# 2019 B+ tree implementation assignment

Course name: Database Systems (ITE2038)

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# 1. Assignment Title

• Implementation of a B+ tree index

# 2. Environment

- OS: Windows
- Language: Java or Python (any version is ok)

# 3. Constraints - Overall

- The B+ tree index should be stored in a single file (index file)
- The file contains all the meta information for the index and also the index nodes
- The internal organization of the file is not considered in grading
- The program should provide following functions:
  - Search
    - ✓ A single key search <u>AND</u> a range search
  - Insertion of a key
  - Deletion of a key
    - ✓ The deleted entry should be completely removed from the index and the file
- Assumption
  - Keys and values are all in the integer type
  - Duplicated keys are not allowed for insertions
  - The keys in a node are stored in an <u>ASCENDING order</u>
- POLICY on COPY DO NOT COPY someone else's program
  - DO NOT USE functions/methods/routines from existing code/library/programs in pre-implemented B+ tree indexes or any other similar tree-based indexes
  - All these actions are regarded as COPY and so will be handled accordingly

# 4. Constraints - Internal Structure

- Each node of a B+ tree index should contain the following data inside:
  - Non-leaf node
    - ✓ m: # of keys
    - ✓ p: an array of b <key, left\_child\_node> pairs
    - $\checkmark$  r: a pointer to the rightmost child node
  - Leaf node
    - ✓ m: # of keys

- ✓ p: an array of b <key, value(or pointer to the value)> pairs
- $\checkmark$  r: a pointer to the right sibling node

# 5. Constraints – Interface

- The program *should support command-line interface*
- The following commands should be implemented:

#### ■ Data File Creation

- ✓ Command: program -c index\_file b
  - program: name of the program (bptree)
  - > index\_file: name of a new index file
  - *b*: size of each node (max. # of child nodes)
- ✓ This command creates a new index file containing an empty index with node size b
  - If the file already exists, it is overwritten
- ✓ Example
  - > java bptree -c index.dat 8

#### ■ Insertion

- ✓ Command: program -i *index\_file data\_file* 
  - data\_file: name of the input data file that has a number of key-value pairs to be inserted
- ✓ This command inserts all the key-value pairs inside the data\_file into the index in the index\_file
  - The insertion causes the modification of the index file
  - ➤ Insertions are performed in the same order of key-value pairs in the data file
- The data file is provided as a .csv file (Comma Separated Values)
  - Each line of the data file contains a key-value pair
    - <key>,<value>\n
  - Data file example (input.csv)

```
26,1290832
```

10,84382

87,984796

86,67945

20,57455

9,87632

00 5000

86,579952 68,97321

84,431142

37,2132

- Example
  - > java bptree -i index.dat input.csv

#### Deletion

- ✓ Command: program -d index\_file data\_file
  - data\_file: name of the input data file that has a number of keys to be deleted

- ✓ This command deletes all the key-value pairs inside the input data file from the index
  - The deletion causes the modification of the index file
  - > Deletions are performed in the same order of keys in the data file
- ✓ The input data file is provided as a .csv file (Comma Separated Values)
  - Each line of the data file contains only a key value
    - <key>\n
- Example
  - java bptree -d index.dat delete.csv

### Single Key Search

- Command: program -s index\_file key
  - ✓ key: key value to be searched
- This command returns a value of a pointer to a record with the key
- Output format
  - ✓ Print output to the *stdout*
  - ✓ While searching, the program prints each non-leaf node in the path that the search passes through
    - > Print all the keys in the node in a single line
    - $\rightarrow$  <key1>,<key2>, $\Box$ ,<keym>\n
  - ✓ When the search reaches the leaf node having the search key, print the value matched with the search key
    - < <value>\n
    - ➤ If not found, print □NOT FOUND□
  - ✓ Example
    - java bptree -s index.dat 125 >java bptree -s index.dat 125 54,356 67,98 65462

### Ranged Search

- Command: program -r index\_file start\_key end\_key
  - ✓ start\_key: lower bound of the range search
  - ✓ end\_key: upper bound of the ranged search
- This command returns the values of pointers to records having the keys within the range provided
- Output format
  - ✓ Print output to the *stdout*
  - ✓ Print all the key-value pairs with the key between *start\_key* and *end\_key* (including start\_key and end\_key)
    - $\rightarrow$  <key1>,<value1>\n<key2>,<value2>\n
  - ✓ Note that *start\_key* and *end\_key* may not be in the index
    - > The program prints only the key-value pairs between them
- Example

✓ java bptree -r index.dat 100 200

> java bptree -r index.dat 100 200

125,65462

169,3728

193,98732

200,164260

### 6. How to turn in

- (1) Write your program
- (2) Write a document (.doc, .docx, or .pdf) that contains:
  - Summary of your algorithm
  - Detailed description of your codes (for each function)
  - Instructions for compiling your source codes at TA's computer (e.g. screenshot) (Important!!)
    - You MUST SUBMIT instructions for compiling your source codes. If TAs read your instructions but cannot compile your program, you will get a penalty Please, write the instructions carefully
  - Any other specification of your implementation and testing
- (3) Zip the codes and the document
  - The filename should follow the format
    - B-tree\_Assignment\_<YOUR\_STUDENT\_NUMBER>.zip
    - Ex.) B-tree\_Assignment\_2010051924.zip
  - The zip file should contain a executable file, all source files, and the document
- (4) Submit it to gitlab (http://hconnect.hanyang.ac.kr/)
  - Due date

Completed before 25 September: 100%
Completed before 2 October: 70%
After 2 October: 0%

You can ask questions about the assignment via class community and/or e-mail YOU WILL GET SERIOUS PENALTIES IF YOU DO COPY OR CHEAT

Good luck!