of the problems that can be solved on a PRAM with a polynomial number of processors $n^{\mathcal{O}(1)}$ in polylogarithmic time $(\log n)^{\mathcal{O}(1)}$, i.e., a time bounded by a polynomial in the logarithm of the instance size n. Furthermore, a class of *P-complete* problems is identified that contains problems that are believed to be among the hardest problems in P. A problem is P-complete if it belongs to P and any other problem in P can be transformed to it using polylogarithmic time and polynomially many processors [Greenlaw, Hoover & Ruzzo, 1995]. It is conjectured that the class of P-complete problems is disjoint from the class NC, and consequently it is believed that P-complete problems cannot be solved in polylogarithmic time.

In practical situations one is often interested in parallel algorithms that solve an instance roughly P times faster when using P processors. Such algorithms may still exist for P-complete problems. So the distinction between P-complete problems and problems in NC is inadequate to capture the informal notion of problems that are amenable for parallel processing. Kruskal, Rudolph & Snir [1990] introduce a class EP of efficient parallel algorithms, in which they measure the performance of parallel algorithms relative to that of sequential algorithms. The usual yardstick is the best existing sequential algorithm for a given problem. EP is defined as follows.

Definition 1.7. Let the time complexity of a sequential algorithm and a parallel algorithm using $P(n)$ processors for instances of size n be given by $t(n)$ and

$T(n)$, respectively. An algorithm is polynomially fast and has constant efficiency, if $T(n) = \mathcal{O}(t(n)^{\epsilon})$ with $\epsilon < 1$ and $T(n) \cdot P(n) = \mathcal{O}(t(n))$. The class that contains these algorithms is called EP. □

Although the class EP does not define strict complexity classes for problems since it depends on a particular sequential algorithm, it provides a practically relevant classification of parallel algorithms.

1.3 Thesis outline

The objective of the research described in this thesis is to design parallel local search algorithms that can handle large problem instances and that are competitive with the best known sequential algorithms both with respect to running time and quality of final solutions. For this, we study general approaches that can be applied to a wide variety of problems, as well as tailored approaches in which problem characteristics are exploited. Computational experiments on parallel machines are considered as an important aspect of this research.

The remainder of this thesis is organized as follows. Chapter 2 deals with local search. In Chapter 3 we introduce the concepts for designing parallel local search algorithms. This chapter is based on [Verhoeven & Aarts, 1995a]. Chapter 4 presents some results for the speed-up that can be obtained by performing several independent runs of a local search algorithm in parallel. These results are based on a probabilistic analysis of local search that can also be found in [Ten Eikelder, Verhoeven, Vossen & Aarts, 1996]. Chapters 5, 6, and 7 discuss tailored approaches to design parallel local search algorithms for some well-studied combinatorial optimization problems. In Chapter 5 we study the traveling salesman problem, in Chapter 6 the Steiner tree problem in graphs, and in Chapter 7 job shop scheduling. Chapter 7, furthermore, discusses sequential and parallel local search for resource-constrained project scheduling. These chapters are based on [Verhoeven, Aarts, Van de Sluis & Vaessens, 1992; Verhoeven & Aarts, 1994; Verhoeven, Aarts & Swinkels, 1995], [Verhoeven, Aarts & Severens, 1995], and [Verhoeven & Aarts, 1995b], respectively.

The main conclusion that can be drawn from the work described in this thesis is that local search algorithms for a wide variety of problems can be sped up significantly using parallelism based on hybrids of the approaches to parallel local search proposed in this thesis.

2

Local Search

Local search algorithms constitute a class of approximation algorithms for hard combinatorial optimization problems that are based on the exploration of neighborhoods of solutions. They have shown to be successfully applicable to a wide range of problems, giving good-quality solutions [Aarts & Lenstra, 1996]. The basic local search algorithm is the so-called *iterative improvement* algorithm. An iterative improvement algorithm starts off with an initial solution constructed by some heuristic. Next, the algorithm repeatedly tries to improve the current solution by replacing it with a lower-cost neighbor. If a solution has been reached that has no neighbors with lower cost, a local minimum has been found. An essential concept in local search algorithms is the notion of a neighborhood structure.

Definition 2.1. Let $(\mathcal{S}, f)$ be an instance of a combinatorial optimization problem. Then a *neighborhood structure* $\mathcal{N} : \mathcal{S} \to \mathcal{P}(\mathcal{S})$ assigns to each solution in $\mathcal{S}$ a set of solutions, called a *neighborhood*. A solution $s \in \mathcal{S}$ is a *local minimum* of $\mathcal{N}$ if $f(s') \geq f(s)$ for each $s' \in \mathcal{N}(s)$. $\square$

The neighbors of a solution are not given explicitly but are to be constructed by a function. Most neighborhoods are based on the replacement of a few elements that constitute a solution. For this we introduce the following definition in which we assume that solutions are sets. The set of *building elements* that constitute solutions in $\mathcal{S}$ is then given by $\mathcal{E} = \{e \mid e \in s \wedge s \in \mathcal{S}\}$.

---

proc *Iterative_Improvement* $(s : \mathcal{S})$
  var $w : \mathcal{P}(\mathcal{S})$;
  begin
    $w := \emptyset$;
    while $\mathcal{N}(s) \setminus w \neq \emptyset$ **do**
      $s' \in \mathcal{N}(s) \setminus w$;
      if $f(s') \geq f(s)$ **then** $w := w \cup \{s'\}$
      else $s := s'; w := \emptyset$ **fi**
    od $\{s$ is a local minimum of $\mathcal{N}\}$
  end

Figure 2.1: An iterative improvement algorithm.

Definition 2.2. Let the *exchange function* $\tau_r : \mathcal{S} \times \mathcal{E}^r \to \mathcal{S}$ be a partial function with $r \in \mathbb{N}$. Then, an r-exchange neighborhood structure $\mathcal{N}_r$ associated with τ_r is defined by $\mathcal{N}_r(s) = \{\tau_r(s, e_1, \ldots, e_r) \mid e_1, \ldots, e_r \in s\}$ for $s \in \mathcal{S}$. □

According to Definition 2.2 neighbors of a solution s can be constructed by applying the exchange function to s and a subset $e_1, \ldots, e_r$ of the building elements of s. Below, we give an example of an exchange function and a neighborhood for one of the best-known combinatorial optimization problems, viz. the traveling salesman problem (TSP). In the TSP a salesman wishes to visit all cities in a given set once and return to the starting point, in such a way that the total distance covered is as short as possible.

Example 2.1 (traveling salesman problem). Given is a complete weighted graph $(V, V \times V)$, where V is a set of N vertices and $d_{ij} \in \mathbb{N}$ gives the distance between each $i, j \in V$. A *tour* t is a set of N directed edges $\{e_1, \ldots, e_N\}$ that constitutes a *Hamiltonian cycle*, a cycle that visits each vertex in this graph precisely once. The solution space $\mathcal{S}$ of a TSP instance is the set of all tours. The cost function f is given by

$$f(t) = \sum_{(i,j) \in t} d_{ij}.$$

The problem is to find a tour $t \in \mathcal{S}$ for which $f(t)$ is minimal. The exchange function $\tau_2 : \mathcal{S} \times (V \times V)^2 \to \mathcal{S}$ gives for each tour a new tour obtained by replacing two edges with two other edges. The well-known 2-exchange neighborhood of Lin [1965] is defined by $\mathcal{N}_2(t) = \{\tau_2(t, e_i, e_j) \mid 1 \leq i < j \leq N\}$. □

Figure 2.1 presents a schematic description of an iterative improvement algorithm. A single iteration of an iterative improvement algorithm, which we call a *step*, consists of the following actions. First, a neighbor is constructed by applying the exchange function to the current solution. Subsequently, the cost of the neighbor is evaluated. If the neighbor has lower cost, it replaces the current solution by effectuating the proposed exchange. Otherwise the search is continued with the old current solution. A *pivoting rule* [Yannakakis, 1990] determines which neighbor will be the new current solution. Well-known pivoting rules are *first improvement*, which replaces the current solution with the first lower-cost neighbor that is found, and *best improvement*, which uses the solution that has lowest cost among all neighbors.

For practical reasons it is often required that each local search step can be done in polynomial time. For this it is necessary that neighbors can be selected in polynomial time, which requires that the exchange function associated with a neighborhood structure has polynomial time complexity, and that the cost difference $f(s') - f(s)$ of two neighboring solutions s and s' can be computed in polynomial time.

2.1 Computational complexity of local search

Unless a neighborhood is *exact*, i.e., each local minimum is a global minimum, it is generally not possible to give a fixed upper bound on the relative excess of local minima. Moreover, for the iterative improvement algorithm of Figure 2.1 it is in general not possible to give a non-trivial upper bound on the number of steps needed to find a local minimum. Examples are known of problem instances for which the number of steps cannot be bounded by a polynomial in the size of the instance. Therefore, the question has been raised whether this is a general property of local search algorithms. Johnson, Papadimitriou & Yannakakis [1988] have formalized the question of the worst-case behavior of local search algorithms by introducing a new complexity class PLS. A *local search problem* Π is a set of problem instances of which each instance is characterized by a triple $(\mathcal{S}, f, \mathcal{N})$, where $\mathcal{S}$ is the set of solutions, f the cost function, and $\mathcal{N}$ the neighborhood structure. The question is to find a local optimum of $\mathcal{N}$. The class PLS is then defined as follows.

Definition 2.3. A local search problem Π is in PLS if polynomially computable functions g and g' exist such that for an instance I of Π, $g(I)$ returns a start solution, $g'(I, s)$ returns a solution $s' \in \mathcal{N}(s)$ with $f(s') < f(s)$ for $s \in \mathcal{S}$ and if no such solution exists, g' returns s, which is then a local minimum of $\mathcal{N}$. $\square$

Informally stated, PLS is the class of local search problems for which each local search step requires polynomial time. The notion of PLS-reducibility has been introduced in order to define PLS-complete problems.

Definition 2.4. A local search problem Π is *PLS-reducible* to a local search problem Π' if there exists a polynomially computable function h that maps instances I of Π onto instances I' of Π' and a polynomial function h' that maps pairs of the form (solution of $h(I)$, I) onto solutions of I such that, if s is a local optimum for $h(I)$, then $h'(s, I)$ is locally optimal for I. □

Definition 2.5. A local search problem $\Pi \in$ PLS is *PLS-complete* if each problem in PLS is PLS-reducible to Π. □

To prove that a local search problem Π is PLS-complete it suffices to show that Π is included in PLS and that there exists a known PLS-complete problem Π' that can be PLS-reduced to Π. The generic PLS-complete problem is FLIP. The class of PLS-complete problems contains the hardest problems in PLS, and if one of these problems can be solved in polynomial time, then all others can. It is, however, conjectured that P_s is a strict subset of PLS, where P_s is the class of search problems that can be solved in polynomial time [Yannakakis, 1996]. Consequently, if this conjecture holds, superpolynomial running times might be required by any algorithm to find a local optimum for the PLS-complete problems.

2.2 Local search variants

A neighborhood structure imposes a directed graph on the solution space. The vertices of this *neighborhood graph* are the solutions, and there is an edge from a vertex to another vertex if this vertex is a neighbor of the first vertex. The course of a local search algorithm is characterized by a walk in this graph. Starting in some vertex, the costs of neighbors are evaluated, and a step to some neighbor is made. This process is repeated until some stop criterion is met. The effectiveness of local search depends on both the structure of the neighborhood graph, which is determined by the neighborhood structure, and the way the neighborhood graph is traversed, which is determined by the type of local search that is used. A neighborhood is *sufficiently connected* if there is a path in the neighborhood graph to an optimal solution from any solution, and it is *completely connected* if there is a path to each solution from any other solution.

A drawback of iterative improvement is that it may get stuck in poor-quality local minima. To overcome this drawback, one can either modify the neighborhood structure, or change the way the neighborhood graph is traversed. The first approach is problem specific, whereas the latter one is less problem dependent.

Most neighborhood structures are based on exchange functions that modify only a few elements, which results in small neighborhoods. In order to avoid the risk of getting stuck in a poor-quality local minimum, one may increase the size of the neighborhood by exchanging more elements. Another option is to exploit characteristics of the problem at hand to tailor the neighborhood to this problem. Examples of this approach are *variable-depth* neighborhood structures [Kernighan & Lin, 1970; Lin & Kernighan, 1973] and the quite similar *ejection chain* approach in [Reeves, 1993]. Variable-depth neighborhoods are based on a simple exchange function. A neighbor is obtained by constructing a sequence of exchanges, instead of a single exchange. An algorithm that uses a variable-depth neighborhood is often no longer regarded as a basic iterative improvement algorithm since it actually generates a sequence of solutions, which may contain solutions with higher cost than a preceding solution. The solution that eventually replaces the current solution is the solution in this sequence with minimal cost. The algorithm stops if this neighbor has higher cost than the current solution.

Another approach to escape from local minima is to accept neighbors with cost higher than that of the current solution. We mention the following local search variants, sometimes called *meta heuristics*, which all traverse the neighborhood graph in a different way to escape from local minima.

Iterated local search [Johnson, 1990; Martin, Otto & Felten, 1991] uses two neighborhoods $\mathcal{N}$ and $\mathcal{N}'$. The search starts with the first neighborhood $\mathcal{N}$ until a local minimum of $\mathcal{N}$ is found. If this local minimum has lower cost than the best solution found so far, this local minimum replaces it. Then, a neighbor of the best solution found so far is chosen from a second neighborhood $\mathcal{N}'$. Subsequently, the search is restarted from this neighbor using the first neighborhood. In this way local search is performed at two levels. At the lower level regular local search steps are made in the neighborhood graph induced by $\mathcal{N}$, but at the higher level large steps between local minima of $\mathcal{N}$ are made.

Tabu search [Glover, 1989; Glover, Taillard & De Werra, 1993] tries to direct the search into unexplored areas of the solution space once a local minimum is found, by memorizing the course of the search. To this end, a finite list of the most recently visited solutions, the *tabu list*, is constructed, whose length is given by the *tabu tenure*. In each step of a tabu search algorithm a neighbor s' is chosen from the neighborhood $\mathcal{N}(s)$ of the current solution s such that s' is not included in the tabu list. Subsequently, s is added to the tabu list. Hence, the set from which a neighbor s' of s is chosen depends on how s is reached, and no s' is chosen that has been visited in the k most recent steps, if the tabu tenure is k. By choosing solutions not on the trajectory to recently visited local minima, tabu search can be directed into other areas of the solution space.

Rather than storing complete solutions that have been visited in the tabu list, building elements associated with exchanges that have led to these solutions are included in the tabu list. If a step is made from a solution s to s' with $s' = \tau(s, e)$, where τ is an exchange function and e a building element of a solution, then an element e' for which $s = \tau(s', e')$ is added to the tabu list. If the tabu list is full, its oldest element is removed. Neighbors are obtained by performing exchanges that act on building elements that do not occur in the tabu list. An exchange is not permissible, or *tabu*, if it occurs on this list. In this way, the implicitly defined set of tabu solutions includes the k most recently visited solutions.

Most tabu search algorithms select the lowest-cost non-tabu neighbor in the neighborhood as the new current solution. If all neighbors are tabu, then the oldest element on the tabu list is removed, until a permissible neighbor can be selected. Since such a pivoting rule can be computationally expensive *candidate sets* can be used. Candidate sets are subsets of neighborhoods in which it is attempted to exclude high-cost neighbors from a neighborhood without exploring them explicitly. *Aspiration criteria* can be used to overrule the tabu status of an exchange when it seems promising in some sense, e.g., if it leads to a solution with lower cost than the best solution found so far.

The effectiveness of many tabu search algorithms is significantly increased by incorporating *intensification* and *diversification* of the search. Intensification of the search around good-quality local minima is used to search promising areas of the solution space more thoroughly. This can be implemented by keeping track of the best solutions found in the course of the search. If no improvement of the best solution is achieved for a number of steps, the search is restarted with one of these best solutions, where the search is forced to follow a different trajectory from this solution by associating a tabu status with each attempted exchange for this solution. This requires storage of tabu lists associated with the best solutions found in the course of the search.

Search diversification is used to ensure that other regions of the solution space are searched as well. Diversification of the search can be achieved by adding penalties to the cost of solutions if some of its elements occur frequently in the current solution or are included in the current solution for a large number of local search steps.

Simulated annealing [Kirkpatrick, Gelatt & Vecchi, 1983; Aarts & Korst, 1989] probabilistically accepts neighbors with higher costs, in addition to lower-cost neighbors, which are always accepted. To this end, an *acceptance probability* is used that is reversely related to the size of the cost increase. The acceptance probability of cost-increasing exchanges is given by

$$\exp(-\frac{(f(s') - f(s))}{c})$$

for solutions s and s' with $f(s') > f(s)$. The *control parameter c* is used to lower the acceptance probability in the course of the search.

A simulated annealing algorithm can be outlined as follows. First, an initial value for the control parameter c is chosen. For each value of c a number of randomly chosen exchanges is attempted, where proposed exchanges are accepted according to the above acceptance probability, after which the control parameter is lowered. If the control parameter c drops below a given value the search is ended. A *cooling schedule* determines how the control parameter c is decremented. In many practical simulated annealing algorithms a *geometric* cooling schedule is used in which c is decremented by multiplication with a constant factor α, with $\alpha < 1$.

Using finite Markov chain theory, it has been proven that simulated annealing converges to a global minimum if the acceptance probability is lowered slowly enough, provided that the employed neighborhood is at least sufficiently connected. This connectivity condition is not hard to meet for many problems.

Genetic local search [Mühlenbein, Gorges-Schleuter & Krämer, 1988; Aarts & Verhoeven, 1996] incorporates local search into genetic algorithms. In genetic local search, a *population* of solutions is explored, instead of only a single solution. A genetic local search algorithm starts with a population of p initial solutions. This population is augmented with q solutions obtained as follows. Two *parent* solutions are selected from the population. Using these two *parent* solutions a new solution is constructed by applying *cross-over*. In a cross-over operation building elements of two solutions are combined to one solution. This newly constructed solution is then improved using a local search algorithm and added to the population. The augmented population is subsequently reduced to its original size using concepts from biological evolution strategies.

Evidently, the cross-over operation is important, since here one must try to take advantage of the availability of more than one local minimum by exploiting their structure. *Mutation* in a traditional genetic algorithms corresponds with the part of the above algorithmic concept in which local search is applied to the solution resulting from the cross-over operation.

Widely recognized advantages of local search are its general applicability, its flexibility, and its ease of implementation. Only a specification of solutions, a cost function, and a neighborhood structure are required, which can easily be defined for many problems. Furthermore, practical experience in scheduling, layouting, and routing applications has revealed that the average-case performance of local search is quite good in the sense that high-quality solutions can be found within acceptable running times. This has led to the conviction that local search is a powerful technique to handle complex real-world combinatorial optimization problems that arise in management science and engineering. For more extensive overviews on local search in which the above variants are discussed in more detail we refer to [Yannakakis, 1990; Reeves, 1993; Aarts & Lenstra, 1996].

3

Concepts of Parallel Local Search

Although local search algorithms may find good-quality solutions, they require substantial amounts of running time for the larger instances of some combinatorial optimization problems; see for instance [Sechen, 1988], who reports running times of 24 hours for his simulated annealing based cell-placement algorithm when applied to real-life problem instances. To cope with this drawback, several approaches have been proposed in the literature to implement these algorithms on parallel machines. Other goals of these parallel approaches to local search are to find better quality solutions in a given amount of time or to enable handling of larger problem instances. In this chapter we try to disclose the underlying concepts on which these approaches are based. In this discussion we try to abstract from implementation issues that depend on the machine model at hand. Furthermore, we study some complexity issues of parallel local search.

Much effort in the field of parallel local search has concentrated on the design of parallel simulated annealing algorithms [Aarts & Korst, 1989; Azencott, 1992]. For an overview of parallel simulated annealing, we refer to [Greening, 1990]. More recently, also parallel approaches to other local search algorithms, such as tabu search and genetic local search, have been reported. Voss [1993] presents a classification of parallel tabu search algorithms based on the use of different search strategies and starting solutions. Crainic, Toulouse & Gendreau [1993b] extended this classification with dimensions based on communication organization and information handling.

We distinguish between *tailored* and *general* approaches. A tailored approach can be applied only to a specific problem, whereas a general approach can be applied to a wide variety of problems. Furthermore, we distinguish between single-walk and multiple-walk parallelism and between asynchronous and synchronous algorithms. In the class of single-walk algorithms, we distinguish between multiple-step and single-step parallelism.

Tailored approaches. A tailored approach to incorporate parallelism in local search requires parallel execution of a problem specific part of the algorithm. Cost computation is a typical problem dependent part of a local search algorithm. Also operations on the data structures used to represent a solution are problem specific. If one of these problem dependent parts is very time consuming, the running time of the local search algorithm can be decreased substantially by efficient parallel execution of this part.

Kravitz & Rutenbar [1987] apply a tailored approach to their local search algorithm for the standard cell placement problem. They compute the cost of a neighbor in parallel. Another example of parallel cost computation is given by Taillard [1994], who uses a parallel longest path algorithm to compute the length of a schedule in the job shop scheduling problem.

Tailored approaches have limited applicability because they strongly depend on the problem at hand. Therefore, we discard these approaches here and concentrate on techniques that can be applied to a wide variety of local search problems.

General approaches. A general approach can be applied to a broad class of local search problems. We can distinguish between two approaches: *single-walk* and *multiple-walk* parallel local search. In a single-walk algorithm only a single walk in the neighborhood graph is carried out, whereas in a multiple-walk algorithm several walks are performed simultaneously.

Within the class of multiple-walk algorithms we can distinguish between algorithms that perform interacting walks and algorithms that perform multiple independent walks. The latter can be considered as the most straightforward approach to introduce parallelism in local search.

In single-walk parallel local search a single walk is carried out in parallel. Typically, this requires some distribution of a solution or elements of a solution over a number of processors. Furthermore, we distinguish between *single-step* and *multiple-step* parallelism. The idea of single-step parallelism is to evaluate neighbors simultaneously and subsequently make a single step. To this end, the neighborhood of a solution is partitioned into subsets that are searched in parallel. In an algorithm with multiple-step parallelism several consecutive steps through the neighborhood graph are made simultaneously. Figure 3.1 shows neighborhood graphs that illustrate the idea of single-step and multiple-step parallelism.

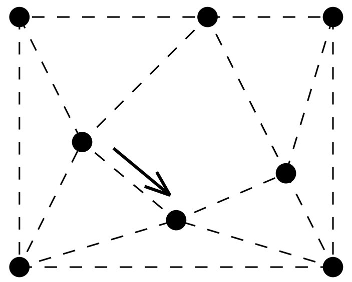

Figure 3.1: Local search with single-step parallelism (left) and multiple-step parallelism (right) in a neighborhood graph. Dotted edges connect neighbors and a solid arc represents a step made by a local search algorithm.

Finally, in both single-walk and multiple-walk parallel local search we distinguish between *synchronous* and *asynchronous* algorithms. In a synchronous algorithm one or more steps of the algorithm are performed simultaneously by all processors. Synchronous parallelism requires a global clocking scheme or token-passing mechanism that guarantees that communication occurs at given points in time. In asynchronous parallelism no such global clocking mechanism exists.

A multiple-walk algorithm performs multiple single walks. Hence, it is possible to combine multiple-walk and single-walk parallelism in the design of parallel local search algorithms. Also tailored approaches can be incorporated into general approaches resulting in all kinds of hybrid approaches, but the items in our classification are the basic concepts of such algorithms.

The remainder of this chapter is organized as follows. In Section 3.1 we discuss multiple-walk parallelism. Section 3.2.1 discusses single-step parallelism, and Section 3.2.2 discusses multiple-step parallelism in more detail. Finally, Section 3.3 discusses some complexity issues of parallel local search.

3.1 Multiple-walk parallelism

Multiple-walk algorithms have inherent parallelism, which can be exploited by distributing the walks over a number of processors and performing them simultaneously. We distinguish between algorithms that perform multiple independent walks and algorithms that perform multiple interacting walks. The first class of algorithms is the simplest one, since no complex parallelization scheme is required. In order to describe algorithms with multiple interacting walks more precisely, we extend the notion of neighborhood structures to hyper neighborhood structures. Neighborhood structures define neighbors for a single solution; hyper neighborhood structures define neighbors for a set of solutions [Vaessens, 1995].

---

proc *Multiple_Walks* $(s_1, \ldots, s_P \in \mathcal{S})$
  begin
    stop_criterion := false;
    while $\neg$ stop_criterion **do**
      par $p \in \Omega$ **do**
        Iterative_Improvement(s_p) ;
        $s_p \in \mathcal{H}(s_{h_p(1)}, \ldots, s_{h_p(M)})$
      rap
    od
  end

Figure 3.2: A multiple-walk parallel local search algorithm.

Definition 3.1. Let $M \geq 1$. Then, a *hyper neighborhood structure* $\mathcal{H} : \mathcal{S}^M \rightarrow \mathcal{P}(\mathcal{S})$ assigns to a set of M solutions in $\mathcal{S}$, a subset of solutions in $\mathcal{S}$. □

Let $\Omega = \{1, \ldots, P\}$ be a set of P processors, and let $h_p : \{1, \ldots, M\} \rightarrow \Omega$ be a function for each $p \in \Omega$, where $h_p(i)$ denotes the i-th processor with which p communicates. h_p gives, for each processor p, M other processors with which interaction takes place in order to construct a hyper neighbor.

Using the concept of hyper neighborhood structures, we can formulate the template presented in Figure 3.2 that describes algorithms with interacting multiple-walk parallelism. The set of current solutions $s_1, \ldots, s_P$ is often called the *population*. A current solution s_p is assigned to processor p. All processors perform simultaneously a walk in the neighborhood graph by executing an iterative improvement algorithm, or some other local search algorithm. After a number of local search steps, a hyper neighbor is constructed, which requires interaction with M other processors $h_p(1), \ldots, h_p(M)$. This process is continued until some stop criterion is met.

3.1.1 Multiple independent walks

A trivial approach to parallel local search is to perform a number of independent walks in parallel, since this requires no communication between different runs. The speed-up one hopes to achieve by using such an approach is based on the following property for random walks in a neighborhood graph.

Theorem 3.1. *Let $Q_p(t)$ be the probability of not having found a solution with relative excess ϵ in t time units with p independent walks, and let $Q_1(t) = e^{-\lambda t}$ with $\lambda \in \mathbb{R}^+$, i.e., Q_1 is distributed exponentially. Then, $Q_p(t) = Q_1(pt)$.* □

Hence, it is possible to achieve linear speed-up with multiple independent walks if the probability to find an optimal (or suboptimal) solution within a given amount of time units is distributed exponentially. An appropriate time measure for local search algorithms is the number of solutions of which the cost has been evaluated. Theorem 3.1 states that the probability to find a (sub)optimal solution in t time units and P processors is equal to the probability to find a (sub)optimal solution in Pt time with a single processor, if this probability is distributed exponentially. So in that case, a linear speed-up and an efficiency equal to one is reached with multiple independent walks.

Several authors have studied the probabilistic behavior of local search algorithms for various problems to investigate whether the condition of Theorem 3.1 holds in practice. Battiti & Tecchiolli [1992] empirically investigate the behavior of tabu search for randomly generated instances of the quadratic assignment problem. They observe that the probability of finding a (sub)optimal solution with a tabu search algorithm is indeed distributed exponentially, provided that the search is started with a local minimum. This indicates that good efficiencies can be obtained with multiple independent parallel walks in tabu search, if the time needed to find the first local minimum is relatively small compared to the time spent in the remainder of the search process. Taillard [1991] also shows that the probability of finding an optimal solution for quadratic assignment problems with a tabu search algorithm fits well with an exponential distribution. Dodd [1990] empirically shows that a similar conclusion also holds for simulated annealing. For the problem and parameter setting that he used an efficiency of one could be obtained with a maximum of 16 processors. Osborne & Gillett [1991] investigate the empirical behavior of simulated annealing for the Steiner tree problem in graphs. They also observe that the probability of finding a near-optimal solution is distributed exponentially. Chapter 4 presents some results for the average-case behavior of iterated local search algorithms for the traveling salesman problem. It is shown that the probability of finding a solution with a small excess over the global minimum using an iterated local search algorithm can be described by a geometrical distribution, which is the discrete counterpart of an exponential distribution, somewhat translated to compensate for the time needed to find the first local minimum. Good efficiencies can be obtained with multiple-walk parallelism but the amount of speed-up depends on the time needed to find the first local minimum, the time needed to find subsequent local minima, and the desired solution quality.

Shonkwiler & Van Vleck [1994] present a theoretical analysis of the speed-up with multiple independent stochastic walks using Markov-chain analysis. They show that the speed-up is given by $\frac{1-\lambda^p}{1-\lambda}s^{p-1}$ where p is the number of walks and s and λ are parameters that are related to the search process at hand. They claim that nearly always $\lambda \approx 1$ so then the speed-up is equal to ps^{p-1}. They present some artificial problems for which they are able to determine the parameters s and λ, and for some of these problems even superlinear speed-ups are achieved with multiple independent stochastic walks. Ferreira & Zerovnik [1993] give another theoretical result which shows that multiple independent walks can be profitable. They show that, after a certain amount of time, the probability to find a (sub)optimal solution with multiple runs of an iterative improvement algorithm is larger than the probability that it is found by a single simulated annealing run in the same amount of time.

The inherent parallelism of genetic algorithms is also based on Theorem 3.1, although in genetic algorithms the walks interact to combine good parts of individual solutions. The population size is therefore an important parameter of a parallel genetic algorithm, because it determines the speed-up and efficiency that can be obtained ultimately.

3.1.2 Multiple interacting walks

Interaction of walks can be used to faster direct the search into promising areas of the solution space. To model the interaction between parallel walks we use the concept of hyper neighborhoods. Hyper neighborhoods can be based on selection of a single solution from a set of solutions. Such hyper neighborhoods assign the same solution to different processors and are useful only if the walks performed by processors follow different paths from this solution. This can be achieved by either introducing randomization in walks or by enforcing different directions for walks. Other hyper neighborhoods are based on combining parts of different solutions to new solutions.

Processors exchange information when walks interact. This information, such as the occurrence of low-cost solutions in a population or the availability of multiple good parts of individual solutions, can be used to speed up the search. A possible way to use the additional information resulting from interaction of walks, is indicated by Lin & Kernighan [1973] and Kirkpatrick & Toulouse [1985], who observe that local minima often have many elements in common. These elements can be identified as the good parts of solutions. So the availability of multiple local minima allows fast identification of good parts of individual solutions that should not be changed in the course of the search. This then leads to a reduction of the sizes of neighborhoods.

Implementation issues. First, we discuss asynchronous and synchronous multiple walks. In synchronous multiple-walk parallelism, walks are synchronized at certain points in time—that is, the construction of a hyper neighbor can only be done in a state that depends on the state of all processors. In asynchronous multiple-walk parallelism construction or selection of a hyper neighbor is not synchronized at a point in time. Here, the number of processors with which a processor interacts to construct a hyper neighbor is typically a subset of all processors. In that case a processor can resume searching as soon as communication with that subset of processors has taken place, and it does not need to wait for other processors.

The amount of communication overhead involved with multiple-walk parallelism depends on how often a hyper neighbor is constructed, the hyper neighborhood structure at hand, the machine architecture, and on the mapping of walks on to processors. It is preferable to map interacting walks on to the individual processors of a given machine such that the resulting communication overhead is minimal. Load balancing is done by specifying an equal amount of work that is to be performed in between two interactions of walks. This can be done by fixing the number of steps between two communications, which typically means that hyper neighbors are not always constructed from local minima, since the number of steps to find a local minimum is not equal in each walk. An advantage of asynchronous walks over synchronous walks is their reduced load imbalance.

Examples. Parallel genetic local search algorithms are typical examples of local search algorithms with multiple-walk parallelism. The following hyper neighborhood is used in genetic algorithms. Two parent solutions, selected from —a subset of— the current population, are used in a cross-over operation to construct a new solution. In a cross-over operation good parts of individual solutions are combined. Both synchronous and asynchronous multiple-walk parallelism can be applied in genetic algorithms. Asynchronous parallel genetic algorithms are mostly based on an *island model* in which parents are chosen from a subset of the population instead of from the entire population. Here, we do not further elaborate on parallel genetic algorithms; for more details we refer the reader to [Jog, Suh & Van Gucht, 1991; Michalewicz, 1992; Mühlenbein, 1992].

Several authors have applied multiple-walk parallelism to simulated annealing and tabu search. A hyper neighborhood of a population of solutions that is often used here is the selection of the best solution in the population. In a tabu search algorithm different paths from this solution should be followed by different processors. This can by done by blocking certain exchanges or by using different search parameters for different processors. Note that in this way an intensification of the search around good solutions is achieved. Additional diversification can be achieved by combining long-term memory of different walks.

Malek, Guruswamy & Pandya [1989] present tabu search and simulated annealing algorithms with synchronous multiple-walk parallelism in which the best solution in the population is communicated to all processors after a fixed number of steps of the simulated annealing or tabu search algorithm. In their algorithm the hyper neighborhood of a solution consists of a single element, viz., the best solution come across in the entire population. Different parameter settings are used to guarantee that different paths are followed by processors. They obtain a good speed-up for seven processors.

Crainic, Toulouse & Gendreau [1995] present various synchronous multiple-walk tabu search algorithms. They propose two hyper neighborhoods: one in which the best solution in the entire population is selected by all processors, and one in which each of the P processors selects a different solution among the P best solutions come across since the previous interaction. Different paths are followed by choosing a different parameter setting for processors. Typically, the best-quality solutions are found by multiple independent walks. Crainic, Toulouse & Gendreau [1993a] also present asynchronous multiple-walk parallel tabu search algorithms. They propose two hyper neighborhoods in which processors interact asynchronously with a central processor that keeps track of either the best solution or the P best solutions found in the search process, where P is the number of processors. In the first hyper neighborhood this overall best solution is selected as hyper neighbor, and in the latter hyper neighborhood a hyper neighbor is randomly chosen from the P best solutions found. Compared to their synchronous algorithms they obtain slightly better final solutions. Although actual running times of the various synchronous and asynchronous algorithms are not presented, the number of iterations after which the overall best solution is found implies that the algorithms display little speed-up.

Aarts, De Bont, Habers & Van Laarhoven [1986] present parallel simulated annealing algorithms based on a combination of multiple interacting walks and single-step parallelism in which the size of the population is gradually decreased in the course of the search. They propose the following two hyper neighborhoods. In the first one the best solution from the entire population is chosen ($M = P$), and in the second one a solution is randomly chosen from two solutions ($M = 2$). The first hyper neighborhood results in synchronous multiple-walk parallelism, the second one in asynchronous multiple-walk parallelism. Similar algorithms using these hyper neighborhoods are implemented by Diekmann, Lüling & Simon [1993], who obtain a speed-up of 85 on a network of 120 transputers for the traveling salesman problem.

Moscato [1993] and Fox [1993] present parallel algorithms that combine genetic algorithms, simulated annealing, and tabu search. They claim a linear speed-up, but no empirical evidence is given. Mahfoud & Goldberg [1995] propose a combination of genetic algorithms and simulated annealing, but so far they do not present results for combinatorial optimization problems.

3.2 Single-walk parallelism

In this section we discuss a template that captures local search algorithms with single-walk parallelism. In an algorithm with single-walk parallelism one or more consecutive steps in the neighborhood graph are made in parallel. To this end, applications of the exchange function, which we call *exchanges*, are proposed that all act on the same solution. Since effectuation of a proposed exchange can prohibit effectuation of other exchanges, the proposed exchanges are combined in such a way that a feasible solution is constructed, which can imply that not all proposed exchanges can be effectuated. Let τ be an exchange function as defined in Definition 2.2. Then a single-walk parallel local search algorithm consists of the following steps.

(1) Partition the domain $\{(s, e_1, \ldots, e_r) \mid e_1, \ldots, e_r \in s\}$ of the exchange function τ for a given current solution $s \in \mathcal{S}$.

(2) Propose exchanges for each subdomain simultaneously.

(3) Effectuate —a subset of— the profitable exchanges found in step (2), which results in a new solution s'.

(4) Replace s by s', and continue steps (1) to (3) until some stopping criterion is met.

Single-step parallel local search, which is based on parallel neighborhood exploration, is a special case of the above concept. In single-step parallelism each processor examines a part of the neighborhood of the current solution s, and only a single exchange is effectuated in step (3). So speed-up is achieved only in step (2). Single-step parallelism implies that, in terms of a walk in the neighborhood graph, only a single step is made as solution s' is obtained from s by applying a single exchange. In an algorithm with multiple-step parallelism several exchanges are effectuated in step (3), which implies that a solution s' is constructed that can only be reached from s by performing multiple steps in the neighborhood graph—that is, multiple exchanges have to be performed to obtain s' from s; see Figure 3.1.

Below, we present a machine-independent template for single-walk parallel local search algorithms. First we introduce the concept of distributed neighborhood structures, which have to deal with the following issues. A distributed neighborhood has to specify how solutions and domains of the exchange function are decomposed. Furthermore, it has to specify how proposed exchanges are combined. These aspects are captured in the following definitions.

Definition 3.2. Let $\Omega = \{1, \ldots, P\}$ be a set of processors, $\mathcal{S}$ the set of solutions, and $\mathcal{U} = \bigcup_{s \in \mathcal{S}} \mathcal{P}(s)$ the set of *partial solutions*. A *solution distribution* δ gives a partial solution δ_p for each processor $p \in \Omega$. A *local neighborhood structure* $\mathcal{L} : \mathcal{U} \mapsto \mathcal{P}(\mathcal{U})$ gives sets of partial solutions for partial solutions. □

Similar to conventional neighborhoods, local neighborhoods are not given explicitly but are constructed using an exchange function. The following definition states how this is done.

Definition 3.3. Let $\tau : \mathcal{U} \times \mathcal{E}^r \mapsto \mathcal{U}$ be an exchange function that can be applied to partial solutions. A *domain distribution* $\lambda_p : \mathcal{U} \mapsto \mathcal{P}(\mathcal{E}^r)$ gives for each processor $p \in \Omega$ and partial solution $u \in U$ a set of arguments, chosen from u, for the exchange function τ. A *local neighborhood* $\mathcal{L}(u)$ is equal to $\{\tau(u, e_1, \ldots, e_r) \mid (e_1, \ldots, e_r) \in \lambda_p(u)\}$ for $u \in \mathcal{U}$. □

The domain distribution describes which elements of partial solutions are the arguments used by the exchange function to construct local neighbors. If solutions are decomposed, exchanges involving elements of different partial solutions cannot occur, and therefore several decompositions need to be examined. This issue is also described in the following definition of a distributed neighborhood.

Definition 3.4. Let $\mathcal{S}$ be a set of solutions, $\mathcal{U}$ the set of partial solutions, Ω a set of P processors, and τ an exchange function. Then, a *distributed neighborhood structure* $\mathcal{D}$ is a triple (Δ, λ, ϕ) defined as follows. A *distribution structure* Δ gives for each solution s and set of processors Ω a set of solution distributions, with $\Delta(s, \Omega) \subseteq \{\delta \in \Omega \to \mathcal{U} \mid \bigcup_{p \in \Omega} \delta_p = s\}$. λ gives for each $p \in \Omega$ a domain distribution $\lambda_p : \mathcal{U} \mapsto \mathcal{P}(\mathcal{E}^r)$ with $r \in \mathbb{N}^+$, where λ gives the arguments of the exchange function τ used to construct the local neighborhood $\mathcal{L}(\delta_p)$ of a partial solution δ_p. A *combination function* $\phi : \mathcal{U}^P \mapsto \mathcal{S}$ combines P partial solutions to a feasible solution. □

A distributed neighborhood structure consists of a distribution structure that specifies how solutions are distributed over processors, a domain distribution that defines local neighborhoods, and a combination function that specifies how proposed exchanges have to be combined. Note that only a subset of the proposed exchanges might be effectuated by the combination function, since effectuating all proposed exchanges might lead to infeasible solutions. Next, we formalize the notion of local optimality for a distributed neighborhood structure.

Definition 3.5. Given are an instance $(\mathcal{S}, f)$, with $f : \mathcal{U} \mapsto \mathbb{Z}$, a distributed neighborhood structure $\mathcal{D} = (\Delta, \lambda, \phi)$, and processor set Ω. Then, a solution $s \in \mathcal{S}$ is a *local minimum* of $\mathcal{D}$ if for each solution distribution $\delta \in \Delta(s, \Omega)$, processor $p \in \Omega$, and partial neighbor $\delta'_p \in \mathcal{L}(\delta_p)$ holds $f(\delta'_p) \geq f(\delta_p)$. □

Using the concept of distributed neighborhood structures we can formulate the template for single-walk parallel local search given in Figure 3.3. In this algorithm first a distribution of the current solution is chosen. Then, all processors

```
proc Single_Walk (s : S)
   var δ : Ω → U; D : P(Ω → U);
   begin
      D := ∅;
      while Δ(s, Ω) \ D ≠ ∅ do
         δ ∈ Δ(s, Ω) \ D;
         par p ∈ Ω do Iterative_Improvement (δ_p) rap;
         if gain found then D := ∅; s := φ(δ_1, ..., δ_P)
         else D := D ∪ {δ} fi
      od {s is a local minimum of D}
   end
```

Figure 3.3: A single-walk parallel local search algorithm.

simultaneously propose exchanges. Subsequently, a subset of the proposed exchanges is effectuated to obtain a new feasible solution. This process is repeated until no solution distribution of the current solution can be improved upon, at which point a local minimum has been found.

To guarantee termination of the algorithm, ϕ should combine a set of neighbors of s in such a way that, if any neighbor has lower cost than s, their combination also has lower cost than s. We define this termination condition as follows.

Definition 3.6. Given are a problem instance $(\mathcal{S}, f)$, a distributed neighborhood structure $\mathcal{D} = (\Delta, \lambda, \phi)$, and a set of processors Ω. Then, the combination function ϕ is called *progressive*, if for all $\delta \in \Delta(s, \Omega)$, $p \in \Omega$, and $s \in \mathcal{S}$ holds that $\exists_{\delta'_p \in \mathcal{L}(\delta_p)} f(\delta'_p) < f(\delta_p) \Rightarrow f(\phi(\delta'_1, \ldots, \delta'_P)) < f(s)$. □

An important issue is raised by the question how we can compare a distributed neighborhood structure $\mathcal{D}$ with a conventional neighborhood structure $\mathcal{N}$. This enables a comparison of the solutions found by a parallel local search algorithm with those found by a sequential local search algorithm. The following definition is useful for such a comparison.

Definition 3.7. Given are a problem instance $(\mathcal{S}, f)$, a neighborhood structure $\mathcal{N}$, and a distributed neighborhood structure $\mathcal{D}$ for this problem. Then, $\mathcal{D}$ is called *isomorphic* with $\mathcal{N}$ if, for any set of processors Ω, holds that each local minimum of $\mathcal{N}$ is a local minimum of $\mathcal{D}$, and vice versa. □

If a distributed neighborhood structure $\mathcal{D}$ is isomorphic with a neighborhood structure $\mathcal{N}$, then the expected average cost of final solutions found by sequential and parallel algorithms, are equal, provided that sequential and parallel algorithms have equal probabilities for finding given local minima.

Implementation issues. First, we discuss synchronous and asynchronous single-walk parallel algorithms. In an algorithm with synchronous single-walk parallelism some step of the algorithm can be performed only in a state dependent on the state of all processors. This can occur for example when global communication has to take place before a next step of the algorithm can be performed. Synchronization is typically required in a decentralized algorithm in which each processor has a copy of the current solution. In order to update all local copies of the current solution after some exchanges have been proposed, global communication is required. If there is a central control mechanism that manages access to the current solution, no inconsistencies in the current solution can occur. So processors can then proceed independently from each other, although it may occur that some of the proposed exchanges cannot be effectuated because the current solution has already been altered by another processor. The choice between synchronous or asynchronous parallelism is often determined by characteristics of the parallel machine at hand, such as the availability of common memory or the configurability of a machine in a master/slave model. As we want to abstract from machine-dependent issues, we do not further elaborate on this subject.

The speed-up obtained with asynchronous and synchronous single-walk parallelism depends on problem characteristics and on the architecture of the target machine. Two important issues to obtain a good speed-up are load balancing and communication overhead, which we briefly address below.

In many single-walk parallel local search algorithms global communication is necessary because effectuation of an exchange has a global impact on the characteristics of solution. If a machine allows direct communication with a central processor, or if it has common memory from which all processors can read efficiently, then the overhead of asynchronous global communication is limited. In many other cases, however, global communication between all processors causes a substantial communication overhead.

Load balancing is needed to reduce the amount of time processors are idle. In synchronous parallelism load balancing must be done by assigning equal amounts of work to all processors in between synchronization points. In asynchronous single-walk parallelism load balancing is needed only for processors that interact.

Finally, the efficiency of single-walk parallelism is also determined by the ratio between the time needed to evaluate the cost of a neighbor and to combinate and effectuate proposed exchanges. For this reason the data structures used to represent a solution are important, since these data structures determine the efficiency of operations that have to be performed to effectuate proposed exchanges.

3.2.1 Single-step parallelism

In the previous section, we have introduced a template to capture single-walk parallel local search algorithms. In order to apply single-step parallelism to a given problem, the distributed neighborhood of Definition 3.4 is instantiated as follows. The distribution structure is chosen as $\Delta(s, \Omega) = \{\delta : \Omega \to \mathcal{S} \mid \delta_p = s$ for all $p \in \Omega\}$ and the combination function is given by $\phi(s_1, \ldots, s_P) \in \{s_i \mid 1 \leq i \leq P\}$. Moreover, it is not difficult to define a domain distribution that partitions a conventional neighborhood, which guarantees that the distributed neighborhood is isomorphic with it. The distributed neighborhood of Definition 3.4 is now straightforward, since a complete solution is assigned to each processor, and the combination function effectuates one of the proposed exchanges. This means that the combination function only has to select a neighbor from the proposed neighbors.

An advantage of single-step parallelism is its general applicability. Moreover, it results in a distributed neighborhood structure that is isomorphic with a conventional neighborhood structure that uses the same exchange function. So, under mild conditions, it is guaranteed that the parallel algorithm finds the same quality solutions as a sequential algorithm.

A disadvantage of single-step parallelism is the amount of speed-up that can be obtained. A local search algorithm with single-step parallelism explores neighbors of a solution simultaneously. Subsequently, one of the neighbors replaces the current solution, which results in a single step in the neighborhood graph. The amount of speed-up that can be obtained by single-step parallelism strongly depends on the number of exchanges that has to be examined before a proposed exchange is accepted. If best improvement is used as pivoting rule, then single-step parallelism gives good speed-up, because finding the neighbor with lowest cost requires scanning of the entire neighborhood of a solution. Best improvement is often used in tabu search, so single-step parallelism can be effectively applied in tabu search algorithms. However, best improvement is not always the most effective pivoting rule since for many problems the same quality solutions can be found with first improvement in smaller running times.

For algorithms with first improvement the amount of speed-up that can be obtained by single-step parallelism depends on the stage of the search process that has been reached by the algorithm. If the current solution has few lower-cost neighbors, a situation arises that is almost similar to best improvement, since then it is more likely that a large proportion of neighbors has to be examined before a solution with lower cost is found. If the proportion of neighbors that would be accepted by a local search algorithm at a certain stage of the search process is given by α, then the maximum speed-up with single-step parallelism is $\frac{1}{\alpha}$, regardless of the number of processors; see also [Roussel-Ragot & Dreyfus, 1990; Azencott, 1992] for a theoretical analysis of the speed-up that can be obtained by single-step parallelism in simulated annealing. Generally, α is large in the beginning of the search process, and consequently the resulting speed-up is small at first. As the cost of the current solution decreases, α increases and thus the resulting speed-up. So for a local search algorithm with first improvement, single-step parallelism can typically be applied efficiently only in a later stage of a local search walk when low-cost solutions are reached.

Examples. One the first examples of single-step parallelism in the literature is the algorithm of Kravitz & Rutenbar [1987] who apply asynchronous single-step parallelism to simulated annealing for the cell placement problem. They report a speed-up of 2.5 on a shared memory machine with four processors. Diekmann, Lüling & Simon [1993] present asynchronous and synchronous single-step parallel simulated annealing algorithms for the traveling salesman problem and the link assignment problem. Their algorithms are implemented on a network of 120 transputers. They obtain a speed-up of 35 for the traveling salesman problem and a speed-up of 70 for the link assignment problem. Kindervater, Lenstra & Savelsbergh [1993] present a local search algorithm with synchronous single-step parallelism for the time-constrained traveling salesman problem. Verification of local optimality can be done in $\mathcal{O}(\log N)$ time with $\mathcal{O}(N^2/\log N)$ processors on a PRAM machine, but no empirical results on a parallel machine are given. Synchronous single-step parallelism is used by Ravikumar [1992] in an iterative best improvement algorithm for the traveling salesman problem.

As already mentioned, many tabu search algorithms also use best improvement as pivoting rule. Synchronous single-step parallelism can then be applied effectively, as is shown by Taillard [1990, 1991] for the flow shop sequencing problem and the quadratic assignment problem. Chakrapani & Skorin-Kapov [1993a, 1993b] use synchronous single-step parallelism in combination with best improvement for the traveling salesman problem and the quadratic assignment problem, respectively. Their algorithm is implemented on a massively parallel Connection Machine with 16,384 processors, but running times are compared to an implementation on a workstation only. They also report that 55 percent of the running time is spent for communication. Li & Pardalos [1992] use synchronous single-step parallelism in a parallel variable depth algorithm for the quadratic assignment problem.

3.2.2 Multiple-step parallelism

Essential to achieve a good speed-up with multiple-step parallelism is that a large proportion of the proposed exchanges can be effectuated in the combination step.

The applicability of multiple-step parallelism therefore strongly depends on the problem at hand, as the problem must allow effectuation of several proposed exchanges for a given solution.

Two approaches to design distributed neighborhood structures can be distinguished based on whether or not solutions are decomposed into partial solutions. If the cost difference between two neighboring solutions can be computed using only information on the removed and inserted parts of a solution, then it might be profitable to decompose a solution and assign each partial solution to a processor. This enables an efficient implementation of the combination function because a new solution can then be constructed without global communication. A disadvantage, however, is that exchanges that involve elements of different partial solutions cannot be proposed. In order to find the same quality solutions as a sequential local search algorithm, different distributions have to be examined, which might lead to a complex distribution structure.

Another approach in the design of distributed neighborhood structures is to provide each processor with a copy of the current solution and to define an appropriate domain distribution. This often leads to a distributed neighborhood that is isomorphic with a conventional neighborhood that uses the same exchange function. To achieve speed-up, however, an effective combination function is essential because effectuation of an exchange can prohibit effectuation of other exchanges to guarantee feasibility of solutions. Furthermore, global communication is often required in the combination function, which can lead to a large communication overhead depending on the target machine's architecture.

Examples. The traveling salesman problem of Example 2.1 is a typical example of a problem for which decomposition of a solution is often used. Here a solution is a Hamiltonian cycle, and the cost difference between neighbors can be computed using only the lengths of removed and inserted edges. Moreover, it is possible to combine proposed exchanges without extensive communication. Most of the examples we discuss below are implemented on a distributed-memory MIMD machine and use synchronous parallelism.

One of the first applications of multiple-step parallelism to the TSP is presented by Felten, Karlin & Otto [1985], who divide a tour in consecutive paths. Each path is assigned to a different processor. A new distribution is obtained by assigning adjacent edges that are assigned to different processors to the same processor. Fiechter [1994] presents a parallel tabu search algorithm for the TSP that uses a similar solution distribution. Partial solutions that are assigned to processors consist of one path in the tour. New distributions of a tour are obtained by assigning different paths to processors.

Allwright & Carpenter [1989] present a parallel simulated annealing algorithm for the TSP based on a distribution of a tour over a linear array of processors. A partial solution in this distribution consists of two non-adjacent paths in the tour. Edges are randomly reassigned to processors to obtain a new distribution. In Chapter 5 we present distributed neighborhood structures for the TSP based on a similar solution distribution. We prove that our distributed neighborhoods are isomorphic with the well-known 2-exchange and 3-exchange neighborhoods. Here, a new distribution is obtained by assigning a single edge to a new processor, which leads to synchronization between processors. In Chapter 5 we also present a parallel Lin-Kernighan algorithm based on domain decomposition. Each processor works on a copy of the entire tour, and the proposed exchanges are subsequently combined to a new tour, where as many as possible exchanges are effectuated in the combination function.

Bachem, Steckemetz & Wottawa [1994] propose a solution distribution based on a geometric partition of the cities in a TSP instance. The partial solutions, which consist of Hamiltonian paths, are combined by a tour construction heuristic. A new distribution is obtained by choosing a new geometric partition.

Applications of multiple-step parallelism can also be found in the placement of cells in circuit layouting. Shahookar & Mazunder [1991] present an overview of this field. In most of the examples simulated annealing is used. For cell placement problems a local change might have a global impact on a solution, since the cost of a solution is determined by the relative position of pairs of cells. If no restrictions are imposed, effectuation of proposed exchanges can lead to erroneous cost computation. Erroneous cost computation changes the convergence behavior of simulated annealing, but various empirical results in the papers below show that simulated annealing for cell placement is rather robust to errors in the cost evaluation. Romeo & Sangiovanni-Vincentelli [1991] present some theoretical results on erroneous cost evaluation in simulated annealing.

Darema, Kirkpatrick & Norton [1987] use a solution distribution in which they assign the cells to processors, so each partial solution consists of a subset of the set of cells, and all proposed exchanges are accepted by the combination function. A new distribution is obtained by randomly reassigning different cells to each processor. They compare two solution distributions: one that guarantees correct cost computations and one that allows errors in the cost computation. A drawback of the first approach is that only a limited number of processors can be used, which prohibits scaling of the algorithm. The second approach does not suffer from this drawback. Their empirical results show that both approaches find the same quality final solutions. Jones & Banerjee [1987] present a similar algorithm in which they also decompose a solution. A new decomposition is obtained by exchanging cells between neighboring processors, and the combination function accepts all proposed exchanges, which may lead to erroneous cost computations. The algorithm is implemented on a distributed MIMD machine. Casotto, Romeo & Sangiovanni-Vincentelli [1987] use a local search heuristic to find a new decomposition of the solution that minimizes the probability of erroneous cost computation. These errors are caused by accepting all proposed exchanges without checking the position of cells in the newly constructed solution. Their algorithm is implemented on a shared-memory parallel machine. A hybrid algorithm that uses both multiple-walk and multiple-step parallelism is presented by Rose, Snelgrove & Vranesic [1988]. They apply multiple-walk parallelism at high temperatures of the simulated annealing algorithm and multiple-step parallelism during low temperature annealing, because at low temperatures erroneous cost computation is less likely to occur due to the low acceptance rate.

Barbosa & Gafni [1989] present a simulated annealing algorithm with multiple-step parallelism that can be applied to problems for which only a single element is replaced in the exchange function of the neighborhood structure. Moreover, such an exchange must have limited interaction with other elements of a solution. They propose a solution decomposition approach that guarantees that no erroneous cost computation can occur when all proposed exchanges are effectuated, at the cost of a limited number of employable processors. As a consequence, the maximum speed-up that can be obtained with their approach depends on characteristics of the problem instance at hand.

Another example of multiple-step parallelism is the algorithm of Savage & Wloka [1991] for the graph partitioning problem, for which they propose a solution distribution based on decomposing a solution. The combination function either accepts all proposed exchanges if the resulting global solution has lower cost, or it rejects all proposed exchanges if the resulting global solution has higher cost. Such a combination function has low computational overhead but it also may effectuate individually deteriorating exchanges.

Boissin & Lutton [1993] present a framework for parallel simulated annealing based on multiple-step parallelism. Exchanges are proposed according to the conventional acceptance criterion of simulated annealing, and the combination function probabilistically accepts or rejects all proposed exchanges—that is, it accepts or rejects the global solution resulting from these exchanges. The algorithm is tested on the quadratic assignment problem and on a 0-1 quadratic function minimization problem, displaying moderate speed-ups when implemented on a connection machine and compared to implementations on a workstation.

Taillard [1993] and Garcia, Potvin & Rousseau [1994] propose parallel tabu search algorithms for vehicle routing. Taillard proposes a solution distribution in which entire, geographically close, routes are preferably assigned to processors that are directly connected in the processor network on which the algorithm is implemented. A new distribution is obtained by assigning elements to neighboring processors. Garcia, Potvin & Rousseau propose a domain distribution for the vehicle routing problem with time windows. Their combination function consecutively effectuates the proposed exchanges in order of descending gain.

3.3 Complexity issues of parallel local search

In this section we discuss some issues related to the parallel complexity of finding local optima. For an overview of the complexity theory of parallel computations, the reader is referred to [Greenlaw, Hoover & Ruzzo, 1995].

If PLS-complete problems require superpolynomial running times, as is conjectured by Yannakakis [1996], then parallel algorithms that find local minima using a polynomially bounded number of processors, which is the case in any realistic machine model, will also run in superpolynomial time. Therefore, we restrict ourselves to the parallel complexity of the *verification problem*, the problem of deciding whether a solution is a local optimum. Consider a problem instance $(\mathcal{S}, f)$ and a neighborhood structure $\mathcal{N}$, then we have the following result.

Theorem 3.2. *Let $|\mathcal{N}(s)| = b(n) > 1$ for all $s \in \mathcal{S}$, where n denotes the size of a solution, and let $c(n)$ be the time complexity of deciding whether $f(s') \geq f(s)$ for a given solution $s' \in \mathcal{N}(s)$. Then, a parallel algorithm for the verification problem exists that uses $b(n)/\log b(n)$ processors on a PRAM and that has a time complexity of $\mathcal{O}(c(n)\log b(n))$ and an efficiency of $\mathcal{O}(1)$.*

Proof. The sequential time complexity for this problem is $\mathcal{O}(c(n)b(n))$. Partition the neighborhood in $b(n)/\log b(n)$ sets that contain $\log b(n)$ solutions. Use an algorithm with synchronous single-step parallelism. Its time complexity is $\mathcal{O}(c(n)\log b(n)) + \mathcal{O}(\log(b(n)/\log b(n)))$. □

In Chapter 1 we have discussed the complexity class EP introduced by Kruskal, Rudolph & Snir [1990] that contains problems for which an efficient parallel algorithm on a PRAM machine exists. Recall that a problem is in EP if there exists a parallel algorithm with speed-up that scales with the instance size and for which the computational effort, the time complexity of the parallel algorithm multiplied by the number of processors, is equal to the time complexity of the sequential algorithm. Theorem 3.2 now leads to the following corollary.

Corollary 3.1. *If the complexity of the problem to decide whether $f(s') \geq f(s)$ for $s \in \mathcal{S}$ and $s' \in \mathcal{N}(s)$ is polynomially bounded in the instance size and if neighborhood sizes are also bounded polynomially, then the verification problem is in the class EP. Moreover, if the former problem is in NC, then the verification problem for $\mathcal{N}$ is in NC.*

Proof. Let $c(n) = \mathcal{O}(n^u)$ be the complexity of the problem to decide whether $f(s') \geq f(s)$ for $s \in \mathcal{S}$ and $s' \in \mathcal{N}(s)$, and let $b(n) = \mathcal{O}(n^v)$ be the neighborhood size for instances of size n. Let T_p be the time required by an algorithm for the verification problem with p processors. Then, $T_1 = \mathcal{O}(b(n)c(n)) = \mathcal{O}(n^{u+v})$ and according to Theorem 3.2 for $p = \frac{b(n)}{\log b(n)}$ holds $T_p = \mathcal{O}(c(n) \log b(n)) = \mathcal{O}(n^u \log n^v)$. It remains to show that $T_p = \mathcal{O}(T_1^\epsilon)$ for $\epsilon < 1$. It holds that $T_p = \mathcal{O}(n^u \log n) = \mathcal{O}(n^u n^\delta)$ for $\delta > 0$. For any δ and ϵ such that $0 < \delta < v$ and $\frac{u+\delta}{u+v} \leq \epsilon < 1$ holds $T_p = \mathcal{O}(n^{u+\delta}) = \mathcal{O}(n^{(u+v)\epsilon}) = \mathcal{O}(T_1^\epsilon)$. Furthermore, $T_p \cdot p = \mathcal{O}(T_1)$. Thus, the verification problem is in EP. Moreover, if $c(n) = (\log n)^{\mathcal{O}(1)}$ on a PRAM, then the verification problem is NC. □

Corollary 3.1 shows that verification of local optimality can be done in polylogarithmic time with a polynomially bounded number of processors, if the cost evaluation of a neighbor can be done in polylogarithmic time. This typically holds when the cost evaluation of a neighbor is $\mathcal{O}(1)$, for example in the 2-exchange neighborhood for the TSP. Corollary 3.1 shows that it is possible to design an efficient parallel algorithm for the verification problem with speed-up that is proportional to the size of the problem. The running time of this parallel algorithm can still be polynomial, but in practical situations one is satisfied with parallel algorithms with a good speed-up that scales with the instance size.

Another important topic is the complexity of finding a local optimum of a distributed neighborhood structure. The notion of PLS-completeness and the definition of the class PLS can also be applied to the problem of finding a local optimum of a distributed neighborhood structure. Using this extension, we obtain the following result.

Theorem 3.3. *Let a distributed neighborhood structure $\mathcal{D}$ be isomorphic with a neighborhood $\mathcal{N}$. If the problem of finding a local optimum of $\mathcal{N}$ is PLS-complete, then the problem of finding a local optimum of $\mathcal{D}$ is PLS-complete.* □

Finally, we discuss upper bounds on the speed-up that can be obtained with the various approaches discussed in this chapter. We assume that we have an infinite number of processors at our disposal. The number of processors that can be employed in multiple-walk parallelism is not bounded by a polynomial in the size of the instance, so here we do not have a polynomial upper bound on the maximum speed-up. In an algorithm with single-walk parallelism the maximum speed-up is at most the size of neighborhoods because at least one neighbor has to be evaluated by each processor that contributes to the speed-up. Most neighborhoods have low-order polynomial sizes. In practice the number of processors of a parallel machine is typically much smaller than the neighborhood size, so the number of available processors is then a trivial upper bound on the maximum speed-up.

4

Multiple Independent Walks

The most straightforward approach to introduce parallelism in local search is to perform multiple independent runs of a local search algorithm simultaneously since this requires no complex parallelization scheme or even dedicated parallel hardware. An important question is therefore to what extend such an approach can be successful. In this chapter we analyze the speed-up that can be obtained by performing several independent walks in the neighborhood graph simultaneously. In this analysis we concentrate on iterated local search for the traveling salesman problem but the conclusions drawn from it can be applied to other problems and local search variants as well. The outline of this chapter is as follows. Section 4.1 presents a theoretical and empirical average-case analysis of a 2-opt algorithm for the TSP. Section 4.2 gives a semi-empirical analysis of the average-case performance of an iterated 2-opt and Lin–Kernighan algorithm. The main results of this chapter are discussed in Section 4.3 in which we show how these results can be used to analyze the speed-up achieved by multiple independent walks.

The performance of local search algorithms can be quantified by the relative excess of the obtained final solutions and the required running time. Empirical results show that local search can find good-quality solutions within low-order polynomial running times. It is, however, conjectured that worst-case running times cannot be bounded polynomially; see also Section 2.1. Furthermore, it is not possible to give theoretical upper bounds on the relative excess of local minima. So there is a considerable difference between the worst-case and empirical average-case behavior of local search algorithms. Theoretical average-case analysis is therefore useful to provide a better understanding of local search algorithms. However, only a few results on average-case analysis of local search are presented in the literature.

The TSP, defined in Example 2.1, belongs to the class of NP-hard problems [Garey & Johnson, 1979]. We consider only *symmetric* TSP instances, in which distances satisfy $d_{ij} = d_{ji}$, for each $i, j \in V$. The average cost of the solutions found by local search strongly depends on the choice of the neighborhood structure. Therefore, various neighborhood structures have been introduced for the TSP, most of which are based on edge exchanges. In the 2-opt neighborhood $\mathcal{N}_2$ of Croes [1958], formally specified in Example 2.1, a tour t' is a neighbor of tour t, if t' can be obtained from t by removing two edges and inserting two edges such that t' is obtained.

For an overview of the worst-case complexity and empirical behavior of local search for the TSP, we refer to [Johnson & McGeoch, 1996]. Others have addressed the theoretical average-case behavior of local search. Stadler & Schnabl [1992] investigate the structure of the 2-opt neighborhood using simplifications for dependencies between neighbors that lead to a flawed model. Kern [1989] showed with a probabilistic analysis that the 2-opt algorithm for Euclidean instances of the TSP has an average-case running time that is polynomially bounded. Other probabilistic models for local search have been studied by Tovey [1985]. In this study artificial problems are considered with special neighborhood graphs consisting of regular structures, e.g., the hypercube. The cost function for these problems is chosen to induce an orientation on this graph. Different cost distributions are considered and for some cases low-order polynomial average-case running times are proved. More recently, Chandra, Karloff & Tovey [1994] obtained similar results for 2-opt and 3-opt algorithms for the TSP, and derived upper bounds on the average cost of the local minima obtained by these algorithms. Most effort is focused on providing upper bounds on the average-case behavior of local search. We are interested in the actual distribution of the cost of local minima and the required number of steps to find them, instead of upper bounds on the average-case behavior only.

4.1 A probabilistic analysis for the 2-opt neighborhood

In this section we discuss the distribution of local minima found by iterative best improvement algorithms with the 2-opt neighborhood and the distribution of the number of steps required to find local minima. The distribution of local minima and the distribution of final solutions found by iterative improvement are not equivalent because not all local minima have the same probability of being found by an iterative improvement algorithm. It may be the case that local minima with low cost have a larger attraction region than local minima with high cost and are therefore more often found by local search.

An instance of the TSP is completely specified by its distance matrix. As the first step in our approach we assume that the distances d_{ij} are independently drawn from a given distribution. Such instances are called *random distance matrix* instances. Such instances are commonly considered in probabilistic analysis; see for instance [Kirkpatrick & Toulouse, 1985; Weinberger, 1991]. However, it should be noted that the assumption of independence is in fact a restriction that excludes Euclidean instances, because then the distances are dependent. It is possible to associate a class of instances with a distribution by letting the edge lengths $\underline{d}_{ij}$ be independent identically distributed random variables with mean μ_l and variance σ_l^2. Consequently, we can also view the cost $\underline{f}(t)$ of a tour t as a random variable, i.e., the sum of n independent, identically distributed edge lengths, where n is the number of cities in an instance. According to the central limit theorem, $\underline{f}(t)$ has approximately a normal distribution with mean $\mu = n\mu_l$ and variance $\sigma^2 = n\sigma_l^2$. The corresponding density of tour lengths is called ω_{tour}.

4.1.1 The distribution of final solutions

Consider a tour $t \in \mathcal{S}_k$, where $\mathcal{S}_k$ is the set of solutions that can be reached in k best improvement steps, starting from an arbitrary initial solution. The analysis of iterative best improvement is based on the computation of the *step probability*

$$g_k(c, c') = \mathbb{P}\{\forall_{t' \in \mathcal{N}_2(t)}\ \underline{f}(t') > c' \mid \underline{f}(t) = c \wedge t \in \mathcal{S}_k\}, \tag{4.1}$$

that is, the conditional probability that all neighbors of tour t have costs larger than c' given that a tour t found after k steps has cost c. The computation of (4.1) is discussed in the next section. Various notions can be expressed in terms of the step probability g defined in (4.1). First, note that $g_0(c, c)$ is the probability that an arbitrary tour with cost c is a local minimum. Hence, the density of local minima is given by

$$A^{-1} g_0(c, c)\omega_{\text{tour}}(c)\ , \tag{4.2}$$

where A is the probability that an arbitrary tour is a local minimum, given by

$$\int_{-\infty}^{\infty} g_0(c, c)\omega_{\text{tour}}(c)dc. \tag{4.3}$$

Consider a tour $t \in \mathcal{S}_k$ that is not a local minimum. Let $\mathcal{S}'$ denote the set of local minima. Recall that in best improvement a step is made to a neighbor with lowest cost. To investigate such a step, we compute for $c' < c$ the probability

$$\mathbb{P}\{\min_{t' \in \mathcal{N}_2(t)} \underline{f}(t') \leq c' \mid \underline{f}(t) = c \wedge t \in \mathcal{S}_k \setminus \mathcal{S}'\} =$$

$$\frac{\mathbb{P}\{\min_{t' \in \mathcal{N}_2(t)} \underline{f}(t') \leq c' \mid \underline{f}(t) = c \wedge t \in \mathcal{S}_k\}}{\mathbb{P}\{t \in \mathcal{S}_k \setminus \mathcal{S}' \mid \underline{f}(t) = c\}} = \frac{1 - g_k(c, c')}{1 - g_k(c, c)},$$

where we have used the fact that t cannot be a local minimum if $\underline{f}(t) = c$ and $\min_{t' \in \mathcal{N}_2(t)} \underline{f}(t') \leq c' < c$. The density corresponding with the above probability is

$$P_k(c, c') = \frac{\partial}{\partial c'} \mathbb{P}\{\min_{t' \in \mathcal{N}_2(t)} \underline{f}(t') \leq c' \mid \underline{f}(t) = c \wedge t \in \mathcal{S}_k \setminus \mathcal{S}'\} = \frac{-\frac{\partial}{\partial c'} g_k(c, c')}{1 - g_k(c, c)}.$$

This expression is the density as function of c' of the cost after $k + 1$ best improvement steps, given that the tour found after k steps has cost c and is not a local minimum.

The density of the local minima found after k best improvement steps can be described by the following recurrence relations. Let ρ_k be the density of the local minima found after at most k steps and let η_k be the density of the remaining tours. If we start the sequence of best improvement steps with a randomly generated tour, then $\eta_0 = \omega_{\text{tour}}$ and $\rho_0(c) = 0$, for all c. After the k^{th} step the density of the residual tours equals η_k. Consider such a tour with cost c. There is a probability $g_k(c, c)$ that it turns out to be a local minimum. This is found out in the $k + 1^{\text{th}}$ step, hence

$$\rho_{k+1}(c) = \rho_k(c) + g_k(c, c)\eta_k(c). \tag{4.4}$$

On the other hand, there is a probability $1 - g_k(c, c)$ that the tour with cost c is not a local minimum. Then, with probability density $P_k(c, c')$, it is transformed into a tour with cost c' in the $k + 1^{\text{th}}$ step. Hence,

$$\eta_{k+1}(c') = \int_{c'}^{\infty} \eta_k(c)(1 - g_k(c, c)) P_k(c, c') dc. \tag{4.5}$$

This set of recurrence relations allows us to compute the densities of the detected local minima and the residual tours after an arbitrary number of steps. Then, $\lim_{k \to \infty} \eta_k = 0$ and $\lim_{k \to \infty} \rho_k = \rho_{\text{fin}}$, the density of the final solutions. Moreover, note that $\rho_k - \rho_{k-1}$ is the density of the local minima found in the k^{th} step. So if $\underline{steps}$ is the random variable describing the number of steps until a local minimum is found, then

$$\mathbb{P}\{\underline{steps} = k\} = \int_{-\infty}^{\infty} g_{k-1}(c, c)\eta_{k-1}(c) dc. \tag{4.6}$$

4.1.2 Evaluation of the step probability

The problem has been reduced to evaluating the step probability (4.1). The joint distribution of $\underline{f}(t_0), \underline{f}(t_1), \ldots, \underline{f}(t_b)$ is needed to compute this probability for an arbitrary tour t_0 with neighboring tours $t_1, \ldots, t_b$, where $b = |\mathcal{N}_2(t_0)| = n \cdot (n-3)/2$. These tours have a large number of edges in common, so the corresponding tour lengths are not independent. Two arbitrary tours s and t that have m edges in common, have a covariance given by

$$\mathbb{E}\{(\underline{f}(t) - \mu) \cdot (\underline{f}(s) - \mu)\} = m\sigma_l{}^2,$$

where $\mu = n\mu_l$ is the mean tour length, μ_l the mean edge length, and $\sigma_l{}^2$ the variance of edge lengths. The tour t_0 and each of its neighbors t_i, with $1 \le i \le b$, have $n-2$ edges in common. However, two neighbors of t_0, say t_i and t_j, can have $n-3$ or $n-4$ edges in common. The case of $n-3$ common edges occurs if, while going from t_0 to t_i respectively t_j, a common edge is removed or inserted. Consequently, the $(b+1) \times (b+1)$ covariance matrix of the random variables $\underline{f}(t_0), \underline{f}(t_1), \ldots, \underline{f}(t_b)$ is rather complicated, which makes it difficult to compute or numerically approximate the right hand side of (4.1) for the 2-opt neighborhood. Therefore, in the remaining part of this section we concentrate on the neighborhood $\mathcal{N}_2'$ that is a restricted version of the 2-opt neighborhood in Example 2.1 defined by

$$\mathcal{N}_2'(t) = \{\tau_2(t, e_i, e_{h(t)}) \mid 1 \le i < h(t) - 1 \vee h(t) + 1 < i \le n\}.$$

So whereas in the conventional 2-opt neighborhood two arbitrary edges can be removed, here one of the edges to be removed from a tour $t = \{e_1, \ldots, e_n\}$ is fixed and is given by $h(t)$ for a function $h : \mathcal{S} \to \{1, \ldots, n\}$. In this neighborhood structure a tour t has only $b = |\mathcal{N}_2'(t)| = n-3$ neighbors, and all these neighbors have $n-3$ edges in common.

If we assume that the probability of (4.1) does not depend on the number of local search steps, so $g_k(c, c') = g(c, c')$ for all k, then the transition properties and several related notions for the best improvement algorithm can be computed for the $\mathcal{N}_2'$ neighborhood structure. Note that $g_0(c, c)$ can be computed without this assumption. If edge lengths are normally distributed with mean μ_l and variance $\sigma_l{}^2$, then the costs $\underline{f}(t_0), \underline{f}(t_1), \ldots, \underline{f}(t_b)$ have a joint normal distribution [Papoulis, 1965] with mean μ and covariance matrix R given by

$$\begin{array}{lll} R_{ii} & = n\sigma_l{}^2 & \text{for } 0 \le i \le b \\ R_{i0} = R_{0i} & = (n-2)\sigma_l{}^2 & \text{for } 1 \le i \le b \\ R_{ij} = R_{ji} & = (n-3)\sigma_l{}^2 & \text{for } 1 \le i < j \le b, \end{array}$$

where R_{ii} gives the variance, R_{i0} the covariance between t_0 and its neighbors, and R_{ij} the covariance between two neighbors of t_0. As (4.1) is a conditional probability, we have to compute the conditional density $\omega_{\text{cond}}(c_1, \dots, c_b \mid c)$, i.e., the density of $\underline{f}(t_1), \dots, \underline{f}(t_b)$ given that $\underline{f}(t_0) = c$. It is a known result (see for instance [Papoulis, 1965]) that ω_{cond} is again a joint normal density, with all means equal to $\mu' = \mu + R_{i0}/R_{00}(c - \mu) = (1 - 2/n)c + (2/n)\mu$. The variances are given by $r_{ii} = R_{ii} - {R_{0i}}^2/R_{ii} = (4 - 4/n){\sigma_l}^2$ and the covariances by $r_{ij} = R_{ij} - R_{i0}R_{j0}/R_{00} = (1 - 4/n){\sigma_l}^2$. Now (4.1) can be rewritten as

$$g(c, c') = \int_{c'}^{\infty} dc_1 \cdots \int_{c'}^{\infty} dc_b \; \omega_{\text{cond}}(c_1, \dots, c_b \mid c).$$

The essential observation is that now all covariances are equal and positive. In this case the b-fold integral can be simplified to a single integral [Stuart & Ord, 1987]. This results into

$$g(c, c') = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^{\infty} e^{-\frac{s^2}{2}} \left(\frac{1}{2} \text{erfc} \left(\frac{c' - (1 - \frac{2}{n})c - 2\mu_l}{\sqrt{6}\sigma_l} - s\sqrt{\frac{1}{6} - \frac{2}{3n}} \right) \right)^b ds,$$

which can be computed numerically.

4.1.3 Empirical results

We validate our model for the density of final solutions ρ_{fin} and for the density (4.6) of the required number of steps by comparing them with empirically obtained densities. We consider instances with 200 and 400 cities in which edge lengths are distributed according to a standard normal distribution, which implies that tour lengths can be negative. Other distributions of edge lengths in which costs of neighbors can be approximated by a joint normal distribution, may also be considered. The results are computed from averages over five instances, although there is little difference between the means of local minima in individual instances since these quantities are self-averaging, which means that these quantities are equal for sufficiently large instances in which edge lengths are samples from the same distribution. For each instance we have sampled 20,000 local minima found by an iterative best improvement algorithm. The fixed edge $h(t)$ in the neighborhood $\mathcal{N}_2'(t)$ is randomly chosen for a tour t.

Figure 4.1 and 4.2 give the results for instances with 200 and 400 cities, respectively. In these figures, the density ρ_{fin} of the costs of final solutions found by a best improvement algorithm is depicted. Furthermore, the density of (4.6) of the number of steps required by a best improvement algorithm to find a local minimum is given. We observe that the theoretical predictions fit well with the empirical results, so our model adequately describes the behavior of local search for these instances.

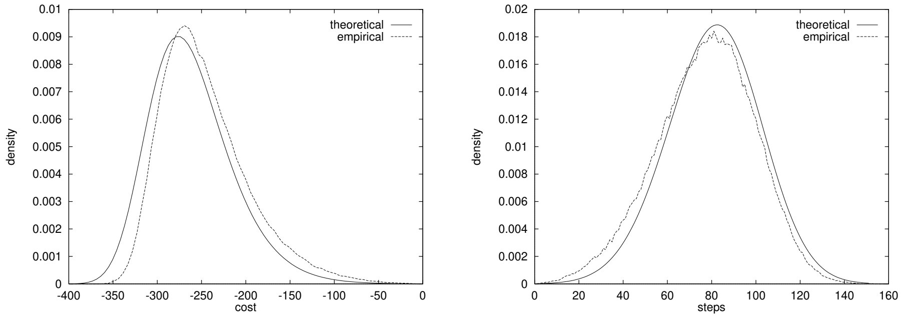

Figure 4.1: Results for instances with 200 cities.

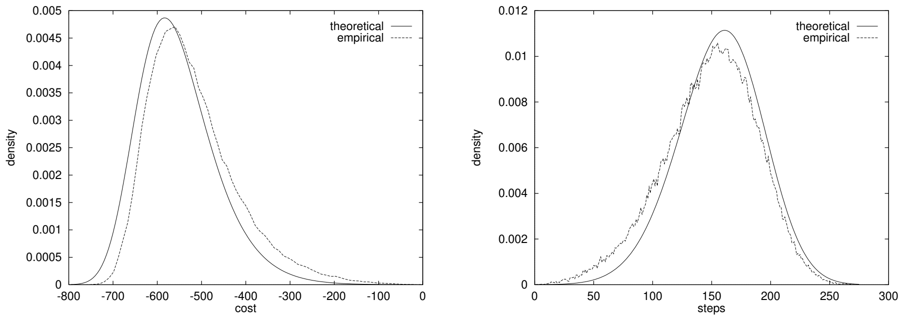

Figure 4.2: Results for instances with 400 cities.

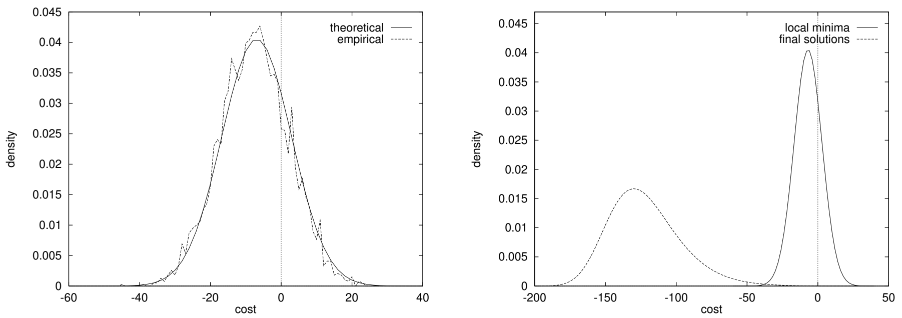

Figure 4.3: Density of local minima.

The probability that an arbitrary tour is a local minimum is given by (4.3). For $n = 100$ this probability amounts to $1.23 \cdot 10^{-4}$. We determined empirically the density of local minima for five instances with 100 cities by randomly generating 37 million tours of which 4410 turned out to be local minima. The lefthand side of Figure 4.3 gives the corresponding empirical density as well as the theoretical density of local minima given by (4.2). Again, we obtain a good fit between theoretical and empirical results.

The righthand side of Figure 4.3 presents the density of local minima as given by (4.2) and the density of final solutions as obtained by iterative best improvement for instances with 100 cities. Typically, the average cost of final solutions is much lower than the average cost of local minima. This implies that low-cost local minima have a much larger attraction region and therefore a much higher probability of being found by local search than high-cost local minima, which is of course an advantageous property for the performance of local search.

Empirical distributions for 2-opt and Lin–Kernighan. Next, we present distributions of final solutions obtained by a first improvement 2-opt algorithm and a Lin–Kernighan algorithm, discussed in more detail in Chapter 5, to show that the distribution shapes observed for the more artificial random distance matrix instances and restricted 2-opt neighborhood we used in our theoretical analysis, are also instructive for real-world Euclidean instances.

Figure 4.4 shows empirical results for a number of real world instances from the TSP library of Reinelt [1991], where the number in an instance name denotes

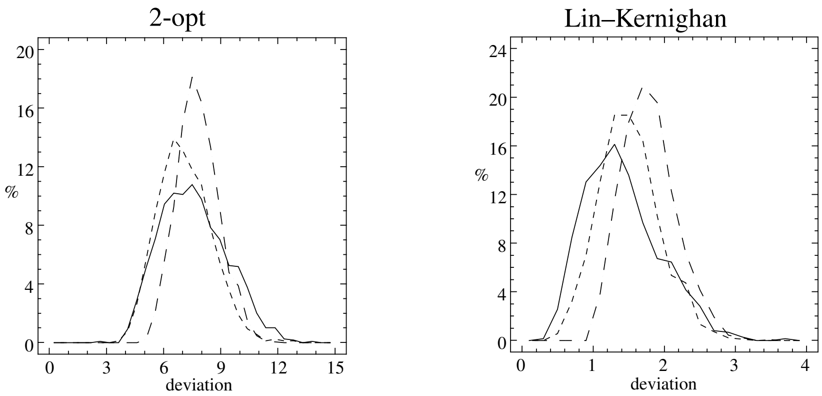

Figure 4.4: The distribution of final solutions. For the 2-opt algorithm, the solid line gives the results for kro200, the short-dashed line for lin318, and the long-dashed line for pcb442. For the Lin–Kernighan algorithm these lines give results for pcb442, u574, and pr1002, respectively.

its size. In these figures, the frequency is plotted against the relative excess over the optimal solution value. Our empirical investigation of the distribution of final solutions obtained by 2-opt and Lin–Kernighan algorithms reveals two interesting properties. First, we have statistically validated that final solutions are distributed according to gamma distributions. Secondly, we have observed that the standard deviation of these distributions decreases when the instance size grows. These characteristics are an important aspect of a neighborhood for iterated local search or other more advanced local search algorithms, because they indicate the additional computational effort required by iterated local search to find lower-cost local minima.

4.2 A semi-empirical analysis of iterated local search

Iterated local search algorithms repeatedly execute an iterative improvement algorithm. Each time the iterative improvement algorithm terminates, the last or best local minimum found is modified and the iterative improvement algorithm is restarted with the modified local minimum [Johnson, 1990; Martin, Otto & Felten, 1991; Boese, Kahng & Muddu, 1994]. The best heuristic to handle the TSP is the iterated Lin–Kernighan algorithm of Johnson [1990]. In Johnson's iterated Lin–Kernighan algorithm the best local minimum found so far is modified by a 4-exchange, which replaces four edges by four new edges, and used as starting solution for the next run of the Lin–Kernighan heuristic.

In this section we approximate the average running times iterated local search algorithms need for finding a solution within a given relative excess over the optimal solution. Time is measured by the number of tours whose costs are evaluated. Our analysis is of a semi-empirical nature, which means that the average number of evaluations is expressed as a function of empirically obtained parameters. These parameters are the distribution of final solutions and the average number of steps needed to find them.

The average number of iterations and the number of evaluations performed per iteration are needed to approximate the average running time. We, furthermore, assume that an iterated local search algorithm samples local minima, which means that we neglect the intermediate solutions generated by the iterative improvement algorithm. Let $\underline{itnr}_{\epsilon}$ be the random variable that gives the number of iterations until a solution with a given relative excess ϵ has been found, then

$$\mathbb{P}\{\underline{itnr}_{\epsilon} \leq i\} = 1 - \prod_{k=1}^{i}(1 - \mathbb{P}\{\underline{f}(s_k) \leq (1+\epsilon) f_{\text{opt}}\}), \tag{4.7}$$

where s_k is the final tour obtained at iteration $k \in \mathbb{N}$.

To determine the average number of iterations needed, we assume that the modification mechanism is primarily a diversification mechanism. Thus, we consider subsequent local minima to be independent, and consequently the probability $\mathbb{P}\{f(s_k) \leq c\}$ is independent of k. This may seem a rather strong assumption since the modification mechanism only changes four edges in a local minimum, but the final solution obtained from this modified local minimum typically differs in much more edges from the original local minimum. Given the density ρ_{fin} of the costs of final solutions, we then have

$$\mathbb{P}\{f(s_k) \leq (1+\epsilon) f_{\text{opt}}\} = \int_{-\infty}^{(1+\epsilon) f_{\text{opt}}} \rho_{\text{fin}}(c) dc.$$

Let p_ϵ denote the above probability, then the probability of (4.7) is given by

$$\mathbb{P}\{\underline{itnr}_\epsilon \leq i\} = 1 - (1 - p_\epsilon)^i. \tag{4.8}$$

Expression (4.8) implies that the number of iterations is distributed according to a geometrical distribution. So the average number of iterations to find a solution within a given relative excess ϵ is equal to $1/p_\epsilon$.

In order to approximate the average number of evaluations needed per iteration, we need the following observations. After the first iteration the iterative improvement algorithm starts with a tour that differs only four edges from a local minimum. Consequently, the average number of evaluations required in subsequent iterations is substantially lower than that required in the first iteration, which is also observed from empirical results. Hence, we differentiate between the following two values.

- k_1, the average number of evaluations needed to reach a local minimum from a start solution.
- k_2, the average number of evaluations needed to reach a local minimum from a solution obtained by modifying a local minimum.

The values of k_1 and k_2 are obtained empirically. They depend strongly on the heuristic used to construct start solutions and on the instance at hand.

Let $\underline{evals}_\epsilon$ denote the random variable that gives the number of evaluations until a solution with a given relative excess ϵ is found. Then, $\underline{evals}_\epsilon = k_1 + (\underline{itnr}_\epsilon - 1)k_2$, and

$$\mathbb{P}\{\underline{evals}_\epsilon \leq j\} = \mathbb{P}\{\underline{itnr}_\epsilon \leq \lceil \frac{j - k_1}{k_2} \rceil\} =$$

$$\begin{cases} 0 & \text{if } j < k_1, \\ 1 - (1 - p_\epsilon)^{\lceil \frac{j-k_1}{k_2} \rceil} & \text{if } j \geq k_1. \end{cases} \tag{4.9}$$

The expected number of evaluations needed to find a solution within a given relative excess ϵ is equal to $k_1 + (\frac{1}{p_\epsilon} - 1)k_2$.