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Keywords parallelism, workload modelling

Abstract

This work presents the novel method for high-level performance estimation of systems consisting of multiple computational units. The goal is to support system designers in the early phases of the system design flow. The focus mainly lies on embedded systems and in this first part of the work, we began from their versions which perform parallel processing with execution units similar to each other. Systems consisting of different types of processors, and the method expansions to support them are also discussed.

The main idea was an attempt to reallocate a single processor's load to multiple simulated processors. The method uses measurements from actual, existing systems and relies on means of simulations with systems under design. Instead of competing with prototyping, the method is supposed to give an estimation of which kind of system architecture would fulfil the desired performance requirements.

In the method, we process the mentioned measurement data automatically, which results in a so-called workload model. The workload model is then executed with a simulated system. This simulation run approximates the proposed system's estimated performance. Due to automation at the modelling phase and a high level of abstraction, the method allows the fast approximation of several different configurations.

The first of the problem areas was to define which type of workload model is suitable and how it can be created. When the workload is measured from a uni-processor system, its parts which can be parallel executed must be discovered, in order to use the model with a multiprocessor system. The second problem area is the modelling of the performance-related parts of the system under design. The larger problem is to study the validity and rationality of the whole method.

We validated the method with two different test cases and both of them gave reasonable results. The first validation consists of a simple threaded application, which uses an inter-thread synchronization mechanism. As the internal functionality of the application is known, the characteristics of the method can be roughly seen. The second validation method is a real-world algorithm, which we will execute in both a simulated and existing two-processor system. The margin for error of the method can be calculated from the latter of the validation cases, by comparing the total execution times of the systems. The margin for error for this case was from 10 to 15 %. It was better than expected for a method with a rather high level of abstraction.

As research results, the work presents the parts needed for the method: an instrumentation for gathering the measurement data, the creation of a workload model out of it, a simulation of a multi-processor system with the workload model, and visualization of the simulation results. In addition, an analysis of these parts and the whole method is presented.

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Avainsanat parallelism, workload modelling

Tiivistelmä

Tämä työ esittelee menetelmän korkean tason suorituskykyarviointiin järjestelmälle, joka koostuu useammasta suoritinyksiköstä. Tarkoituksena on tukea suunnittelijoita järjestelmän määrittelyvaiheessa. Menetelmä on tarkoitettu ensisijaisesti sulautetuille järjestelmille, ja tässä laajemman työn ensimmäisessä vaiheessa mielenkiinto oli sellaisissa versioissa, joissa rinnakkainen suoritus tapahtuu samanlaisia suorittimia käyttäen. Työssä käsitellään myös keskenään erityyppisistä suorittimista koostuvia järjestelmiä ja menetelmän laajennusta tukemaan myös niiden analyysiä.

Tärkeimpänä osa-alueena menetelmässä on yrittää jakaa yhden prosessorin kuorma useammalle simuloidulle prosessorille. Menetelmä käyttää mitattua dataa olemassa olevista järjestelmistä ja tukeutuu simulointiin suunnitteluvaiheessa olevien järjestelmien kanssa. Menetelmää ei ole tarkoitettu kilpailemaan prototyyppien tekemisen kanssa vaan antamaan arvio siitä, minkälainen arkkitehtuuri täyttäisi halutut suorituskykyvaatimukset.

Olemassa olevista järjestelmistä mitattua kuormitusdataa prosessoidaan automaattisesti, ja tulosta kutsutaan kuormamalliksi. Tätä mallia käytetään syötteenä simulointivaiheelle, joka jäljittelee suunniteltavana olevan järjestelmän käytöstä. Simuloinnin tulokset antavat informaatiota järjestelmän ennustetusta suorituskyvystä. Esimerkiksi tietyn kuorman kokonaissuoritusaika on yksinkertainen suorituskyvyn mitta. Mallinnuksen automaatiosta sekä menetelmän korkeasta abstraktiotasosta johtuen eri arkkitehtuurivaihtoehtojen arviointi on nopeaa.

Työn ensimmäinen ongelma-alue oli sopivan kuormamallin löytäminen. Jotta kuormamalli soveltuisi moniprosessorijärjestelmille, sen luonnollisesti tulee pystyä erottelemaan rinnakkaiseen suoritukseen soveltuvat osat, kuten myös riippuvuudet eri osien välillä. Seuraava ongelmaalue on mallintaa suorituskykyyn liittyvät osa-alueet suunnittelussa olevasta järjestelmästä. Isompana kokonaisuutena olivat koko menetelmän järkevyyden ja oikeellisuuden tarkastelut.

Menetelmä validoitiin kahta erilaista lähestymistapaa käyttäen. Ensimmäinen validointitapa toteutettiin yksinkertaisella säikeistetyllä ohjelmalla, joka käytti säikeidenvälistä synkronointia. Koska ohjelman sisäinen rakenne on nyt tunnettu, menetelmän toiminnallisuus voidaan nähdä karkeasti. Toinen validointitapa on todellinen algoritmi, joka suoritettiin sekä simuloidulla että olemassa olevalla kaksiprosessorijärjestelmällä. Jälkimmäisestä validointitavasta pystyttiin laskemaan menetelmän virhemarginaali vertaamalla molempien ajojen kokonaissuoritusaikoja. Virhemarginaaliksi tälle tapaukselle saatiin noin 10–15 %. Tämä virhemarginaali oli odotettua parempi menetelmän korkea abstraktiotaso huomioon ottaen.

Tutkimustuloksina esitellään menetelmään tarvittavat osa-alueet: instrumentointi mittausdatan saamiseksi, kuormamallin muodostaminen tästä datasta, moniprosessorijärjestelmien simulointi edellä mainitun mallin avulla sekä tulosten visualisointi. Lisäksi esitellään menetelmän ja sen osa-alueiden analysointi.

Preface

This thesis, required for a diploma, was made as a part of the TWINS project at the VTT Research Centre of Finland. The TWINS project is a jointly funded project in the Information Technology for European Advancement (ITEA) programme. There are 24 research and industrial partners from five different European countries. The project aims to enhance the hardware-software co-design flow for software intensive system development. The performed work, discussed in this thesis, provides one specific solution for the early system design phases. The first words of the thesis saw the daylight in autumn 2007 and the thesis was finished in spring 2008.

At first, the given task – a very loosely defined one – seemed almost impossible to complete, but through the course of time and by dividing the overall problem into logical sub areas, the solution was shaped into its current form. The gained knowledge and experience of project working will be very valuable in the future.

I would like to thank my direct superior, Mika Hongisto, firstly for hiring me for this interesting job, and secondly for acting as a local supervisor for the thesis.

I thank Professor Tapio Seppänen and Professor Olli Silvén from the University of Oulu for supervising my thesis.

I thank my colleagues, from the TWINS project and also from my team. Without any help from these professionals, many problems would have remained unsolved. Special thanks go to the three persons who gave the most important practical support: Markku Pollari from the project and my team members Tuukka Miettinen and Anton Yrjönen.

Last but not least, compliments go to my parents, who have made all this possible throughout all these years.

Oulu, Finland, 23rd May, 2008

Marko Jaakola

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Abbreviations

1.1 Designing performance

Through the history of computers, there has always been one clearly distinctive factor between products from two different generations – the performance. The performance can have different interpretations depending on the system's use. For example, a low response time on user actions, a large throughput of data or the handling of several simultaneous requests can be thought as representing good performance. Since an increase in the performance is nearly always considered as an obligatory requirement when computer systems evolve, its analysis has received more and more attention. Acquiring a good performance for the final product naturally begins from the early phases of the design process.

One certain type of computer-based system, called embedded system, usually has both hardware and software affiliated in its design process. Likewise, both hardware and software have an effect on the total performance of an embedded system. Traditionally, the hardware part of an embedded system has been developed first, and then the software engineers must try to fit in proper software, in order to complete the system. The current trend in the embedded systems' development is to design both hardware and software components simultaneously. This way of action is called co-design, and the general codesign flow is presented in figure 1 (figure 2 in [1]).

Figure 1. Activities in the co-design of an embedded system.

System requirements development, also known as Requirements engineering (RE), is the phase which aims at obtaining requirements for the system under design. We define requirements as early stage specifications of what should be implemented, and they can be descriptions of a system's behaviour, properties, attributes, constraints and compatibility issues [2, p. 4]. The following are examples of system requirements [2, p. 4]:

- Features provided for the user (e.g. a spell checker in a word processor)
- General system properties (e.g. the securing of the personal information)
- Constraints for the functionality of the system (e.g. a polling interval for a sensor)
- Constraints in the development of the system (e.g. defining the programming language to be used).

Correspondingly, we define Requirements engineering (RE) to include all of the activities related to discovering, documenting and maintaining a set of requirements for a computerbased system [2, p. 5]. In this work, we will concentrate on performance requirements. The performance requirements belong to the requirements class, which is called either non-functional requirements (NFR) or sometimes quality attributes. In information processing systems, the most often used performance requirements relate to throughput and response times [3]. The typical problems in performance requirements – some already discussed above – are the following [3]:

- Interactions and conflicts with other requirements: for example, accuracy and speed may form a trade-off in some systems
- Effect of choosing the development technique an aspect which is costly to fix after the development has started
- Global impact to the whole system: in the worst case, optimizing performance can necessitate modifications in every system module
- Characteristics of the performance vary between different organizations and different systems.

System architecture development is sometimes also considered as a separate activity in the specification phase; it contains the consideration of available hardware components and the possible constraints set by software components. [1]

Allocation and flow-down means a decision of which functionalities should be performed with hardware, and which with software. Roughly speaking, adding more hardwarebased functionalities means more speed and unfortunately more costs. However, the adding of software-based solutions also has obligatory costs, such as more software development, larger read-only memories (ROMs), and in the worst case the changing of the selected processor for a more powerful and more expensive version, which also causes higher energy consumption. Relying on hardware also has problems with a possible need for redesigning, which is much more complicated on regarding hardware than with software. Usually, the partitioning decision should be done at the latest possible time. This is due to the fact that the understanding of the problem evolves continuously, as the process advances. A drawback is that debugging of software is more complicated before the actual hardware is done; early-stage testing must be done with evaluation boards or so called stub codes, which mimic the behaviour of the hardware. [4, p. 49]

Nowadays the partition decision is becoming more difficult, because the advancement of integrated circuits makes the implementation of very complex algorithms possible with reasonable costs [4, p. 50]. The distinction between software and hardware is also blurring, as hardware designs and software designs look quite alike, they are just used with different compilers [4, pp. 55–56].

A product's performance is substantially affected already during these presented steps forming the *co-specification* phase. The following phases are *co-development* and *coverification*. In co-development, the actual development work of hardware and software, as well as the proper interface between them, are performed. In co-verification, these parts are integrated and tested. The coordination work of these phases is called *co-management*. [1]

Our focus is to propose a new method for the described, early phases of the design process. To be more specific, we are interested in the performance, as it can be seen in the next section.

1.2 Motivation

The work aims to provide a new, rather highly abstracted method for system designers in a multi-processor domain. The method's purpose is to aid architecture decisions before prototyping. We will provide a *performance simulation* for estimating the feasibility of different design alternatives. The method will attempt to help the selection of the most favourable ones to be validated with prototypes. After all, we are doing the simulation in a non-detailed and coarse way. We are favouring simulation, because it is an applicable way to support the design flow due its modest amount of work. Figure 2 (based on figure 1.1 on [5]) shows possible ways to examine a given system.

Naturally, the most straightforward way to examine the system is to *experiment with the actual system* itself. If the system is however under design, it probably does not exist yet. This means that the actual system must be replaced with a temporary implementation, in other words, one has to *experiment with a model of the system.* Some systems can be modelled with a *physical model*, for example a scale model, but this approach is hardly feasible in the domain of computer systems. A more practical physical model for computer systems is a prototype, but its development can be a heavy and time-consuming process. The alternative to a physical model is a *mathematical model*, which means presenting the system's essential or interesting parts with the help of mathematics. The mathematical model can have an *analytical solution*, but unfortunately the solution can be impossible to find because of the complexity of the modelled system. This leaves *simulation* as the only alternative for the mathematical model's solution. As we are focusing on system architecture development, allocation and flow-down phases where the actual system or prototype does not exist yet, the selected method in our approach was the simulation.

A simulation intends the execution of a system model. A system model represents some existing or planned real-world system or process and its features, which can include logical, mathematical or structural properties. The detail level of the model should be selected according to the application's need. A simulation model differs from a generic system model by also associating time and changes that occur over time into the model. The models can be classified by their relation to time: the state of a discrete model can change only at discrete points in time, whereas a continuous model can change its state at an arbitrary moment [6]. Computer systems are almost always simulated with discrete simulation models, obviously due to the discrete nature of computers.

Figure 2. Different ways to study a system under interest.

A simulation has some major benefits when compared to prototyping. A simulation model can be executed under arbitrary projected operation conditions, as the control of experimental conditions is better in a simulated environment. Alternative designs or operation policies can usually be examined and compared with small changes and even in a single simulation run, whereas in prototyping, there may be the need for one prototype per one design proposal. Control over the simulation time is also a remarkable benefit: a simulation can be driven either in compressed or in expanded time. [5, pp. 76–77]

Our method also includes experimentation with actual systems: we are using data gathered from existing versions to support our performance estimations for the next generation product. The processing of this data is automated and therefore some of the required steps in a common simulation study are performed without a user interaction in our method.

1.3 Approach and research questions

In this work, the focus was to study whether our simulation process, shown in figure 3, would be a feasible way to give a coarse estimate of performance of a proposed multiprocessor system. The figure shows the main steps in our performance analysis. The method uses *measurement data* from an existing device, which is possibly previous development version with a single processor, and the next one is planned to use more processors. The acquired data is utilized in a *modelling* phase, where we build a so-called workload model, which is an abstract presentation of the essential characteristics of a rather large amount of data. We will also develop a simulation model of the system under design. The simulation model is modelled in sufficient detail to represent both hardware architectures, mostly the amount of processors, and how the workload is allocated be-

tween these processors by scheduling policies. The workload model is used as an input to the simulation model, when the *simulation* itself is executed. The simulation's output, for example the total execution time for a certain task set, is the desired information and the goal of the method. Further *analysis* of the results is done with a three-dimensional visualization tool named the PerVisGL [7].

Figure 3. The main steps of our performance analysis by using simulation.

The challenge is to find out if the idea of performance requirements fulfilment estimation could be done with this type of method. If the method is feasible, it will be a considerable addition to the tool range of a system designer.

The research questions we are going to address in this work, are as follows:

- How is the workload modelling performed, especially the exploration of dependencies and parallel parts?
- How to model systems' from the performance perspective?
- How these models and the whole method can be validated?
- How designers can make decisions based on the provided novel method?

After this introduction part, the rest of the thesis is organized into chapters giving prerequisites on embedded systems, describing our approach, presenting the obtained results and finally discussing these. We are roughly following the steps of the scientific method, which consists of four sequential phases [8, p. 167]:

- Analysis
- Hypothesis
- Synthesis
- Validation.

The analysis phase includes gaining an understanding on components of the problem domain, and the formulation of a problem description [8, p. 182]. In this work, we will prepare for the actual approach by reviewing the embedded system's performance related aspects from both hardware and software sides, in chapters 2. and 3.. The hypothesis phase aims to propose a solution to achieve the task objective [8, p. 200]. The synthesis phase is about implementing the task method [8, p. 220]. Both of these are presented in the approach chapter (the chapter 4.). The validation phase decides whether the objective has been achieved [8, p. 280]. Our validation results are presented in the results chapter (the chapter 5.).

This chapter introduces the hardware related areas, which are connected to this work and thereby gives a basis for our method. At first, we will briefly discuss the special characteristics of embedded systems' hardware, and then focus on multi-processor architectures as well as the processing units forming them. The chapter is wrapped up by discussing performance analysis by simulation methods.

2.1 Characteristics of an embedded hardware

We will begin familiarizing ourselves with the hardware of embedded systems by comparing it with a normal personal computer (PC). A PC's general-purpose processor has support for a wide range of different level computational applications; on an embedded system, processing power is ordinarily fitted for a specific task. This is possible due to the overflowing amount of microprocessor/microcontroller choices for embedded systems. For personal computers, there are less current processor architecture alternatives. Embedded systems have stricter power constraints than desktop PCs, firstly because they are often powered by batteries, and secondly because their size may set restrictions on cooling. Due to power saving, an embedded system may be completely interrupt driven; the processor is in sleep mode and wakes up only upon timer ticks. Environmental conditions, which an embedded system must tolerate, are sometimes also totally different than for a PC. Aircraft, space and military applications must stand extreme heat or cold, humidity, dust and vibration. The amount of resources, for example buses, main memory and mass storage space, is in practice always smaller in embedded systems, when compared to a PC. [4, pp. xviii–xxiv]

2.2 Multi-processor systems

When a computer system has more than one "independent" computational unit, it is called as a multi-processor system. The three main reasons for an ascending popularity of multiprocessor systems are the following: Performance improvements are achieved in a logical way by connecting multiple processors together, when the capacity of a single processor is not sufficient. This is probably more cost-efficient that designing a new, more powerful one. The second reason is the uncertainty about how long the advancement rate can be kept up by increasing the complexity, silicon and power of a single processor. The third reason comes from the viewpoint of the software; the trend seems to be towards parallel operations. Server and embedded areas have a particularly natural parallelism and utilizing multi-processor hardware is a real advantage there. [9, p. 528]

2.3 Classification of multi-processors

Multi-processor systems can be divided into different classes with at least three separate classification rules. The division can be based on the relation between the system's instruction and data streams, the system's memory architecture or the architecture's hierarchy.

2.3.1 Division by input and output

The following classic, simple model to categorize computer systems has four classes, which are still valid despite the model's high age, although some multi-processors are hybrids of more than one class [9, 10, p. 529]:

- 1. Single instruction stream, single data stream (SISD): This category includes normal uni-processors, which process single data element with a single instruction.
- 2. Single instruction stream, multiple data streams (SIMD): Instructions are dispatched by a control processor to multiple processors, which have their own data memories. One example of these kinds of systems is a vector processor, which operates multiple data elements with a single instruction. Some multimedia extensions in current general-use processors can also be considered as usage of the SIMD method.
- 3. Multiple instruction streams, single data stream (MISD): A comprehensive MISDstyle commercial processor is not yet built, but some stream processors can be loosely classified as this since single data stream is operated with successive functional units.
- 4. Multiple instruction streams, multiple data streams (MIMD): Everything from instruction fetching to data operating is handled by each of the multiple processors themselves. This category is the most used nowadays in general-purpose multiprocessor systems, in contrast to the early multi-processors which applied SIMD.

The benefits of the MIMD systems are cost-efficiency and flexibility. Cost-efficiency is achieved due to the building of the system with a set of normal microprocessors. Flexibility appears in cases, where the same system can run a single application at a top performance by utilizing all the processors for the same task, and secondly by running several tasks simultaneously, when required. [9, pp. 529–530]

2.3.2 Division by memory architecture

MIMD multi-processors can be further classified by their memory organization. The first class is centralized *shared-memory architectures*. Figure 4 (based on figure 6.1 on [9, p. 531]) shows one example system of this approach. This type has at most a few dozen processors, which have a shared single centralized memory, as well as a shared I/O system. On the other hand, the processors have their own caches, which means one or more levels of very fast and relatively small memory. A bus connects the processors and the main memory. The relationship to the memory is symmetric for all processors, and the system can therefore be called a *symmetric shared-memory multi-processor* (SMP) and the whole architecture a *uniform memory access* (UMA), since the memory access time is identical for each processor. Inter-processor communication is easy due to shared memory. [9, pp. 530–531]

The second class of MIMD multi-processors, *distributed memory architectures*, has a physically distributed memory, like in figure 5 (based on figure 6.2 on [9, p. 532]). This type is better for larger systems, as there will be problems with bus bandwidth in sharedmemory systems, when the number of processors is increased. This distributed-memory architecture can be thought to consist of nodes, where every node includes a processor, memory, sometimes an input/output (I/O) functionality and an interface to the interconnection network. Besides containing one processor, a single node can be a small symmetric multi-processor system itself. The advantages of the distributed-memory multiprocessor architecture are that smaller memory bandwidth is sufficient, if it is assumed that most of the memory accesses will relate to the local memory, and the memory access latency is also lower. The inter-processor communication is not as simple as with the shared-memory architectures. [9, pp. 531–532]

Figure 4. A centralized shared-memory multi-processor.

Figure 5. A distributed-memory multi-processor.

The communication in distributed-memory multi-processors can be managed by two different techniques. The first method is distributed shared-memory access (DSM), which means that the memory is not shared but the address space is: the same physical address in different processors refers to the same memory location, and thereby the memory access latency is different depending on which memory is referenced – the local memory ver-

sus the other node's memory. This architecture can also be called non-uniform memory access (NUMA). Still, a processor cannot access every memory location of the system; some parts require proper access rights. The second communication method has memories, which are totally isolated from the other nodes' processors and the architecture is called a message-passing multi-processor. Converted into real terms, the same physical address in different processors refers to different memory, respectively. When there is a need for inter-processor communication, the processor sends a request for some data operation to other processor, which can be thought of being a remote procedure call (RPC). The destination processor receives the message via polling or interrupts, performs the requested operation and sends the response. The requesting processor waits for the reply before it continues, so the passing of the message is synchronous. Asynchronous messaging is also possible: the writer of some data is aware that other processors require that data too, it sends the data directly without waiting for any requests and immediately continues after the messages have been sent. The nodes in the message-passing multi-processors can be thought to be separate computers, and the architecture is thereby sometimes called a multi-computer. A multi-computer can be built with completely separate computers connected to a local area network, if the amount of the communication is small. [9, p. 533]

2.3.3 Division by the architecture's hierarchy

Multi-processor systems can also be divided into *heterogeneous* and *homogeneous* systems. Homogeneous systems have computational units similar to each other, whereas heterogeneous systems consist of different types of processors: usually one or more central processing units and also one or more application specific hardware components. Embedded systems, related to digital signal processing (DSP), are usually heterogeneous. Figure 6 (based on figure 1 on [11]) shows the so called Garp architecture, which is a small heterogeneous multi-processor system where a standard processor is supported by a reconfigurable slave computational unit. In heterogeneous systems, the central processing unit may be for example, a microcontroller or a programmable digital signal processor, and the hardware processing element may be an application-specific integrated circuit (ASIC) or some reconfigurable logic such as a field programmable gate array (FPGA). One example of an embedded system, which could utilize this kind of configuration, is a device which performs video or audio decoding, like a present-day mobile phone. [12, p. 1]

One currently interesting heterogeneous processor architecture is the Cell Broadband Engine. It provides single-chip multi-processing with two different core types which are called power processor elements (PPEs) and synergistic processor elements (SPEs). PPEs are responsible for system-wide services such as virtual memory management, handling exceptions and thread scheduling. SPEs are responsible for most of the data processing. SPEs perform the computation in an SIMD-style. For example, the configuration of a Cell chip can consist of one PPE and eight SPEs, like in figure 7 (based on figure on the page 3 on [13]). The figure shows how each SPE consists of a synergistic processing unit (SPU) and a synergistic memory flow controller (MFC). Furthermore, each SPU consists of a synergistic execution unit (SXU) and a local storage (LS). Each SPE has its own memory flow controller which handles the communication into an element interconnect bus (EIB) which is used to transfer data between the system memory and local storages. A power processor unit (PPU) with a level two cache forms a power processor element.

Figure 6. A block diagram of a heterogeneous multi-processor system, called as Garp.

PPU consists of a PowerPC execution unit (PXU) and level one cache. In addition, the system has a memory interface controller (MIC) and a bus interface controller (BIC). [13]

Figure 7. One possible Cell multi-processor configuration.

When two or more homogeneous processor cores are located in the same chip, the architecture is called a *multi-core processor* and the whole technique is *chip multi-processing* (CMP). The cores can also share some resources with each other, for example regarding Intel's Core Duo architecture; each core has a private level one cache memory but they

share a level two cache. On the other hand, Intel's Pentium D architecture also has private level two caches for each of the cores; however it is still called a multi-core processor, as the cores reside on the same physical chip. [14, p. 249]

2.4 Building blocks for multi-processor systems

As stated in section 2.3.3, a multi-processor system can be assembled from a set of different kinds of operational units. One significant difference between these units is their usability for varying tasks – as we move from central processing units to digital signal processors, field-programmable gate arrays and finally to application-specific integrated circuits, software-dependency and flexibility decline at the same time. This is illustrated in figure 8. The above-mentioned units are presented shortly next.

Figure 8. A comparison between three different computational units. Efficiency on the x-axle means the energy-efficiency of the unit, when it is performing some task. Flexibility on the y-axle means the suitability for varying tasks of the unit, including those which are not known of during the unit's design-time.

2.4.1 Central processing unit

A central processing unit (CPU), sometimes also called just a processor, or nowadays a microprocessor when the whole CPU is located in a single integrated circuit, covers a control unit (CU), arithmetic and logic unit (ALU), input/output interface and internal memory in the form of registers. The actual random access memory (RAM) is located externally from the chip as its own component. Interpreting program instructions and processing data by arithmetic or logic operations are tasks which the CPU handles. The actual tasks are performed by using software, which utilizes basic operations supported by the processor – this provides versatility but also causes fairly modest energy efficiency. [15, pp. 83–84]

2.4.2 Microcontroller

A microcontroller differs from a microprocessor by including a CPU, memory, which can be random-access memory (RAM) or read-only memory (ROM), and other peripherals like I/O-functionality in the same chip. In other words, a microcontroller is designed to fulfil a minimum complement of external parts, and this compact package has some advantages over microprocessor-based systems. A higher level of integration causes lower cost, as one part replaces many parts. Fewer packages and fewer interconnections enhance reliability. Since system components are optimized for their environment, and signals can remain on the same chip, an improved performance is usually achieved. Fast signals do not radiate from a large board and thereby the so called radio frequency signature is lower. Due to these aspects, microcontrollers are common and even dominant in embedded systems. [4, pp. 24–25].

In a multi-processor system, usually either a microprocessor or a microcontroller is in control of the whole system, and other components can be considered to be slaves to them.

2.4.3 Digital signal processor

A digital signal processor (DSP) is a processor with enhancements and optimizations for digital signal processing and data transfer oriented tasks. It usually follows the Harvard architecture, where instruction codes and data have separate memories and at least one dedicated bus for both. Besides classification into fixed-point and floating-point DSPs, they can be categorized as general purpose and special purpose DSPs. Special purpose DSPs can be further classified as algorithm-specific processors performing for example digital filtering or fast Fourier transforms, and application-specific processors performing some telecommunications or audio related tasks. [16, pp. 615–618].

When compared with the von Neumann architecture, where the instructions and data are located in the same memory, the Harvard architecture allows instruction codes to be of a different size than the data, as well as their addresses. Parallel instruction fetching and data reading is also possible. [17].

2.4.4 Application-specific integrated circuit

Application-specific integrated circuit (ASIC) is a component, which is designed to constantly perform a specific task by hardware. It has a static structure, where logic gates and their interconnections are permanently decided on manufacturing. The advantages of ASICs are their performance, good energy efficiency and small size due to the fact, that it has only the required amount of logic gates. The disadvantages are a lack of flexibility, costs for small production batches, long turn-around time, which is the time between the order and the delivery, and a verification which should be performed before the actual production. [18]

2.4.5 Field-programmable gate array

Obviously, it would be good if the benefits of the ASICs and general purpose processors could be combined. In other words, there is a need for a device, which relies on hardware in performing different tasks but maintains versatility. These types of components are called programmable logic devices (PLD). One subtype of PLDs is the fieldprogrammable gate array (FPGA), which has configurable wiring and logic elements on one layer, and a personalization memory on the second layer. This personalization memory is used to configure wiring and logic elements, according to the components' programming. These types of FPGAs are re-programmable, but there are also one-time programmable versions, where the customization is based for instance on fuses. However, a FPGA has a substantial cost overhead when compared to an ASIC: both the personalization memory and configurable logic elements, with their interconnections, have a remarkable amount of transistors, and depending on which kind and the amount of logic elements really required, part of the transistors are unused in the current implementation. [19].

Since the microprocessor's relatively slow instruction fetch-decode-execute cycle is not needed, the FPGA performs faster and consumes less energy. Dynamically reconfigurable systems can switch their configuration during the run-time. This is analogous to a microprocessor, when it changes a software program under execution. [20]

2.5 Performance gauging – the hardware perspective

Traditionally, processors and whole systems are compared by their performance. The performance may be thought absolute or relative with other systems. Two classic units of measurement in the area of performance are MIPS and Dhrystone. The first benchmark, MIPS, which stands for a million instructions per second, originates from the VAX 11/780 minicomputer, which was the first system which was advertised to perform one MIPS. A single instruction, however, does not have much to do with actual work performance; the same work scales into different instruction counts on different architectures. Therefore, the MIPS is mostly a helpful unit only when comparing different versions of the same architecture. A more valuable benchmark is the Dhrystone, which is a simple C program that compiles to about 2000 lines of assembly code. One Dhrystone corresponds to the execution of 1757 program loops per second. Similarly to the MIPS, the calibration is inherited from the same VAX machine, which could execute the aforementioned 1757 loops per second. The program is independent of operating system services, but also has some weaknesses. For example, if this small program fits an on-chip cache of an embedded system totally, the performance results are naturally skewed. The benchmark also has difficulties with exploiting parallel performance in the proper way, and compiler optimizations towards favourable Dhrystone performance are also possible. [4, pp. 26– 27]

2.6 Performance simulations – the hardware perspective

Performance simulators are used for predicting the performance of the given system. In a computer domain these simulators are almost always software programs written with high-level languages. Analyzing the performance of a computer system would be naturally easiest to do with direct measurements – however, this kind of method is a postdesign step and rarely contributes directly to the design process of future systems. Predicting the performance can be completed with analytical models or the performance simulators, which are usually more detailed and therefore give more valuable information to designers. For example, current microprocessors alone are enormously complex systems and so are the simulators mimicking their performance. The most accurate simulators work on a so called register transfer level and they simulate the functionality of basic logic circuits. [21]

A few rather detailed performance simulation solutions are presented next. Although they actually simulate the complete combination of hardware and software, their low-level orientation justifies their positioning in the hardware part of this chapter.

Rsim can be used to simulate various non-uniform memory access shared memory multiprocessors (see section 2.3.2). The exploitation of instruction-level parallelism (see section 3.1) is one of the main interests in the Rsim. The simulator itself is a discrete event-driven simulator. The events are used to model processor pipelines, caches, memories and the network connecting the multi-processor architecture. The Rsim models the competition over system resources and inter-processor synchronization in multi-processor systems plus speculative execution also in uni-processor systems. Actual program executables, compiled and linked for the Sparc V9 systems, can be used as an input for the simulator; the gathered instructions are processed in a fetch-decode-retire style, which means that the instruction is dismissed in the third step, in contrast to a real system which would perform the actual execution as the last step. The output of the simulation has statistics on the total execution cycles, how many instructions per cycle were achieved, the usage rate of different functional units in the processor and several readings on the cache, memory and network operations. [22]

Simics is a simulation platform simulating several different processor architectures at the instruction-set level. The level of detail is sufficient to run unmodified operating systems at the top of the simulation platform. The focus is to simulate the full system consisting of both hardware and the actual software rather than the test code, or even a distributed system consisting of several nodes and each one of them are simulated. In the above-mentioned distributed case, simulated nodes could be situated in several hosts or just in a single one; the network connections in a distributed system can also be simulated when required. The provided device models are accurate enough to utilize the real firmware and device drivers. [23]

SimpleScalar is a flexible instruction-set level processor simulator, which can be used in varying detail. The simplest and fastest model only simulates the instruction set, whereas the most detailed microarchitectural model has features such as dynamic scheduling, speculative execution and a multilevel memory system. The SimpleScalar is an eventdriven simulation which uses actual program binaries as its input. [24]

Asim provides modularity to performance modelling. It is a framework where the total performance model consists of reusable software modules presenting physical components. When the user has created the desired performance model by selecting proper ready-made modules or writing their own modules, the simulation can be executed in three different ways. A static trace, acquired from another performance model, or a real system can be fed into the simulator. A dynamic trace works in almost the same way, but the trace is delivered forward simultaneously while measuring it from another model, thus saving storage capacity and time. The instructions can then finally be fed into the simulator from a program binary. [25]

This chapter discusses topics from the software side of embedded systems. Once again, the focus is placed on areas affecting the performance. Without underestimating the actual software applications' effect on the system's performance, in this work we will maintain quite a neutral sentiment towards them and focus more on the operating system level. Translated into real terms, we will handle programs as processes and threads and give special attention to their scheduling. We will finally discuss performance from the software perspective and how it can be estimated by the means of workload modelling and simulation.

3.1 Concepts of processes and threads

A process consists of a running program itself, and its state. A process is independent from other processes, although processes can exchange information. Thereby, utilizing several processors with several processes is a quite straightforward matter. The concept of a process is also widely used in uni-processor systems, when programs share a single uniprocessor computer. Processes are executed in short time slices, known as time-sharing, so it seems to the user that they are running simultaneously. Process state information is used when a process switch or a context switch occurs: the state of the preceding process is saved and the following is correspondingly restored. [9, p. 469].

The three most essential states for a process are running, ready and blocked. A running process is logically a process that is currently under execution. A ready process is all set for execution and waiting for processor assignment. A blocked process is waiting for some event, an I/O-event completion for example. [26, p. 55].

Processes, which share their code and most of their address space, are called threads, as shown in figure 9. Threads are also used with uni-processor systems, but a multi-processor system can utilize them perfectly. For a multi-processor system with n processors, there must be at least n processes, threads or a combination of these two, in order to system run at full throughput. Threads are usually created by the programmer, or in some cases the compiler can optimize a code by generating threads automatically. The parallelism here is called thread-level parallelism (TLP), and when compared to instruction-level parallelism (ILP), the ILP is handled mostly by hardware and it relates to a single instruction at a time, whereas thread-level parallelism relates to a substantial amount of instructions. [9, p. 272]

The utilization of threads within a single processor, called multi-threading, is divided into different approaches. Fine-grained multi-threading is a version which switches between threads on each instruction and skips threads, which are not ready to run at that precise moment. Practically, this kind of multi-threading requires support for thread switching at every clock cycle. The main advantage is that the system can execute other threads when some of them stall. On the other hand, an individual thread's execution is constantly interfered with by other threads. Coarse-grained multi-threading does switch between threads but only on costly long stalls. An execution of an individual thread is more continuous, but the throughput of the system is affected by shorter stalls. In simultaneous multi-threading (SMT), both the TLP and ILP are exploited simultaneously. Various functional units of a processor are used to execute instructions from different threads in a single clock cycle. The SMT demands the ability to fetch instructions from different threads and also sufficient buffer spaces. [9, pp. 608–610]

Some operating systems, for example Linux, do not draw a major difference between

Figure 9. The greatest difference between processes and threads is the sharing of address spaces.

processes and threads. Linux threads have all the features such as normal processes, however part of their resources are shared and therefore they are called as lightweight processes. [27, p. 80]

3.2 Scheduling levels and objectives

The assignment of physical processors to processes, in order to processes accomplish work, is called processor scheduling [26, p. 249]. Respectively, a scheduling algorithm is a set of rules that determine the task to be executed at a particular moment [28]. One can say that the distributed system resource here, the CPU time, is a type of renewable one. However, its handling is not a trivial task as we will see in the following sections.

The scheduling, performed by an operating system, can be divided into three different levels. High-level or long-term scheduling, also called job or admission scheduling, is used to determine jobs which shall be allowed to compete actively for the system resources. Once a job is admitted to the system, it becomes a process or a group of processes. Intermediate-level or medium-term scheduling determines which processes shall be allowed to compete for the CPU and brought to the main memory by swapping them from the mass storage; the method used is suspending and activating processes depending on fluctuations in the system's load. The benefit of this kind of scheduling is a smoother system operation and contribution to certain performance goals. Low-level or short-term scheduling determines the assignment of ready processes to the CPU. This is called dispatching and the module responsible for low-level scheduling is correspondingly called the dispatcher. The relationships between different process states (see section 3.1) and scheduling levels are presented in figure 10 (based on figure 9.2 on [29, p. 396]). [26, 29, pp. 249–250, pp. 394–398]

Scheduling has several general objectives. Scheduling should be fair – all processes should be treated likewise. The number of serviced processes per time unit should be maximized, whereas the response times should be minimized. The load of the system should not have an effect on the scheduling, and the overheads caused by the scheduling should be minimal. Resource utilization should be balanced, and the given priorities

Figure 10. Scheduling levels.

obeyed. [26, pp. 250–251]

Scheduling is said to be non-pre-emptive, if the currently executing process cannot be forced to stop and release the CPU. Process switching takes place when the executing process decides to release the CPU. In contrast, pre-emptive scheduling can interrupt a currently running process, for example, when another task with a higher priority is ready to run. In real-time and also in interactive systems, pre-emption is used to guarantee proper response times. The disadvantage in pre-emption is the overhead, which is firstly caused by additional context switching related tasks, and secondly by the need to keep non-running processes in the main storage in order to get them running quickly. Nonpre-emptive systems have more predictable response times, as the incoming jobs do not replace running processes. On the other hand, short jobs have to wait for longer jobs to finish. [26, pp. 252–253]

The mechanism, which an operating system uses to reclaim the CPU from a user process, is called an interrupt clock or an interval timer. The interrupt clock generates an interrupt after a certain period of time, which causes the operating system to run and decide which process is dispatched next. Of course, it is possible that the process releases the CPU voluntarily before the clock interrupt, or some other interrupt takes the CPU. The advantages of using the interrupt clock are enhanced response times and the prevention of the system becoming stuck because some process is looping infinitely. [26, p. 253]

The priorities for tasks can be assigned with few a different tactics. With the static way, the priorities are assigned to all the tasks once. Dynamic priorities can change during

scheduling, and a mixed scheduling algorithm contains varying priorities for some tasks and static for the rest. [31]. The assignment can be performed by the operating system or it may come from outside the system. Static priorities have a low overhead, but they do not respond to changes which would require adjustments. An overhead caused by dynamic priorities is usually compensated for by improved responsiveness. [26, p. 253]

3.3 Scheduling algorithms from a uni-processor viewpoint

Uni-processor scheduling strategies are discussed in this section, in order to ease the understanding of their multi-processor versions. Scheduling can be implemented with several different algorithms. *First-in-first-out* (FIFO), also known as *first-come-first-served* (FCFS) scheduling, is a simple non-pre-emptive discipline which dispatches processes based on their arrival time to a ready queue. It has quite predictable response times, but long jobs force short jobs to wait, as well as unimportant jobs make more important jobs wait. FIFO scheduling can be used as a part of other scheduling algorithms, for example on decisions among processes with the same priority. [26, 29, pp. 254–255, pp. 403–406]

Round robin (RR) scheduling works partially like the FIFO, but it only gives a slice of the CPU time, so called quantum to each process at a time. Processes, which do not finish their execution before the quantum has passed, are pre-empted and placed at the back of the ready list. Round robin has reasonable response times in time-sharing environments. The quantum size can be fixed or variable; a too small quantum size causes the context switching overhead to grow larger than useful work, whereas a too long quantum makes the RR to function as with FIFO scheduling. Figure 11 clarifies the policy of this algorithm. When a new task arrives (event 1 in the figure), it is placed at the back of the queue. When a task under execution has spent all of its time slice, it will also be placed at the back of the queue (event 2 in the figure). The next task to be executed is the task which has spent the longest time in the queue (event 3 in the figure). [26, 29, pp. 255–256, pp. 406–408]

Figure 11. An example scenario of a round robin scheduling.

Shortest job first (SJF), or *shortest process next* (SPN) is a non-pre-emptive scheduling, which always selects the process with the lowest estimated run-time-to-completion. The favouring of short jobs causes the number of waiting processes in the system to decrease fast and the average waiting time is also minimized. The problem is where to gain the estimates – in environments, where the same jobs come into execution in a periodic way, there may be good estimates, but the proper estimates are often impossible to obtain. SJF does not apply to time-sharing systems because it is non-pre-emptive. [26, 29, p. 257, pp. 408–410]

The pre-emptive counterpart of the SJF is called *shortest remaining time* (SRT) scheduling. The SRT may replace a currently running process with a process, which has a smaller run-time estimate. The replacement of a nearly completed process with only a slightly shorter job can be avoided by setting a threshold value, which guarantees that processes closing their completion can continue to execute uninterrupted. This algorithm requires the recording of elapsed execution times, which causes some overhead. [26, 29, pp. 257– 258, p. 410]

Highest response ratio next (HRN or HRRN) is a non-pre-emptive scheduling which calculates a priority-ratio for processes as a sum of the elapsed waiting time and the required service time, which is divided by the service time. The denominator causes shorter jobs to be preferred, but the increase in waiting time, called aging, increases the ratio and therefore guarantees that longer jobs will also be executed. The weakness of the HRRN is the need for service time estimates. [26, 29, p. 258, p. 412]

A scheduling algorithm called *multilevel feedback queues* consists of a queuing network, where a new process is placed at the back of the highest level queue, which has the highest priority. When a process goes through this queue in FIFO-style, and does not complete its execution in the given time quantum, it is dropped into the lower level and lower priority queue. Long processes go through the whole network, until they are finally completed in the lowest level, which is usually implemented in RR style. This method is based on imposing a penalty on processes which have run too long, although the time quantum can increase at the lower levels. The methods advantage is that service time estimates are not required. [26, 29, pp. 259–261, pp. 412–414].

3.4 Multi-processor scheduling

Scheduling can be thought to be the most essential factor, when a user reviews the performance of an interactive system. When we are moving from a uni-processor system to a multi-processor version, its importance is emphasized even more. Particularly, keeping all the processors as utilized as possible is one of the common scheduling problems in a multi-processor domain. Scheduling related aspects therefore play an important role in this work.

When scheduling multi-processor systems is considered, in addition to the actual dispatching of a process and the use of time-sharing on an individual processor, the assignment of processes to processors must be handled. A static assignment is the simpler alternative: a process is assigned to one processor for its whole execution span. The disadvantage is that one processor may be idle, while other processors have lot of assigned tasks waiting. A dynamic assignment bypasses this problem because the execution for the same process can take place in an arbitrary processor; however the repetition of the assignment during scheduling creates some overhead. Implementation for the assignment can be performed with master/slave or peer architectures. The master/slave architecture uses a particular processor to execute the key kernel functions of the operating system, such as scheduling. In this approach, the master processor can become a bottleneck for the whole system's performance and its failure will also halt other processors. Peer architecture can execute these kernel functions on any processor and therefore the processors

self-schedule their tasks. This approach must have some kind of synchronization and conflict resolution, as processors may compete for same processes. An intermediate solution is to perform scheduling on a subset of processors. [29, pp. 440–441]

Methods for the actual implementation of the multi-processor scheduling include for example *load sharing*, *gang scheduling*, *dedicated processor assignment* and *dynamic scheduling*. Load sharing uses a global queue of ready threads, which causes an even distribution of work to all processors, and also bypasses the problem of idle processors if there is work available. Scheduling decisions are made in a non-centralized way; the operating system is run on an available processor to select the next thread for it. Arranging the queue can be based on basic FCFS, or jobs with the smallest number of unscheduled threads can be alternatively executed first. The problems in this method are the synchronization of the global ready queue, extensive swapping of the cached data because pre-empted threads are often forced to continue their execution on different processor, and a distortion on the execution times of dependent threads, which would need to be executed concurrently. [29, pp. 444–445]

The performance bottleneck of the load sharing methods is caused by interacting threads or processes, when they are executed in separate time slots. Repetitive and unnecessary context switches occur because the synchronization points are reached at different times. This can be avoided through gang scheduling, also called group scheduling, which schedules closely related processes and threads to be executed in parallel. A performance gain comes from reduced context switching, which is consequence of reduced synchronization blocking – threads have a possibility to catch up to synchronization points in a much shorter time interval due to parallel execution. The scheduling related overhead is smaller, because decisions are made for a group of jobs at the one time. Resource allocation is also more efficient, because the cooperating threads are probably interested in the same resource, which can be accessed by several threads via their cooperation. [29, 30, pp. 446–447]

A dedicated processor assignment does not use pre-emption or multiprogramming at all; instead, a group of processors is dedicated to the application. In this approach, applications must be able to dynamically control the number of processes, depending on the amount of the assigned processors. Processes and threads run on the same processor until their completion – an absence of context switching causes improved performance. A lack of multiprogramming is justified in larger systems, because the single processor's blocking does not substantially affect the efficiency of the whole system. [29, 31, p. 447]

Equally, in dynamic scheduling, the number of threads is dynamically controllable by applications. A dynamic allocator primarily fulfils new requests with unallocated processors. If there are not any, allocated but currently unused processors are used next. The last possibility is to enforce equipartition by pre-empting processors from an application with the largest amount of processors. [29, 32, p. 449]

3.4.1 Algorithms especially for embedded systems

We will next look at multi-processor scheduling models, which are particularly utilized in embedded systems. They focus on task-level parallelism. All of these models are restricted to fulfil non-pre-emptive scheduling: tasks are executed as long as they wish once they are started; there is no forced context switching (see section 3.2). This restriction is justified with the implementation overhead caused by the pre-emption, which is unwanted in embedded time-critical systems. A computational task is called an actor; actors

are types of sub-functions, which jointly carry out some larger function. The scheduling of these actors starts from the processor assignment step, which means choosing the processor where an actor will be executed. The next task is the actor ordering step, which means the order that these actors are to execute when considering just a single processor. Finally, the execution start time, called firing, is determined for each actor. Different strategies can be classified based on which of the above-mentioned tasks are performed at the compile-time and which at the run-time. A *fully-static strategy* does each of the three tasks at the compile time. A *self-timed strategy* determines the start times at run-time, by communication between the processors in order to get tasks synchronized, but the ordering and allocation to processors are decided during the compiling. A *static allocation scheduling* only performs processor assignment step at the compile time. If all scheduling decisions are made at the run-time, the strategy is called *fully dynamic*. These are the four main classes; there are also two more modifications of these. A *quasi-static technique* has a small amount of run-time control, which handles ordering in the data-depended parts of the execution. An *ordered-transaction strategy* is almost like a self-timed approach, but the inter-processor communication order is determined at the compile time. [12, 33, pp. 55–67]

A *fully-static* (FS) strategy is used in systolic arrays and in very long instruction word (VLIW) processors. The timing of operations is known at the compile time, and this is enforced during the run-time, either by the programs themselves or by a program sequencer. The goal is usually to minimize the total schedule length, the make-span of the schedule, which also minimizes the idle periods for each processor. This strategy can further be divided into blocked and overlapping schedules. A blocked schedule completes the whole iteration before proceeding into the next one; the dependencies between successive iterations can thereby be ignored. The length of the critical path – which is the longest delay free path – specifies the minimum iteration period T for a blocked schedule. Some of the FS scheduling heuristics also take into account inter-processor communication (IPC), which causes costs when the processors need to communicate with each other. The enhancements to blocked schedules are unfolding and retiming. The unfolding strategy improves blocking schedules by scheduling the N iterations together. The number of replicated iterations N, which is called the blocking factor, also increases usage of the program memory by the same factor. In the retiming strategy, delays are manipulated in order to shorten the critical path, which directly affects the minimum of the iteration period. The overlapping strategy interleaves successive iterations, and is capable of lower iteration periods than blocking schedules, even when unfolding and retiming are used. [12, pp. 57–62]

When actor execution times are not known beforehand, in other words when execution times are varying, the fully-static strategy can be used with worst-case execution times. This is obviously an inefficient way, and secondly, the precise worst-case estimates may be unknown. In a *self-timed* scheduling (ST), the fully-static schedule is first built based on execution time estimates. Timing information is discarded; processor assignment and the ordering of the actors are instead used as such. Some of the actors are so called communication actors, performing send or receive operations; these are executed after all the input data is available. Sending actors are also blocked if the target buffer is full; correspondingly, receiving actors are blocked if the source buffer is empty. The senderreceiver synchronization is thus performed at the run-time; the disadvantage is a higher interprocessor communication cost due to this synchronization, when compared to the FS

strategy. In FS scheduling, there are typically only a few processor cycles required for the IPC, whereas in ST scheduling, the amount is dozens, unless special hardware for flow control is used. On the other hand, ST scheduling simplifies the compiler software because it does not have to adjust exact timing. [12, pp. 62–64]

A *fully dynamic* approach makes all the scheduling related decisions at the run-time. It has a very general applicability for various kinds of tasks; however it also has disadvantages such as the cost of performing these run-time decisions by special purpose hardware, like in superscalar processors, or by kernels running on one or more processors. Another disadvantage is that a dynamic scheduler usually only performs locally optimal decisions. [12, pp. 64–65]

A *quasi-static* scheduling is feasible with data-dependent execution times, which is generally caused by data-depended control structures, such as conditionals and datadependent iterations [12, pp. 65–67]. The firing of all nodes is decided at the compile time, but they are adjusted, when absolutely necessary, also at the run-time. Execution and scheduling progresses normally, until it reaches some data-dependent control structure. At this point, for example in the case of a branch with two unequal execution times, a run-time scheduler generates two different schedules, one for each branch. The difference here is that the initial pattern of processor availability is arbitrary. Obtained schedules are adjusted by padding them with idle periods, in order to make the processor availability patterns symmetric. If some other kind of synchronization between the processors is present, padding may not be required. If there is probability information available, about which branch will be executed, the higher probability branch can be scheduled first, and the second branch will be padded to fulfil the same pattern of processor availability. The absolute time values do not have to be identical, just the pattern. Finally, when this datadependent part of the schedule is completed, the original schedule can continue normally. [34].

3.4.2 Scheduling in shared memory systems

Fixed-priority scheduling algorithms in shared-memory multi-processor systems can also be divided between two methods: partitioned and non-partitioned. With the partitioned method, the execution of all instances of a task takes place on the same processor, which is already chosen before the run-time, by a partitioning algorithm. The non-partitioned version, which is also known as "dynamic binding" and "global scheduling", allows the task to be executed on any processor. This also includes the case when the task is continued after pre-emption. The non-partitioned method has less pre-emptions, since high-priority processes can utilize possible idle processors, rather than replace the executing task on the same processor. Since the partitioned method is guided by worst-case execution time estimates; whenever the actual execution times may be substantially lower than this, the utilization of processors remain low and would be better with a non-partitioned method. [35]

3.5 Performance gauging – the software perspective

In addition to the aspects mentioned in section 2.5, real benchmarking includes applicationspecific measuring. In real-time embedded systems, the fluency of context-switching and interrupt handling are two components required for a good real-time performance. The best results can be achieved with benchmark suites based on real-world algorithms. [4, pp. 28–31]

The over-designing of hardware could usually be avoided with proper software tuning, and the applicable points of the software can be pointed out with performance testing. The basic operation in performance testing is the measurement of the time consumed in the execution of a function. The execution is generally a nondeterministic process, factors such as the contents of instruction and data caches, operating system task loading, interrupts and other exceptions and finally data-processing requirements in the function cause diversity in the execution time. Therefore, the statistical measuring of minimum, maximum, average, and cumulative execution times is recommended. The execution times can basically be measured by identifying the memory addresses of the entry and exit points of the desired functions from the link map file. By observing the address bus, and recording the times when the corresponding addresses are detected, the execution time can be calculated. The difficulties in this technique include the calls of other functions inside the interesting one, with possible recursion, and interrupts whilst the function is executing; both aspects must be noticed in order to obtain correct results. Fortunately, the commercial tools built for performance testing can usually handle these kinds of problems. On-chip counters (see section 3.7.4) can be also used as performance measurement tools, for example to count elapsed time. [4, pp. 201–205]

3.6 Performance simulations – the software perspective

As explained in section 2.6, the division between hardware- and software-oriented performance simulators is a matter of opinion, as the performance is affected by both of them. This section wields simulators more from the side of software.

One approach in performance simulations is to have separate models for the applications to be executed, and for the hardware itself. It is even possible to use two different modelling languages to represent these two models; for example the Unified modelling language (UML) for the applications and SystemC modelling language for the hardware. The applications can be modelled by utilizing algorithm descriptions or the source code. To allow the execution of the applications modelled in a different language than the system, a proper interface between the models must be implemented. [36]

It is also possible to estimate the performance with a much higher level of abstraction. This usually means more abstraction to the hardware part. One practical and efficient way to simulate the performance of a computer-based system is to abstract their load somehow and then to consume it in an abstract simulated environment. The input data processing – called workload modelling – is an important step in this kind of simulation.

Simulation naturally has also its disadvantages and pitfalls. The development of a simulation model takes time and resources, the output of the simulation is just an estimate of the true system's characteristics and often the simulation results are given much more confidentiality than justified. [5, p. 77]. Especially in cases where a modelled system is only just at the proposal phase, its validation cannot be performed against any real world system, which causes an uncertainty in the results.

3.7 Workload modelling

Workload modelling can be described as an attempt to create a simple and general model, which can be used to generate synthetic workloads for systems that are under some kind of evaluation. One typical case, where a workload is required, is a performance analysis of computer-based systems. The workload presents the software that is meant to be executed in the actual system. A synthetic workload should be as similar as possible with the

real workload. A good workload model also makes it possible to generate various kinds of workloads with slight modifications. The workload modelling is usually based on measurements from the system of interest; the record or data log of workload-related events is often called a trace. An example of a single trace entry could consist of the arrival time of a job, where it was executed and what resources it required. This type of measurement-based modelling began strongly as late as the 1990's, when it was noticed that mathematical analyses, based on assumptions, differ notably from real workloads. [37, p. 10]

Workload modelling is about generalization and simplification. Measurements are always limited, the measuring process may be inconvenient or costly to realize, and the measurement instrumentation may cause an overhead and therefore disturb the results. The collected data is valid under strict conditions, for example, data which is collected from a 128-node supercomputer does not apply directly to 64- or 256-node supercomputers, but with a valid model based on the mentioned measurement, this limit can be overtaken. Models can also be executed with different seeds for random numbers, and therefore the statistical conditions remain the same, as required in the calculation of confidence intervals. This is not possible within a single trace. [37, pp. 10–12]

3.7.1 Alternative approaches in workload modelling

There are two different approaches to analyze or evaluate a system design with a measured workload. A simulation can be driven with a traced workload directly or alternatively with a model, which has been created from the trace. When using a trace directly, which means the measured workload, the system is going through a realistic test as the workload is accurate, and the person performing the analysis does not have to know every detail about the workload. On the other hand, a generalization to other systems or other system configurations may be a problem, because different systems with different architectures are probably unable to use the same trace at all. Information on the circumstances, when the trace was collected, should also be known. [38]

Workload models have some advantages, when compared to traces [38]:

- Models provide full information of workload characteristics to the modeller, for example, a correlation of different parameters is visible
- Model parameters can be changed one at a time, whereas manipulating traces with such an accuracy is challenging at the very least
- Models are not affected by the policies and constraints of the recording site
- Models enable the easier cleaning of so called bogus data, which can be for example killed jobs, which are executed multiple times before completion
- Workload modelling also increases understanding on the system requirements and can lead to new designs.

3.7.2 Creating a workload model

The creation of a workload model is based on data analysis; instead of trusting just the statistical methods, which are the primary tools in this task, it is desirable to also use graphical methods and common sense. Depending on the planned usage of the workload model one can choose between two different model types [37, pp. 15–16]:
- A descriptive model tries to describe just the measured phenomenon
- A generative model tries to articulate the process that originally generated the measured workload.

Descriptive modelling usually relies on creating a statistical summary of the observed workload. In most cases, the observation period should be as long as possible. Once the model is created, the synthetic workload can be obtained by sampling the distribution functions of the model. The proper distribution can be selected not by just fitting its shape, but also by some other feature, such as moments. Capturing a certain feature from the observed workload is sometimes sufficient, but when the effects of other features are unknown, it is safer to try the modelling as complete as possible. The validity of the model is dependent on many factors; if one attribute is modelled with great precision and other ones are formed by baseless assumptions, the validity is clearly not the best. The advantage of generative, which is also called indirect modelling, is that the workloadgeneration process can be manipulated, in order to obtain different types of workloads, which are still correct. [37, pp. 15–16]

The structure of a workload model depends on its purpose; the following examples are given in [37, pp. 5–6]. For scheduling problems, the relevant, sufficient attributes for the model are the arrival and running times for each job – there is no need for very specific details on what happens during the execution of the job. If the memory aspects of some system are also being examined, there is usually the need for total memory consumption and also a locality of reference. When I/O-processes are taken into consideration, the appropriate attributes are a distribution of I/O sizes and how they interleave with the execution. A typical case is that either computation or I/O is the scope of the evaluation, and the other one is modelled in a very abstracted way – just spending some time between the more interesting tasks. Parallel jobs can also have the number of processors used as an additional parameter. In contrast, when new microprocessor architecture is evaluated, the workload model must be much more specific. This type of model requires details such as instruction mix, instruction count between branches, loop sizes and dependencies between the instructions for instruction-level parallelism [39].

3.7.3 Types of workload

The workload itself can be classified into two different types [37, pp. 7–8]:

- A static workload has a certain amount of work, it gets completed at the end
- A dynamic workload contains jobs that are arriving all the time, and it will never get finished.

Because dynamic workloads include an arrival process, it must be also characterized and implemented properly which causes extra effort. A dynamic workload model should include all possible jobs and also their realistic frequency of occurrence. The difference between input distribution and "live system snapshot" must be noticed: even if the input has substantially more shorter jobs, a random snapshot may give an erroneous impression that longer jobs would be more common. This originates from the fact that the longer jobs naturally stay in the system for a longer period of time. The state of the system may affect the evaluation; a clean system may have a better performance than a system which has been executing a dynamic workload. Aging may also be present in a system; for example,

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memory leaks are one source for this kind of property in computer systems. Therefore, a dynamic workload model is usually the best representation of a real workload. [37, pp. 7–8]

3.7.4 Gathering data for modelling

As mentioned before in section 3.7, workload modelling is typically based on measured data. In some cases, data may be already available: for example, larger server machines usually log requests and other activities. However, these logs may not have a sufficient level of detail. If the data must be collected, it can be done by instrumenting the system in a way that its activities can be recorded. The required facilities should not modify the behaviour of the system at all; however, this can be problematic to implement. Depending on the instrumentations interaction with the actual system, the instrumentations can be divided into the following classes [37, pp. 18, 25–26]:

- A passive instrumentation does not modify the system itself
- An active instrumentation is integrated into the system, either at the design phase or afterwards.

Passive instrumentations are implemented with external components, which monitor system activity without interfering with it. For example, listening nodes can be added to the network in communication networks. These can record logs about the activity of the network, without sending any data into it. An example of an active instrumentation could be a microprocessor, which includes counters used to write down the amount of multiplication, division, load and store operations. These can be utilized when the workload is being modelled. [37, pp. 18, 25–26]

With active instrumentations, their interference should be as minimal as possible, and data buffering is one possible technique to reduce it. It means that a note of an event is stored in an internal buffer, and its output takes place later, in order to minimize the overhead caused by the data transfer [37, pp. 18, 26–27]. The so called instrumentation uncertainty principle has the following aspects [40]:

- Instrumentation causes perturbation into the system state
- Execution phenomena and instrumentation have a logical connection
- Volume and accuracy are contrary to each other.

Another method to manage instrumentations interference is to create models of perturbations, and then use them when approximating the traces, as they would be without instrumentation. This is called as perturbation analysis, and it consists of two phases: execution timing analysis and event trace analysis. The goal of the execution timing analysis is to adjust the trace event times, so that perturbations are removed. This is based on the measured costs of instrumentation. In event trace analysis, the sequence of events is adjusted based on event dependencies, in order to remove reordering caused by the instrumentation. [40]

One possible interference reduction technique is dynamically scalable instrumentation. The instrumentation can be inserted and altered during the execution of a program. This is implemented by modifying the binary image of the program while it is running. The volume of collected data is controlled by collecting only the information required at the moment, and also by adjusting the sampling rate. Instrumentation can be added in defined locations of application, called points. Instrumentation operations, which alter counters and timers, are called primitives. Boolean expressions, which affect instrumentation execution, are called predicates. Manual invocation is also possible. [41]

If the amount of collected data becomes too great, it is usually recommended to take samples over a longer period of time, than shorten the duration of the whole process. Anyway, sampling has some problems: choosing a constant interval between the samples can cause an aliasing effect on periodic data, and on the other hand, random sampling can also hide some characteristics of the data. If the data includes some internal structure, the sampling strategy should be constructed so that such structures are preserved. [37, p. 27]

One solution, providing dynamic instrumentation support for Linux operating systems, is named SystemTap. If a Linux kernel supports SystemTap and has proper debugging information included, the core of the operating system can be instrumented dynamically during the run-time. Probes are written with relatively simple scripting language. After compilation of a probe, it is injected into a kernel binary image. Besides the function entry and exit points, arbitrary statements are opportune probe positions. Whereas the probe's own variables can be used for example to count the amount of calls to a particular kernel function, whilst local variables in the target system can also be traced. [42]

This chapter describes the researched and implemented method during this work.

4.1 Our performance simulation process

The proposed simulation process can be divided into the following five steps, which are also presented in figure 12:

- 1. Workload modelling, which includes the instrumentation of the desired system, measuring performance data from it and finally creating a workload model from the gathered data
- 2. System modelling, which includes exploring the performance related parts of the system, like the degree of co-processing and scheduling, and presenting these with a programming language
- 3. Validation, if possible at all, can be done by executing the simulation model of some existing system and then comparing the simulation results with the real-world system
- 4. Simulation, which means the actual execution of the simulation models with the workload model, and gathering the desired results
- 5. Analysis, which can include three-dimensional visualization of the simulation results and possible comparison with the given performance requirements

Figure 12. Coarse representation of the main tasks of performance simulation process.

4.2 Workload modelling

The workload modelling process in our case was formed out of three different questions:

- What should we measure?
- How should we measure it?
- How should we process it?

Section 4.2.1 attempts to answer the first question. In section 4.2.2, possible solutions to the second question are discussed. Finally, section 4.2.3 covers the third question. In addition, we should also ask how our processed workload is validated to remain valid, however, in this work; we saw the postponement of the validation for a little bit as useful option. We will validate it together with the simulated system.

4.2.1 Proper data and its sources

An essential part of our method is to form appropriate input data for the performance simulator. We use measurements from existing systems, but not directly. Instead, we utilize the acquired data by creating a workload model out of it (see section 3.7). Due the nature of these simulations, approximating how the load of the uni-processor-system would execute in a multi-processor-environment, would require more information than with a typical execution trace log. The most interesting required additional information is related to the dependencies between processes and threads. The workload model will contain details about executed tasks, CPU times they have used and dependencies between various parts. These dependencies will have a substantial role in the execution in a multi-processor system, although the same also applies for uni-processor systems. In the extreme case, a performance gain cannot be achieved by increasing the amount of processors, if the workload is strictly sequential. Some kind of enhancement is usually possible, but its ratio to higher hardware costs must be taken into account.

The goal is to expose thread-level parallelism as much as possible (see section 3.1), and on the other hand, try to keep the parallelism accurate. In this work, the chosen approach was to research whether the scheduler of the operating system would be a feasible place to expose parallelism. The migration of tasks between ready and blocked queues, or corresponding task state changes, was thought to be a possible approach, in order to seek for dependencies and possible parallel parts. One deficiency – known already beforehand – is that a critical dependency between two processes can already be fulfilled before the latter process is executed at all. These processes should naturally be executed chronologically in the right order, but such dependency will remain unseen in this approach. Therefore, these tasks could be executed in an inverse order in our method. This all means that some parts of the workload models may be too optimistic. In order to bypass the described problem, we would need a great deal of additional information from the operating system. For starters, we should measure every possible synchronization mechanism, and so on. In addition, we also see only the last event which causes a task to be activated – a task can certainly have more than one of the prerequisites before its execution, but this approach is unable to find them.

Besides dependencies, we are also interested in the amount of work performed in the system. The amount of execution can be represented by a few alternatives. Firstly, it may be simply represented as a time since most systems allow easy measurement of the

system time at the beginning and ending points of desired event. The system time usually proceeds as a normal "wall clock" and therefore its comprehensibility for humans is good. However, this kind of representation is comparable only within identical systems, since devices with different computational capacities naturally do different amounts of work within an equal time interval. When these kinds of measurements are used in simulationrelated tasks, where the architecture under interest probably differs from the original one, this causes additional work and therefore reduces the usefulness of time-based representations.

By switching the unit from the time into processor clock cycles, some of the previously mentioned problems can be bypassed. The clock cycles are comparable between identical processor architectures running constantly or even dynamically in different frequencies. There may be a difference between different architectures, on how many cycles the processors require in order to execute the same instruction. The average of this processorspecific attribute is called cycles per instruction (CPI) [9, p. 42]. It also depends on the current workload, in other words what type of distribution of low-level instructions it has, but in the long run and with varying task sets, processor-specific values can be approximated. By dividing the measured workload in cycles with this value, the value is converted into an approximated total amount of instructions. The workload is now comparable between different architectures, as long as they implement the same instruction set architecture (ISA). An instruction set architecture is the interface between the hardware and the software [9, pp. 8–9]. If we do not want to take architecture modifications into account at all, they can be simply skipped by using equal values for the workload and a simulated system.

In Linux, the basic operating system scheduling concepts (see section 3.2) are used with minor modifications. Traditionally, the Linux scheduler has used a data structure called a run queue to gather together all the tasks which are ready to run or currently running on the CPU. These tasks have their state information set to running and when deciding the next task to run, the scheduler does its selection within this queue. With large number of tasks in the system, the scheduling decision will however have a substantial overhead. This naturally affects the performance of the whole system. Another deficiency was that multi-processor systems will have processors queuing for the next task, since the single queue must be locked, while one of the processors is choosing its task. The enhancements for the problems were per-CPU and per-priority run queues, which overcame both of the problems: per-CPU run queues obviously solve the problem related to locking, and storing runnable tasks into per-priority lists changes the rather complicated scheduling decision into the scheduler just knowing the highest priority level which has tasks [27, pp. 266–268]. There are 140 different priority levels in total and also 140 lists respectively. Moving a task into the run queue is called activating a task, and correspondingly, deactivating a task means its removal from the run queue. So, instead of having both running and ready queues, Linux uses the same queue for currently running tasks and for those which are waiting to receive processor time. The centralized blocked queue, discussed in section 3.2, is also split into several smaller per-event or per-resource entities in Linux; these are called wait queues and tasks in these are said to be suspended. The state of these kinds of tasks is set to interruptible or uninterruptible, depending on their reaction to possibly arriving signals [27, pp. 81–82].

Linux kernel versions 2.6.23 and 2.6.24, which were the current official mainline versions during this work, have a so called completely fair scheduler (CFS). It does not use run queues as with former versions; runnable tasks are stored instead into a data structure called a red-black tree. However, from our viewpoint, its functionality is the same as that of a run queue because the relevant information is the temporal relations and dependencies between tasks. The activation of a task is handled with one centralized function, as well as its deactivation. All context switches are also progressed through a centralized place in the kernel code. We will look at these three places in more detail in section 4.2.2. In summary, the Linux operating systems are evidently providing the data that we are interested of.

4.2.2 Instrumentation

The tracing part in this work evolved through two different approaches. The first implemented instrumentation into the Linux scheduler requires kernel source code modifications. It consists of actual logging functions and a simple controlling functionality via the Linux Procfs virtual file system, to begin and end the measurements as we do not want to normally measure all the time. Each of the logging points stores the system time, which is provided at nanosecond accuracy by the kernel, and selected relevant kernel variable values. An existing Linux feature, called kernel messages, was found to be feasible for the trace logging task and no modifications or new features were therefore needed to that functionality. The kernel message log is primarily saved into a size-limited virtual file and at intervals into a physical file. The physical file is useful, when a desired data is transferred into the next phases of the workload modelling process.

The second tracing approach was the utilization of SystemTap framework (see section 3.7.4). Due to its dynamical instrumentation abilities, no kernel code modifications and recompilation are required. This helps particularly in tracing the development phase and in principle, it gives an easier conversion between different kernel versions. When tracing function entry and exit points, modifications are not needed as long as both the kernels still have the same functions. However, in our approach we also had probe positions which had to be placed inside functions by using source code line numbers and therefore the probes must be fine-tuned for each kernel version separately.

During the latter approach, the unit of time measurement was also changed from nanoseconds to processor clock cycles. Because the workload is finally presented as an amount of processor instructions, measurements based on the system time must be first converted to clock cycles and then scaled with a processor-specific CPI ratio. Performing the tracing directly with clock cycles reduced one unnecessary step from the modelling.

The selected points, already mentioned briefly in the section 4.2.1, were three functions responsible for task activations, task deactivations and actual scheduling. Both activation and deactivation are short simple functions and the probing point inside these does not matter much. In the actual scheduling function, the probing point was selected just before the actual context switching, in order to ensure that the function is really going to switch tasks. Thus, we have three probing points and three different message types, respectively. All of the messages begin with a timestamp, which was already mentioned to be in the CPU cycles. The message then has a type identifier: ACTI, DEAC or COSW for activation, deactivation or context switch messages respectively. The following parts are message-specific and we will look at them next.

Table 1 presents a message from the probe located in the task activation function. The message has an activated task's ID as its third field. The fourth field denotes the previous state of this task, which is the state before activation. In Linux, 0 is the only executable

state and numbers above 0 are different classes of blocked or otherwise non-executable states. Because the task's state is inevitably 0, after an activation, we do not have to include it into this message. As the last field, the message has the activated task's priority.

Table 1. The form of a task activation message with an example

			Timestamp Msg type Task ID Previous state Priority	
387000567	ACTI	3959		120

Table 2 presents a corresponding message from the deactivation function. Respectively, it has a deactivated task's ID as its third field. The fourth field denotes the new state of the task. Deactivation of a task is about switching its state to something above 0. The last field is the task's priority.

Table 2. The form of a task deactivation message with an example

Timestamp Msg type Task ID New state Priority			
387051332	DEAC	3959	120

Table 3 presents a message denoting a context switch. In this message, the third field means the previous task, which is about to be removed from the processor. The fourth field denotes its state $-$ if it is 0, the task would still be runnable, otherwise it has been deactivated and we presumably already have a deactivation message on that event. The fifth field is the priority of the previous task. The sixth field denotes the next task, which the system is about to execute. The last field represents the priority of the next task.

Table 3. The form of a context-switch message with an example

Timestamp Msg type Prev. Prev. state Prev. prio. Next Next prio.				
$ 387009901 $ COSW $ 3963 $		120	$ 3959\rangle$	120

With the described message types, the instrumentation generates an execution log. Figure 13 shows what kind of output we get from the instrumentation. In the figure, a task numbered 3959 is emphasized in order to clarify how its execution progresses.

4.2.3 Creating the workload model

The workload model presents the workload as bipartite per-task queues. The two elements forming these queues are *executable* and *waiting elements*, and due bipartite structure, an executable element is always followed by a waiting element and vice versa. Either of the elements can be the first element in a queue, depending on the initial state of the corresponding task. An executable element includes a finite amount of instructions which must be "executed" before the element is deleted from the head of the queue. Additionally, it can have one or more of *activations*, which activate other, possibly blocked tasks. A waiting element is removed from the head of the queue, when a dependency related to it is fulfilled. Figure 14 shows the structure of a simple workload. The white rectangles present executable parts. They are followed by hexagonal waiting elements. Small rectan-

Figure 13. Short clip from a log produced by the scheduler instrumentation.

gles inside executable parts are activations, which become realized during the execution of executable parts.

Figure 14. Our workload model's structure.

The described model is created by reading our tracing log (presented in section 4.2.2) line by line. Figure 15 presents the high level structure of the developed algorithm responsible for the model creation. Each timestamp of the message is used to increase the the amount of processing of the currently running task. During a context switch, the next task will gain a new executable element in its queue's tail, if there is not one already. If the the state of the previous task is blocked, its queue will be lengthened with a new waiting element

with a unique dependency stamp. An activation message causes the tail of the queue of the activated task to be examined; the number of the dependency it has been waiting for is fetched and a new activation event is associated into currently running task's element. A deactivation message causes the deactivated task's executable element to be tagged as deactivated, in order to handle it properly during the next context switch.

```
switch Message type do
    case Context switch
        switch Task type do
            case Next
                if No executable element in the tail then
                 Create a new executable element;
                end
            end
            case Previous
                Add the amount of processing for this task;
                if State==blocked then
                    Create a new waiting element with a new dependency stamp;
                 \mathbb{R}end
            end
        end
    end
    case Activation
        Add the amount of processing for the currently running task;
        Fetch the dependency stamp from the tail of the activated task;
        Create a new activation event into the currently running task;
    end
    case Deactivation
        Add the amount of processing for the currently running task;
        Tag the deactivated task as deactivated;
   end
end
```
Figure 15. Pseudo code of the workload modeller.

The deactivation message is redundant for most of the time, as only the currently running task will be typically deactivated, and the very same deactivation information could be acquired during the next context switch from the previous task's state. There are however occasions, where tasks use kernel-provided macros in order to access their state information directly and thereby passing normal activation and deactivation functions. These kinds of "invisible" activations and deactivations will disturb the algorithm and in the worst case they cause a deadlock in the model. Therefore, when handling context switches, we treat cases where state information is accessed directly in a special way. Since no deactivation message is found, it is also a sign of a missing activation message in the near future. We therefore process these cases, as the deactivated task would remain runnable. The amount of errors caused by this was empirically discovered to be insignificant; since these cases seem to be quite rare and on the other hand, these tasks are usually returned to the execution just in a couple of context switches anyhow.

The current version also includes the idle task as a normal, executable work. The reason

for this is that the idle can often represent the time when the system is waiting for user interactions. The response times of the user must naturally also be taken into account also in the simulated system, and the easiest way to ensure it is to let the processors also execute the idle sections.

4.3 System modelling and simulation

In our case, the system modelling means the development of an abstract consumer for the generated workload model. Obviously, the workload should also be consumed equally such as with the real system, which we are trying to model. The abstraction level of the simulator, for the actual execution part of the tasks, is very high – basically, we just perform a repetitive subtraction operation, at a speed defined by the simulated processors' cycles-per-instruction value. The scheduling is modelled more accurately, and the dependencies, presented in the workload model, are naturally strictly obeyed when the tasks are executed.

The implemented performance simulator has some modularity, and therefore different scheduling algorithms and system architecture alternatives should be easy to add afterwards. The first system implemented and discussed in this work is a homogeneous architecture with a centralized scheduler. This means that the processors use the scheduler individually to fetch the next task, and the scheduling can only take place for one processor at a time. Other processors will be forced to wait if there is a concurrent attempt for scheduler access. This causes an extra overhead to the system, besides the normal scheduling overhead. The main memory is also centralized and shared in the model (see section 2.3.2), which eases the simulation as memory access times can be thought to be same as in the original system. The simulator implements the finite state machine presented in figure 16.

The simulator is fed with the workload model (see section 4.2.3). First, the model is checked to be deadlock-free, because a part of the workload will otherwise remain unexecuted. The simulator begins by forming initial run and wait queues. Initially runnable tasks of the workload model are naturally placed into the run queue, and non-runnable tasks respectively into the wait queue. The processors are modelled in dedicated threads, which are synchronously proceeding simulation steps. The simulation step is freely adjustable, beginning from one clock cycle. The first actual state after the initial preparations is the *idle* state. However, a flag denoting a need for rescheduling is initially up and each processor therefore calls the scheduler. One of the processors manages to access the scheduler, whilst others are forced to wait. *Waiting for the scheduler* is modelled as a dedicated state in the state machine. As previously mentioned, this is one factor that lowers the efficiency of systems with a centralized scheduler, as the processors are unable to do useful work, while they are waiting for the scheduler. After successful access to the scheduler, either by waiting for or accessing it directly, the simulation models the *scheduling overhead*, which presents the overhead related to the scheduling procedures. Selecting the next task from a group of several tasks causes some extra computation. Additionally, scheduling is usually followed by a context switch, which means transferring the data of the next task to the processor. This also causes some overhead, but in our case, the measured workload already includes both the scheduling and context switching overhead. The amount of the overhead can be set to represent an estimate of the additional overhead caused by the multi-processor scheduling. As already said, queuing the scheduler access is taken into account in this simulation and it does not need to be approximated in the

Figure 16. A finite state machine, with rather few states, is used to simulate a homogeneous architecture with a centralized scheduler.

scheduling overhead.

In the worst case, a context switch may require a reloading of the whole cache memory. This situation is called a cold cache. A cold cache is the consequence of a "new" task's scheduling to a processor: its cache is occupied by the previous task's data and the cache must be thereby cleared and then loaded with the proper data. This means that the tasks should be kept running within the same processor as much as possible. If the cache has the correct data ready for a task, the cache is respectively called a hot cache. Multiprocessor scheduling algorithms can utilize knowledge on the tasks' previously allocated processors and thereby, they can attempt to avoid movement of the tasks between different processors. The implemented algorithm in the simulator did not include this kind of feature, as the measured workload also includes the cache overhead from the original system.

After the scheduling overhead has been consumed, the next state is usually the actual execution of a task, in a state called as *processing a task*. The only exceptions to this are the cases when a rescheduling flag has been raised during the scheduling overhead consumption, which causes the processor to schedule again, or that there is no task available to the processor, which causes it to enter the idle state. Without an immediate rescheduling, the simulation begins to execute the scheduled task. Depending on the set CPI value, one instruction usually takes more than one, for example 1.5 cycles to complete. As with the workload modelling, we do not use an exact instruction-specific CPI value but an estimated average of it. During the execution of tasks, the dependencies presented in the workload model become fulfilled. The fulfilment triggers the following operations in the simulator: an update in a dependency matrix, a migration of corresponding tasks from the wait queue to the run queue and a possible rescheduling. The dependency matrix is

used for tracking fulfilled dependencies – a waiting point in the workload model can be simply skipped if the required dependency is already filled in the simulated run. The task execution continues until a need for reschedule flag is raised, the task uses its entire time slice or the runnable part of the task is consumed totally. All of these cause the processor to schedule, and depending on the scheduler's availability, the next possible states are waiting the scheduler or processing the scheduler overhead.

The implemented scheduling algorithm has a centralized run-queue, which furthermore consists of separate round-robin (see section 3.3) sub-queues for each priority. In general terms, this algorithm is a simplified and centralized version of the algorithm of the previous Linux versions (see section 4.2.1). Because the highest priority process with the longest waiting time is always selected, and on the other hand a rescheduling occurs every time tasks are activated, the algorithm assures that N runnable processes with the highest priorities are always executing in a system with N processors. When looking more closely at the simulators internal functionality, in addition to the task itself, the scheduler also returns the scheduling overhead and time slice values, as shown in figure 17. As this type of per-priority run queue scheduler has a virtually constant overhead, and the workload model itself includes a scheduling overhead from the original system, we did not develop any kind of approximation function for the scheduling overhead. Instead, the used scheduler returns a small constant amount of cycles. Of course, the overhead could be also accurately modelled based on the actual computation needed in picking the next task. In the current version, the given time slice is also modelled in a coarse way: it varies stochastically between the values that we observed with measurements from the real-world system. The real operating systems calculate this value premising on characteristics of the task, such as the load of the system.

Figure 17. The scheduler returns a structure, which consists of the whole task event queue, scheduling overhead and time slice values.

4.4 Analysis with visualization

A 3D-visualization tool called PerVisGL, originally created by Yrjönen [7], was extended during this work to support multiple data logs, in order to visualize multi-processor systems. In addition, we also developed other minor features for it, mostly related to easier

task identification. PerVisGL allows the use of three-dimensional space freely, to represent arbitrary data after a data-specific reader has been developed. In this case, we selected adjacent 100 x 10 x 10 rectangular boxes to present the execution. Each of the described boxes denotes one processor. The longest dimension is aligned with the x-axis and is 100 units. This denotes the time, and despite of the actual duration of the log, the whole log is scaled to this interval. The y-axis, up to 10 units, denotes the relative load of the CPU in a linear fashion. The z-axis, up to 10 units as well, is relative to the priorities of the tasks, but not linearly. Normal non-real-time tasks use a scale from 0 to 8 units, real-time tasks use the next distance from 8 to 9 units and so called background load, which is the scheduling overhead, is the longest in the z-direction, reaching a full 10 units. Thus, the "thickness" of the graph presents the importance of the work; the background load may not be esteemed, but it is obligatory and at least important to notify.

We also use colours as an additional dimension. Tasks can be identified by their colours; the colour value for a task is calculated from its task number. The function, calculating this value, uses a simple hash method in order to clearly present different colours for consecutive task numbers. One can also give desired task numbers, which will be the only ones to be coloured. This notably emphasizes the graph, when there are dozens or hundreds of tasks presented simultaneously.

Instead of drawing the tasks strictly in a sequential order along the time-axis, the application splits the space into smaller blocks. This is due to the overwhelming amount of context switches present in a typical log; the graph would be as hard to read as the original, raw text-form log itself. For example, if we want to present our data in ten blocks, the first block represents the distribution of computation in the first tenth of the log, and its dimensions are 10 x 10 x 10 respectively. The boxes presenting individual tasks are drawn in this space, one upon the other. Tasks with the highest priorities go to the bottom, because they were also the thickest ones. The block now fills the x-axis for one tenth of the total distance, but the y- and the z-axes have varying sizes in different positions. The height of a box presents the corresponding task's share of computation during the block interval, and because the boxes are piled vertically, the top of the topmost box also shows the total relative computation. Any free space over the topmost box correspondingly represents the amount of idle. The z-dimension is also task-wise and represents the priority, as already mentioned in this section.

Figure 18 is used here as a practical example of the functionality of the visualization. The figure has roughly ten blocks visible. In the very first block, three different tasks, drawn in blue, red and orange, have been executed, and all of them have different priorities. "The orange task" had the highest priority and therefore it is drawn at the bottom with the thickest box. Furthermore, these three tasks, together, used roughly 55 percent of the available CPU time during this block's interval, since there are 4.5 units of free space at the top in the y-axis. In the next blocks, only red and blue tasks have been running, and finally in the last blocks visible, there is only a green task running.

Figure 18 also shows how the visualization tool draws grids to help in the determining of load and priority values. The distance between two grid lines is one unit, so in the direction of the y-axis, the lines correspond to a change of 10 % in the load of the system. In the direction of the x-axis, the scale is not linear since most of it, the values from 0 to 8 units, were defined for non-real-time tasks as we are usually more interested of them. Non-real-time tasks in Linux have only 40 priorities against the 100 priorities of real-time tasks [27, p. 265].

Figure 18. A simple visualization to clarify the visualization applications functionality.

The interface between the simulator and the visualization tool is a set of log files: one file per simulated processor. One line in a log represents a set time interval, however a context switch, always triggers the logging so that data on each task's execution is acquired properly. Each line has a time stamp, the amount of overhead in the interval, the amount of the actual processing in the interval, a task number and a priority. Because the time stamp in the log file is in processor cycles, the simulated device's clock rate must be entered into the visualization tool, in order to present the correct "wall clock" time values in the graph.

5. Results

This chapter presents the results achieved during the work. Firstly, the performed validations are discussed and then we will analyze the whole method.

5.1 Validation of the models

As explained in section 3.6 the simulation models can be hard to validate completely. In this work, two somewhat different validation methods were applied. The first one was more coarse and meant for early phase testing, whereas the second one gives useful results about the accuracy and feasibility of the approach.

The first validation was performed with a simple self-made application, which runs a selected amount of threads using an inter-thread synchronization mechanism. We will measure the execution of this application, model it and finally simulate the acquired workload with different multi-processor systems. As the internal functionality of the application is known, it is possible to see if our performance simulation method gives too optimistic or too pessimistic results for the given workload.

The second validation method is more general and clearly more challenging. The workload was measured from a dual-core PC with operating system which was forced to use only one core. The measured workload is then simulated in modelled dual-processor architecture and the simulation results are compared to a run of the same tasks in a PC which is using the both cores. The modelled and real-world systems should now be effectively the same. The result of this is that the difference in the results directly presents the accuracy of the whole method.

In other words, we are validating the workload model, the simulation model and the whole method at the same time.

5.1.1 Threads with barrier synchronization

The self-made application for this validation purpose performed repetitive memory operations with three child threads. These child threads each had 250 iterations and the iterations were synchronized with the so called barrier synchronization, which holds the threads in a barrier point, until they have all reached it. In our case, with three threads, the first and second threads to reach the barrier are blocked temporarily. The third thread continues its execution directly without blocking at the barrier, and this event also releases the two other ones to continue.

This test case was selected to be inconvenient for our method. Since, one of the threads in every iteration proceeds without blocking, the method will not see that dependency at all, although there clearly is one by knowing the test application's internal structure. Thereby, the method could produce results that are too optimistic. However, in the long run, the threads will encounter an equal amount of blocking when compared to each other, and therefore they are still clearly synchronized with each other, although one dependency per iteration is missed. This means that the method should be able to also simulate the execution in a roughly synchronized way.

Figure 19 presents the visualization of the trace achieved from the described test application. Only the threads affiliated to the test application, one parent, number 11786, and three child threads, numbers 11787, 11788 and 11789, are coloured. Particularly in the middle part, the three child threads can be seen to execute quite equally.

Table 4 presents the original trace and the results for four different simulated platforms. The amount of total execution is given in millions of CPU cycles, and the relative perfor-

Figure 19. Visualization of the test application's trace.

mance, which is the ratio of execution times, is scaled to be 1 for the measured device. As we knew that there is thread-level parallelism between the three threads, one could expect a linear gain in performance with two and three processors. The CPI values were the same for all of the systems including the measured one, so we do modelled similar architectures with different degrees of parallel processing. The dual-processor gets very close to doubling the performance, and the system with three processors is not so far from triplication. So far, the results are within the presumptions.

So it seems as if the method performed quite well in exposing sufficient parallelism for the simulated architectures with two and three processors. Figure 20 presents a visualization of the simulation run with two processors available. The utilization for both of the processors can be seen to be quite high, as there are runnable threads available all the time.

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Figure 20. The visualization of the test application run in a simulated dual-processor system.

In figure 21, the test application is simulated to use three processors. The workload manages to keep all the available processors relatively busy, which is still reasonable since we know that the three threads should be able to run concurrently.

Figure 21. The visualization of the test application run in a simulated triple-processor system.

When we add one more processor to the system, we should not get a significant performance gain, because there is not enough parallelism available. The small exceeding of the relative performance of 3 with four processors can be explained with other small tasks present in the workload. In figure 22, we can see how the execution is divided among four processing units. The graphical analysis also confirms that there are idle slots clearly visible, and the utilization of the system is not the best possible. Figure 23 demonstrates the scalability of the simulation also in addition to the visualization. The simulated system has eight processors and therefore it could be a type of super-computer. The figure clearly shows that the amount of parallelism in the workload was not sufficient for this kind of

architecture and therefore the processors are idling a lot.

Figure 22. The visualization of the test application run in a simulated quad-processor system.

Besides the total execution times and 3D-graphs, we can also receive additional information about the execution. Table 5 presents per-processor characteristics about the simulated architectures. We had four states in the state machine representing one processor (see section 4.3). The simulator counts how many cycles each processor has spent in each of the states. The numbers are averaged here from the individual statistics of the processors. All of the units are in millions of cycles. The amounts of processing and idle are a partially repetition of what we have already discussed, but the scheduling statistics have not been reviewed yet. As expected, the overhead, caused by having a single scheduler for multiple processors, rises quickly, as the amount of processors increases. On the other hand, the per-processor scheduling overhead decreases. This is due to the fact that with a static amount of workload and an increasing number of processors, one processor has less scheduling occasions.

System	Processing	Idle	Waiting for scheduler	Scheduling overhead
Dual	7882	93.1		2.3
Triple	5254	298		
Quad	3941	993	2.6	1.8
Octa	1971	2790	5.1	.6

Table 5. Per-processor details about the simulation runs

Figure 23. The visualization of the test application run in a simulated super-computer, which has eight processors in total.

5.1.2 Threaded video encoding

The selected application for the second part of the validation was FFmpeg [43], which is a multi-threaded video encoding software for Linux. Normally the application selects a suitable amount of threads to use depending on the hardware, naturally one thread per one processor, but in this case, it was forced to use two threads for encoding. In addition to the two threads performing the encoding, the application's "main thread" is also running and probably performing some kind of control tasks, but it does not utilize the CPU as heavily as the two other threads. The encoder's threads also have a lot of dependencies between them; threads are almost always blocked when they are switched from the execution. This is a good situation for our validation purposes. For example, a threaded file compression was initially planned as a validation case, but its threads were noticed to remain runnable when they were switched out. Due to this fact, the validation would have been too trivial, as the dependencies and their fulfilments would only have had a small effect.

The operating system was Fedora Core 8 Linux with a self-built kernel version 2.6.24.3. Self-building of the kernel was required due to the SystemTap instrumentation. The operating system was forced to use only one of the two cores provided by the Core 2 Duo workstation, running at 2.16 GHz. The test case was to encode a motion JPEG AVI file into a DVD PAL MPEG-2 format. The size of the original file was 10.0 megabytes and the output file enlarged to 23.7 megabytes. This encoding process was measured and then fed into the combination of the workload modeller and simulator. Figure 24 shows that there were also plenty of other tasks involved. However, most of the processing was done by the three threads of the video encoder.

The video encoding was also executed in the described workstation with both cores in

Figure 24. The visualization of the trace, from the encoding process, which was used when the workload model was created.

use, in order to provide reference results. The total process of the uni-core measurement, the simulated two-processor execution and the real dual-core execution, was repeated five times, as there was some deviations in the measurements. This is due the fact that we used a normal operating system with obligatory background tasks running. There will also be a number of other tasks running at both the heads and tails of logs, which must be subtracted before the results are comparable to each other. In other words, we are interested in the part, which begins when the video encoder was launched, and ends when the video encoder exits. Table 6 presents the results. Both of the workload statistics are given in millions of CPU cycles. The relative performance is scaled to be 1 for the real uni-core system. For this workload, the simulation gave an average of 12 % too pessimistic results. However, by taking standard deviations into account, the results are overlapping with the real system, so the results in the outline were quite much as expected. Once again, we used identical CPI values in order to only focus on the validation of the parallelism at first.

System	Average		Standard deviation Relative performance
Real uni-core	83241	5333	
Real dual-core	37576	3648	2.21
Simulated dual-core	42093	2874	198

Table 6. Results of the second part of the validation

Figure 25 shows the visualization of the output of the simulator. After arbitrary tasks at the beginning, it can be seen that the video encoder is receiving CPU time quite equally from both processors. As the current implementation of the simulated scheduler does not have any policy for keeping the same threads running in the same processors, the threads are migrating a lot between the processors. The workload model actually already has some extra variables for additional information, and one of them is the last place of execution. So, a more sophisticated scheduling algorithm could use this information when it decides the next task. Although this figure has other tasks present at both ends of the graphs, the values given in table 6 were calculated by using the first and the last occurrence of the encoder's threads.

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Figure 25. The visualization of the encoding simulation results with all the tasks shown equally.

Figure 26 has only the encoder's threads emphasized, for easier understanding. This figure also shows that, in the beginning, there is a group of other tasks causing the load to the system. After that, the visualization reveals that the encoding process is able to roughly utilize 90 % of the available processing power in our simulated two-processor architecture. This type of information can be a very useful to a system designer, and on the other hand, obtaining the value from raw text-form logs would probably be troublesome or inconvenient. In addition to this broad overview, the visualization also shows more detailed statistics. For example, some higher priority processes, longer in the Z-direction and drawn with black and grey, are also occasionally receiving small partitions of the CPU time.

Figure 26. The visualization of the encoding simulation results, where only the threads belonging to the encoding task are emphasized.

In figure 27, we have selected a short interval of about 1.3 million CPU cycles from the first processor's graph under closer examination. We can see that two threads performing the actual encoding, numbers 3961 and 3963, are responsible for nearly two thirds of the CPU's utilization. Since the visualization tool had only the three encoding threads emphasized, the numeric summary also only gives detailed information only about these threads, and all the other 15 threads have just their combined CPU time visible.

Figure 27. The visualization tool allows to pick arbitrary region under closer examination.

5.2 Analysis of the whole method

We have already analyzed the individual cases, and in this section we will look more at the whole method. Even though the synchronization mechanism in the first validation case was intentionally selected to be difficult, the results were logical. Despite the periodical loss of "invisible" dependencies during the measurement, the acquired workload model seemed to be feasible in the longer run. The simulated architecture's scaling to a virtually unlimited quantity of processors was also found to be functional.

In the latter validation case, the encoder's relative performance was well over 2 in the real two-processor architecture. Our method has a theoretical upper limit of 2 for dualprocessor simulations, as it always executes everything that we have measured, and at its best with double speed, but not any faster. Although we simulated a centralized scheduler, whereas the real dual-core platform used dedicated run queues for both of the processors and load balancing between the queues when needed, these differences were not relevant. The simulator spent less than 8 million of CPU cycles waiting for the scheduler in the runs averaging 42 billion of cycles, which is less than 0.02 %. The video encoding algorithm seems to have some kind of "synergy" benefits, when it is running in multiple cores. Some reasons for this can be seen by processing the measured log files a little. One event causing the unexpected speed-up is the fact that by counting the context switches related to these encoding threads, the amount was 37 % smaller on average in the real-world dualcore system than its uni-core configuration. Because the current version of the workload modelling also includes the context switching overhead from the measured system as an executable workload, the simulation is unable to achieve this kind of performance gains.

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In summary, some factors of a parallel processing performance gain are unobtainable for our method's current functionality. The instrumentation also has its effect on the measured workload, which we will look closer at in section 6.2.

The greatest disappointment this time was not found in the method's characteristics, but from the practical implementation of its one part. The performance simulator itself is very slow to execute, and despite an adjustable simulation step, the simulation takes significantly more time than the corresponding real-world process. The code of the simulator should be optimized.

Despite the promise of two cases, we have to remember that we performed several validation steps at once, including the workload model, the system model and the method in general. The scope of these two cases in the domain of computer systems, where workloads can be virtually anything, is also far from sufficient. However, in summary, we are satisfied with the obtained results.

6. Discussion

We will review here the strengths and weaknesses of the method. As the work was just the first phase of a larger work, we will also cover some possible future scenarios.

6.1 Advantages of the method

This section presents the aspects that we found particularly successful during the work. We will look first at the subtopics and then discuss the whole method.

The *instrumentation* provides an easy completion of measurements, since we only need information from three different points of an operating system: task activations, task deactivations and context switches. If the states of the tasks can be reliably checked during the context switches, a previous task can be found out to be deactivated from the context switch's probing point. Probing the deactivations becomes effectively redundant and thereby only two points of the operating system must be instrumented. The instrumentation's functionality is also simple and needs only a few lines of additional code. Due to these facts, all operating systems, implementing somewhat the same kind of centralized active and blocked task handling, should be possible to instrument with the given guidelines.

The *workload modelling* step is automated and does not require user interaction, excluding possible CPI value settings. When compared to some other methods, the user does not need to be familiar with the workload in detail.

Due to the *simulation's* accurate modelling of the scheduling, it could be used in a uniprocessor domain as well – in addition to parallel processing, one could test uni-processor scheduling algorithms. As the simulator uses workloads which are measured from realworld devices, the algorithms would be under realistic testing.

Our *visualization* approach is commonly thought to be easy to understand. It can handle varying amounts of data and visualize them in an equally understandable way. Emphasizing the desired parts of the execution, and giving statistics on the selected areas were noticed to be useful resources.

Despite the *whole method's* high level of abstraction, the cases selected for validation at least gave reasonable and comparatively accurate results. The autonomous processing of measurements from real-world devices, and their linkage into simulated domains was also found to be a working solution. This removes the demand for manual workload creation present in several other methods. The most important breakthrough during the work was the exploration of parallelism and dependencies by instrumenting the scheduling queues. This idea also proved to be feasible and relatively straightforward to implement.

6.2 Considerations of the method

We already had knowledge of some of the method's restrictions at the beginning of the work, and we also encountered some new ones during the work. This section declares these aspects. Once again, we will first look at the individual areas and then the complete method.

Despite the small code size of the *instrumentation*, we are probing a very sensitive part of an operating system. The target events are also recurring very frequently. This causes an unavoidable probing effect, which makes the workloads to seem larger than they really are. Equally, the measured system seems to be slower, which also affects the simulated system. Table 7 presents statistics on five test runs, where the video encoding task from the second validation case was run with and without instrumentation, in order

6. Discussion

to estimate the amount of the probing effect. The duration of the encoding process was measured in seconds. The amount of the overhead, caused by the instrumentation, seems to be 5.1 % in this case. The value depends on the task switching pace of the workload, as every context switch, task activation and task deactivation cause an additional logging event. By measuring this kind of "calibration value" for a given process, and then scaling the amounts of the execution during the workload model creation, the method's accuracy could be refined further. On the other hand, the relative performance gain between the instrumented original system and the simulated system should be roughly the same as between the uninstrumented original system and the real-world system which we are trying to model. This is due to the fact that the instrumentation slows down both the measured and simulated system at the same ratio. In summary, one could focus more on the relative performance gain than the absolute duration of the execution, in order to bypass the error caused by the instrumentation.

Table 7. Test runs of a real system with and without instrumentation

System		Average Standard deviation
Uni-core without instrumentation \vert 35.237 s		1.3043 s
Uni-core with instrumentation	37.031 s	1.2602 s

The sizes of the logs are also large, as there are plenty of necessary events. Some kind of sampling – but still conserving the dependencies – would be useful if we desire to measure longer executions. In this work, we focused only on quick example runs. However, the results can be also easily extrapolated for long lasting cases.

Because we are gain dependencies only by observing task activations and deactivations, we do not know what the class of the dependency is. For example, when a task is blocked, we do not know whether it is waiting for a user input, waiting for data from some device, or is it synchronized to proceed exactly at the same rate as its sibling threads. By spreading instrumentation into other places of an operating system, we could also gain this kind of data, but also compromise on the instrumentation's adaptability in other operating systems. As already mentioned in the section 4.2.1, the method has also some "blind spots" for detecting dependencies. The method will not recognize dependencies, which were fulfilled before their appearance. In the case of several prerequisites, only the last one which actually activates some task is noticed.

The measurements could also be cleared out of the scheduling, context switching and cache reloading overheads. This could be performed by more wide-ranging kernel instrumentation. For example, instead of one probe in the middle of the main scheduling function, there could be probes at the beginning and at the end of the function. With pure amounts of execution, we would have to take more details into the simulation; it would however also improve the method's accuracy.

In order to measure multi-processor systems, for example dividing the load from a dualprocessor system to a simulated quad-processor version, the instrumentation would only require a processor ID number to its messages. The implemented Linux-instrumentation already views all events from each of the processors, but because our focus was to measure uni-processor systems, the current version does not separate the events of different processors in any way.

Our *simulation* process was found to be slower to execute, than the process which it is

mimicking. Simulation studies are typically performed in accelerated time for systems, whose internal working does not need to be known in a very detailed way [5, p. 77]. This would be advantageous for us, in order to flexibly examine several possible architectures, however the current implementation does not provide this kind of quickness. For example, the encoding tasks presented in section 5.1.2 were under 40 and 20 seconds when executed in uni- and dual-core systems. The duration of the simulation run for this workload was nearly 60 minutes. The utilization of rather heavy-weighted data structures such as C++ vectors and maps in the workload models, as well as in the run and waiting queues is probably the main reason for the slowness.

The *visualization* application does not give any kind of information on dependencies at its present state. This kind of extension is easy to implement at a code level, however, finding a practical and understandable graphical form is more difficult.

Because the *method* is seeking thread-level parallelism (see section 3.1) among the already existing threads, we do not gain much new information out of the workloads which are not using threads. However, the method will find out the independent processes and allocate them to different processors when appropriate, but that is an assignment which could be essentially completed by using just common sense, instead of any kind of simulations. For example, we can say that executions of a word processor and an Internet browser – two clearly uncorrelated applications – could be successfully allocated to different processors in order to get a performance gain. In addition, some multi-threaded applications are adaptive enough to execute only a single thread in a single processor system, which would obviously harm measurements in our method. The method does not try to find parallelism inside applications written sequentially.

6.3 Future work

During this work, we only focused on the homogeneous systems. A simulation of heterogeneous systems would be technically a lightweight update. On the other hand, knowledge on the workload's parts, which could be executed in non-general-purpose processors, is hard to automate. Another difficulty is to approximate the amount of the scaled workload in such processors. One possible way would be to manually seek for tasks executing some algorithms, which are known to be feasible to implement with more hardwareoriented solutions. The ID numbers of these tasks would then be fed to the simulator with a configuration file. The scheduler would give these tasks only to co-processors, when they are runnable. Another possible approach would require more details on the workload model. The next detail level could be a representation of work with very basic operations, such as memory reads, memory writes and executions [36]. With characteristics modelled for these operations per each computational unit, the heterogeneous system would be able to consume the workload in a proper way.

7. Conclusion

This thesis presented a novel method for exposing thread-level parallelism from an existing device's workload and simulating its execution in a multi-processor environment. The work confirmed that the original, high-level idea of a modelling load from existing devices to new, more parallel architectures is doable.

We also presented a validation of the method with two practical cases: the first case with an "ad-hoc" test application and the second one with a more common video encoding process. According to the case studies, the approach seems to be valid. The achieved estimation of the method's inaccuracy was from 10 to 15 %. On the other hand, more wide-ranging validation cases will be still required before the true characteristics of the method are known.

The method supports system designers in the early phases of the system design flow, when the required amount of co-processing for the new system is being decided. The most important contributions provided by this work are:

- An automatic generation of workload models from measurements
- Finding parallelism from a few simple locations of an operating system
- Highly abstracted simulation models
- Easily understandable visualizations of multi-processor systems' load.

As our method has the automated workload generation, the amount of required work is small when compared to several, more detailed simulation solutions. The results are still useful, especially if the alternative is just an educated guess about the designed system's performance. The method however does not give support for analysing a single processor's internal architecture changes.

The method exposes parallelism at the level of an operating system, and therefore does not require a detailed knowledge on the workload's internal structure. Our method provides support for many kinds of workloads without modifications, although the most obvious way would be to attempt to find parallelism directly from the applications and algorithms running on the measured system – and thereby lose the versatility. The restriction is that the workload must be a threaded one. The generalization of the method for other source systems is good, because the required instrumentation was kept intentionally as minimal as possible. This is naturally aiding adaptability to other operating systems.

The high level of abstraction with the models mimicking the systems under design also provides easiness for the system designers. The method's scalability for a wide range of number of parallel processors in the simulated system was found to be reasonable. The extension of the method for heterogeneous system support was also discussed and seems doable.

The method also provides support for the analysis of results, as our 3D-visualizations were proved to be more informative than typical raw data logs. The designer can have a quick overview of the distribution of the load in the system at a glance, whereas it would not be possible with unprocessed text-form data.

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Title

Performance Simulation of Multi-processor Systems based on Load Reallocation

Abstract

This work presents the novel method for high-level performance estimation of systems consisting of multiple computational units. The goal is to support system designers in the early phases of the system design flow. The focus mainly lies on embedded systems and in this first part of the work, we began from their versions which perform parallel processing with execution units similar to each other. Systems consisting of different types of processors, and the method expansions to support them are also discussed.

The main idea was an attempt to reallocate a single processor's load to multiple simulated processors. The method uses measurements from actual, existing systems and relies on means of simulations with systems under design. Instead of competing with prototyping, the method is supposed to give an estimation of which kind of system architecture would fulfil the desired performance requirements.

In the method, we process the mentioned measurement data automatically, which results in a so-called workload model. The workload model is then executed with a simulated system. This simulation run approximates the proposed system's estimated performance. Due to automation at the modelling phase and a high level of abstraction, the method allows the fast approximation of several different configurations.

The first of the problem areas was to define which type of workload model is suitable and how it can be created. When the workload is measured from a uni-processor system, its parts which can be parallel executed must be discovered, in order to use the model with a multi-processor system. The second problem area is the modelling of the performance-related parts of the system under design. The larger problem is to study the validity and rationality of the whole method.

We validated the method with two different test cases and both of them gave reasonable results. The first validation consists of a simple threaded application, which uses an inter-thread synchronization mechanism. As the internal functionality of the application is known, the characteristics of the method can be roughly seen. The second validation method is a real-world algorithm, which we will execute in both a simulated and existing twoprocessor system. The margin for error of the method can be calculated from the latter of the validation cases, by comparing the total execution times of the systems. The margin for error for this case was from 10 to 15 %. It was better than expected for a method with a rather high level of abstraction.

As research results, the work presents the parts needed for the method: an instrumentation for gathering the measurement data, the creation of a workload model out of it, a simulation of a multi-processor system with the workload model, and visualization of the simulation results. In addition, an analysis of these parts and the whole method is presented.

Julkaisun sarja, numero ja raporttikoodi

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Nimeke

Suorituskykysimulaatio moniprosessorijärjestelmille kuorman uudelleenjakamisen avulla

Tiivistelmä

 I _{ICDN}

Tämä työ esittelee menetelmän korkean tason suorituskykyarviointiin järjestelmälle, joka koostuu useammasta suoritinyksiköstä. Tarkoituksena on tukea suunnittelijoita järjestelmän määrittelyvaiheessa. Menetelmä on tarkoitettu ensisijaisesti sulautetuille järjestelmille, ja tässä laajemman työn ensimmäisessä vaiheessa mielenkiinto oli sellaisissa versioissa, joissa rinnakkainen suoritus tapahtuu samanlaisia suorittimia käyttäen. Työssä käsitellään myös keskenään erityyppisistä suorittimista koostuvia järjestelmiä ja menetelmän laajennusta tukemaan myös niiden analyysiä.

Tärkeimpänä osa-alueena menetelmässä on yrittää jakaa yhden prosessorin kuorma useammalle simuloidulle prosessorille. Menetelmä käyttää mitattua dataa olemassa olevista järjestelmistä ja tukeutuu simulointiin suunnitteluvaiheessa olevien järjestelmien kanssa. Menetelmää ei ole tarkoitettu kilpailemaan prototyyppien tekemisen kanssa vaan antamaan arvio siitä, minkälainen arkkitehtuuri täyttäisi halutut suorituskykyvaatimukset.

Olemassa olevista järjestelmistä mitattua kuormitusdataa prosessoidaan automaattisesti, ja tulosta kutsutaan kuormamalliksi. Tätä mallia käytetään syötteenä simulointivaiheelle, joka jäljittelee suunniteltavana olevan järjestelmän käytöstä. Simuloinnin tulokset antavat informaatiota järjestelmän ennustetusta suorituskyvystä. Esimerkiksi tietyn kuorman kokonaissuoritusaika on yksinkertainen suorituskyvyn mitta. Mallinnuksen automaatiosta sekä menetelmän korkeasta abstraktiotasosta johtuen eri arkkitehtuurivaihtoehtojen arviointi on nopeaa.

Työn ensimmäinen ongelma-alue oli sopivan kuormamallin löytäminen. Jotta kuormamalli soveltuisi moniprosessorijärjestelmille, sen luonnollisesti tulee pystyä erottelemaan rinnakkaiseen suoritukseen soveltuvat osat, kuten myös riippuvuudet eri osien välillä. Seuraava ongelma-alue on mallintaa suorituskykyyn liittyvät osa-alueet suunnittelussa olevasta järjestelmästä. Isompana kokonaisuutena olivat koko menetelmän järkevyyden ja oikeellisuuden tarkastelut.

Menetelmä validoitiin kahta erilaista lähestymistapaa käyttäen. Ensimmäinen validointitapa toteutettiin yksinkertaisella säikeistetyllä ohjelmalla, joka käytti säikeidenvälistä synkronointia. Koska ohjelman sisäinen rakenne on nyt tunnettu, menetelmän toiminnallisuus voidaan nähdä karkeasti. Toinen validointitapa on todellinen algoritmi, joka suoritettiin sekä simuloidulla että olemassa olevalla kaksiprosessorijärjestelmällä. Jälkimmäisestä validointitavasta pystyttiin laskemaan menetelmän virhemarginaali vertaamalla molempien ajojen kokonaissuoritusaikoja. Virhemarginaaliksi tälle tapaukselle saatiin noin 10–15 %. Tämä virhemarginaali oli odotettua parempi menetelmän korkea abstraktiotaso huomioon ottaen.

Tutkimustuloksina esitellään menetelmään tarvittavat osa-alueet: instrumentointi mittausdatan saamiseksi, kuormamallin muodostaminen tästä datasta, moniprosessorijärjestelmien simulointi edellä mainitun mallin avulla sekä tulosten visualisointi. Lisäksi esitellään menetelmän ja sen osa-alueiden analysointi.

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