# Database Management Systems

Lecture 7
Evaluating Relational Operators
Query Optimization

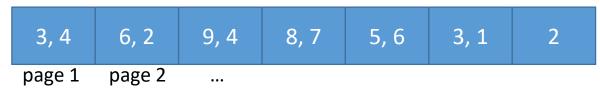
- running example schema
  - Students (SID: integer, SName: string, Age: integer)
  - Courses (CID: integer, CName: string, Description: string)
  - Exams (SID: integer, CID: integer, EDate: date, Grade: integer, FacultyMember: string)
  - Students
    - every record has 50 bytes
    - there are 80 records / page
    - 500 pages of Students tuples
  - Courses
    - every record has 50 bytes
    - there are 80 records / page
    - 100 pages of Courses tuples

- Exams
  - every record has 40 bytes
  - there are 100 records / page
  - 1000 pages of Exams tuples

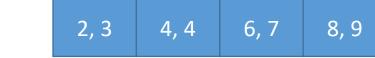
- Simple Two-Way Merge Sort: buffer pages are not used effectively
  - for instance, if 200 buffer pages are available, this algorithm still uses only 2 input buffers for passes 1, 2, ...
- generalize the Two-Way Merge Sort algorithm to effectively use the available main memory and minimize the number of passes
- input file to be sorted: N pages
- B buffer pages are available
- <u>pass 0</u>:
  - use B buffer pages
  - read in B pages at a time and sort them in memory
    - $=> \left| \frac{N}{B} \right|$  runs of B pages each (except for the last one, which may be

smaller)

consider again the input file in the previous example:



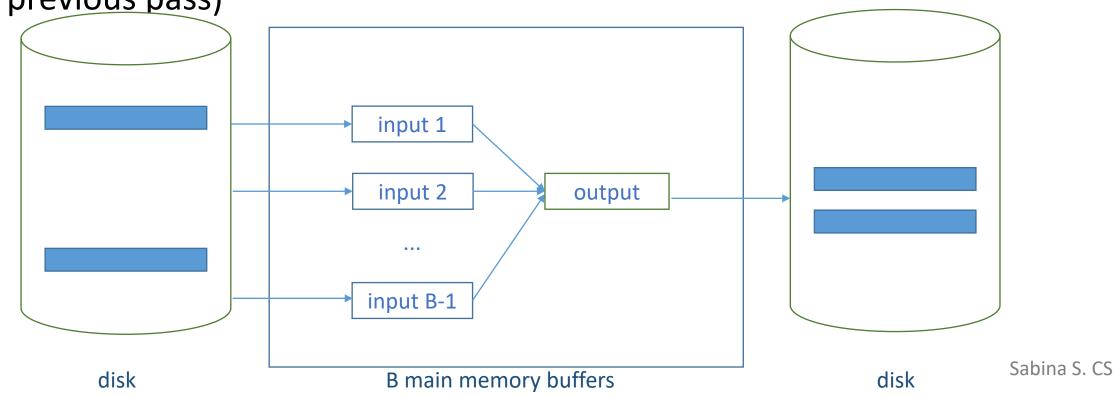
- N = 7 (number of pages in the file)
- B = 4 (there are 4 available buffer pages)
- pass 0 produces  $\left[\frac{N}{B}\right] = \left[\frac{7}{4}\right] = 2$  runs:
  - use all 4 buffer pages
  - read in 4 pages: 3, 4 6, 2 9, 4 8, 7
  - sort the pages in memory, write to disk a run that is 4 pages long:



- read in remaining 3 pages: 5,6 3,1 2
- sort pages in memory, write to disk run of 3 pages: 1,2

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- input file to be sorted: N pages
- B buffer pages are available
- pass 1, 2 ...:
  - use B-1 pages for input, and one page for output
  - perform a (B-1)-way merge in each pass (i.e., merge B-1 runs from the previous pass)



runs at the end of pass 0:

2, 3 4, 4 6, 7 8, 9

1, 2 3, 5 6

- pass 1
  - read in & merge the first B-1 = 4-1 = 3 runs from pass 0
  - pass 0 produced only 2 runs in this example; read in and merge these 2 runs:

=> run 1,2 2,3 3,4 4,5 6,6 7,8 9

- another example:
  - 5 buffer pages B = 5
  - sort file with 108 pages N = 108

#### pass 0

- use all 5 buffer pages
- read in the first 5 pages of the file, sort them in memory, write the resulting run to disk (5 pages long)
- read in the next 5 pages of the file, sort them in memory, write the resulting run to disk (5 pages long)
- •
- read in the remaining 3 pages of the file, sort them in memory, write the resulting run to disk (3 pages long)
- 21 runs are 5 pages long; 1 run is 3 pages long

- another example: B = 5, N = 108
- pass 0
  - at the end of pass 0 there are  $\left[\frac{N}{B}\right] = \left[\frac{108}{5}\right] = 22 \text{ runs}$
- pass 1
  - use B-1 = 5-1 = 4 pages for input, and one page for output
  - do a 4-way merge: read in and merge 4 runs from the previous pass
  - read in the first 4 runs from pass 0 (each run into an input buffer)
    - merge the runs and write to the output buffer
    - write the output buffer to disk one page at a time
    - => a run that is 20 pages long (4 runs from pass 0 times 5 pages per run)
  - read in the next 4 runs from pass 0; merge the runs and write to the output buffer; write the output buffer to disk one page at a time

=> another run (20 pages long)

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- another example: B = 5, N = 108
- pass 0
  - at the end of pass 0 there are 22 runs
- pass 1
  - read in the last 2 runs from pass 0 (one has 5 pages, the other one has 3 pages)
    - merge the runs and write to the output buffer; write the output buffer to disk one page at a time
    - => the last run (8 pages long)
  - at the end of pass 1 there are  $\left[\frac{22}{4}\right]$  = 6 runs
  - 5 runs are 20 pages long; 1 run is 8 pages long

- another example: B = 5, N = 108
- pass 1
  - at the end of pass 1 there are 6 runs
- pass 2
  - 4-way merge
  - read in the first 4 runs from pass 1
    - merge the runs and write to the output buffer; write the output buffer to disk one page at a time
    - => a run that is 80 pages long (4 runs from pass 1 times 20 pages per run)
  - read in the remaining 2 runs from pass 1 (20 and 8 pages, respectively)
     => a run that is 28 pages long
  - at the end of pass 2 there are  $\left[\frac{6}{4}\right] = 2$  runs

- another example: B = 5, N = 108
- pass 2
  - at the end of pass 2 there are 2 runs
- pass 3
  - read in the 2 runs from pass 2 and merge them
     => a run that is 108 pages long, representing the sorted file

- cost
  - N number of pages in the input file, B number of available pages in the buffer
  - in each pass: read / process / write each page
  - number of passes:  $\lceil log_{B-1}[N/B] \rceil + 1$
  - total cost:  $2 * N * \left( \left\lceil log_{B-1} \left\lceil \frac{N}{B} \right\rceil \right\rceil + 1 \right) I/Os$
- previous example: B = 5 and N = 108, with 4 passes over the data
  - cost:

• 
$$2*108*\left(\left\lceil log_{5-1}\left\lceil \frac{108}{5}\right\rceil\right\rceil + 1\right) = 216*\left(\left\lceil log_422\right\rceil + 1\right) = 216*4 = 864 \text{ I/Os}$$

- B buffer pages
- sort file with N pages

# Simple Two-Way Merge Sort

number of passes = 
$$\lceil log_2 N \rceil + 1$$

pass 
$$0 \Rightarrow \left[\frac{N}{B}\right]$$
 runs

number of passes = 
$$\left[log_{B-1}\left[\frac{N}{B}\right]\right] + 1$$

- External Merge Sort reduced number of:
  - runs produces by the 1<sup>st</sup> pass
  - passes over the data
- B is usually large => significant performance gains

# External Merge Sort – number of passes for different values of N and B

N	B = 3	B = 5	B = 9	B = 17	B = 129	B = 257
100	7	4	3	2	1	1
1,000	10	5	4	3	2	2
10,000	13	7	5	4	2	2
100,000	17	9	6	5	3	3
1,000,000	20	10	7	5	3	3
10,000,000	23	12	8	6	4	3

100,000,000

1,000,000,000

- equality join, one join column:  $E \otimes_{i=j} S$  (i<sup>th</sup> column's value in  $E = j^{th}$  column's value in S)
- sort E and S on the join column (if not already sorted):
  - for instance, by using External Merge Sort
  - => partitions = groups of tuples with the same value in the join column
- merge E and S; look for tuples e in E, s in S such that  $e_i = s_i$ :
  - while current e<sub>i</sub> < current s<sub>i</sub>
    - advance the scan of E
  - while current e<sub>i</sub> > current s<sub>i</sub>
    - advance the scan of S
  - if current  $e_i$  = current  $s_j$ 
    - output joined tuples  $\langle e, s \rangle$ , where e and s are in the current partition (i.e., they have the same value in the i<sup>th</sup> and j<sup>th</sup> column, respectively)
    - there could be multiple tuples in E with the same value in the i<sup>th</sup> column as the current tuple *e* (same is true for S)

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• partitions are illustrated on tables Students and Exams below (join column SID in both tables):

SID	SName	Age
20	Ana	20
30	Dana	20
40	Dan	20
45	Daniel	20
50	Ina	20

SID	CID	<b>EDate</b>	Grade	<b>FacultyMember</b>
30	2	20/1/2018	10	Ionescu
30	1	21/1/2018	9.99	Рор
45	2	20/1/2018	9.98	Ionescu
45	1	21/1/2018	9.98	Рор
45	3	22/1/2018	10	Stan
50	2	20/1/2018	10	Ionescu

 during the merging phase, E is scanned once; every partition in S is scanned as many times as there are matching tuples in the corresponding partition in E

	 i <sup>th</sup> column	 	j <sup>th</sup> column		
	1		2		
	3		3	tion D	S
E	3		3 Parti	tion P	
	3		4		
	8				

- for instance, partition P in the above table S is scanned 3 times, once per matching tuple in the corresponding partition in E
- there are 6 output joined tuples <e, s> for partition P
- this algorithm avoids the enumeration of the cross-product: tuples in a partition in E are compared only with the S tuples in the same partition!
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- <u>cost</u>:
  - sorting E
    - cost: O(MlogM)
  - sorting S
    - cost: O(NlogN)
  - cost of merging: M + N I/Os, assuming partitions in S are scanned only once
    - worst-case scenario: O(M \* N) I/Os (when all records in E and S have the same value in the join column)

\* E - M pages; S - N pages\*

# Sort-Merge Join (Exams $\bigotimes_{Exams,SID=Students,SID}$ Students)

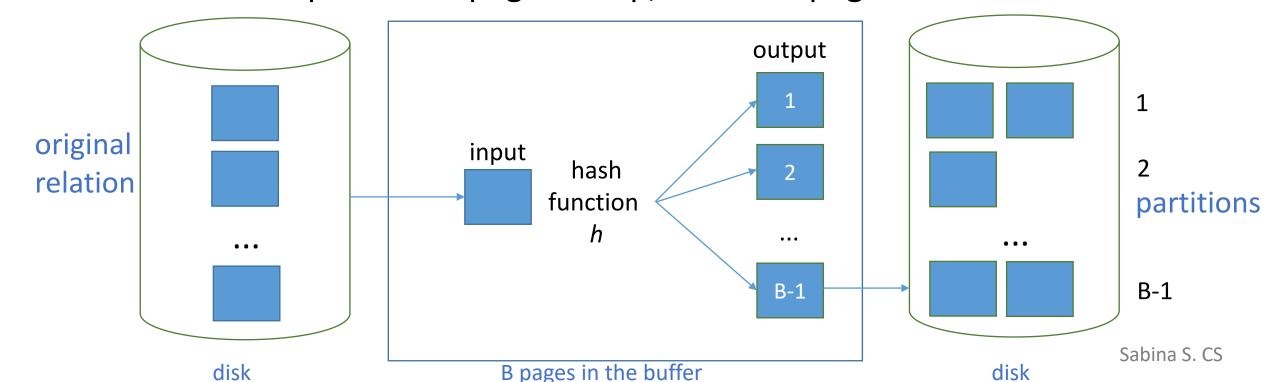
- 100 buffer pages
  - sort Exams
    - 2 passes => cost: 2 \* 2 \* 1000 = 4000 I/Os
  - sort Students
    - 2 passes => cost: 2 \* 2 \* 500 = 2000 I/Os
  - merging phase
    - cost: 1000 + 500 = 1500 I/Os
  - total cost: 4000 + 2000 + 1500 = 7500 I/Os
    - similar to the cost of Block Nested Loops Join
- <u>35 buffer pages</u>, <u>300 buffer pages</u> cost remains unchanged (need 2 passes to sort Exams, 2 passes to sort Students)
  - ex: compute cost of BNLJ and compare
- \* E M pages, p<sub>F</sub> records / page \* \* 1000 pages \* \* 100 records / page\*

<u>Hash Join</u> - equality join, one join column:  $E \otimes_{i=j} S$ 

• <u>phases</u>: partitioning (building phase) & probing (matching phase)

- partitioning phase:
  - there are B pages available in the buffer:
    - use one page as the input buffer page
    - and the remaining B-1 pages as output buffer pages
  - choose a hash function h that distributes tuples uniformly to one of B-1 partitions
  - hash E and S on the join column (the i<sup>th</sup> column of E, the j<sup>th</sup> column of S)
    with the same hash function h

- hash E on the join column with hash function h (similarly for S):
  - for each tuple e in E, compute  $h(e_i)$  ( $e_i$ : the value of the  $i^{th}$  column in tuple e)
  - add tuple e to the output buffer page that it is hashed to by h (buffer page  $h(e_i)$ )
  - when an output buffer page fills up, flush the page to disk

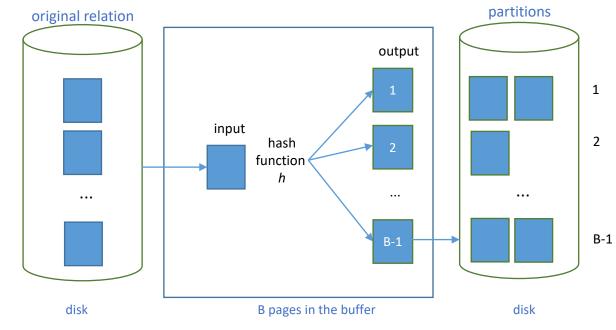


- partitioning phase => partitions of E ( $E_1$ ,  $E_2$ , etc.) and S ( $S_1$ ,  $S_2$ , etc.) on disk
- <u>partition</u> = collection of tuples that have the same hash value
- tuples in partition  $E_1$  can only join with tuples in partition  $S_1$  (they cannot join with tuples in partitions  $S_2$  or  $S_3$ , for instance, since these tuples have a different hash value)

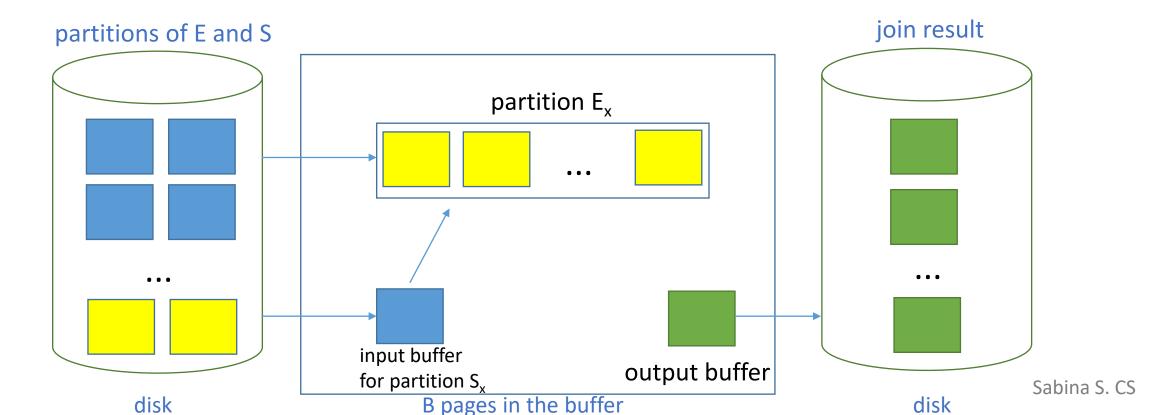
so to compute the join, we need to scan E and S only once (provided any

partition of E fits in main memory)

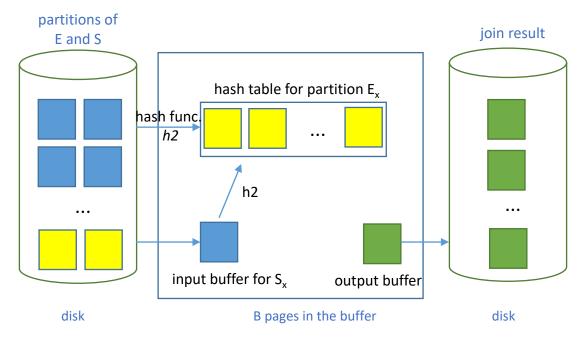
when reading in a partition  $E_k$  of  $E_k$  we must scan only the corresponding partition  $S_k$  of S to find matching tuples (compare tuples e in  $E_k$  with tuples s in  $S_k$  to test the join condition value of  $i^{th}$  column in E = value of  $i^{th}$  column in S)



- probing phase:
  - read in a partition of the smaller relation (e.g., E) and scan the corresponding partition of S for matching tuples
  - use one page as the input buffer for S, one page as the output buffer, and the remaining pages to read in partitions of E



- probing phase:
  - in practice, to reduce CPU costs, an in-memory hash table is built, using a different function *h2*, for the E partition
- consider a partition E<sub>x</sub> of E
- build in-memory hash table for E<sub>x</sub> using hash function h2 (the function is applied to the join column of E)
- for each tuple s in partition  $S_x$ , find matching tuples in the hash table using the hash value  $h2(s_i)$



- result tuples <e, s> are written to output buffer
- once partitions  $E_x$  and  $S_x$  are processed, the hash table is emptied (to prepare for the next partition)

- <u>cost</u>:
  - partitioning: both E and S are read and written once => cost: 2\*(M+N) I/Os
  - probing: scan each partition once => cost: M+N I/Os
  - => total cost: 3\*(M+N) I/Os
    - assumption: each partition fits into memory during probing
  - 3\*(1000 + 500) = 4500 I/Os
- \* E M pages, p<sub>F</sub> records / page \* \* 1000 pages \* \* 100 records / page\*
- partition overflow an E partition does not fit in memory during probing: apply hash join technique recursively:
  - divide E, S into subpartitions
  - join subpartitions pairwise
  - if subpartitions don't fit in memory, apply hash join technique recursively

- memory requirements objective: partition in E fits into main memory (S - similarly)
  - B buffer pages; need one input buffer => maximum number of partitions: B-1
  - size of largest partition: B-2 (need one input buffer for S, one output buffer)
  - assume uniformly sized partitions => size of each E partition: M/(B-1)
  - => M/(B-1) < B-2 => we need approximately B >  $\sqrt{M}$
  - if an in-memory hash table is used to speed up tuple matching => need a little more memory (because the hash table for a collection of tuples will be a little larger than the collection itself)

\* E - M pages, p<sub>F</sub> records / page \* \* 1000 pages \* \* 100 records / page\*

#### general join conditions

- <u>equalities</u> over several attributes
  - E.SID = S.SID AND E.attrE = S.attrS
    - index nested loops join
      - Exams inner relation:
        - build index on Exams with search key <SID, attrE> (if not already created)
        - can also use index on SID or index on attrE
      - Students inner relation (similar)
    - sort-merge join
      - sort Exams on <SID, attrE>, sort Students on <SID, attrS>
    - hash join
      - partition Exams on <SID, attrE>, partition Students on <SID, attrS>
    - other join algorithms
      - essentially unaffected

#### general join conditions

- inequality comparison
  - E.attrE < S.attrS</li>
    - index nested loops join: B+ tree index required
    - sort-merge join: not applicable
    - hash join: not applicable
    - other join algorithms: essentially unaffected
- \* no join algorithm is uniformly superior to others
- choice of a good algorithm depends on:
  - size(s) of:
    - joined relations
    - buffer pool
  - available access methods

#### Q:

```
SELECT *
FROM Exams E
WHERE E.FacultyMember = 'Ionescu'
```

- use information in the selection condition to reduce the number of retrieved tuples
- e.g., |Q| = 4 (result set has 4 tuples), there's a B+ tree index on FacultyMember
  - it's expensive to scan E (1000 I/Os) to evaluate the query
  - should use the index instead
- selection algorithms based on the following techniques:
  - iteration, indexing

- simple selections
  - $\sigma_{E.attr\ op\ val}(E)$
- no index on attr, data not sorted on attr
  - must scan E and test the condition for each tuple
  - access path: file scan
  - => cost: M I/Os = 1000 I/Os
- no index, sorted data (E physically sorted on attr)
  - binary search to locate 1<sup>st</sup> tuple that satisfies condition and
  - scan E starting at this position until condition is no longer satisfied
  - access method: sorted file scan

Review lecture notes on *Relational Algebra, Indexes, DB – Physical Structure (Databases* course)

- simple selections
  - $\sigma_{E.attr\ op\ val}(E)$
- no index, sorted data (E physically sorted on attr)=> cost:
  - binary search: O(log<sub>2</sub>M)
  - scan cost: varies from 0 to M
  - binary search on E
    - $\log_2 1000 \approx 10 \text{ I/Os}$

- simple selections
  - $\sigma_{E.attr\ op\ val}(E)$
- B+ tree index on attr
  - \* search tree to find 1<sup>st</sup> index entry pointing to a qualifying E tuple
    - cost: typically 2, 3 I/Os
  - \* scan leaf pages to retrieve all qualifying entries
    - cost: depends on the number of qualifying entries
  - \* for each qualifying entry retrieve corresponding tuple in E
    - cost: depends on the number of tuples and the nature of the index (clustered / unclustered)

- simple selections
  - $\sigma_{E.attr\ op\ val}(E)$
- B+ tree index on attr
  - assumption
    - indexes use a2 or a3
    - a1-based index => data entry contains the data record => the cost of retrieving records = the cost of retrieving the data entries!
  - access path: B+ tree index
    - clustered index:
      - best access path when op is not equality
      - good access path when op is equality

- simple selections:  $\sigma_{E.attr\ op\ val}(E)$
- B+ tree index on attr

```
Q
SELECT *
FROM Exams E
WHERE E.FacultyMember < 'C%'
```

- names uniformly distributed with respect to 1<sup>st</sup> letter
- $\Rightarrow$  |Q|  $\approx$  10,000 tuples = 100 pages
- clustered B+ tree index on FacultyMember
- => cost of retrieving tuples: ≈ 100 I/Os (a few I/Os to get from root to leaf)
- non-clustered B+ tree index on FacultyMember
- => cost of retrieving tuples: up to 1 I/O per tuple (worst case) => up to 10.000 I/Os
- \* E M pages, p<sub>E</sub> records / page \* \* 1000 pages \* \* 100 records / page \* sabina \$ 0

- simple selections:  $\sigma_{E.attr\ op\ val}(E)$
- B+ tree index on attr

```
SELECT *
FROM Exams E
WHERE E.FacultyMemger < 'C%'
```

- non-clustered B+ tree index on FacultyMember
  - refinement sort rids in qualifying data entries by page-id
     => a page containing qualifying tuples is retrieved only once
    - cost of retrieving tuples: number of pages containing qualifying tuples (but such tuples are probably stored on more than 100 pages)
- range selections
  - non-clustered indexes can be expensive
  - could be less costly to scan the relation (in our example: 1000 I/Os)

- general selections
  - selections without disjunctions
- C CNF condition without disjunctions
  - evaluation options:
  - 1. use the most selective access path
    - if it's an index I:
      - apply conjuncts in C that match I
      - apply rest of conjuncts to retrieved tuples
    - example
      - c < 100 AND a = 3 AND b = 5</li>
        - can use a B+ tree index on c and check a = 3 AND b = 5 for each retrieved tuple
        - can use a hash index on a and b and check c < 100 for each retrieved tuple

- general selections selections without disjunctions
  - evaluation options:
  - 2. use several indexes when several conjuncts match indexes using a2 / a3
    - compute sets of rids of candidate tuples using indexes
    - intersect sets of rids, retrieve corresponding tuples
    - apply remaining conjuncts (if any)
    - example: *c* < 100 AND *a* = 3 AND *b* = 5
      - use a B+ tree index on c to obtain rids of records that meet condition  $c < 100 \, (R_1)$
      - use a hash index on a to retrieve rids of records that meet condition a = 3 ( $R_2$ )
      - compute  $R_1 \cap R_2 = R_{int}$
      - retrieve records with rids in  $R_{int}$  (R)
      - check b = 5 for each record in R

- general selections
  - selections with disjunctions
- C CNF condition with disjunctions, i.e., some conjunct *J* is a disjunction of terms
  - if some term *T* in *J* requires a file scan, testing *J* by itself requires a file scan
    - example:  $a < 100 \lor b = 5$ 
      - hash index on b, hash index on c
    - => check both terms using a file scan (i.e., best access path: file scan)
  - compare with the example below:
    - $(a < 100 \lor b = 5) \land c = 7$
    - hash index on b, hash index on c
  - => use index on c, apply  $a < 100 \lor b = 5$  to each retrieved tuple (i.e., most selective access path: index)

- general selections
  - selections with disjunctions
- C CNF condition with disjunctions
  - every term T in a disjunction matches an index
  - => retrieve tuples using indexes, compute union
  - example
    - $a < 100 \lor b = 5$
    - B+ tree indexes on a and b
    - use index on a to retrieve records that meet condition  $a < 100 (R_1)$
    - use index on b to retrieve records that meet condition  $b = 5 (R_2)$
    - compute  $R_1 \cup R_2 = R$
    - if all matching indexes use a2 or a3 => take union of rids, retrieve corresponding tuples

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