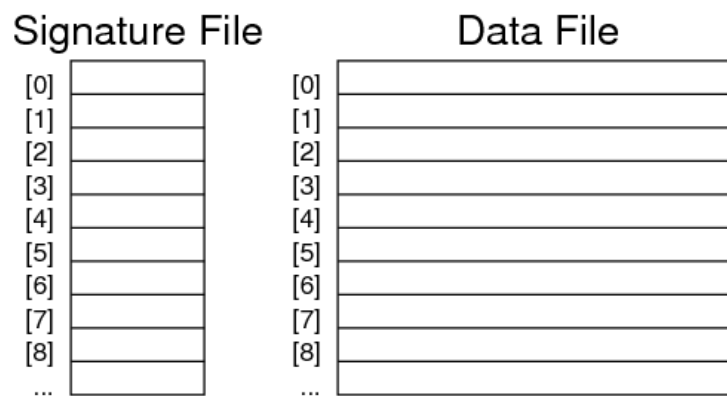


SIMC Indexing

- Signature-based indexing
- Superimposed Codewords (SIMC)
- SIMC Example
- SIMC Queries
- Example SIMC Query
- SIMC Parameters
- Query Cost for SIMC
- Page-level SIMC
- Bit-sliced SIMC
- Comparison of Approaches
- Signature-based Indexing in PostgreSQL

❖ Signature-based indexing

Reminder: file organisation for signature indexing (two files)



One signature slot per tuple slot; unused signature slots are zeroed.

❖ Superimposed Codewords (SIMC)

In a superimposed codewords (simc) indexing scheme

- a tuple descriptor is formed by overlaying attribute codewords
- each codeword is m bits long and has k bits set to 1

A tuple descriptor $desc(t)$ is

- a bit-string, m bits long, where $j \leq nk$ bits are set to 1
- $desc(t) = cw(A_1) \text{ OR } cw(A_2) \text{ OR } \dots \text{ OR } cw(A_n)$

Method (assuming all n attributes are used in descriptor):

```
bits desc = 0
for (i = 1; i <= n; i++) {
    bits cw = codeword(A[i], m, k)
    desc = desc | cw
}
```

❖ SIMC Example

Consider the following tuple (from bank deposit database)

Branch	AcctNo	Name	Amount
Perryridge	102	Hayes	400

It has the following codewords/descriptor (for $m = 12$, $k = 2$)

A_i	$cw(A_i)$
Perryridge	010000000001
102	000000000011
Hayes	000001000100
400	000010000100
$desc(t)$	010011000111

❖ SIMC Queries

To answer query q in SIMC

- first generate $desc(q)$ by OR-ing codewords for known attributes
- then attempt to match $desc(q)$ against all signatures in sig file

E.g. consider the query (**Perryridge**, ?, ?, ?).

A_i	$cw(A_i)$
Perryridge	010000000001
?	000000000000
?	000000000000
?	000000000000
$desc(q)$	010000000001

❖ SIMC Queries (cont)

Once we have a query descriptor, we search the signature file:

```
pagesToCheck = {}  
// scan  $r$  signatures  
for each descriptor  $D[i]$  in signature file {  
    if (matches( $D[i]$ , desc( $q$ ))) {  
        pid = pageOf(tupleID( $i$ ))  
        pagesToCheck = pagesToCheck  $\cup$  pid  
    }  
}  
// then scan  $b_{sq} = b_q + \delta$  pages to check for matches
```

Matching can be implemented efficiently ...

```
#define matches(sig, qdesc) ((sig & qdesc) == qdesc)
```

❖ Example SIMC Query

Consider the query and the example database:

Signature	Deposit Record
0 1 000000000 1	(Perryridge,?,?,?)
10010100100 1	(Brighton,217,Green,750)
0 1 001100011 1	(Perryridge,102,Hayes,400)
10100100100 1	(Downtown,101,Johnshon,512)
1011000000 1	(Mianus,215,Smith,700)
0 1 010101010 1	(Clearview,117,Throggs,295)
1001010100 1	(Redwood,222,Lindsay,695)

Gives two matches: one **true match**, one **false match**.

❖ SIMC Parameters

False match probability p_F = likelihood of a false match

How to reduce likelihood of false matches?

- use different hash function for each attribute (h_i for A_i)
- increase descriptor size (m)
- choose k so that \approx half of bits are set

Larger m means larger signature file \Rightarrow read more signature data.

Having k too high \Rightarrow increased overlapping.

Having k too low \Rightarrow increased hash collisions.

❖ SIMC Parameters (cont)

How to determine "optimal" m and k ?

1. start by choosing acceptable p_F
(e.g. $p_F \leq 10^{-4}$ i.e. one false match in 10,000)
2. then choose m and k to achieve no more than this p_F .

Formulae to derive m and k given p_F and n :

$$k = 1/\log_e 2 \cdot \log_e (1/p_F)$$

$$m = (1/\log_e 2)^2 \cdot n \cdot \log_e (1/p_F)$$

Formula from Bloom (1970), "Space/Time Trade-offs in Hash Coding with Allowable Errors", Communications of the ACM, 13 (7): 422-426

❖ Query Cost for SIMC

Cost to answer pmr query: $Cost_{pmr} = b_D + b_{sq}$

- read r descriptors on b_D descriptor pages
- then read b_{sq} data pages and check for matches

$$b_D = \text{ceil}(r/c_D) \text{ and } c_D = \text{floor}(B/\text{ceil}(m/8))$$

$$\text{E.g. } m=64, B=8192, r=10^4 \Rightarrow c_D = 1024, b_D=10$$

b_{sq} includes pages with r_q matching tuples and r_F false matches

Expected false matches = $r_F = (r - r_q) \cdot p_F \approx r \cdot p_F$ if $r_q \ll r$

E.g. Worst $b_{sq} = r_q + r_F$, Best $b_{sq} = 1$, Avg $b_{sq} = \text{ceil}(b(r_q + r_F)/r)$

❖ Page-level SIMC

SIMC has one descriptor per tuple ... potentially inefficient.

Alternative approach: one descriptor for each data page.

Every attribute of every tuple in page contributes to descriptor.

Size of page descriptor (PD) (clearly larger than tuple descriptor):

- use above formulae but with $c.n$ "attributes"

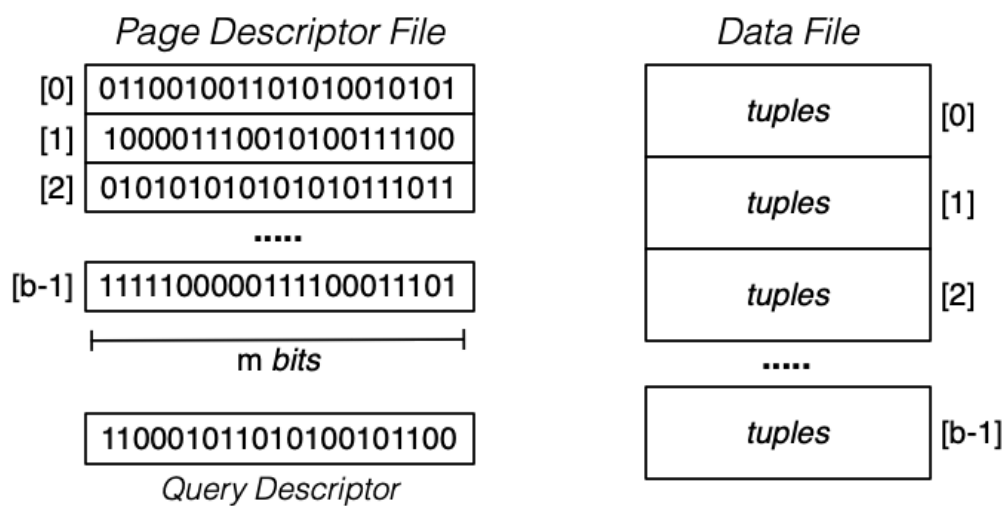
E.g. $n = 4, c = 64, p_F = 10^{-3} \Rightarrow m_p \approx 3680\text{bits} \approx 460\text{bytes}$

Typically, pages are 1..8KB $\Rightarrow 8..64$ PD/page (c_{PD}).

E.g. $m_p \approx 460, B = 8192, c_{PD} \approx 17$

❖ Page-level SIMC (cont)

File organisation for page-level superimposed codeword index



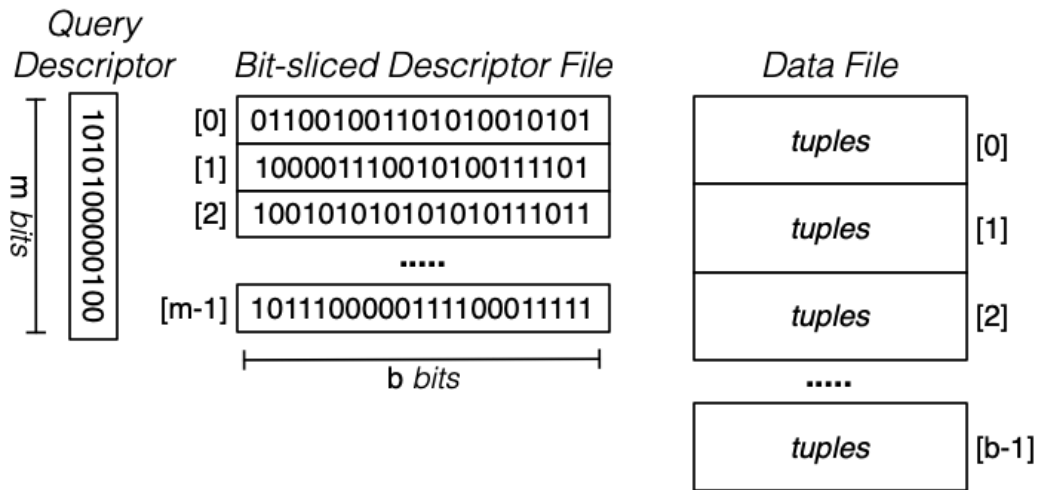
❖ Page-level SIMC (cont)

Algorithm for evaluating *pmr* query using page descriptors

```
pagesToCheck = {}  
// scan  $b_{m_p}$ -bit page descriptors  
for each descriptor  $D[i]$  in signature file {  
    if (matches( $D[i]$ , desc( $q$ ))) {  
        pid = i  
        pagesToCheck = pagesToCheck  $\cup$  pid  
    }  
}  
// read and scan  $b_{sq}$  data pages  
for each pid in pagesToCheck {  
    Buf = getPage(dataFile, pid)  
    check tuples in Buf for answers  
}
```

❖ Bit-sliced SIMC

Improvement: store b m -bit page descriptors as m b -bit "bit-slices"



❖ Bit-sliced SIMC (cont)

Algorithm for evaluating *pmr* query using bit-sliced descriptors

```
matches = ~0    //all ones
// scan m r-bit slices
for each bit i set to 1 in desc(q) {
    slice = fetch bit-slice i
    matches = matches & slice
}
for each bit i set to 1 in matches {
    fetch page i
    scan page for matching records
}
```

Effective because *desc(q)* typically has less than half bits set to 1

❖ Comparison of Approaches

Tuple-based

- r signatures, m -bit signatures, k bits/attribute
- read all pages of signature file in filtering for a query

Page-based

- b signatures, m_p -bit signatures, k bits/attribute
- read all pages of signature file in filtering for a query

Bit-sliced

- m signatures, b -bit slices, k bits/attribute
- read less than half of the signature file in filtering for a query

All signature files are roughly the same size, for a given p_F

❖ Signature-based Indexing in PostgreSQL

PostgreSQL supports signature based indexing via the **bloom** module

(Signature-based indexes like this are often called **Bloom filters**)

Creating a Bloom index

```
create index Idx on R using bloom (a1,a2,a3)
with (length=64, coll=3, col2=4, col3=2);
```

Example: 10000 tuples, query = select * from R where a1=55 and a2=42, no matching tuples, random numeric values for attributes

No indexes ... execution time 15ms

B-tree index on all attributes ... execution time 12ms

Bloom index ... execution time 0.4ms

For more details, see PostgreSQL doc, Appendix F.5

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