

# Bash Datalog: Answering Datalog Queries with Unix Shell Commands

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**Abstract.** Dealing with large tabular datasets often requires extensive preprocessing. This preprocessing happens only once, so that loading and indexing the data in a database or triple store may be an overkill. In this paper, we present an approach that allows preprocessing large tabular data in Datalog – without indexing the data. The Datalog query is translated to Unix Bash and can be executed in a shell. Our experiments show that, for the use case of data preprocessing, our approach is competitive with state-of-the-art systems in terms of scalability and speed, while at the same time requiring only a Bash shell on a Unix system.

## 1 Introduction

**Motivation.** Many data analytics tasks work on tabular data. Such data can take the form of relational tables or TAB-separated files. Even RDF knowledge bases can be seen abstractly as tabular data of a subject, a predicate, an object, and an optional graph id. Quite often, such data has to be preprocessed before the analysis can be made. We focus here on preprocessing in the form of select-project-join-union operations with recursion – removing superfluous columns, selecting rows of interest, recursively finding all instances of a class, etc. The defining characteristic of such a preprocessing step is that it is executed only once on the data in order to constitute the dataset of interest for the later analysis. This one-time preprocessing is the task that we are concerned with.

Databases or triple stores can obviously help. However, loading large amounts of data into these systems may take hours or even days (Wikidata [33], e.g., contains 267GB of data). Another possibility is to use systems such as DLV [18], Souffle [26], or RDFox [22], which work directly on the data. However, these systems load the data into memory. While this works well for small datasets, it does not work for larger ones (as we show in our experiments) Large-scale data processing systems such as BigDataLog [28], Flink [5], Dryad [15], or NoDB [3] can help. However, these require the installation of particular software, the knowledge of particular programming languages, or even a particular distributed infrastructure. Installing and getting to run such systems can take several hours. The user may not have the necessary knowledge and infrastructure to do this (think of a researcher in the Digital Humanities who wants to preprocess a file of census data; or of an engineer in a start-up who has to quickly join log files on a common column; or of a student who wants to extract a subgraph of Wikidata).

**Our Proposal.** In this paper, we propose a method to preprocess tabular data files without installing any particular software. We propose to express the preprocessing steps in Datalog [1]. For example, assume that there is a file `facts.tsv` that contains RDF facts in the form of TAB-separated subject-predicate-object triples. Assume that we want to recursively extract all places located in the United States. The Datalog program in our dialect would be:

```
fact(X, R, Y) :~ cat facts.tsv
locatedIn(X, Y) :- fact(X, "locatedIn", Y) .
locatedIn(X, Y) :- locatedIn(X, Z), fact(Z, "locatedIn", Y) .
main(X) :- locatedIn(X, "USA") .
```

This program prints the file `facts.tsv` into a predicate `fact`. The following two lines are the recursive definition of the `locatedIn` predicate. The `main` predicate is a predefined predicate that acts as the query. Our rationale for choosing Datalog is that it is a particularly simple language, which has just a single syntactic construction, and no reserved keywords. Yet, Datalog is expressive enough to deal with joins, unions, projections, selections, negation, and recursivity. In particular, it can deal with  $n$ -ary tables ( $n > 3$ ). If the user deals primarily with RDF data, our approach can also be used with N-Triples files as the A-Box, a subset of OWL 2 RL [21] as the T-Box, and SPARQL [14] for the query.

To execute the Datalog program, we propose to compile it automatically to Unix Bash Shell commands. We offer a Web page to this end: <https://www.thomasrebele.org/projects/bashlog>. The user can just enter the Datalog program, and click a button to obtain the following Bash code (simplified):

```
awk '$2 == "locatedIn" {print $1 "\t" $3}' facts.tsv > li.tmp
awk '$2 == "USA" {print $0}' li.tmp | tee full.tmp > delta.tmp
while
  join li.tmp delta.tmp | comm -23 - full.tmp > new.tmp
  mv new.tmp delta.tmp
  sort -m -o full.tmp full.tmp delta.tmp
  [ -s delta.tmp ];
do continue; done
cat full.tmp
```

The Bash code can be copy-pasted into a Unix Shell and run. Such a solution has several advantages. First, it does not require any software installation. It just requires a visit to a Web site. The resulting Bash code runs on any Unix-compatible system out of the box. Second, the Bash shell has been around for several decades, and the commands are not just tried and tested, but actually continuously developed. Modern implementations of the `sort` command, e.g., can split the input into several portions that fit into memory, and sort them individually. Finally, the Bash shell allows executing several processes in parallel, and their communication is managed by the operating system.

**Contribution.** We propose to compile a Datalog program automatically into Bash commands. Our method optimizes the Datalog program with relational algebra

optimization techniques, re-uses previously computed intermediate results, and produces a highly parallelized Shell script. For this purpose, our method employs pipes and process substitution. Our experiments on a variety of datasets and preprocessing tasks show that this method is competitive in terms of runtime with state-of-the-art database systems, Datalog engines, and triple stores.

We start with a discussion of related work in Section 2. Section 3 introduces preliminaries. Section 4 presents our approach, and Section 5 evaluates it.

## 2 Related Work

**Data Processing Systems.** Relational Databases such as Oracle, IBM DB2, Postgres, MySQL, MonetDB [4] and NoDB [3] can handle tabular data of arbitrary form, while the triple stores such as OpenLink Virtuoso [8], Stardog, and Jena [6] target RDF data. HDT [11] is a binary format for RDF, which can be used with Jena. RDFSlice [20] can preprocess RDF datasets. Reasoners such as Pellet [23], Hermit [27], RACER [13], Fact++ [30] and Jena [6] can perform OWL reasoning on RDF data. Datalog systems such as DLV [18], Souffle [26], BigDatalog [28], RDFox [22] and others [16,34,35] can efficiently evaluate Datalog queries on large data. Distributed Data Processing systems such as Dryad [15], Apache Tez [25], SCOPE [37], Impala [9], Apache Spark [36], and Apache Flink [5] provide advanced features such as support for SQL or streams.

All of these systems can be used for preprocessing data. However, the vast majority of these systems require the installation of software. The parallelized systems also require a distributed infrastructure. Our approach, in contrast, requires none of these. It just requires a visit to a Web page. The Bash script that we produce runs in a common shell console without any further prerequisites. Interestingly, our approach still delivers comparable performance to the state of the art, as we shall see in the experiments.

Only very few systems do not require a software installation beyond downloading a file (e.g., RDFSlice [20], Stardog, and DLV [18]). Yet, as we shall see in the experiments, these systems do not scale well to large datasets.

**Other Work.** Linked Data Fragments [32] aim to strike a balance between downloading an RDF data dump and querying it on a server. The method thus addresses a slightly different problem from ours. AI planning with softbots [10] aims to answer queries on an incomplete and evolving database. In our problem setting, however, we have access to all the information. NoSQL Databases such as Cassandra, HBase, and Google’s BigTable [7] target non-tabular data. Our method, in contrast, aims at tabular data.

## 3 Preliminaries

**Datalog.** A Datalog rule with negation [1] takes the form

$$H :- B_1, \dots, B_n, \neg N_1, \dots, \neg N_m.$$

Here,  $H$  is the head atom,  $B_1, \dots, B_n$  are the positive body atoms, and  $N_1, \dots, N_m$  are the negated body atoms. Each atom is of the form  $r(x_1, \dots, x_k)$ , where  $r$  is a relation name and  $x_1, \dots, x_k \in \mathcal{V} \cup \mathcal{C}$ , where  $\mathcal{V}$  is a set of variables and  $\mathcal{C}$  is a set of constants. Intuitively, such a rule says that  $H$  holds if  $B_1, \dots, B_n$  and none of the  $N_1, \dots, N_m$  holds. We consider only *safe rules*, i.e., each variable in the head or in a negated atom must also appear in a positive body atom. A *Datalog program* is a set of Datalog rules. A set  $M$  of atoms is a *model* of a program  $P$ , if the following holds:  $M$  contains an atom  $a$  iff  $P$  contains a rule  $H :- B_1, \dots, B_n, \neg N_1, \dots, \neg N_m$ , such that there exists a substitution  $\sigma : \mathcal{V} \rightarrow \mathcal{C}$  with  $\sigma(B_i) \in M$  for  $i \in \{1, \dots, n\}$  and  $\sigma(N_i) \notin M$  for  $i \in \{1, \dots, m\}$  and  $a = \sigma(H)$ . We consider only *stratified* Datalog programs [1], which entails that there exists a unique minimal model.

OWL RL [21] is a subset of the OWL ontology language. Since every OWL RL ontology can be translated to Datalog [22], we deal with Datalog as the more general case in all of the following.

**Relational Algebra.** *Relational algebra* [1] provides the semantics of relational database operations on tables. For our purposes, we use the unnamed relational algebra with the operators select  $\sigma$ , project  $\pi$ , join  $\bowtie$ , anti-join  $\triangleright$ , and union  $\cup$  (see [24] for their definitions). We also use the *least fixed point (LFP) operator*  $\mu$  [2]. For a function  $f$  from a table to a table,  $\mu_x(f(x))$  is the least fix point of  $f$  for the  $\subseteq$  relation. With this, our algebra has the same expressivity as safe stratified Datalog programs, i.e., the translation of safe Datalog with stratified negations to relational algebra is sound and complete [1].

*Example 2 (Relational Algebra):* The following expression computes the transitive closure of a two-column table *subclass*:

$$\mu_x(\text{subclass} \cup \pi_{1,4}(x \bowtie_{2=1} x))$$

$\mu$  computes the least fix point of a function. Here, the function is given by the argument of the  $\mu$ -operator. To compute the least fix point, we execute the function first with the empty table,  $x = \emptyset$ . Then the function returns the *subclass* table. Then we execute the function again on this result. This time, the function will join *subclass* with itself, project the resulting table on the first and last column, and add in the original *subclass* table. We repeat this process until no more changes occur.

**Unix.** Unix is a family of multitasking computer operating systems, which are widely used on servers, smartphones, and desktop computers. One of the characteristics of Unix is that “Everything is a file”, which means that files, pipes, the standard output, the standard input, and other resources can all be seen as streams of bytes. Here, we are interested only in *TAB-separated* byte streams, i.e., streams that consist of several *rows* (sequences of bytes separated by a new-line character), which each consist of the same number of *columns* (sequences of bytes separated by a tabulator character).

The Bourne-again shell (Bash) is a command-line interface for Unix-like operating systems. A *Bash command* is either a built-in keyword, or a small pro-

gram. We will only use commands of the POSIX standard. One of them is the `awk` command, which we use as follows:

```
awk -F$'\t' 'p' b
```

This command executes the program  $p$  on the byte stream  $b$ , using the TAB as a separator for  $b$ . We will discuss different programs  $p$  later in this paper.

**Pipes.** When a command is executed, it becomes a *process*. Two processes can communicate through a pipe, i.e., a byte stream that is filled by one process, and read by the other one. If the producing process is faster than the receiving one, the pipe buffers the stream, or blocks the producing process if necessary. In Bash, a pipe between process  $p_1$  and process  $p_2$  pipes can be constructed by stating  $p_1 \mid p_2$ . A pipe can also be constructed “on the fly” by a so-called *process substitution*, as follows:  $p_1 < ( p_2 )$ . This construction pipes the output of  $p_2$  into the first argument of  $p_1$ .

## 4 Approach

### 4.1 Our Datalog Dialect

In our Datalog dialect, predicates are alphanumeric strings that start with a lowercase character. Variables start with an uppercase letter. Constants are strings enclosed by double quotes. Constants may not contain quotation marks, TAB characters, or newline characters. For our purposes, the Datalog program has to refer to files or byte streams of data. For this reason, we introduce an additional type of rules, which we call *command rules*. A command rule takes the following form:

$$p(x_1, \dots, x_n) \text{ :}\sim c$$

Here,  $p$  is a predicate,  $x_1, \dots, x_n$  are variables, and  $c$  is a Bash command. Semantically, this rule means that executing  $c$  produces a TAB-separated byte stream of  $n$  columns, which will be referred to by the predicate  $p$  in the Datalog program. In the simplest case, the command  $c$  just prints a file, as in `cat facts.tsv`. However, the command can also be any other Bash command, such as `ls -l`.

Our goal is to compute a certain output with the Datalog program. This output is designated by the distinguished head predicate `main`. An *answer* of the program is a grounded variant of the head atom of this rule that appears in the minimal model of the program. See again Figure 1 for an example of a Datalog program in our dialect. Our dialect is a generalization of standard Datalog, so that a normal Datalog program can be run directly in our system.

Our approach can also work in “RDF mode”. In that mode, the input consists of an OWL 2 RL [21] ontology, a SPARQL [14] query, and an N-Triples file  $F$ . We build a `main` predicate for the SPARQL query, and we use a small AWK program that converts  $F$  into a TAB-separated byte stream (see our technical report [24] for details). Much like in RDFox [22], we convert the OWL ontology into Datalog rules (see again [24]). For now, we support only a subset of OWL 2 RL: Like RDFox [22], we assume that all classes and properties axioms are provided by

the ontology and will not be queried by the SPARQL query. We also do not yet support OWL axioms related to literals. Our SPARQL implementation supports basics graph patterns, property paths without negations, OPTIONAL, UNION and MINUS.

## 4.2 Loading Datalog

Next, we build a relational algebra expression for the `main` predicate of the Datalog program. Our algorithm is similar to existing approaches [31]. Algorithm 1 takes as input a predicate  $p$ , a cache, and a Datalog program  $P$ . The algorithm is initially called with  $p=\text{main}$ ,  $\text{cache}=\emptyset$ , and the Datalog program that we want to translate. The cache stores already computed relational algebra plans. This allows us to re-use the same sub-plan multiple times in the final plan, thus the algorithm builds a directed acyclic graph (DAG) instead of a tree.

Our algorithm first checks whether  $p$  appears in the cache. In that case,  $p$  is currently being computed in a previous recursive call of the method, and the algorithm returns a variable  $x$  indexed by  $p$ . This is the variable for which we compute the least fix point.

Then, the algorithm traverses all rules with  $p$  in the head. For every rule, the algorithm recursively retrieves the plan for the body atoms. The algorithm (anti-)joins the sub-plans, adding selections  $\sigma_{j=k}$  if necessary. Finally, it puts the resulting formula into a project-node that extracts the relevant columns.

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### Algorithm 1: Translation from Datalog to relational algebra

---

```

1 fn mapPred ( $p$ , cache,  $P$ ) is
2   if  $p \in \text{cache}$  then return  $x_p$  ;
3    $\text{plan} \leftarrow \emptyset$ ;  $\text{newCache} \leftarrow \text{cache} \cup \{p\}$ 
4   foreach  $p(H_1, \dots, H_{n_h}) :- r_1(X_1^1, \dots, X_{n_1}^1), \dots, \neg q_1(Y_1^1, \dots, Y_{m_1}^1), \dots$  in  $P$  do
5      $\text{bodyPlan} \leftarrow \{\}$ 
6     foreach  $r_i(X_1^i, \dots, X_{n_i}^i)$  do
7        $\text{atomPlan} \leftarrow \text{mapPred}(r_i, \text{newCache}, P)$ 
8       foreach  $(X_j^i, X_k^i) \mid X_j^i = X_k^i, j \neq k$  do
9          $\text{atomPlan} \leftarrow \sigma_{X_j^i = X_k^i}(\text{atomPlan})$ 
10       $\text{bodyPlan} \leftarrow \text{bodyPlan} \bowtie \text{atomPlan}$ 
11     foreach  $\neg q_i(Y_1^i, \dots, Y_{m_i}^i)$  do
12        $\text{atomPlan} \leftarrow \text{mapPred}(q_i, \emptyset, P)$ 
13       foreach  $(Y_j^i, Y_k^i) \mid Y_j^i = Y_k^i, j \neq k$  do
14          $\text{atomPlan} \leftarrow \sigma_{Y_j^i = Y_k^i}(\text{atomPlan})$ 
15       $\text{bodyPlan} \leftarrow \text{bodyPlan} \triangleright \text{atomPlan}$ 
16      $\text{plan} \leftarrow \text{plan} \cup \pi_{H_1, \dots, H_{n_h}}(\text{bodyPlan})$ 
17 foreach  $\text{rule } p(H_1, \dots, H_{n_h}) \sim c$  in  $P$  do  $\text{plan} \leftarrow \text{plan} \cup \pi_{H_1, \dots, H_{n_h}}(c)$  ;
18 return  $\mu_{x_p}(\text{plan})$ 

```

---

*Example 3 (Datalog Translation):* Assume that there is a two-column TAB-separated file `subclass.tsv`, which contains each class with its subclasses. Consider the following Datalog program  $P$ :

- (1) `directSubclass(x,y) :~ cat subclass.tsv`
- (2) `main(x,y) :- directSubclass(x,y).`
- (3) `main(x,z) :- directSubclass(x,y), main(y,z).`

Our algorithm will go through all rules with the head predicate `main`. These are Rule 2 and Rule 3. For Rule 2, the algorithm will recursively call itself and return  $\mu_{x_{\text{directSubclass}}}(\emptyset \cup [\text{cat subclass.tsv}])$ . Since the argument of  $\mu$  does not contain the variable  $x_{\text{directSubclass}}$ , this is equivalent to `[cat subclass.tsv]`. For the first body atom in Rule 3, the algorithm returns `[cat subclass.tsv]` just like before. For the second body atom, the algorithm returns  $x_{\text{main}}$ , because `main` is in the cache. Thus, Rule 3 yields  $\pi_{1,4}([\text{cat subclass.tsv}] \bowtie_{2=1} x_{\text{main}})$ . Finally, the algorithm constructs

$$\mu_{x_{\text{main}}}([\text{cat subclass.tsv}] \cup \pi_{1,4}([\text{cat subclass.tsv}] \bowtie_{2=1} x_{\text{main}}))$$

### 4.3 Producing Bash Commands

The previous step has translated the input Datalog program to a relational algebra expression. Now, we translate this expression to a Bash command by the function  $b$ , which is defined as follows:

$$b([c]) = c$$

An expression of the form  $[c]$  is already a Bash command, and hence we can return directly  $c$ .

$$b(e_1 \cup \dots \cup e_n)$$

We translate a union into a sort command that removes duplicates:

$$\text{sort -u } \langle(b(e_1)) \dots \langle(b(e_n))$$

$$b(e_1 \bowtie_{x=y} e_2)$$

A join of two expressions  $e_1$  and  $e_2$  on a single variable at position  $x$  and  $y$ , respectively, gives rise to the command

$$\begin{aligned} &\text{join -t\$'\t' -1x -2y} \\ &\quad \langle(\text{sort -t\$'\t' -kx } \langle(b(e_1))) \\ &\quad \langle(\text{sort -t\$'\t' -ky } \langle(b(e_2))) \end{aligned}$$

This command sorts the byte streams of  $b(e_1)$  and  $b(e_2)$ , and then joins them on the common column.

$$b(e_1 \bowtie_{x=y, \dots} e_2)$$

The Bash `join` command can perform the join on only one column. If we want to join on several columns, we have to add a new column to each of the byte streams. This new column concatenates the join columns into a single column. This can be achieved with the following AWK program, which we run on both  $b(e_1)$  and  $b(e_2)$ :

$$\{ \text{print } \$0 \text{ FS } \$j_1 \text{ s } \$j_2 \text{ s } \dots \text{ s } \$j_n \}$$

Here, the indices  $j_1, \dots, j_n$  are the positions of the join columns in the input byte stream, FS is the field separator, and  $s$  is a special separation character (we use ASCII character 2, but any other one can be used as well). Once we have done this with both byte streams, we can join them on this new column as described above. This join will also remove the additional column.

$b(e_1 \triangleright_x e_2)$

Just as a regular join, an anti-join becomes a `join` command. We use the parameter `-v1`, so that the command outputs only those tuples emerging from  $e_1$  than cannot be joined with those from  $e_2$ . We deal with anti-joins on multiple columns in the same way as with multi-column joins.

$b(\pi_{i_1, \dots, i_n}(e))$

A projection becomes the following AWK program, which extracts the given columns from the input byte stream  $b(e)$ :

```
{ print $i_1 FS ... FS $i_n }
```

$b(\pi_{i,a}(e))$

A constant introduction becomes the following AWK program, which produces a TAB-separated byte stream that inserts the constant in column  $i$  of the input byte stream  $b(e)$ :

```
{ print $1 FS ... $(i-1) FS a FS $i FS ... $n }
```

$b(\sigma_{i=v}(e))$

A selection node gives rise to the following AWK program, which selects the corresponding rows from the input byte stream  $b(e)$ :

```
 $i$ =="v" { print $0 }
```

This command can be generalized easily to a selection on several columns.

Several of these translations produce process substitutions. In such cases, Bash starts the parent process and the inner process in parallel. The parent process will block while it cannot read from the inner processes. Thus, only the innermost processes run in the beginning. Every process is run asynchronously as soon as input and CPU capacity is available. Thus, our Bash program is not subject to the forced synchronization that appears in Map-Reduce systems.

#### 4.4 Recursion

We have just defined the function  $b$  that translates a relational algebra expression to a Bash command. We will now see how to define  $b$  for the case of recursion. A node  $\mu_x(f(x))$  becomes

```
echo -n > delta.tmp; echo -n > full.tmp
while
  sort b(f(delta.tmp)) | comm -23 - full.tmp > new.tmp;
  mv new.tmp delta.tmp;
  sort -u -m -o full.tmp full.tmp <(sort delta.tmp);
  [ -s delta.tmp ];
do continue; done
cat full.tmp
```



This code uses 3 temporary files to compute the least fix point of  $f$ : `full.tmp` contains all facts inferred until the current iteration. `delta.tmp` contains newly added facts of an iteration. `new.tmp` is used as swap file. The code first creates `delta.tmp` and `full.tmp` as empty files. It then runs  $f$  on the delta file. The `comm` command compares the sorted outcome of  $f$  to the (initially empty) file `full.tmp`, and writes the new lines to the file `new.tmp`. This file is then renamed to `delta.tmp`. This procedure updates the file `delta.tmp` to contain the newly added facts. The `comm` command cannot write directly to `delta.tmp`, because this file also serves as input to the command produced by  $b(f(\text{delta.tmp}))$ .

The following `sort` command merges the new lines into `full.tmp`, and writes the output to `full.tmp` (unlike the `comm` command, the `sort` command can write to a file that also serves as input). Now, all facts generated in this iteration have been added to `full.tmp`. The `[...]` part of the code lets the loop run while the file `delta.tmp` is not empty, i.e., while new lines are still being added. If no new lines were added, the code quits the loop, and prints all facts. Note that, due to the monotonicity of our relational algebra operators, and due to the stratification of our programs, we can afford to run  $f$  only on the newly added lines.

#### 4.5 Materialization

**Materialization nodes.** To avoid re-computing an algebra expression that has already been computed, we introduce a new type of operator to the algebra, the materialization node. A materialization node  $\square(m, (\lambda y : p))$  has two sub-plans:  $m$  is the plan that is used multiple times, and that we will materialize. The lambda function  $(\lambda y : p)$  is the main plan, and takes the materialized plan as parameter. The variable  $y$  replaces all occurrences of  $m$  in the original plan (see [24] for details).

**Bash translation.** The translation to Bash is shown on the right.  $b$  is the Bash translation function defined in Section 4.3.  $t$  is a temporary file name. The code first creates a named pipe called `lock.t`. Commands that use  $t$  have to wait until  $b(m)$  finishes. We ensure this by making these commands read from `lock.t`. Since this pipe contains no data, the commands block. When  $b(m)$  finishes, the two `exec` commands close the named pipe, thus unblocking the commands that need  $t$ . There can be a rare race condition:  $b(m)$  may finish before any process that listens on the pipe was started. In that case, the two `exec` commands will try to close a pipe that has no listeners. In such cases, the `exec` command will block. We solve this problem by reading from the pipe with a `cat` command that runs in the background. This way, the pipe has at least one listener, and the `exec` commands will close the pipe. This, however, brings a second problem: If the processes that listen on the pipe were still not started, they will try to listen to a closed pipe. To avoid this problem, we rename the pipe from `lock.t` to `done.t`. Such a renaming does not affect any processes that

```
mkfifo lock_t
(
  b(m) > t
  mv lock_t done_t
  cat done_t &
  exec 3> done_t
  exec 3>&-
) &
by→t(p)
rm t
```

already listen on the pipe, but it will prevent any new processes from listening on the pipe under the old name.

$b_{y \rightarrow t}$  extends  $b$  as follows:  $b_{y \rightarrow t}(y)$  generates the bash code `cat  $t$` , and all plan nodes  $p_i$  that have a child  $y$  generate the bash code

```
cat lock_ $t$  2> /dev/null ; b( $p_i$ )
```

As explained above, the `cat` command blocks the execution until  $t$  is materialized. The part “`2> /dev/null`” removes the error message in case `cat` is executed when the pipe was already renamed.

## 4.6 Optimization

**Query Optimizations.** We apply the usual optimizations on our relational algebra expressions: we push selection nodes as close to the source as possible; we merge unions; we merge projects; we apply a simple join re-ordering. Additionally, we remove an occurrence of a LFP variable, if it cannot contribute new facts; we remove the LFP when there is no recursion; and we extract from a LFP node the non-recursive part of the inner plan (so that it is computed only once at the beginning of the fixed point computation). We also optimize fix point computations in the same way as in the semi-naive Datalog evaluation [1] (see again [24] for details).

**Bash Optimizations.** We collect different AWK commands that select or project on the same file into a single AWK command. This command runs only once through the file, and writes out all selections and projections into several files, one for each original AWK command. We replace multiple comparisons with constants on the same columns by a hash table lookup. We detect nested `sort` commands, and remove redundant ones. We run `sort -u` on the final output to make all results unique. We estimate the number of concurrently running sort commands, and assign each of them an equal amount of memory, if the buffer size parameter is available. Finally, we force all commands to use the same character set and sort order by adding the command `export LC_ALL=C` to our program.

## 5 Experiments

We ran our method on several datasets, and compared it to several competitors. All our experiments were run on a laptop with Ubuntu 16.04, an Intel Core i7-4610M 3.00 GHz CPU, 16 GB of RAM, 16 GB of swap space, and 3.8 TB of hard disk space. We used GNU Bash 4.3.48, mawk 1.3.3 for AWK, and GNU coreutils 8.25 for the other POSIX commands.

We emphasize that our goal is not to be faster than each and every system that currently exists. For this, the corpus of related work is simply too large (see Section 2). This is also not the purpose of Bash Datalog. The purpose of Bash Datalog is to provide a preprocessing tool that runs without installing any additional software besides a Bash shell. This is an advantage that no competing approach offers. Our experiments then serve mainly to show that our approach is generally comparable in terms of scalability with the state of the art.

Bash	DLV	Souffle	RDFox	Jena TDB	Jena HDT	Stardog	Virtuoso	Postgres*		NoDB*	MonetDB*		RDF- Slice*
								-I	+I		-I	+I	
<b>0.7</b>	9.6	7.8	2.2	25.7	26.4	12.8	11.7	4.8	27.5	>600	1.8	2.6	12.6
<b>1.3</b>	9.3	119.4	2.2	281.3	>600	13.6	11.8	-	-	-	-	-	-
<b>0.9</b>	9.2	8.8	2.2	26.7	27.0	12.7	11.5	7.8	30.5	292.9	1.9	2.7	-
<b>1.9</b>	9.3	11.9	2.2	>600	>600	13.2	12.2	14.7	37.4	>600	2.3	3.1	-
<b>1.4</b>	9.3	10.3	2.2	>600	>600	12.9	-	28.9	51.6	-	2.2	3.0	-
<b>1.9</b>	9.4	11.1	2.4	>600	>600	17.6	-	21.2	43.9	-	3.9	4.7	-
2.4	9.5	56.3	<b>2.2</b>	>600	>600	13.4	-	21.6	44.3	>600	3.0	3.9	-
2.5	9.3	12.9	<b>2.3</b>	>600	>600	15.3	-	-	-	-	-	-	-
3.1	9.4	>600	<b>2.3</b>	>600	>600	13.4	-	71.1	93.8	>600	25.5	26.8	-
<b>2.0</b>	9.3	11.6	2.2	>600	>600	13.5	-	23.0	45.7	>600	5.8	7.1	-
<b>0.9</b>	9.3	7.8	2.2	25.3	35.7	13.0	11.8	-	-	-	-	-	-
<b>1.4</b>	9.2	10.9	2.2	>600	>600	13.1	-	-	-	-	-	-	-
<b>1.4</b>	9.2	10.1	2.2	>600	>600	12.9	-	8.4	31.1	>600	4.3	5.4	-
<b>0.8</b>	9.5	6.7	2.3	34.5	24.8	13.5	12.0	4.8	27.5	19.1	1.9	2.7	3.7
of which loading:				16.8	7.4	11.0	5.9	4.4	24.3		1.7	2.5	

**Table 1.** Runtime for the 14 LUBM queries with 10 universities (155 MB), in seconds.

\* = no support for querying with a T-Box. We folded the T-Box into the query.

+/-I = with/without indexes. A dash means that the query is not supported.

## 5.1 Lehigh University Benchmark

**Setting.** The Lehigh University Benchmark (LUBM) [12] is a standard dataset for Semantic Web repositories, which models universities. It is parameterized by the number of universities, and hence its size can be varied. LUBM comes with 14 queries, which are expressed in SPARQL. We compare our approach to Stardog<sup>1</sup>, Virtuoso [8], RDFSlice [20], Jena [6], and Jena with the binary triple format HDT [11]. For RDFox [22], Souffle [26], DLV [18], we translated the queries to Datalog in the same way that we translate the queries to Datalog for our own system (Section 4.2). For NoDB [3], Postgres<sup>2</sup>, and MonetDB [4], we translated the Datalog queries first to an algebra expression, and then to SQL. In this process, we applied the relational algebra optimizations of Section 4.6. In this way, the T-Box of the LUBM queries is folded into the SQL query. Not all systems support all types of queries. MonetDB does not support recursive SQL queries. Postgres supports only certain types of recursive queries [24]. The same applies to NoDB. Virtuoso currently does not support intersections. RDFSlice aims at the slightly different problem of RDF-Slicing. It supports only a specific type of join. Also, it does not support recursion.

We ran every competitor on all queries that it supports, and averaged the runtime over 3 runs. We checked whether the query results were correct.

**LUBM10.** Table 1 shows the runtimes of all queries on LUBM with 10 universities. The runtimes include the loading and indexing times. For systems where

<sup>1</sup> <https://www.stardog.com/>, v. 5.2.0

<sup>2</sup> <https://www.postgresql.org/>, v. 10.1

we could determine these times explicitly, we noted them in the last row of the table. Since most systems finished in a matter of seconds, we aborted systems that took longer than 10 minutes. Among the 4 triple stores (Jena+TDB, Jena+HDT, Stardog, and Virtuoso), only Stardog can finish on all queries in less than 10 minutes. RDFSlice can answer only 2 queries, and runs a bit faster than Stardog. The 5 database competitors (Postgres, NoDB, and MonetDB – with and without indexes) are generally faster. Among these, MonetDB is much faster than Postgres and NoDB. Postgres and MonetDB are fastest without indexes, which is to be expected when running the query only once. Among the best performing systems are two Datalog systems (Bash Datalog, and RDFox). RDFox shines with a very short and nearly constant time for all queries. We suspect that this time is given by the loading time of the data, and that it dominates the answer computation time. Nevertheless, Bash Datalog is faster than RDFox on nearly all queries on LUBM 10.

Query	LUBM 500 (7.8 GB)							LUBM 1000 (16 GB)					
	Bash	RDFox	Stardog	Virtuoso	MonetDB (no indices)	MonetDB (indices)	RDFSlice	Bash	RDFox	Stardog	MonetDB (no indices)	MonetDB (indices)	RDFSlice
1	<b>27</b>	131	582	1577	83	97	229	<b>75</b>	273	1955	185	210	1042
2	<b>53</b>	132	683	1580				<b>118</b>	278	2030			
3	<b>35</b>	131	609	1578	88	101		<b>89</b>	276	1955	186	217	
4	<b>95</b>	129	583	1579	118	131		307	<b>273</b>	1962	522	471	
5	<b>62</b>	131	498		290	364		<b>168</b>	278	1956	894	793	
6	<b>93</b>	137	1011		866	797		354	<b>287</b>	2361	2066	1934	
7	<b>122</b>	134	673		898	753		544	<b>279</b>	2005	1809	2016	
8	151	<b>132</b>	768					447	<b>274</b>	1967			
9	250	<b>136</b>	749		2669	3064		712	<b>283</b>	2018	3275	3090	
10	<b>95</b>	132	678		587	491		334	<b>277</b>	1959	1845	1834	
11	<b>28</b>	130	498	1576				<b>64</b>	273	1957			
12	<b>56</b>	130	682					<b>164</b>	273	1959			
13	<b>63</b>	132	669		312	287		<b>174</b>	277	1969	908	955	
14	<b>28</b>	136	787	1595	85	99	74	<b>63</b>	284	2069	181	217	334
of which load:		489	1575	72	92					1946	160	194	

**Table 2.** Runtime for the LUBM queries, in seconds.

**LUBM500 and LUBM1000.** For the larger LUBM datasets, we chose the fastest systems in each group as competitors: RDFox for the Datalog systems, Stardog and Virtuoso for the triple stores, MonetDB for the databases, and RDFSlice as its own group. Table 2 shows the sizes of the datasets and the runtimes of the systems. Our system performs best on more than half of the queries. The only system that can achieve a similar performance is RDFox. As before, RDFox always needs just a constant time to answer a query, because it loads the dataset into main memory. This makes the system very fast. However, this technique will not work if the dataset is too large, as we shall see next.

## 5.2 Reachability

**Setting.** Our next datasets are the social networks LiveJournal [19], com-orkut [19], and friendster [17]. Table 3 shows the sizes of these datasets. We used a single query, which asks for all nodes that are reachable from a given node. As competitors, we chose again RDFox, Stardog, and Virtuoso. We could not use MonetDB or RDFSlice, because the reachability query is recursive. As an additional competitor, we chose BigDatalog [28]. This system was already run on the same LiveJournal and com-orkut graphs in the original paper [28]. We chose 3 random nodes (and thus generated 3 queries) for LiveJournal and com-orkut. We chose one random node for Friendster.

**Results.** Table 3 shows the average runtime for each system. Virtuoso was the slowest system, and we aborted it after 25min and 50min, respectively. We did not run it on the Friendster dataset, because Friendster is 20 times larger than the other two datasets. Stardog performs better. Still, we had to abort it after 10 hours on the Friendster dataset. BigDatalog performs well, but fails with an out of space error on the Friendster dataset. The fastest system is RDFox. This is because it can load the entire data into memory. This approach, however, fails with the Friendster dataset. It does not fit into memory, and RDFox is unable to run. Bash Datalog runs 50% slower than RDFox. In return, it is the only system that can finish in reasonable time on the Friendster dataset (4:32h).

dataset	Nodes	Edges	Bash	RDFox	BigDatalog	Stardog	Virtuoso
LiveJournal	4.8 M	69 M	117	<b>70</b>	532	941	>1500
orkut	3.1 M	117 M	225	<b>121</b>	1838	1123	>3000
friendster	68 M	2 586 M	<b>16306</b>	OOM	OOS	>36000	

**Table 3.** Runtime for the reachability query, in seconds (OOM=Out of memory; OOS=Out of space).

## 5.3 YAGO and Wikidata

**Setting.** Our final experiments concern the knowledge bases YAGO [29] and Wikidata [33]. The YAGO data comes in 3 different files, one with the 12 M facts (814 MB), one with the taxonomy with 1.8 M facts (154 MB), and one with the 24 M type relations (1.6 GB in size). Wikidata is a single file of 267 GB with 2.1 B triples. We designed 4 queries that are typical for such datasets (Table 1), together with a T-Box (Table 2).

**Results.** Table 4 shows the results of RDFox and our system on both datasets. On YAGO, RDFox is much slower than our system, because it needs to instantiate all rules in order to answer queries. On Wikidata, the data does not fit into main memory, and hence RDFox cannot run at all. Our system, in contrast, scales to the larger sizes of the data. One may think that a database system such as Postgres may be better adapted for such large datasets. This is, however, not the case. Postgres took 104 seconds to load the YAGO dataset, and 190 seconds

```

query1(X) :- subclassOf(X, "person") .
query1(X) :- subclassOf(X, Y), query1(Y) .
query2(X) :- hasParent(X, "Louis XIV") .
query3(X) :- hasAncestor(X, "Louis XIV") .
query4(X) :- hasBirthPlace(X, Y), isLocatedIn(Y, "Andorra") .

```

Fig. 1. Knowledge Base queries

```

hasParent(X,Y) :- hasChild(Y,X) .
hasAncestor(X,Y) :- hasParent(X,Y) .
hasAncestor(X,Z) :- hasAncestor(X,Y), hasParent(Y,Z) .
isLocatedIn(X,Y) :- containsLocation(Y,X) .
containsLocation(X,Y) :- isLocatedIn(Y,X) .
isLocatedIn(X,Y) :- isLocatedIn(X,Y), isLocatedIn(Y,Z) .

```

Fig. 2. Knowledge Base rules

to build the indexes. In this time, our system has already answered nearly all the queries.

**Discussion.** All of our experiments evaluate only the setting that we consider in this paper, namely the setting where the user wants to execute a single query in order to preprocess the data. Our experiments show that Bash Datalog can preprocess tabular data without the need to install any particular software.

Our approach has some limitations. For example, we could not implement a disk-based hash-join efficiently in Bash commands. Another limitation is the heuristic join reordering. It sometimes introduces large intermediate results, resulting in a less efficient query execution.

Overall, however, our approach is competitive in both speed and scalability to the state of the art. We attribute this to the highly optimized POSIX commands, and to our optimizations described in Section 4.6. Furthermore, the startup cost of our system is quite low, as it consists mainly of translating the query to a Bash script.

query	YAGO		Wikidata	
	Bash	RDFox	Bash	RDFox
1	<b>8</b>	483	<b>2259</b>	OOM
2	<b>5</b>	483	<b>2254</b>	OOM
3	<b>293</b>	483	<b>10171</b>	OOM
4	<b>5</b>	481	<b>2270</b>	OOM

**Table 4.** Runtime for the Wikidata/YAGO benchmark in seconds. (OOM = out of memory error)

## 6 Conclusion

In this paper, we have presented a method to compile Datalog programs into Unix Bash programs. This allows executing Datalog queries on tabular datasets without installing any software. We show that our method is competitive in terms of speed with state-of-the-art systems. Our system can be used online at <https://www.thomasrebele.org/projects/bashlog>. The Web interface also provides an API which allows to translate Datalog to Bash via an HTTP request. The source code is available at <https://github.com/thomasrebele/bashlog>. For future work, we aim to explore extensions of this work such as adding support of numerical comparisons to the Datalog language.

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