



ALGORITHMS OF BIOINFORMATICS

Googling Genomes

18 December 2025

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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:

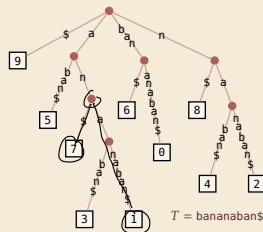
The *longest common extension (LCE)* data structure:

- ▶ **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 !

- ▶ In \mathcal{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(!) time**.

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

and suffix tree construction inside ...



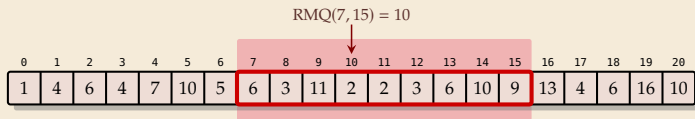
\rightsquigarrow for now, use $O(1)$ LCA as black box.

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

7.1 Range-Minimum Queries

Range-minimum queries (RMQ)

- array / numbers don't change
- ▶ **Given:** Static array $A[0..n)$ of numbers
 - ▶ **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*:

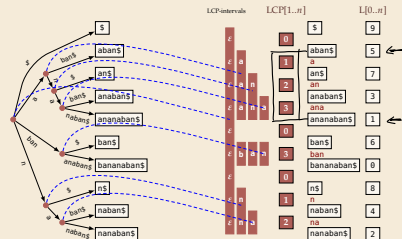
- $R[i] = r \iff L[r] = i$ $L = \text{leaf array}$
- \iff there are r suffixes that come before T_i in sorted order
- $\iff T_i$ has (0-based) *rank* $r \rightsquigarrow$ call $R[0..n]$ the *rank array*

i	$R[i]$	T_i		r	$L[r]$	$T_{L[r]}$
0	6 th	bananabans		0	9	\$
1	4 th	ananabans		1	5	abans
2	9 th	nanabans		2	7	ans
3	3 th	anabans		3	3	anabans
4	8 th	nabans		4	1	ananabans
5	1 th	abans		5	6	ban\$
6	5 th	ban\$		6	0	bananabans
7	2 th	an\$		7	8	n\$
8	7 th	n\$		8	4	nabans
9	0 th	\$		9	2	nanabans

sort suffixes

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LCP array and internal nodes



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

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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)
- $\nwarrow \rightsquigarrow$ space usage $\leq P(n)$
- ▶ Write $\langle P(n), Q(n) \rangle$ **time solution** for short

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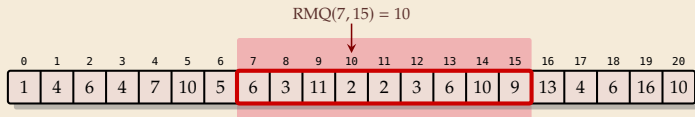
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RMQ Implications for LCE

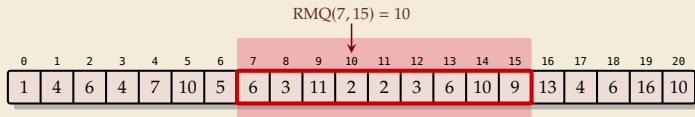
- ▶ Recall: Can compute (inverse) suffix array and LCP array in $O(n)$ time
- $\rightsquigarrow \langle P(n), Q(n) \rangle$ time RMQ data structure implies
 $\langle P(n) + O(n), Q(n) \rangle$ time LCE data structure

Trivial Solutions



- Two easy solutions show extreme ends of scale:

Trivial Solutions



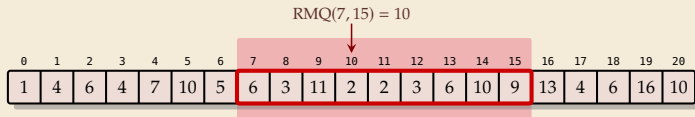
- ▶ Two easy solutions show extreme ends of scale:

1. Scan on demand

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

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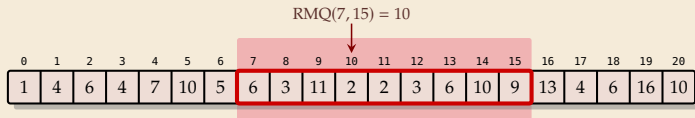
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- ▶ Precompute all answers in a big 2D array $M[0..n][0..n]$
 - ▶ queries simple: $\text{RMQ}(i, j) = M[i][j]$
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- ▶ Preprocessing can reuse partial results $\rightsquigarrow \langle O(n^2), O(1) \rangle$

7.2 RMQ – Sparse Table Solution

Sparse Table

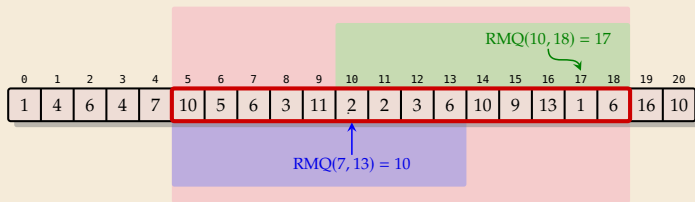
- ▶ **Idea:** Like “precompute-all”, but keep only *some* entries
- ▶ store $M[i][j]$ iff $\ell = j - i + 1$ is 2^k .
 - ↪ $\leq n \cdot \lg n$ entries
 - ↪ Can be stored as $M'[i][k]$

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- ▶ How to answer queries?

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 - $\leadsto \leq n \cdot \lg n$ entries
 - \leadsto Can be stored as $M'[i][k]$
- How to answer queries?



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 $\text{rmq}_1 = \text{RMQ}(i, i + 2^k - 1)$
 - $\text{rmq}_2 = \text{RMQ}(j - 2^k + 1, j)$

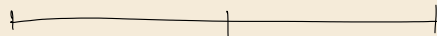
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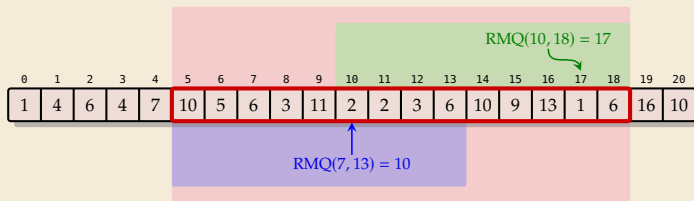
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with $\text{rmq}_1 = \text{RMQ}(i, i + 2^k - 1)$
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► Preprocessing can be done in $O(n \log n)$ times

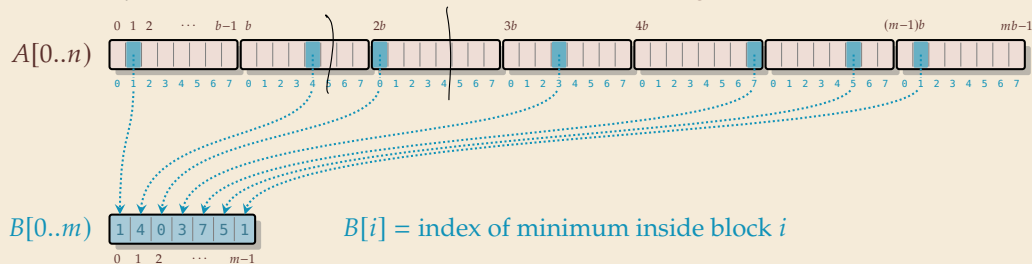
↪ $\langle O(n \log n), O(1) \rangle$ time solution!

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, $O(m \log m) = O(n)$!

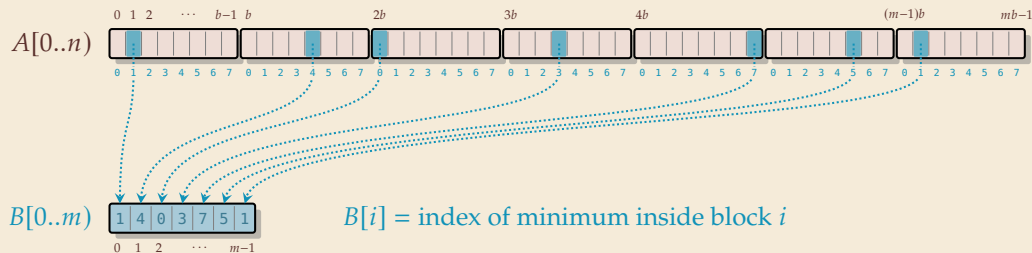
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- ▶ Create array of block minima $B[0..m)$ for $m = \lceil n/b \rceil = O(n/\log n)$



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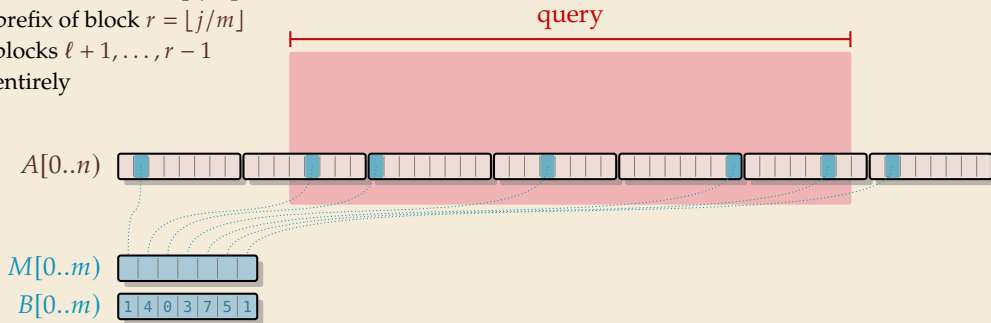


\rightsquigarrow Use sparse table solution for B .

\rightsquigarrow Can solve RMQs in $B[0..m]$ in $\langle O(n), O(1) \rangle$ time

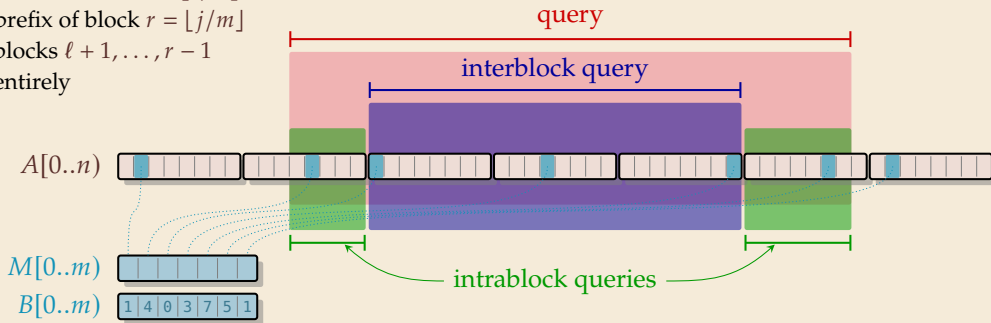
Query decomposition

- ▶ Query $\text{RMQ}_A(i, j)$ covers
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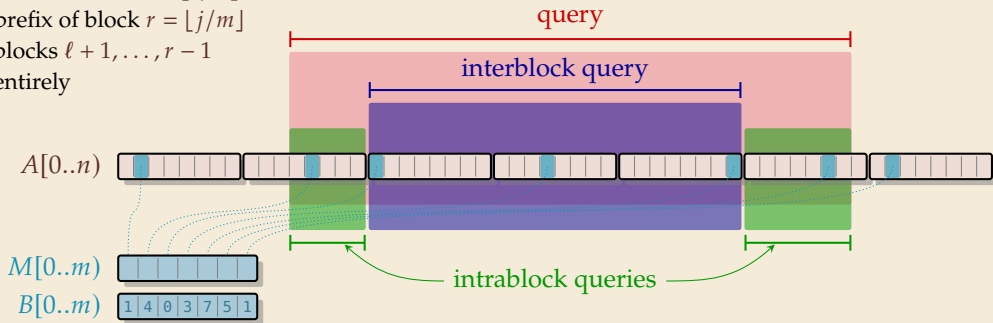
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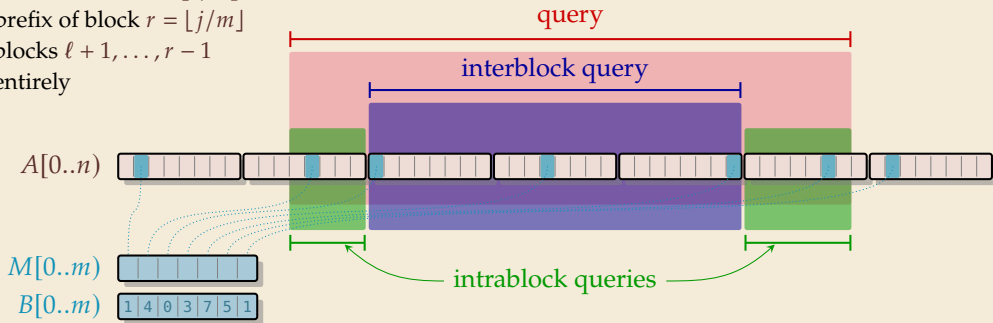
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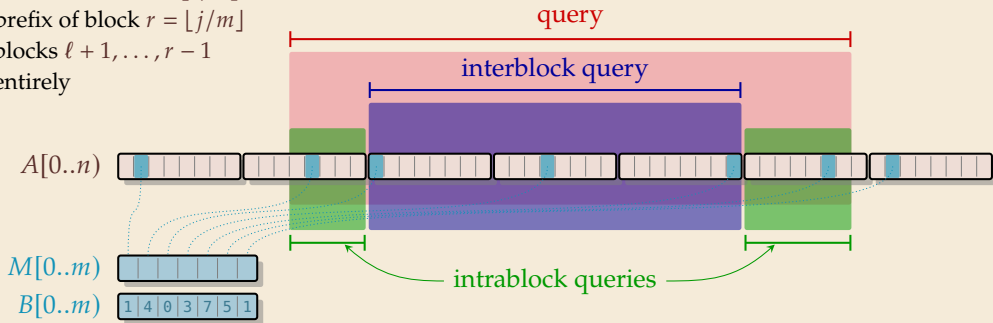


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if **intrablock** and **interblock** queries known

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