

**DCGI**

**KATEDRA POČÍTAČOVÉ GRAFIKY A INTERAKCE**

# **MODERN ALGORITHMS (not only in computational geometry)**

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<https://cw.felk.cvut.cz/doku.php/courses/a4m39vg/start>

Based on [Kolingerova], [Brönnimann], and [Muthukrishnan]

Version from 2.1.2019

# Modern algorithms

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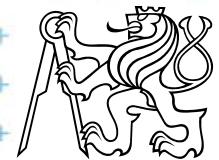
1. Computational geometry today
2. Space efficient algorithms  
(In-place / in situ algorithms)
3. Data stream algorithms
4. Randomized algorithms



# Computational geometry today

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- Popular: beauty as discipline, wide applicability
- Started in 2D with linear objects (points, lines,...), now 3D and nD, hyperplanes, curved objects,...
- Shift **from** purely mathematical approach and asymptotical optimality ignoring singular cases
- **to** practical algorithms, simpler data structures and robustness => **algorithms and data structures provable efficient in realistic situations** (application dependent)



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# Space efficient algorithms



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Felkel: Computational geometry

(4)



# Space efficient algorithms

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- output is in the same location as the input and
- need only a small amount of additionally memory
  - *in-place* –  $O(1)$  extra storage  
sometimes including  $O(\log n)$  bits for indice
  - *in situ* –  $O(\log n)$  extra storage



# Space efficient algorithms - practical advantages

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- Allow for processing larger data sets
  - Algorithms with separate input and output need space for  $2n$  points to store –  $O(n)$  extra space
  - Space efficient algs. –  $n$  points +  $O(1)$  or  $O(\log n)$  space
- Greater locality of reference
  - Practical for modern HW with memory hierarchies (e.g., main RAM – ram on chip – registers, caches, disk latency, network latency )
- Less prone to failure
  - no allocation of large amounts of memory, which can fail at run time
  - good for mission critical applications

■ Less memory => faster program



# Ex: String reverse

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```
function reverse(a[0..n])  
    allocate b[0..n]  
    for i from 0 to n  
        b[n-i] = a[i]  
    return b
```

×

```
function reverseInPlace(a[0..n])  
    for i from 0 to floor(n/2)  
        swap (a[n-i], a[i])
```



# In-place sorting

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- In array – continuous block in memory
  - $n^{\text{th}}$  element in  $O(1)$  time
  - Select sort, insert sort ... in-place,  
 $O(1)$  additional memory,  $O(n^2)$  time
  - Heapsort – in-place,  $O(1)$  add. memory,  $O(n \log n)$  time
  - Quicksort – in-situ,  $O(\log n)$  add. memory for recursion
  - Mergesort – not in-place, not in-situ,  $O(n)$  add. memory
- In list – linked lists in dynamical memory
  - $n^{\text{th}}$  element in  $O(n)$  time
  - Mergesort – in-situ,  $O(\log n)$  add. memory,  $O(n \log n)$  time





# Graham in-place algorithm

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Graham-InPlaceScan( $S, n, d$ )

*Input:*  $S$  – index to array of length  $n$  with points in plane,  $d = \pm 1$  direction

*Output:* Convex Hull in clockwise order

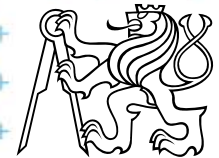
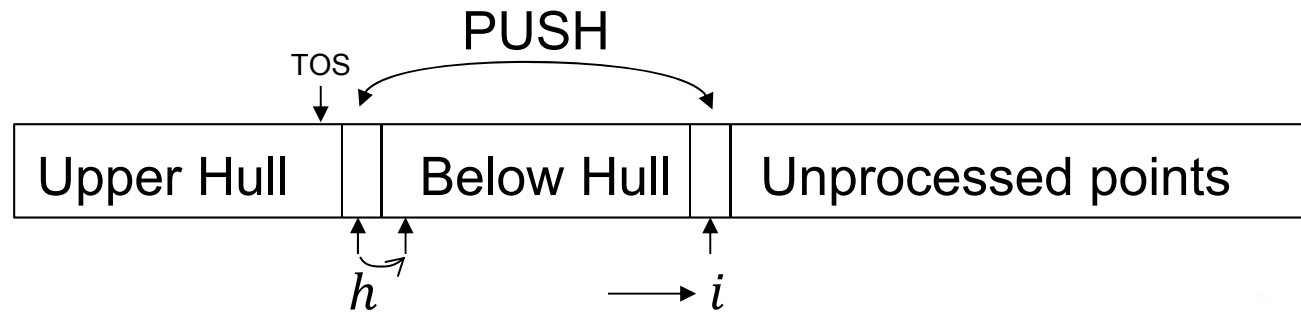
- //  $d$  controls the sort direction:
1. InPlace-Sort( $S, n, d$ ) //  $d = 1$  sort ascending for upper hull
  2.  $h \leftarrow 1$  // empty stack //  $d = -1$  sort descending for lower hull
  3. for  $i \leftarrow 1 \dots n - 1$  do
  4.     while  $h \geq 2$  and not right turn(  $S[h - 2]$ ,  $S[h - 1]$ ,  $S[i]$  ) do
  5.          $h \leftarrow h - 1$  // pop top element from the stack
  6.         swap  $S[i] \leftrightarrow S[h]$  // push the new point to the stack
  7.          $h \leftarrow h + 1$  // increment stack length
  8. return  $h$  // end of convex hull (the first point above the stack)

The array:  $S$  = offset of the sub-array (index of its first point)  
 $h$  = index of the first point above the **stack** (offset to  $S$ )  
 $i$  = index of the **current point**



# Graham in-place algorithm

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# Graham in-place algorithm

Graham-InPlaceHull( $S, n$ )

*Input:*  $S$  – an array of length  $n$  with points in plane

*Output:* Convex Hull in clockwise order (CW)

1.  $h \leftarrow$  Graham-InPlaceScan( $S, n, 1$ ) // 1 = ascending – CW upper hull
2. for  $i \leftarrow 0 \dots h - 2$  do
3.     swap  $S[i] \leftrightarrow S[i + 1]$  // bubble  $a$  to the right  $O(h)$
4.  $h' \leftarrow$  Graham-InPlaceScan( $S + h - 2, n - h + 2, -1$ ) // lower hull
5. return  $h + h' - 2$

sort direction

$O(n \log n)$

CW upper hull

$O(h)$

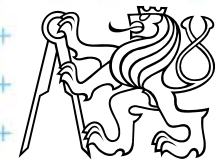
lower hull

Principle:

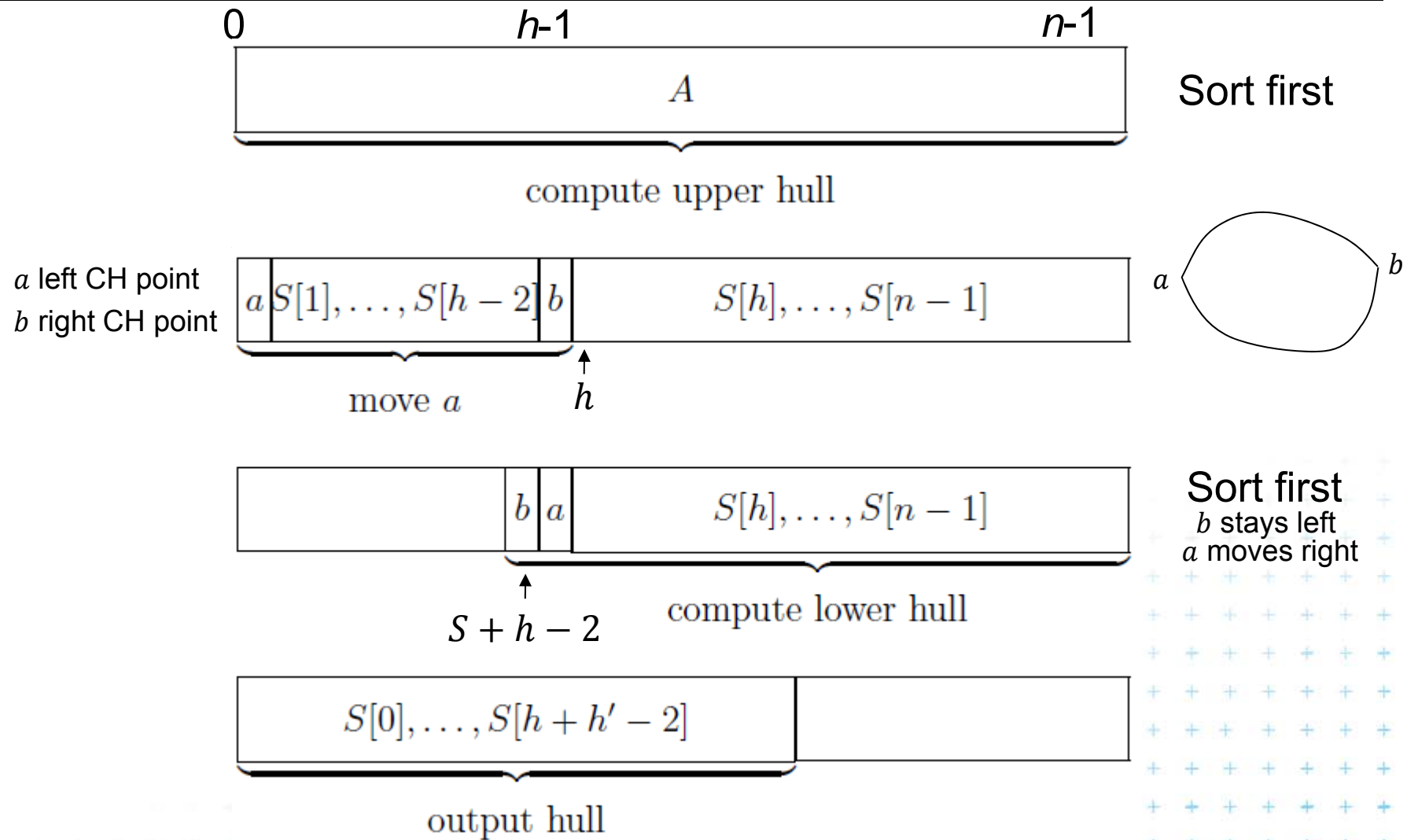
Stack at the beginning of the array  $S$  on indices  $[0 \dots h - 1]$

Exchange by swap operation

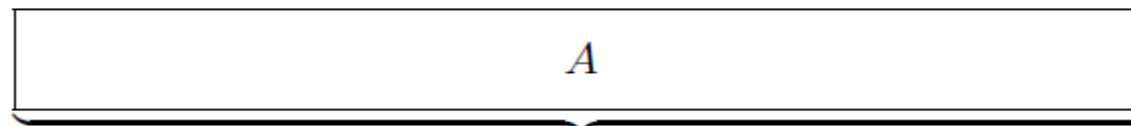
We need the in-place sort



# Graham in-place algorithm



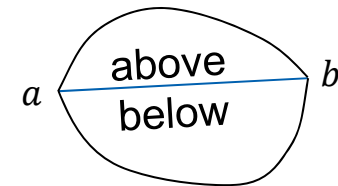
# Optimized Graham in-place algorithm



partition

above  $a, b$

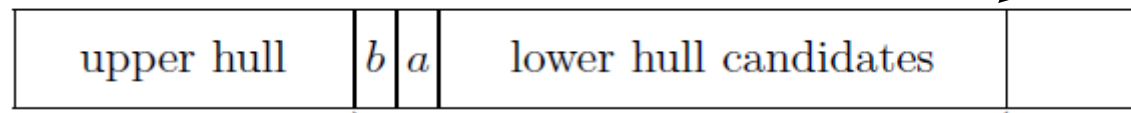
below  $a, b$



compute upper hull

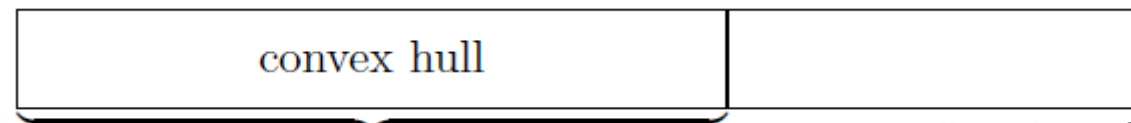


shift



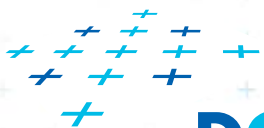
Sort first  
 $b$  stays left  
 $a$  moves right

compute lower hull



[BrönnimannC]

output hull

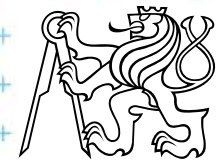


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# Data stream algorithms



- Data stream = a massive sequence of data
  - Too large to store (on disk, memory, cache,...)
- Examples
  - Network traffic
  - Database transactions
  - Sensor networks
  - Satellite data feeds
  - ...
- Approaches
  - Ignore it (CERN ignores 9/10 of the data)
  - Develop algorithms for dealing with such data



# Motivation example

[Muthukrishnan]

- Paul presents numbers  $x = \{1 \dots n\}$  in random order, one number missing
- Carole must determine the missing number but has only  $O(\log n)$  bits of memory

Any idea?

- Compute the sum of the numbers and subtracts the incoming numbers one by one.

$$\text{missing number} = \frac{n(n+1)}{2} - \sum_{i < n} x[i]$$

- The missing number “remains”





# Motivation example

[Muthukrishnan]

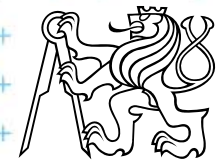
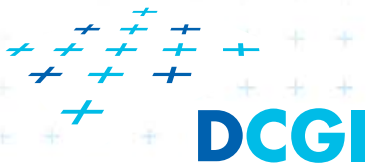
- And two missing numbers  $i, j$  ?
- Store sum of numbers  $s$  and sum of squares  $s'$

$$i + j = \frac{n(n + 1)}{2} - s$$
$$i^2 + j^2 = \frac{n(n + 1)(2n + 1)}{6} - s'$$

(this principle is applicable for  $k$ -missing numbers)



- Single pass over the data:  $a_1, a_2, \dots, a_n$ 
  - Typically  $n$  is known
- Bounded storage (typically  $n^\alpha$  or  $\log^c n$  or only  $c$ )
  - Units of storage: bits, words, or elements (such as points, nodes/edges, ...)
  - Impossible to store the complete data
- Fast processing time per element
  - Randomness is OK (in fact, almost necessary)
  - Often sub-linear time for the whole data
  - Often approximation of the result



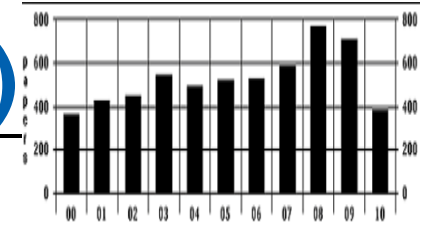
# Data stream models classification

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- Input stream  $a_1, a_2, \dots, a_n$ 
  - arrives sequentially, item by item
  - describes an **underlying signal  $A$** ,  
a 1D function  $A: [1..N] \rightarrow R$
- Models differ on how the input  $a_i$ 's describe the signal  $A$  for increasing  $i$   
(in increasing order of generality):
  - a) Time series model -  $a_i$  equals to signal  $A[i]$
  - b) Cash register model-  $a_i$  are increments to  $A[j]$ ,  $I_i > 0$
  - c) Turnstile model -  $a_i$  are updates to  $A[j]$ ,  $U_i \in R$



# a) Time series model (časová řada)



- Stream elements  $a_i$  are equal to  $A[i]$  ( $a_i$ 's are **samples** of the signal)
- $a_i$ 's appear in **increasing order of  $i$**  ( $i \sim$  time)

## ■ Applications

- Observation of the traffic on IP address each 5 minutes
- NASDAQ volume of trades per minute



## b) Cash register model (*pokladna*)



- $a_i$  are **increments** to signal  $A[j]$ 's
- Stream elements  $a_i = (j, I_i)$ ,  $I_i \geq 0$  to mean

+ only

$I_i$  = Increment

$$A_i[j] = A_{i-1}[j] + I_i$$

where

( $i \sim$  time,  $j \sim$  bucket)

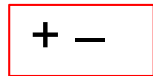
- $A_i[j]$  is the state of the signal after seeing  $i$ -th item
- multiple  $a_i$  can increment given  $A[j]$  over time
- A **most popular** data stream model
  - IP addresses accessing web server (histogram)
  - Source IP addresses sending packets over a link
  - access many times, send many packets,...



## c) Turnstile model (*turniket*)



- $a_i$  are **updates** to signal  $A[j]$ 's
- Stream elements  $a_i = (j, U_i)$ ,  $U_i \in R$  to mean



$U_i =$  Update

$$A_i[j] = A_{i-1}[j] + U_i$$

where

( $i \sim$  time,  $j \sim$  bucket, turnstile)

- $A_i$  is the state of the signal after seeing  $i$ -th item
- $U_i$  may be **positive or negative**
- multiple  $a_i$  can update given  $A[j]$  over time
- A **most general** data stream model
  - Passengers in NY subway arriving and departing
  - Useful for completely dynamic tasks
  - Hard to get reasonable solution in this model





## c) Turnstile model variants (for completeness)

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- **strict** turnstile model –  $A_i[j] \geq 0$  for all  $i$ 
  - People can only exit via the turnstile they entered in
  - Databases – delete only a record you inserted
  - Storage – you can take items only if they are there
- **non-strict** turnstile model –  $A_i[j] < 0$  for some  $i$ 
  - Difference between two cash register streams
  - ( $A_i[j] < 0$  ... negative amount of items for some  $i$ )

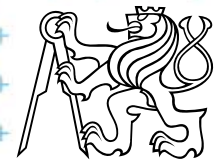
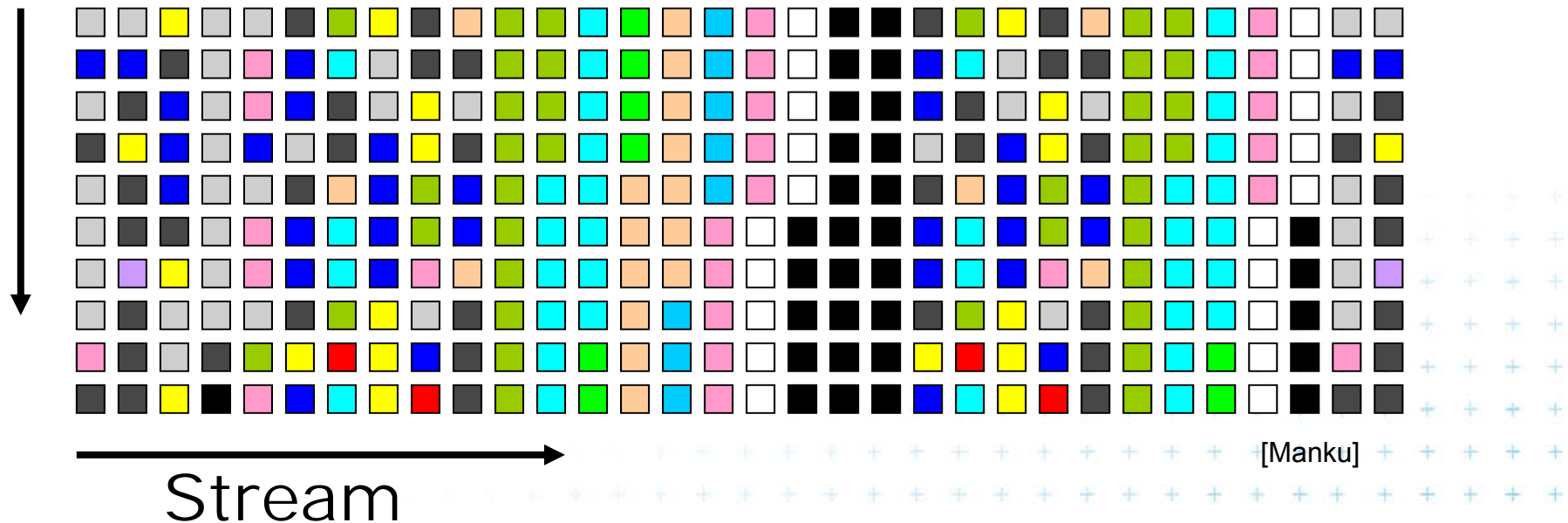


# Examples: Iceberg queries

[Manku]

- Identify all elements whose current frequency  $f$  exceeds support threshold  $s = 0.1\%$

$$f \geq sN$$





# Ex: Iceberg queries – a) ordinary solution

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The ordinary solution in two passes (not data stream)

## 1. Pass – identify frequencies (count the hashes)

- a set of **counters** is maintained. Each incoming item is **hashed** onto a counter, which is incremented.
- These counters are then **compressed into a bitmap**, with a 1 denoting a large counter value.

## 2. Pass – count exact values for large counters only

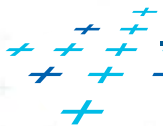
- **exact frequencies counters** for only those elements which hash to a value whose **corresponding bitmap value is 1**

- Hard to modify for data stream – unknown frequencies after only 1<sup>st</sup> pass



## Ex: Iceberg queries – data stream definition

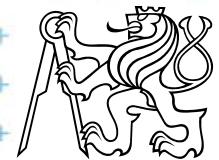
- Input: threshold  $s \in (0,1)$ , error  $\epsilon \in (0,1)$ , length  $N$
- Output: list of items and frequencies  $\epsilon \ll s$
- Guarantees:
  - No item omitted (reported all items with frequency  $> sN$ )
  - No item added (no item with frequency  $< (s - \epsilon)N$ )
  - Estimated frequencies are not less than  $\epsilon N$  of the true frequencies
- Ex:  $s = 0.1\%$ ,  $\epsilon = 0.01\%$   $\rightarrow \epsilon$  about  $\frac{1}{10}$  to  $\frac{1}{20}$  of  $s$ 
  - All elements with freq.  $> 0.1\%$  will output
  - None of element with freq.  $< 0.09\%$  will output
  - Some elements between 0.09% and 0.1% will output



# Ex: Iceberg queries – b) sticky sampling

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- Probabilistic algorithm, given threshold  $s$ , error  $\epsilon$  and probability of failure  $\delta$ 
  - Data structure  $S$  of entries  $(e, f)$ , //  $S$  =subset of counters  
 $e$  element,  $f$  estimated frequency,  
 $r$  sampling rate, sampling probability  $\frac{1}{r}$
- $S \leftarrow \emptyset, r \leftarrow 1$
- If  $e \in S$  then  $(e, f++)$  //count, if the counter exists  
else insert  $(e, f)$  into  $S$  with probability  $\frac{1}{r}$
- $S$  sweeps along the stream as a magnet, attracting all elements which already have an entry in  $S$



# Ex: Iceberg queries – b) sticky sampling

---

- $r$  changes over the stream,  $t = \frac{1}{\epsilon} \log \left( \frac{1}{s\delta} \right)$ ,  $|S| < 2t$ 
  - $2t$  elements  $r = 1$
  - next  $2t$  elements  $r = 2$
  - next  $4t$  elements  $r = 4 \dots$
- whenever  $r$  changes, we update  $S$ 
  - For each entry  $(e, f)$  in  $S$  // random decrement of counters
    - toss a coin until successful (head) // with probability  $1/2$
    - if not successful (tail), decrement  $f$
    - if  $f$  becomes 0, remove entry  $(e, f)$  from  $S$
- Output: list of items with threshold  $s$   
i.e. all entries in  $S$  where  $f \geq (s - \epsilon)N$



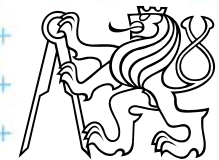
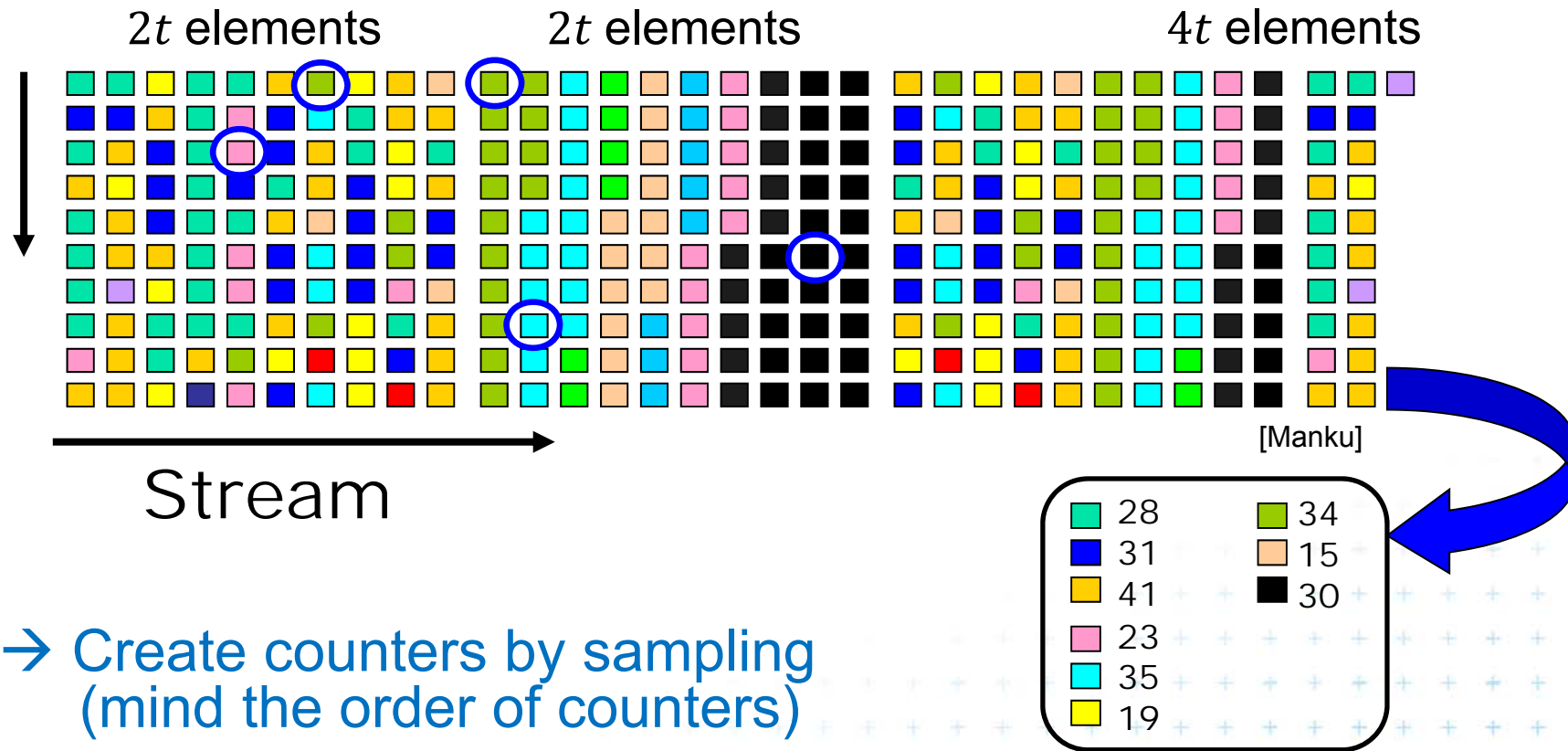
# Ex: Iceberg queries – b) sticky sampling

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- Space complexity is independent on  $N$
- For
  - support threshold  $s = 0.1\%$ ,
  - error  $\epsilon = 0.01\%$ ,
  - and probability of failure  $\delta = 1\%$
- Sticky sampling computes results
  - with  $(1 - \delta) = 99\%$  probability
  - using at most  $2t = 80\,000$  entries
  - $t = \frac{1}{\epsilon} \log\left(\frac{1}{s\delta}\right) = 40\,000, |S| < 2t$



# Ex: Iceberg queries – b) sticky sampling



# Ex: Iceberg queries – c) lossy counting

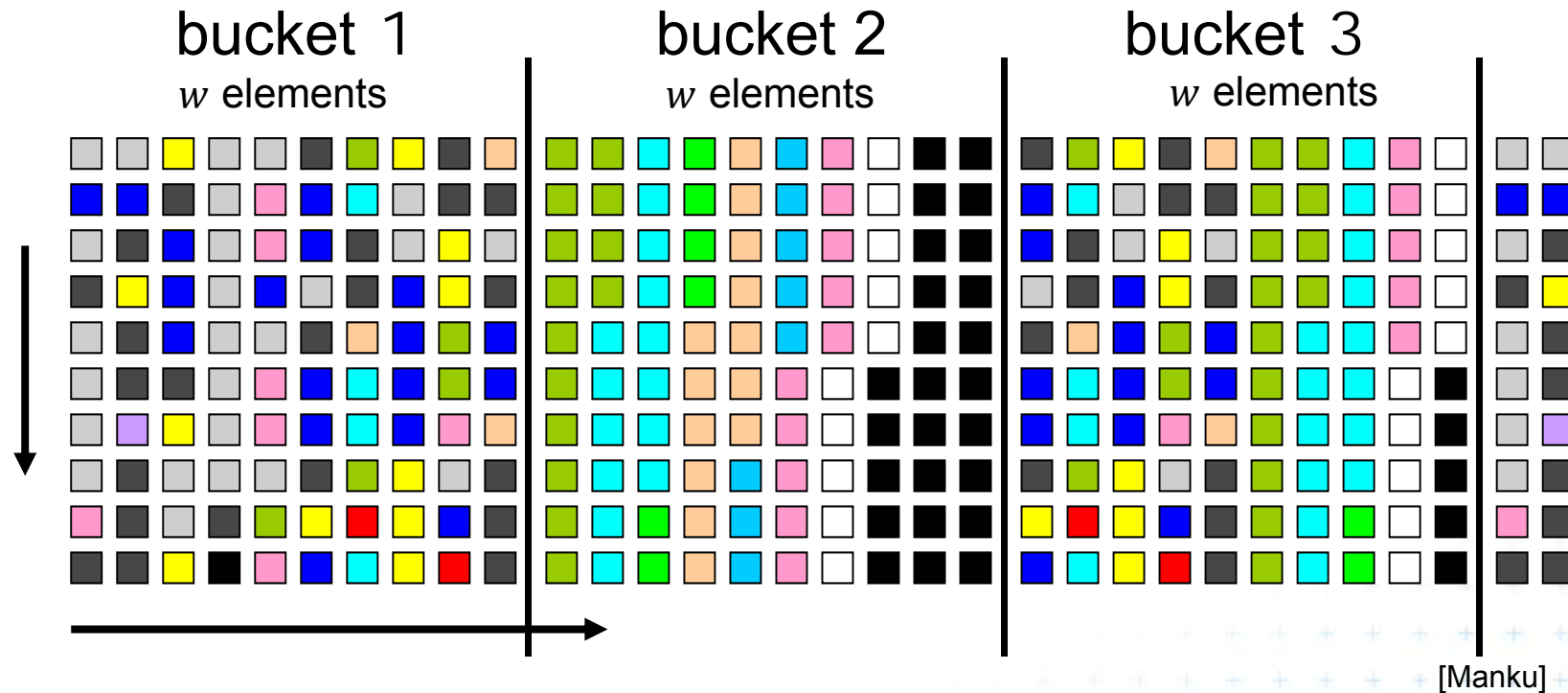
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- **Deterministic algorithm** (user specifies error  $\varepsilon$  and threshold  $s$ )
- **Stream conceptually divided into buckets**
  - With bucket size  $w = \lceil 1/\varepsilon \rceil$  items each
  - Numbered from 1, current bucket id is  $b_{current}$
- **Data structure  $D$  of entries  $(e, f, \Delta)$ ,**
  - $e$  element,
  - $f$  estimated frequency,
  - $\Delta$  maximum possible error of  $f$ ,  $\Delta = b_{current} - 1$   
(max number of occurrences in the previous buckets)
- **At most  $\frac{1}{\varepsilon} \log(\varepsilon N)$  entries**





# Ex: Iceberg queries – c) lossy counting



- Divide the stream into buckets
- Keep exact counters for items in the buckets
- Prune entries at bucket boundaries  
(remove entries for which  $f + \Delta \leq b_{current}$ )





# Ex: Iceberg queries – c) lossy counting alg.

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- $D \leftarrow \emptyset$
- New element  $e$ 
  - If  $e \in D$  then increment its  $f$
  - If  $e \notin D$  then
    - Create a new entry  $(e, 1, b_{current} - 1)$
    - If on the bucket border, i.e.,  $N \bmod w = 0$  then delete entries with  $f + \Delta \leq b_{current}$
    - i.e., with zero or one occurrence in each of the previous buckets
  - New  $\Delta = b_{current} - 1$  is maximum number of times  $e$  could have occurred in the first  $b_{current} - 1$  buckets
- Output: list of items with threshold  $s$   
i.e. all entries in  $S$  where  $f \geq (s - \epsilon)N$



# Comparison of sticky and lossy sampling

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- Sticky sampling performs worse
  - Tendency to remember every unique element
  - The worst case is for sequence without duplicates
- Lossy counting
  - Is good in pruning low frequency elements quickly
  - Worst case for pathological sequence which never occurs in reality



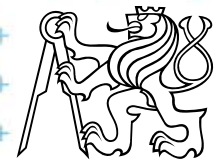
# Number of mutually different entries

1/2

- Input: stream  $a_1, a_2, \dots, a_n$ , with repeated entries
- Output: Estimate of number  $c$  of different entries
- Appl: # of different transactions in one day

## a) Precise deterministic algorithm:

- Array  $b[1..U]$ ,  $U = \text{max number of different entries}$
- Init by  $b[i] = 0$  for all  $i$ , counter  $c = 0$
- for each  $a_i$ 
  - if  $b[a_i] = 0$  then  $\text{inc}(c)$ ,  $b[i] = 1$
- Return  $c$  as number of different entries in  $b[]$
- $O(1)$  update and query times,  $O(U)$  memory



## b) Approximate algorithm

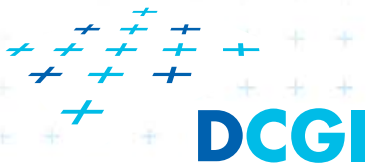
- Array  $b[1 \dots \log U]$ ,  $U = \text{max number of different entries}$
- Init by  $b[i] = 0$  for all  $i$
- Hash function  $h: \{1..U\} \rightarrow \{0..\log U\}$
- For each  $a_i$ 
  - Set  $b[h(a_i)] = 1$
- Extract probable number of different entries from  $b[]$



# Sublinear time example

$$O(\text{alg}) < O(n)$$

- Given mutually different numbers  $a_1, a_2, \dots, a_n$
- Determine any number from upper half of values
- Alg: select  $k$  numbers equally randomly
  - Compute their maximum
  - Return this estimation as solution
- Probability of wrong answer = probability of all selected numbers are from the lower half =  $\left(\frac{1}{2}\right)^k$
- For error  $\epsilon$  take  $\log \frac{1}{\epsilon}$  samples
- Not useful for MIN, MAX selection



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# Randomized algorithms



# Randomized algorithms

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## Motivation

- Array of elements, half of char "a", half of char "b"
- Find "a"
- Deterministic alg:  $n/2$  steps of sequential search (when all "b" are first)
- Randomized:
  - Try random indices
  - Probability of finding "a" soon is high regardless of the order of characters in the array (Las Vegas algorithm – keep trying up to  $n/2$  steps)

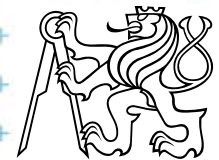




# Randomized algorithms

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- May be **simpler** even if the same worst time
- Deterministic algorithm
  - is **not known** (prime numbers)
  - does **not exist**
- Randomization
  - can **improve the average running time** (with the same worst case time), while
  - the worst time **depends on our luck** – **not on the data distribution**  
(It is “hard” to prepare killing datasets)





# Randomized algorithms

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- a) Incremental algorithms  
(insert something in random order)
  - Linear programming (random plane insertion)
  - Convex hulls
  - Intersections, space subdivisions
  
- b) Divide and conquer  
(split in random place)
  - Random sampling
  - Nearest neighbors, trapezoidal subdivisions



# Another classification

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## ■ Monte Carlo

- We **always** get an answer, often not correct
- **Fast** solution with risk of an error
- It is **not possible to determine**, if the answer is **correct**  
→ run multiple times and compare the results
- Output can be understood as a **random variable**
- Example: prime number test
  - Task: Find  $a \in \left(2, \frac{n}{2}\right)$  such as  $n$  is divisible by  $a$
  - Algorithm: Sample 10 numbers from the given interval, answer

## ■ Las Vegas

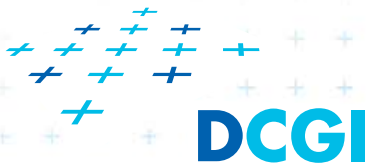


# Las Vegas algorithms

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## Las Vegas

- We **always** get a **correct answer**
- The **run time is random** (typically  $\leq$  deterministic time)
- **Sometimes fails**  $\rightarrow$  perform restart
- Example: Randomized quicksort
  - No median necessary
  - Simpler algorithm
  - Independent on data distribution
  - Return a correct result
  - The result will be ready in  $\theta(n \log n)$  time with a high probability
  - Bad luck – we select the smallest element  $\rightarrow$  Selection sort



# Randomized quicksort (Las Vegas alg.)

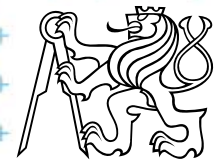
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RQS( $S$ ) = Randomized Quicksort

*Input:* sequence of data elements  $a_1, a_2, \dots, a_n \in S$

*Output:* sorted set  $S$

1. Step 1: choose  $i \in \langle 1, n \rangle$  in random
2. Step 2: Let  $A$  is a multiset  $\{a_1, a_2, \dots, a_n\}$ 
  - if  $n = 1$  then output( $S$ )
  - else – create three subsets of  $S_{<}, S_{=}, S_{>}$ 
$$S_{<} = \{b \in A: b < a_i\}$$
$$S_{=} = \{b \in A: b = a_i\}$$
$$S_{>} = \{b \in A: b > a_i\}$$
3. Step 3:  $RQS(S_{<})$  and  $RQS(S_{>})$
4. Return:  $RQS(S_{<}), S_{=}, RQS(S_{>})$



# Conclusion on randomized algs.

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- Randomized algorithms are often experimental
- We would not get perfect results, but nicely good
- We use randomized algorithm if we do not know how to proceed



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