

DB 2

09 – Ordered Indexes (B+Trees)

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Torsten Grust
Universität Tübingen, Germany

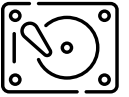
1 | Q_8 — Filtering an Indexed Table

Sequential scan (**Seq Scan**) and interpreted predicate evaluation go a long way. Large input tables call for significantly more **efficient support for value-based row access**:

```
SELECT i.b, i.c
FROM   indexed AS i
WHERE  i.a = 42 [i.c = 0.42] -- either filter on i.a or i.c
```

Assume column **a** is **primary key** in table **indexed**: expect query workload that frequently identifies rows via predicates **a = k**. **Indexes** can support such queries.

Primary Key and Indexes



```
CREATE TABLE indexed (a int PRIMARY KEY, -- ⇒ NOT NULL
                       b text,
                       c numeric(3,2));    -- ± d.dd
```

DBMS expects predicates $a = k$ and creates an **index on column a** —a data structure associated with and maintained in addition to table **indexed**—to speed up evaluation:

```
CREATE INDEX indexed_a ON indexed USING btree (a);
```

1. Whenever possible/promising, index **indexed_a**¹ is (also) consulted when table **indexed** is queried. 🚀
2. When **indexed** is updated, **indexed_a** is maintained. ⚙️

¹ PostgreSQL chooses index name **indexed_pkey** but let's follow a $\langle table \rangle_ \langle column \rangle$ naming scheme here.



EXPLAIN VERBOSE

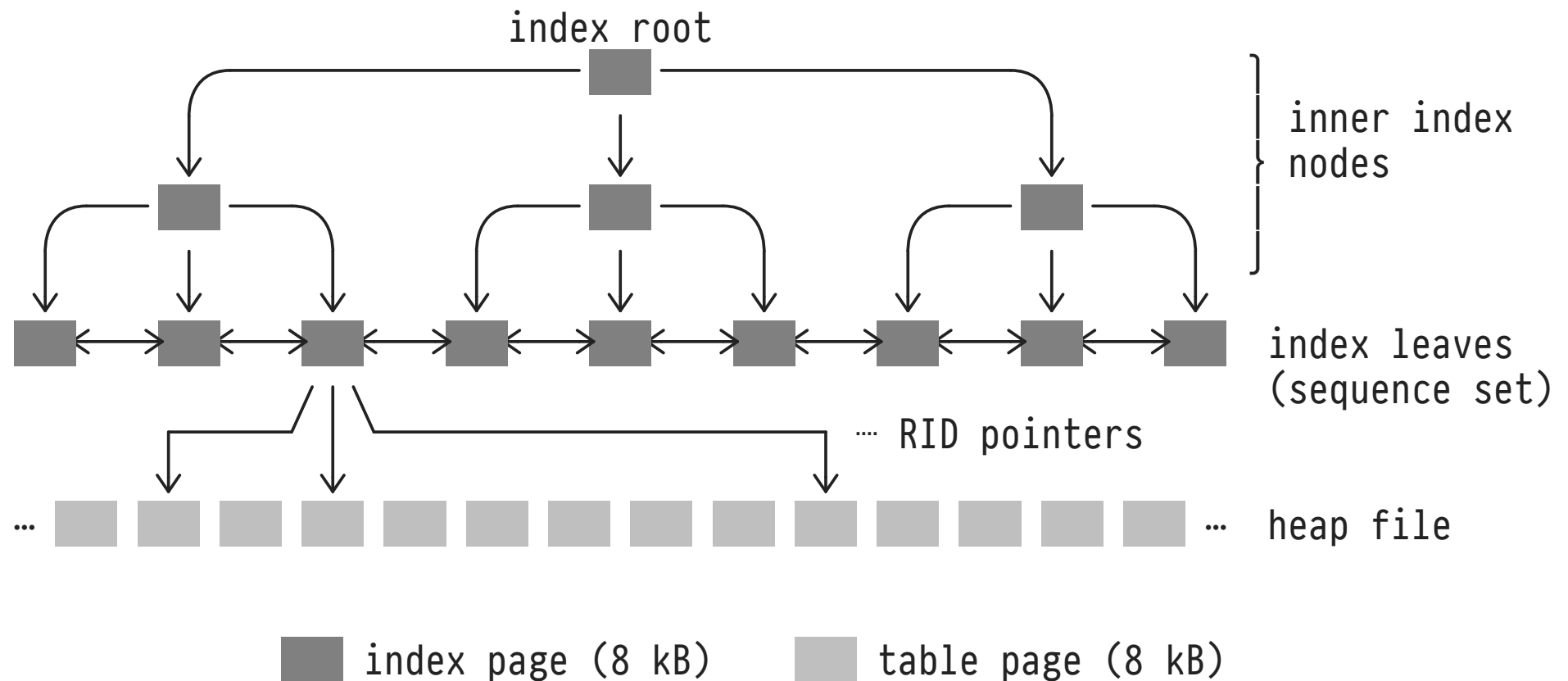
```
SELECT i.b, i.c
FROM   indexed AS i    -- 106 rows
WHERE  i.a = 42;       -- selection on key column a ⇒ ≤ 1 row will qualify
```

QUERY PLAN

```
Index Scan using indexed_a on indexed i (cost=0.42..8.44 rows=1 ...)
  Output: b, c
  Index Cond: (i.a = 42)
```

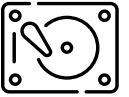
- DBMS uses **Index Scan** (instead of **Seq Scan**), index scan will evaluate predicate $i.a = k$.
- System expects small result of a single row (**rows=1**), i.e., the predicate is assumed to be *very selective*.

2 | B+Trees: Ordered Indexes



Anatomy of a B+Tree

B+Trees: Ordered Indexes

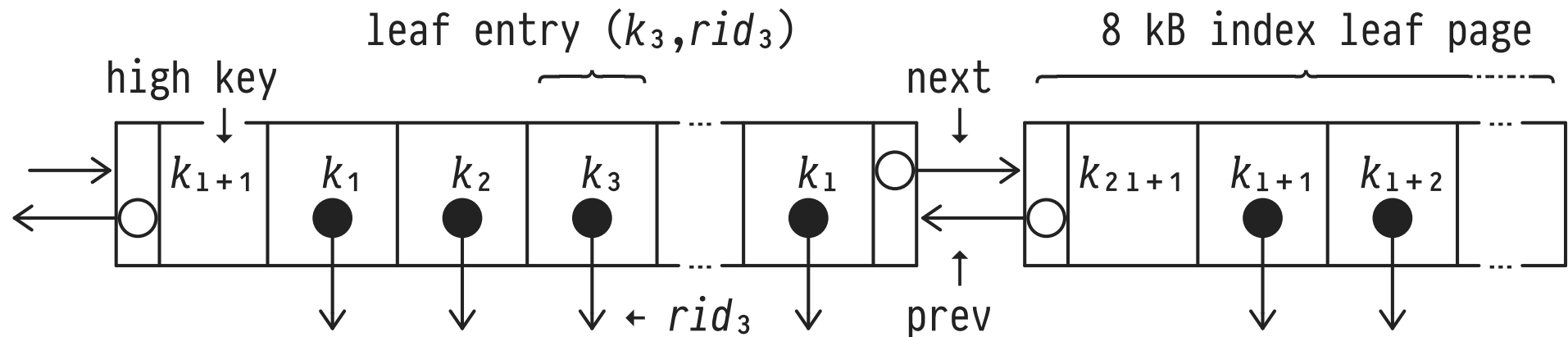


Notes on B+Tree anatomy:

- A **B+Tree²** index I on column $T(a)$ is an *ordered*, n -ary ($n \gg 2$), *balanced*, *block-oriented*, *dynamic* search tree.
- Inner nodes and leaves are formed by 8 kB **index pages**.
- Each inner node holds $n-1$ values of column a (**separators**) that allow to navigate the search tree structure.
- Leaves form a bidirectional chain, the **sequence set**.
- **Leaves use RIDs to point to rows** in the heap file of table T : besides a column values, I holds no data of T .

² Invented by Bayer and McCreight (1969) at Boeing Labs. The “B” in “B+Tree” does *not* stand for Bayer, binary, balanced, block, or Boeing. (We tried to find out, but Rudolf Bayer wouldn't say.)

B+Trees: Inside a Leaf Node



- Uses pointers **prev/next** to form the chained **sequence set**.
- **Leaf entries are ordered** by index keys k_i : $k_i \leq k_{i+1}$.
- RID rid_i points to a row t of T with $t.a = k_i$.
- The **high key** holds smallest key of *next* leaf (if any).

B+Trees: How to Find Rows t With $t.a = k$?

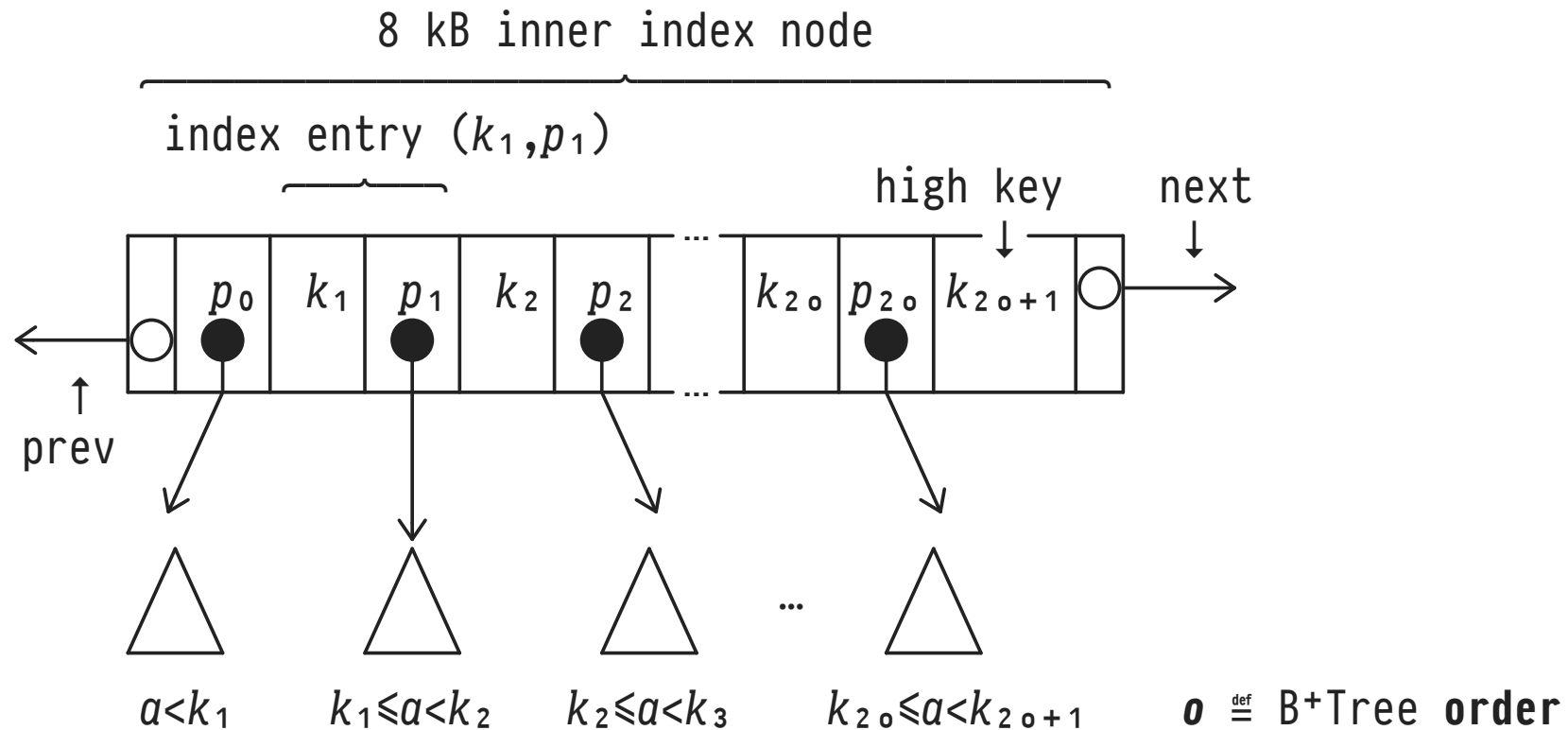
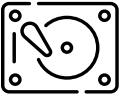


As described, a B+Tree is a **dense** index structure: every row t of T is represented by one leaf entry.

- The sequence set is ordered by keys $k_i \Rightarrow$ a *binary search* for a key $k = k_i$ may sound viable, **BUT** the search would
 1. need to inspect $\log_2(|T|)$ keys in the sequence set and access just as many pages 🗨, and
 2. “jump around” the sequence set in an unpredictable fashion, thus leading to random I/O. 🗨

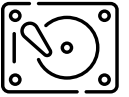
B+Trees exploit the sequence set ordering and erect an **n -ary search tree structure** (n large!) atop the leaf entries.

B+Trees: Inside an Inner Node



- The **separator** keys k_i are ordered: $k_i \leq k_{i+1}$.
- Page pointers p_j point to index (leaf or inner) nodes.

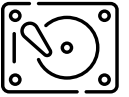
B+Trees: Notes on Inner Nodes



- Space in inner nodes is used economically: in a B+Tree of **order** o , any inner node—but the root node—is guaranteed to hold between o and $2 \times o$ ($\stackrel{\text{def}}{=} \mathbf{fan-out } F$) index entries.
- Given predicate $t.a = k$, perform **binary search inside node** to find B+Tree subtree with $k_i \leq k < k_{i+1}$.
- B+Tree is **balanced**: subtrees Δ are of identical height.
- **Path length** s from B+Tree root to leaf node **predictable**:

$$\underbrace{|T| \times 1/F \times \dots \times 1/F}_{s \text{ times}} = 1 \Leftrightarrow s = \log_F(|T|)$$

3 | Index Scan



A B+Tree is *the* index structure to support the evaluation of these kinds of conditions:³

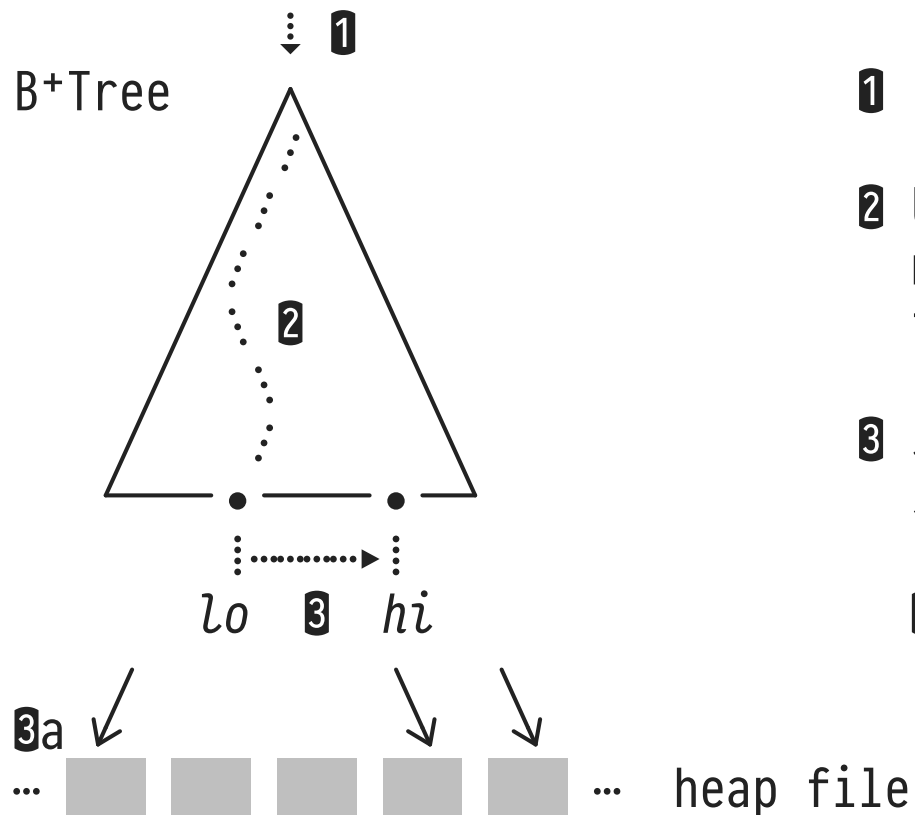
1. **Range predicates:** $lo \leq a \leq hi$
 2. **Half-open ranges:** $lo \leq a$ or $a \leq hi$
 3. **Equality predicates:** $a = lo$
- An **Index Scan** on index I for column $T(a)$ is parameterized by such a condition (PostgreSQL **EXPLAIN: Index Cond**).
 - **Index Scan** uses lo to navigate the search tree structure and locate the start of relevant sequence set section.

³ Half-open ranges are special range predicates where $hi = \infty$ ($lo = \infty$). Equality predicates are special range predicates where $lo = hi$.

Index Scan for Condition $lo \leq a \leq hi$



An **index scan** accesses the B+Tree index *and* the heap file:



- 1 Enter at B+Tree root page
- 2 Use key lo to **navigate the inner nodes** (search tree) until we reach the leaf level
- 3 Scan leaf entries in the sequence set section $lo \leq a \leq hi$, **extract RIDs**
 - 3a For each RID, **access heap file for table T** and return matching row

Navigating the Inner Nodes



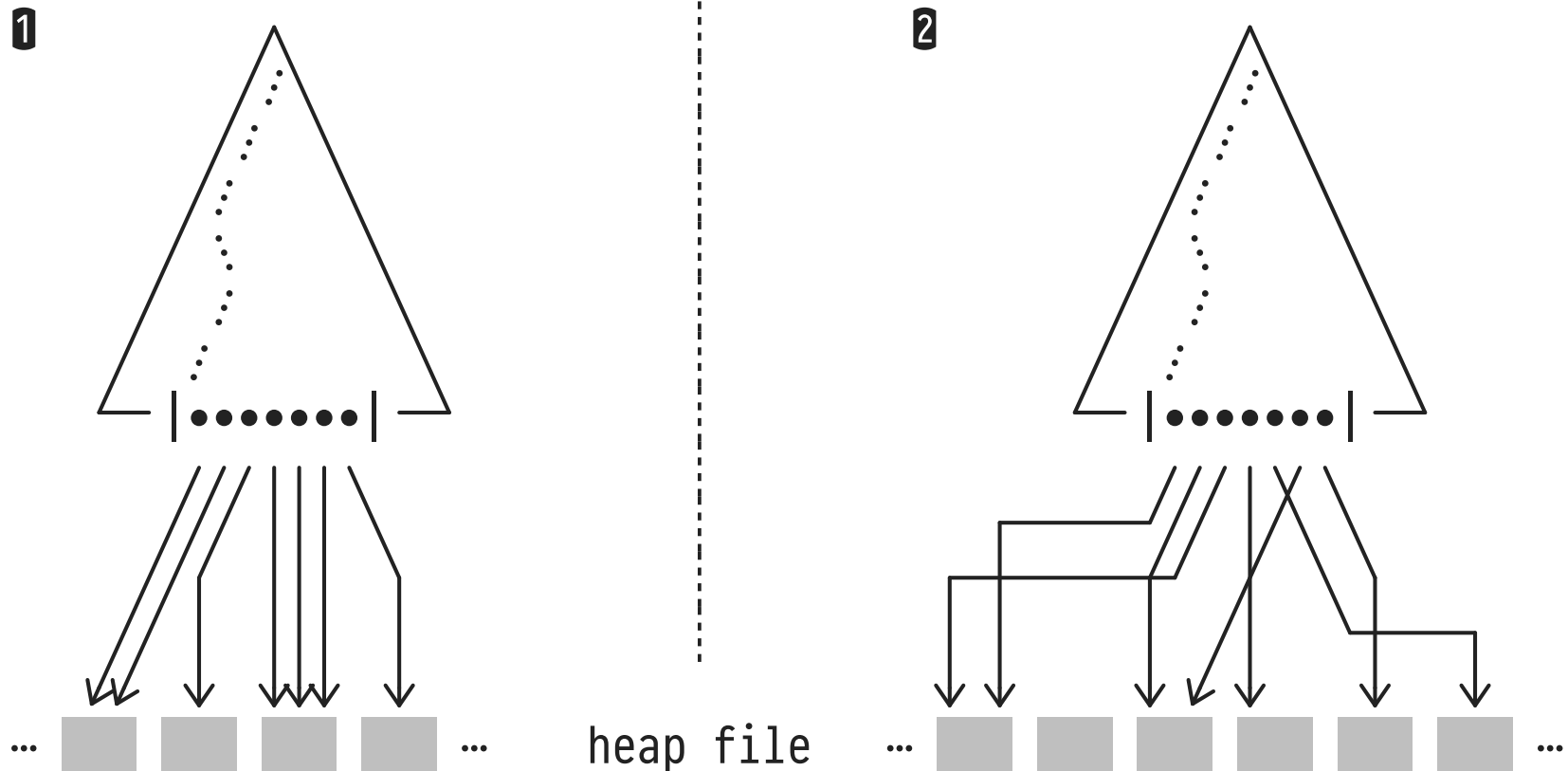
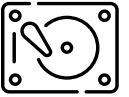
Phase ② runs a vanilla traversal of a $2 \times o$ -way search tree:

```
Search(lo):  
    return TreeSearch(lo, root);  
  
TreeSearch(lo, node):  
    if (node is a leaf)  
        | return node;  
    switch lo  
        | case  $lo < k_1$   
        |   | return TreeSearch(lo,  $p_0$ );  
        | case  $k_i \leq lo < k_{i+1}$   
        |   | return TreeSearch(lo,  $p_i$ );  
        | case  $k_{2o} \leq lo$   
        |   | return TreeSearch(lo,  $p_{2o}$ );
```

} returns entry point
for scan of
sequence set

} use binary search
to implement
subtree choice

4 | Order of Leaf Entries vs Order of Table Rows



- ① Order of leaf entry keys k_i \equiv row order in heap file. 👍
- ② Order of k_i in sequence set and row order do *not* match.

Clustered Indexes



Index I for column $T(a)$ is **clustered** if the order of leaf entries coincides with T 's row order (i.e., both I 's sequence set and T 's heap file are ordered by a):

Given entries $\langle k_i, p_i \rangle$ and $\langle k_j, p_j \rangle$, $k_i \leq k_j \Rightarrow p_i \leq p_j$.

- An **Index Scan** over a *clustered* index
 1. collects matching rows from adjacent heap file pages (\Rightarrow sequential I/O 👍),
 2. will find many matching rows on each accessed heap file page (\Rightarrow less page I/O 👍).

Non-Clustered Indexes

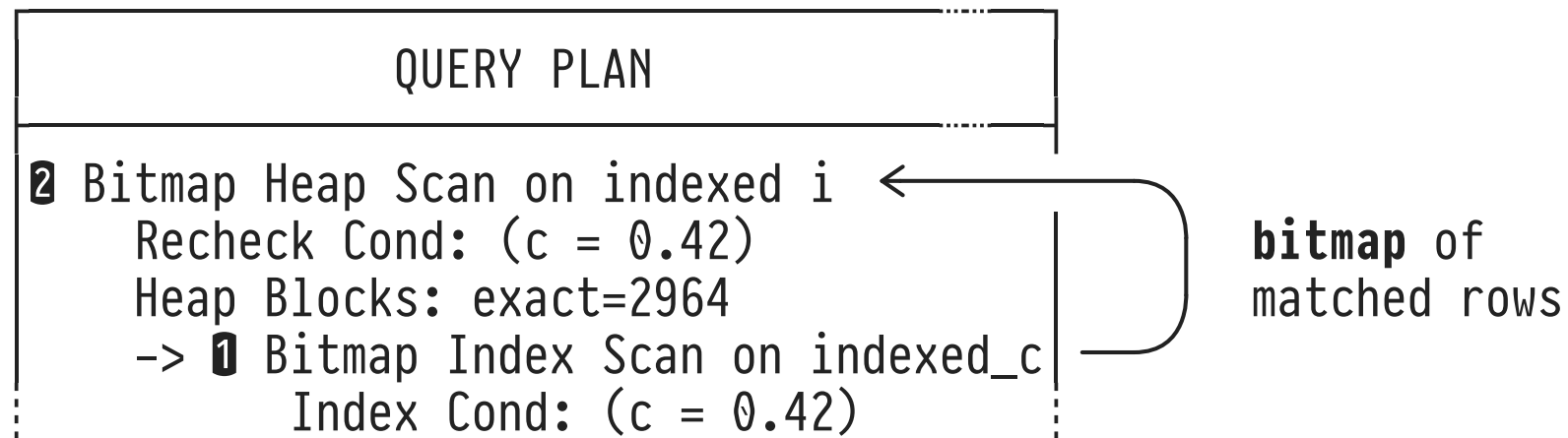
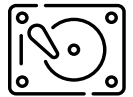


Sad fact: only *one*—among the many possible—indexes for a table may be clustered. Most indexes are non-clustered.

- An **Index Scan** over a *non-clustered* index
 1. will find matching rows potentially scattered across all heap file pages (\Rightarrow random I/O 🗨),
 2. will find few matching rows on each accessed heap file page and may access the same page more than once (\Rightarrow as many page I/Os as matching rows 🗨).

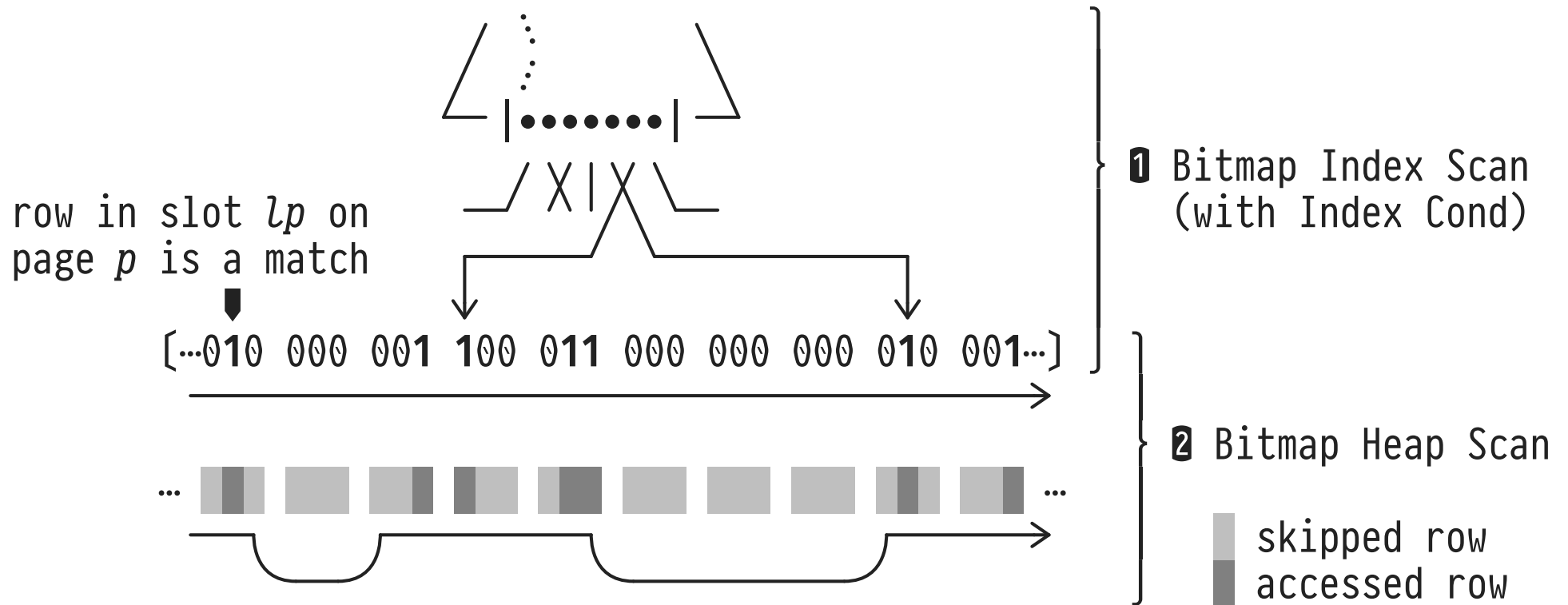
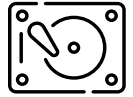
PostgreSQL addresses this challenge through **RID sorting**, implemented via **Bitmap Index Scan** & **Bitmap Heap Scan**.

Bitmap Index Scan & Bitmap Heap Scan



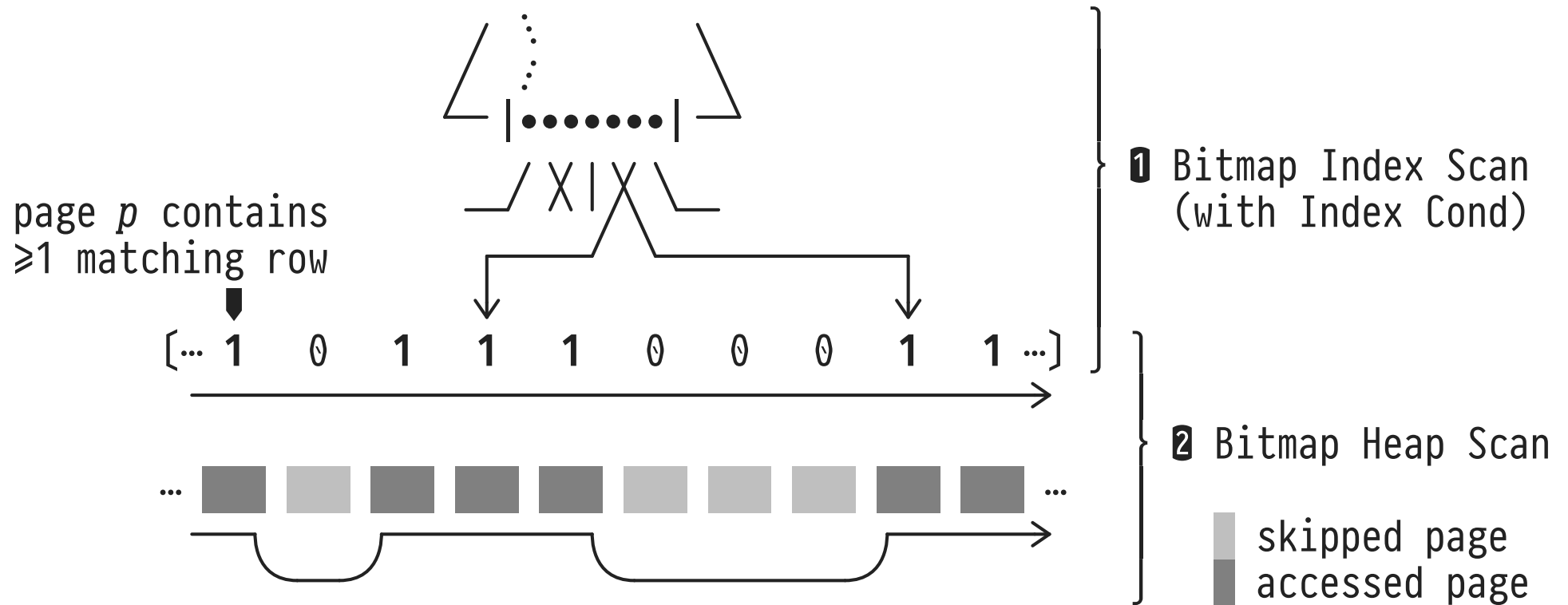
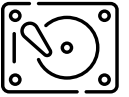
- 1 Bitmap Index Scan:** perform Index Scan and create **bitmap** that encodes *heap file locations* of rows matching the **Index Cond**. Do *not* access rows in heap file yet.
- 2 Bitmap Heap Scan:** scan heap file once, only access those rows (pages) that have been marked **1** in the bitmap.

Bitmap Index Scan & Bitmap Heap Scan: Row-Level Bitmap



Bitmap Heap Scan performs one sequential scan (with skips) of the heap file, regardless of RID order in sequence set.

Bitmap Index Scan & Bitmap Heap Scan: Page-Level Bitmap



Working memory tight \Rightarrow build **page-level** bitmap. ⚠ In ②, need to **recheck condition** for all rows on accessed pages.

5 | CLUSTERing Based on an Index



If the workload depends on top performance of particular predicates supported by *non-clustered* index *I*, we may

physically reorder the rows of underlying table's *T* heap file to coincide with the key order in *I*'s sequence set (i.e., *I* will become a *clustered* index⁴):

```
CLUSTER [VERBOSE] <T> USING <I>;  
CLUSTER <T>;    -- re-cluster once T's rows get out of order
```

- ⚠ Subsequent updates on *T* can destroy the perfect clustering. (May need to re-cluster *T* in intervals.)

⁴ At a price, of course: formerly *clustered* indexes on *T* will turn into *non-clustered* indexes.



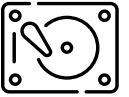
B+Trees...

1. **economically utilize space** in inner/leaf nodes (minimum node occupancy 50%, typical fill factor 67%),
2. are **balanced** trees and thus require a **predictable number of page I/Os** to traverse from root to sequence set—enables query optimizer to forecast B+Tree access cost.

DBMSs maintain properties 1. and 2. when rows are **inserted into/deleted** from an B+Tree-indexed table.⁵

⁵ Some real B+Tree implementations of row deletion deviate from the textbook to keep things simpler.

B+Tree Insertion for New Entry $\langle k, rid \rangle$ (Sketch)

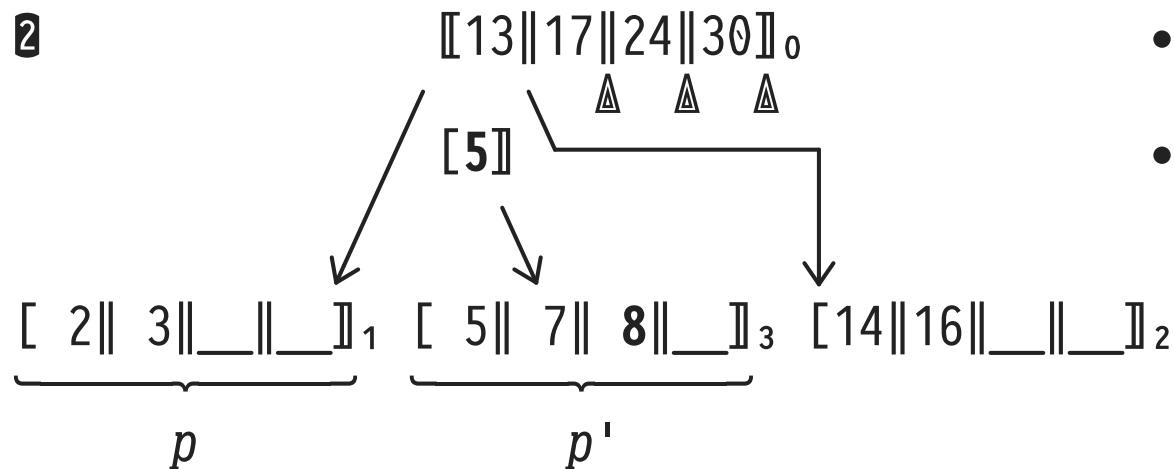
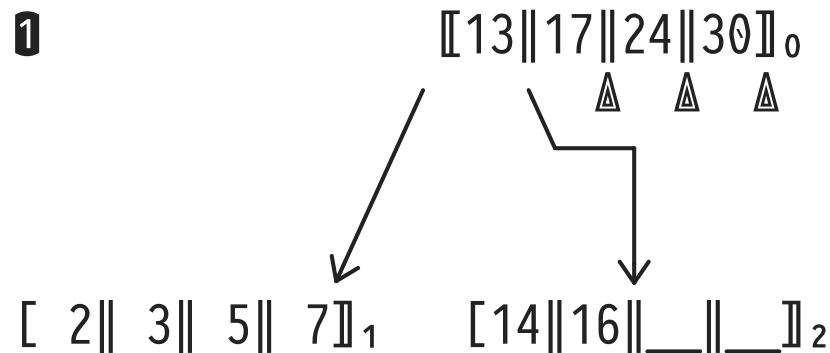


1. Use $\text{Search}(k)$ to **find leaf page** p which should hold the entry for k .
2. If p has **enough space** to hold new entry (i.e., at most $2 \times o - 1$ entries in p), **simply insert** $\langle k, rid \rangle$ into p .
3. Otherwise, node p must be **split** into p and p' and a new **separator** has to be inserted \bigcirc into the parent of p .

Splitting happens recursively \bigcirc and may eventually lead to a split of the root node (increasing B+Tree height).

- **Distribute** the entries of p and new entry $\langle k, rid \rangle$ onto pages p and p' .

B+Tree Insertion and Leaf Node Split

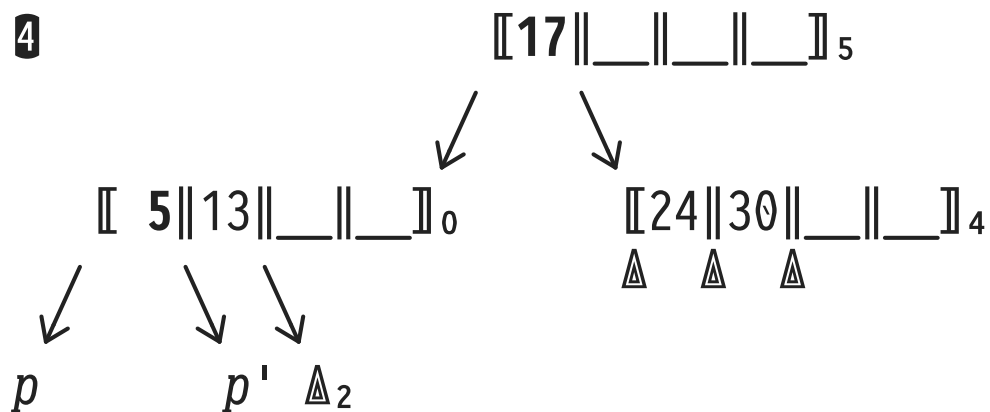
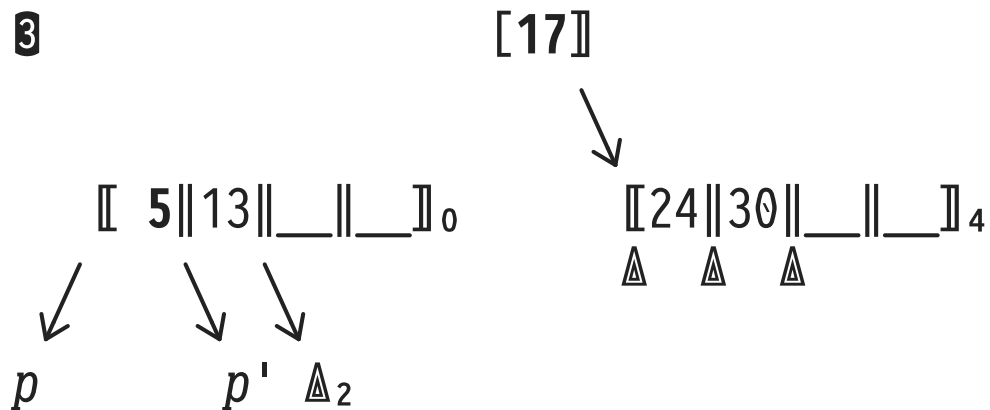


- 1** Insert new entry $\langle 8, rid \rangle$
- Search(8) returns leaf $p = 1$
 - Leaf 1 is full \Rightarrow split

- 2** Leaf 1 split into leaves $p = 1$ and $p' = 3$
- Distribute $\{2, 3, 5, 7, 8\}$ between leaves 1 and 3
 - **Copy** new separator [5] into parent node 0

p'

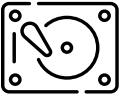
B+Tree Insertion and Inner Node Split



- 3** Inner node 0 (here: root) is full \Rightarrow split
- Inner node 0 splits into old node 0 and new $p'' = 4$
 - Distribute {5, 13, 24, 30} Δ between nodes 0 and 4
 - **Move** new separator [17] into parent of node 0
- ↓
- p''

- 4** Split node 0 has been the old root
- Create new root node 5, has [17] as only entry
 - B+Tree height has increased

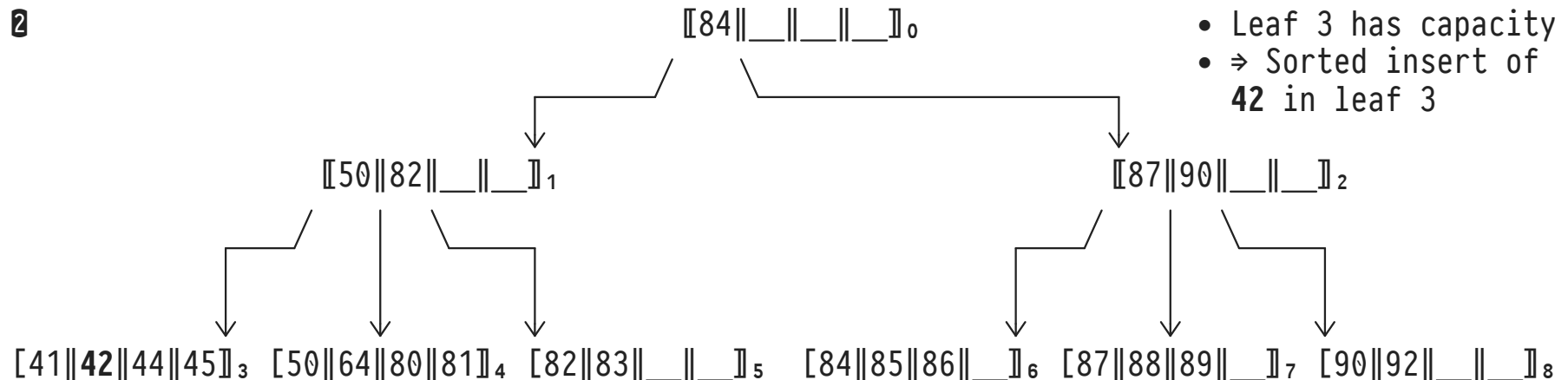
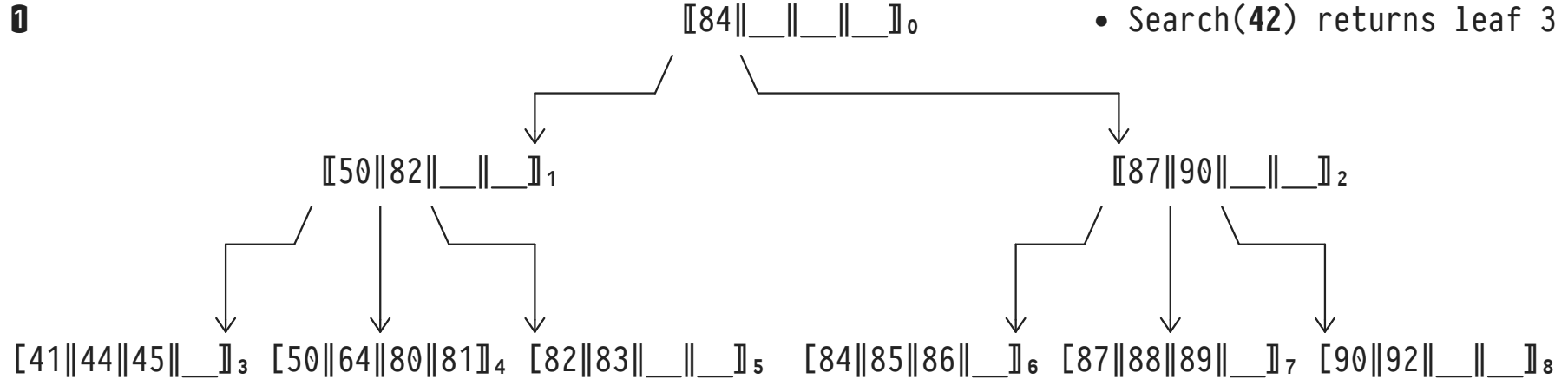
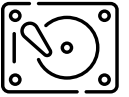
B+Tree Insertion Notes



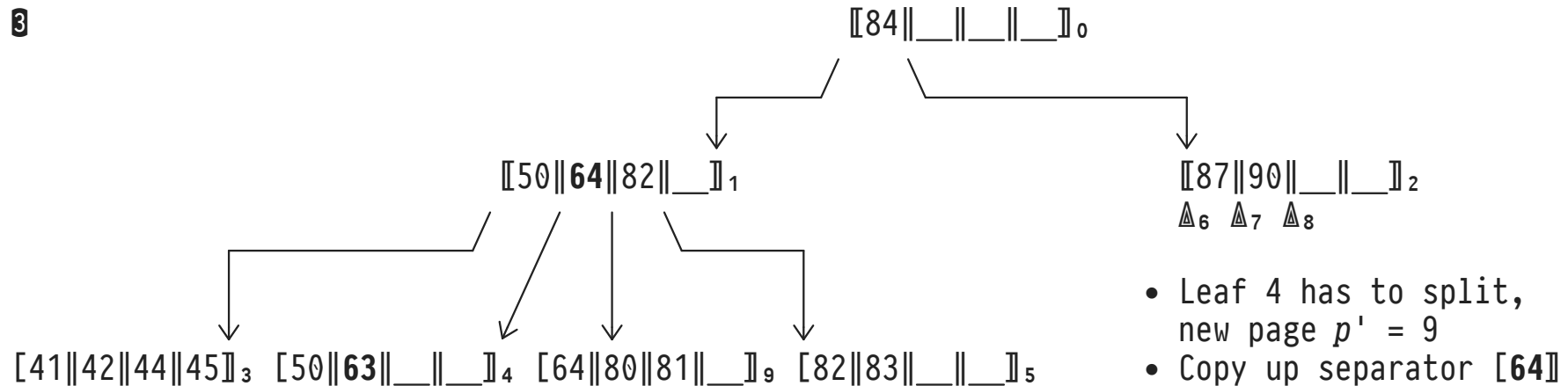
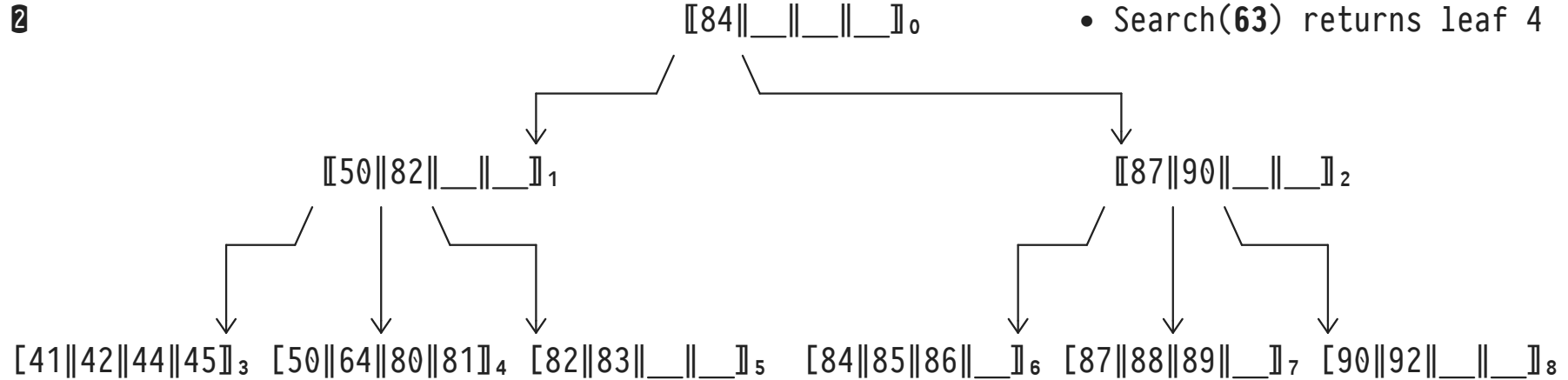
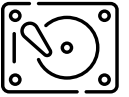
- Splitting starts at the leaf level and continues upward as long as inner index nodes are fully occupied (holding $2 \times o$ entries).
- ⚠ Unlike during a *leaf* split, an *inner* node split **moves**⁶ the new separator [**sep**] discriminating between p and p' upwards and recursively inserts it into the parent. Q: Why?
- Q: How often do you expect a root node split to happen?

⁶ A leaf node split **copies** the new separator upwards, i.e., the entry [**sep**] also remains at the leaf level.

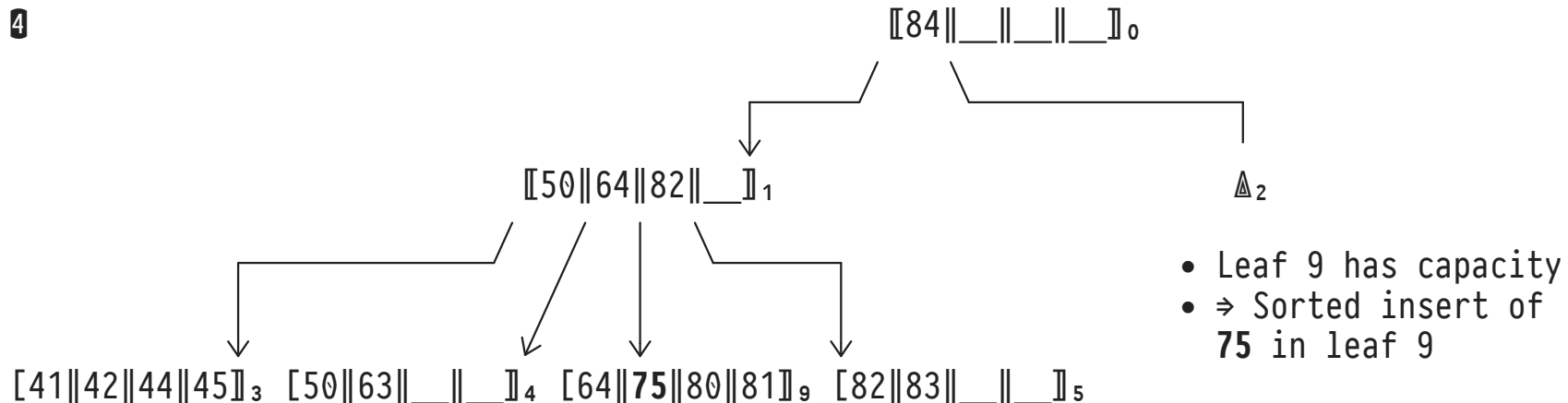
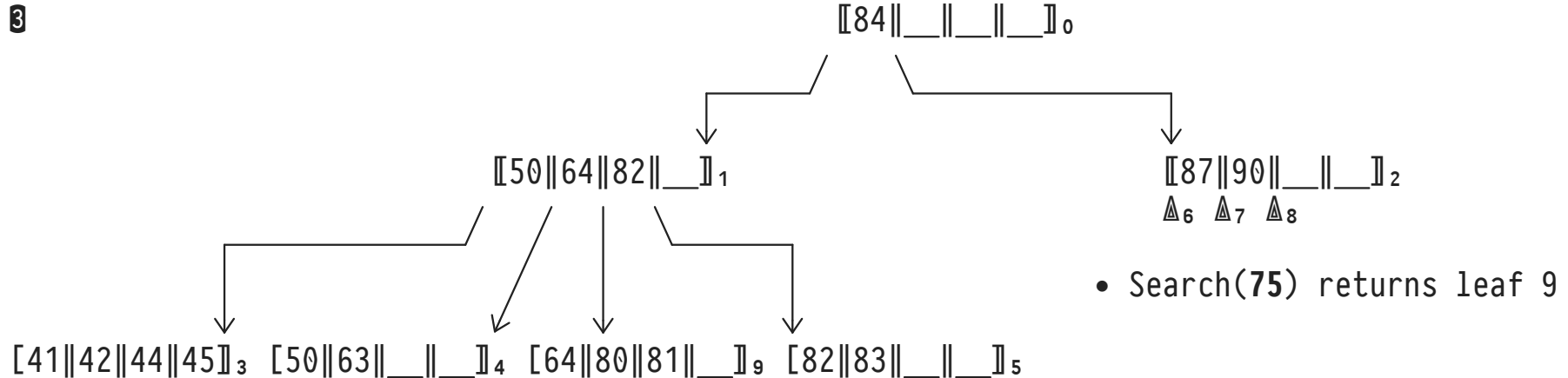
B+Tree Insertion Example: Insert *<42,rid>*



B+Tree Insertion Example: Insert $\langle 63, \text{rid} \rangle$



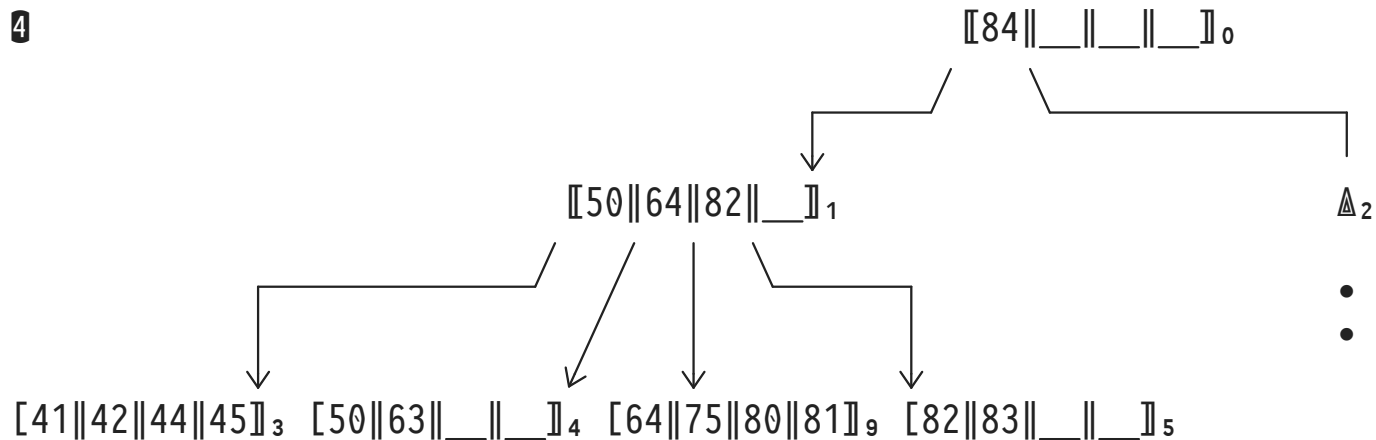
B+Tree Insertion Example: Insert $\langle 75, rid \rangle$



B+Tree Insertion Example: Insert $\langle 77, \text{rid} \rangle$

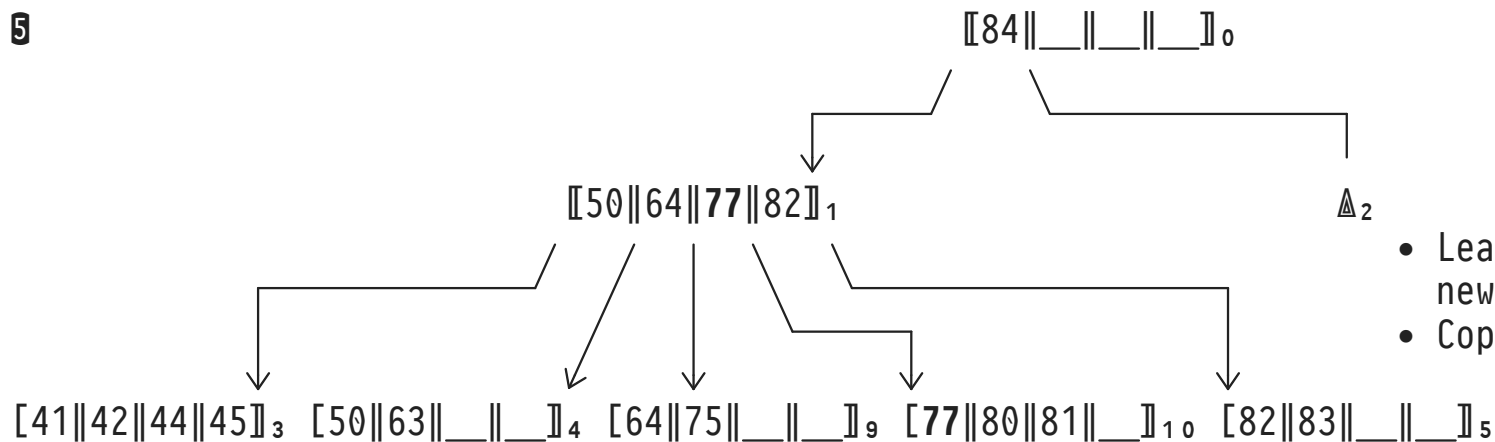


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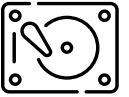
- Search(77) returns leaf 9
- Leaf 9 is full (already holds $2 \times 0 = 4$ entries)

5



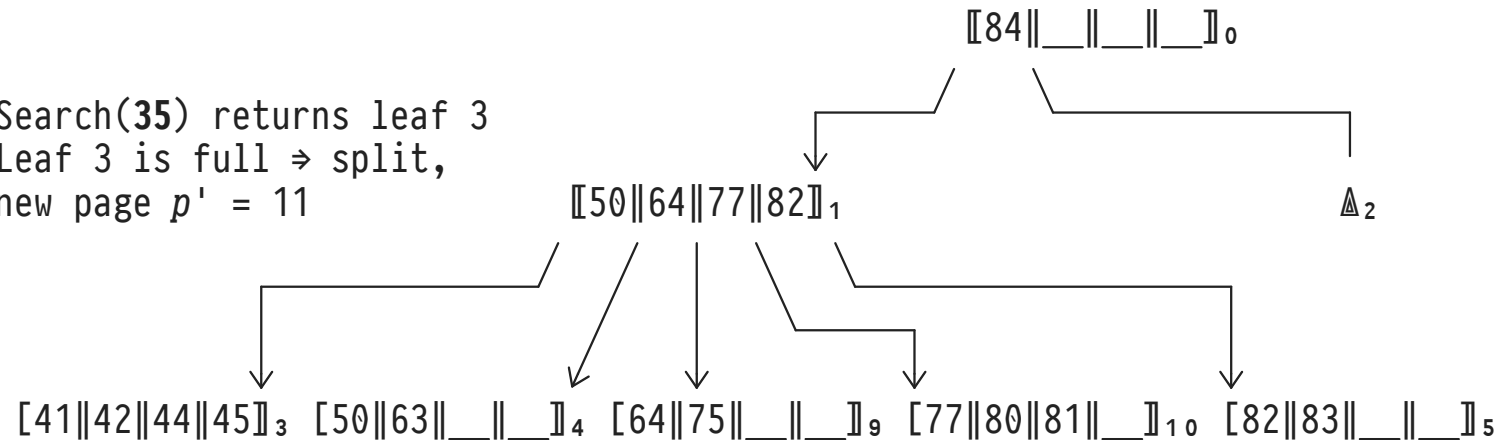
- Leaf 9 has to split, new page $p' = 10$
- Copy up separator [77]

B+Tree Insertion Example: Insert $\langle 35, \text{rid} \rangle$



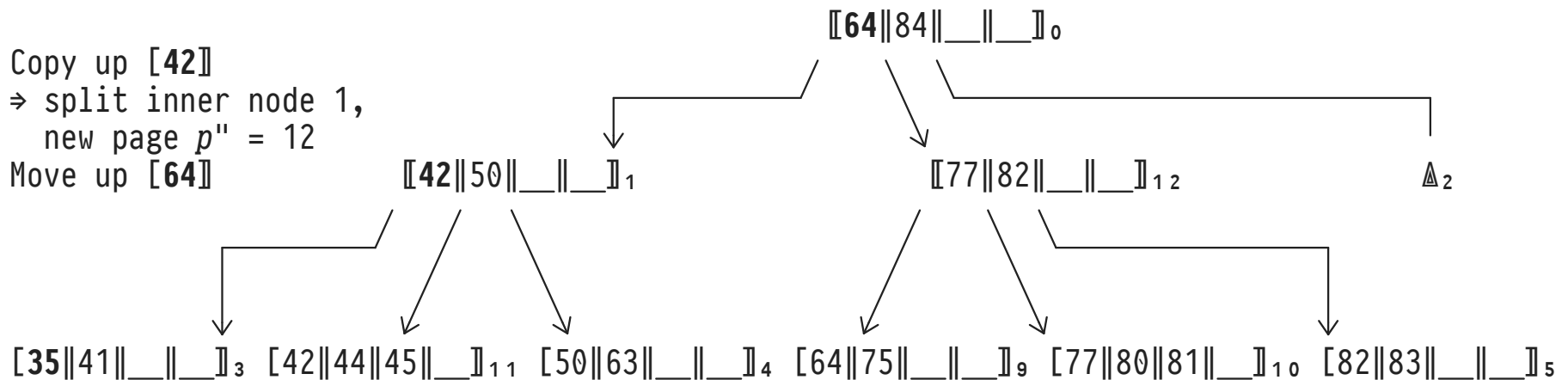
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- Search(35) returns leaf 3
- Leaf 3 is full \Rightarrow split, new page $p' = 11$



6

- Copy up [42]
 \Rightarrow split inner node 1, new page $p'' = 12$
- Move up [64]



⁷ Note: $\langle sep, ptr \rangle \equiv [sep]$ in our discussion above.

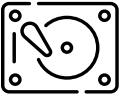
B+Tree Insertion Algorithm (2)



```
LeafInsert( $\langle k, rid \rangle, node$ ):  
  if ( $node$  has  $< 2 \times o$  entries)  
    | insert  $\langle k, rid \rangle$  into  $node$ ;  
    | return  $\langle 1, 1 \rangle$ ; }  $\langle 1, \_ \rangle \equiv$  no upwards split required  
  else  
    |  $p' \leftarrow$  allocate leaf page;  
    |  $[\langle k_1, rid_1 \rangle, \dots, \langle k_{2o+1}, rid_{2o+1} \rangle] \leftarrow$  entries of  $node \cup \langle k, rid \rangle$ ;  
    |  $node \leftarrow [k_1 | rid_1 | \dots | k_o | rid_o | \_ || \_ ]$ ;  
    |  $p' \leftarrow [k_{o+1} | rid_{o+1} | \dots | k_{2o+1} | rid_{2o+1} | \_ || \_ ]$ ;  
    | return  $\langle k_{o+1}, p' \rangle$ ; } new separator to be copied upwards
```

- **Copy upwards:** entry $\langle k_{o+1}, rid_{o+1} \rangle$ remains in leaf p' .

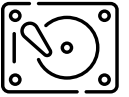
B+Tree Insertion Algorithm (3)



```
InnerInsert(<sep, ptr>, node):  
  if (node has < 2×o entries)  
    | insert <sep, ptr> into node;  
    | return <1, 1>; } <1, _> ≡ no upwards split required  
  else  
    | p' ← allocate inner node page;  
    | [p0, <k1, p1>, ..., <k2o+1, p2o+1>] ← entries of node ∪ <sep, ptr>;  
    | node ← [p0 | k1 | p1 | ... | ko | po | __ || __];  
    | p' ← [po+1 | ko+2 | po+2 | ... | k2o+1 | p2o+1 | __ || __];  
    | return <ko+1, p'>; } new separator to be moved upwards
```

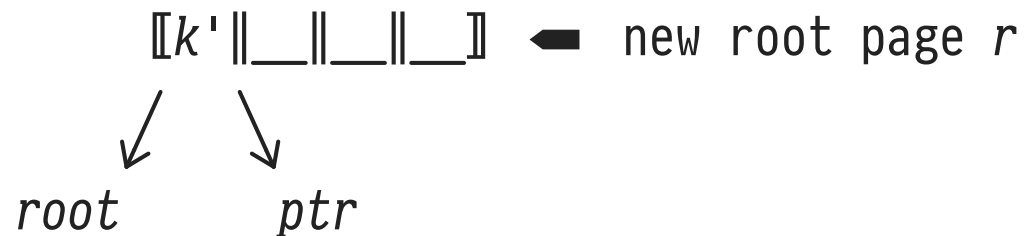
- **Move upwards:** new entry $\langle k_{o+1}, p' \rangle$ returned for insertion at parent. No entry $\langle k_{o+1}, _ \rangle$ remains at level of $node/p'$.

B+Tree Insertion Algorithm (Top Level)



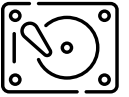
`Insert(<k,rid>)` is the top-level B+Tree insertion routine:

```
Insert(<k,rid>):  
  <k',ptr> ← TreeInsert(<k,rid>,root); } root ≡ old root page  
  if (k' ≠ ⊥)  
  |   r ← [root|k'|ptr|__||__||__];      } r ≡ new root page  
  |   root ← r
```

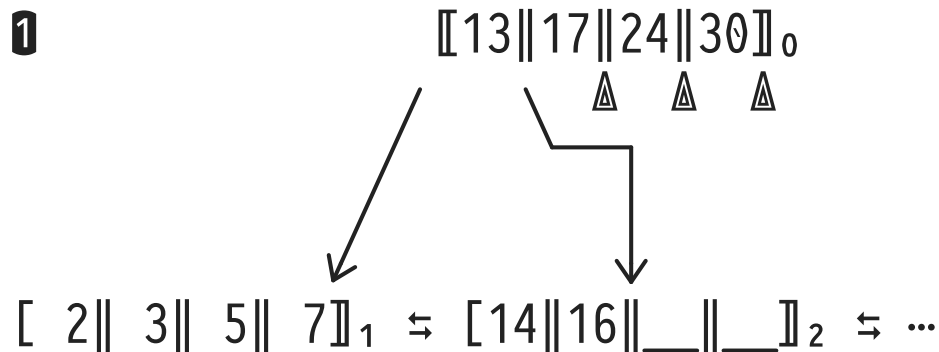


- Note: `Insert()` may leave us with a new root node that violates the minimum occupancy rule. $\text{~}\backslash(\text{ツ})/\text{~}$

B+Tree Insertion: Redistribution (1)



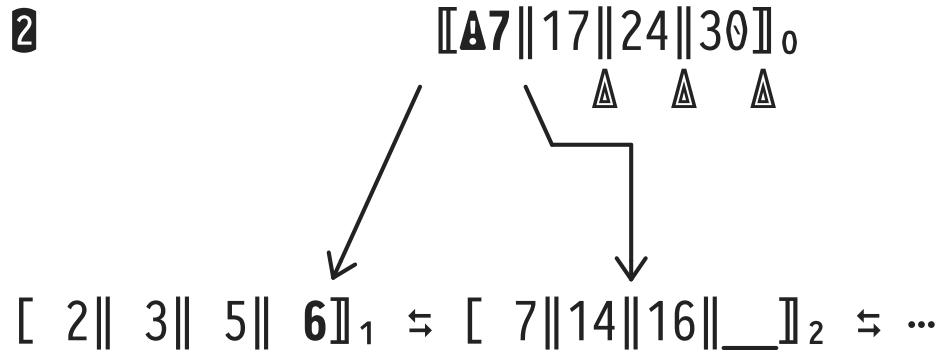
Can improve average occupancy and delay height increase on B+Tree insertion through **redistribution**:



- 1 Insert new entry $\langle 6, rid \rangle$
- Search(6) returns leaf 1
 - Leaf 1 is full, but its right **sibling** 2 has capacity

- Use sequence set chain pointers (\Leftrightarrow) to inspect **sibling** nodes for spare capacity.
- **Push** entry from overflowing node to sibling and **!** **update separator in parent node** to reflect this redistribution.

B+Tree Insertion: Redistribution (2)



- 2 Push entry $\langle 7, rid' \rangle$ to leaf 2
- Place $\langle 6, rid \rangle$ in leaf 1
 - Update separator (13 \rightarrow 7) in parent node 0
 - B+Tree remains at height 1

- Inspecting node sibling involves additional page I/O. 🗨
- Actual implementations use redistribution on the index leaf level only (if at all).

7 : B+Tree Deletion of Entry With Key k (Sketch)

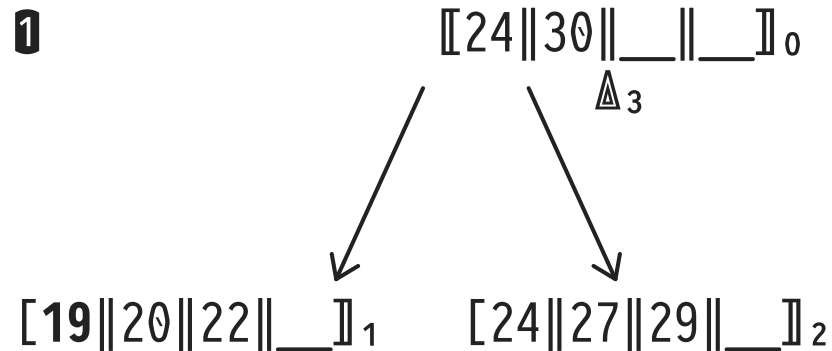
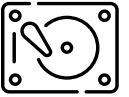


1. Use $\text{Search}(k)$ to **find the leaf** p holding entry $\langle k, \text{rid} \rangle$.
2. **Simply delete** $\langle k, \text{rid} \rangle$ from p .⁸
3. If p now holds $< o$ entries, leaf p **underflows**. Any sibling of p with spare entries?
 - Yes, use **redistribution** to move an entry into p .
 - No, **merge** p and a sibling leaf p' of o entries.
Delete \bigcirc the now obsolete separator of p and p' in their parent node.

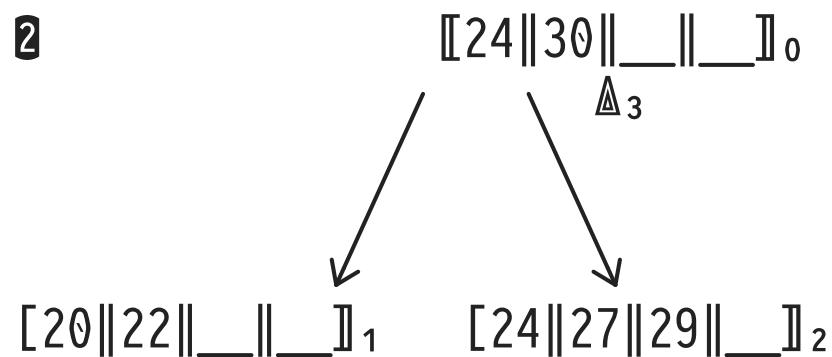
Deletion propagates upwards and may eventually leave the root node empty (decreases B+Tree height).

⁸ **Q:** If $\langle k, \text{rid} \rangle$ is the leftmost entry in p , do we need to update the associated separator entry in p 's parent node? Why not?

B+Tree Deletion (No Underflow)

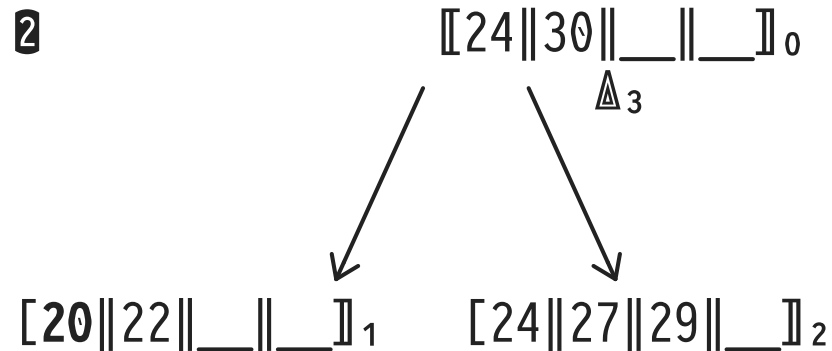


- 1 Delete entry with key $k = 19$
- Search(**19**) returns leaf 1
 - Leaf 1 has > 0 entries, node will not underflow

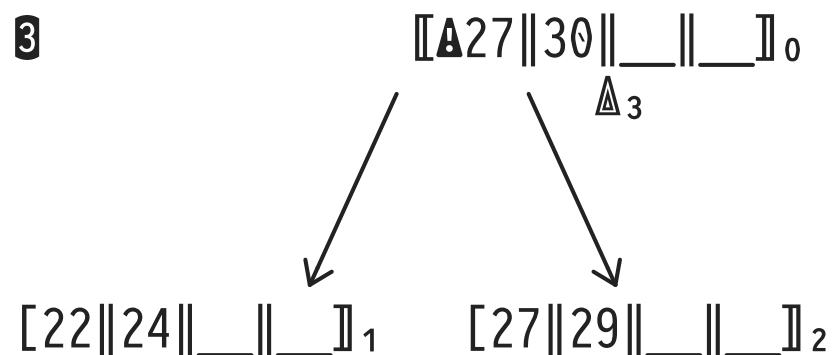


- 2 Simply delete entry $\langle 19, rid \rangle$ from leaf 1

B+Tree Deletion and Redistribution

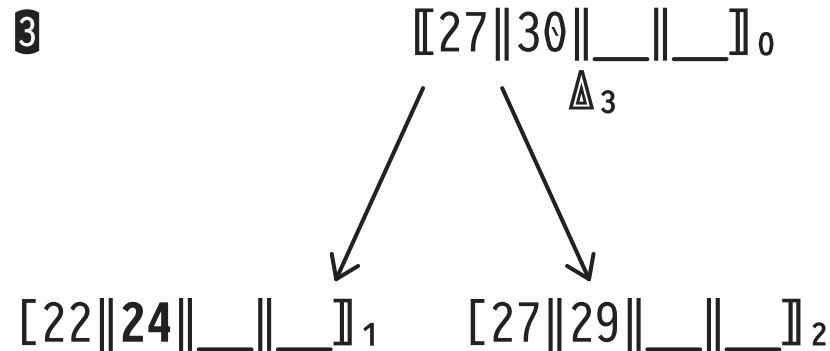


- 2 Delete entry with key $k = 20$
- Search(20) returns leaf 1
 - Leaf 1 has minimum occupancy of 0 entries \Rightarrow will underflow

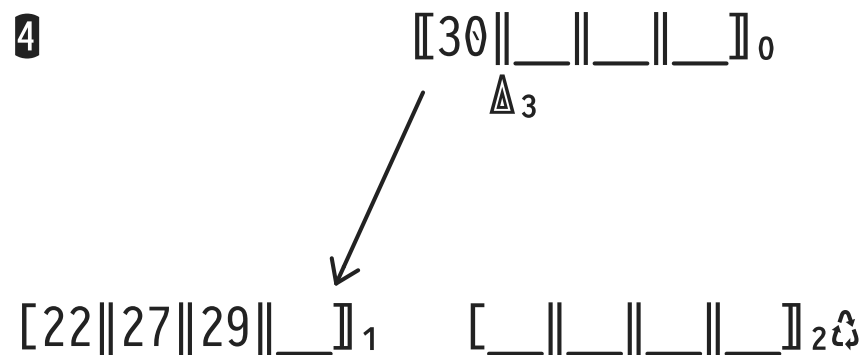


- 3 Sibling $p' = 2$ has one entry to spare \Rightarrow redistribution
- Move entry $\langle 24, rid' \rangle$ from leaf 2 to leaf 1
 - Update separator (24 \rightarrow 27) in parent node 0

B+Tree Deletion and Leaf Node Merging

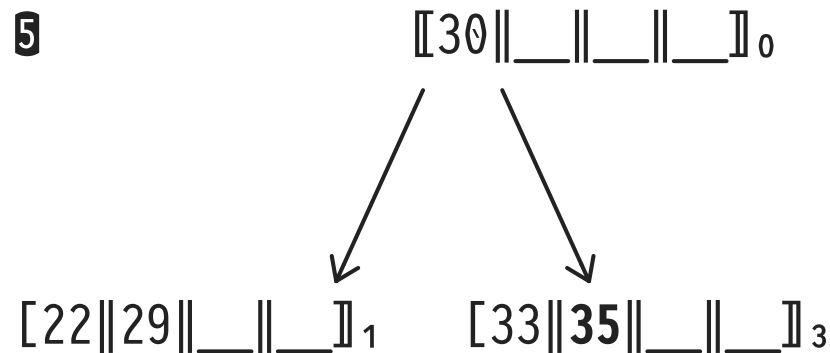
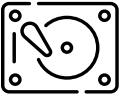


- 3 Delete entry with key $k = 24$
- Search(24) returns leaf 1
 - Leaf 1 has minimum occupancy, no sibling with spare entries

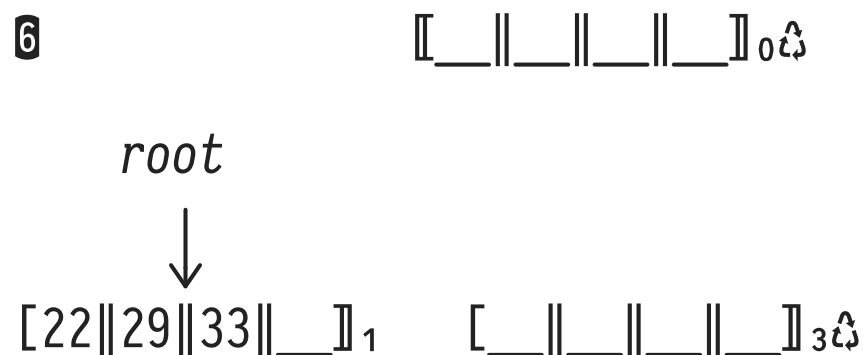


- 4 Merge leaf nodes 1 and 2, mark empty page 2 as garbage
- In parent 0, delete obsolete separator [27]

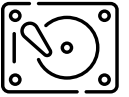
B+Tree Deletion and Leaf Node Merging (Empty Root)



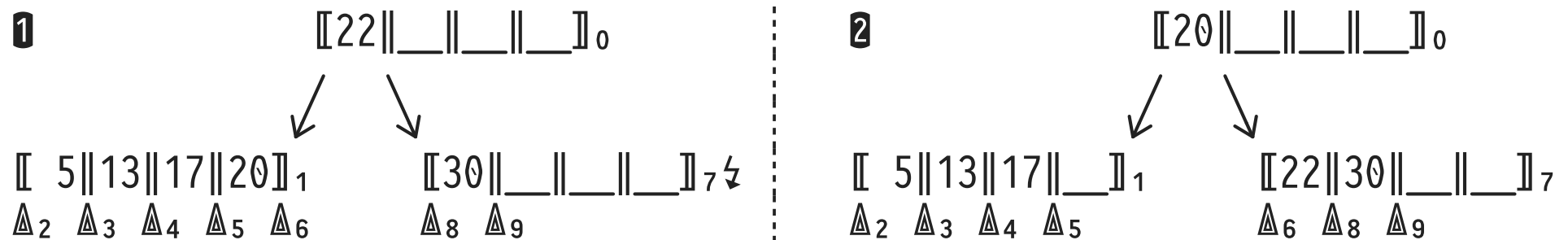
- 5 Delete entry with key $k = 35$
- Search(35) returns leaf 3
 - Leaf 3 has minimum occupancy, no sibling with spare entries



- 6 Merge leaf nodes 1 and 3, mark empty page 3 as garbage
- In parent 0, delete obsolete separator [30]
 - Old root empty (\Rightarrow garbage), mark page 1 as the new root
 - B+Tree height decreases



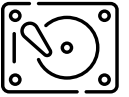
- **Redistribution** is also defined for **inner nodes**. Suppose we encounter underflow ❶ during \odot deletion propagation:



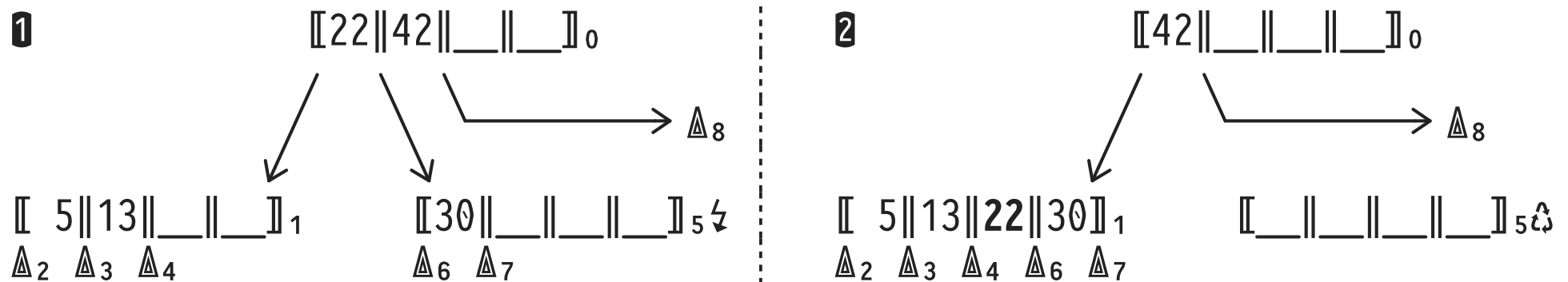
- Inner node 1 has two spare entries. “Rotate entry [20] through parent” to underflowed inner node 7.

NB: Semantics of subtree A_6 (holds index entries with $k \geq 20 \wedge k < 22$) are preserved.

B+Tree Deletion and Inner Node Merging



- Likewise, **inner nodes** may also be **merged**. The underflow in ❶ cannot be handled by redistribution:



- Note how the separator **22** has been **pulled down** from the parent to discriminate between subtrees Δ₄ and Δ₆:
 - Δ₄: $k \geq 13 \wedge k < 22$
 - Δ₆: $k \geq 22 \wedge k < 30$

8 | B+Trees: Key Compression



The higher the **fan-out** F , the more index entries fit in a B+Tree of fixed height. How to maximize F ?

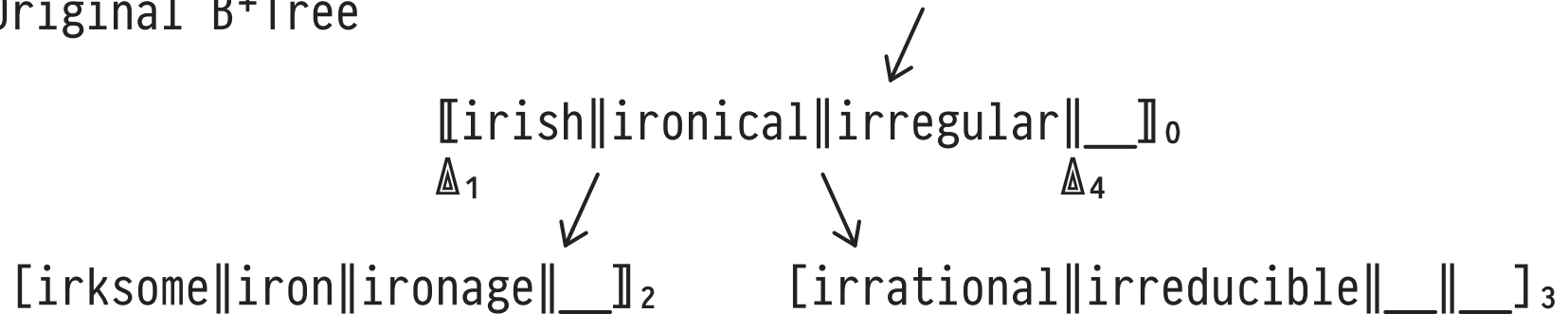
- For entries $\langle k, p \rangle$ in indexes over **text/char** columns, we may have $|k| \gg |p|$.⁹ Can we reduce the size of k ?
- 💡 **Search()** and **TreeInsert()** do *not* inspect the actual key values but only use $</\leq$ to direct tree traversals.
 - \Rightarrow May **shorten (truncate) string keys** as long as the ordering relation is preserved.
 - This applies to index entries in inner nodes only. Leaf level keys remain as is.

⁹ The implementation (thus size) of page pointers p is prescribed by the DBMS. Nothing to win here.

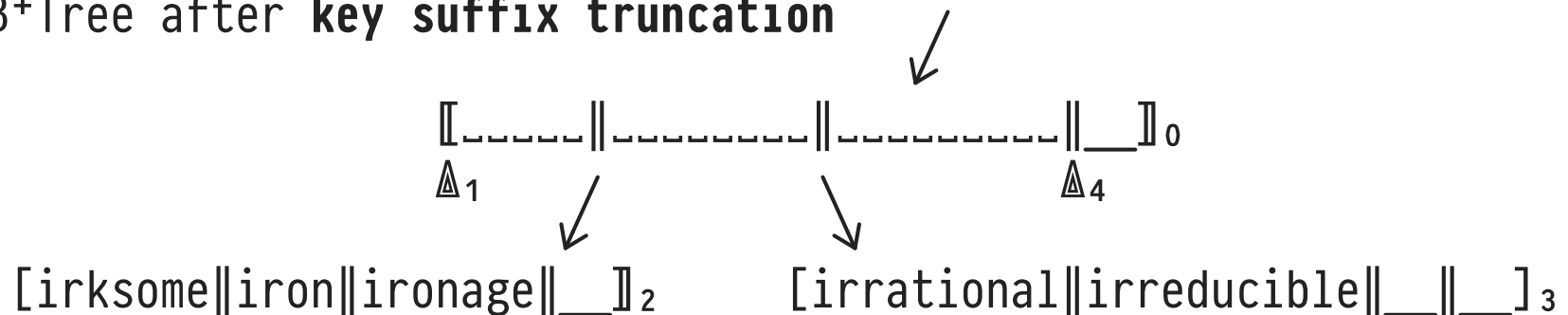
B+Trees: Key Suffix Truncation



1 Original B+Tree

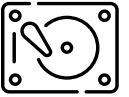


2 B+Tree after key suffix truncation



! While truncating, preserve the **separator** semantics.

B+Trees: Key Prefix Compression



Observation: string keys within a B+Tree inner node often **share a common prefix**.

- 💡 Store common prefix only once (e.g., as " k_0 ").
- Violating the 50% occupancy rule can help compression.

1 Original B+Tree

↓
[[irish||ironical||irregular||__]]₀
△₁ △₂ △₃ △₄

2 B+Tree after **key prefix compression**

↓
[ir+||ish||onical||regular||__]]₀
 k_0 △₁ △₂ △₃ △₄

9 | B+Tree Bulk Loading



Grab a hot cup of ☕ and start a war on Stack Overflow:¹⁰

Q: Which order of operations is better?

```
❶ CREATE TABLE T (...);  
❷ INSERT INTO T VALUES (<5 × 106 rows>);  
❸ CREATE INDEX I ON T USING btree (...);
```

or

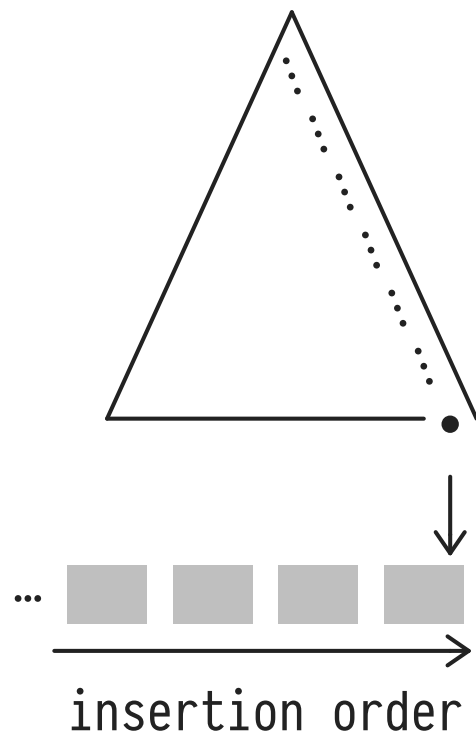
```
❶ CREATE TABLE T (...);  
❸ CREATE INDEX I ON T USING btree (...);  
❷ INSERT INTO T VALUES (<5 × 106 rows>);
```

¹⁰ See, for example, <https://stackoverflow.com/questions/5910486/indexes-on-a-table-database>

B+Tree Bulk Loading



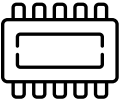
If insertions happen in index key order (i.e., ascending values of k), we observe a particular B+Tree access pattern:



- `TreeInsert()` will always traverse path \therefore , will always hit the rightmost leaf.
- ⇒ Fix rightmost leaf in buffer, insert next entry right there (*no* traversal from root). Node splits only occur along path \therefore .
- We effectively create a clustered index.

...  ... heap file (sorted on keys k)

insertion order

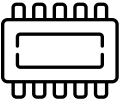


```
SELECT i.b, i.c  
FROM   indexed AS i  
WHERE  i.a = 42 [i.c = 0.42] -- either filter on i.a or i.c
```

Indexes in MonetDB play a secondary role and are *not* organized in tree shapes.

MMDBMSs try to exploit that data resides in directly-addressable memory and primarily aim to avoid access to separate index data structures (to avoid pointer chasing and potential cache misses).

Using **EXPLAIN** on Q_8 : Filter on Column a

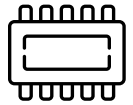


```
sql> EXPLAIN SELECT i.b, i.c
      FROM indexed AS i
      WHERE i.a = 42;

:
indexed :bat[:oid] := sql.tid(sql, "sys", "indexed");
a0       :bat[:int] := sql.bind(sql, "sys", "indexed", "a", 0:int);
p1       :bat[:oid] := algebra.thetaselect(a0, indexed, 42:int, "=="); ← ≡ a = 42
c0       :bat[:sht] := sql.bind(sql, "sys", "indexed", "c", 0:int);
c        :bat[:sht] := algebra.projection(p1, c0);
b0       :bat[:str] := sql.bind(sql, "sys", "indexed", "b", 0:int);
b        :bat[:str] := algebra.projection(p1, b0);
:
```

- MonetDB uses `algebra.thetaselect(..., 42:int, "==")` to implement the predicate filter.

Using **EXPLAIN** on Q_8 : Filter on Column c ¹¹

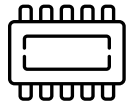


```
sql> EXPLAIN SELECT i.b, i.c
      FROM indexed AS i
      WHERE i.c = 0.42;
:
indexed :bat[:oid] := sql.tid(sql, "sys", "indexed");
c0       :bat[:sht] := sql.bind(sql, "sys", "indexed", "c", 0:int);
p1       :bat[:oid] := algebra.thetaselect(c0, indexed, 42:sht, "=="); ← ≡ c = 0.42
c        :bat[:sht] := algebra.projection(p1, c0);
b0       :bat[:str] := sql.bind(sql, "sys", "indexed", "b", 0:int);
b        :bat[:str] := algebra.projection(p1, b0);
:
```

- Plan is nearly identical (modulo access to the **a** BAT).
- MonetDB *appears* to use the same `algebra.thetaselect(..., 42:sht, "==")` MAL operation.

¹¹ Note how MonetDB maps the domain of type `numeric(3,2)` of column `c`, i.e., the set $N_{3,2} \equiv \{-9.99, \dots, 9.99\}$ with $|N_{3,2}| = 1999$, to a 16-bit value of type `:sht`. Nifty.

BAT Tail Properties

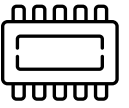


When MonetDB constructs a BAT t , a family of tail column **properties** $prop(t)$ is derived/maintained:¹²

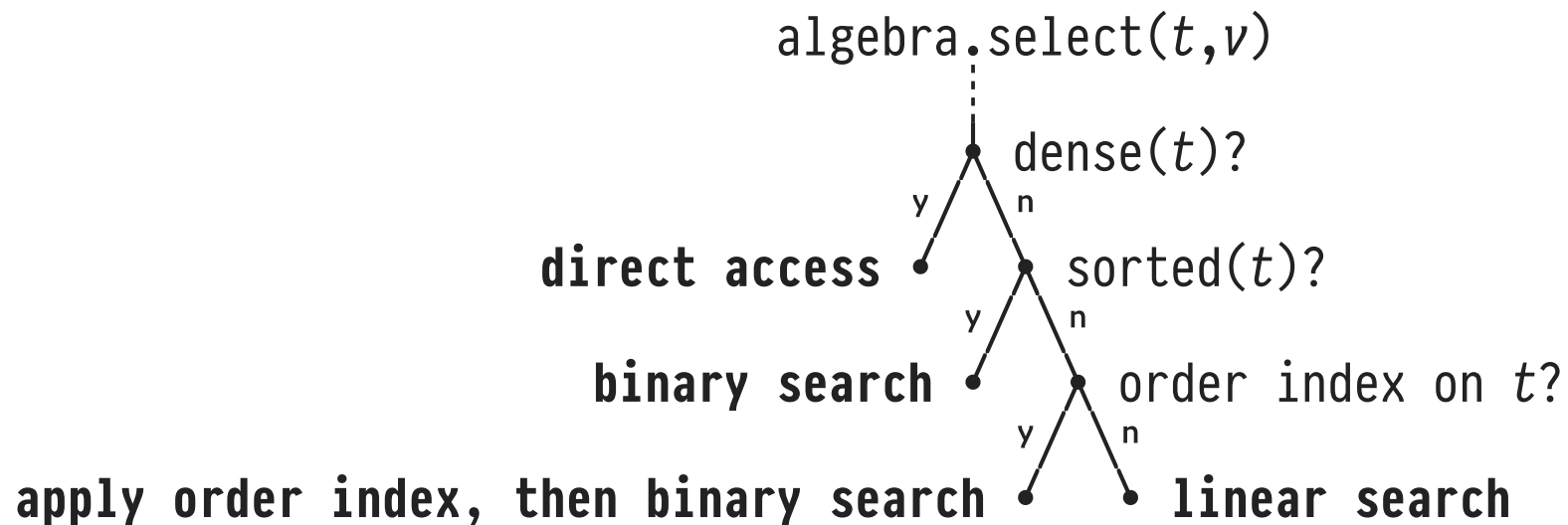
BAT Property $prop(t)$	Description
dense (tails of type <code>:oid</code> only)	ascending values, no gaps
key	unique values
sorted	ascending values
revsorted	descending values
nil/nonil	at least one/no <code>nil</code> value

- Use `bat.info(t)` to inspect current properties of t .
- ⚠ Incomplete: t 's tail may be sorted although `sorted(t) = false` (\Rightarrow but not \Leftrightarrow).

¹² Additional properties `nokey`, `nosorted`, `norevsorted` give “proofs” (tail positions) why property does not hold. Example: `nosorted = 3` \equiv tail value for row `300` < tail value for row `200`.

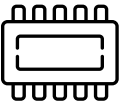


MAL operations inspect BAT properties at *query runtime*,
select one of several efficient implementations:

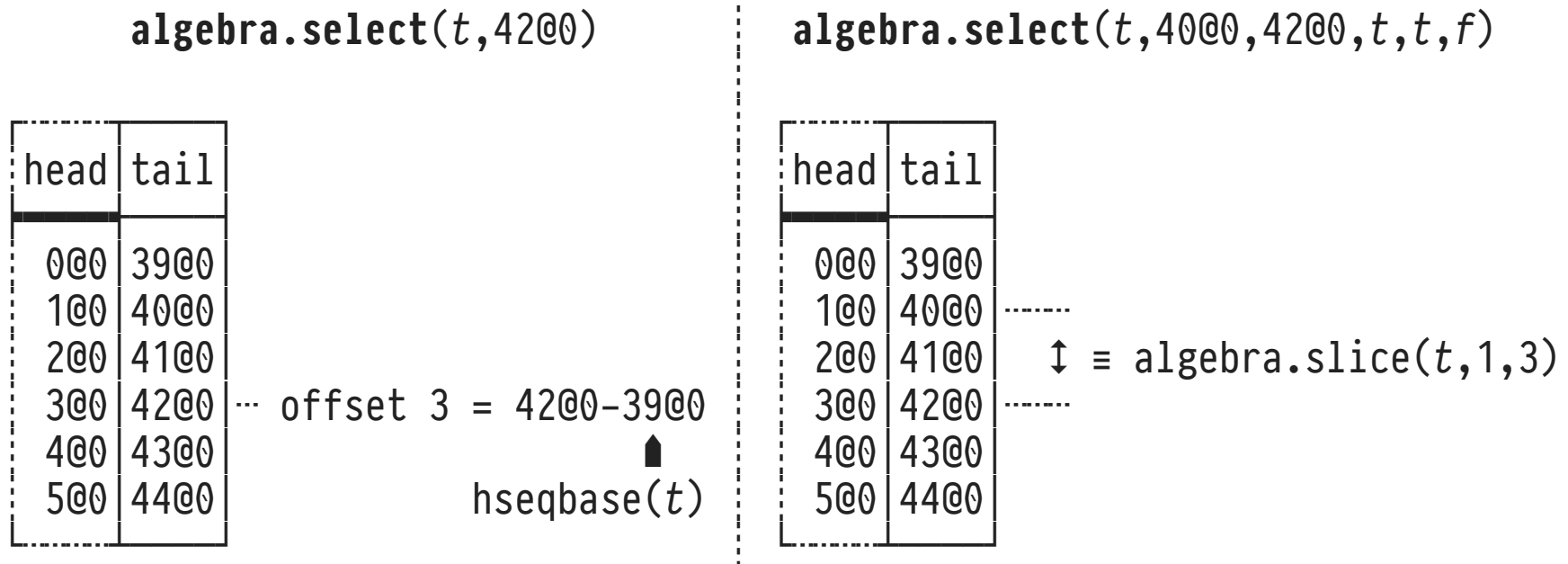


- This is coined **tactical optimization** (as opposed to strategical query optimization at *query compile time*).

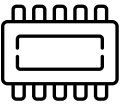
The Tactics of `algebra.select: dense(t)`



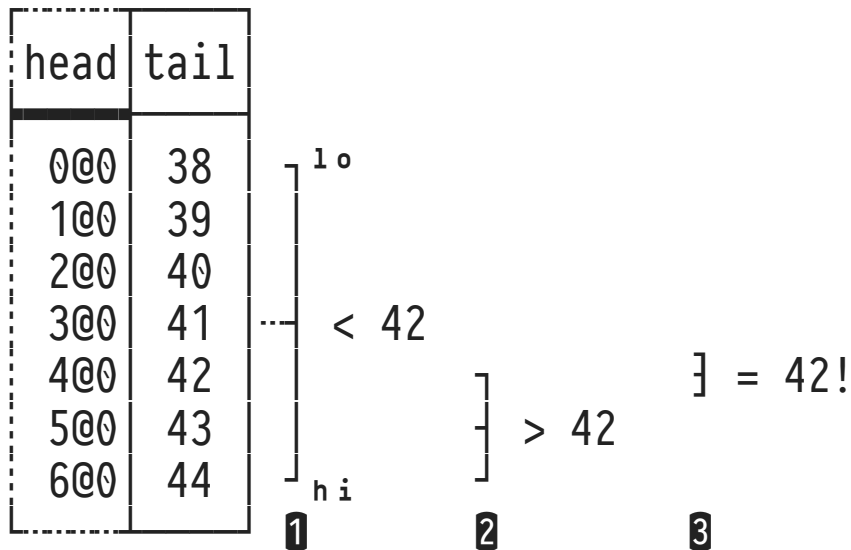
If input BAT `t` is **dense**, use **positional access** and **slicing** to evaluate equality and range selections:



The Tactics of `algebra.select: sorted(t)`



`algebra.select(t,42)`

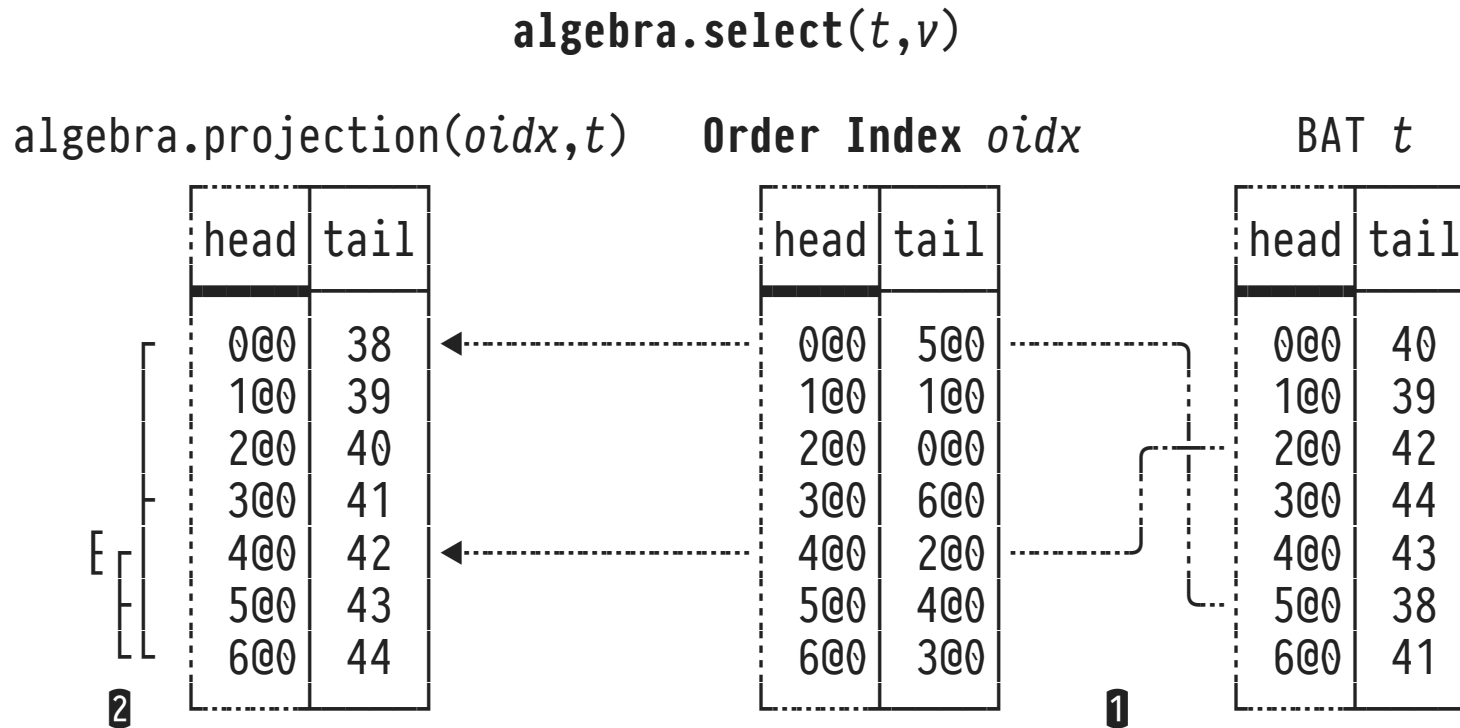
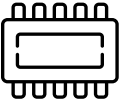


Binary Search:

- Test middle value (pivot) between limits lo and hi
- Recurse into upper or lower partition based on test
- Finishes in $\log_2(|t|)$ steps

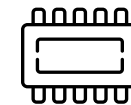
- **NB:** Unpredictable branches ($\leq 42?$) and jumps of pivot position less than ideal for CPU.

The Tactics of `algebra.select`: Order Indexes



- Row $[i@0, j@0] \in oidx$: value at offset *j* is *i*th largest in tail. Tactic: ① Apply *oidx*, ② then use binary search.

Creating Order Indexes (On the Fly)



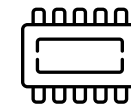
MonetDB may *automatically* create a temporary order index to support predicates $lo \leq a \leq hi$ or other order-sensitive queries (e.g., **ORDER BY**, **GROUP BY**).

- Check current properties of column BATs and presence of indexes in MonetDB system table **sys.storage**:

```
sql> SELECT column, sorted, revsorted, "unique", orderidx  
      FROM    sys.storage('sys', 'indexed');
```

column	sorted	revsorted	unique	orderidx
a	true	null	true	0
b	null	null	null	0
c	false	false	null	0

Creating Order Indexes (Manually)

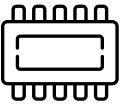


If this seems beneficial for the **query workload**, clients may *manually* create an order index.

- ⚠ Order indexes are **static** (i.e., not maintained under updates—costly) \Rightarrow underlying table must be *read-only*:

```
<create and populate table T>  
sql> ALTER TABLE T SET READ ONLY;  
sql> CREATE ORDERED INDEX I ON T(a);
```

- Order index *I* is made persistent (in a `*.torderidx` disk file) and will be used by future `algebra.select()`s on column *a*.

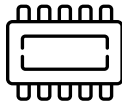


With **column cracking**,¹³ MonetDB introduced a **self-organizing** (partially) ordered index structure.

- A **cracker index** for column a is created/updated as a **by-product of processing range predicates** $lo \leq a \leq hi$.
 - In the cracker index, the a values $\in [lo, hi]$ are stored physically contiguous (yet unordered, in general).
- If the **query workload** focuses only on a subset of column a , that part is indexed with fine granularity (while the other parts remain largely non-indexed).

¹³ “Database Cracking”, S. Idreos, M. Kersten, S. Manegold. Proc. CIDR, Asilomar (CA, USA), 2007.

Column Cracking As a By-Product of Query Processing



1 BAT a

head	tail
0@0	17
1@0	3
2@0	8
3@0	6
4@0	2
5@0	15
6@0	13
7@0	4
8@0	12

2 Cracker BAT (Index)

head	tail
0@0	4
1@0	3
2@0	2
3@0	6
4@0	8
5@0	15
6@0	13
7@0	17
8@0	12

3 Cracker BAT (Index)

head	tail
0@0	2
1@0	3
2@0	4
3@0	6
4@0	8
5@0	12
6@0	13
7@0	17
8@0	15

$\xrightarrow{Q_i}$

$\xrightarrow{Q_j}$

$\leq 5 \quad s_1$

$> 5 \quad s_2$

$\geq 10 \quad s_3$

$\leq 3 \quad s_4$

$> 3 \quad s_5$

$> 5 \quad s_6$

$\geq 10 \quad s_7$

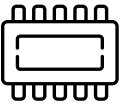
$\geq 14 \quad s_8$

• Q_i : ... **WHERE** $a > 5$ **AND** $a < 10$

• Q_j : ... **WHERE** $a > 3$ **AND** $a < 14$

Result: slice s_2

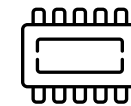
Result: slices $s_5 + s_6 + s_7$



- $\forall x \in s_i, y \in s_{i+1}: x < y$: a fully cracked column ($\forall i |s_i| = 1$) is completely ordered. This is uncommon (workload skew).
- First cracking step (❶→❷) copies source BAT. All further steps physically reorganize the cracker BAT.
- MonetDB implements slicing in terms of *views*¹⁴ of the cracker BAT, no data copying involved. Cost free.
- Physical cracker index reorganization (“tail shuffling”) can be efficiently performed *in-situ*.

¹⁴ A possible BAT view: (source BAT, first row, last row).

Cracker Index Reorganization For Predicate $a < hi$



Reorganize column vector $a[]$ between row offsets $start$ and end , relocate its elements *in-situ*:

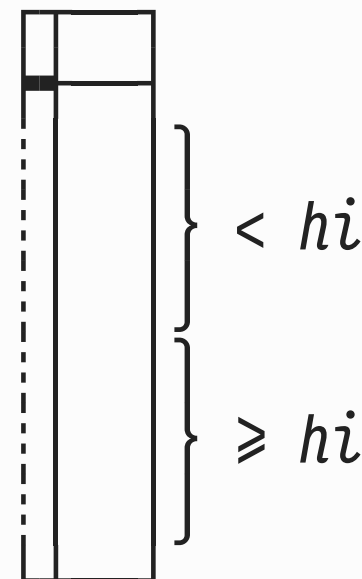
```
CrackInTwo( $a, start, end, hi$ ):
```

```

while ( $start < end$ )
    if ( $a[start] < hi$ )
        |  $start \leftarrow start + 1$ ;
    else
        while ( $a[end] \geq hi \wedge end > start$ )
            |  $end \leftarrow end - 1$ ;
        swap( $a[start], a[end]$ );  $**$ 
        |  $start \leftarrow start + 1$ ;
        |  $end \leftarrow end - 1$ ;

```

Result



- $**$ Either $a[start] \geq hi \wedge a[end] < hi$ or $start = end$.