





# Índice

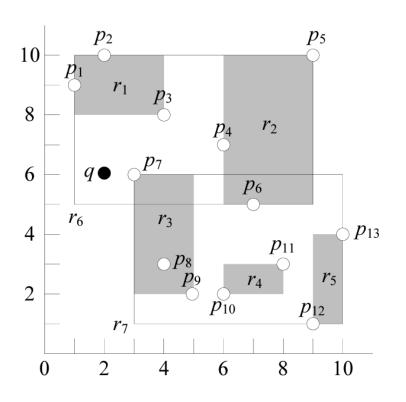
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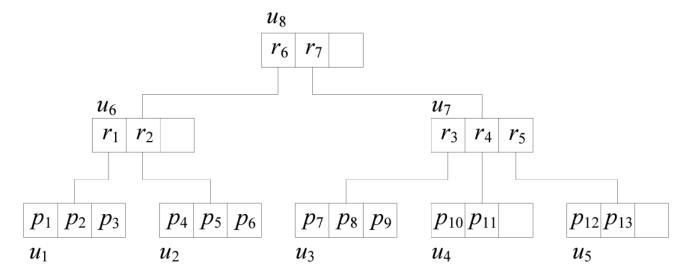




#### Branch and bound

```
algorithm BaB(u, q)
/* u is the node being accessed, q is the query point;
   p_{best} is a global variable that keeps the NN found so far;
   the algorithm should be invoked by setting u to the root */
   if u is a leaf node then
       if the NN of q in u is closer to q than p_{best} then
2.
3.
           p_{best} = the NN of q in u
   else
4.
       sort the MBRs in u in ascending order of their mindists to q
5.
       /* let r_1, ..., r_f be the sorted order */
       for i = 1 to f
6.
           if mindist(q, r_i) < ||q, p_{best}|| then
8.
               Bab(u_i,q)
               /* u_i is child node of r_i */
```





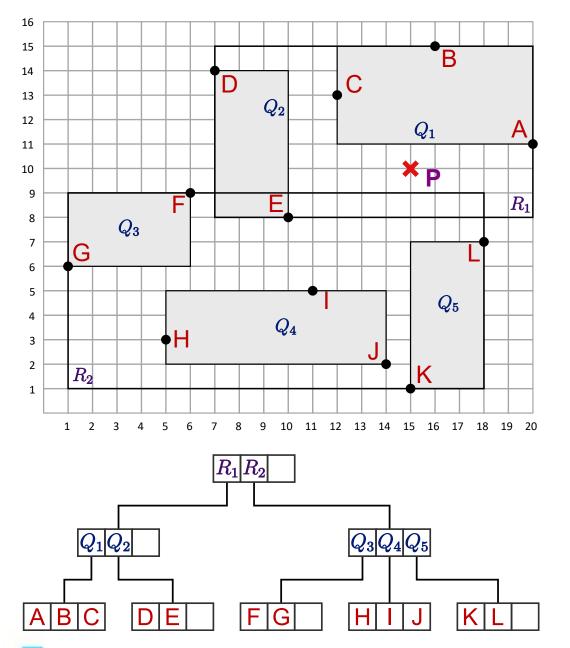


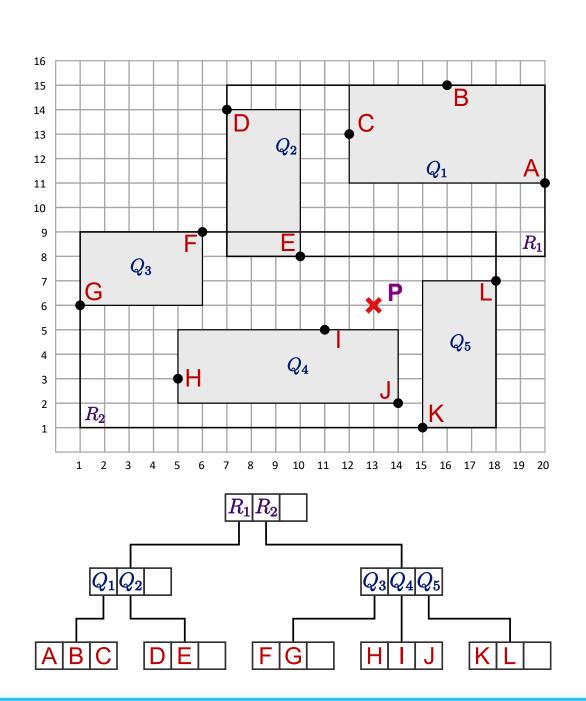
## Branch and bound

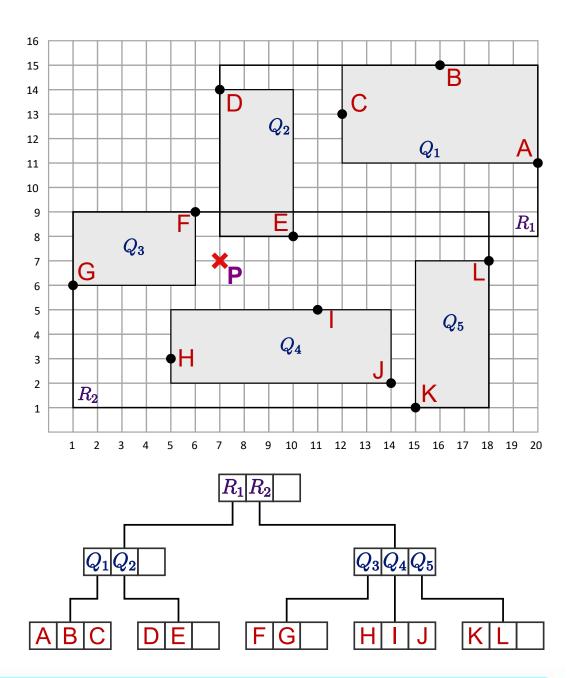
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#### Branch and bound





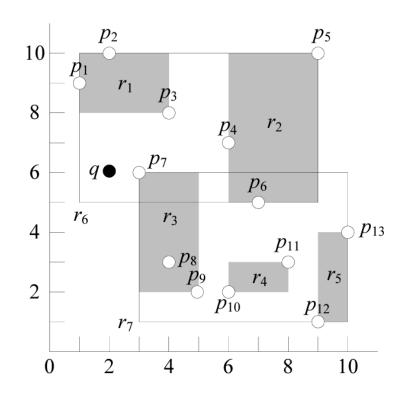


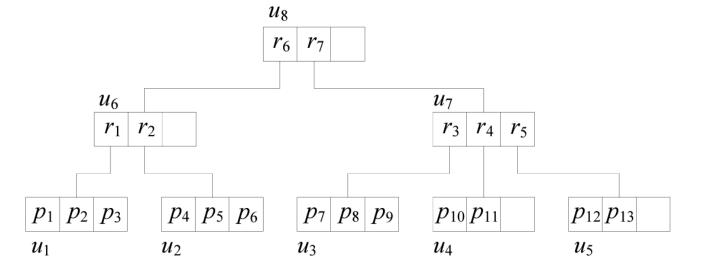




#### **Best** First

```
algorithm BF(q)
/* in the following H is a sorted list where each entry is an MBR
   whose sorting key in H is its mindist to q;
   p_{best} is a global variable that keeps the NN found so far. */
1. insert the MBR of the root in H
2. while ||q, p_{best}|| is greater than the smallest mindist in H
    /* if p_{best} = \emptyset, \|q, p_{best}\| = \infty */
       remove from H the MBR r with the smallest mindist
3.
       access the child node u of r
4.
       if u is an intermediate node then
           insert all the MBRs in u into H
6.
7.
       else
8.
           if the NN of q in u is closer to q than p_{best} then
               p_{best} = the NN of q in u
9.
```



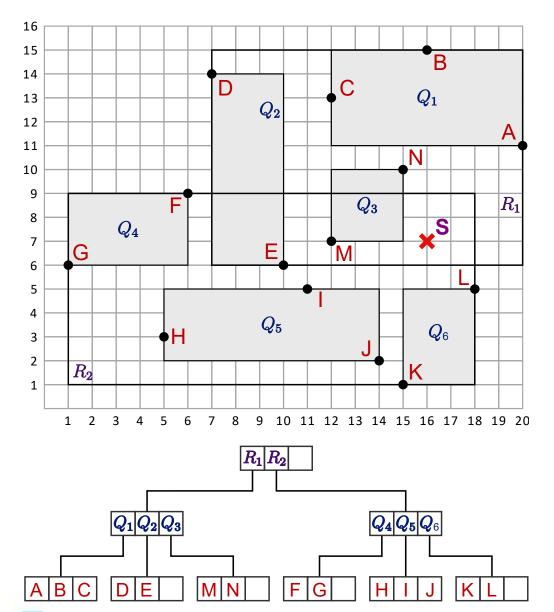


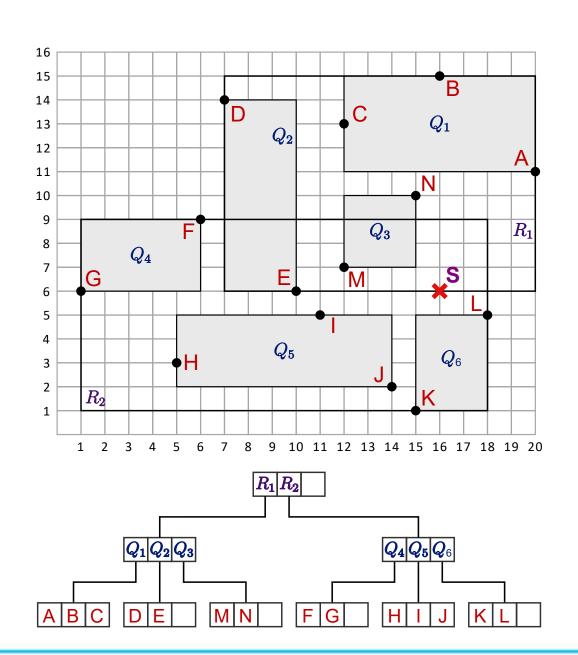


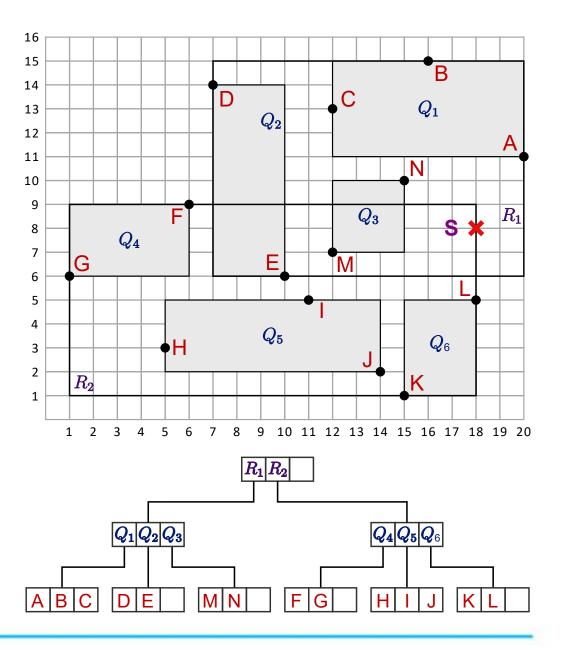
#### **Best** First



#### **Best** First











## R\*-Tree: split

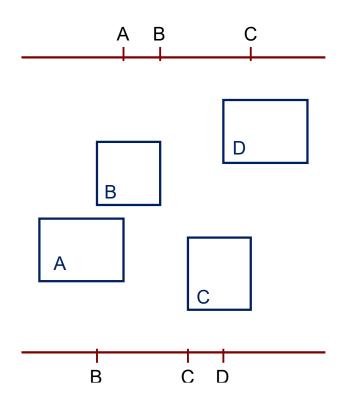
#### **Split**

- Seleccionar eje

a) Eje x

- Limite inferior
- Limite superior

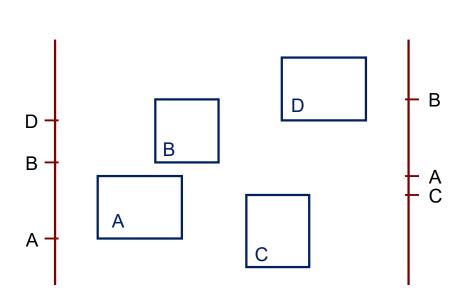
Suma de semiperímetros de las regiones



b) Eje y

- Limite inferior
- Limite superior

Suma de semiperímetros de las regiones



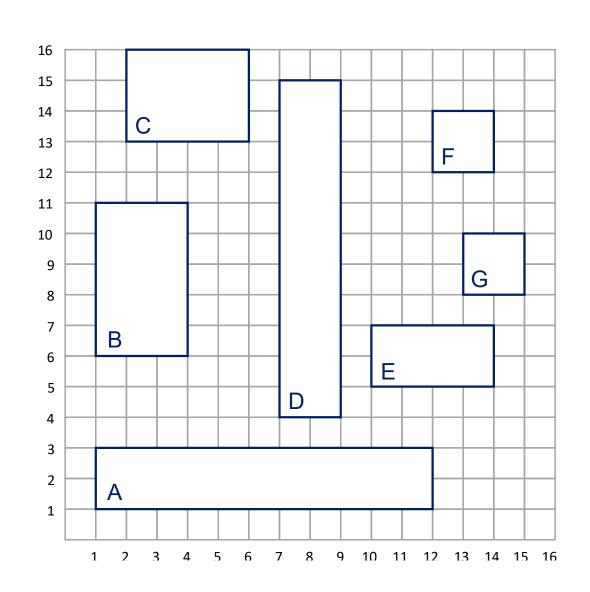


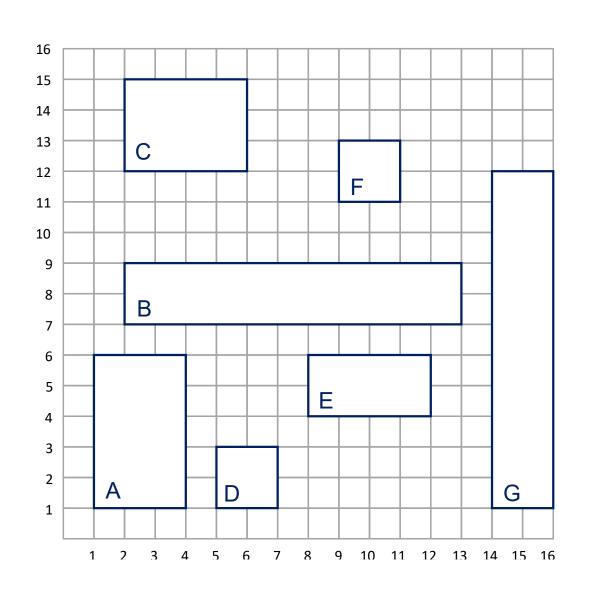
# R\*-Tree: split

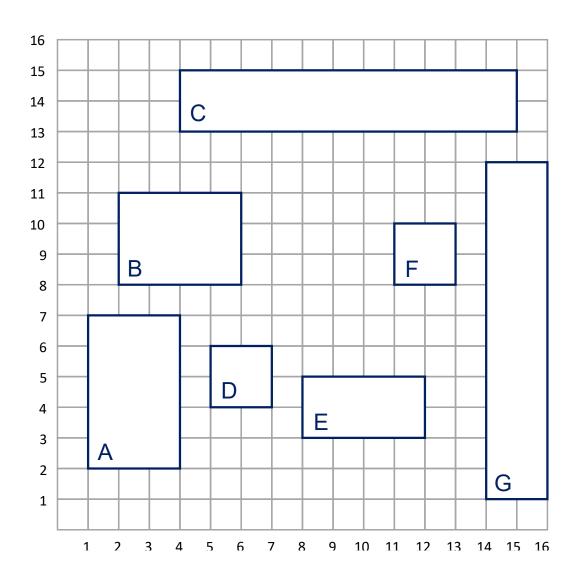
1 2 3



# R\*-Tree: split



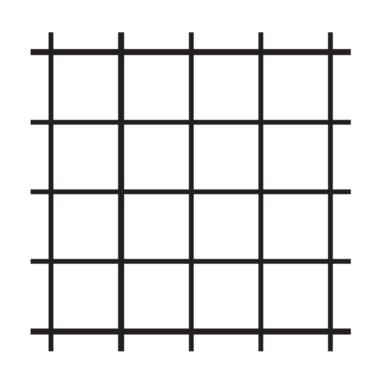








## Celdas/mosaico



Notación [4<sup>4</sup>]

Regular tiling Similar tiling

Mosaico ilimitado La celda atómica se puede descomponer

Número de adyacencia: 2

Orientación uniforme



## Celdas/mosaico

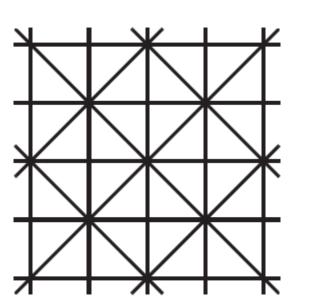
1

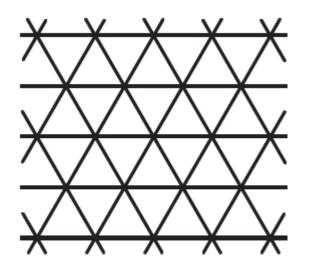
2

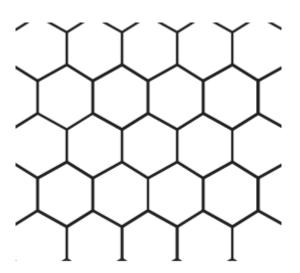
3



## Celdas/mosaico







Notación

Similar tiling?

Mosaico limitado?

Número de adyacencia:

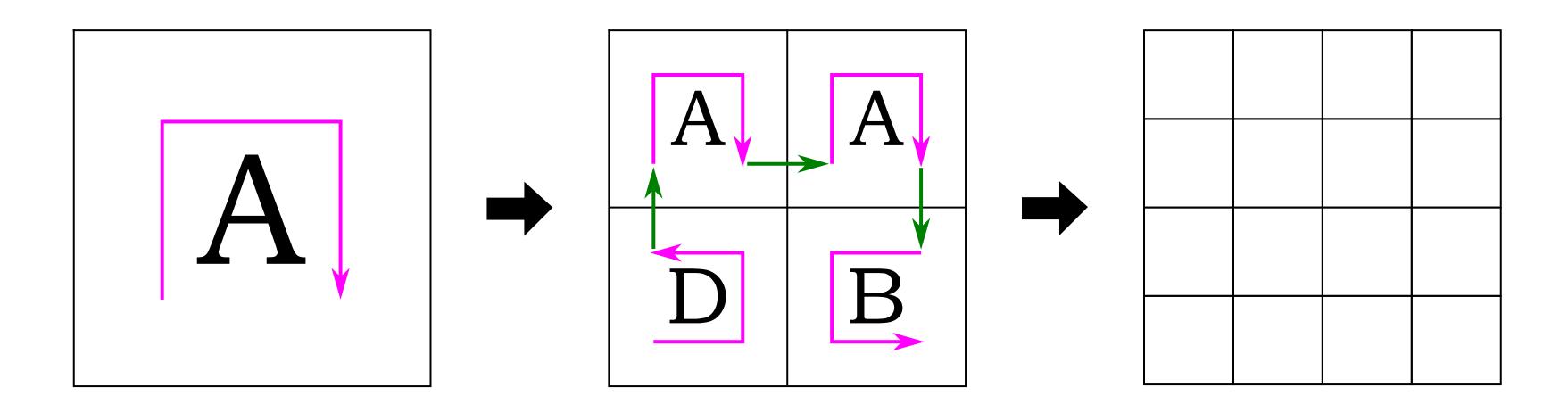
Orientación uniforme?:





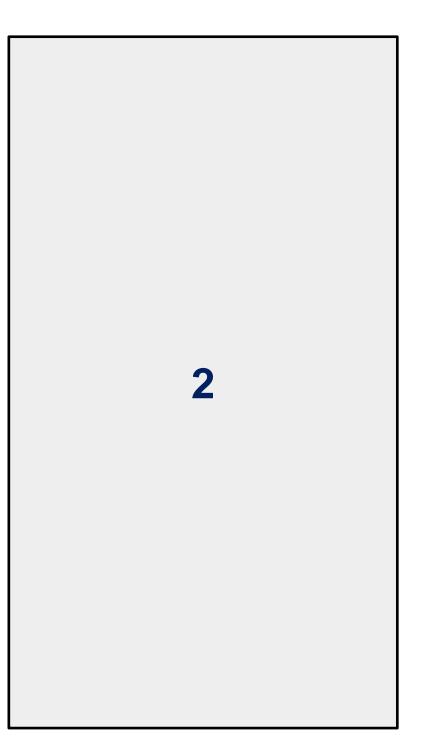


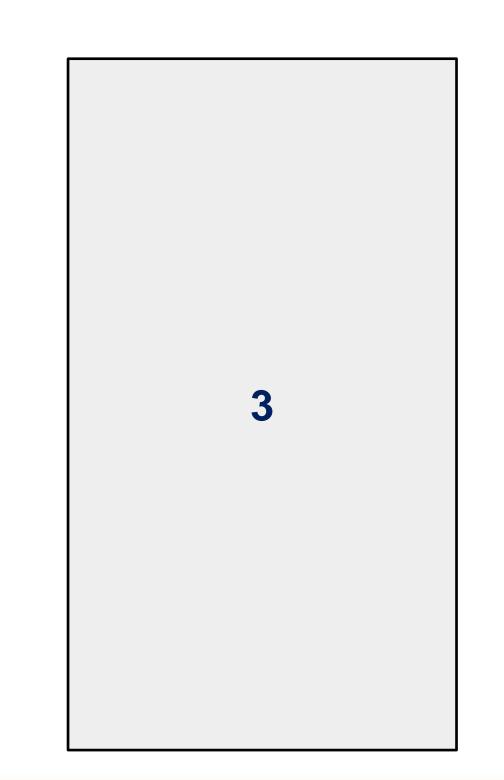
## Hilbert-curve





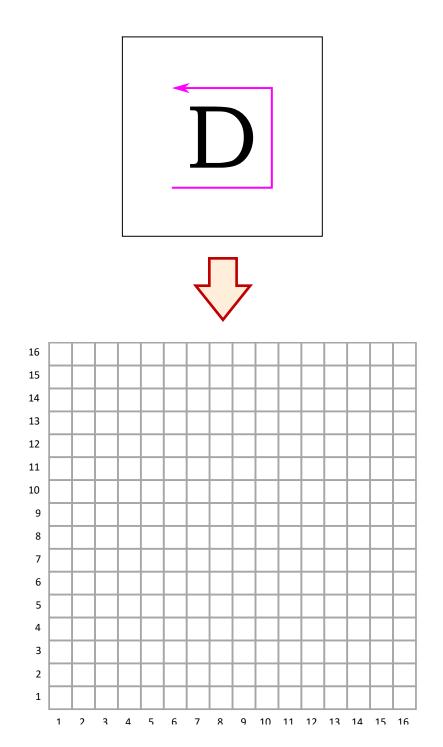
## Hilbert-curve

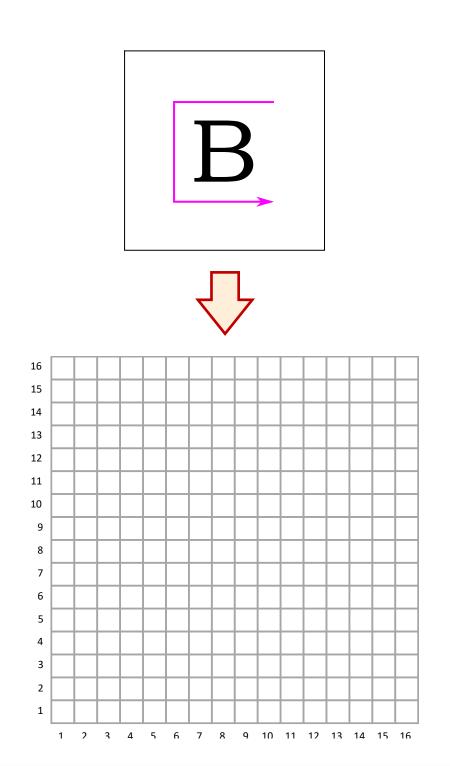


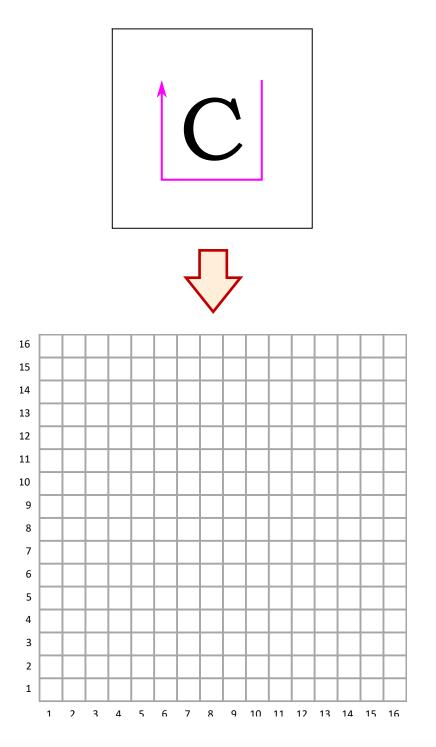




## Hilbert-curve











### **Z-**curve

	x: 0 000	1 001	2 010	3 011	1 1 4 1 100	5 101	6 110	7 111
y: 0 000	000000	000001	000100	000101	010000	010001	010100	010101
1 001	000010	000011	000110	000111	010010	010011	010110	010111
2 010	001000	001001	001100	001101	011000	011001	011100	011101
3 011	001010	001011	001110	001111	011010	011011	011110	Ollin
4 100	100000	100001	100100	100101	110000	110001	110100	110101
5 101	100010	100011	100110	100111	110010	110011	110110	110111
6 110	101000	101001	101100	101101	111000	111001	111100	111101
7 111	101010	101011	101/10	101111	111010	111011	111110	May



#### **Z-**curve

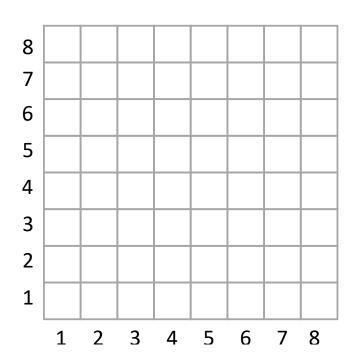
1

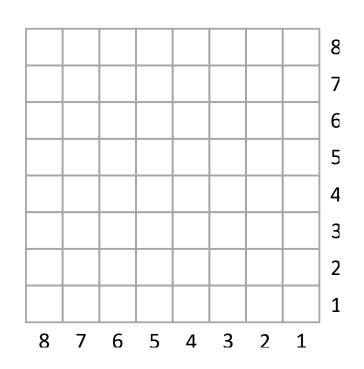
2

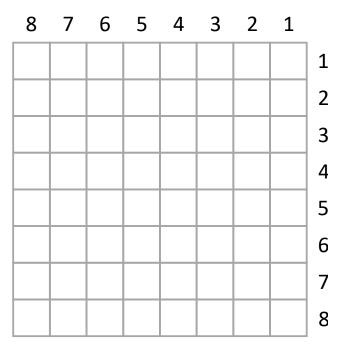
3



### **Z-**curve



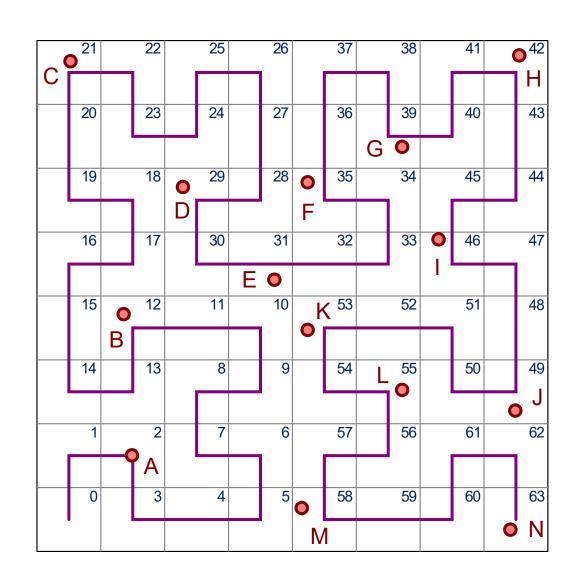


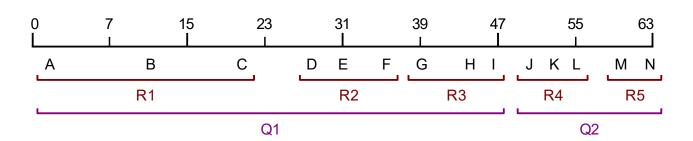


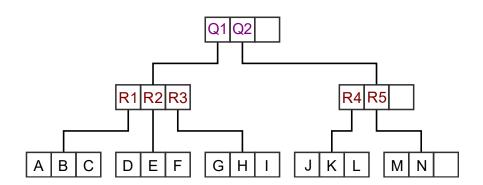




#### Hilbert R-Tree







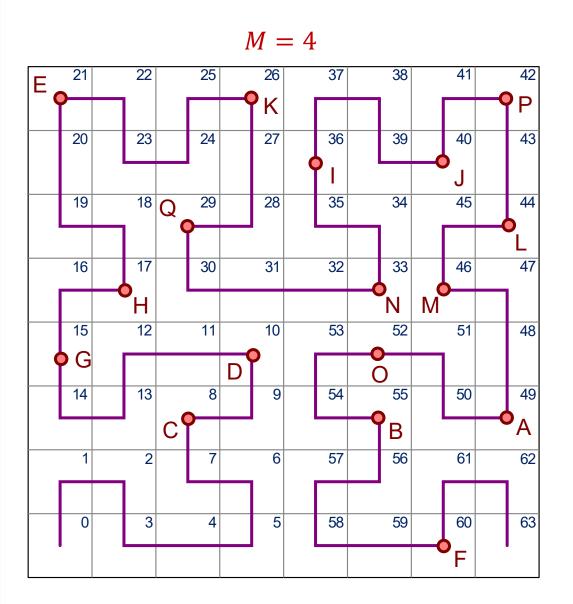


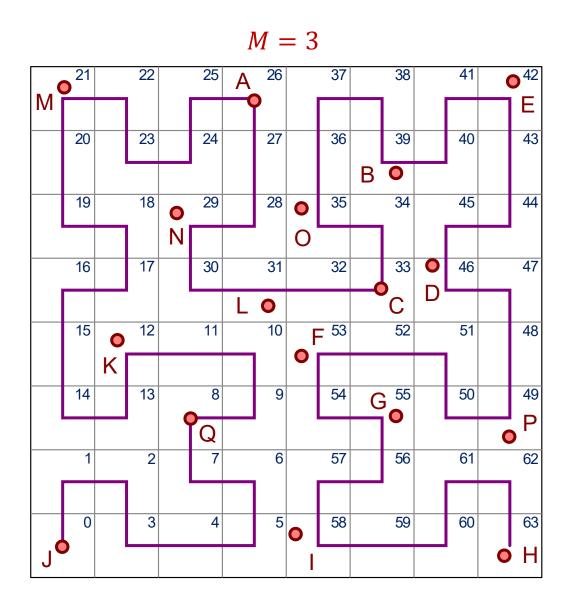
## Hilbert R-Tree

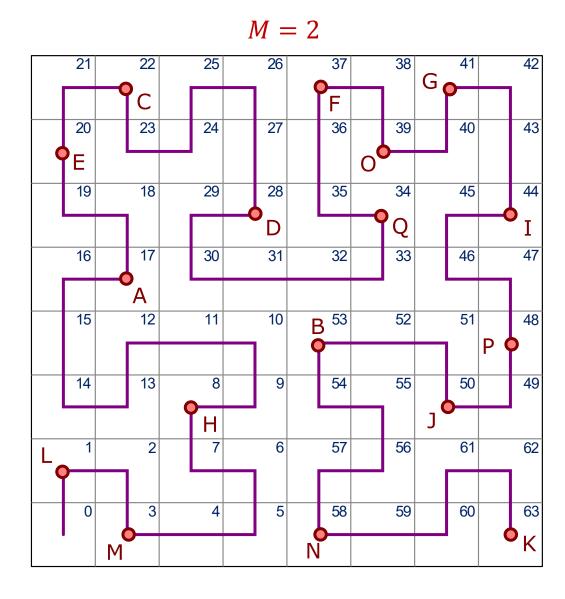




#### Hilbert R-Tree





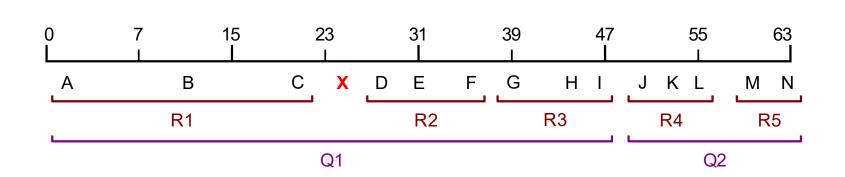


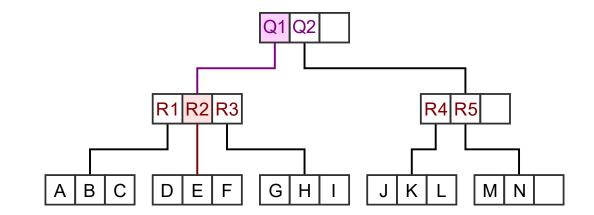




## Dynamic Hilbert R-Tree

Durante la inserción, se selecciona el nodo con LHV mínimo que supera al h-index del nuevo dato.





**Sobrecarga** Buscamos apoyo del hermano izquierdo.



Repartimos todos los datos entre **todos** los nodos

Pero, que hacemos si todos los nodos izquierdos están llenos?

Creamos nuevo nodo

... y repartimos todos los datos entre **todos** los nodos



# **Dynamic Hilbert** R-Tree





# **Dynamic Hilbert** R-Tree

