



ALGORITHMS OF BIOINFORMATICS



Googling Genomes

18 December 2025

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Outline

7 Googling Genomes

- 7.1 Range-Minimum Queries
- 7.2 RMQ – Sparse Table Solution
- 7.3 RMQ – Cartesian Trees
- 7.4 String Matching in Enhanced Suffix Array
- 7.5 The Burrows-Wheeler Transform
- 7.6 Inverting the BWT
- 7.7 The LR Mapping

Recall Unit 6

Application 4: Longest Common Extensions

- We implicitly used a special case of a more general, versatile idea:

The *longest common extension (LCE)* data structure:

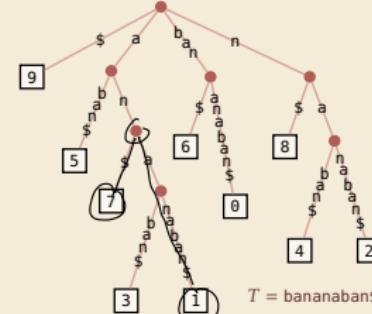
- Given: String $T[0..n]$
- Goal: Answer LCE queries, i.e.,
given positions i, j in T ,
how far can we read the same text from there?
formally: $\text{LCE}(i, j) = \max\{\ell : T[i..i + \ell] = T[j..j + \ell]\}$

↔ use suffix tree of T !

(length of) longest common prefix
of i th and j th suffix

- In T : $\text{LCE}(i, j) = \text{LCP}(T_i, T_j) \rightsquigarrow$ same thing, different name!
 $=$ string depth of
lowest common ancestor (LCA) of
leaves \boxed{i} and \boxed{j}

- in short: $\text{LCE}(i, j) = \text{LCP}(T_i, T_j) = \text{stringDepth}(\text{LCA}(\boxed{i}, \boxed{j}))$



Recall Unit 6

Efficient LCA

How to find lowest common ancestors?

- ▶ Could walk up the tree to find LCA $\rightsquigarrow \Theta(n)$ worst case 
- ▶ Could store all LCAs in big table $\rightsquigarrow \Theta(n^2)$ space and preprocessing 



Amazing result: Can compute data structure in $\Theta(n)$ time and space that finds any LCA in **constant(l)** time.

- ▶ a bit tricky to understand
- ▶ but a theoretical breakthrough
- ▶ and useful in practice

\rightsquigarrow for now, use $O(1)$ LCA as black box.

\rightsquigarrow After linear preprocessing (time & space), we can find LCEs in $O(1)$ time.

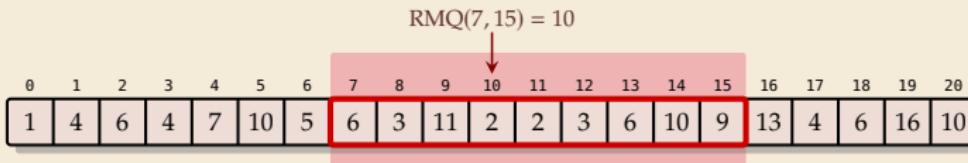


and suffix tree construction inside ...

7.1 Range-Minimum Queries

Range-minimum queries (RMQ)

- Given: Static array $A[0..n]$ of numbers
array/numbers don't change
- Goal: Find minimum in a range;
 A known in advance and can be preprocessed



- Nitpicks:
 - Report *index* of minimum, not its value
 - Report *leftmost* position in case of ties

Finally: Longest common extensions

- In Unit 6: Left question open how to compute LCA in suffix trees
- But: Enhanced Suffix Array makes life easier!

$$\text{LCE}(i, j) = \text{LCP}[\text{RMQ}_{\text{LCP}}(\min\{R[i], R[j]\} + 1, \max\{R[i], R[j]\})]$$

Inverse suffix array: going left & right

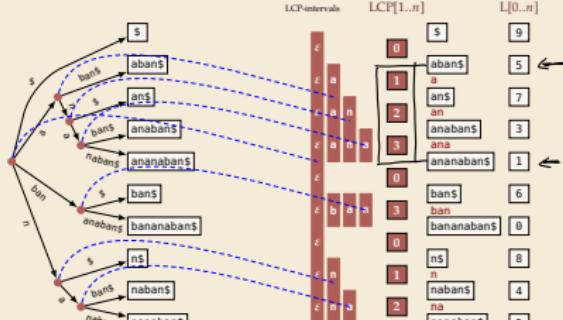
► to understand the fastest algorithm, it is helpful to define the *inverse suffix array*:

- $R[i] = r \iff L[r] = i$ L = leaf array
- \iff there are r suffixes that come before T_i in sorted order
- $\iff T_i$ has (0-based) **rank** r \rightsquigarrow call $R[0..n]$ the **rank array**

i	$R[i]$	T_i	r	$L[r]$	$T_{L[r]}$
0	6 th	bananaban\$	0	9	\$
1	4 th	ananaban\$	1	5	aban\$
2	9 th	nanaban\$	2	7	an\$
3	3 th	anaban\$	3	3	anaban\$
4	8 th	naban\$	4	1	anabanan\$
5	1 st	aban\$	5	6	ban\$
6	5 th	ban\$	6	8	bananaban\$
7	2 nd	an\$	7	8	n\$
8	7 th	n\$	8	4	naban\$
9	0 th	\$	9	2	nanaban\$

sort suffixes

LCP array and internal nodes



\rightsquigarrow Leaf array $L[0..n]$ plus LCP array $LCP[1..n]$ encode full tree!

Rules of the Game

- ▶ For the following, consider RMQ on arbitrary arrays
- ▶ comparison-based \rightsquigarrow values don't matter, only relative order
- ▶ Two main quantities of interest:
 1. Preprocessing time: Running time $P(n)$ of the preprocessing step
 2. Query time: Running time $Q(n)$ of one query (using precomputed data)
- ▶ Write $\langle P(n), Q(n) \rangle$ **time solution** for short

\rightsquigarrow space usage $\leq P(n)$

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RMQ Implications for LCE

- ▶ Recall: Can compute (inverse) suffix array and LCP array in $O(n)$ time
- \rightsquigarrow $\langle P(n), Q(n) \rangle$ time RMQ data structure implies $\langle P(n) + O(n), Q(n) \rangle$ time LCE data structure

Trivial Solutions

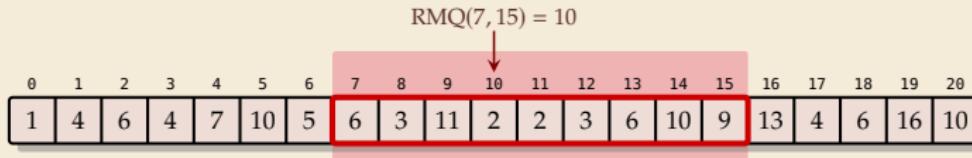
0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
1	4	6	4	7	10	5	6	3	11	2	2	3	6	10	9	13	4	6	16	10

$$\text{RMQ}(7, 15) = 10$$



- ▶ Two easy solutions show extreme ends of scale:

Trivial Solutions

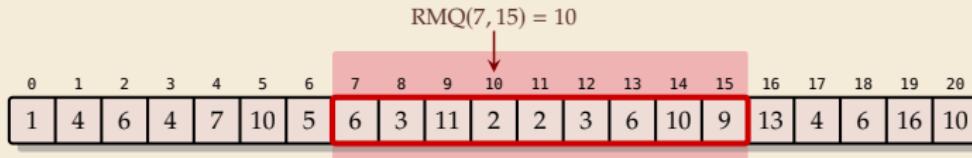


- ▶ Two easy solutions show extreme ends of scale:

1. Scan on demand

- ▶ no preprocessing at all
- ▶ answer $\text{RMQ}(i, j)$ by scanning through $A[i..j]$, keeping track of min
 $\rightsquigarrow \langle O(1), O(n) \rangle$

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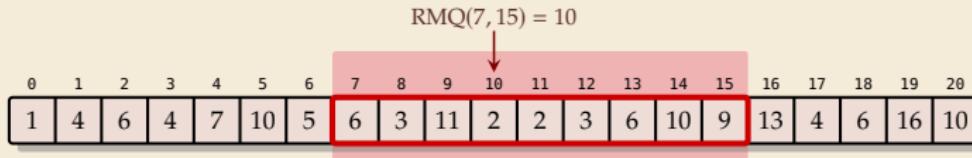
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2. Precompute all

- ▶ Precompute all answers in a big 2D array $M[0..n][0..n]$
- ▶ queries simple: $\text{RMQ}(i, j) = M[i][j]$
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- ▶ Preprocessing can reuse partial results $\rightsquigarrow \langle O(n^2), O(1) \rangle$

7.2 RMQ – Sparse Table Solution

Sparse Table

- ▶ **Idea:** Like “precompute-all”, but keep only *some* entries
- ▶ store $M[i][j]$ iff $\ell = j - i + 1$ is 2^k .
 - ~~ $\leq n \cdot \lg n$ entries
 - ~~ Can be stored as $\underline{M'[i][k]}$

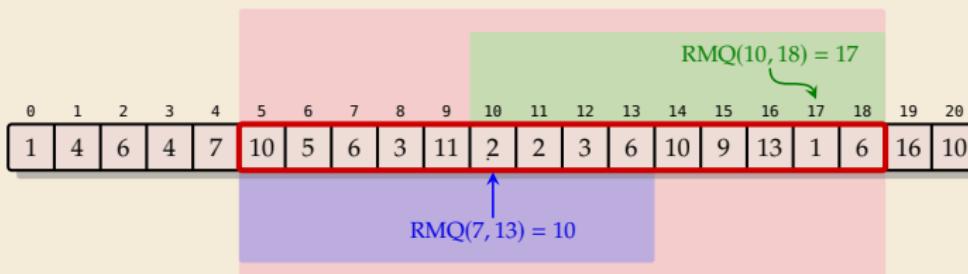
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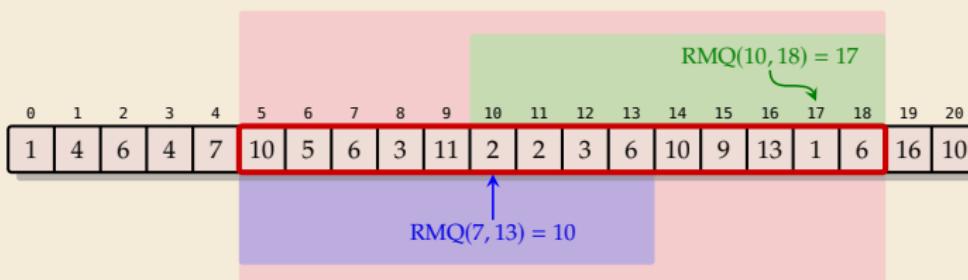
1. Find k with $\ell/2 \leq 2^k \leq \ell$
2. Cover range $[i..j]$ by
 2^k positions right from i and
 2^k positions left from j
3. $\text{RMQ}(i, j) = \arg \min\{A[\text{rmq}_1], A[\text{rmq}_2]\}$
with $\text{rmq}_1 = \text{RMQ}(i, i + 2^k - 1)$
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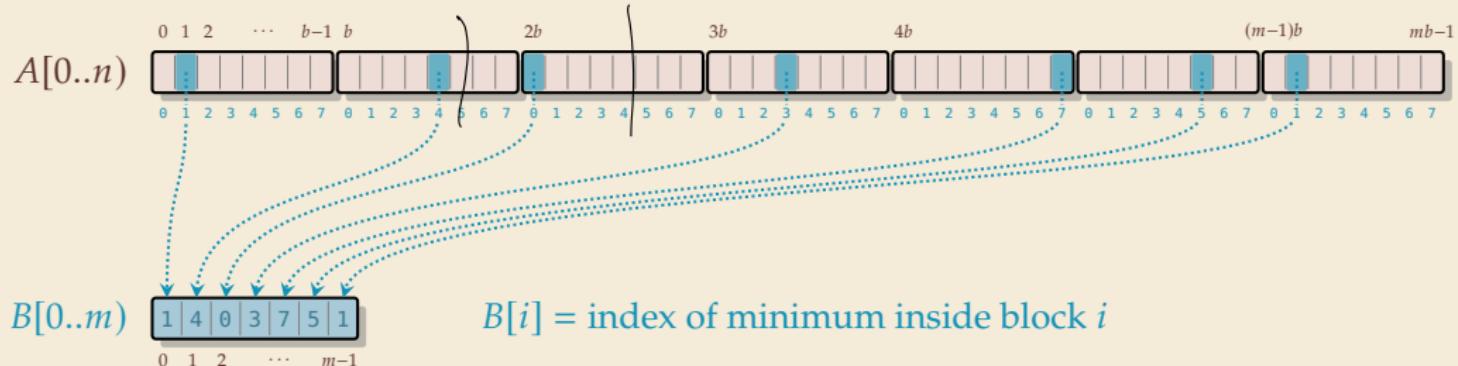
- Preprocessing can be done in $O(n \log n)$ times
 - ~ $\langle O(n \log n), O(1) \rangle$ time solution!

Bootstrapping

- ▶ We know a $\langle O(n \log n), O(1) \rangle$ time solution
- ▶ If we use that for $m = \Theta(n/\log n)$ elements, $O(m \log m) = O(n)!$

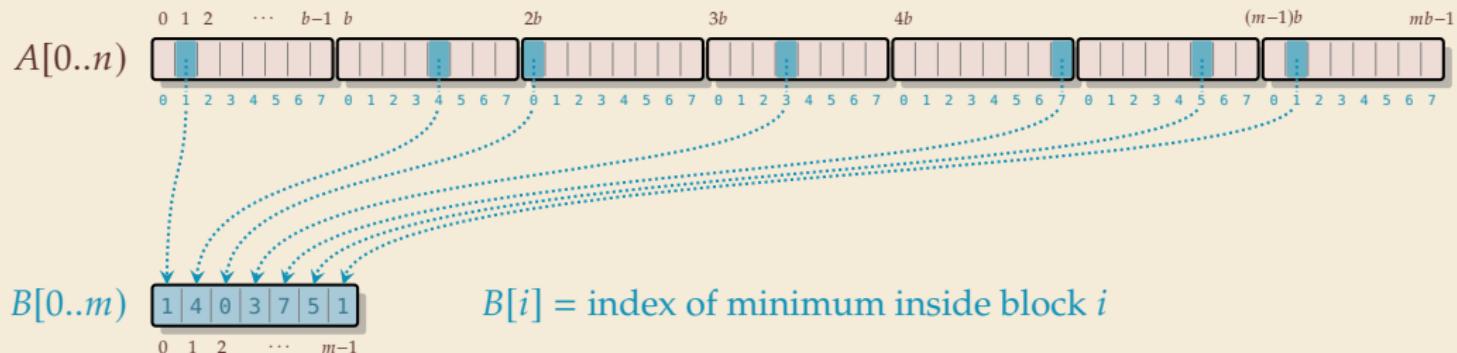
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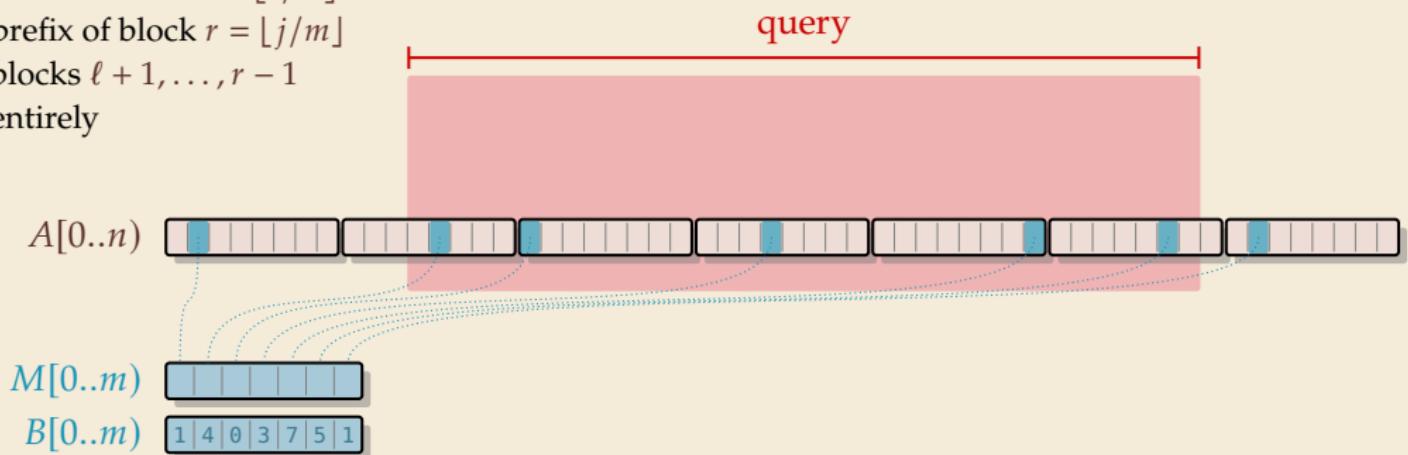
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- ~~ Use sparse table solution for B .
- ~~ Can solve RMQs in $B[0..m)$ in $\langle O(n), O(1) \rangle$ time

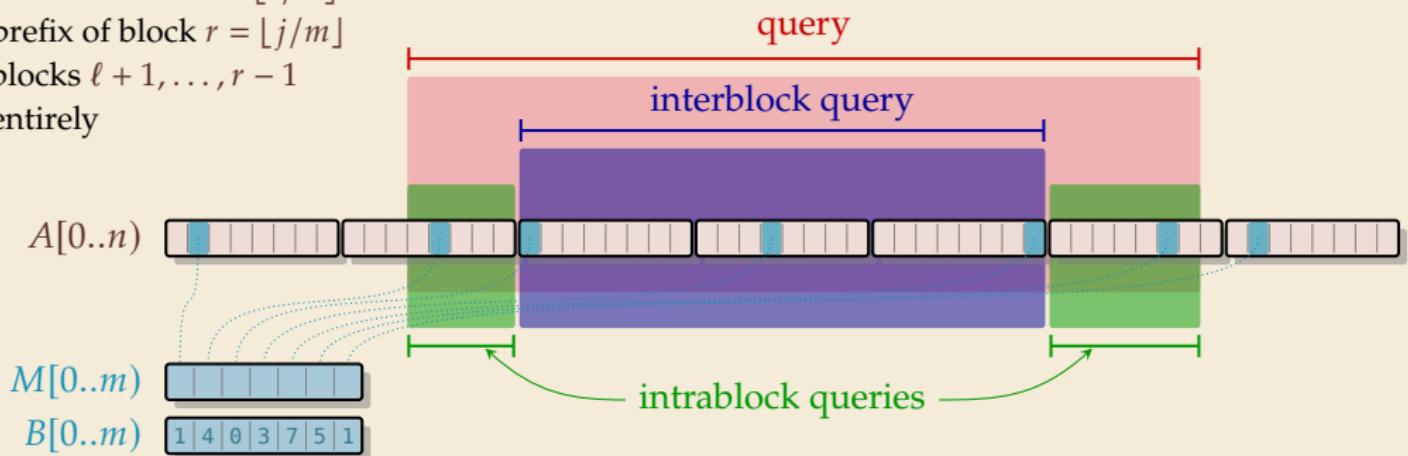
Query decomposition

- ▶ Query $\text{RMQ}_A(i, j)$ covers
 - ▶ suffix of block $\ell = \lfloor i/m \rfloor$
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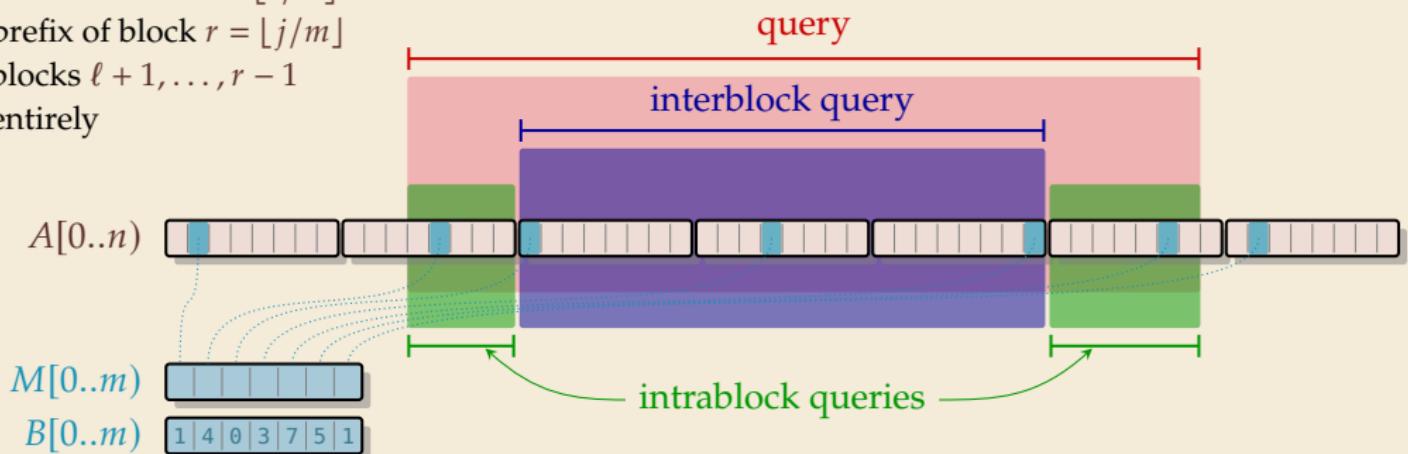
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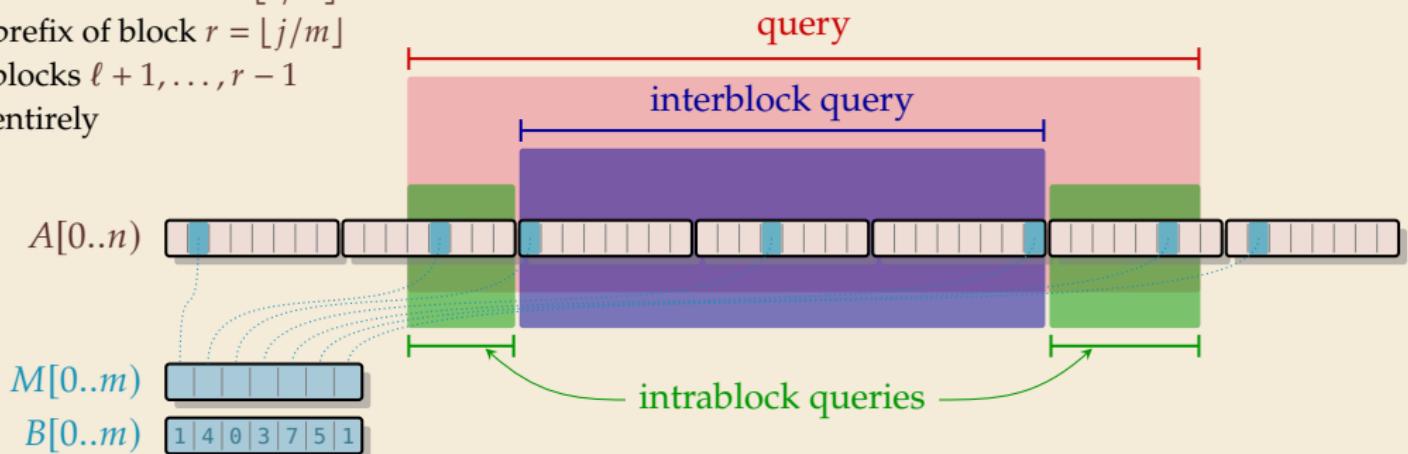
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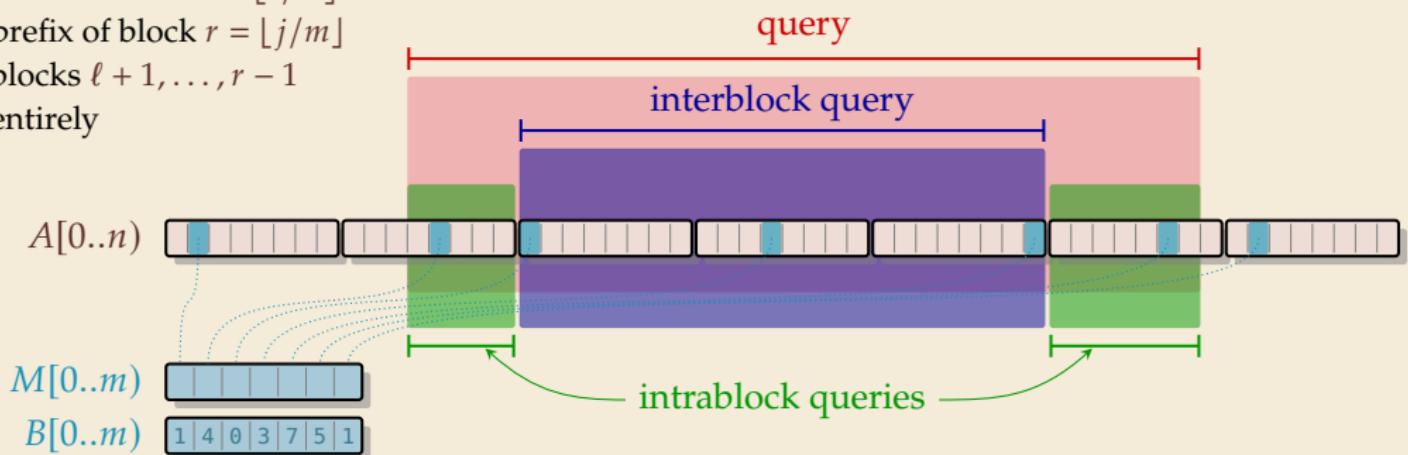
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 $O(\log n)$

$\langle O(n), O(\log n) \rangle$