# Towards Adjusting Mobile Devices to User's Behaviour

Peter Fricke<sup>1</sup>, Felix Jungermann<sup>1</sup>, Katharina Morik<sup>1</sup>, Nico Piatkowski<sup>1</sup>, Olaf Spinczyk<sup>2</sup>, and Marco Stolpe<sup>1</sup>

> <sup>1</sup> Technical University of Dortmund, Artificial Intelligence Group Baroper Strasse 301, Dortmund, Germany

{fricke,jungermann,morik,piatkowski,stolpe}@ls8.cs.tu-dortmund.de, http://www-ai.cs.tu-dortmund.de

> <sup>2</sup> Technical University of Dortmund, Embedded System Software Group Otto-Hahn-Strasse 16, Dortmund, Germany

> > olaf.spinczyk@tu-dortmund.de, http://ess.cs.uni-dortmund.de

Abstract. Mobile devices are a special class of resource-constrained embedded devices. Computing power, memory, the available energy, and network bandwidth are often severely limited. These constrained resources require extensive optimization of a mobile system compared to larger systems. Any needless operation has to be avoided. Timeconsuming operations have to be started early on. For instance, loading files ideally starts before the user wants to access the file. So-called prefetching strategies optimize system's operation. Our goal is to adjust such strategies on the basis of logged system data. Optimization is then achieved by predicting an application's behavior based on facts learned from earlier runs on the same system. In this paper, we analyze system-calls on operating system level and compare two paradigms, namely server-based and device-based learning. The results could be used to optimize the runtime behaviour of mobile devices.

Keywords: Mining system calls, ubiquitous knowledge discovery

# 1 Introduction

Users demand mobile devices to have long battery life, short application startup time, and low latencies. Mobile devices are constrained in computing power, memory, energy, and network connectivity. This conflict between user expectations and resource constraints can be reduced, if we tailor a mobile device such that it uses its capacities carefully for exactly the user's needs, i.e., the services, that the user wants to use. Predicting the user's behavior given previous behavior is a machine learning task. For example, based on the learning of most often used file path components, a system may avoid unnecessary probing of files and could intelligently prefetch files. Prefetching those files, which soon will be accessed by the system, leads to a grouping of multiple scattered I/O requests to a batched one and, accordingly, conservation of energy.

The resource restrictions of mobile devices motivate the application of machine learning for predicting user behavior. At the same time, machine learning dissipates resources. There are four critical resource constraints:

- Data gathering: logging user actions uses processing capacity.
- Data storage: the training and test data as well as the learned model use memory.
- Communication: if training and testing is performed on a central server, sending data and the resulting model uses the communication network.
- Response time: the prediction of usage, i.e., the model application, has to happen in short real-time.

The dilemma of saving resources at the device through learning which, in turn, uses up resources, can be solved in several ways. Here, we set aside the problem of data gathering and its prerequisites on behalf of operation systems for embedded systems [13] [22] [3]. This is an important issue in its own right. Regarding the other restrictions, especially the restriction of memory, leads us to two alternatives.

- Server-based learning: The learning of usage profiles from data is performed on a server and only the resulting model is communicated back to the device. Learning is less restricted in runtime and memory consumption. Just the learned model must obey the runtime and communication restrictions. Hence, a complex learning method is applicable. Figure 1 shows this alternative.
- **Device-based learning:** The learning of usage profiles on the device is severely restricted in complexity. It does not need any communication but requires training data to be stored. Data streaming algorithms come into play in two alternative ways. First, descriptive algorithms incrementally build-up a compact way to store data. They do not classify or predict anything. Hence, in addition, simple methods are needed that learn from the aggregated compact data. Second, simple online algorithms predict usage behavior in realtime. The latter option might only be possible if specialized hardware is used, e.g., General Purpose GPUs. Figure 2 shows this alternative.

In this paper, we want to investigate the two alternatives using logged system calls. Server-based learning is exemplified by predicting file-access patterns in order to enhance prefetching. It is an open question whether structural models are demanded for the prediction of user behavior on the basis of system calls, or simpler models such as Naive Bayes suffice. Should the sequential nature of system calls be taken into account by the algorithm? Or is it sufficient to encode the sequences into the features? Or should features as well as algorithm be capable of explicitly addressing sequences? We investigate the use of two extremes,

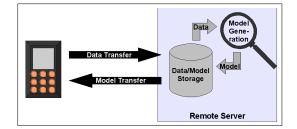


Fig. 1. Server-based Architecture

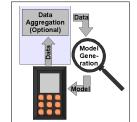


Fig. 2. Device-based Architecture

Conditional Random Fields (CRF) and Naive Bayes (NB). In particular, we inspect their memory consumption and runtime, both, for training and applying the learned function. Section 2 presents the study of server-based learning for ubiquitous devices. We derive the learning task from the need of enhancing prefetching strategies, describe the log data used, and present the learning results together with resource consumptions of NB and CRF.

Device-based learning is exemplified by recognizing applications from system calls in order to prevent fraud. We apply the data streaming algorithm Hierarchical Heavy Hitters (HHH) yielding a compact data structure for storage. Using these, the simple kNN method classifies systems calls. In particular, we investigate how much HHH compress data. Section 3 presents the study of device-based learning using a streaming algorithm for storing compact data. We conclude in Section 4 by indicating related and future work.

# 2 Server-based Learning

In this section we present the first case-study, where log data are stored and analyzed on a server (data are described in Section 2.2). Learning aims at predicting file access in order to prefetch files (see Section 2.1). The learning methods NB and CRF are introduced shortly in Section 2.3 and Section 2.4, respectively. The results are shown in Section 2.5.

## 2.1 File-access pattern prediction

A prediction of file-access patterns is of major importance for the performance and resource consumption of system software. For example, the Linux operating system uses a large "buffer cache" memory for disk blocks. If a requested disk block is already stored in the cache (*cache hit*), the operating system can deliver it to the application much faster and with less energy consumption than otherwise (*cache miss*). In order to manage the cache the operating system has to implement two strategies, block replacement and prefetching. The *block replacement* strategy is consulted upon a cache miss: a new block has to be inserted into the cache. If the cache is already full, the strategy has to decide which block has to be replaced. The most effective victim is the one with the longest forward distance, i.e. the block with the maximum difference between now and the time of the next access. This requires to know or guess the future sequence of cache access. The *prefetching* strategy proactively loads blocks from disk into the cache, even if they have not been requested by an application, yet. This often pays off, because reading a bigger amount of blocks at once is more efficient than multiple read operations. However, prefetching should only be performed if a block will be needed in the near future. For both strategies, block replacement and prefetching, a good prediction of future application behavior is crucial.

Linux and other operating systems still use simple heuristic implementations of the buffer cache management strategies. For instance, the prefetching code in Linux [2] continuously monitors read operations. As long as a file is accessed sequentially the read ahead is increased. Certain upper and lower bounds restrict the risk of mispredictions. This heuristics has two flaws:

- No prefetching is performed *before* the first read operation on a specific file, e.g., after "open", or even earlier.
- The strategy is based on assumptions on typical disk performance and buffer cache sizes, in general. However, these assumptions might turn out to be wrong in certain application areas or for certain users.

Prefetching based on machine learning avoids both problems. Prefetching can already be performed when a file is opened. It only depends on the prediction that the file will be read. The prediction is based on empirical data and not on mere assumptions. If the usage data change, the model changes, as well.

## 2.2 System Call Data for Access Prediction

We logged streams of system calls of type FILE, which consist of various typical sub-sequences, each starting with an **open-** and terminating with a **close**-call, like those shown in Figure 3. We collapsed such sub-sequences to one observation and assign the class label

- full, if the opened file was read from the first seek (if any) to the end,
- read, if the opened file was randomly accessed and
- zero, if the opened file was not read after all.

We propose the following generalization of obtained filenames. If a file is regular, we remove anything except the filename extension. Directory names are replaced by "DIR", except for paths starting with "/tmp" – those are replaced by "TEMP". Any other filenames are replaced by "OTHER". This generalization of filenames yields good results in our experiments. Volatile information like thread-id, process-id, parent-id and system-call parameters is dropped, and consecutive observations are compound to one sequence if they belong to the same process. The resulting dataset consists of 673887 observations in 80661 sequences, a snippet<sup>3</sup> is shown in Table 1.

 $<sup>^{3}</sup>$  The final dataset is available at:

http://www-ai.cs.tu-dortmund.de/PUBDOWNLOAD/MUSE2010

```
1,open,1812,179,178,201,200,firefox,/etc/hosts,524288,438,7 : 361, full
2,read,1812,179,178,201,200,firefox,/etc/hosts,4096,361
3,read,1812,179,178,201,200,firefox,/etc/hosts,4096,0
4,close,1812,179,178,201,200,firefox,/etc/hosts
```

**Fig. 3.** A sequence of system calls to *read* a file. The data layout is: timestamp, syscall, thread-id, process-id, parent, user, group, exec, file, parameters (optional) : read bytes, label (optional)

user	group	exec	file	label
201	200	firefox-bin	cookies.sqlite-	zero
			journal	
201	200	$\operatorname{firefox-bin}$	default	zero
201	200	$\operatorname{firefox-bin}$	hosts	full
201	200	$\operatorname{firefox-bin}$	hosts	full
201	200	multiload-	mtab	full
		apple		
102	200	kmail	png	zero

**Table 1.** Snippet of the preprocessed dataset (the marked row corresponds to the open call of Fig. 3).

predicted	true	full	zero	read

full	0	2	1
zero	5	0	4
read	4	2	0
Table 2.	Cost n	natrix	

exec	file	label
?_firefox-bin	?_?_cookies.sqlite-journal	zero
firefox-bin_firefox-bin	?_cookies.sqlite-journal_default	zero
firefox-bin_firefox-bin	$cookies.sqlite-journal\_default\_hosts$	full
firefox-bin_firefox-bin	default_hosts_hosts	full
?_multiload-apple	?_?_mtab	full
?_kmail	?_?_png	zero

 Table 3. Snippet of the final dataset using two features.

We used two feature sets for the given task. The first encodes information about sequencing as features, resulting in 24 features, namely  $f_t$ ,  $f_{t-1}$ ,  $f_{t-2}$ ,  $f_{t-2}/f_{t-1}$ ,  $f_{t-1}/f_t$ ,  $f_{t-2}/f_{t-1}/f_t$ , with  $f \in \{user, group, exec, file\}$ . The second feature set simply uses two features  $exec_{t-1}/exec_t$  and  $file_{t-2}/file_{t-1}/file_t$  as its only features – an excerpt of the dataset using these two features is shown in Table 3.

Errors in predicting the types of access result in different degrees of failure. Predicting a partial caching of a file, if just the rights of a file have to be changed, is not as problematic as predicting a partial read if the file is to be read completely. Hence, we define a cost-matrix (see Table 2) for the evaluation of our approach. For further research the values used in this matrix might have to be readjusted based on results of concrete experiments on mobile devices or simulators.

#### 2.3 Naive Bayes Classifier

The Naive Bayes classifier [10] assigns labels  $y \in Y$  to example  $x \in X$ . Each example is a vector of m attributes written here as  $x_i$ , where i = 1...m. The probability of a label given an example is according to the Bayes Theorem:

$$p(Y|x_1, x_2, ..., x_m) = \frac{p(Y)p(x_1, x_2, ..., x_m|Y)}{p(x_1, x_2, ..., x_m)}$$
(1)

Domingos and Pazzani [7] rewrite eq. (1) and define the *Simple Bayes Classifier* (SBC):

$$p(Y|x_1, x_2, ..., x_m) = \frac{p(Y)}{p(x_1, x_2, ..., x_m)} \prod_{j=1}^n p(x_j|Y)$$
(2)

The classifier delivers the most probable class Y for a given example  $x = x_1 \dots x_m$ :

$$\arg\max_{Y} p(Y|x_1, x_2, ..., x_m) = \frac{p(Y)}{p(x_1, x_2, ..., x_m)} \prod_{j=1}^m p(x_j|Y)$$
(3)

The term  $p(x_1, x_2, ..., x_m)$  can be neglected in eq. (3) because it is a constant for every class  $y \in Y$ . The decision for the most probable class y for a given example x just depends on p(Y) and  $p(x_i|Y)$  for i = 1...m. These probabilities can be calculated after one run on the training data. So, the training runtime is  $\mathcal{O}(n)$ , where n is the number of examples in the training set. The number of probabilities to be stored during training are  $|\mathcal{Y}| + (\sum_{i=1}^{m} |\mathcal{X}_j| * |\mathcal{Y}|)$ , where  $|\mathcal{Y}|$  is the number of classes and  $|\mathcal{X}_i|$  is the number of different values of the *i*th attribute. The storage requirements for the trained model are  $\mathcal{O}(mn)$ .

It has often been shown that SBC or NBC perform quite well for many data mining tasks [7, 11, 8].

# 2.4 Linear-chain Conditional Random Fields

Linear-chain Conditional Random Fields, introduced by Lafferty et al. [12], can be understood as discriminative, sequential version of Naive Bayes Classifiers. The conditional probability for an actual sequence of labels  $\mathbf{y_1}, \mathbf{y_2}, ..., \mathbf{y_m}$ , given a sequence of observations  $\mathbf{x_1}, \mathbf{x_2}, ..., \mathbf{x_m}$  is modeled as an exponential family. The underlying assumption is that a class label at the current timestep t just depends on the label of its direct ancestor, given the observation sequence. Dependency among the observations is not explicitly represented, which allows the use of rich, overlapping features. Equation 4 shows the model formulation of linear-chain CRF

$$p_{\lambda}\left(Y=\mathbf{y}|X=\mathbf{x}\right) = \frac{1}{Z\left(\mathbf{x}\right)} \prod_{t=1}^{T} \exp\left(\sum_{k} \lambda_{k} f_{k}\left(y_{t}, y_{t-1}, \mathbf{x}\right)\right)$$
(4)

with the observation-sequence dependent normalization factor

$$Z(\mathbf{x}) = \sum_{\mathbf{y}} \prod_{t=1}^{T} \exp\left(\sum_{k} \lambda_k f_k(y_t, y_{t-1}, \mathbf{x})\right)$$
(5)

The sufficient statistics or *feature functions*  $f_k$  are most often binary indicator functions which evaluate to 1 only for a single combination of class label(s) and attribute value. The parameters  $\lambda_k$  can be regarded as weights or scores for this feature functions. In linear-chain CRF, each attribute value usually gets  $|\mathcal{Y}| + |\mathcal{Y}|^2$ parameters, that is one score per state-attribute pair as well as one score for every transition-attribute triple, which results in a total of  $\sum_{i=1}^{m} |\mathcal{X}_i| (|\mathcal{Y}| + |\mathcal{Y}|^2)$ model parameters, where  $|\mathcal{Y}|$  is the number of classes, m is the number of attributes and  $|\mathcal{X}_i|$  is the number of different values of the *i*th attribute. Notice that the feature functions explicitly depend on the whole observation-sequence rather than on the attributes at time t. Hence, it is possible and common to involve attributes of preceding as well as following observations from the current sequence into the computation of the total score  $\exp(\sum_k \lambda_k f_k(y_t, y_{t-1}, \mathbf{x}))$  for the transition from  $y_{t-1}$  to  $y_t$  given  $\mathbf{x}$ .

The parameters are usually estimated by the maximum-likelihood method, i.e., maximizing the conditional likelihood (Eq. 6) by quasi-Newton [14], [19], [15] or stochastic gradient methods [25], [17], [18].

$$\mathcal{L}(\lambda) = \prod_{i=1}^{N} p_{\lambda}(Y = \mathbf{y}^{(i)} | X = \mathbf{x}^{(i)})$$
(6)

The actual class prediction for an unlabeled observation-sequence is done by the Viterbi algorithm known from Hidden Marcov Models [21], [16].

Although CRF in general allow to model arbitrary dependencies between the class labels, efficient exact inference can solely be done for linear-chain CRF. This is no problem here, because they match the sequential structure of our system-call data, presented in section 2.2.

#### 2.5 Results of Server-based Prediction

Comparing the prediction quality of the simple NB models and the more complex CRF models, surprisingly, the CRF are only slightly better when using the two best features (see Tables 4 and 6). CRF outperforms NB when using all features (see Tables 5 and 7). These two findings indicate that the sequence information is not as important as we expected. Neither encoding the sequence into features

nor applying an algorithm which is made for sequential information outperforms a simple model. The Tables show that precision, recall, accuracy, and misclassification cost are quite homogeneous for CRF, but vary for NB. In particular, the precision of predicting "read" and the recall of class "zero" differs from the numbers for the other classes, respectively. This makes CRF more reliable.

Inspecting resource consumption, we stored models of the two methods for both feature sets and for various numbers of examples to show the practical storage needs of the methods. Table 10 presents the model sizes of the naive Bayes classifier on both feature sets and for various example set sizes. We used the popular open source data mining tool  $RapidMiner^4$  for these experiments. Table 10 also shows the model sizes of CRF on both feature sets and various example set sizes.

predicted\true	full	zero	read	prec.
full	1427467	19409	3427	98.43
zero	12541	2469821	40258	97.91
read	80872	217380	2467695	89.22
recall	93.86	91.25	98.26	

Table 4. Result of Naive Bayes Classifier on best Table 5. Result of Naive Bayes two features, 10x10-fold cross-validated, accuracy: Classifier on all 24 features,  $94.45 \pm 0.00$ , missclassification costs:  $0.152 \pm 0.001$  10x10-fold cross-validated, accu-

predicted\true	full	zero	read	prec.
full	1446242	7123	29051	97.56
zero	19452		133007	
read	55186	60390	2349322	95.31
recall	95.09	97.51	93.55	

full	zero	read	prec.
1426858	21562	22717	96.99
15392	2371009	97566	95.45
78630	314039	2391097	85.89
93.82	87.60	95.21	

racy:  $91.84 \pm 0.00$ , missclassification costs: 0.218  $\pm$  0.002

read prec. 25629 97.71 126403 94.93 2359348 95.36 93.95

read	prec.	full	zero
29051	97.56	1450147	8335
133007	94.54	14563	2639724
2349322	95.31	56170	58551
93.55		95.35	97.53

Table 6. Result of HMM-like CRF on the best two features, 10x10-fold cross-validated, accuracy:  $95.49 \pm 0.00$ , missclassification costs:  $0.150 \pm 0.000$ 

Table	7.	Resu	ılt	of	HM	M-like
CRF on	all	24  fea	atui	res,	10x1	0-fold
cross-va	lida	ated,	ace	cura	acy:	95.70
$\pm 0.00$	, m	isscla	assif	icat	tion	costs:
$0.143 \pm$	0.0	000				

read

5022

95 51

2662966 107859 95.75 39527 2398499 96.69

prec

99.38

zero

4117

98.39

predicted\true	full	zero	read	prec.	full
full	1467440	4733	7503	99.17	146809
zero	10883	2659294	108340	95.71	10306
read	42557	42583	2395537	96.57	42479
recall	96.49	98.25	95.39		96.53

Table 8. Result of linear-chain CRF on the best Table 9. Result of linear-chain two features, 10x10-fold cross-validated, accuracy: CRF on all 24 features, 10x10-fold  $96.79 \pm 0.00$ , missclassification costs:  $0.112 \pm 0.000$  cross-validated, accuracy: 96.89

 $\pm$  0.00, missclassification costs:  $0.110 \pm 0.000$ 

 $<sup>^4</sup>$  Rapid Miner is available at: http://www.rapidminer.com

#Att.\#Seq.	0	67k	135k	202k	270k	337k	404k	472k	539k	606k	674k
2 nB	244	248	251	253	256	255	256	257	257	256	256
24 nB	548	561	571	577	582	585	588	590	590	585	585
2 CRF++ (HMM)	5	247	366	458	490	512	569	592	614	634	649
24 CRF++ (HMM)	12	615	878	1102	1170	1216	1367	1420	1463	1521	1551
2 CRF++	6	523	776	978	1043	1089	1213	1260	1299	1345	1378
24 CRF++	19	1339	1914	2415	2559	2652	2988	3095	3184	3303	3365

**Table 10.** Storage needs (in kB) of the naive Bayes (nB) classifier model produced by *RapidMiner*, the HMM-like CRF (CRF++ (HMM)) and the linear-chain CRF (CRF++) on different numbers of sequences and attributes.

#Att. #Seq.	0	67k	135k	202k	270k	337k	404k	472k	539k	606k	674k
2 nB	< 1	< 1	< 1	< 1	1	< 1	< 1	< 1	< 1	< 1	< 1
24 nB	< 1	< 1	< 1	1	< 1	1	1	1	1	2	1
2 CRF++ (HMM)											
24 CRF++ (HMM)	< 1	27.92	55.9	103.24	153.53	160.33	230.7	273.29	232.84	309.19	317.62
2 CRF++	< 1	16.69	50.23	85.18	113.21	145.96	173.56	200.98	234.65	260.56	325.54
24  CRF++	< 1	41.06	105.29	156.67	296.31	300.83	343.28	433.03	440.88	463.84	632.96

**Table 11.** Training time (in seconds) of the naive Bayes (nB) classifier model produced by RapidMiner, the HMM-like CRF (CRF++ (HMM)) and the linear-chain CRF (CRF++) on different numbers of sequences and attributes.

We used the open source CRF implementation  $CRF++^5$  with  $L_2$ -regularization,  $\sigma = 1$  and L-BFGS optimizer in all CRF experiments. Obviously, the storage needs for a model produced by a NB classifier are lower than those for a CRF model. This is the price to be paid for more reliable prediction quality. CRF don't scale-up well. Considering training time, the picture becomes worse. Table 11 shows the training time of linear-chain or HMM-like CRF consuming orders of magnitude more time than NB.

# 3 Device-based Learning

In this section, we present the second case-study, where streams of log data are processed in order to store patterns of system use. The goal is to aggregate the streaming system data. A simple learning method might then use the aggregated data. The method of Hierarchical Heavy Hitters (HHH) is defined in Section 3.1. The log data are shown in Section 3.2. For the comparison of different sets of HHH, we present a distance measure that allows for clustering or classifying sets of HHH. In addition to the quality of our HHH application, its resource consumption is presented in Section 3.3.

#### 3.1 Hierarchical Heavy Hitters

The heavy hitter problem consists of finding all frequent elements and their frequency values in a data set. According to Cormode [4], given a (multi)set S

<sup>&</sup>lt;sup>5</sup> CRF++ is available at: http://crfpp.sourceforge.net/

of size N and a threshold  $0 < \phi < 1$ , an element e is a *heavy hitter* if its frequency f(e) in S is not smaller than  $\lfloor \phi N \rfloor$ . The set of heavy hitters is then  $HH = \{e | f(e) \ge \lfloor \phi N \rfloor\}$ .

If the elements in S originate from a hierarchical domain D, one can state the following problem [4]:

**Definition 1 (HHH Problem).** Given a (multi)set S of size N with elements e from a hierarchical domain D of height h, a threshold  $\phi \in (0, 1)$  and an error parameter  $\epsilon \in (0, \phi)$ , the Hierarchical Heavy Hitter Problem is that of identifying prefixes  $P \in D$ , and estimates  $f_p$  of their associated frequencies, on the first N consecutive elements  $S_N$  of S to satisfy the following conditions:

 $\begin{array}{l} - \ accuracy: \ f_p^* - \varepsilon N \leq f_p \leq f_p^*, \ where \ f_p^* \ is \ the \ true \ frequency \ of \ p \ in \ S_N. \\ - \ coverage: \ all \ prefixes \ q \not\in P \ satisfy \ \phi N > \sum f(e) : (e \preceq q) \land (\not\supseteq p \in P : e \preceq p). \end{array}$ 

Here,  $e \leq p$  means that element e is generalizable to p (or e = p). For the extended multi-dimensional heavy hitter problem introduced in [5], elements can be multi-dimensional d-tuples of hierarchical values that originate from d different hierarchical domains with depth  $h_i$ ,  $i = 1, \ldots, d$ . There exist two variants of algorithms for the calculation of multi-dimensional HHHs: Full Ancestry and Partial Ancestry, which we have both implemented. For a detailed description of these algorithms, see [6].

#### 3.2 System Call Data for HHH

FILE	COMM	PROC	INFO	DEV
open read write lseek _llseek writev fcntl fcntl64 dup dup2 dup3	recvmsg recv send sendfile sendto rt_sigaction pipe pipe2 socket accept	PROC mmap2 munmap brk clone fork vfork mprotect unshare execve futex nanosleep	access getdents getdents64 clock_gettime gettimeofday time uname poll fstat fstat64 lstat	DEV
cntl cntl64 up up2	rt_sigaction pipe pipe2 socket	mprotect unshare execve futex	uname poll fstat fstat64	
			<pre>stat64 inotify_init inotify_init1 readlink select</pre>	

**Table 12.** We focus on 54 system call types which are functionally categorized into five groups. FILE: file system operations, COMM: communication, PROC: process and memory management, INFO: informative calls, DEV: operations on devices.

The kernel of current Linux operating systems offers about 320 different types of system calls to developers. Having gathered all system calls made by

several applications, we observed that about 99% of all calls belonged to one of the 54 different call types shown in Tab. 12. The functional categorization of system calls into five groups is due to [20]. We focus on those calls only, since the remaining 266 call types are contained in only 1% of the data and therefore can't be frequent.

HHHs can handle values that have a hierarchical structure. We have utilized this expressive power by representing system calls as tuples of up to three hierarchical feature values. Each value originates from a taxonomy (type, path or sequence) that either can be derived dynamically from the data itself or has to be defined explicitly by the user. The groups introduced in Tab. 12 form the top level of the taxonomy for the hierarchical variable type (see Fig. 4). The socket call is a child of group COMM and FILE is the parent of calls like open and fcnt164. Subtypes of system calls can be defined by considering the possible values of their parameters. For example, the fcnt164 call which operates on file descriptors has fd, cmd and arg as its parameters. We have divided the 16 different nominal values of the *cmd* parameter into seven groups — notify, dflags, duplicate, sig, lock, fflags and lease — that have become the children of the fcntl64 system call in our taxonomy (see Fig. 4). One may further divide fcntl64 calls of subtype fflags by the values F\_SETFL and F\_GETFL of the arq parameter. In the same way, we defined parents and children for each of the 54 call types and their parameters.

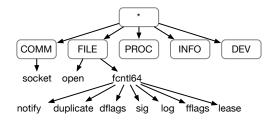


Fig. 4. Parts of the taxonomy we defined for the hierarchical variable type.

The *path* variable is filled whenever a system call accesses a file system path. Its hierarchy comes naturally along with the given path hierarchy of the file system. The *sequence* variable expresses the temporal order of calls within a process. The directly preceding call is the highest, less recent calls are at deeper levels of the hierarchy.

We collected system call data from eleven applications (like Firefox, Epiphany, NEdit, XEmacs) with the *strace* tool (version 4.5.17) under Ubuntu Linux (kernel 2.6.26, 32 bit). All child processes were monitored by using option -f of *strace*. For each application, we logged five times five minutes and five times ten minutes of system calls if they belonged to one of the 54 types shown in Tab. 12, resulting in a whole of 110 log files comprising about 23 million of lines (1.8 GB).

#### 3.3 Resulting Aggregation through Hierarchical Heavy Hitters

We have implemented the Full Ancestry and Partial Ancestry variants of the HHH algorithm mentioned in Section 3.1. The code was integrated into the RapidMiner data mining tool. Regarding run-time, all experiments were done on a machine with Intel Core 2 Duo E6300 processor with 2 GHz and 2 GB main memory.

		Memory			Run-time		Similarity		
		Min	Max	Avg	Min	Max	Avg	Avg	Dev
	Т	19	151	111	16	219	79	0.997	0.006
FA	TP	25	9,971	5,988	31	922	472	0.994	0.003
	TPS	736	73,403	48,820	78	14,422	6,569	0.987	0.008
	Т	7	105	70	15	219	74	0.985	0.010
PA	TP	7	4,671	2,837	31	5,109	2,328	0.957	0.017
	TPS	141	18,058	10,547	78	150,781	74,342	0.921	0.026

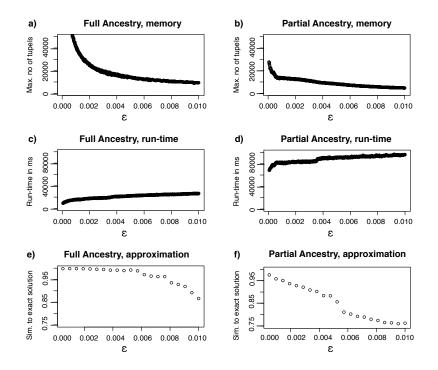
**Table 13.** Memory consumption (number of stored tupels), run-time (milliseconds) and similarity to exact solution of the Full Ancestry (FA) and Partial Ancestry (PA) algorithms ( $\varepsilon = 0.0005$ ,  $\phi = 0.002$ ). Minimum (Min), maximum (Max) and average (Avg) values were calculated over measurements for the first log file of all eleven applications with varying dimensionality of the element tupels (T = type hierarchy, P = path hierarchy, S = sequence hierarchy).

Since we want to aggregate system call data on devices that are severely limited in processing power and available memory, measuring the resource usage of our algorithms was of paramount importance. Table 13 shows the run-time and memory consumption of the Full Ancestry and Partial Ancestry algorithms using only the *type* hierarchy, the *type* and *path* hierarchy, or the *type*, *path*, and *sequence* hierarchy. Minimum, maximum and averages were calculated over a sample of the ten gathered log files for each of the eleven application by taking only the first log file for each application into account.

Memory consumption and run-time increase with the dimensionality of the elements, while at the same time approximation quality decreases. Quality is measured as similarity to the exact solution. Full Ancestry has a higher approximation quality in general. The results correspond to observations made by Cormode and are probably due to the fact that Partial Ancestry outputs bigger HHH sets, which was the case in our experiments, too. Note that approximation quality can always be increased by changing parameter  $\varepsilon$  to a smaller value at the expense of a longer run-time.

Even for three-dimensional elements, memory consumption is quite low regarding the number of stored tuples. The largest number of tuples (73,403), only equates to a few hundred kilobytes in main memory! The longest run-time of 150,781 ms for Partial Ancestry in three dimensions relates to the size of the biggest log file (application Rhythmbox).

Figure 5 shows the behaviour of our algorithms on the biggest log file (application Rhythmbox) for three dimensions with varying  $\varepsilon$  and constant  $\phi$ . Memory consumption and quality decrease with increasing  $\varepsilon$ , while the run-time increases. So the most important trade-off involved here is weighting memory consumption against approximation quality — the run-time is only linearly affected by parameter  $\varepsilon$ . Again, Full Ancestry shows a better approximation quality in general.



**Fig. 5.** Memory consumption (a, b), run-time (c, d) and similarity to exact solution (e, f) of HHH algorithms (three-dimensional) with varying  $\varepsilon$ ,  $\phi = 0.001$  on biggest log file of application Rhythmbox.

**Classification results** For the 110 log files of all applications, we determined the HHHs, resulting in sets of frequent tupels of hierarchical values. Interpreting each HHH set as an example of application behaviour, we wanted to answer the question if the profiles could be separated by a classifier. So we estimated the expected classification performance by a leave-one-out validation for kNN.

Therefore, we needed to define a distance measure for the profiles determined by HHH algorithms. The data structures of HHH algorithms contain a small subset of prefixes of stream elements. The estimated frequencies  $f_p$  are calculated from such data structure by the output method and compared to  $\phi$ , thereby generating a HHH set. The similarity measure DSM operates not on the HHH sets, but directly on the internal data structures  $D_1, D_2$  of two HHH algorithms:

$$\sin(D_1, D_2) = \frac{\sum_{p \in P_1 \cap P_2} \operatorname{contrib}_{DSM}(p)}{|P_1 \cup P_2|}$$

Be  $f_p^i$  the estimated frequency of prefix p for data structure  $D_i$  as normally calculated by the HHH output method. The contribution of individual prefixes to overall similarity can then be defined as

contrib<sub>DSM</sub>(p) = 
$$\frac{2 \cdot \min(f_p^1, f_p^2)}{\min(f_p^1, f_p^2) + \max(f_p^1, f_p^2)}$$
.

The so defined similarity measure is independent from the choice of  $\phi$ , as no HHH sets need to be calculated.

The classification errors for different values of k, hierarchies and distance measures are shown in Tab. 14. The new DSM distance measure which is independent of parameter  $\phi$  shows the lowest classification error in all validation experiments. As a baseline, we also determined the relative frequencies (TF, term frequencies) of call types per log file and classified them using kNN (with Euclidean distance). The error for profiling by HHH sets is significantly lower than for the baseline.

		т		$\mathbf{TS}$		
k	DSM	TF	DSM	TF		
3	10.3	17.0	7.7	17.0		
5	12.7	18.7	8.7	18.7		
7	14.0	21.7	8.7	21.7		
9	14.0	21.0	9.0	21.0		

**Table 14.** Results for kNN (k = 3, 5, 7, 9),  $\varepsilon = 0.0005, \phi = 0.002$  and distance measures DSM and TF, when only the *type* hierarchy or *type* and *sequence* hierarchy together are used.

# 4 Conclusion

Server-based and device-based learning has been investigated regarding resource constraints. Further experiments to measure resource consumption and prediction accuracy will be conducted on real mobile devices, like Android mobile phones, whose operating system is also based on the Linux kernel.

Aggregation using HHH worked successfully for the classification of applications. Further work will exploit HHH aggregation for other learning tasks and inspect other data streaming algorithms. Concerning server-based learning, we may now answer the questions from the introduction, whether structural models are demanded for the prediction of user behavior on the basis of system calls, or simpler models such as Naive Bayes suffice. Should the sequential nature of system calls be taken into account by the algorithm? Or is it sufficient to encode the sequences into the features? Or should features as well as algorithm be capable of explicitly addressing sequences? We have compared CRF and NB with respect to their model quality, memory consumption, and runtime. Neither encoding the sequence into features nor applying an algorithm which is made for sequential information (i.e., CRF) outperforms a simple model (i.e., NB).

This is in contrast with studies on intrusion detection, where it was shown advantageous to take into account the structure of system calls, utilizing Conditional Random Fields (CRF) [9] and special kernel functions to measure the similarity of sequences [23]. Structured models in terms of special tree kernel functions outperformed n-gram representations when detecting malicious SQL queries [1]. Possibly, for prefetching strategies, the temporal order of system calls is not as important as we expected it to be. In the near future the resulting improvements in terms of cache hit rate and file operation latencies will be evaluated systematically based on a cache simulator and by modifying the Linux kernel.

Given regular processors, CRF are only applicable in server-based learning. Possibly, the integration of special processors into devices and a massively parallel training algorithm could speed up CRF for device-based learning. Further work will implement CRF on a GPGPU (general purpose graphic processing unit). GPGPUs will soon be used by mobile devices. It has been shown that their energy efficiency is advantagous [24].

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