

# SimpleRA

*A laughably-minimalist, integer-only Relational Database Management System that makes the author question why they ever bothered to write it up!*

***Data Systems, Monsoon 2021***

17th August, 2021

Tuesday

# Important Features

- Relational Algebra Operators
- Integers Only
- No nested queries
- No transaction management
- Single thread programming only
- No identifiers should have spaces in them

# Commands

There are 2 kinds of commands in this database.

- Assignment statements
- Non-assignment statements

*Note: Not all operators have been implemented, some have been omitted for you to implement in later phases*

# Non-Assignment Statements

Non-assignment statements do create a new table (except load which just loads an existing table) in the process

- LOAD
- LIST
- PRINT
- RENAME

# Non-assignment Statements

- EXPORT
- CLEAR
- QUIT

# LOAD

Syntax:

```
LOAD <table_name>
```

- To successfully load a table, there should be a csv file names <table\_name>.csv consisiting of comma-seperated integers in the data folder
- None of the columns in the data file should have the same name
- every cell in the table should have a value

Run: `LOAD A`

# LIST TABLES

## Syntax

```
LIST TABLES
```

This command lists all tables that have been loaded or created using assignment statements

Run: `LIST TABLES`

Run: `LOAD B, LIST TABLES`

# PRINT

## Syntax

```
PRINT <table_name>
```

- Displays the first PRINT\_COUNT (global variable) rows of the table.
- Less number of rows can be printed if the table has only a few rows

Run: `PRINT B`



# EXPORT

## Syntax

```
EXPORT <table_name>
```

- All changes made and new tables created, exist only within the system and will be deleted once execution ends (temp file)
- To keep changes made (RENAME and new tables), you have to export the table (data)

Run: `EXPORT B`

# QUIT

## Syntax

```
QUIT
```

- Clear all tables present in the system (***WITHOUT EXPORTING THEM***) (temp file - empty)

Run: 

```
QUIT
```

# Assignment Statements

- All assignment statements lead to the creation of a new table.
- Every statement is of the form

```
<new_table_name> <- <assignment_statement>
```
- Naturally in all cases, <new\_table\_name> shouldn't already exist in the system

# Assignment Statements

- CROSS
- PROJECTION
- SELECTION

The following haven't been implemented

- JOIN
- SORT
- GROUP\_BY

# SELECTION

## Syntax

```
<new_table_name> <- SELECT <condition> FROM <table_name>
```

Where <condition> is of either form

```
<first_column_name> <bin_op> <second_column_name>  
<first_column_name> <bin_op> <int_literal>
```

Where <bin\_op> can be any operator among {>, <, >=, <=, =>, =<, ==, !=}

- The selection command only takes one condition at a time

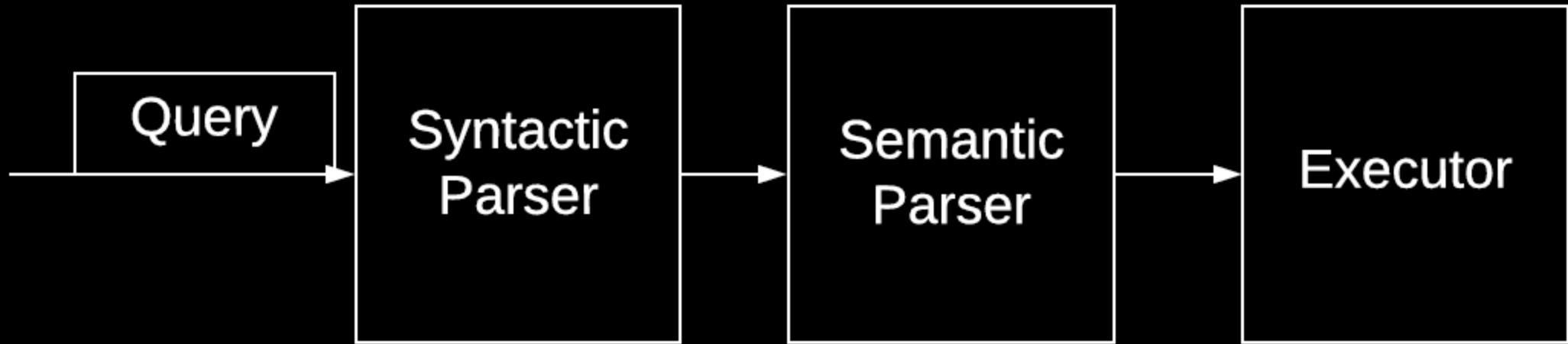
Run: `R <- SELECT a >= 1 FROM A`

`S <- SELECT a > b FROM A`

# Internals

- Buffer Manager
- Cursors
- Tables
- Executors

# Command Execution Flow



Run: `LOAD A` with debugger

see: `load.cpp`



# Syntactic Parser

- Splits the query into query units

see: syntacticParser.h syntacticParser.cpp

# Semantic Parser

- Makes sure your query makes semantic sense

see: semanticParser.h semanticParser.cpp

# Executors

Every command(COMMAND) has a file in the executors directory, within that directory you'll find 3 functions

```
syntacticParseCOMMAND  
semanticParseCOMMAND  
executeCOMMAND
```

# Buffer Manager

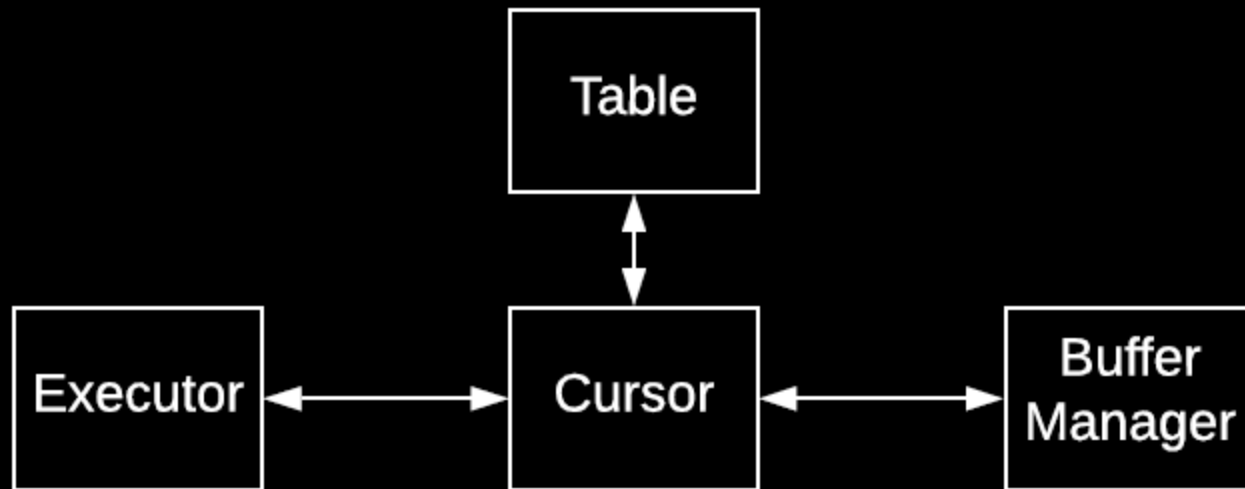
- Load splits and stores the table into blocks. For this we utilise the Buffer Manager
- Buffer Manager follows a FIFO paradigm. Essentially a queue

# Table Catalogue

- The table catalogue is an index of tables currently loaded into the system

# Cursors

A cursor is an object that acts like a pointer in a table. To read from a table, you need to declare a cursor.



Run: `R <- SELECT a == 1 FROM A` with debugger

# Logger

Every function call is logged in file names "log"

# Project\*

- Phase 1: Code Familiarity
- Phase 2: 2 Phase Merge Sort
- Phase 3: Indexing
- Phase 4: Optimization

\* Tentative

# Project Rules

- **Plagiarism: F**
- Not sticking to submission guidelines will lead to penalties and at times to scoring 0
- Project phases build on top of each other, failing to do one phase may hinder the rest
- If for any reason you fail to complete the project on time, please mail the Prof directly for extensions and not the TAs, the TAs have no jurisdiction in these cases



# Administrative Rules

- Moodle is the preferred platform. Informal contact is discouraged, do so at your own expense.
- A "doubts document" will be put up on Moodle.
- If you need to contact the TAs for matters that don't concern the whole class you may mail us here -  
[datasystems\\_ta\\_m21@IIITAPhyd.onmicrosoft.com](mailto:datasystems_ta_m21@IIITAPhyd.onmicrosoft.com)

# References

- GitHub Repo - [SimpleRA](#)
- Build and run instructions will be provided later