R-TREE IMPLEMENTATION

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1. Program execution requirements

1.1. Program environment

OS Environment	MacOS		
CPU	Apple M2 Pro		
RAM	16GB		
Language	Python		
Version	Python 3.11.4		
Software	Visual Studio Code		

1.2. Input files and parameters

- Input files:
- "R_tree_Contruction.txt": dataset (points) file
- "200Range.txt": range query file
- The setting of the directory: in one folder

2. Program documentation

2.1. Program organisation

File Name	Description			
	The file script processes spatial range queries on a set of			
	data points. It reads points from			
	"R_tree_construction.txt" and 200 queries from			
"Squential_Scan.py"	"200Range.txt", performs a sequential scan to count			
	points within each query range, and outputs the results,			
	along with total and average query processing times, to			
	"Sequential_output.txt".			
	The file reads points from "R_tree_construction.txt" and			
(D)	200 queries from "200Range.txt". Then the data is split			
"Rtree.py"	into two sections to construct two separate R-trees using			
	point-based division. For each specified range in			

	"200_range.txt", it traverses the respective R-trees to			
	count the points within these ranges, outputting the totals			
	to "Rtree_output.txt".			
	Same with "Rtree.py". However, in Divide and Conquer,			
	the point set is divided into two halves, and then two			
"Rtree_DC.py"	separate R-trees are constructed for these subsets,			
	allowing for more efficient organization and query			
	processing of spatial data.			

Class	Description			
"Node"	Class(Node) in an R-tree serves as a building block for the tree structure, encapsulating attributes such as a unique identifier, a collection of child nodes, and the node's Minimum Bounding Rectangle (MBR). For leaf nodes, it additionally holds a list of data points, while all nodes maintain a reference to their parent node, facilitating efficient tree traversal.			
"RTree"	Class(RTree) encompasses functions crucial for constructing and managing the R-tree structure, including methods to insert data points, handle node overflows, and compute Minimum Bounding Rectangles (MBRs) for efficient spatial querying.			

2.2. Function description

Function Name (parameters)	Description (detailed information)		
	Initialize a new instance of the "Node" or		
init (self)	"Rtree".		
	Calculates and returns the half perimeter of		
perimeter(self)	the node's MBR.		
	Check if the node has more child nodes or		
	data points than a specified limit "B=4"		
	For leaf nodes: check if the length of		
	"data_points" exceeds "B=4".		
	• For internal nodes: check if the		
	length of "child_nodes" exceeds		
is_overflow(self)	"B=4".		
	Determine if the node is root by checking if		
	its parent is "None". A node is the root if it		
is_root(self)	has no parent ("self.parent" is "None").		
	Identify if the node is a leaf node, indicated		
is_leaf(self)	by having no child nodes.		
	Execute a query on the R-Tree starting from		
	the given note.		
	• For leaf nodes: check each point		
	directly.		
	• For internal nodes: check if the		
	child's MBRs intersect with the		
query(self, node, query)	query range.		
	Check if a given "point" lies within bounds		
is_covered(self, point, query)	of a "query" rectangle.		

	Determine if the MBR of a node intersects
	with a query rectangle, using center points
is_intersect(self, node, query)	and dimensions of the rectangles.
	Insert a data point "p" into the R-tree,
	starting at node "u". If a node overflows
	because of the insertion, it triggers overflow
insert(self, u, p)	handling.
	Select the most appropriate child node of
	"u" to insert a new point "p" to minimize the
choose_subtree(self, u, p)	increase in perimeter after insertion.
	Calculate the increase in the perimeter of a
	node's MBR if a new point "p" is added to
peri_increase(self, node, p)	it.
	Handle the overflow condition in a node "u"
	by splitting it into two nodes ("u1" and
	"u2")
	If "u" is the root, a new root is created
handle_overflow(self, u)	If "u1" and "u2" are added as child to parent
	Split an overflowing node "u" into two
	separate nodes ("best_s1" and "best_s2)
	If "u" is a leaf node, its data points are sorted
	by x and y coordinates.
	If "u" is an internal node, it sorts
split(self, u)	"child_nodes" by the corners of MBR
	Add a child node to a given node and
	updates the MBR of the parent node
add_child(self, node, child)	accordingly
add_data_point(self, node,	Add a data point to a node and update the
data_point)	node's MBR to encompass the new point.

	Update the MBR of a node, recalculating it		
	based on the current child nodes on data		
Update_mbr(self, node)	points.		

2.3. Program results

Sequential-Based Method:

```
Total time: 3.33951997756958 seconds
Average time per query: 0.0166975998878479 seconds
Points count:
12
10
5
13
13
7
10
3
12
8
6
8
6
6
```

R-Tree Based Method:

```
    ■ Rtree_output.txt

     The time-cost of building up the R-Tree is: 18.826796770095825 seconds.
     The query processing time is: 0.02881312370300293 seconds.
     The average query processing time per point is: 0.00014406561851501465 seconds.
     Points count:
    12
 7 10
     5
     6
10 13
     13
12 7
14 10
15 3
     8
   12
     6
    6
20 8
     6
     8
     6
25 6
```

Divide-and-Conquer:

```
■ Rtree_DC_output.txt

     The time-cost of building up the R-Tree is: 19.630218982696533 seconds.
     The query processing time is: 0.018118858337402344 seconds.
     The average query processing time per point is: 9.059429168701172e-05 seconds.
     Points count:
     9
     12
     10
     5
     6
     13
     13
     7
     10
     3
     8
     12
     6
     6
     8
     6
     7
     8
     6
     6
```

Utilizing an R-tree-based approach for search operations results in an average query processing speed that is 116 times quicker than the Sequential-Based Method. Moreover, incorporating Divide and Conquer into the R-tree-based search further reduces the average query time by 37% compared to using the R-tree method alone.

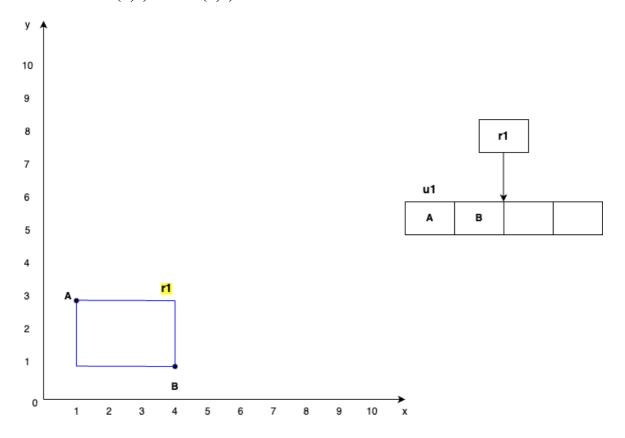
3. Analysing the Working of R-Tree

3.1. R-Tree establishment

3.1.1 R-Tree Insertion

Take B = 4 so we have 4 points in each node

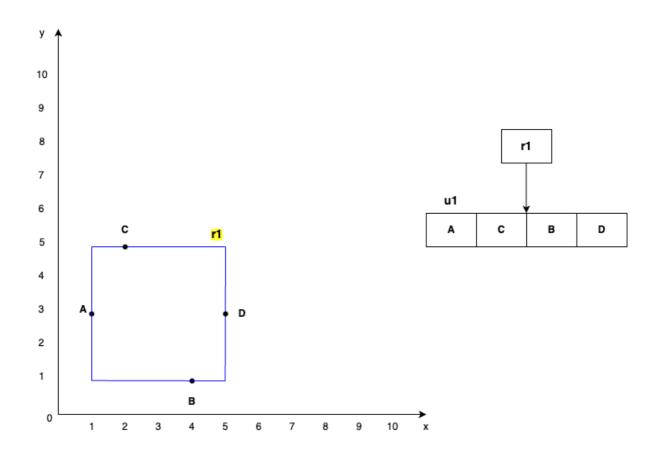
• Insert A (1,3) and B (4,1)



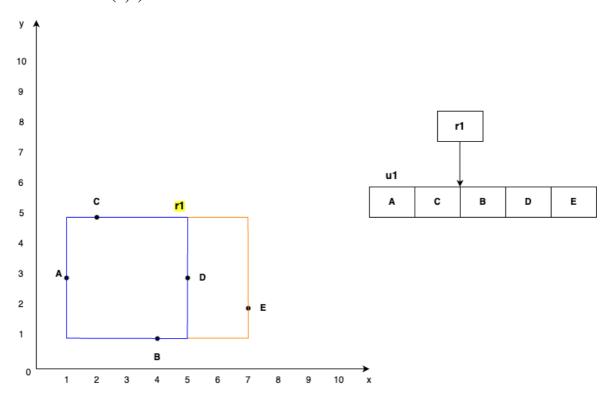
• Insert C (2,5) and D (5,3)

Before the R-tree establishment, the points are sorted based on their X and Y dimensions:

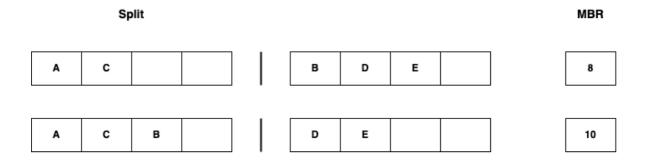
- Based on X-dimension: $A(1,3) \rightarrow C(2,5) \rightarrow B(4,1) \rightarrow D(5,3)$
- Based on Y-dimension: since yA = yD = 3, we do not consider to Y-dimension.



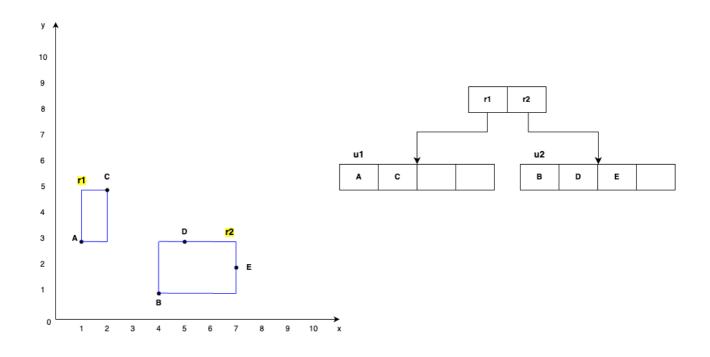
• Insert E (7,2)



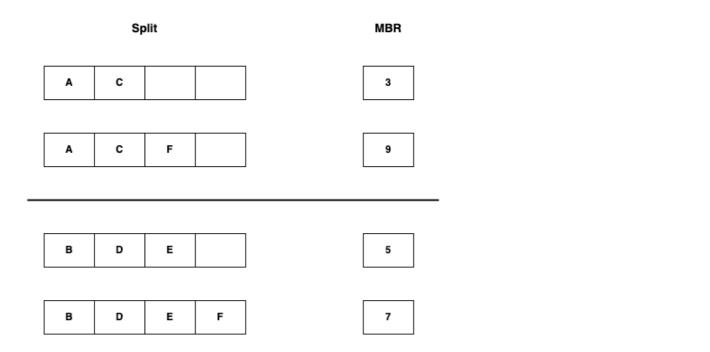
Due to the parameter B=4, the node experiences an overflow. By applying the rule 0.4B, which equates to 2, it's determined that each node must contain a minimum of 2 points.



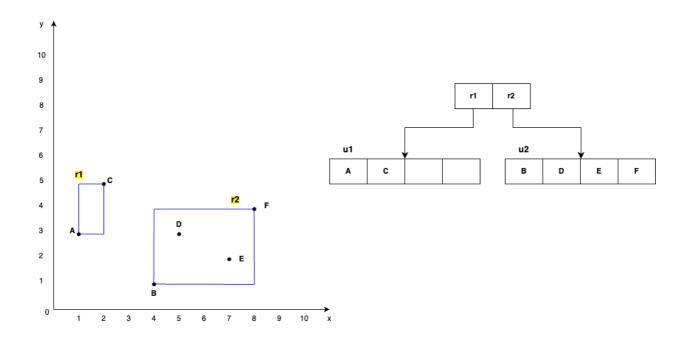
Between the two splits, the first option $(A, C) \mid (B,D, E)$ has a lower total perimeter of 8 compared to the second option's 10. Therefore, splitting the nodes into (A, C) and (B,D,E) after the insertion of E(7,2) would be the more efficient choice based on the perimeter criteria.



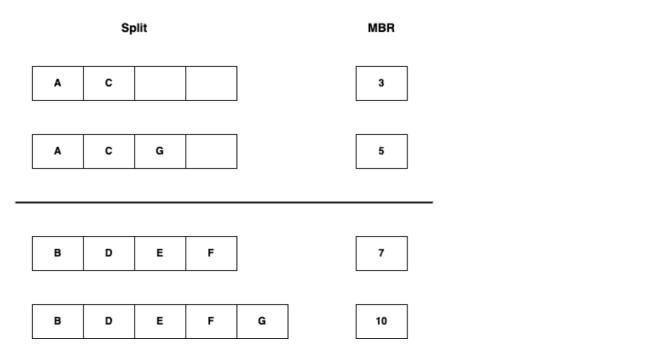
• Insert F (8,4)



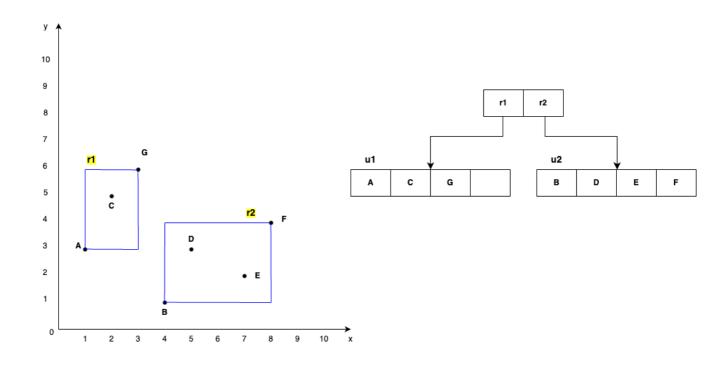
In this case, it's more efficient to add F to the group (B, D, E) since the MBR perimeter for (B, D, E, F) is smaller.



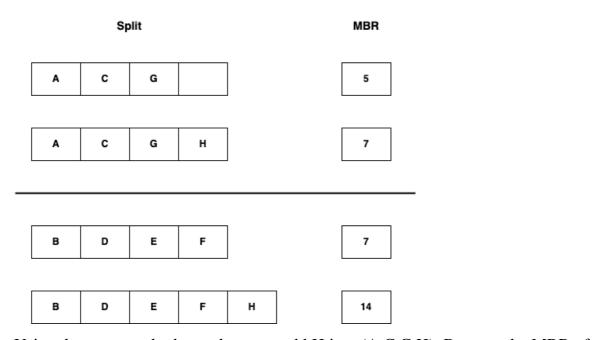
• Insert G (3,6)



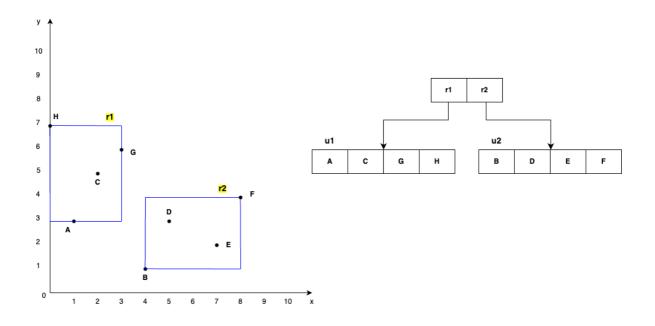
Using the same method, we can add G into node u1 (A, C, G) since MBR for (A, C, G) is smaller than that of (B, D, E, F, G).



• Insert H(0,7)



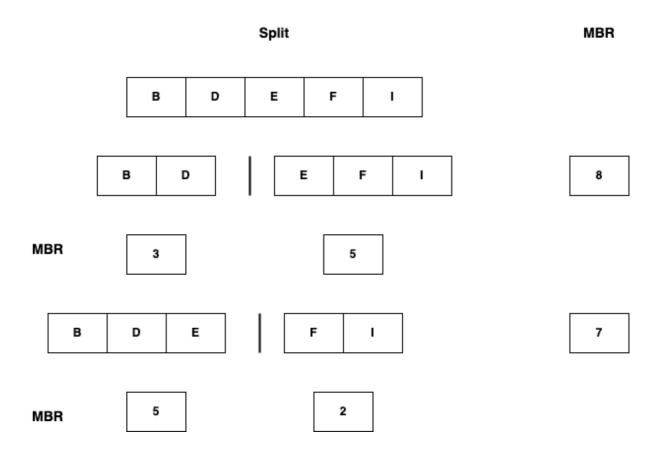
Using the same method, we choose to add H into (A,C,G,H). Because the MBR of (A, C, G, H) is 7 while MBR of (B, D, E, F, H) is 13.



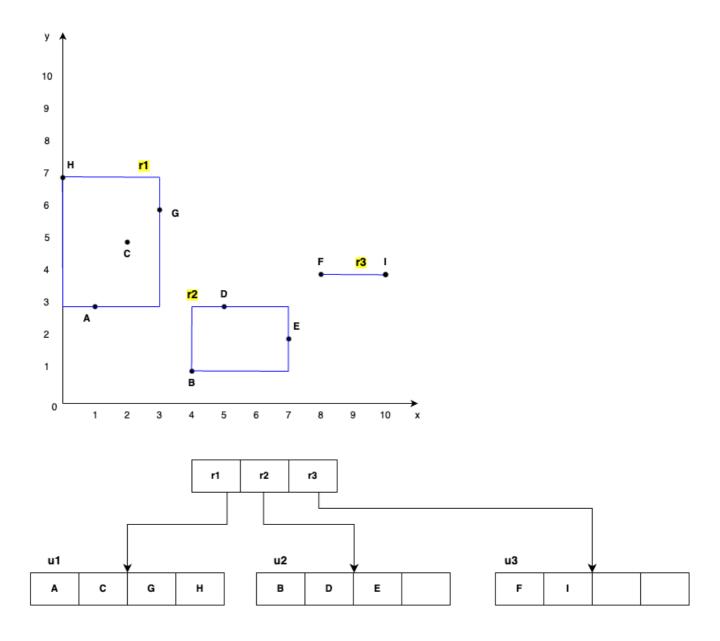
• Insert I (10,4)

	Sp	olit			MBR
А	С	G	н		7
А	С	G	н	ı	14
В	D	E	F		7
В	D	E	F	I	9

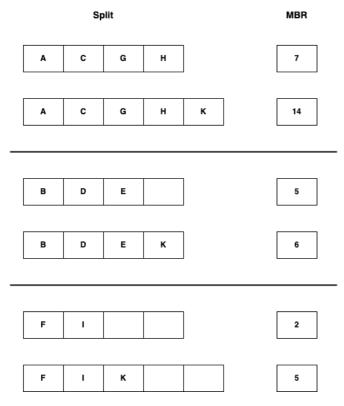
Using the same method, we choose the third split (B, D, E, F, I) which has a smaller MBR. However, (B, D, E, F, I) has 5 points while B = 4, leading to an overflow. Therefore, we need to split the node (B, D, E, F, I) into 2 nodes which have at least 2 points in each.



Using the same method, we choose the second split (B, D,E) and (F,I) with the smaller MBR.

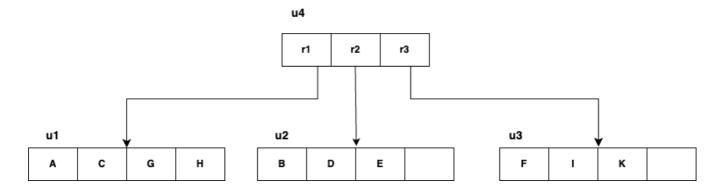


• Insert K (8,1)

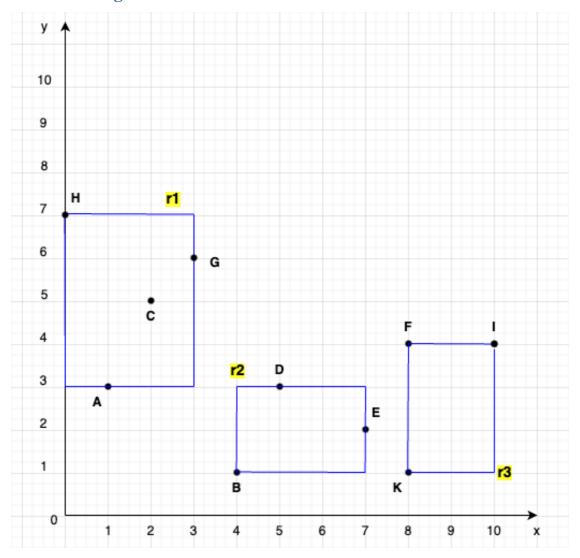


Using the same method, we get (F, I, K) has the smallest MBR.

This is the final R-tree



3.1.2 Bounding Boxes



3.2. Sequential-based search and R-Tree based search

3.2.1. Sequential-based search

For each point, check if its "x" coordinate lies between "x1" and its "y" coordinate lies between "y1" and "y2" of the query rectangle. If the condition is met, increment the count by 1.

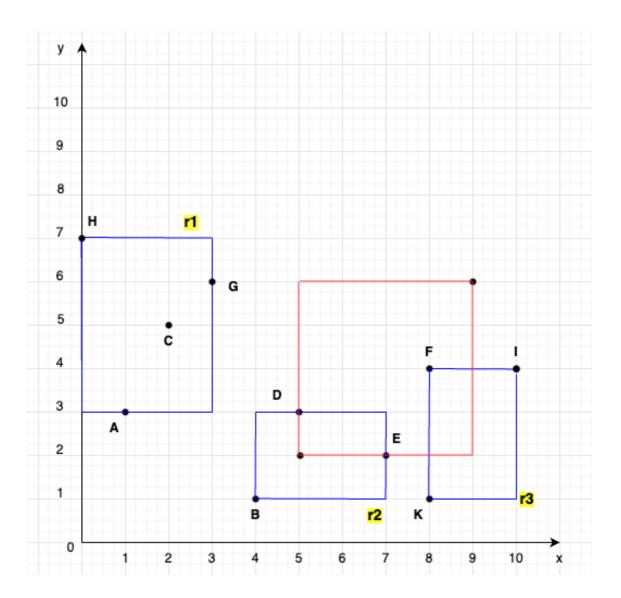
For the query range "x1 = 5", "y1 = 2", "x2 = 9", "y2 = 6", we would check each point:

Points	Check condition	Result
A (1, 3)	1 < 5	Not satisfy
B (4,1)	4 < 5	Not satisfy
C (2, 5)	2 < 5	Not satisfy
D (5, 3)	5 ≤5 ≤ 9	Satisfy
	$2 \le 5 \le 6$	
E (7, 2)	$5 \le 7 \le 9$	Satisfy
	$2 \le 2 \le 6$	
F (8, 4)	5 ≤ 8 ≤ 9	Satisfy
	$2 \le 4 \le 6$	
G (3, 6)	3 < 5	Not satisfy
H (0, 7)	0 < 5	Not satisfy
I (10, 4)	10 > 9	Not satisfy
K (8, 1)	$5 \le 8 \le 9$	Not satisfy
	1 < 2	

From the above checks, only the points D (5, 3), E (7, 2) and F (8, 4) lie within the query box. Therefore, the total count of points within the query box is 3.

3.2.2. R-Tree based search

Based on the created bounding boxes, we check the intersection of the query range in the bounding boxes.



The R-tree-based search provides an optimized approach to spatial querying, as illustrated by the **Query Range** (red box) delineated by:

$$x1 = 5$$
, $y1 = 2$, $x2 = 9$, $y2 = 6$

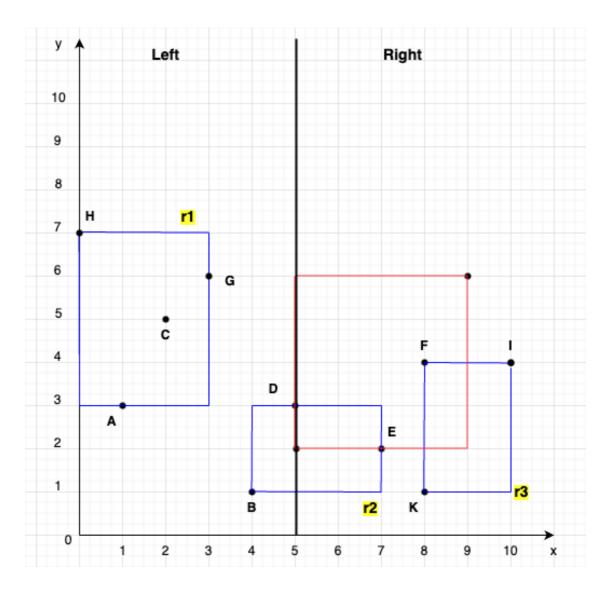
- For r1:
- Bounded by x1 = 0, y1 = 3, x2 = 3, y2 = 7
- The query does not overlap with r1 since the maximum x-coordinate of r1 is less than the minimum x-coordinate of the query range.
- ⇒ A, C, G, and H are excluded from the results.A,C,G,H do not lie within the query range

- For r2:
- Bounded by x1 = 4, y1 = 1, x2 = 7, y2 = 3
- \Rightarrow The query intersects r2 at its upper right boundary.
- \Rightarrow D (5,3) is a part of the query result.
 - For r3:
 - Bounded by x1 = 8, y1 = 1, x2 = 10, y2 = 4
 - The query encompasses a significant part of r3
- \Rightarrow E (7, 2) and F (8, 4) are encapsulated within the query range.

Therefore, D (5,3), E (7, 2) and F (8, 4) lie within the query range.

Thus, when using the R-tree-based search, the algorithm efficiently identifies the bounding boxes (MBRs) that intersect with the query range. By examining only the points within these intersecting MBRs, we've quickly identified that points D, E, and F are the ones that lie within the given query range. This process showcases the efficiency and speed advantage of using an R-tree based search for spatial queries compared to a full Sequential-based search across all points.

3.3. Divide-and-Conquer in R-Tree-based search



Given a dataset with 10 data points and a maximum x-value of 10, we utilize the Divide and Conquer approach in the R-tree-based search. The dataset is bifurcated at the midpoint x=5.

- Left Subset: Points with x < 5, namely A, B, C, G, H.
- **Right Subset:** Points with $x \ge 5$, specifically D, E, F, I, K.

Considering the provided **Query Range** (red box) defined by:

- x1 = 5, y1 = 2
- x2 = 9, y2 = 6

- \Rightarrow The query range exclusively falls within the "Right" subset, given x1 = 5 (inclusive) and x2=9.
- ⇒ Consequently, points in the "Left" subset are not within the query range.
- \Rightarrow Evaluating the "Right" subset, data points D (5, 3), E (7, 2), and F (8, 4) are encapsulated by the query range.