

- The divide and conquer algorithm design technique
- Analysing a divide and conquer algorithm: Mergesort
- Counting inversions
- Closest pair of points
- Convex hull

The divide and conquer algorithm design technique

- Suppose you have n items of input and the simplest technique to process it would be two nested for loops with a $\Theta(n^2)$ running time
- If n is small then this is fine
- With divide and conquer we instead aim at:
 - Divide in linear time the problem into two subproblems with $n/2$ items
 - Solve each subproblem
 - Combine the solutions to the subproblems in linear time into a solution for the n item problem
- The resulting running time becomes $\Theta(n \log n)$
- We will next study Mergesort

4 GHz modern CPU

n	n	$n \log n$	n^2	n^3	1.5^n	2^n	$n!$
10	2.5 ns	8.3 ns	25.0 ns	250.0 ns	14.4 ns	256.0 ns	907.2 μ s
11	2.8 ns	9.5 ns	30.2 ns	332.8 ns	21.6 ns	512.0 ns	10.0 ms
12	3.0 ns	10.8 ns	36.0 ns	432.0 ns	32.4 ns	1.0 μ s	119.8 ms
13	3.2 ns	12.0 ns	42.2 ns	549.2 ns	48.7 ns	2.0 μ s	1.6 s
14	3.5 ns	13.3 ns	49.0 ns	686.0 ns	73.0 ns	4.1 μ s	21.8 s
15	3.8 ns	14.7 ns	56.2 ns	843.8 ns	109.5 ns	8.2 μ s	5 min
16	4.0 ns	16.0 ns	64.0 ns	1.0 μ s	164.2 ns	16.4 μ s	1 hour
17	4.2 ns	17.4 ns	72.2 ns	1.2 μ s	246.3 ns	32.8 μ s	1.0 days
18	4.5 ns	18.8 ns	81.0 ns	1.5 μ s	369.5 ns	65.5 μ s	18.5 days
19	4.8 ns	20.2 ns	90.2 ns	1.7 μ s	554.2 ns	131.1 μ s	352.0 days
20	5.0 ns	21.6 ns	100.0 ns	2.0 μ s	831.3 ns	262.1 μ s	19 years
30	7.5 ns	36.8 ns	225.0 ns	6.8 μ s	47.9 μ s	268.4 ms	10^{15} years
40	10.0 ns	53.2 ns	400.0 ns	16.0 μ s	2.8 ms	5 min	10^{31} years
50	12.5 ns	70.5 ns	625.0 ns	31.2 μ s	159.4 ms	3.3 days	10^{47} years
100	25.0 ns	166.1 ns	2.5 μ s	250.0 μ s	3 years	10^{13} years	10^{141} years
1000	250.0 ns	2.5 μ s	250.0 μ s	250.0 ms	10^{159} years	10^{284} years	huge
10^4	2.5 μ s	33.2 μ s	25.0 ms	4 min	huge	huge	huge
10^5	25.0 μ s	415.2 μ s	2.5 s	2.9 days	huge	huge	huge
10^6	250.0 μ s	5.0 ms	4 min	8 years	huge	huge	huge
10^7	2.5 ms	58.1 ms	7 hour	10^4 years	huge	huge	huge
10^8	25.0 ms	664.4 ms	28.9 days	10^7 years	huge	huge	huge
10^9	250.0 ms	7.5 s	8 years	10^{10} years	huge	huge	huge

Mergesort

- Mergesort is a stable sort algorithm
- Running time $\Theta(n \log n)$
- See `mergesort.c` e.g. in the book

Recurrence relation

- Swedish differensekvation or rekursionsekvation
- A **recurrence relation** or just **recurrence** is a set of equalities or inequalities such as

$$T(n) = \begin{cases} 0, & n = 1 \\ 2T(n/2) + n, & n > 1 \end{cases}$$

- The value of $T(n)$ is expressed using smaller instances of itself and a boundary value.
- To analyze the running time of a divide and conquer algorithm, recurrences are very natural
- But we want to have an expression for $T(n)$ in **closed form**
- Closed form means an expression only involving functions and operations from a generally accepted set — i.e. "common knowledge".
- Closed form can also be called **explicit form**
- So our next goal is to rewrite $T(n)$ into closed form

Mergesort recurrence

- $T(n)$ = max comparisons to mergesort n items
- Mergesort recurrence:

$$T(n) \leq \begin{cases} 0, & n = 1 \\ T(\lceil n/2 \rceil) + T(\lfloor n/2 \rfloor) + n, & n > 1 \end{cases}$$

- This is a simplification as can be seen if compared with the source code, but it is sufficiently accurate.
- We ignore ceil and floor:

$$T(n) \leq \begin{cases} 0, & n = 1 \\ 2T(n/2) + n, & n > 1 \end{cases}$$

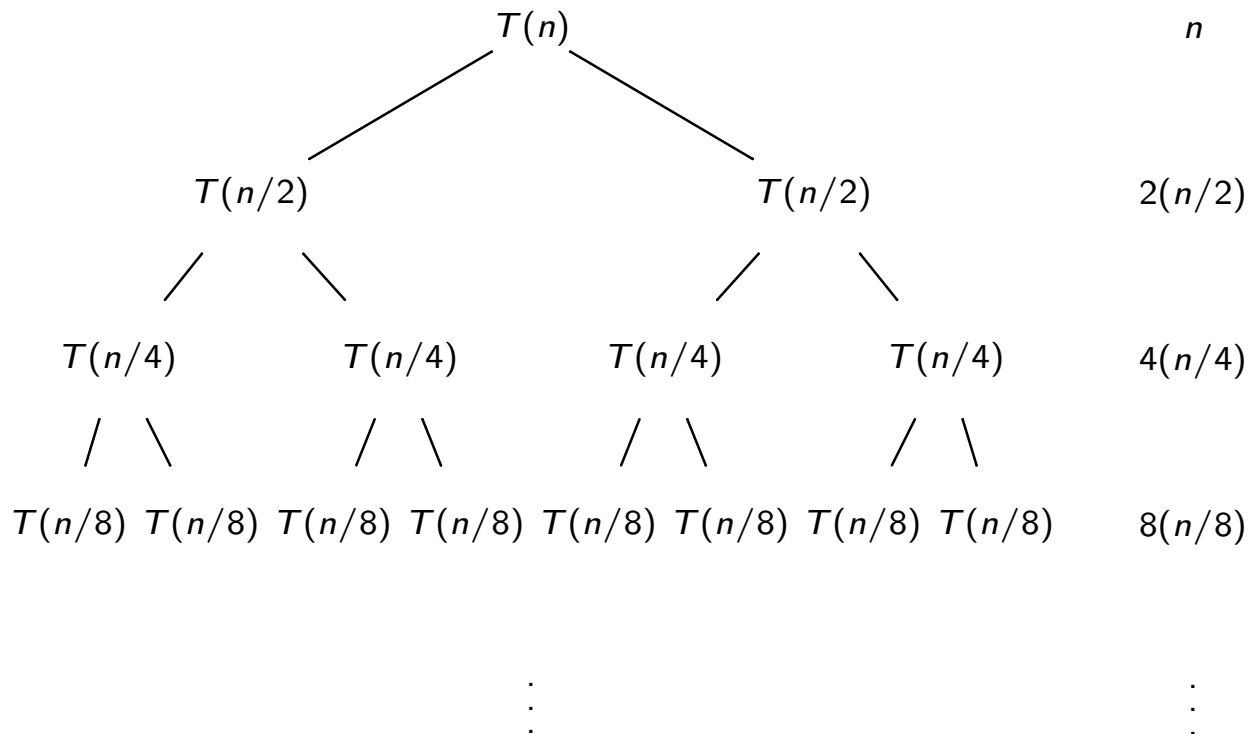
- We also assume n is a power of 2
- In the book it is shown that these simplifications do not affect our running time analysis

Rewriting a recurrence to closed form

- The easiest way to understand what the closed form is, may be to "expand" or "unroll" the recurrence and simply see what is happening
- Another way is to look at small inputs and try to guess the closed form
- When we have a guess which works for the small inputs, we then prove by induction that our guess is correct
- In both cases we prove our closed form by induction
- We will start with expanding $T(n)$

Expanding the recurrence and count

$$T(n) \leq \begin{cases} 0, & n = 1 \\ 2T(n/2) + n, & n > 1 \end{cases}$$



- Assume n is power of 2
- $\log_2 n$ levels
- n comparisons per level
- In total $n \log n$ comparisons
- $T(n) = n \log n$

Proof by induction

Lemma

The recurrence

$$T(n) = \begin{cases} 0, & n = 1 \\ 2T(n/2) + n, & n > 1 \end{cases}$$

has the closed form $T(n) = n \log_2 n$.

Proof.

- Recall $\log ab = \log a + \log b$, so $\log_2 2n = \log_2 n + \log_2 2 = \log_2 n + 1$, and $\log_2 n = \log_2 2n - 1$
- Induction on n .
- Base case: $n = 1$: $T(1) = 1 \log_2 1 = 0$
- Induction hypothesis: assume $T(n) = n \log_2 n$
- $T(2n) = 2T(n) + 2n = 2n \log_2 n + 2n = 2n(\log_2 n + 1) = 2n(\log_2 2n - 1 + 1) = 2n \log_2 2n$



Remark about previous proof

- Normally we assume $S(i)$ is true and prove $S(i + 1)$
- On previous slide we did not increment by one but rather doubled our variable
- We could have stated the lemma in terms of $S(i)$ and let $n = 2^i$
- Then we use induction on i and assume $S(i)$ and prove $S(i + 1)$

Looking at small inputs

$$T(n) \leq \begin{cases} 0, & n = 1 \\ 2T(n/2) + n, & n > 1 \end{cases}$$

- Let us try out some small values:

n	1	2	4	8	16	32	64
$T(n)$	0	2	8	24	64	160	384

- Can we identify a pattern?

n	1	2	4	8	16	32	64
$T(n)$	0	2	8	24	64	160	384
$T(n)/n$	0	1	2	3	4	5	6

$= \log_2(n)$

- So $T(n) = n \log_2 n$ is tempting to try to prove by induction, which we already know is true

The master theorem (MSc thesis by Dorothea Haken)

- There is a nice formula for finding $T(n)$ for many recursive algorithms:

$$\begin{aligned}T(1) &= 1 \\T(n) &= aT(n/b) + n^s.\end{aligned}$$

- There are three closed form solutions (for details, see the book):

$$T(n) = \begin{cases} O(n^s) & \text{if } s > \log_b a \\ O(n^s \log n) & \text{if } s = \log_b a \\ O(n^{\log_b a}) & \text{if } s < \log_b a. \end{cases}$$

- $T(n) = 2T(n/2) + n$. With $a = b = 2$ and $s = 1$, we have $\log_b a = \log_2 2 = 1 = s$, so $T(n) = O(n \log n)$.
- $T(n) = 2T(n/2) + \sqrt{n}$. With $a = b = 2$ and $s = 0.5$, we have $\log_b a = \log_2 2 = 1 > s$, so $T(n) = O(n)$.
- $T(n) = 4T(n/3) + n^2$. We have $\log_b a = \log_3 4 = \frac{\log_{10} 4}{\log_{10} 3} = 1.26 < s = 2$, so $T(n) = O(n^2)$.

Finding people with similar tastes

- Consider a category such as text editor, programming language, preferred tab width, or the 22 Mozart operas
- To compare how similar tastes within a category three people have, they can rank a list of say 5 operas A-E

Tintin: A D C E B

Captain Haddock: A C B D E

Bianca Castafiolen: A B D C E

- All agree opera A is best
- Who have most similar tastes?

Inversions

- Tintin: A D C E B
- Captain Haddock: A C B D E
- Bianca Castafiolen: A B D C E
- We have 5 positions in each list
- Start with Tintin's list and label each item 1, 2, ..., 5:
 Tintin: A D C E B
 Tintin: 1 2 3 4 5
- Then we put these labels according to Captain Haddock's ranking:
 Captain Haddock: 1 3 5 2 4
 a_1 a_2 a_3 a_4 a_5
- i and j are **inverted** if $i < j$ and $a_i > a_j$
- Inversions: (3,2), (5,2), and (5,4)
- The fewer inversions, the more similar tastes

Counting inversions

```
for (c = i = 0; i < n; i += 1)
    for (j = i+1; j < n; j += 1)
        if (a[i] > a[j])
            c += 1;
```

```
printf("%d inversions\n", c);
```

- Running time is $O(n^2)$
- How can we use divide and conquer to achieve $O(n \log n)$?
1 3 | 5 2 4
- Count inversions in left part
- Count inversions in right part
- Somehow combine these parts and add number of inversions...???

What can we do to simplify the problem?

- 1 3 | 5 2 4
- Assume you know there are no inversions in the left part and two in the right part
- It is OK to "destroy" the array, such as sorting it, if that helps...
- If modifying the array is forbidden, we can always make a copy and work with the copy instead
- Copying the array is fine since that is faster than $O(n \log n)$
- Copying the array is $O(n)$ but memory allocation can be costly so don't do it too much
- For Mergesort, it is non-trivial to not use a second array

Sorting the array

- By subarray is meant the part our recursive subproblem is going to work with

- Sorting the subarray **after** counting the inversions may help

- 1 3 | 5 2 4

- After having counted in the subarrays we have: 1 3 | 2 4 5

- Combining two sorted parts can be done in linear time as in

Mergesort

3	2	4	5	1
3		4	5	1 2

The 2 was inverted with each remaining in left part — only the 3 in this example so one inversion is counted when the parts are combined

	4	5	1	2	3
		5	1	2	3 4
			1	2	3 4 5

- In total 3 inversions

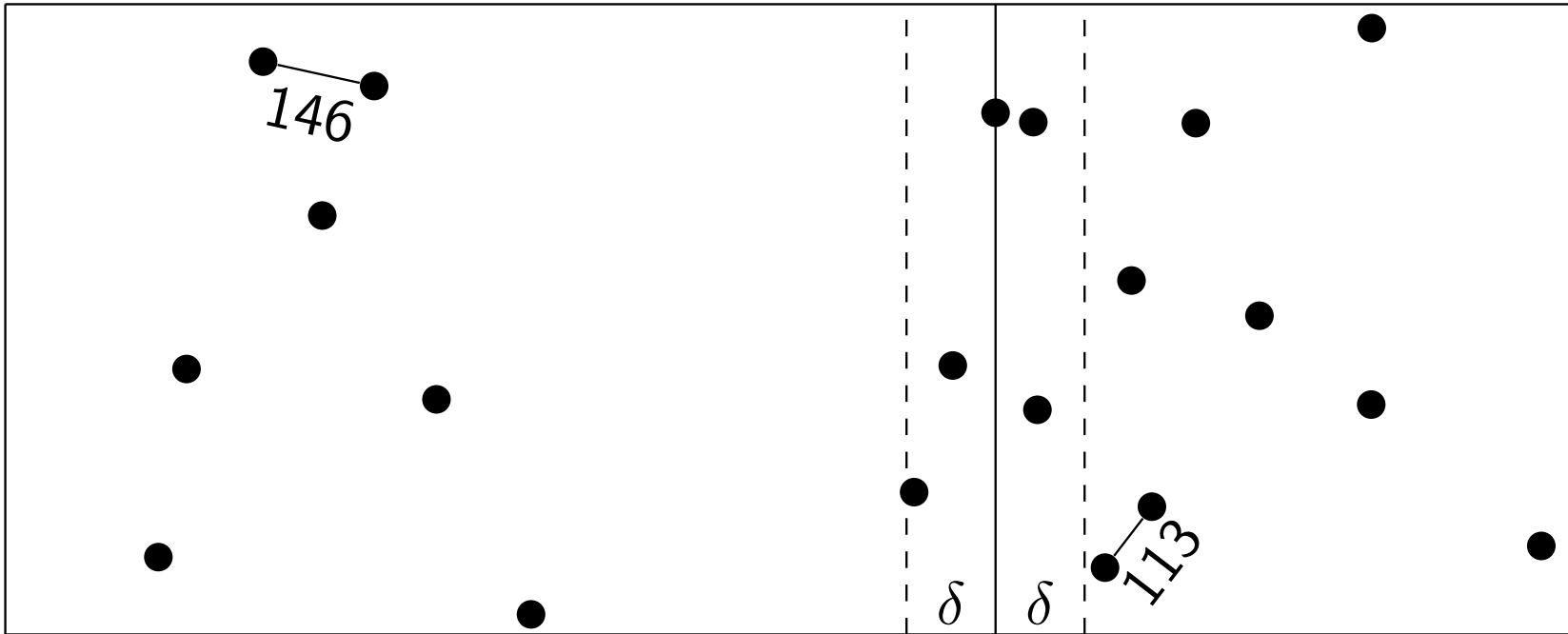
Implementing the $n \log n$ algorithm

- As always: first make a simple reference implementation that can be used to verify the correctness of a faster implementation
- In this case the n^2 algorithm is ideal if used with small inputs

Closest points in a plane

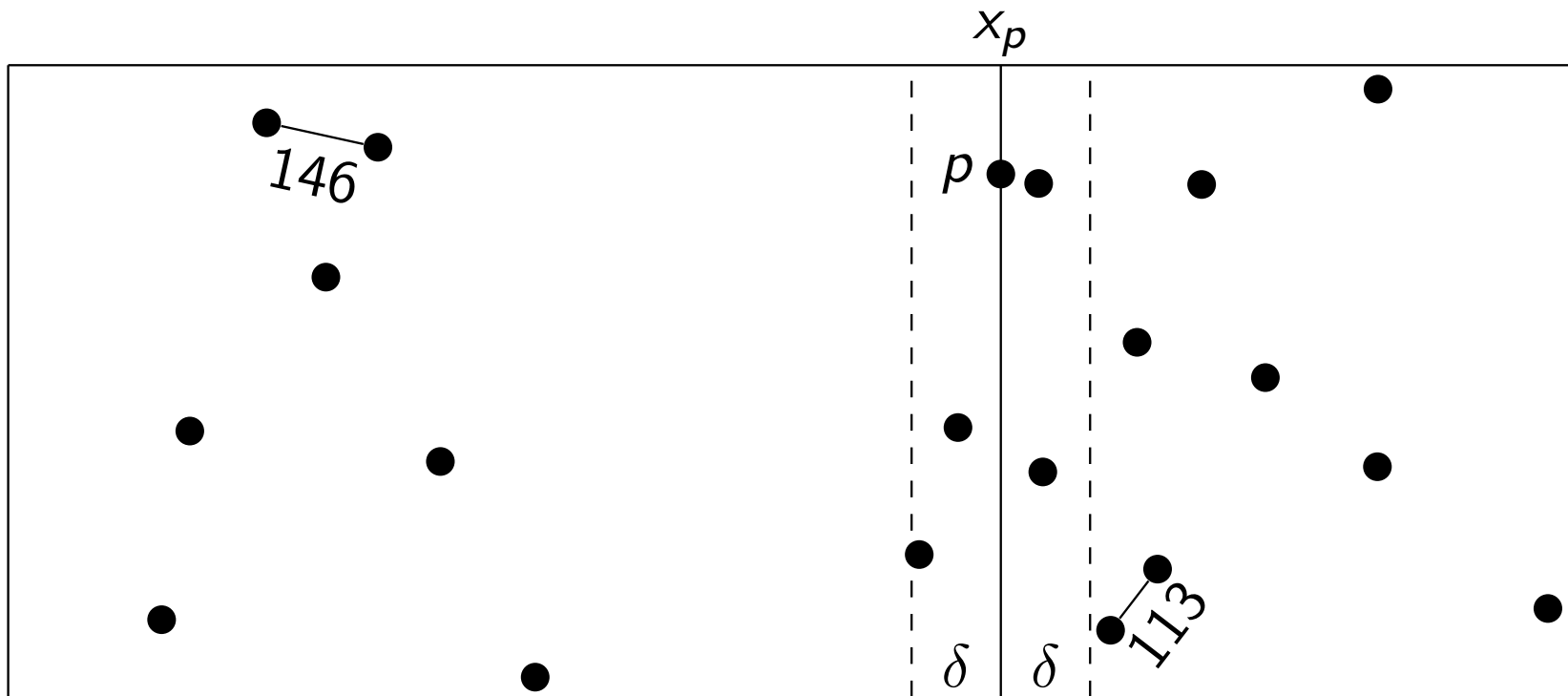
- This field is called **computational geometry**
- Consider n points (x_i, y_i) in a plane
- We want to find which points are closest
- Comparing all points with each other in an n^2 algorithm is simple
- But comparing points "obviously" far from each other is a waste
- How can divide and conquer be used to find an $n \log n$ algorithm?
- We cut the plane in two halves and find closest points in each half
- We have then three categories of point pairs which can be closest:
 - 1 Point pairs in the left half
 - 2 Point pairs in the right half
 - 3 Point pairs with one point in the left and the other in the right half
- Can we find close points from the last category in linear time???

An example



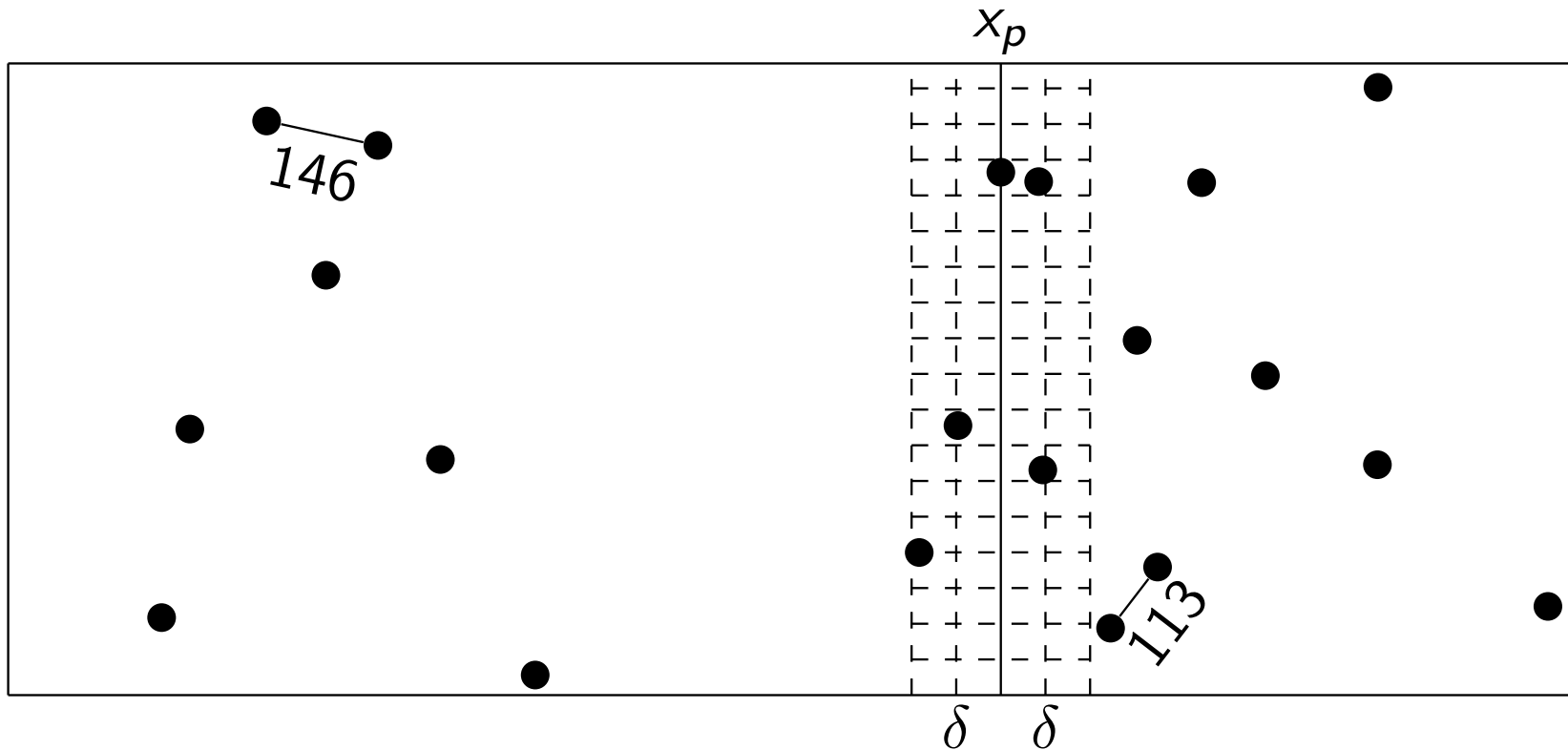
- We cut the plane in two halves with 10 points in each half
- We compute the nearest points in each half
- $\delta = \min(146, 113)$
- We only have to consider points within δ from the vertical line
- If there are none, then δ is the answer
- If there are, then they must be checked with points from the other side which also must be within δ from the vertical line, of course

Combining



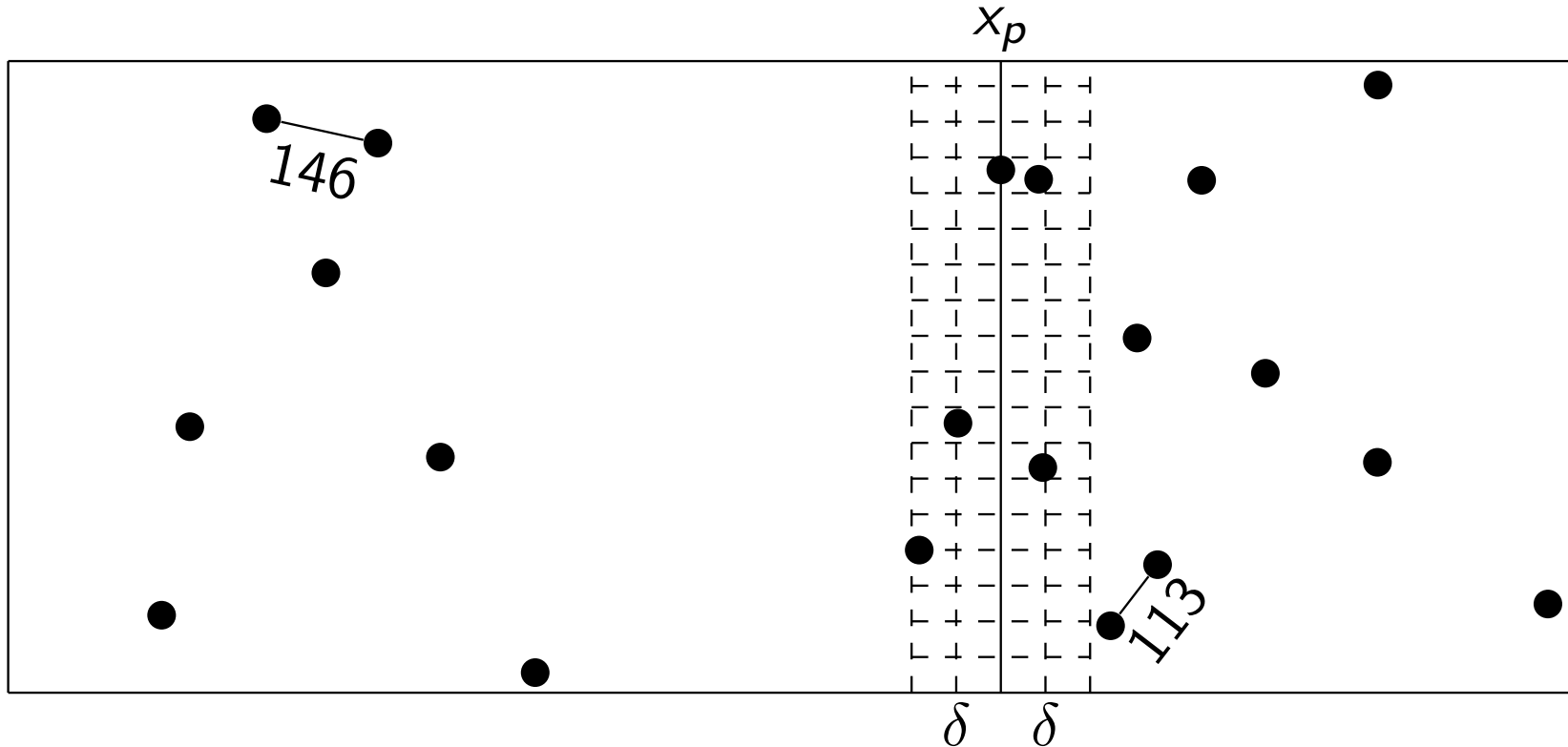
- The point p on the vertical line x_p belongs to the left half but there could also be points in the right half with the same x-coordinate
- Let the set S consist of all points with a distance within δ from the line x_p , (5 points here)
- Clearly it is sufficient to compare only points q and r from S such that p comes from the left half and q from the right part

Combining



- Each dashed box has a side of $\delta/2$
- How many points can each such box contain at most?
- The diagonal of a dashed box is $\sqrt{2} \times \delta/2 < \delta$
- With two points in a dashed box, their distance would be less than δ so at most one point

Combining

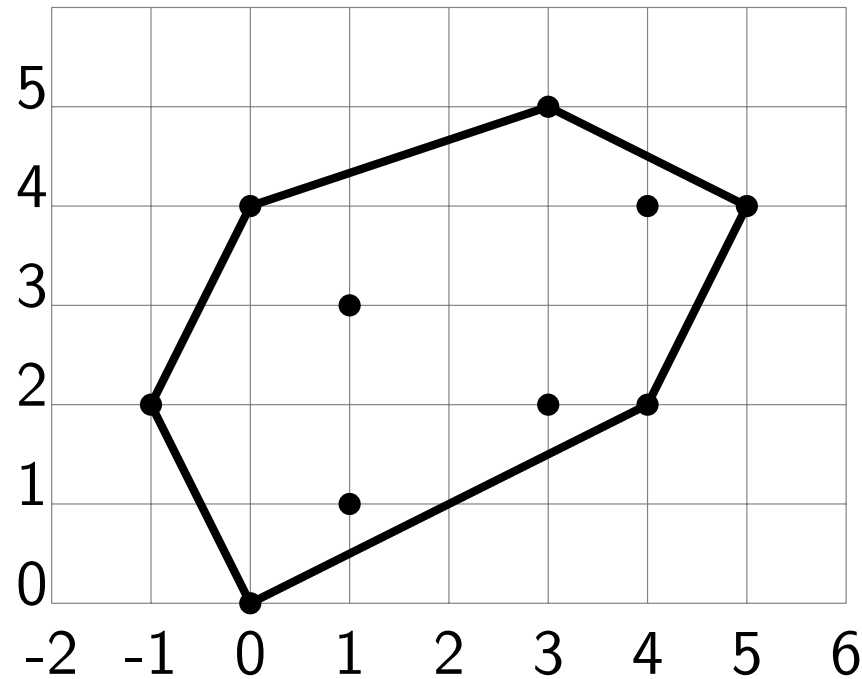


- With at most one point per dashed box, we can do as follows.
- Let S be sorted on y-coordinates
- Each point $p \in S$ is inspected at a time.
- The distances from p to each of the next six points on the other side in S (according to y-coordinates) are checked to see if it less than the shortest distance found so far

Algorithm outline

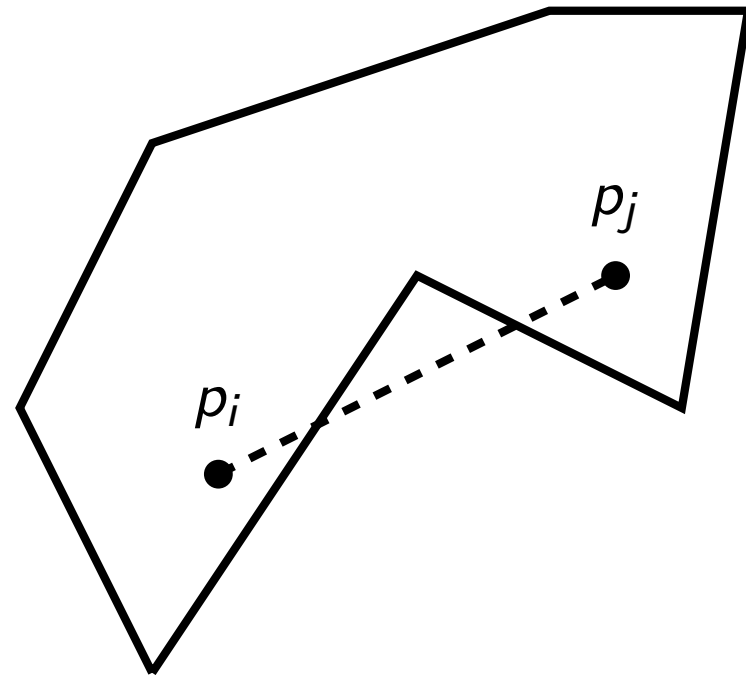
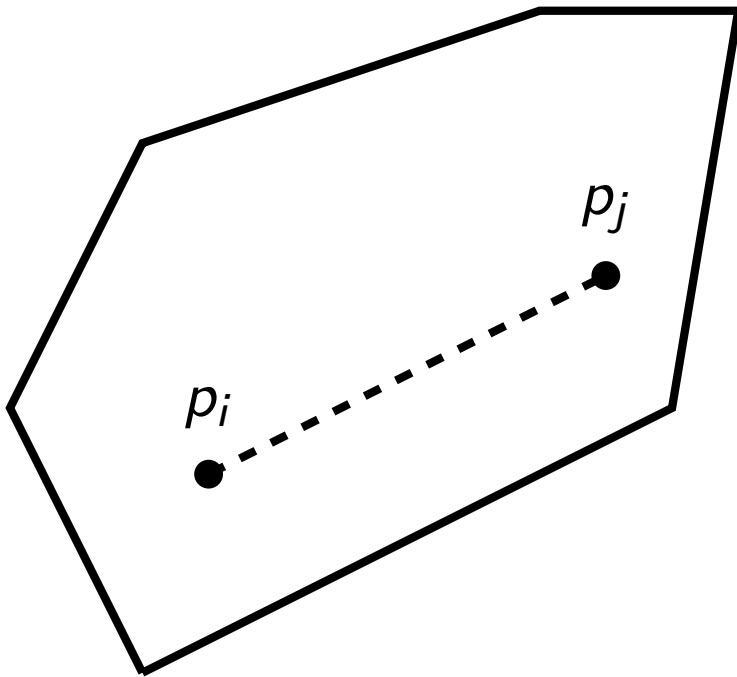
- What do we need for this?
- Input is a set of n points P
- We produce two sorted arrays P_x and P_y before starting our recursion
- We divide P_x into two arrays L_x and R_x (left and right)
- We divide P_y into two arrays L_y and R_y
- We solve the two subproblems $(L_x, L_y, n/2)$ and $(R_x, R_y, n/2)$
- Then we compute δ as the minimum from these subproblems
- Then we create the set S_y from P_y
- All dividing and combining can be done in linear time, so we solve this in $\Theta(n \log n)$ time

A set of points P and its convex hull $\text{CH}(P)$

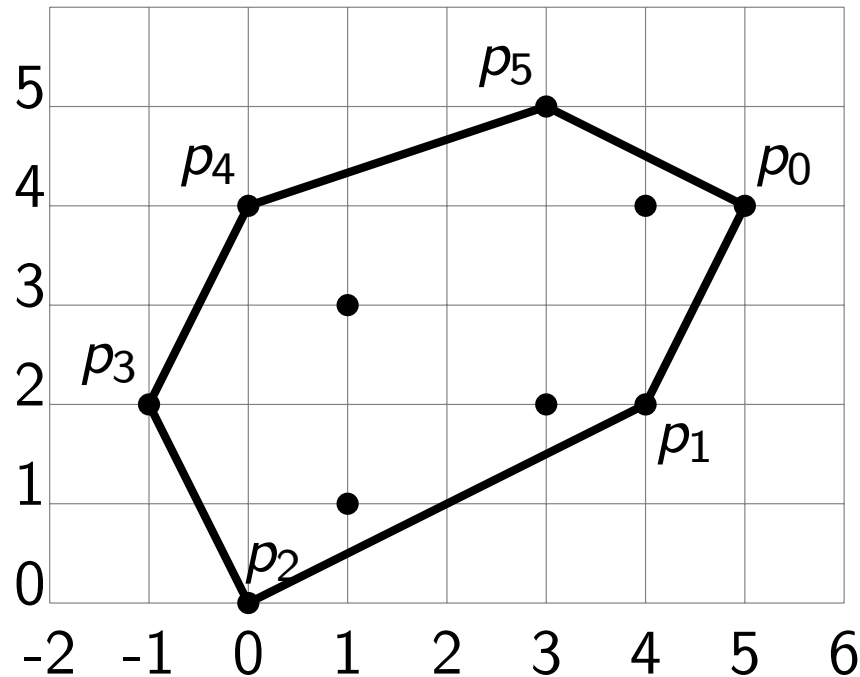


- The convex hull should have a minimal number of points.
- For example a point at $(2, 1)$ would not be in the convex hull.

A convex and a non-convex region



Clockwise order of CH(P)

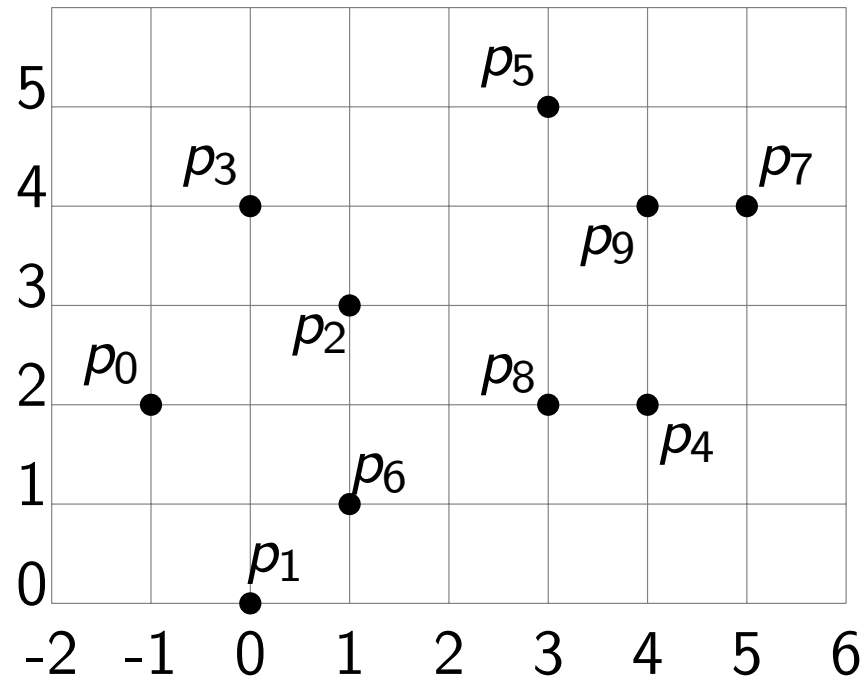


- Not necessary to select the rightmost point as p_0 in general
- In Lab 4 (2023) we did that however, since that is what the divide-and-conquer algorithm does.

Three algorithms for computing the convex hull

- Jarvis march, or gift wrapping
- Graham scan
- Preparata-Hong
- Always a good approach to implement a non-trivial algorithm: start with something simpler and use the simple (and hopefully correct) as a reference

Jarvis march

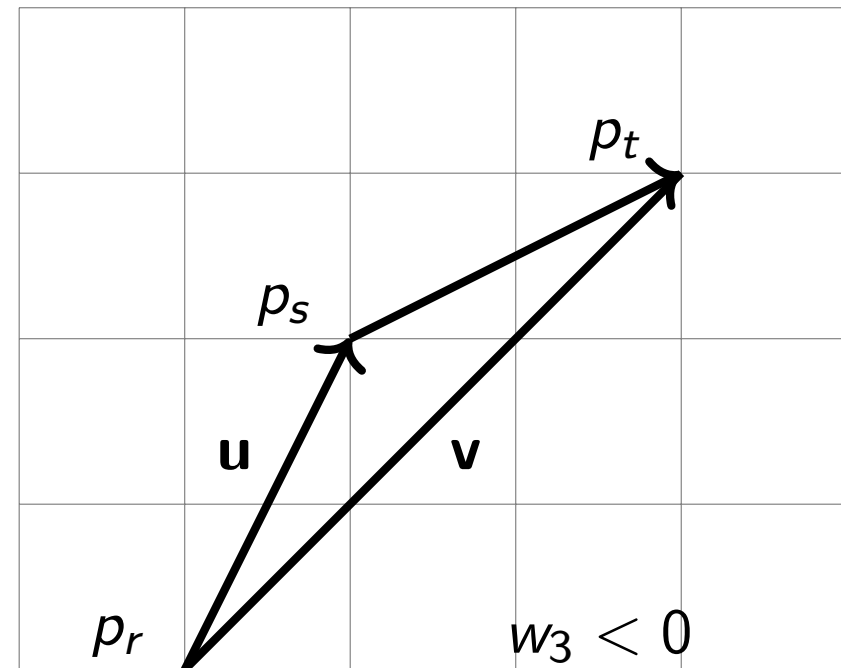
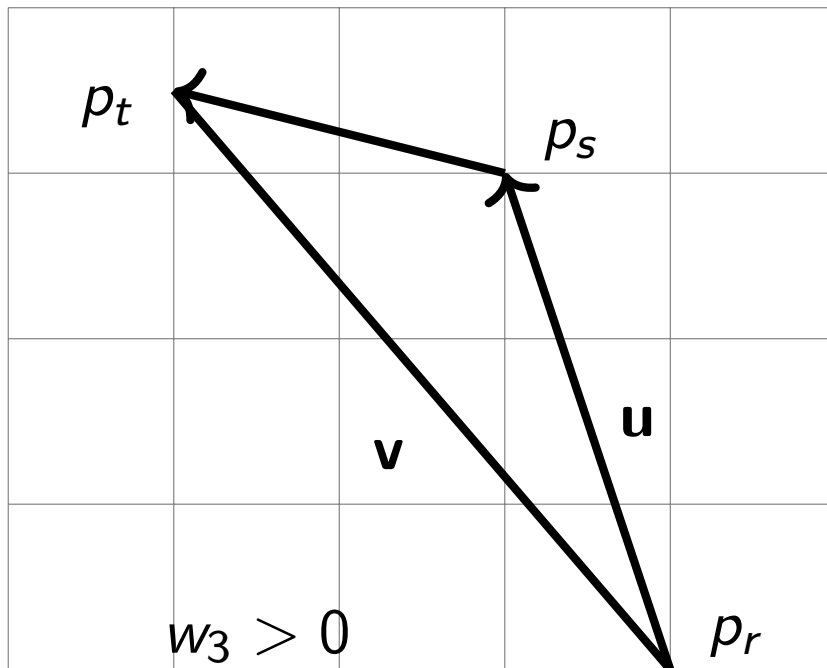


- Start at any point known to be in CH such as the leftmost: p_0 , $i \leftarrow 0$
- Select the next point p_j as the one through which we only make left turns when going from p_i through p_j to p_k for every other point p_k
- Right turn with p_0 , p_2 , p_1 so don't take p_2 next
- Always left turn with p_0 , p_1 , p_k so take p_1 as next
- Continue until back at p_0

Review of vector products

- Also called cross product (vektorprodukt or kryssprodukt)
- Given two vectors in \mathbb{R}^3 , \mathbf{u} and \mathbf{v} , the vector product, $\mathbf{w} = \mathbf{u} \times \mathbf{v}$ is a vector with the following properties:
 - 1 \mathbf{w} is perpendicular to both \mathbf{u} and \mathbf{v} , i.e., the dot products $\mathbf{w} \cdot \mathbf{u}$ and $\mathbf{w} \cdot \mathbf{v}$ are both zero,
 - 2 $|\mathbf{w}| = |\mathbf{u}||\mathbf{v}| \sin \theta$ where θ is the angle between \mathbf{u} and \mathbf{v} ,
 - 3 \mathbf{u} , \mathbf{v} and \mathbf{w} are positively oriented, i.e. according to the right-hand rule.
- $\mathbf{u} = u_1\mathbf{e}_1 + u_2\mathbf{e}_2 + u_3\mathbf{e}_3$ and $\mathbf{v} = v_1\mathbf{e}_1 + v_2\mathbf{e}_2 + v_3\mathbf{e}_3$
- $\mathbf{w} = (u_1\mathbf{e}_1 + u_2\mathbf{e}_2 + u_3\mathbf{e}_3) \times (v_1\mathbf{e}_1 + v_2\mathbf{e}_2 + v_3\mathbf{e}_3) = (u_2v_3 - u_3v_2)\mathbf{e}_1 + (u_3v_1 - u_1v_3)\mathbf{e}_2 + (u_1v_2 - u_2v_1)\mathbf{e}_3 = w_1\mathbf{e}_1 + w_2\mathbf{e}_2 + w_3\mathbf{e}_3.$
- Since our points are in \mathbb{R}^2 , their \mathbf{e}_3 coordinates are zero and so w_1 and w_2 also become zero
- $\mathbf{w} = w_3\mathbf{e}_3 = (u_1v_2 - u_2v_1)\mathbf{e}_3.$

Left or right direction at p_s determined with $\mathbf{w} = \mathbf{u} \times \mathbf{v}$



- To find the direction from p_r through p_s to p_t we let
- $\mathbf{u} = \overrightarrow{p_r p_s}$,
- $\mathbf{v} = \overrightarrow{p_r p_t}$, and
- $\mathbf{w} = \mathbf{u} \times \mathbf{v}$. If $w_3 > 0$ it is a left turn, if $w_3 < 0$ it is a right turn, and otherwise the three points are on the same line.

Jarvis march

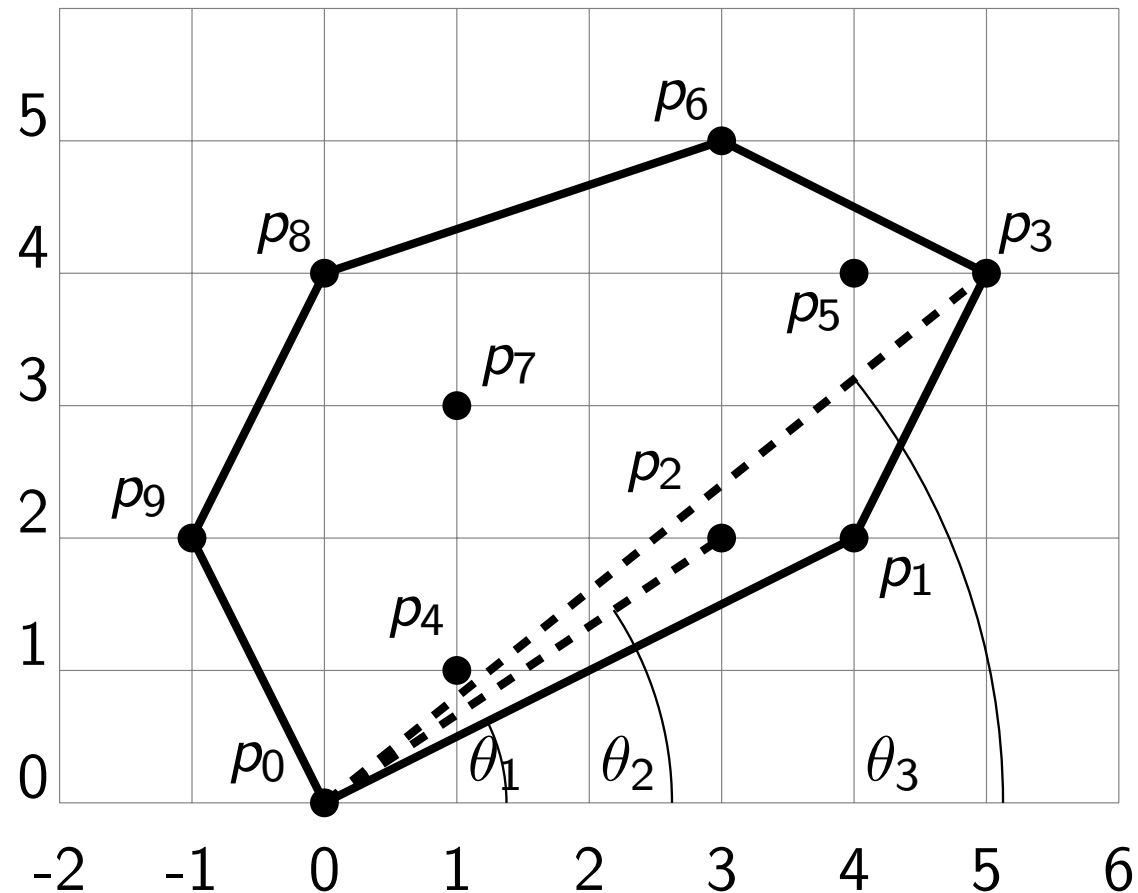
```
function jarvis_march( $p$ )
begin
   $n \leftarrow |p|$ 
   $i \leftarrow$  index of point in  $p$  with minimum  $x$  coordinate
  swap  $p_0$  and  $p_i$ 
   $r \leftarrow 0$ 
  while (1) {
     $s \leftarrow (r + 1) \bmod n$ 
    for  $t \leftarrow 0; t < n; t \leftarrow t + 1$  {
      if  $s = t$  then
        continue
       $\mathbf{u} = \overrightarrow{p_r p_s}$ 
       $\mathbf{v} = \overrightarrow{p_r p_t}$ 
       $\mathbf{w} = \mathbf{u} \times \mathbf{v}$ 
       $(w_1, w_2, w_3) \leftarrow \mathbf{w}$ 
      // right turn or  $p_s$  between  $p_r$  and  $p_t$  on a line?
      if  $w_3 < 0$  or  $w_3 = 0$  and  $|\mathbf{v}|^2 > |\mathbf{u}|^2$  then
         $s \leftarrow t$ 
    }
     $r \leftarrow r + 1$ 
    if  $s = 0$  then
      break
    swap  $p_s$  and  $p_r$ 
  }
  return  $r$  // number of points in  $CH(P)$ 
end
```


Time complexity of Jarvis march

- Time complexity is $O(n \cdot h)$ with h points in the convex hull.
- We increment r once for every point in the convex hull.
- Since some convex regions consist of all their points, the worst case is $O(n^2)$
- For example a regular polygon ("circle" but not exactly round...)
- Regelbunden polygon
- We will next look at Graham scan which is $O(n \log n)$ due to all points must be sorted first

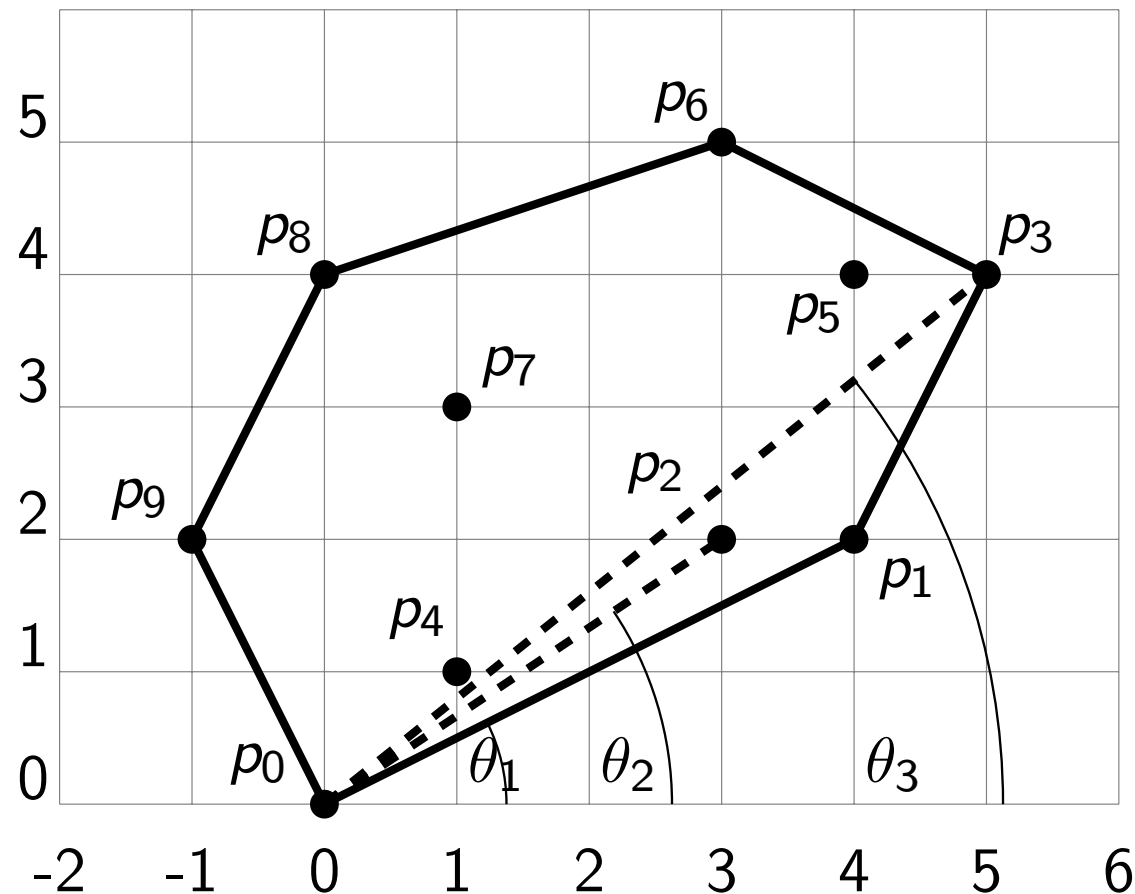
Graham scan

- First a point p_0 with minimal y -coordinate is made a new origo.
- One can make an angle between the x -axis, p_0 , and every other point
- The points are sorted by these angles θ_i , $1 \leq i \leq p_{n-1}$



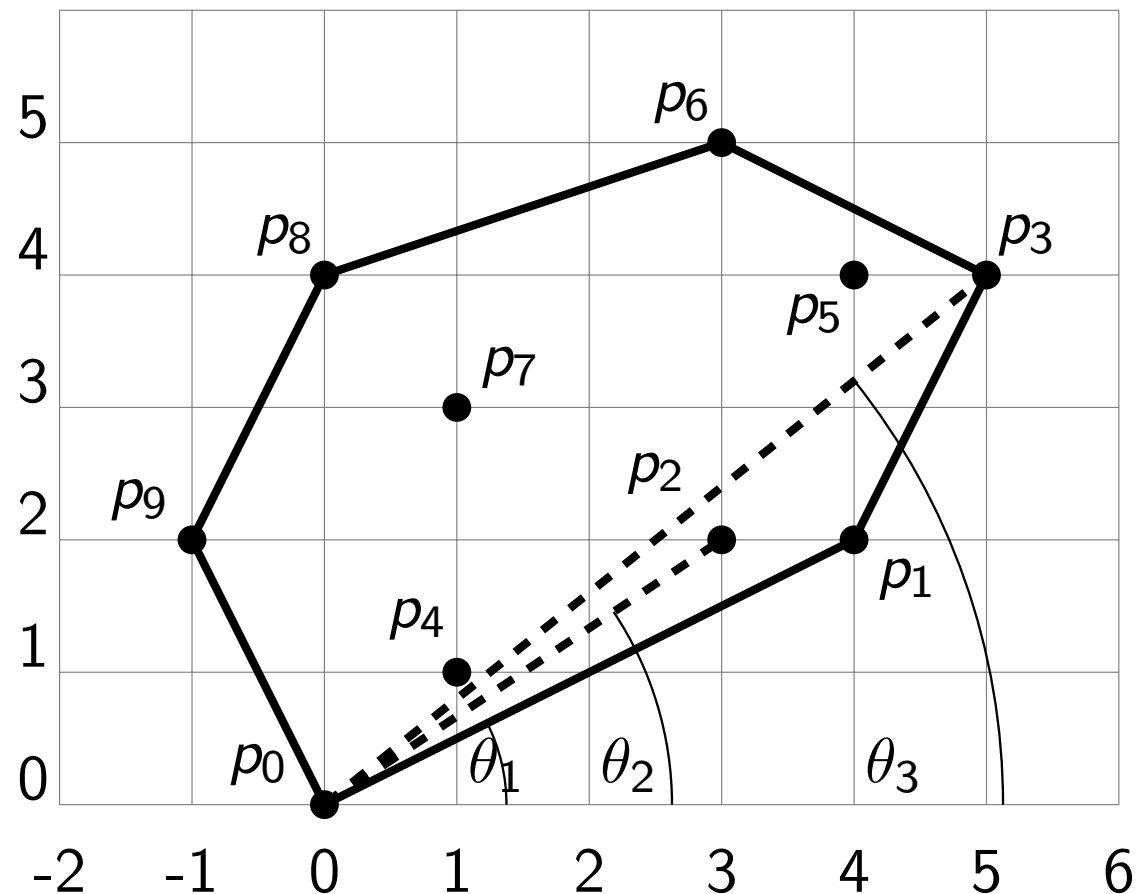
Including points in the convex hull

- The points p_0, p_1, p_2 are pushed to a stack with p_2 at the top
- Call the point at the top of the stack p_s (initially p_2)
- Call the point just below the top of the stack p_r (initially p_1)
- Call the "next point" p_t (initially p_3)



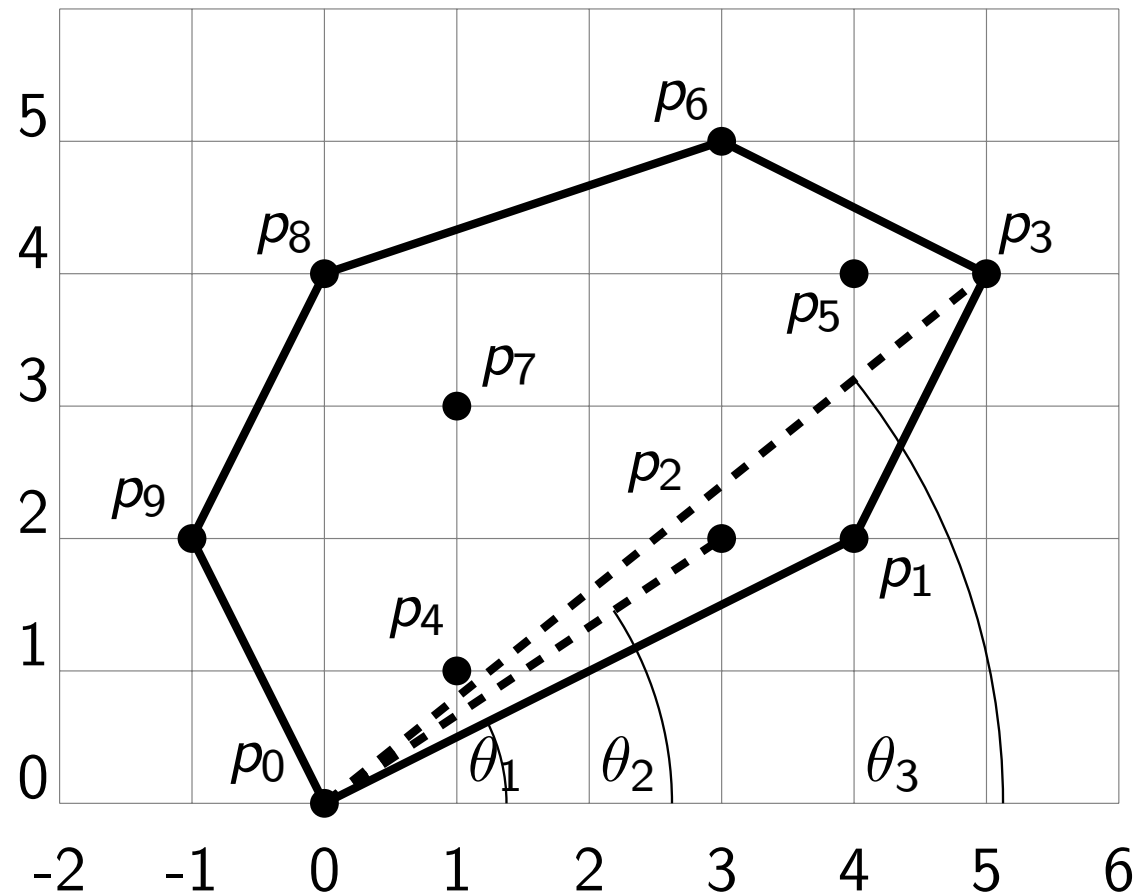
Excluding points from the convex hull

- Consider going from p_r , through p_s and to p_t initially p_1, p_2, p_3
- If the direction through p_s is straight or right, then p_s is not in CH
- In that case it is popped from the stack



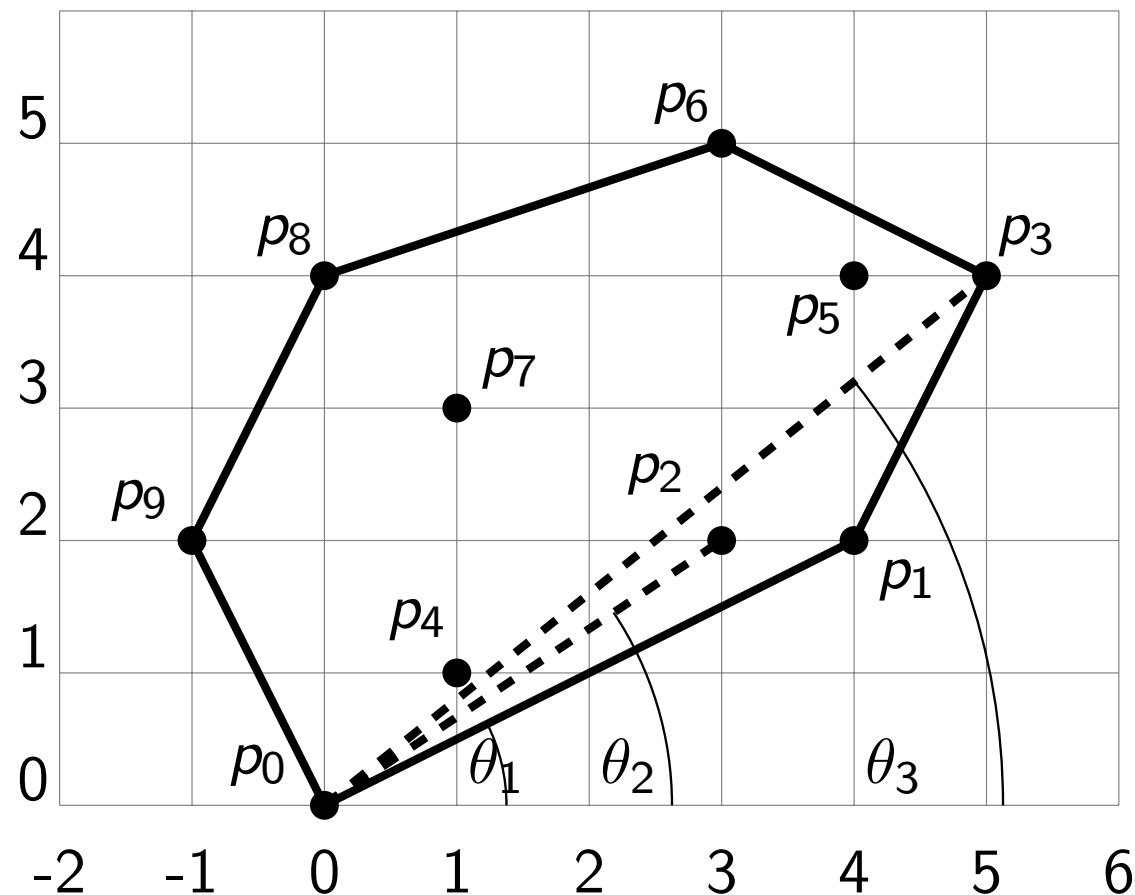
Excluding points from the convex hull

- Consider going from p_r , through p_s and to p_t
- After p_2 was popped, p_1 becomes new p_s and p_0 new p_r
- Any more non-left turns results in a pop
- Then p_t is pushed so $p_r = p_1$ and $p_s = p_3$



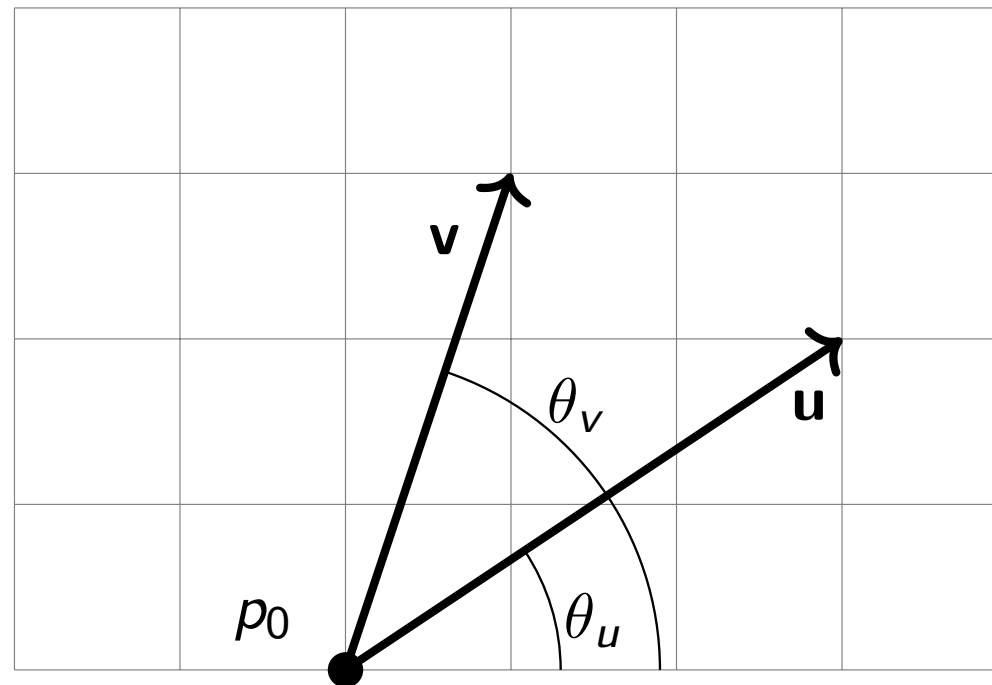
Excluding points from the convex hull

- $p_r = p_1$ and $p_s = p_3$
- $p_t = p_4$ with a left turn from p_r and p_s so p_4 is pushed
- After that p_5 will cause p_4 being popped
- In the end, all points remaining on the stack are the convex hull



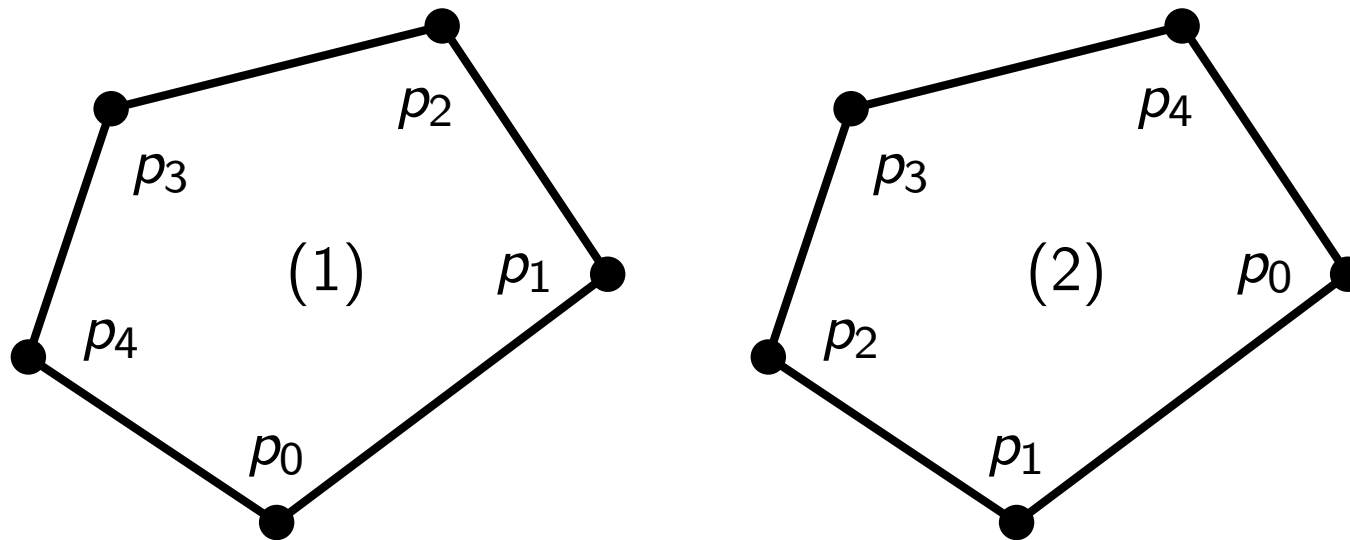
Relative sizes of θ is sufficient

- Compare angles with $\mathbf{u} \times \mathbf{v}$
- If $\theta_u = \theta_v$ then how should they be ordered?
- We want the point nearest origo on the stack first so the other can pop it



Output for Lab 4 2023

- The `check_solution.sh` script expects output as in (2)
- The reason is that Preparata-Hong produces that output.

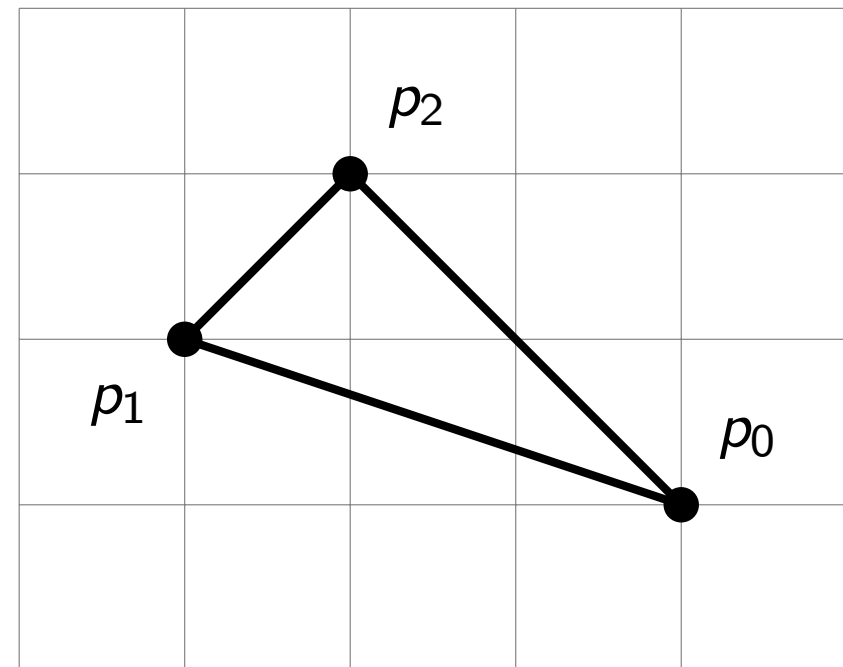
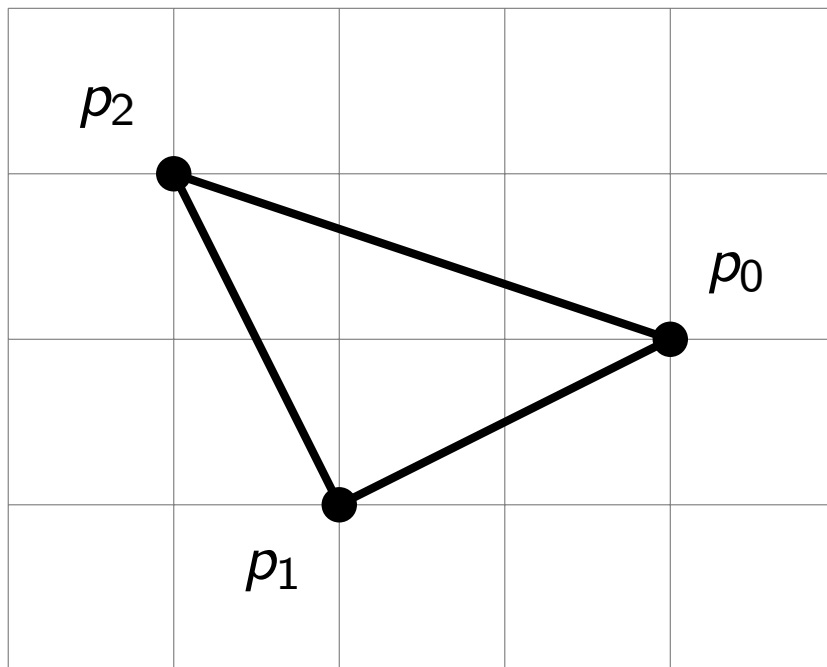


Graham scan

```
function graham_scan(p)  
begin  
  // p is an array with n points.  
   $n \leftarrow |p|$   
   $i \leftarrow$  index of point in p with minimum y coordinate  
  swap  $p_0$  and  $p_i$   
   $t = p_0$   
  subtract the coordinates of t from every point  
  sort elements  $1..n - 1$  of p by  $\theta_i$   
  h  $\leftarrow$  new stack  
  push(h,  $p_0$ )  
  push(h,  $p_1$ )  
  push(h,  $p_2$ )  
  for ( $k \leftarrow 3$ ;  $k < n$ ;  $k = k + 1$ ) {  
    //  $p_s$  is the top of h  
    //  $p_r$  is below  $p_s$  on h  
     $p_t \leftarrow p_k$   
    while direction(next_top(h), top(h),  $p_t$ ) is not left  
      pop(h)  
    push(h,  $p_t$ )  
  }  
  add the coordinates of t to every point  
   $n \leftarrow$  number of points on the stack  
  copy the points in h to p, and deallocate h  
  return n  
end
```

Preparata-Hong output

- The sequence that is the convex hull
- The number of points in the convex hull
- The index of the leftmost point



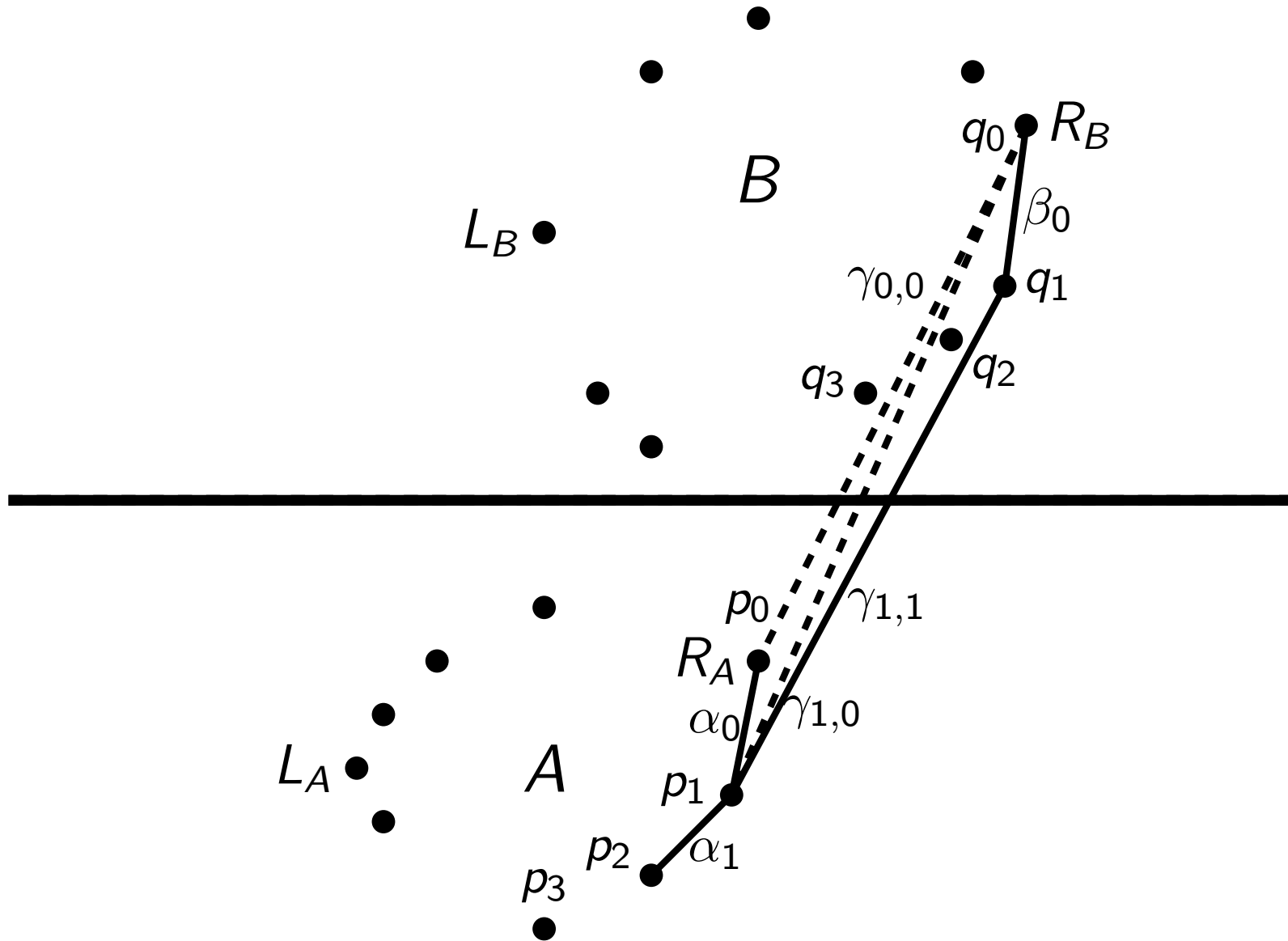
Preparata-Hong algorithm

- It relies on lines expressed as $y = k \cdot x + m$
- Sort all points n in order of increasing y -coordinates
- With $n \leq 3$ solve directly and return
- Divide in two approximately equal parts A and B
- All points in A must have a y -coordinate lower than any in B
- Find $CH(A)$ and $CH(B)$
- Merge $CH(A)$ and $CH(B)$ which is simplified by knowing that they are in clockwise order

- n_a points in lower convex hull
- n_b points in upper convex hull
- The lower points are called $p_0..p_{n_a-1}$
- The upper points are called $q_0..q_{n_b-1}$
- The inner points from A and B are not needed for anything
- $y = k \cdot x + m$
- We need to compute the k -values from p_0 to p_1 , from p_1 to p_2 etc
- The k value of the line segment from p_i to $p_{i+1 \bmod n_a}$ is called α_i
- The k value of the line segment from q_i to $q_{i+1 \bmod n_b}$ is called β_i
- For merging $CH(A)$ and $CH(B)$ we start with computing α and β

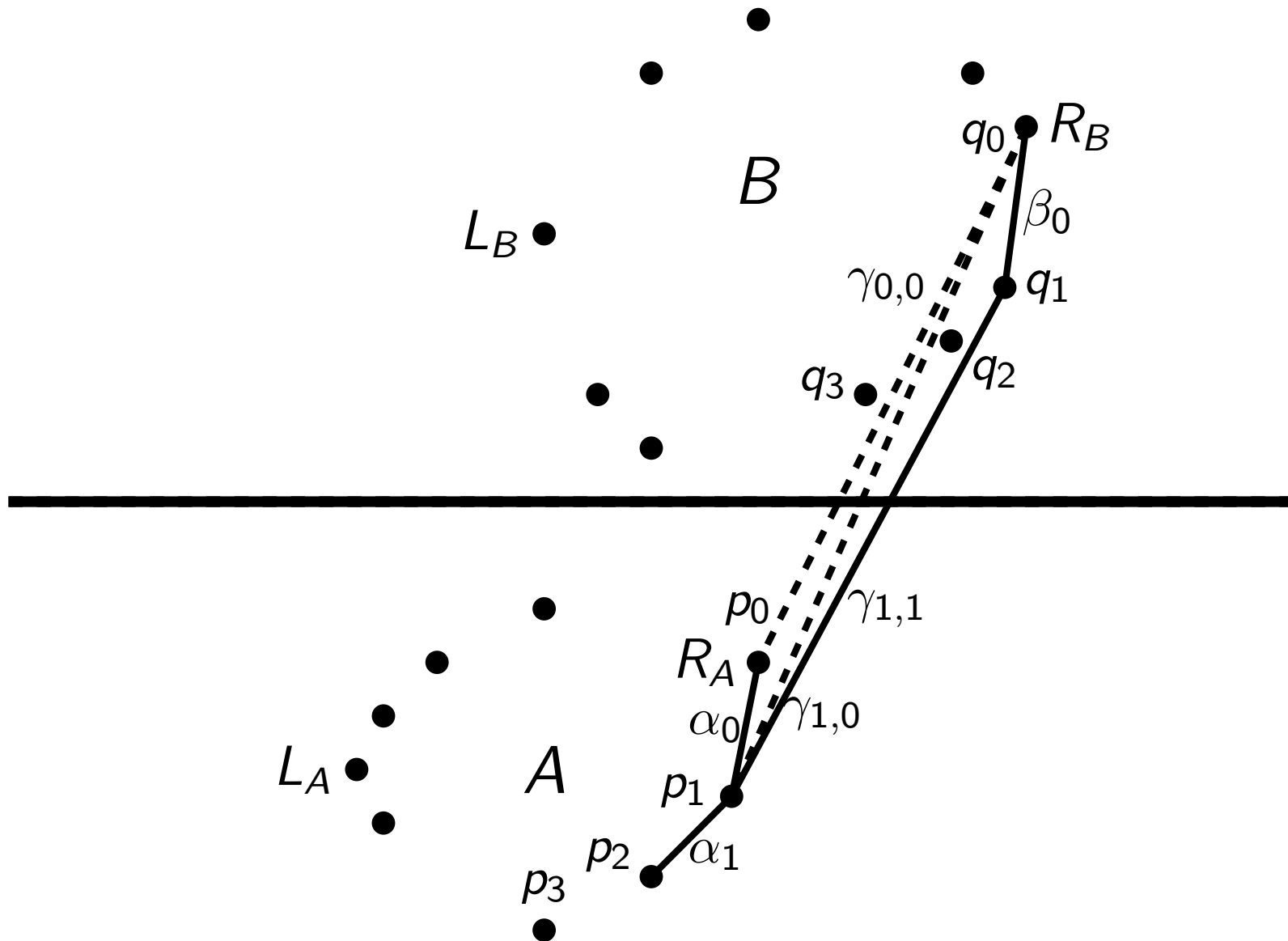
Selecting points on right side: q_0, q_1 and p_1, p_2, p_3, p_4

- We want to find first $i_R^* = p_1$ and last $j_R^* = q_1$ and skip q_2, \dots and p_0



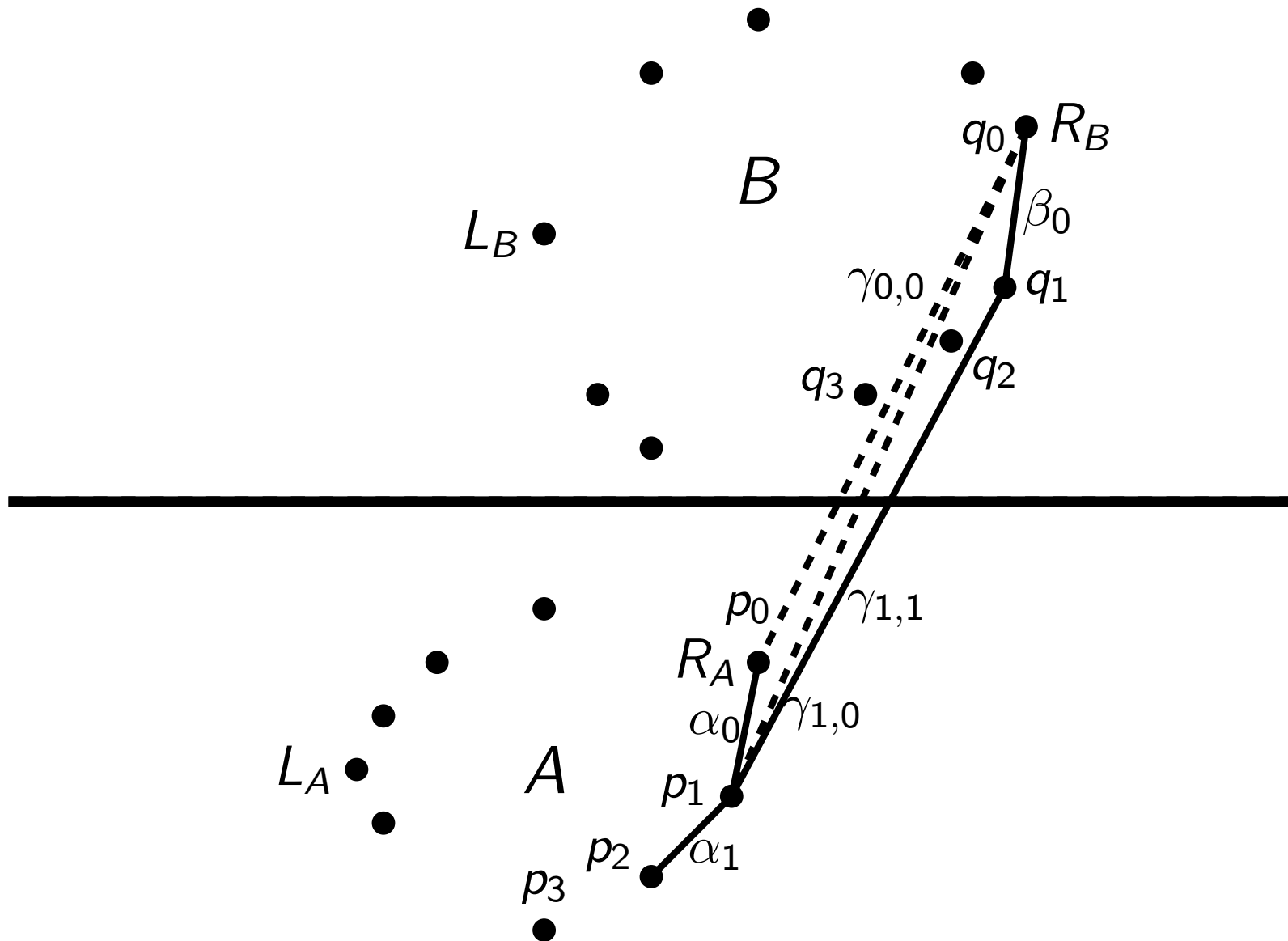
Exclude p_0

- First compare α_0 and $\gamma_{0,0}$. Since $\alpha_0 > \gamma_{0,0}$ we exclude p_0



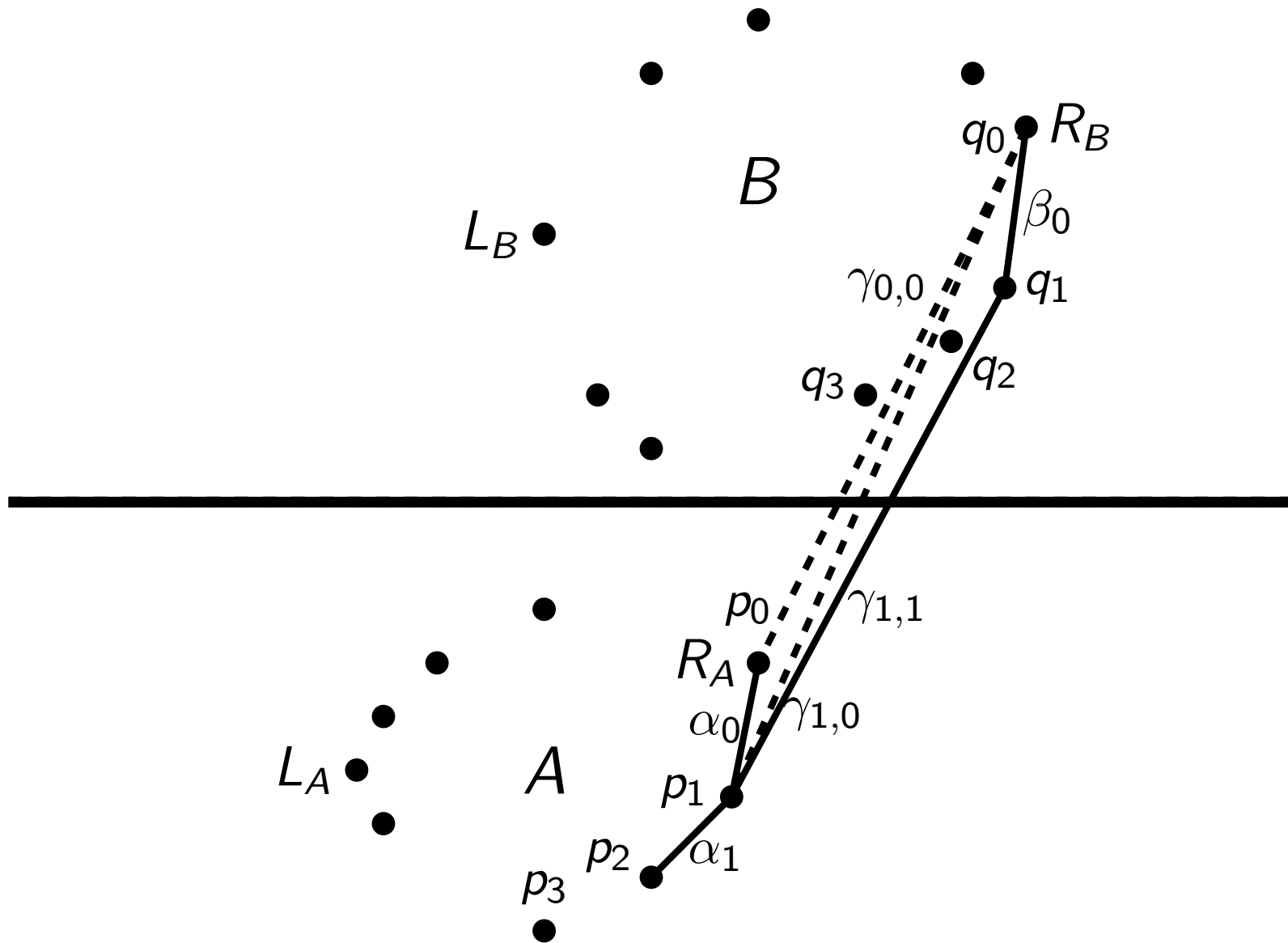
Include q_1

- Compare α_1 and $\gamma_{1,0}$. Since $\alpha_1 \leq \gamma_{1,0}$ we check β_0 and include q_1 .



We are done at right side

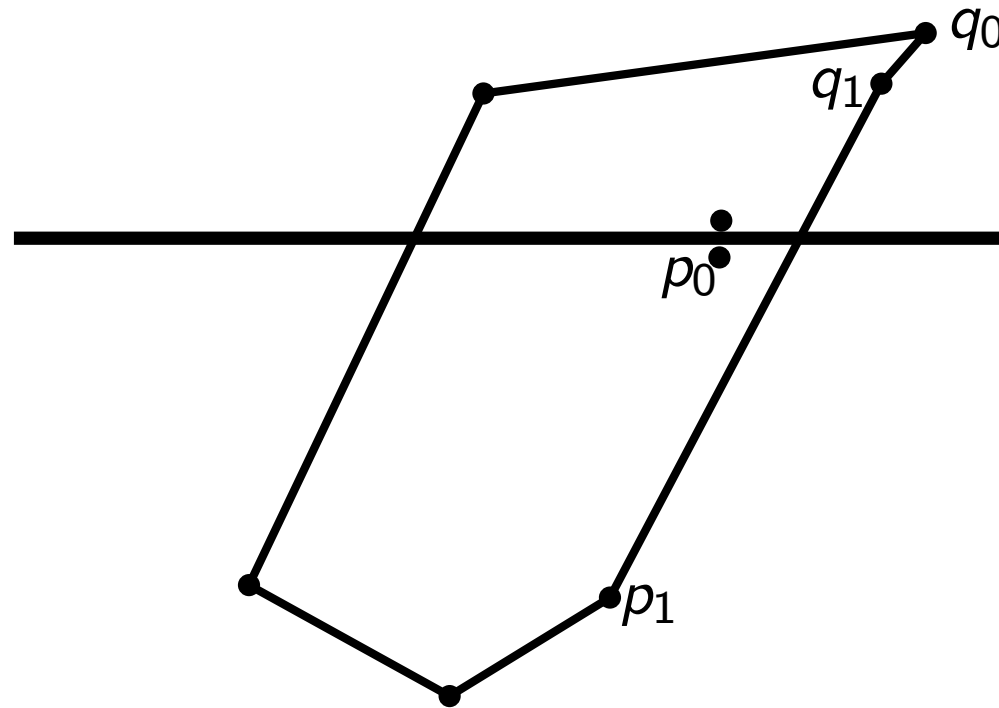
- Since both α_1 and β_1 are less than $\gamma_{1,1}$ we have found i_R^* and j_R^*



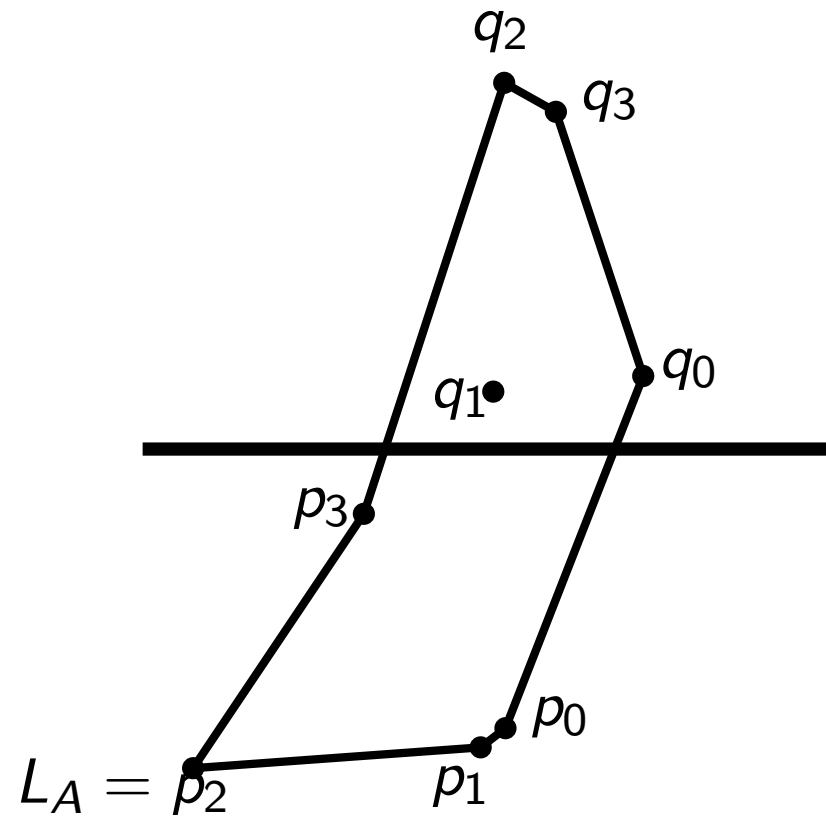
Next we do the same on the left side

- On the left side we want to find i_L^* and j_L^*
- Then we add lower points from i_R^* up to and including i_L^* to the output
- And add upper points from j_L^* up to and including j_R^* to the output
- Does it matter which of α_i and β_j we first compare with $\gamma_{i,j}$?

Wrong order on right side: comparing β_j before α_i

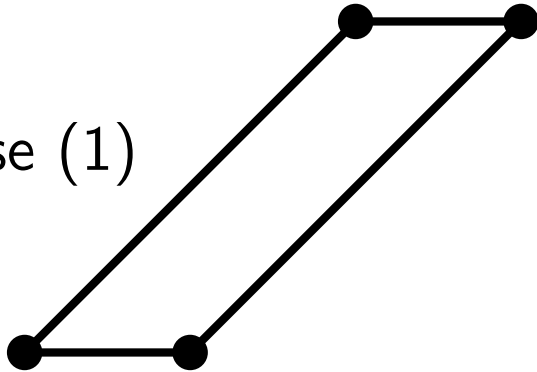


Wrong order on left side: comparing α_i before β_j



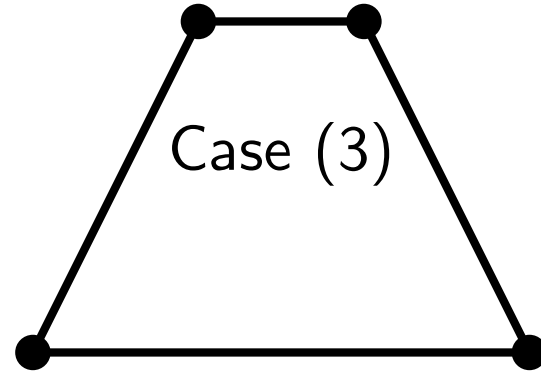
Four cases

Case (1)



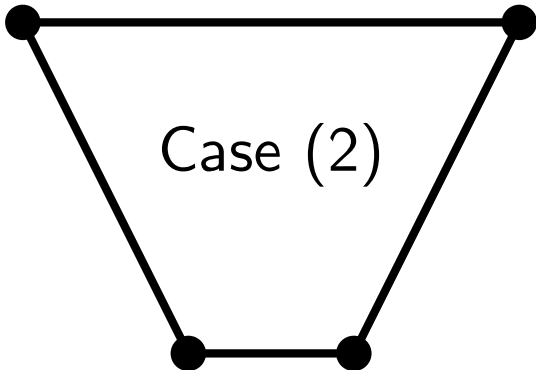
$$L_A < L_B \text{ and } R_A < R_B$$

Case (3)



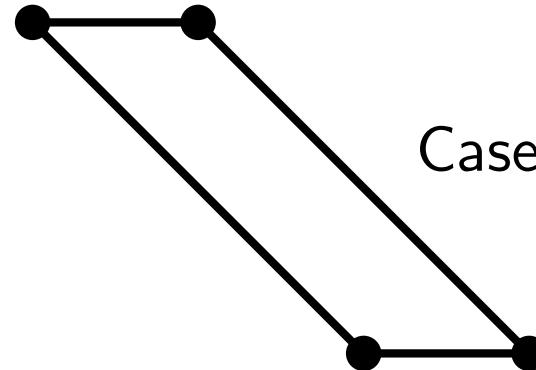
$$L_A < L_B \text{ and } R_A \geq R_B$$

Case (2)



$$L_A \geq L_B \text{ and } R_A < R_B$$

Case (4)



$$L_A \geq L_B \text{ and } R_A \geq R_B$$

Some hints

```
function  $k(p, q)$   
begin  
    return  $(q.y - p.y)/(q.x - p.x)$   
end
```

```
function  $compute\_k(p)$   
begin  
     $n \leftarrow |p|$   
     $\alpha \leftarrow \text{new double } [n]$   
    for  $i = 0; i < n; i \leftarrow i + 1$   
         $\alpha[i] \leftarrow k(p[i], p[i + 1 \bmod n])$   
    return  $\alpha$   
end
```

```
function  $add(k, from, to, q, p, n)$   
begin  
     $j \leftarrow from$   
    do {  
         $q[k] \leftarrow p[j]$   
         $k \leftarrow k + 1$   
         $i \leftarrow j$   
         $j \leftarrow (j + 1) \bmod n$   
    } while  $i \neq to$   
    return  $k$   
end
```

Avoid middle point when three points are on a line

```
function include_points(j, q, p, n)
begin
  for k  $\leftarrow$  0; k < n; k  $\leftarrow$  k + 1 {
    u  $\leftarrow$  (n + k + j - 1) mod n // point before v
    v  $\leftarrow$  (n + k + j - 0) mod n // point that may be excluded
    w  $\leftarrow$  (n + k + j + 1) mod n // point after v

    if not line_segment(q[v], q[u], q[w]) {
      p[i]  $\leftarrow$  q[v]
      i  $\leftarrow$  i + 1
    }
  }
  return i
end
```

Case 1

```
function case_1( $a, n_a, b, n_b, \alpha, \beta, i_L, j_L$ )
   $i \leftarrow 0$ 
   $j \leftarrow 0$ 
  while (1) {
     $\gamma_{i,j} \leftarrow k(a_i, b_j)$ 
    if ( $\alpha_i > \gamma_{i,j}$  or  $\alpha_i == -\infty$ ) and  $i < i_L$  then
       $i \leftarrow i + 1$ 
    else if ( $\beta_j > \gamma_{i,j}$  or  $\beta_j == -\infty$ ) and  $j < j_L$  then
       $j \leftarrow j + 1$ 
    else
      break
  }
   $i_R^* \leftarrow i$ 
   $j_R^* \leftarrow j$ 
   $i \leftarrow i_L$ 
   $j \leftarrow j_L$ 
  while (1) {
     $\gamma_{i,j} \leftarrow k(a_i, b_j)$ 
    if  $\beta_j > \gamma_{i,j}$  and  $j \neq 0$  then
       $j \leftarrow (j + 1) \bmod n_b$ 
    else if  $\alpha_i > \gamma_{i,j}$  and  $i \neq 0$  then
       $i \leftarrow (i + 1) \bmod n_a$ 
    else
      break
  }
   $i_L^* \leftarrow i$ 
   $j_L^* \leftarrow j$ 
  return ( $i_R^*, i_L^*, j_L^*, j_R^*$ )
```

Case 2

```
function case_2( $a, n_a, b, n_b, \alpha, \beta, i_L, j_L$ )  
   $i \leftarrow 0$   
   $j \leftarrow 0$   
  while (1) {  
     $\gamma_{i,j} \leftarrow k(a_i, b_j)$   
    if ( $\alpha_i > \gamma_{i,j}$  or  $\alpha_i == -\infty$ ) and  $i < i_L$  then  
       $i \leftarrow i + 1$   
    else if ( $\beta_j > \gamma_{i,j}$  or  $\beta_j == -\infty$ ) and  $j < j_L$  then  
       $j \leftarrow j + 1$   
    else  
      break  
  }  
   $i_R^* \leftarrow i$   
   $j_R^* \leftarrow j$   
   $i \leftarrow i_L$   
   $j \leftarrow j_L$   
  while (1) {  
     $\gamma_{i,j} \leftarrow k(a_i, b_j)$   
     $a_k \leftarrow (n_a + i - 1) \bmod n_a$   
     $b_k \leftarrow (n_b + j - 1) \bmod n_b$   
    if isfinite( $\alpha_{a_k}$ ) and  $\alpha_{a_k} < \gamma_{i,j}$  and  $i \neq 0$  then  
       $i \leftarrow a_k$   
    else if  $\beta_{b_k} < \gamma_{i,j}$  and  $j \neq 0$  then  
       $j \leftarrow b_k$   
    else  
      break  
  }  
   $i_L^* \leftarrow i$   
   $j_L^* \leftarrow j$   
  return ( $i_R^*, i_L^*, j_L^*, j_R^*$ )
```


Case 3

```
function case_3( $a, n_a, b, n_b, \alpha, \beta, i_L, j_L$ )  
   $i \leftarrow 0$   
   $j \leftarrow 0$   
  while (1) {  
     $\gamma_{i,j} \leftarrow k(a_i, b_j)$   
     $a_k \leftarrow (n_a + i - 1) \bmod n_a$   
     $b_k \leftarrow (n_b + j - 1) \bmod n_b$   
    if  $\beta_{b_k} < \gamma_{i,j}$  and  $j < j_L$  then  
       $j \leftarrow b_k$   
    else if  $\alpha_{a_k} < \gamma_{i,j}$  and  $i \neq i_L$  then  
       $i \leftarrow a_k$   
    else  
      break  
  }  
   $i_R^* \leftarrow i$   
   $j_R^* \leftarrow j$   
   $i \leftarrow i_L$   
   $j \leftarrow j_L$   
  while (1) {  
     $\gamma_{i,j} \leftarrow k(a_i, b_j)$   
    if  $\beta_j > \gamma_{i,j}$  and  $j \neq 0$  then  
       $j \leftarrow (j + 1) \bmod n_b$   
    else if  $\alpha_i > \gamma_{i,j}$  and  $i \neq 0$  then  
       $i \leftarrow (i + 1) \bmod n_a$   
    else  
      break  
  }  
   $i_L^* \leftarrow i$   
   $j_L^* \leftarrow j$   
  return ( $i_R^*, i_L^*, j_L^*, j_R^*$ )
```

Case 4

```
function case_4( $a, n_a, b, n_b, \alpha, \beta, i_L, j_L$ )  
   $i \leftarrow 0$   
   $j \leftarrow 0$   
  while (1) {  
     $\gamma_{i,j} \leftarrow k(a_i, b_j)$   
     $a_k \leftarrow (n_a + i - 1) \bmod n_a$   
     $b_k \leftarrow (n_b + j - 1) \bmod n_b$   
    if  $\beta_{b_k} < \gamma_{i,j}$  and  $j \neq j_L$  then  
       $j \leftarrow b_k$   
    else if  $\alpha_{a_k} < \gamma_{i,j}$  and  $i \neq i_L$  then  
       $i \leftarrow a_k$   
    else  
      break  
  }  
   $i_R^* \leftarrow i$   
   $j_R^* \leftarrow j$   
   $i \leftarrow i_L$   
   $j \leftarrow j_L$   
  while (1) {  
     $\gamma_{i,j} \leftarrow k(a_i, b_j)$   
     $a_k \leftarrow (n_a + i - 1) \bmod n_a$   
     $b_k \leftarrow (n_b + j - 1) \bmod n_b$   
    if isfinite( $\alpha_{a_k}$ ) and  $\alpha_{a_k} < \gamma_{i,j}$  and  $i \neq 0$  then  
       $i \leftarrow a_k$   
    else if isfinite( $\beta_{b_k}$ ) and  $\beta_{b_k} < \gamma_{i,j}$  and  $j \neq 0$  then  
       $j \leftarrow b_k$   
    else  
      break  
  }  
   $i_L^* \leftarrow i$   
   $j_L^* \leftarrow j$ 
```

More hints

- You don't need to split the code into four cases if you don't want to (simpler though, in my opinion, and likely faster)
- If you fail a test case it is a good idea to print the points in Matlab or to a pdf-file
- Implementing the Preparata-Hong algorithm is harder than Graham scan so why would you want to do it?
- An advantage of divide-and-conquer algorithms is that they are easier to parallelize
- Compute $CH(A)$ and $CH(B)$ by different threads at the same time
- If you have e.g. 80 hardware threads, you can easily let 64 work in parallel
- The machine `power.cs.lth.se` has just 80 hardware threads

Is p on a line segment between q and r in a plane?

- We want to know if p is between two points on a line in which case p should not be in the convex hull
- $\mathbf{u} = \overrightarrow{qp}$
- $\mathbf{v} = \overrightarrow{qr}$
- $\mathbf{w} = \mathbf{u} \times \mathbf{v}$.
- $\mathbf{w} = w_3 \mathbf{e}_3 = (u_1 v_2 - u_2 v_1) \mathbf{e}_3$ due to a plane.
- If $w_3 \neq 0$ then p is not on the line through q and r .
- Assume instead $w_3 = 0$ so the points are colinear.
- Is p between q and r ?
- Compute the dot products $\mathbf{v} \cdot \mathbf{v}$ and $\mathbf{u} \cdot \mathbf{v}$
- If $\mathbf{u} \cdot \mathbf{v} < 0$ then \mathbf{u} and \mathbf{v} have opposite directions so p is not between
- Otherwise if $\mathbf{u} \cdot \mathbf{v} > \mathbf{v} \cdot \mathbf{v}$ then p is also not between them