# RANGE TREES AND PRIORITY SEARCH TREES 

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## RANGE TREES

- Balanced binary search tree
- All data stored in the leaf nodes
- Leaf nodes linked in sorted order by a doubly-linked list
- Searching for [B: E]

1. find node with smallest value $\geq B$ or largest $\leq B$
2. follow links until reach node with value $>E$

- $O\left(\log _{2} N+F\right)$ time to search, $O\left(N \cdot \log _{2} N\right)$ to build, and $O(N)$ space for $N$ points and $F$ answers
- Ex: sort points in 2-d on their $x$ coordinate value



## 2-D RANGE TREES

- Binary tree of binary trees
- Sort all points along one dimension (say $x$ ) and store them in the leaf nodes of a balanced binary tree such as a range tree (single line)
- Each nonleaf node contains a 1-d range tree of the points in its subtrees sorted along $y$ (double lines)
- Ex:

- Actually, don't need the 1-d range tree in $y$ at the root and at the sons of the root
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## SEARCHING 2-D RANGE TREES ([bx:Ex],[By:Ey])

1. Search tree $T$ for nodes $B X$ and EX

- find node $L X$ with a minimum value $\geq B X$
- find node RX with a maximum value $\leq$ EX

2. Find their nearest common ancestor $Q$
3. Compute $\left\{L_{i}\right\}$ and $\left\{R_{i}\right\}$, the sequences of nodes forming the paths from $Q$ to $L X$ and $R X$, respectively (including LX and RX but excluding Q)

- LEFT (P) and RIGHT (P) are sons of $P$
- mIDRANGE(P) discriminates on $x$ coordinate value
- RANGE_TREE(P) denotes the 1-d range tree stored at $P$

4. For each element in the sequences $\left\{L_{i}\right\}$ and $\left\{R_{i}\right\}$ do


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- if $P$ and $\operatorname{LEFT}(P)$ are in $\left\{L_{\mathrm{j}}\right\}$, then look for [BY,EY] in RANGE_TREE(RIGHT(P))



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- if $P$ and RIGHT (P) are in $\left\{\mathrm{R}_{\mathrm{j}}\right\}$, then look for [BY,EY] in RANGE_TREE(LEFT(P))

5. Check if $L X$ and $R X$ are in ([BX:EX],[BY:EY])


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- if $P$ and RIGHT (P) are in $\left\{\mathrm{R}_{\mathrm{j}}\right\}$, then look for [BY,EY] in RANGE_TREE(LEFT(P))

5. Check if $L X$ and $R X$ are in ([BX:EX],[BY:EY])

- Total $O\left(\log _{2} 2 N+F\right)$ time to search and $O\left(N \cdot \log _{2} N\right)$ space and time to build for $N$ points and $F$ answers



## EXAMPLE OF SEARCH IN A 2-D RANGE TREE

- Find all points in ([25:85],[8:16])



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1. Find nearest common ancestor - ie., A
2. Find paths to $L X=25$ and $R X=85$

## EXAMPLE OF SEARCH IN A 2-D RANGE TREE

- Find all points in ([25:85],[8:16])


1. Find nearest common ancestor - ie., A
2. Find paths to $L X=25$ and $R X=85$
3. Look in subtrees

- B and B's left son D are in path, so search range tree of B's right son E and report $(52,10)$


## EXAMPLE OF SEARCH IN A 2-D RANGE TREE

- Find all points in ([25:85],[8:16])


1. Find nearest common ancestor - ie., A
2. Find paths to $L X=25$ and $R X=85$
3. Look in subtrees

- B and B's left son D are in path, so search range tree of B's right son E and report $(52,10)$
- C and c's right son G are in path, so search range tree of C's left son F and report none


## EXAMPLE OF SEARCH IN A 2-D RANGE TREE

- Find all points in ([25:85],[8:16])


1. Find nearest common ancestor - i.e., A
2. Find paths to $L X=25$ and $R X=85$
3. Look in subtrees

- B and B's left son D are in path, so search range tree of B's right son E and report $(52,10)$
- C and C's right son G are in path, so search range tree of C's left son F and report none

4. Check boundaries of $x$ range (ie., $(27,35)$ and $(85,15)$ ) and report $(85,15)$

## PRIORITY SEARCH TREES

- Sort all points by their $x$ coordinate value and store them in the leaf nodes of a balanced binary tree (i.e., a range tree)
- Starting at the root, each node contains the point in its subtree with the maximum value for its $y$ coordinate that has not been stored at a shallower depth in the tree; if no such node exists, then node is empty
- $O(N)$ space and $O\left(N \cdot \log _{2} N\right)$ time to build for $N$ points
- Result: range tree in $x$ and heap (i.e., priority queue) in $y$

- Good for semi-infinite ranges - i.e., ([BX:EX],[BY: $\infty]$ )
- Can only perform a $2-\mathrm{d}$ range query if find ([BX:EX],[BY:©]) and discard all points $(x, y)$ such that $y>E Y$
- No need to link leaf nodes unless search for all points in range of $x$ coordinate values


## SEMI-INFINITE RANGE QUERY ON A PRIORITY SEARCH TREE ([bx:Ex],[BY: $\infty]$ )

- Procedure

1. Descend tree looking for the nearest common ancestor of $B X$ and $E x$ - i.e., $Q$

- associated with each examined node $T$ is a point $P$
- exit if $P$ does not exist as all points in the subtrees have been examined and/or reported
- exit if $P_{y}<B Y$ as $P$ is point with maximum $y$ coordinate value in $T$
- otherwise, output $P$ if $P_{X}$ is in $[B X: E X]$

2. Once $Q$ has been found, process left and right subtrees applying the tests above to their root nodes $T$

- Tin left (right) subtree of Q:
a. check if $B X(E X)$ in $\operatorname{LEFT}(T)(\operatorname{RIGHT}(T))$
b. yes: all points in $\operatorname{RIGHT}(\mathrm{T})(\operatorname{LEFT}(\mathrm{T})$ ) are in $x$ range - check if in $y$ range
- recursively apply to LEFT(T) (RIGHT(T))
c. no: recursively apply to RIGHT(T) (LEFT(T))
- $O\left(\log _{2} N+F\right)$ time to search for $N$ points and $F$ answers
- Ex:



## SEMI-INFINITE RANGE QUERY ON A PRIORITY SEARCH TREE ([bx:Ex],[by: $\infty]$ )

- Procedure

1. Descend tree looking for the nearest common ancestor of $B X$ and $E X$ - i.e., $Q$

- associated with each examined node $T$ is a point $P$
- exit if $P$ does not exist as all points in the subtrees have been examined and/or reported
- exit if $P_{y}<B Y$ as $P$ is point with maximum $y$ coordinate value in $T$
- otherwise, output $P$ if $P_{X}$ is in [ $\left.B X: E X\right]$

2. Once $Q$ has been found, process left and right subtrees applying the tests above to their root nodes T

- T in left (right) subtree of Q:
a. check if $B X(E X)$ in $\operatorname{LEFT}(T)(\operatorname{RIGHT}(T))$
b. yes: all points in RIGHT(T) (LEFT(T)) are in $x$ range - check if in $y$ range
- recursively apply to LEFT(T) (RIGHT(T))
c. no: recursively apply to RIGHT(T) (LEFT(T))
- $O\left(\log _{2} N+F\right)$ time to search for $N$ points and $F$ answers
- Ex: Find all points in ([35:80],[50: $\infty$ ])



## EXAMPLE OF A SEARCH IN A PRIORITY SEARCH TREE

- Find all points in ([35:83],[50: $\infty$ ])



## EXAMPLE OF A SEARCH IN A PRIORITY SEARCH TREE

- Find all points in ([35:83],[50: $\infty$ )


1. Find nearest common ancestor - i.e., A

- output toronto $(62,77)$ since 62 is in $[35: 80]$ and $77 \geq 50$


## EXAMPLE OF A SEARCH IN A PRIORITY SEARCH TREE

- Find all points in ([35:83],[50: $\infty$ ])


1. Find nearest common ancestor - i.e., A

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2. Process left subtree of $A$ (i.e., $B$ )

- cease processing as $45<50$


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- Find all points in ([35:83],[50: $\infty$ )


1. Find nearest common ancestor - i.e., A

- output toronto $(62,77)$ since 62 is in $[35: 80]$ and $77 \geq 50$

2. Process left subtree of $A$ (i.e., $B$ )

- cease processing as $45<50$

3. Process right subtree of A (i.e., C )

- output $(82,65)$ as $65 \geq 50$ and 82 is in [35:83]


## EXAMPLE OF A SEARCH IN A PRIORITY SEARCH TREE

- Find all points in ([35:83],[50: $\infty$ )


1. Find nearest common ancestor - ie., A

- output toronto $(62,77)$ since 62 is in $[35: 80]$ and $77 \geq 50$

2. Process left subtree of A (i.e., B)

- cease processing as $45<50$

3. Process right subtree of A (i.e., c)

- output $(82,65)$ as $65 \geq 50$ and 82 is in $[35: 83]$

4. Examine midrange value of c which is 84 and descend left subtree of c (i.e., F)

- cease processing since no point is associated with F meaning all nodes in the subtree have been examined


## RANGE PRIORITY TREES

- Variation on priority search tree
- Inverse priority search tree: heap node stores point with minimum $y$ coordinate value that has not been stored in a shallower depth in the tree (instead of maximum)
- Structure

1. sort all points by their $y$ coordinate value and store in leaf of a balanced binary tree such as range tree (single lines)

- no need to link leaf nodes unless search for all points in range of $x$ coordinate values

2. nonleaf node left sons of their father contains a priority search tree of points in subtree (double lines)
3. nonleaf node right sons of their father contains an inverse priority search tree of points in subtree (double lines)

- $O\left(N \cdot \log _{2} N\right)$ space and time to build for $N$ points

- Procedure

1. find nearest common ancestor of BY and EY - ie., Q
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SEARCHING A RANGE PRIORITY TREE ([bx:Ex],[by:Ey])

- Procedure

1. find nearest common ancestor of $B Y$ and $E Y$ - i.e., $Q$
2. all points in LEFT(Q) have $y$ coordinate values $<E Y$

- want to retrieve just the ones $\geq B Y$
- find them with ([BX:EX],[BY: $\infty]$ ) on priority tree of LEFT(Q)
- priority tree is good for retrieving all points with a specific lower bound as it stores an upper bound and hence irrelevant values can be easily pruned


SEARCHING A RANGE PRIORITY TREE ([bx:Ex],[by:ey])

## - Procedure

1. find nearest common ancestor of $B Y$ and $E Y$ - i.e., $Q$
2. all points in LEFT(Q) have $y$ coordinate values $<E Y$

- want to retrieve just the ones $\geq B Y$
- find them with ([BX:EX],[BY: $\infty$ ]) on priority tree of LEFT(Q)
- priority tree is good for retrieving all points with a specific lower bound as it stores an upper bound and hence irrelevant values can be easily pruned

3. all points in RIGHT(Q) have $y$ coordinate values $>B Y$

- want to retrieve just the ones $\operatorname{SEY}$
- find them with ([BX:EX],[->:EY]) on the inverse priority tree of RIGHT(Q)
- inverse priority tree is good for retrieving all points with a specific upper bound as it stores a lower bound and hence irrelevant values can be easily pruned



## SEARCHING A RANGE PRIORITY TREE ([bx:Ex],[bY:Ey])

- Procedure

1. find nearest common ancestor of BY and EY - i.e., Q
2. all points in LEFT(Q) have $y$ coordinate values $<E Y$

- want to retrieve just the ones $\geq B Y$
- find them with ([BX:EX],[BY: $\infty]$ ) on priority tree of LEFT(Q)
- priority tree is good for retrieving all points with a specific lower bound as it stores an upper bound and hence irrelevant values can be easily pruned

3. all points in RIGHT(Q) have $y$ coordinate values >BY

- want to retrieve just the ones $\leq E Y$
- find them with ([BX:EX],[->:EY]) on the inverse priority tree of RIGHT(Q)
- inverse priority tree is good for retrieving all points with a specific upper bound as it stores a lower bound and hence irrelevant values can be easily pruned
- $O\left(\log _{2} N+F\right)$ time to search for $N$ points and $F$ answers



## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])



## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

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1. Find nearest common ancestor of 15 and 45 - i.e., A

## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])


1. Find nearest common ancestor of 15 and 45 - i.e., A
2. Search for ([25:60],[15: $\infty$ ]) in priority tree hanging from left son of A - i.e., B (all with $y \leq 45$ since a range tree in $y$ and in left subtree of a node with $y$ midrange value of 39)

## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])


1. Find nearest common ancestor of 15 and 45 - i.e., A
2. Search for ([25:60],[15: $\infty$ ]) in priority tree hanging from left son of A - i.e., B (all with $y \leq 45$ since a range tree in $y$ and in left subtree of a node with $y$ midrange value of 39)

- output $(27,35)$ as in range


## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])


1. Find nearest common ancestor of 15 and 45 - i.e., A
2. Search for ([25:60],[15: $\infty$ ]) in priority tree hanging from left son of A - i.e., B (all with $y \leq 45$ since a range tree in $y$ and in left subtree of a node with $y$ midrange value of 39)

- output $(27,35)$ as in range
- reject left subtree as 10 < lower limit of $y$ range


## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])


1. Find nearest common ancestor of 15 and 45 - i.e., A
2. Search for ([25:60],[15: $\infty$ ]) in priority tree hanging from left son of A - i.e., B (all with $y \leq 45$ since a range tree in $y$ and in left subtree of a node with $y$ midrange value of 39)

- output $(27,35)$ as in range
- reject left subtree as 10 < lower limit of $y$ range
- reject items in right subtree as out of $x$ range


## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])


1. Find nearest common ancestor of 15 and 45 - i.e., A
2. Search for ([25:60],[15: $\infty$ ]) in priority tree hanging from left son of A - i.e., B (all with $y \leq 45$ since a range tree in $y$ and in left subtree of a node with $y$ midrange value of 39)

- output $(27,35)$ as in range
- reject left subtree as 10 < lower limit of $y$ range
- reject items in right subtree as out of $x$ range

3. Search for ([25:60],[->:45]) in inverse priority tree hanging from right son of A - i.e., c (all with $y \geq 15$ since in right subtree of a node with $y$ midrange value of 39)

## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])


1. Find nearest common ancestor of 15 and 45 - i.e., A
2. Search for ([25:60],[15: $\infty$ ]) in priority tree hanging from left son of A - i.e., B (all with $y \leq 45$ since a range tree in $y$ and in left subtree of a node with $y$ midrange value of 39)

- output $(27,35)$ as in range
- reject left subtree as 10 < lower limit of $y$ range
- reject items in right subtree as out of $x$ range

3. Search for ([25:60],[ $-\infty: 45]$ ) in inverse priority tree hanging from right son of A - i.e., c (all with $y \geq 15$ since in right subtree of a node with $y$ midrange value of 39)

- output $(35,42)$ as in range


## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])


1. Find nearest common ancestor of 15 and 45 - i.e., A
2. Search for ([25:60],[15: $\infty$ ]) in priority tree hanging from left son of A - i.e., B (all with $y \leq 45$ since a range tree in $y$ and in left subtree of a node with $y$ midrange value of 39)

- output $(27,35)$ as in range
- reject left subtree as 10 < lower limit of $y$ range
- reject items in right subtree as out of $x$ range

3. Search for ([25:60],[- -45$]$ ) in inverse priority tree hanging from right son of A - i.e., c (all with $y \geq 15$ since in right subtree of a node with $y$ midrange value of 39)

- output $(35,42)$ as in range
- reject unreported items in left subtree as out of $x$ range


## EXAMPLE OF A SEARCH IN A RANGE PRIORITY TREE

- Find all points in ([25:60],[15:45])


1. Find nearest common ancestor of 15 and 45 - i.e., A
2. Search for ([25:60],[15: $\infty$ ]) in priority tree hanging from left son of A - i.e., B (all with $y \leq 45$ since a range tree in $y$ and in left subtree of a node with $y$ midrange value of 39)

- output $(27,35)$ as in range
- reject left subtree as 10 < lower limit of $y$ range
- reject items in right subtree as out of $x$ range

3. Search for ([25:60],[->:45]) in inverse priority tree hanging from right son of A - i.e., c (all with $y \geq 15$ since in right subtree of a node with $y$ midrange value of 39)

- output $(35,42)$ as in range
- reject unreported items in left subtree as out of $x$ range
- reject right subtree as 65 > upper limit of $y$ range

