

Parametric sequence alignment

(pairwise) sequence alignment

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- a pair of sequences $S=\{s_1,s_2\}$ with lengths m and n , and
- an alignment objective function

find an $2 \times L$ matrix

- where $\max(m,n) < L < m+n$,
- each row represents one sequence from the set with inserted gaps, and
- is optimal under the objective function.



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$O(mn)$
running time



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When we look at it as a whole, we are defining the whole alignment score $f(\mathbb{A})$:

- with Needleman-Wunch it was

$$f_{\vec{\sigma}}(\mathbb{A}) = \sum_{a,b \in \Sigma \cup \{-\}} \sigma(a,b) \times \#(\mathbb{A}, a, b)$$

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$$f_{\alpha,\beta,\gamma}(\mathbb{A}) = \alpha \sum_{a \in \Sigma} \#(\mathbb{A}, a, a) + \beta \sum_{a \neq b \in \Sigma} \#(\mathbb{A}, a, b) + \gamma \left(\sum_{a \in \Sigma} \#(\mathbb{A}, a, '- ') + \sum_{b \in \Sigma} \#(\mathbb{A}, '- ', b) \right)$$

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$$f_{\alpha,\beta,\gamma}(\mathbb{A}) = \alpha \cdot \text{mt}_{\mathbb{A}} + \beta \cdot \text{ms}_{\mathbb{A}} + \gamma \cdot \text{id}_{\mathbb{A}}$$

- what about when we add affine gaps?

Alignment objective function

$$f_{\alpha,\beta,\gamma,\delta}(A) = \alpha \cdot \mathbf{mt}_A - \beta \cdot \mathbf{ms}_A - \gamma \cdot \mathbf{id}_A - \delta \cdot \mathbf{gp}_A$$

- \mathbf{mt}_A -- number of columns where both characters match
- \mathbf{ms}_A -- number of columns where their characters are different (mismatches)
- \mathbf{id}_A -- number of gap characters (indels)
- \mathbf{gp}_A -- number of gaps

An example

$s_1 = \text{AACCCG}$

$s_2 = \text{AAGGCC}$

A_1 **AA--CCCG**
AAGGCC--

	A_1
mt	4
ms	0
id	4
gp	2

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	A_1	A_2	A_3	A_4
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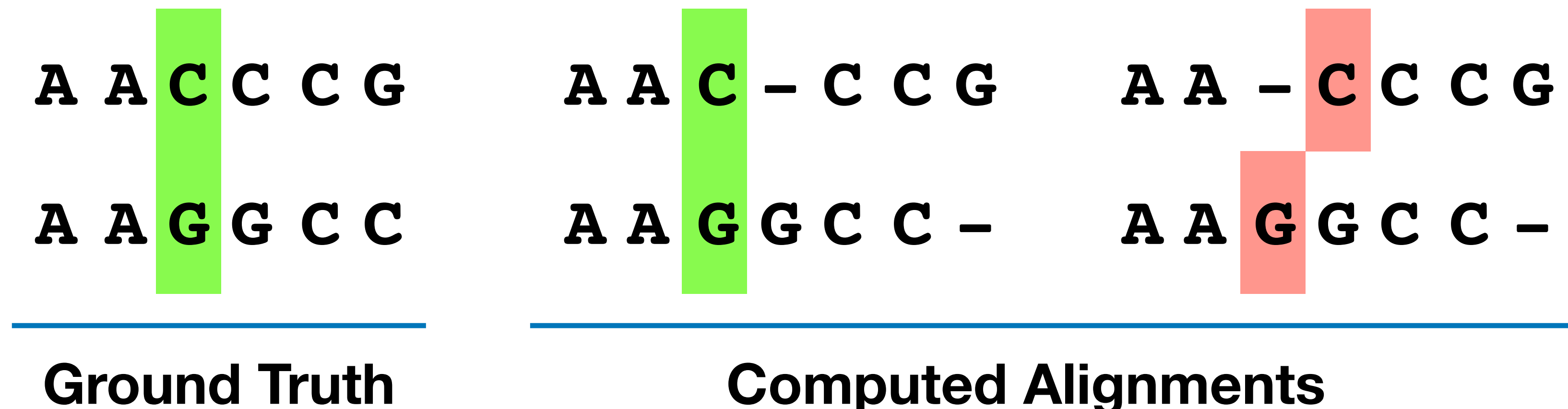
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What do we even mean by "best"?

A Digression on Accuracy

How would we know how accurate an alignment was if we knew the right answer?

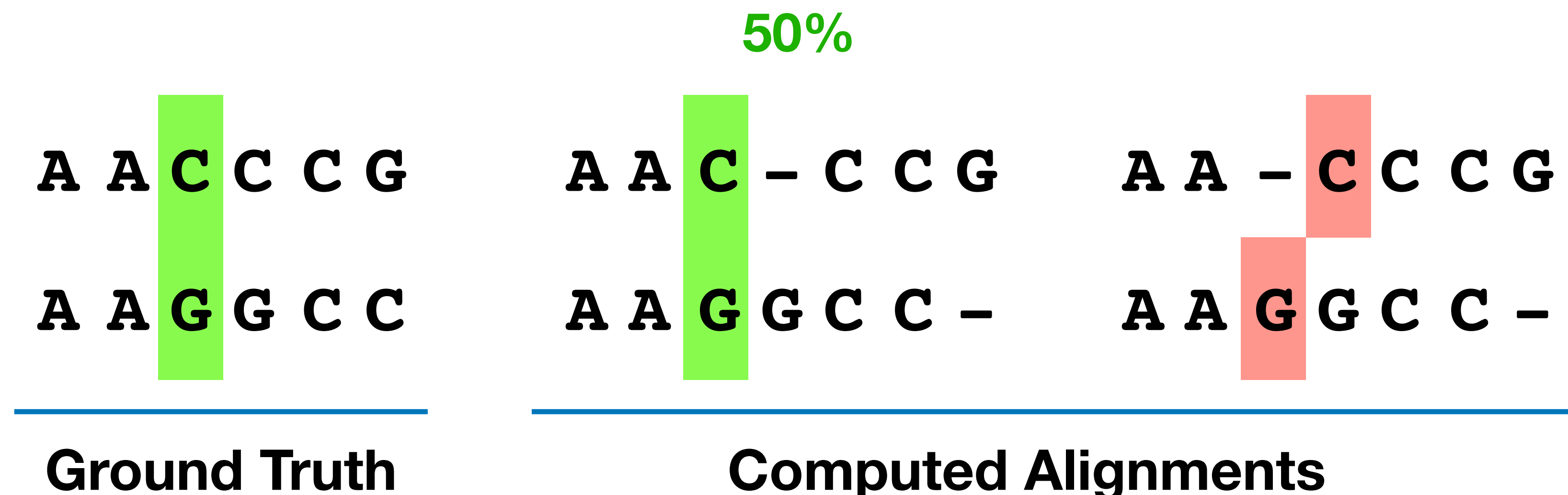
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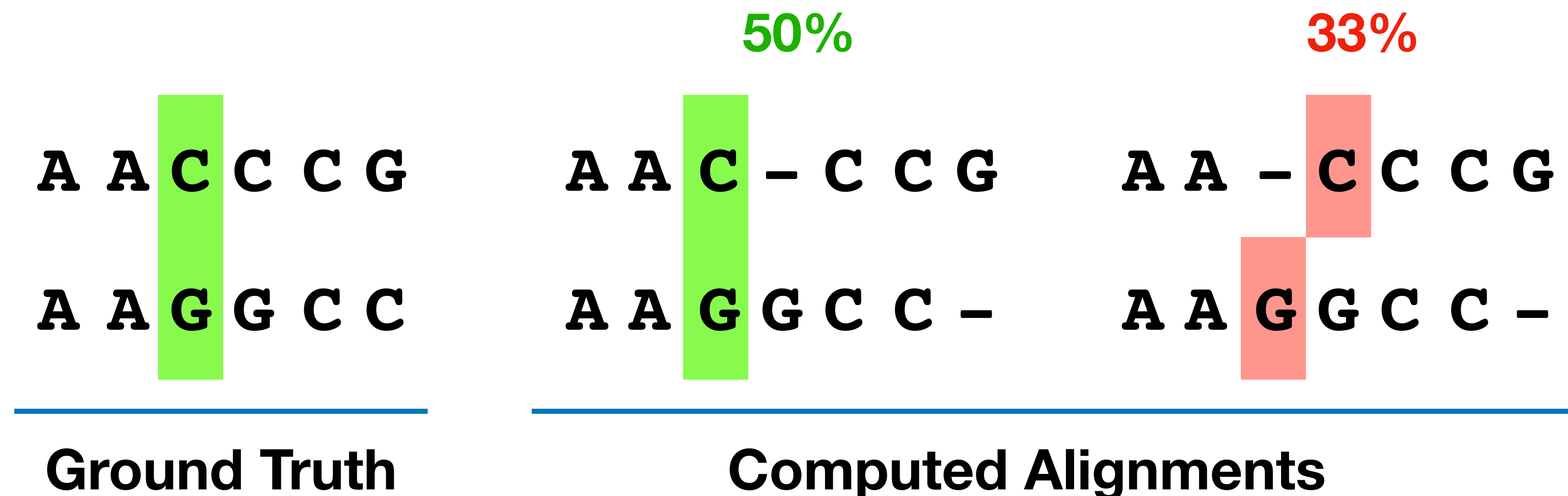
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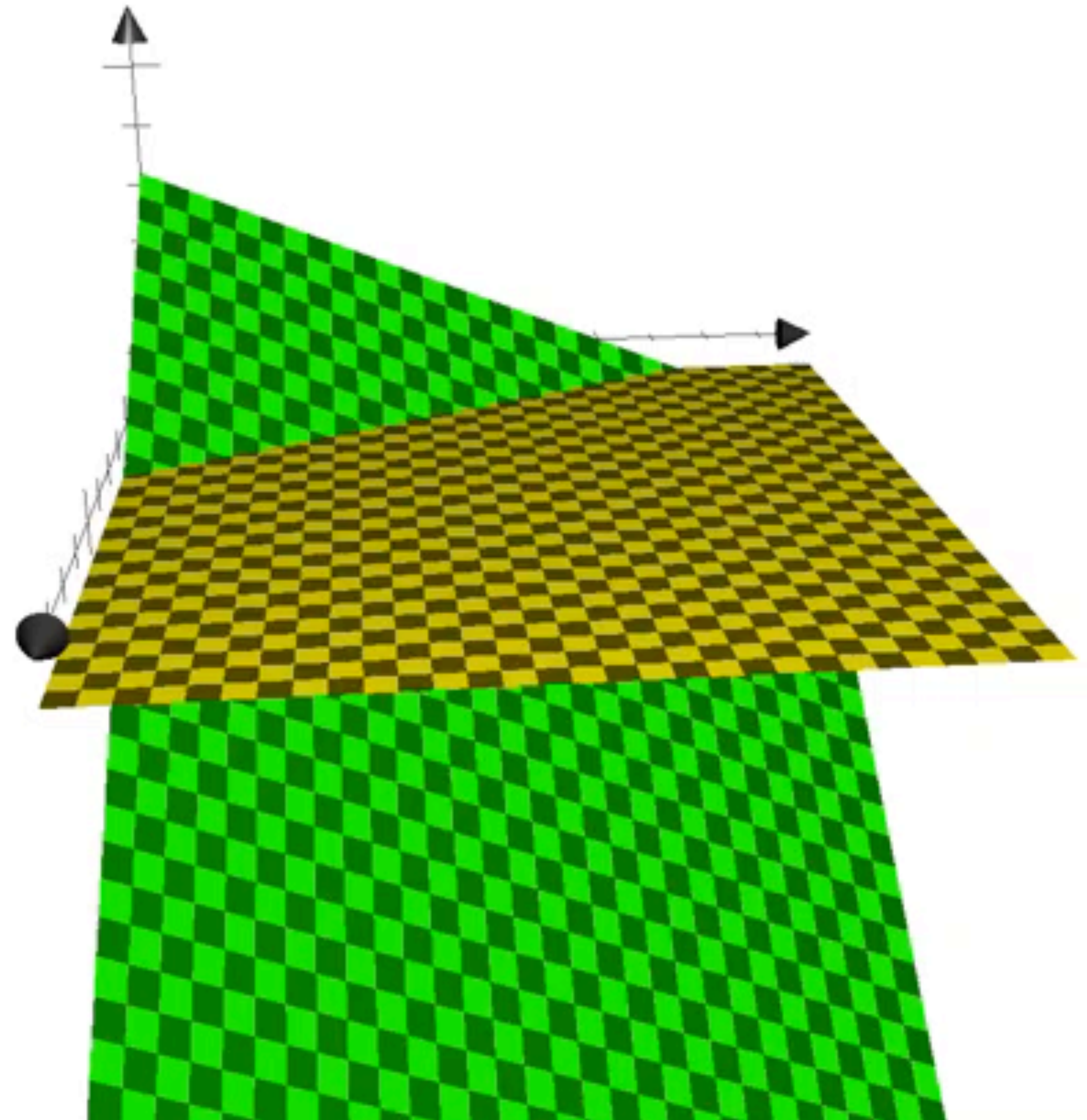
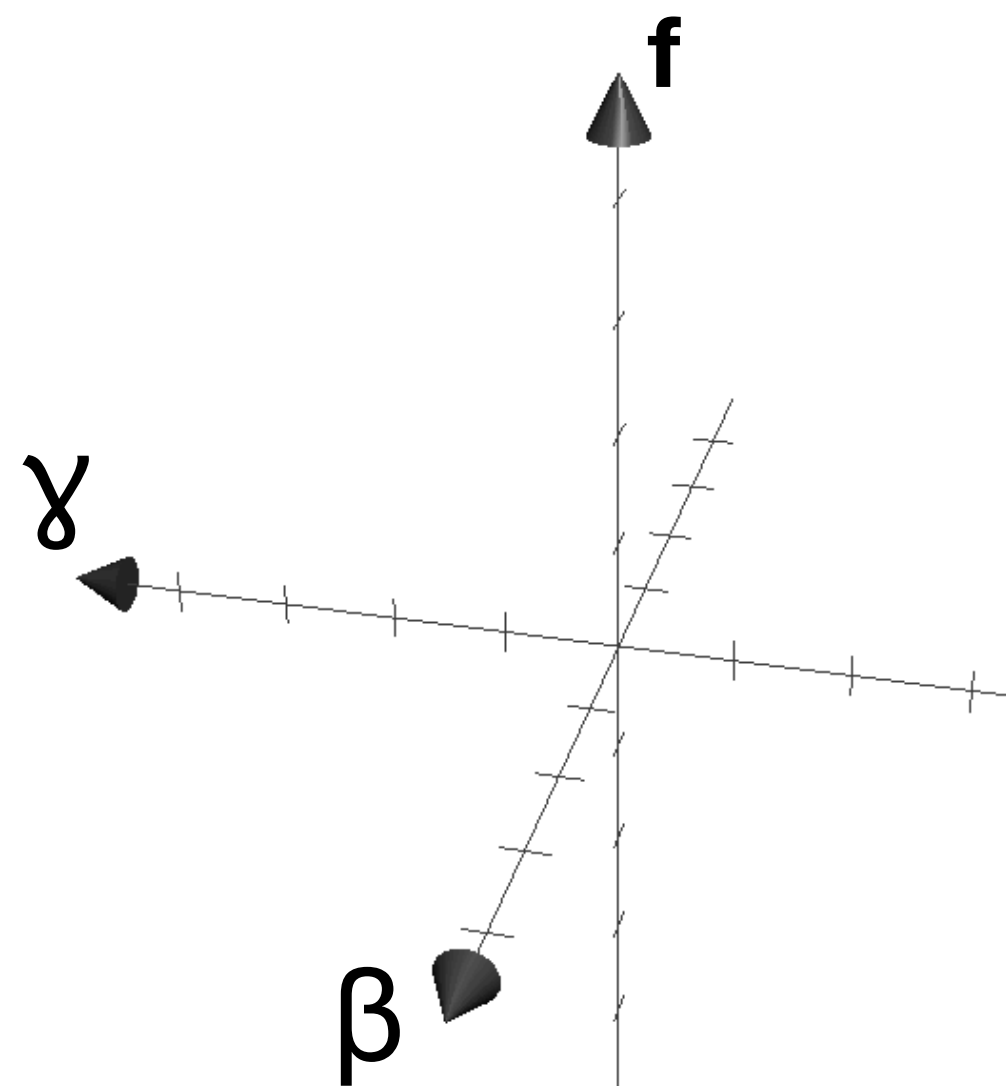
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Pairs of alignments in parameter space

Each alignment can be represented as a plane in the (γ, δ, f) -space.

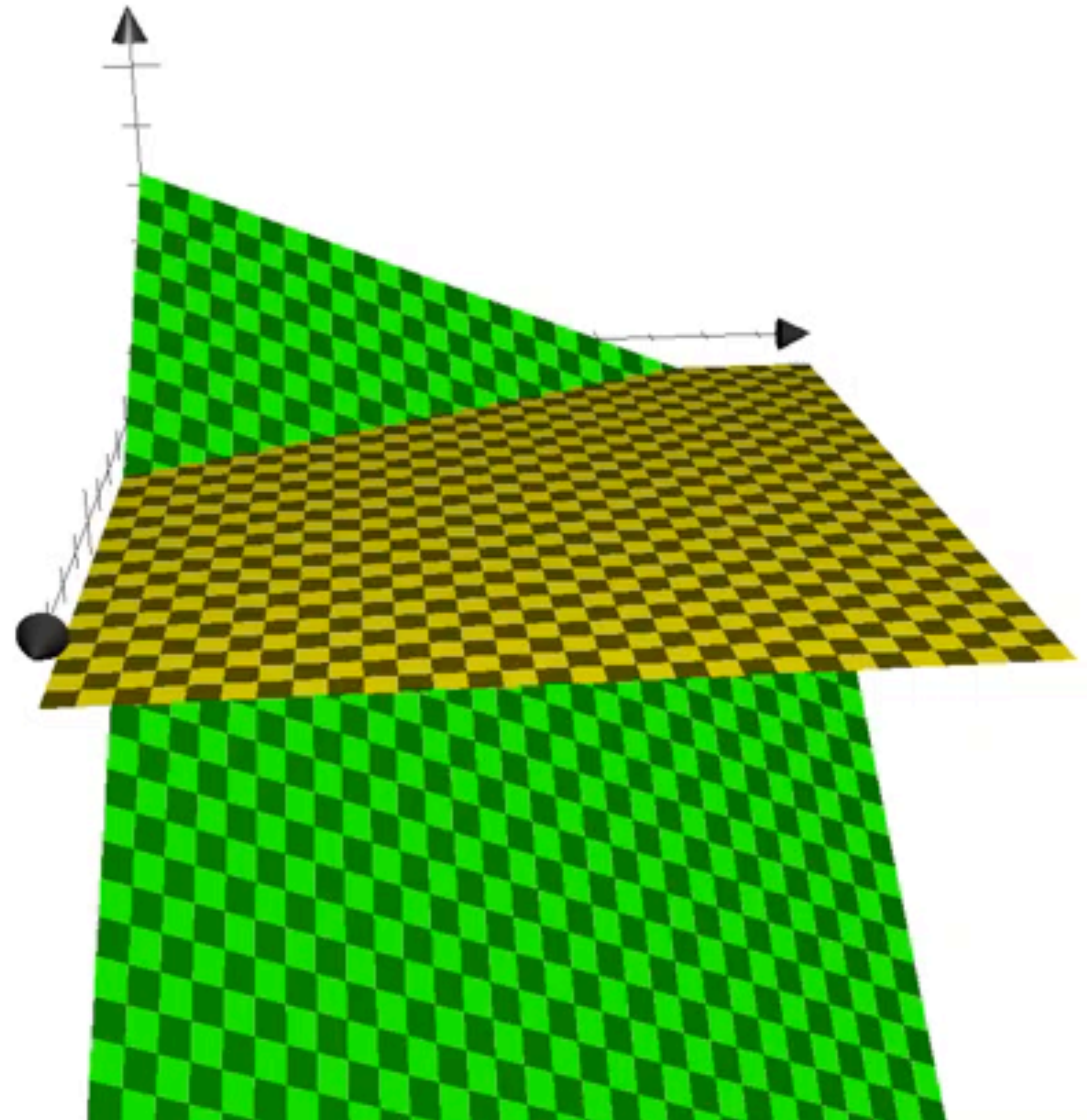
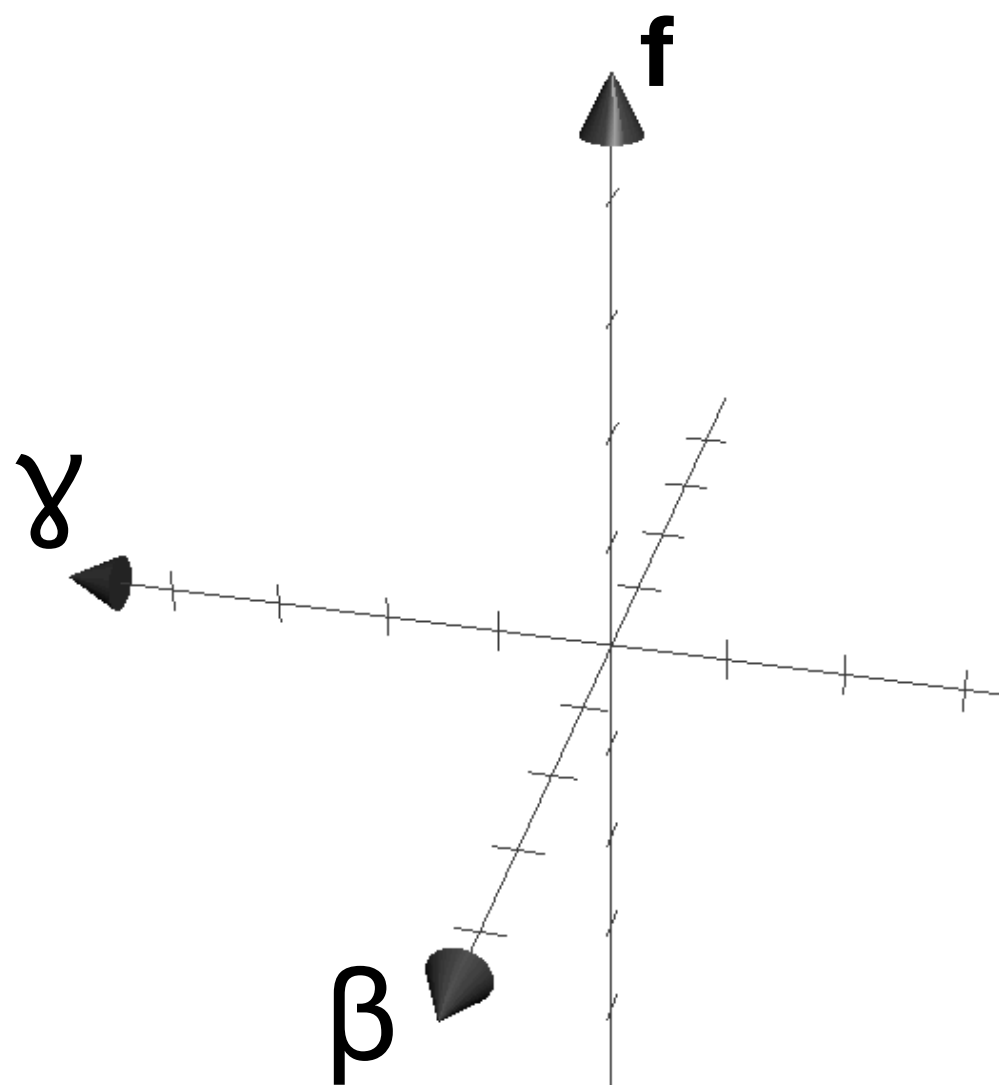
If the planes of alignments \mathbb{A} & \mathbb{A}' intersect, and are distinct, then there is a line L in (γ, δ, f) -space along which \mathbb{A} & \mathbb{A}' have the same objective value. If the planes don't intersect then one alignment had a larger objective value at all assignments of γ & δ .



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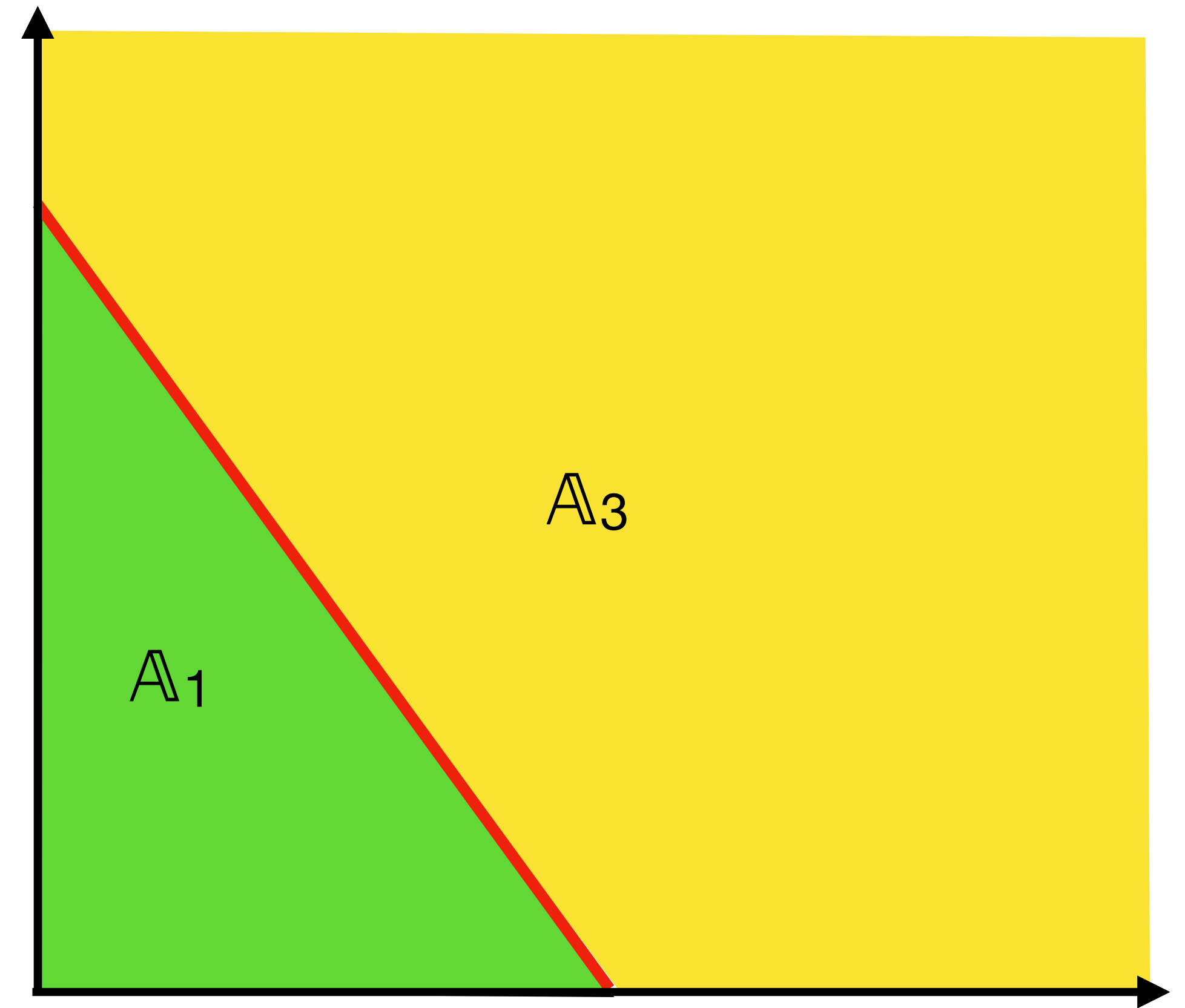


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If the planes of alignments \mathbb{A} & \mathbb{A}' intersect, and are distinct, then there is a line L in (γ, δ) -space along which \mathbb{A} & \mathbb{A}' have the same objective value; \mathbb{A} has a larger value on one half plane and \mathbb{A}' on the other. If the planes don't intersect then one alignment had a larger objective value at all assignments of γ & δ .

When projected to the (γ, δ) -plane, we can designate regions for which $f(\mathbb{A}) > f(\mathbb{A}')$ and vice versa

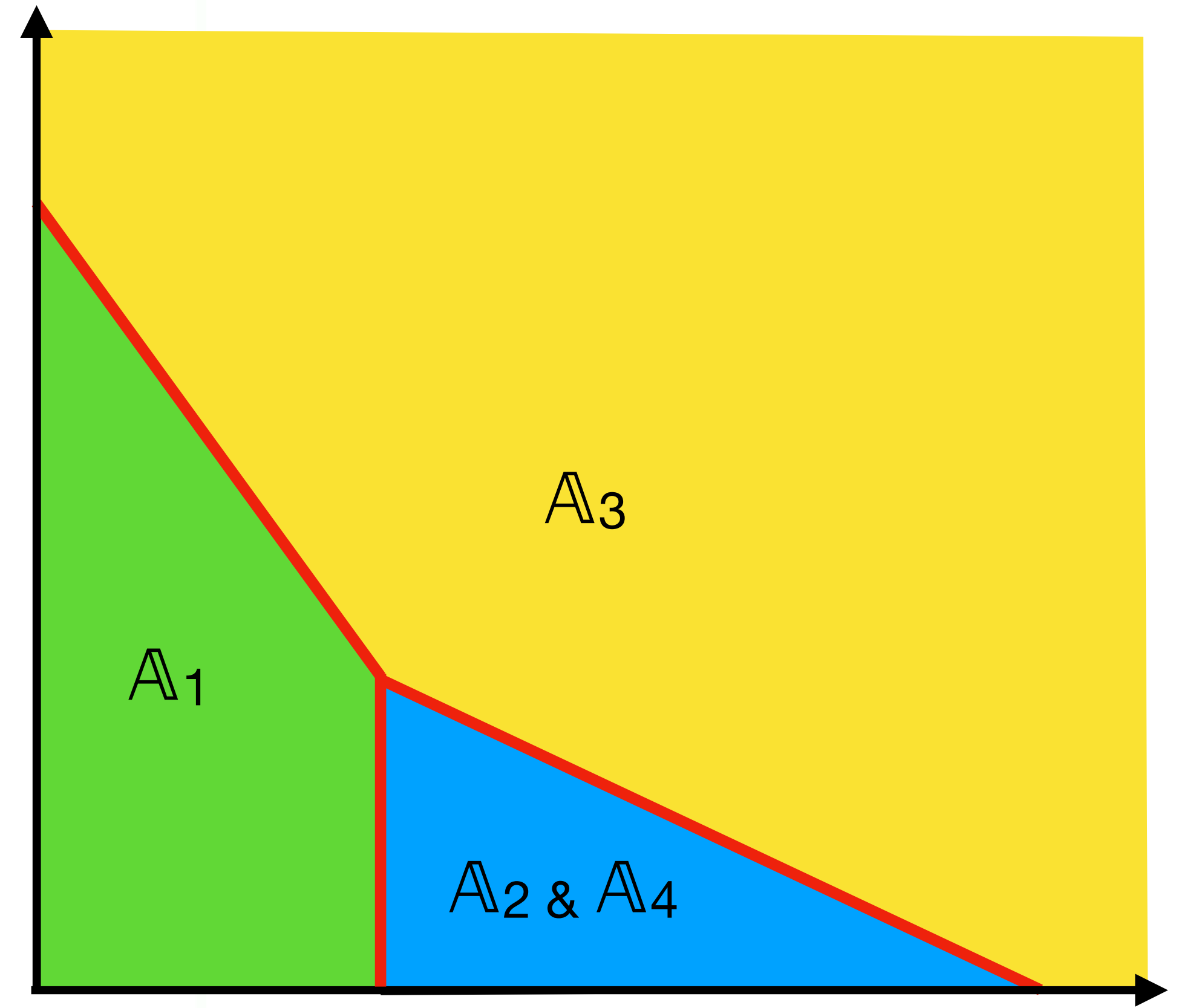
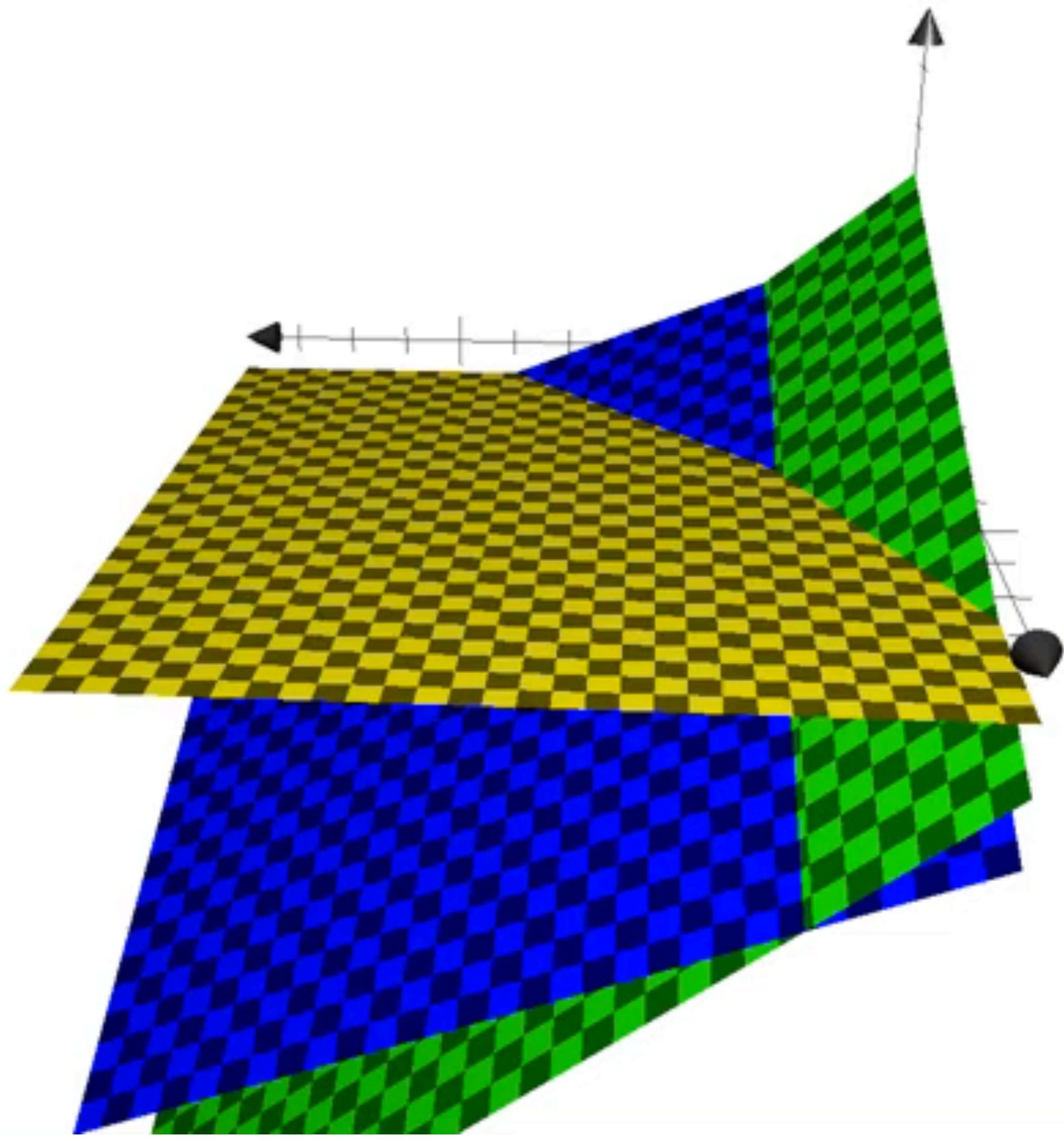


Things we know so far

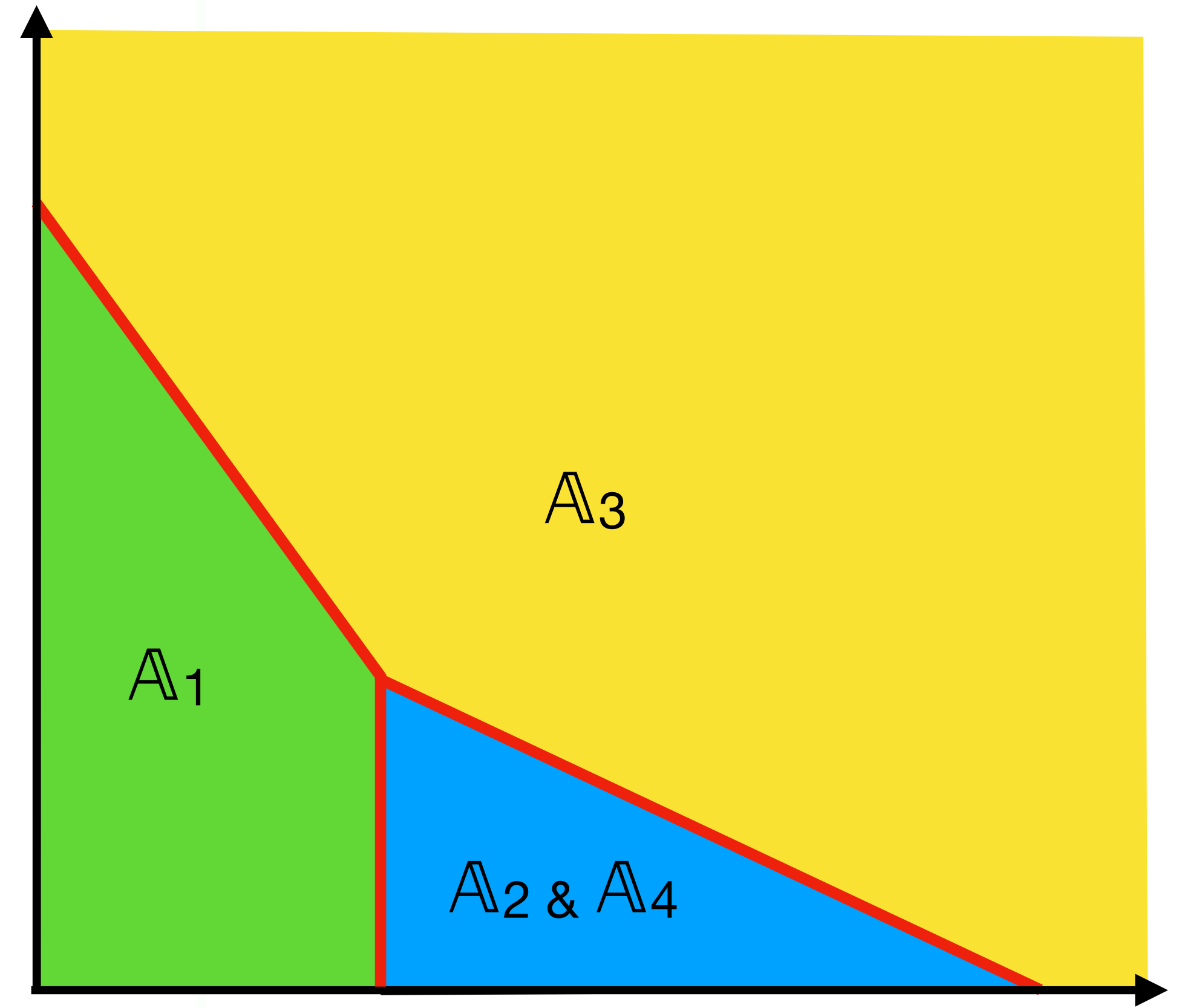
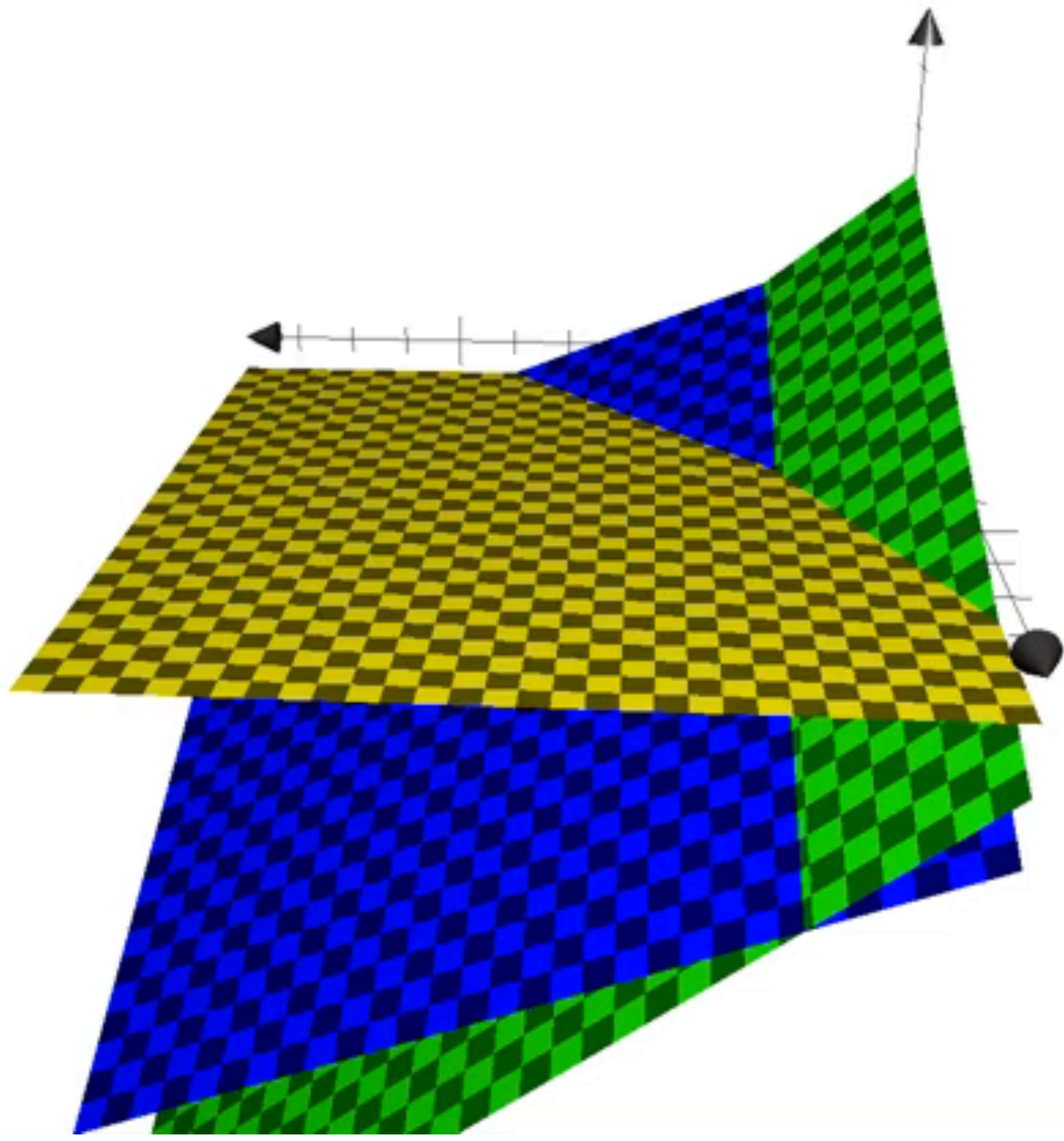
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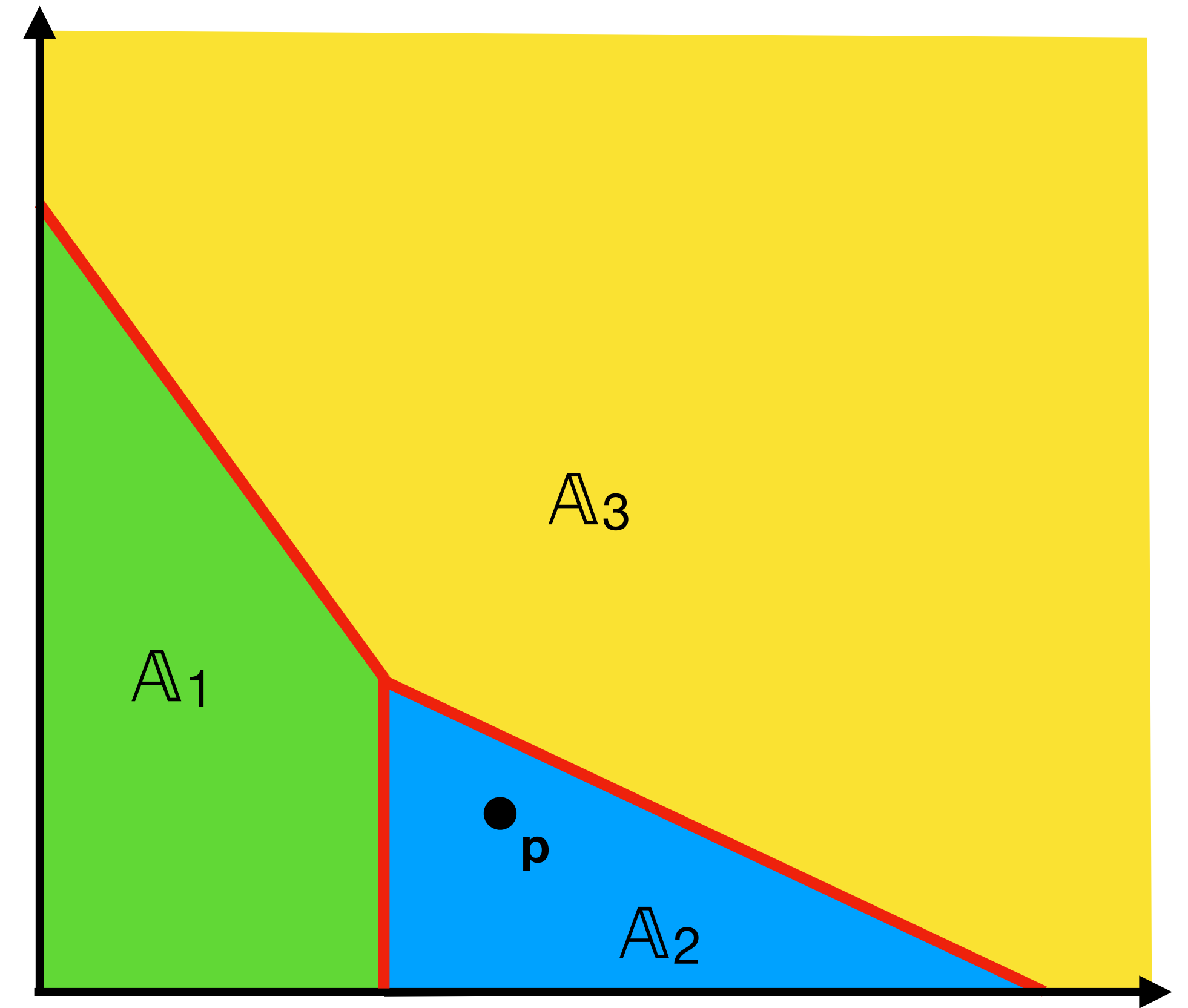


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If \mathbb{A} is optimal at some point p , it is on the correct side of the line that separates \mathbb{A} from all \mathbb{A}' .

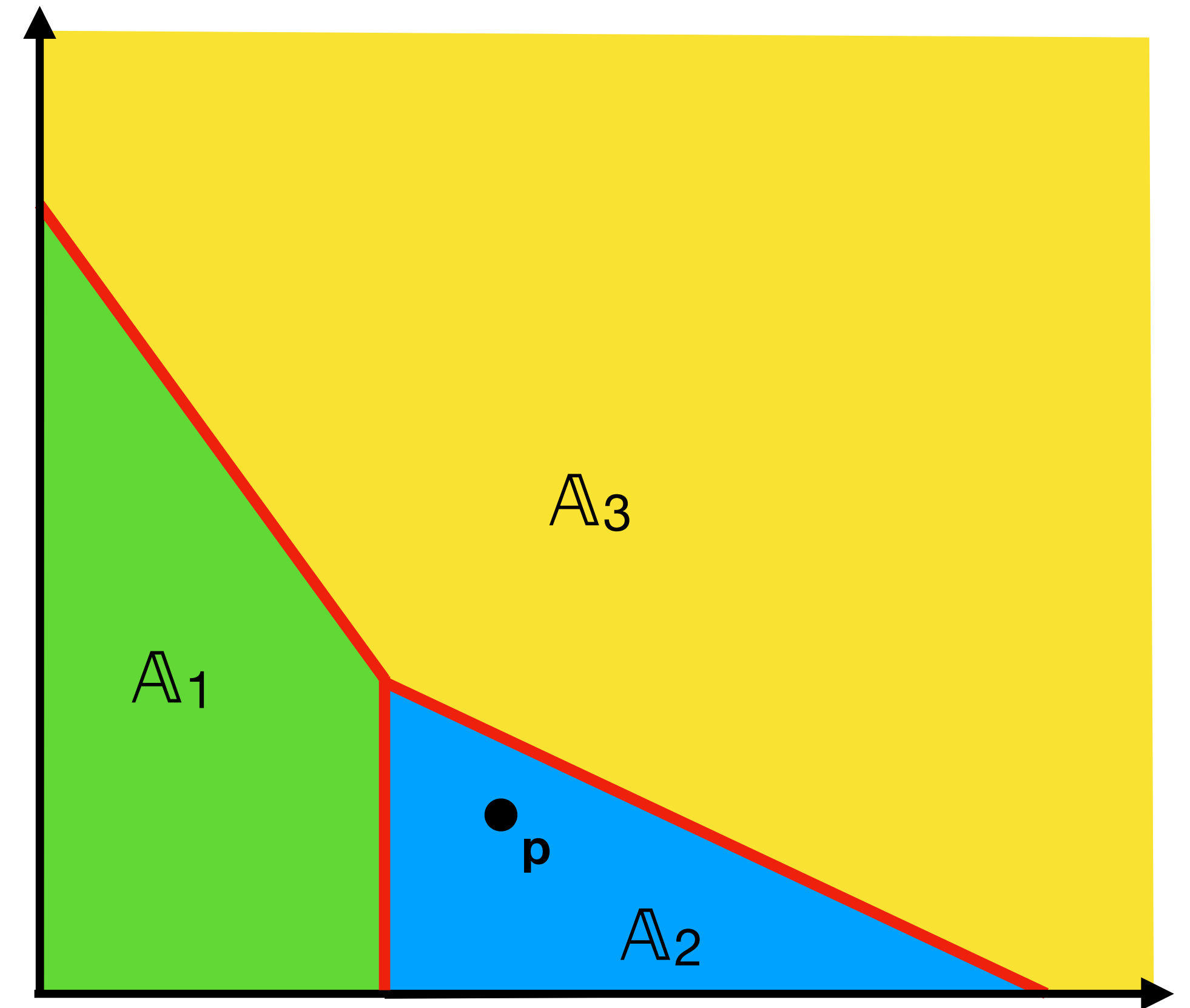


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If \mathbb{A} is optimal for at least 1 point p in the (γ, δ) -space then it is optimal for:

- (1) only point p ,
- (2) only a line segment that contains p , or
- (3) a convex polygon that contains p



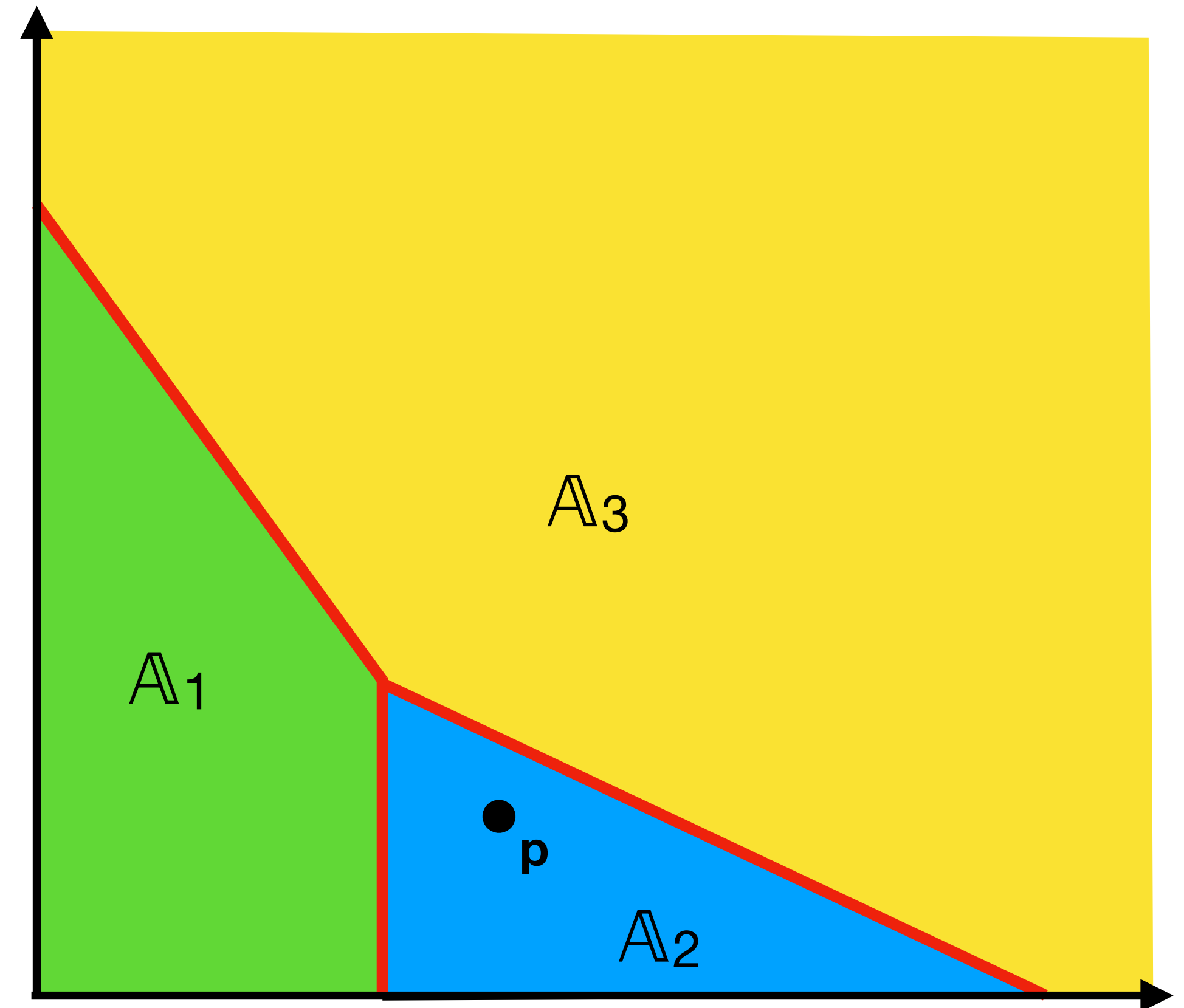
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Given two strings s_1 and s_2 the (γ, δ) -space decomposes into convex polygons such that any point in the interior of the polygon P is optimal for all points in P



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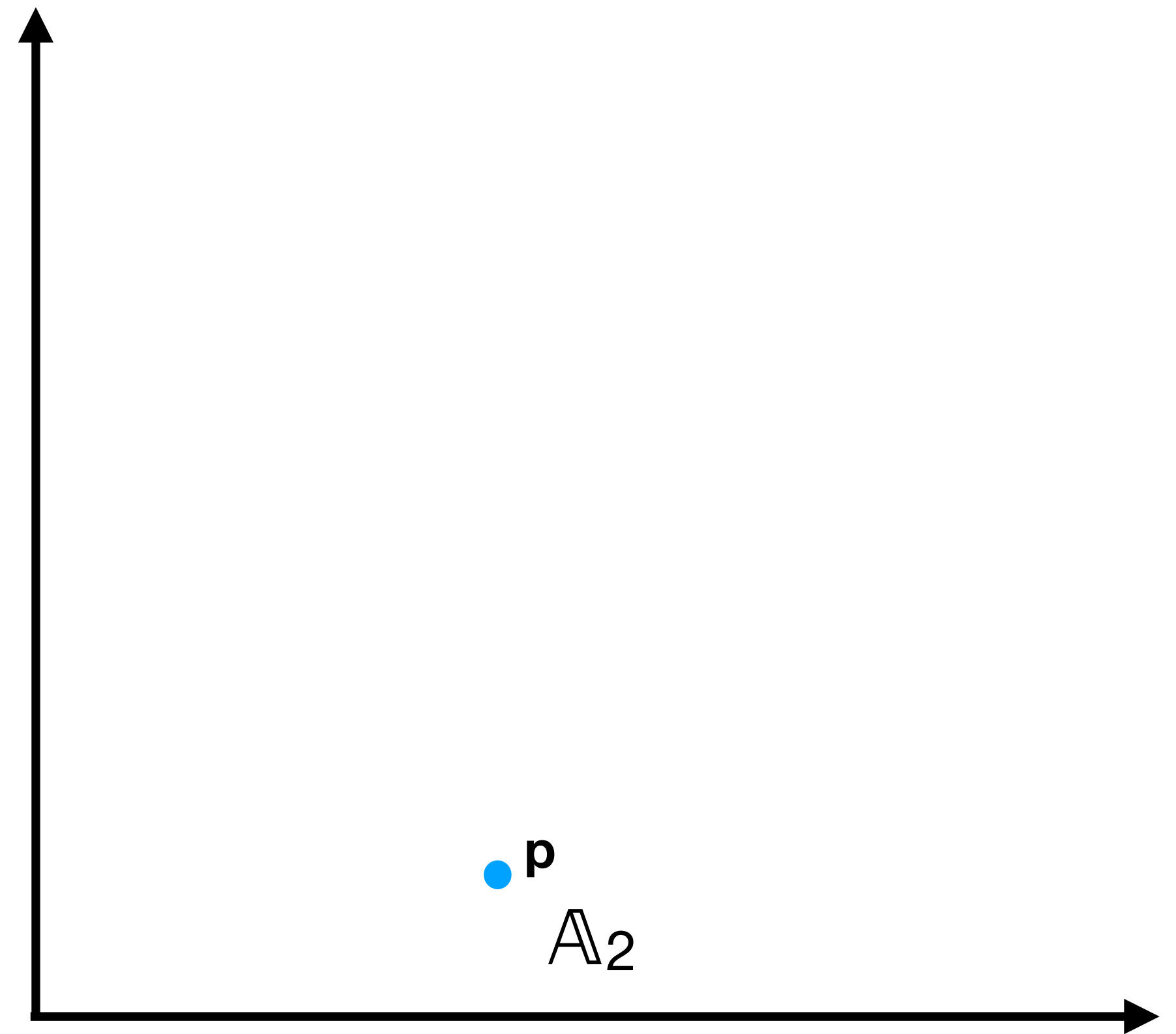
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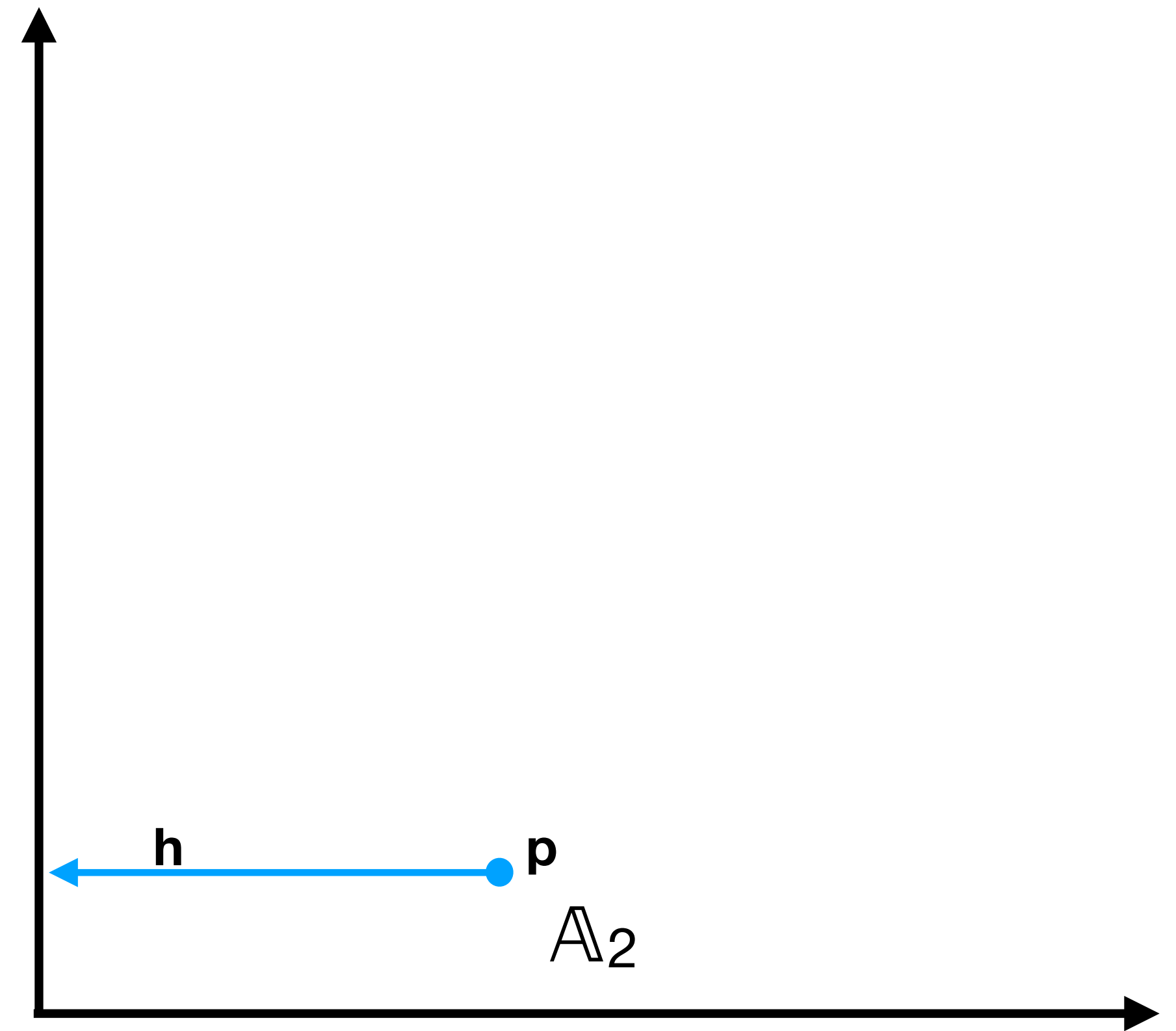
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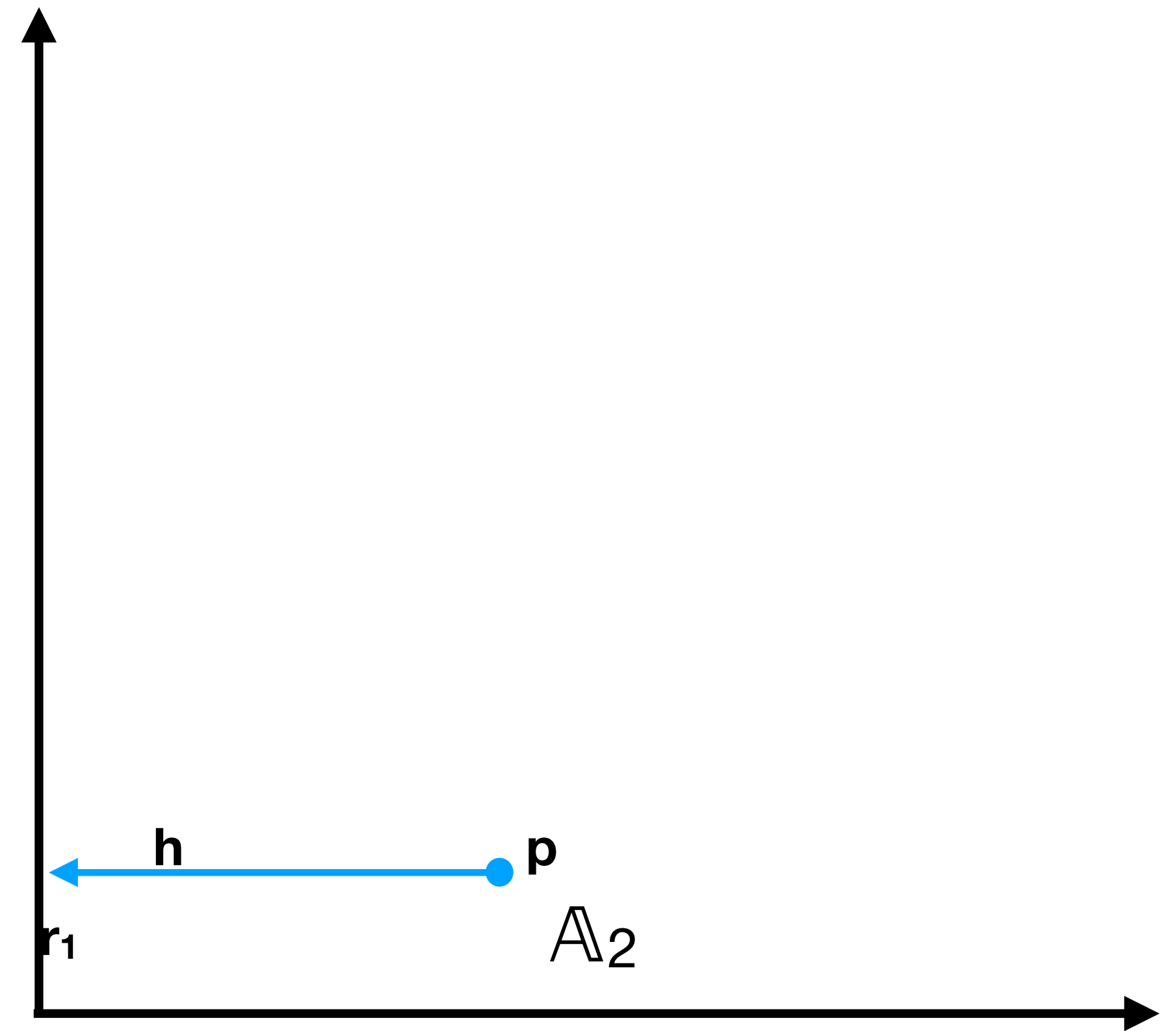
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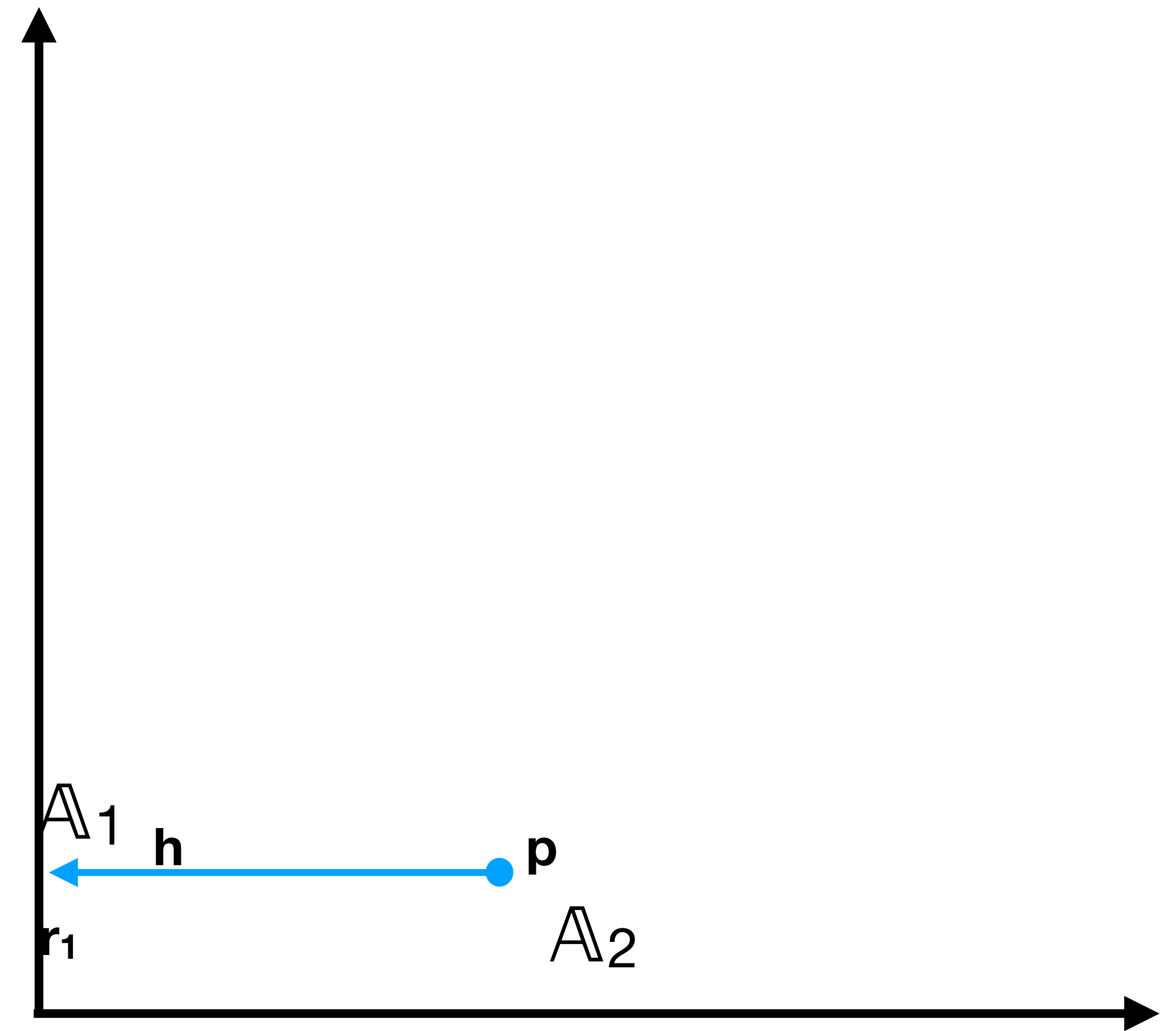


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Find alignment A' that is optimal at r_1 .



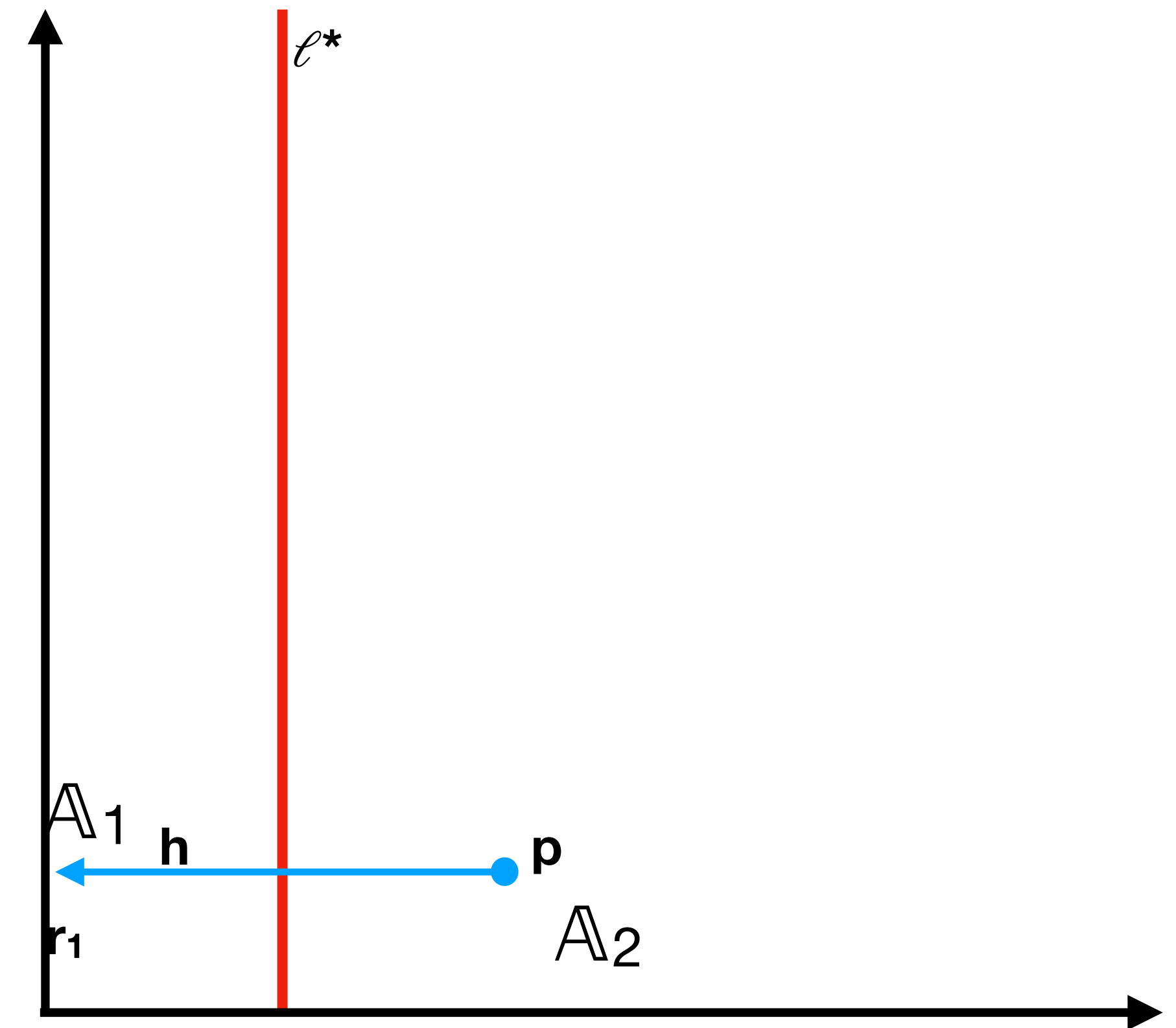
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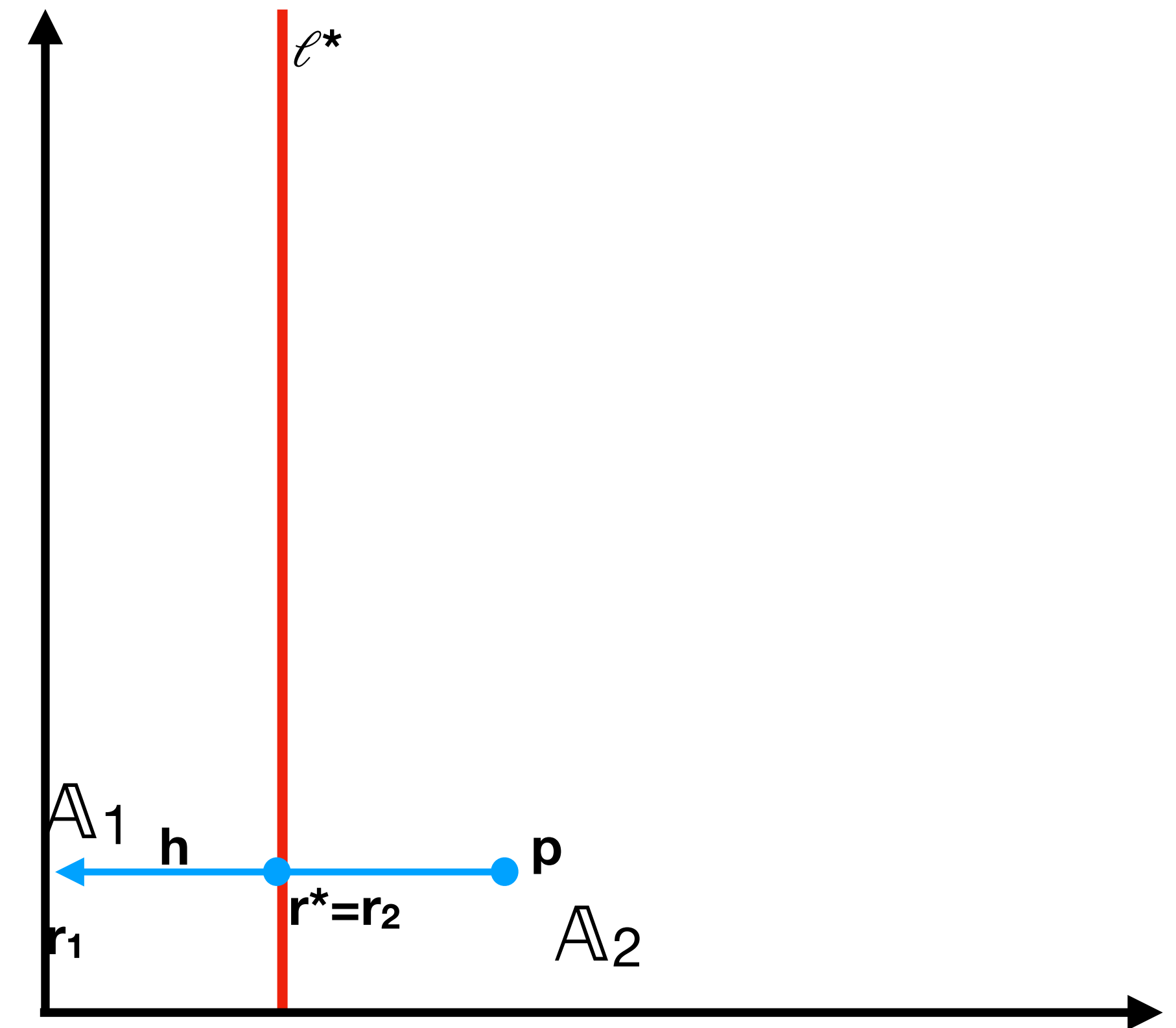
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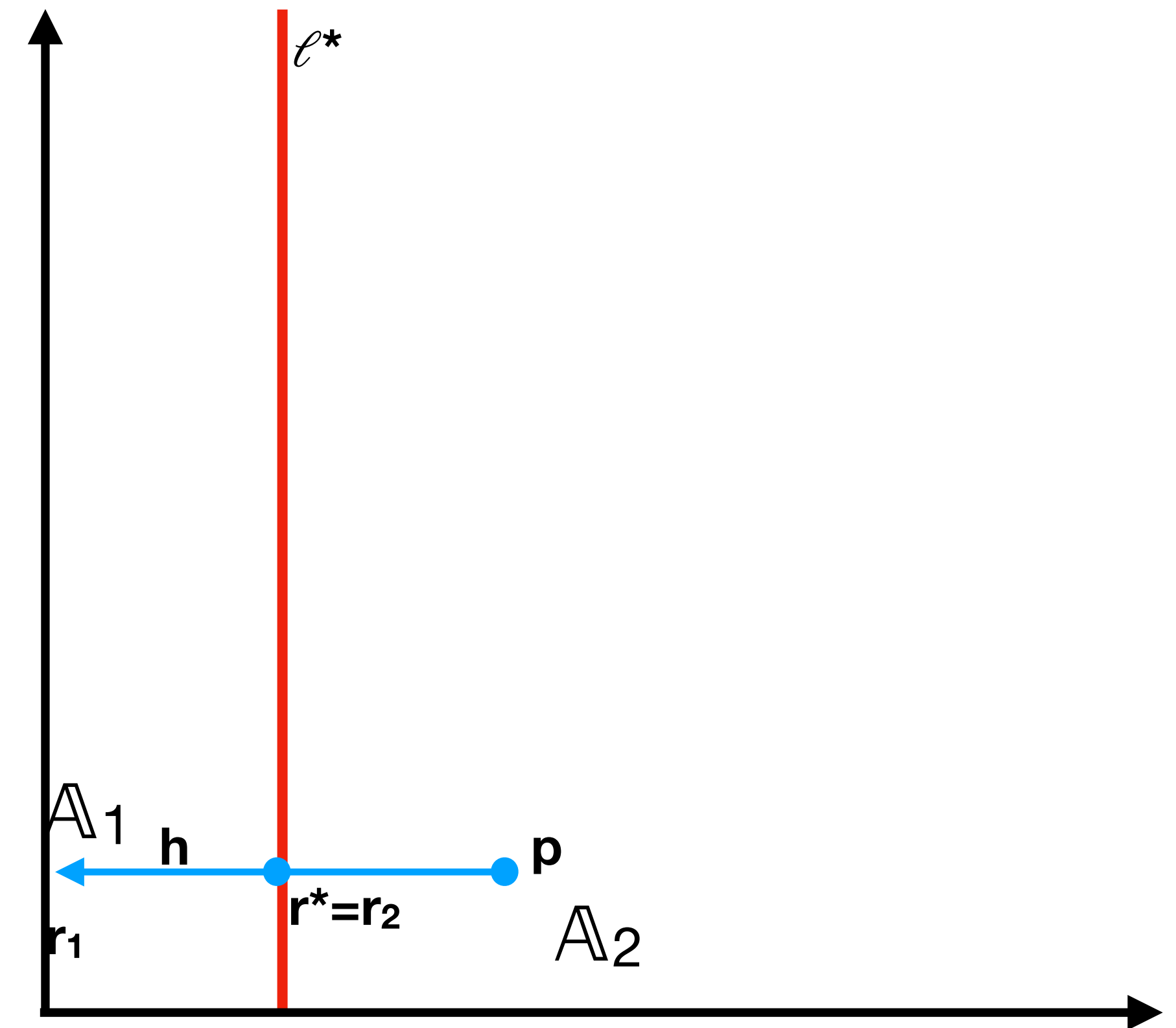
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The boundary for the polygon in which p resides is a segment of that line.



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note: it follows that any polygon P intersected by h , a single ray search computes alignments at no more than 2 points of P .

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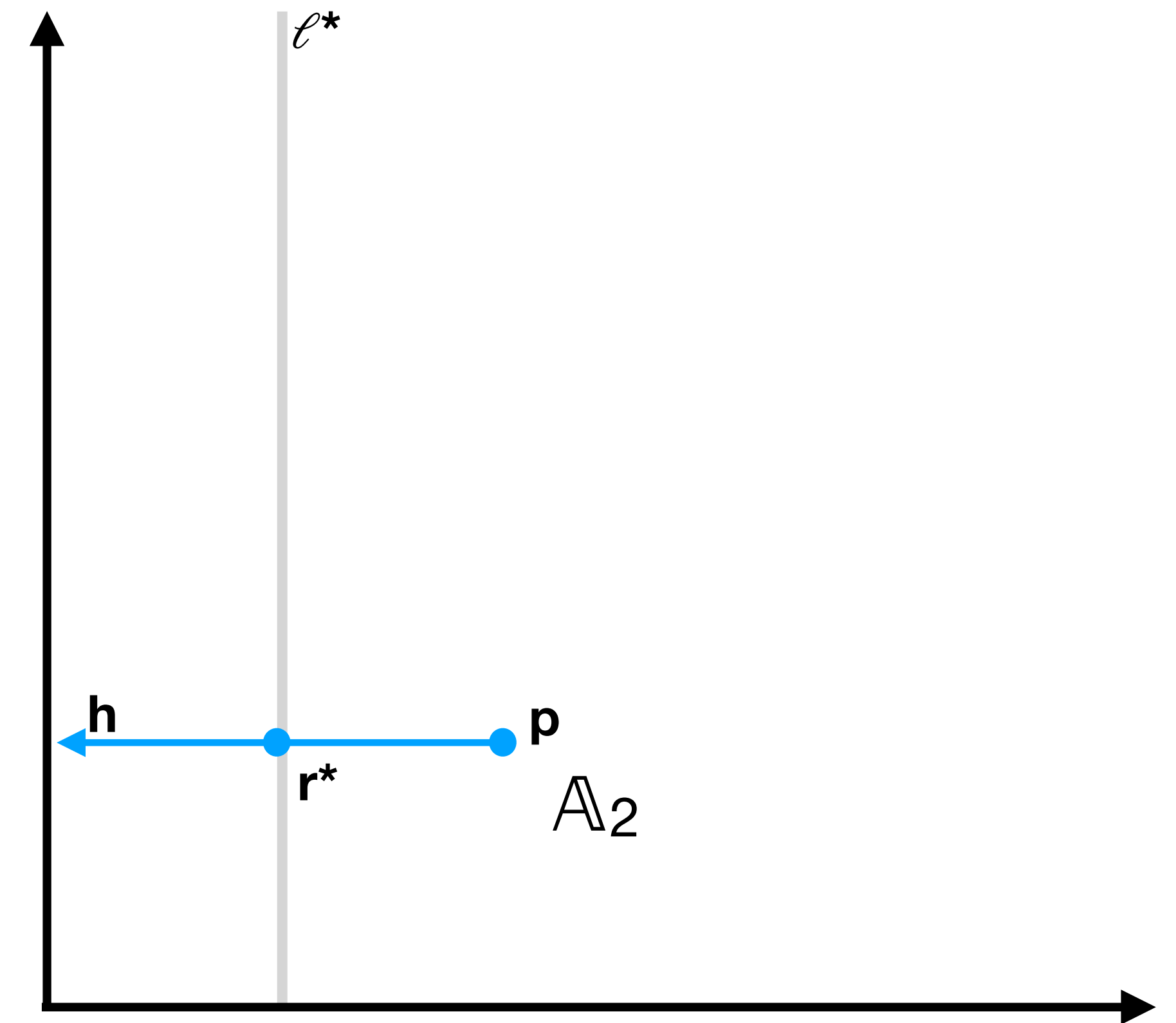
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**How many regions are there?
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Finding an edge of the polygon

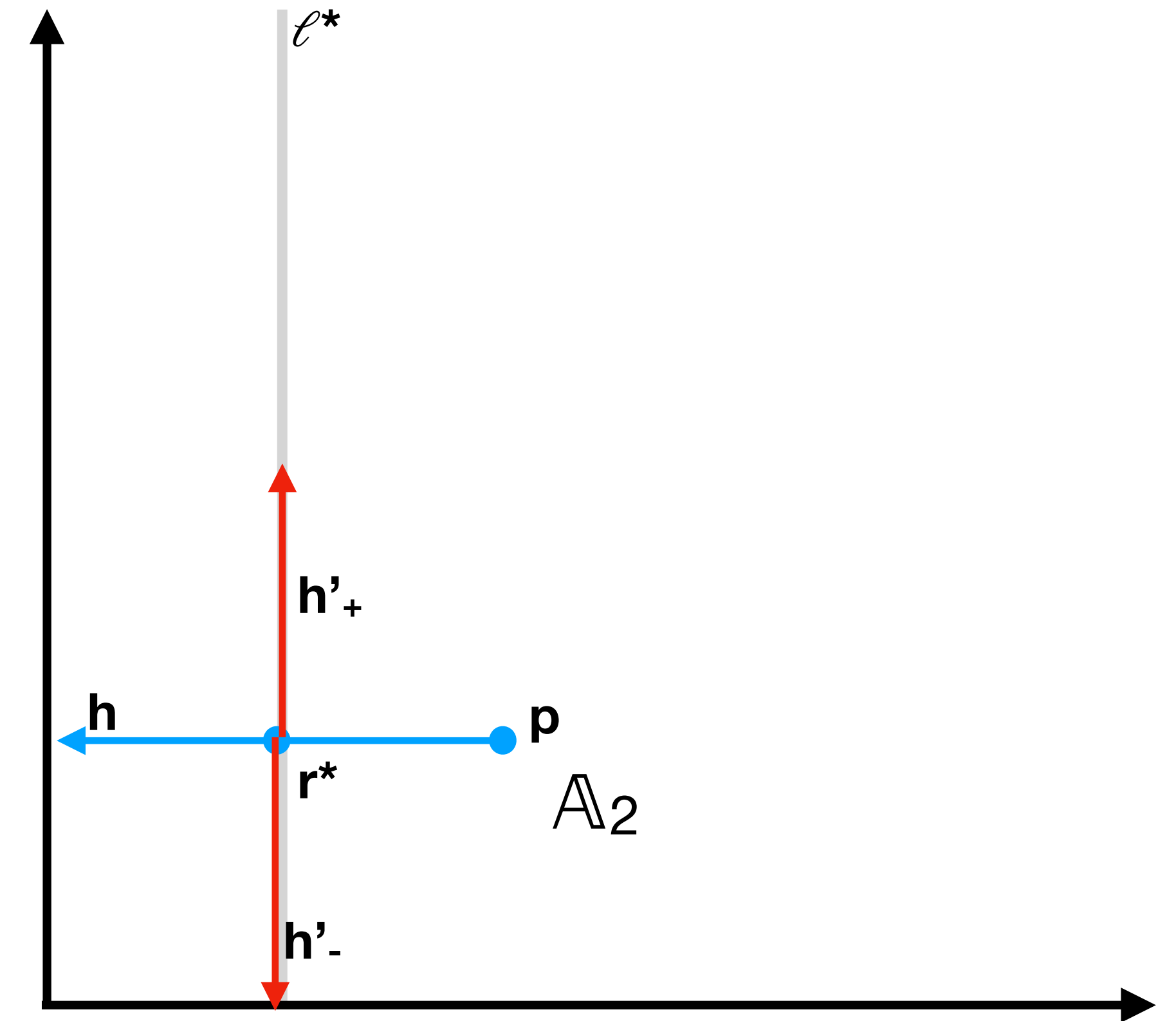
Given p , h , r^* , and the line that separates A from the next optimal alignment on h . (assume p is inside a polygon)



Finding an edge of the polygon

Given p , h , r^* , and the line that separates \mathbb{A} from the next optimal alignment on h . (assume p is inside a polygon)

Perform a ray search in both directions from r^* , along the line.

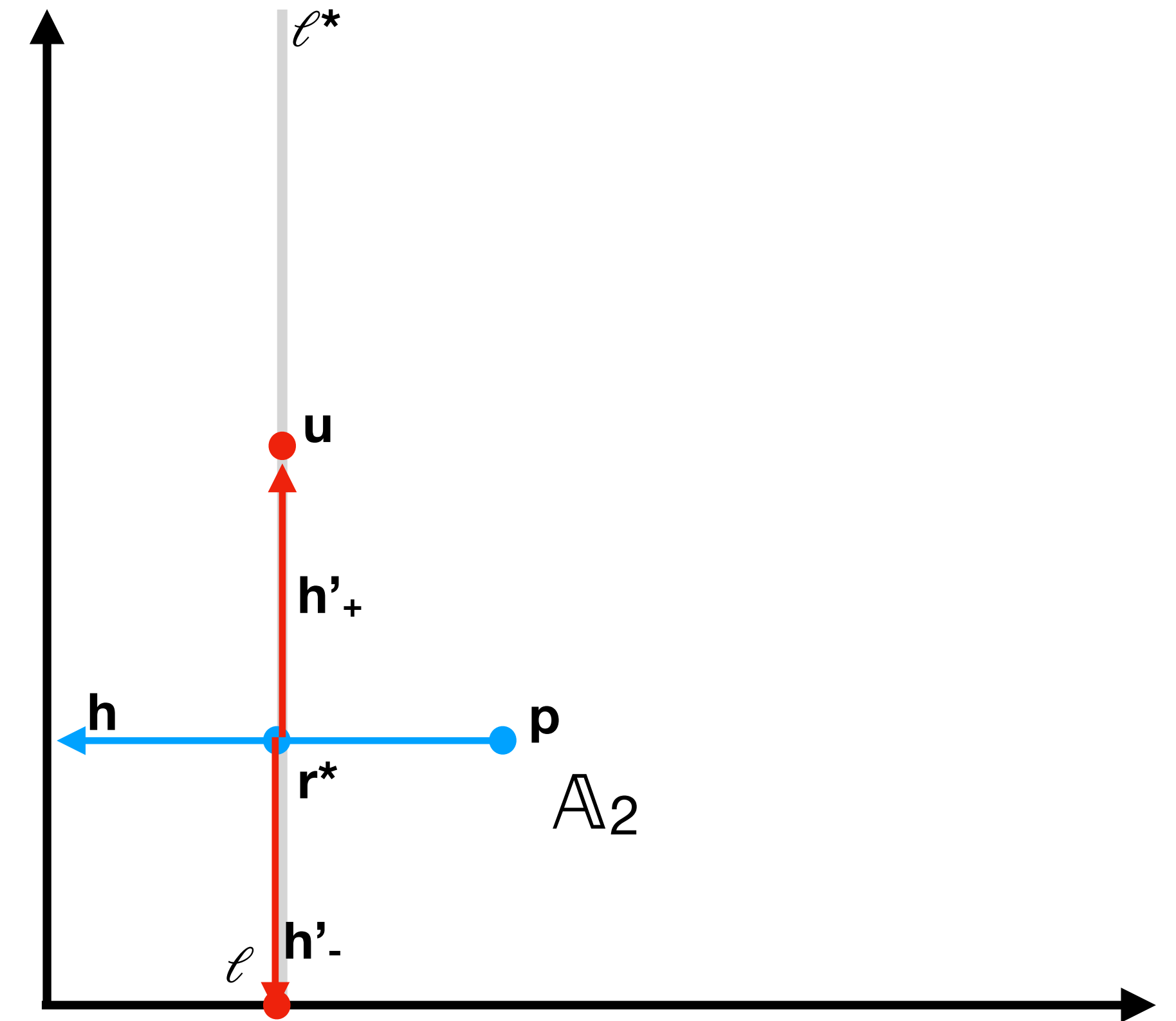


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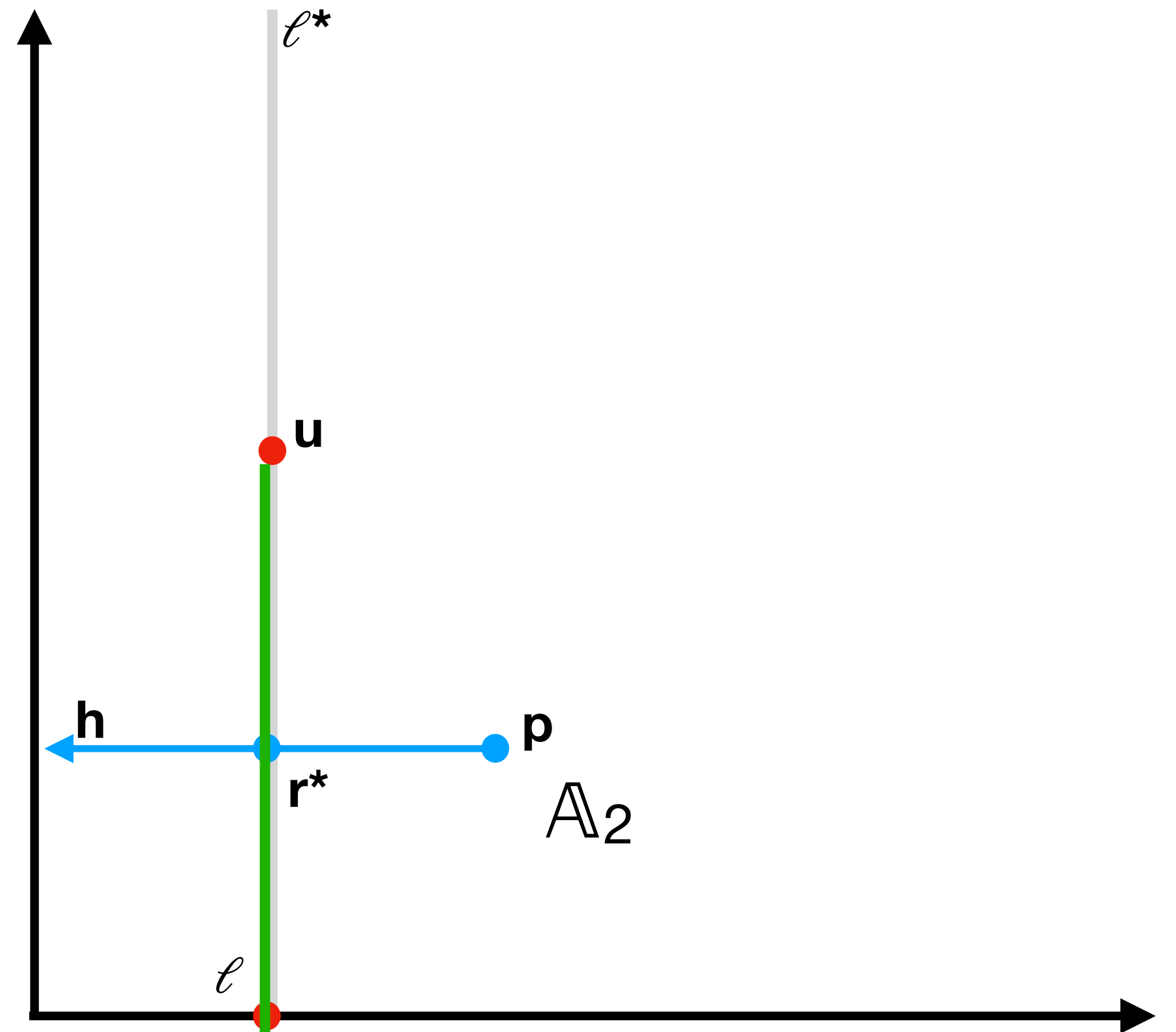
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The line segment (l,u) is a face of the polygon for \mathbb{A} .



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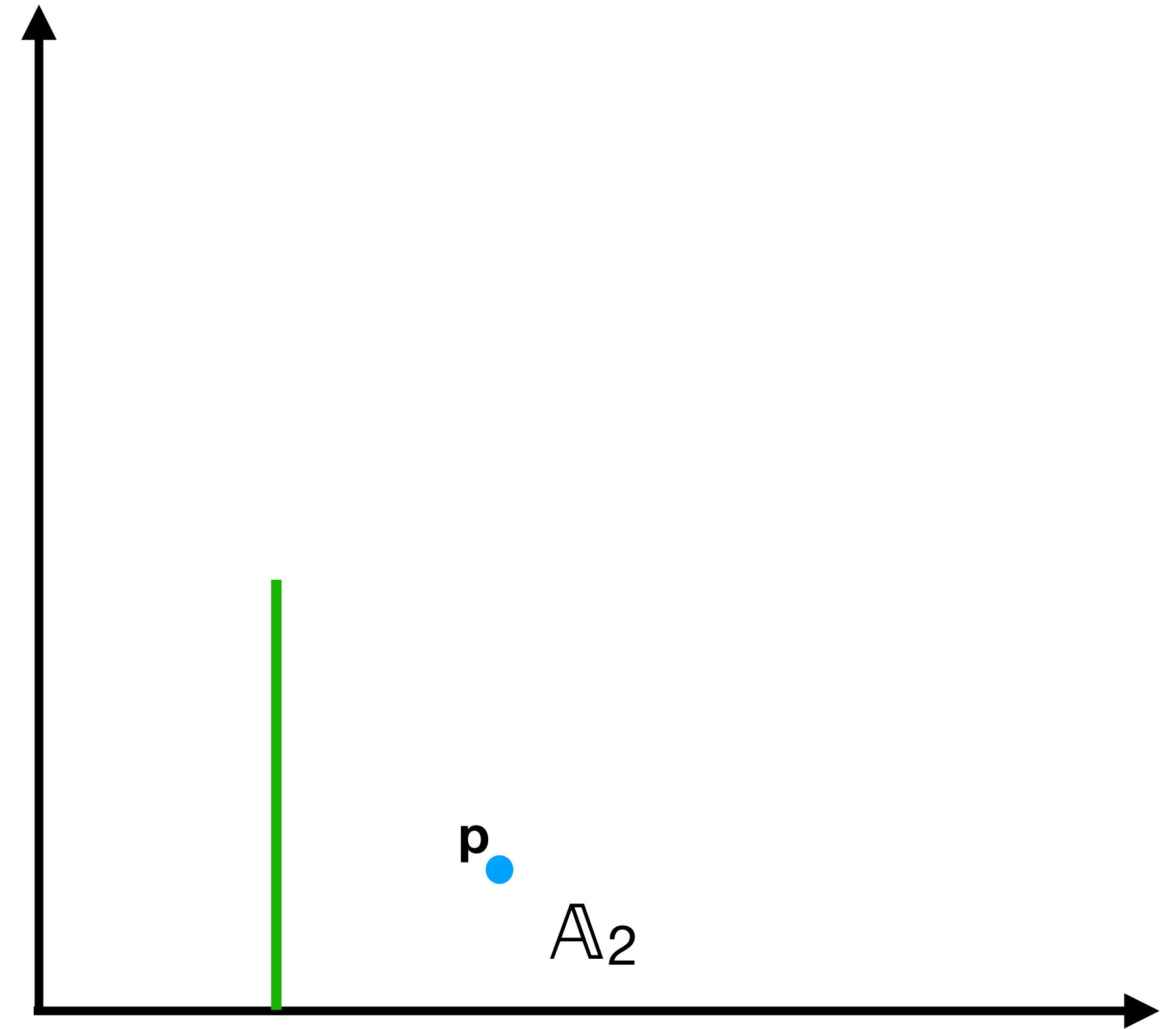
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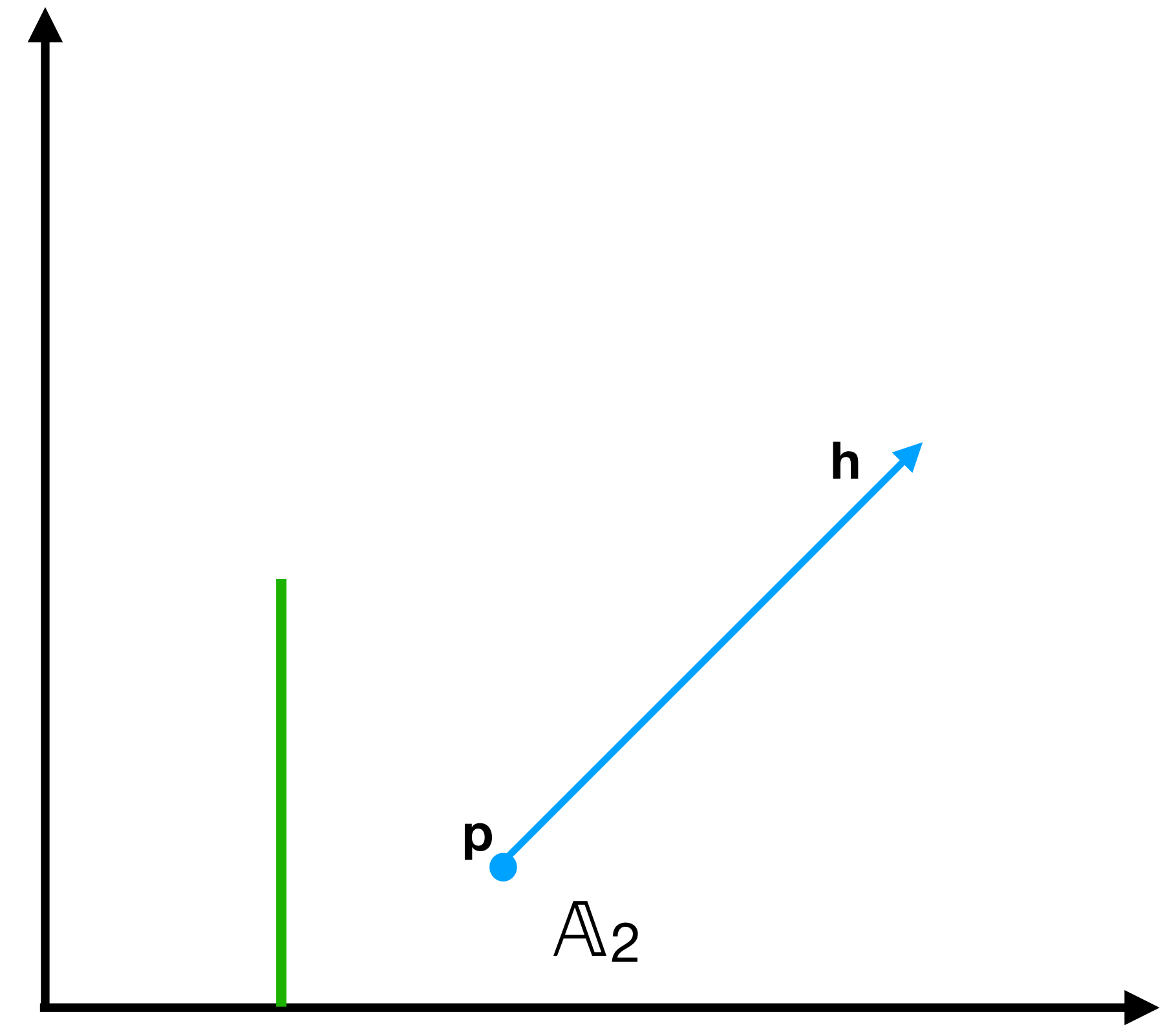
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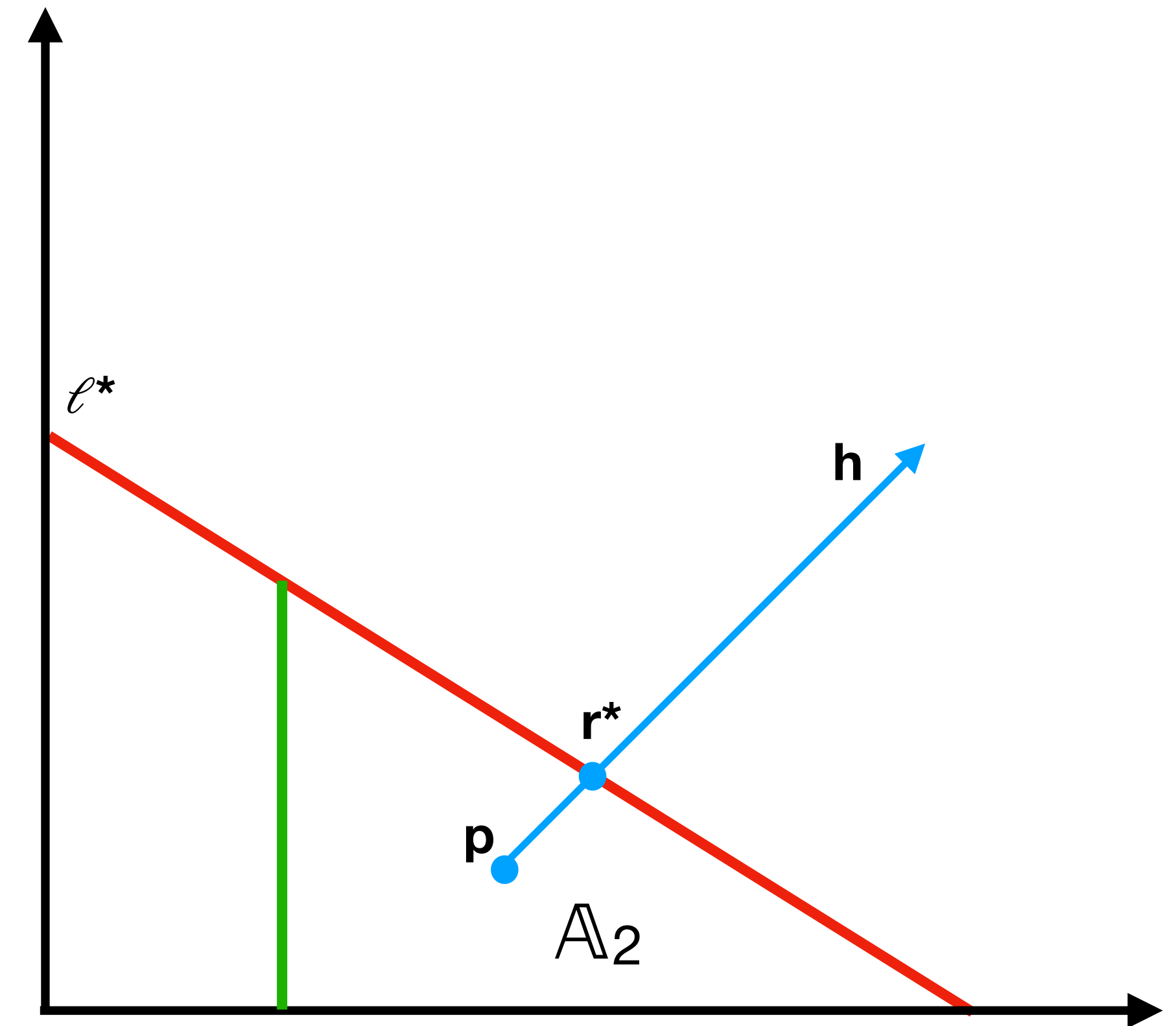


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Apply the ray finding algorithm to find r^* .



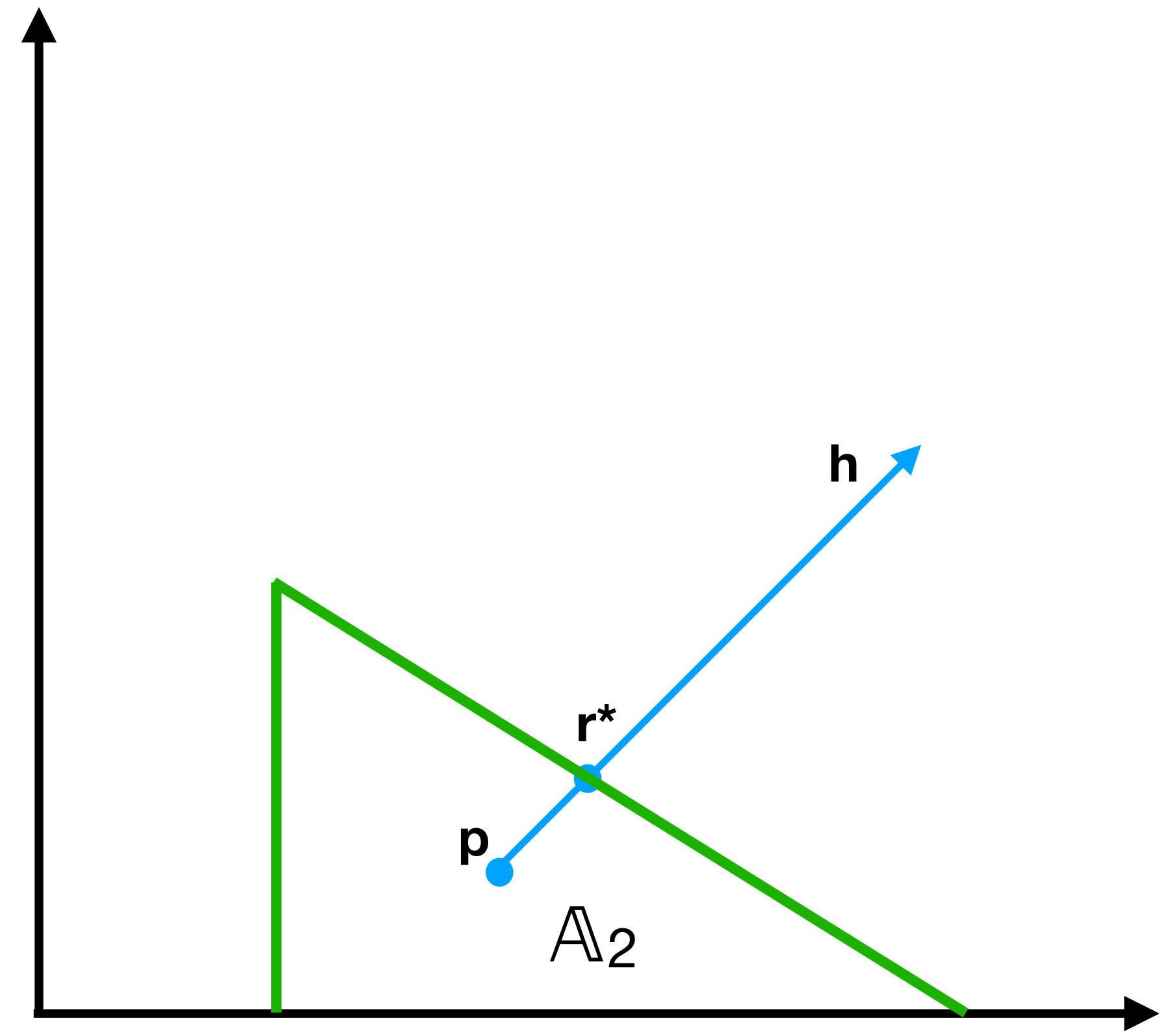
Finding other faces

Given a point p , and a subset of the faces of the polygon in which p resides. (assume p is inside a polygon).

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Apply the ray finding twice to find the face of the polygon that intersects r^* .



Completing the polygon

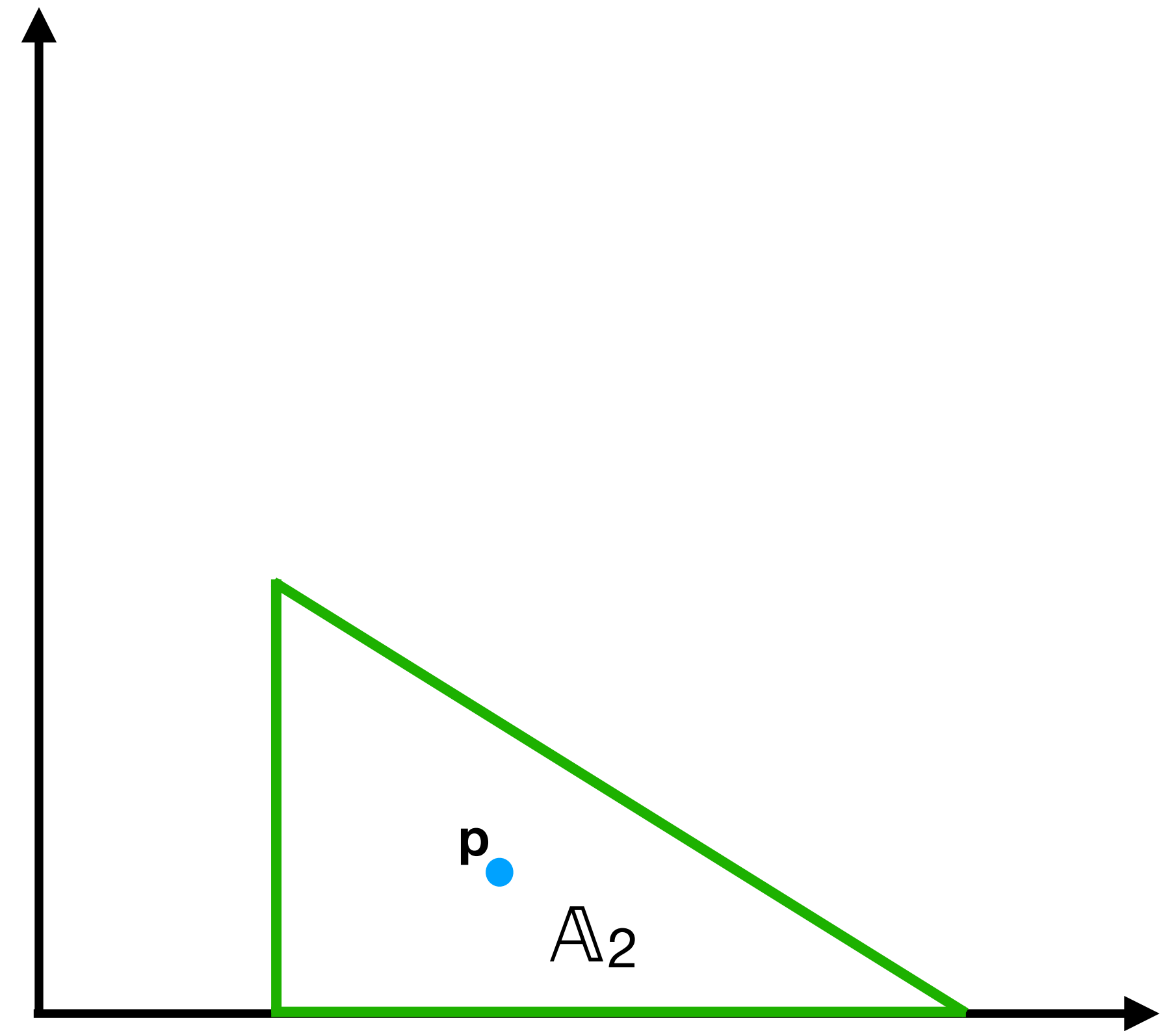
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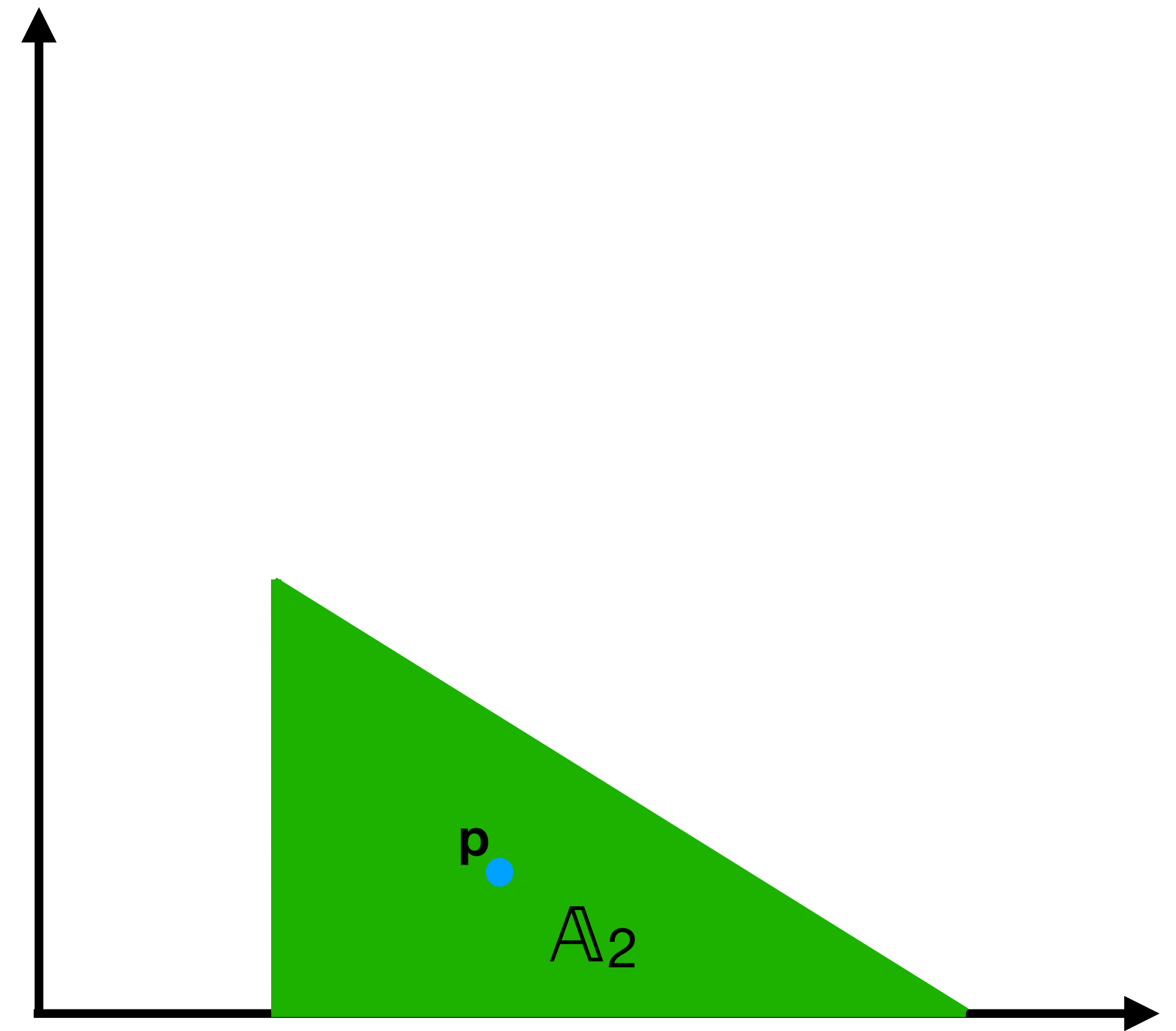
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r^* is a vertex of the polygon

- one of the ray searches along l^* will not find any point beyond r^* .
- Stop and use another h that avoids the current r^* .

Things we know so far

For a parameter setting, we can find the optimal alignment.

Two alignments will have a line in (γ, δ) -space where they are co-optimal*.

For any point, the optimal alignment is optimal for a point, a line, or a region.

There are a limited number of regions for a fixed input.

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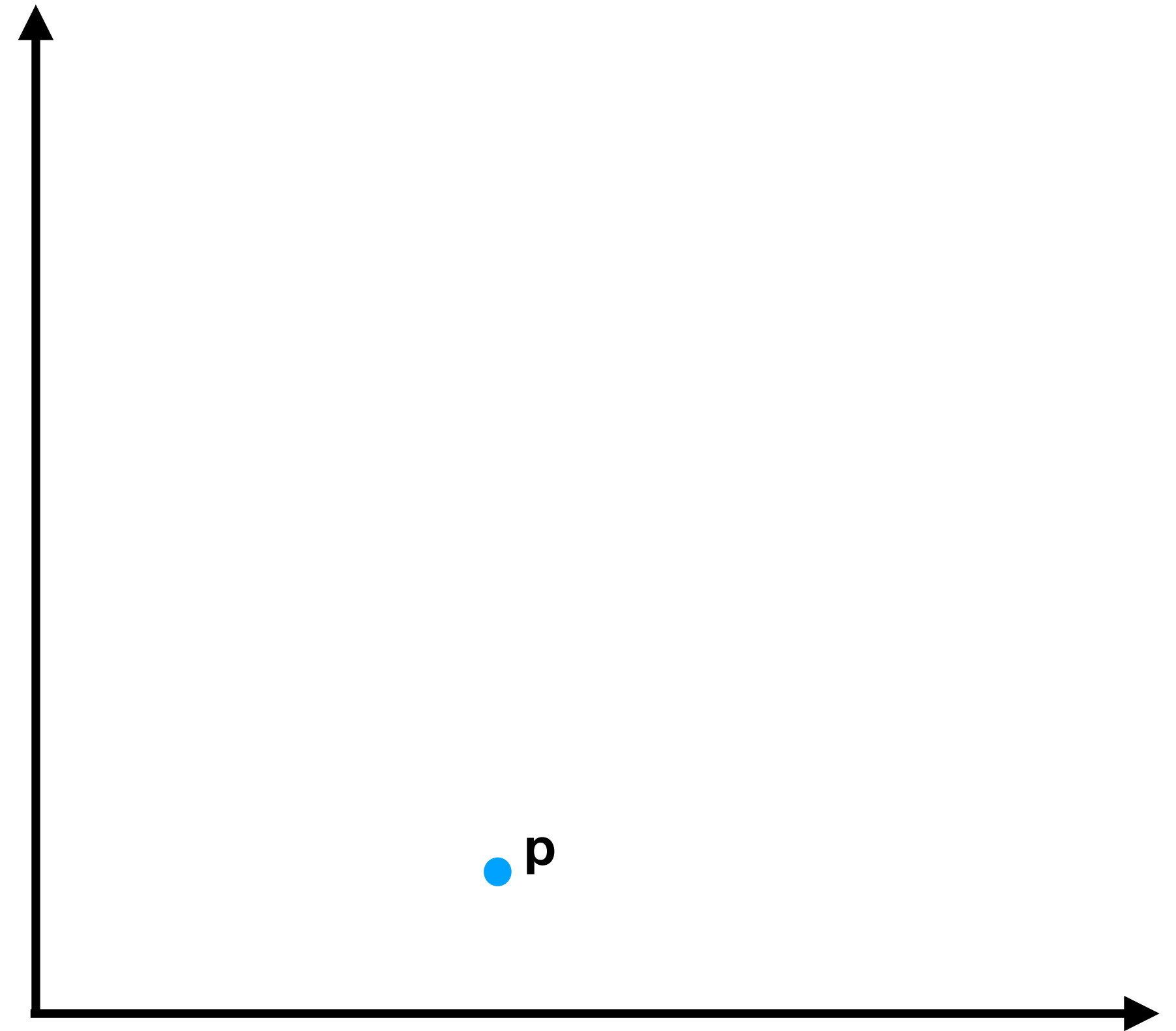
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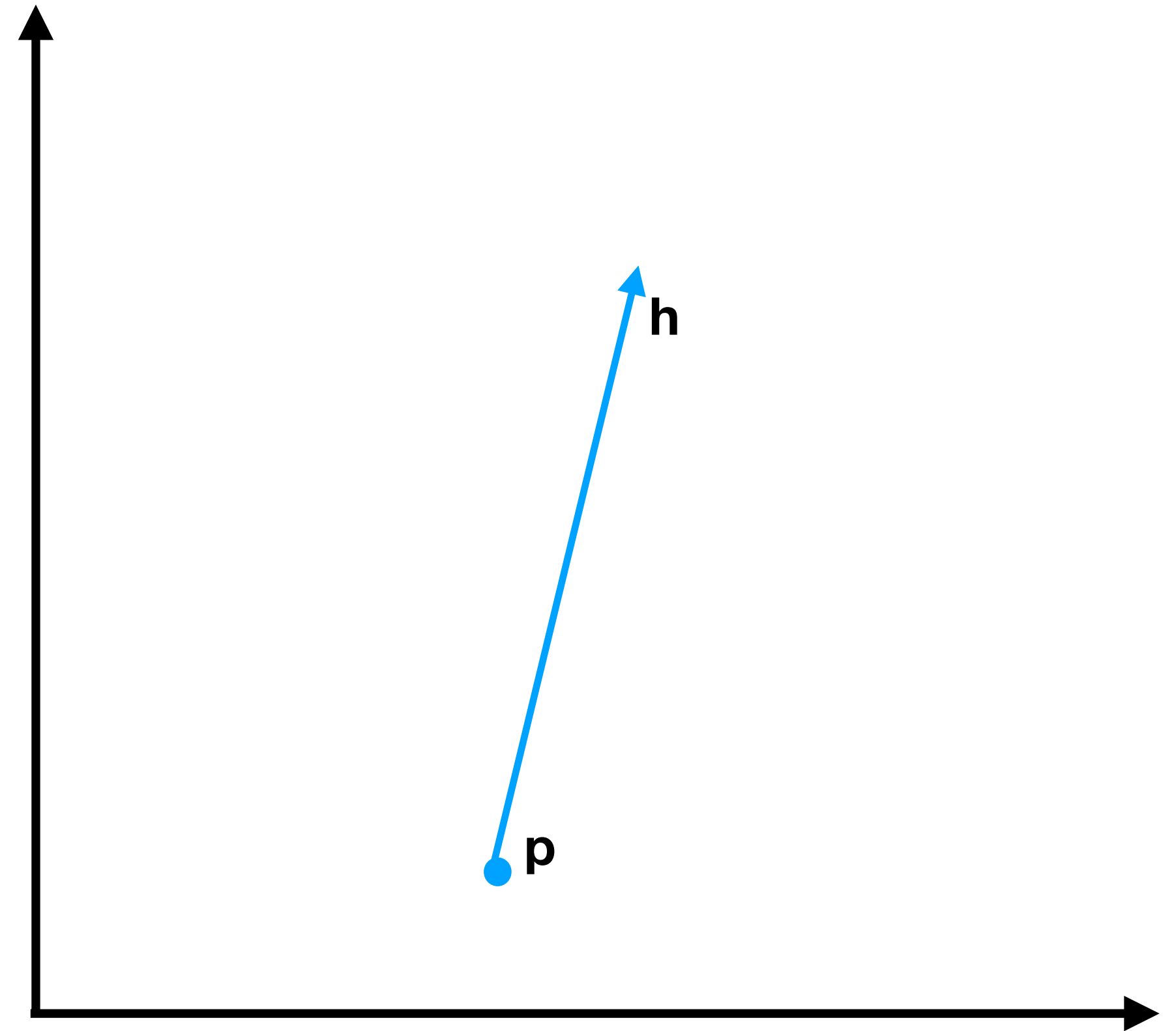
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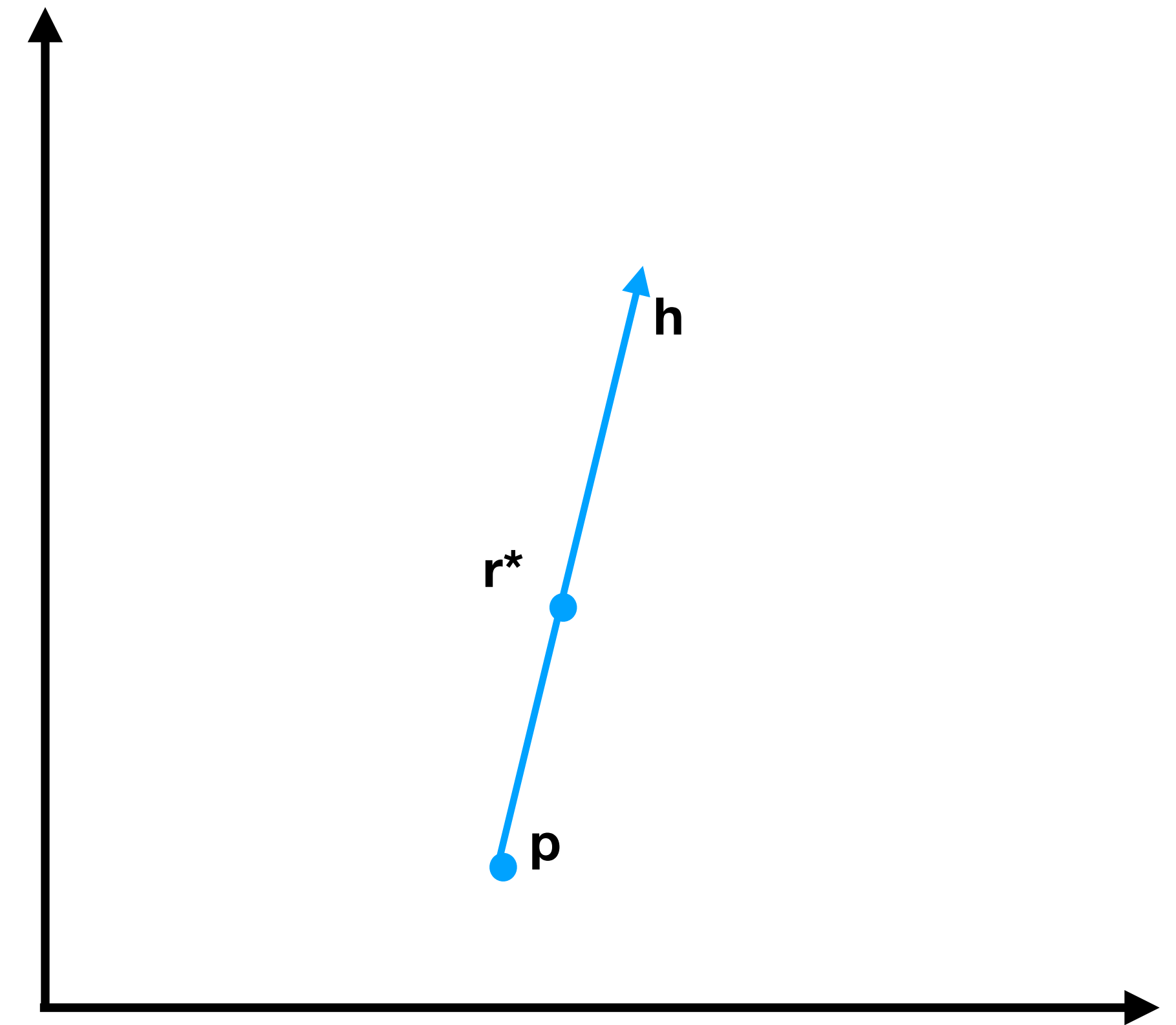


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If $r^* = p$ --- A^* is optimal for some non-zero distance along h .

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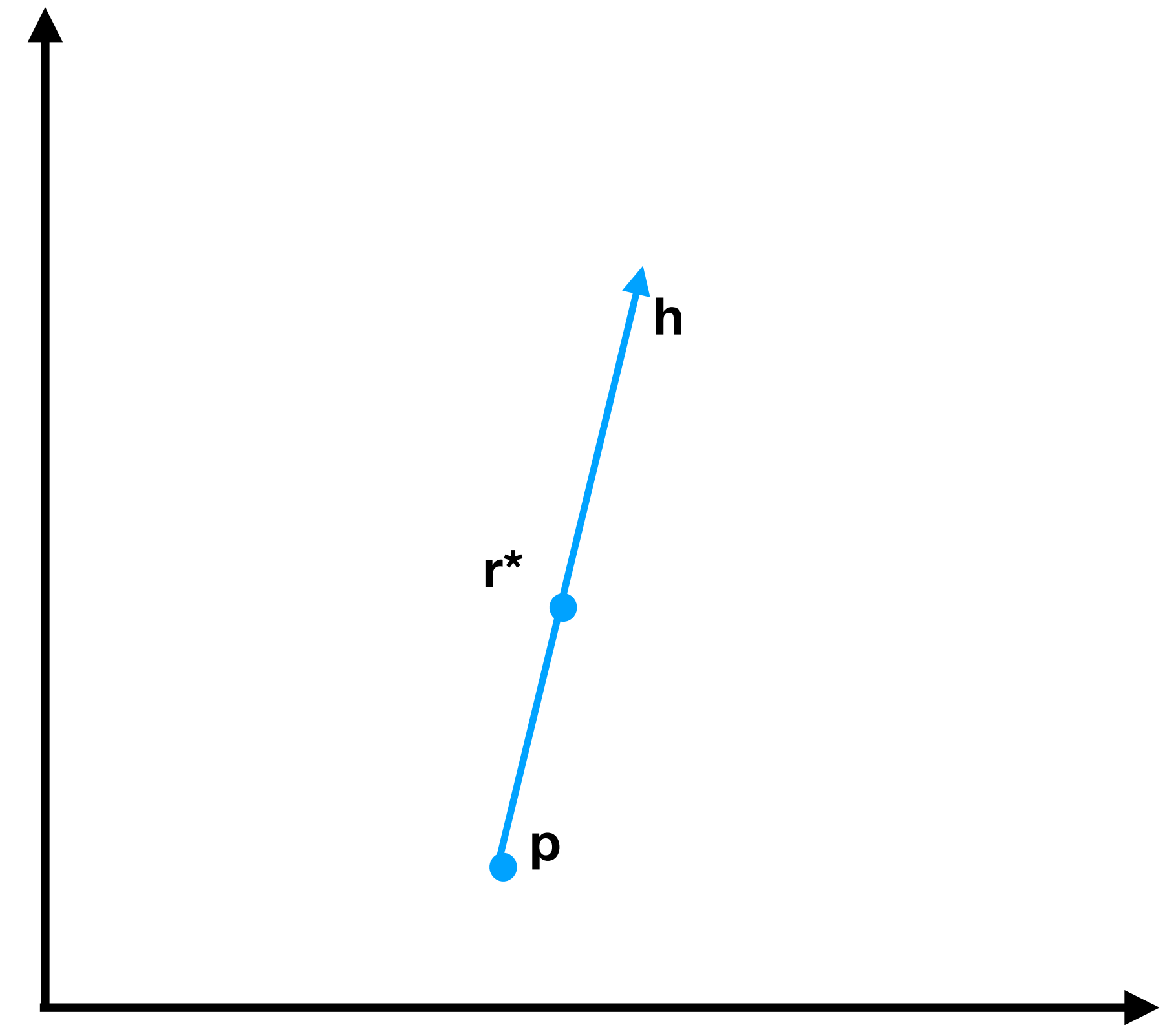
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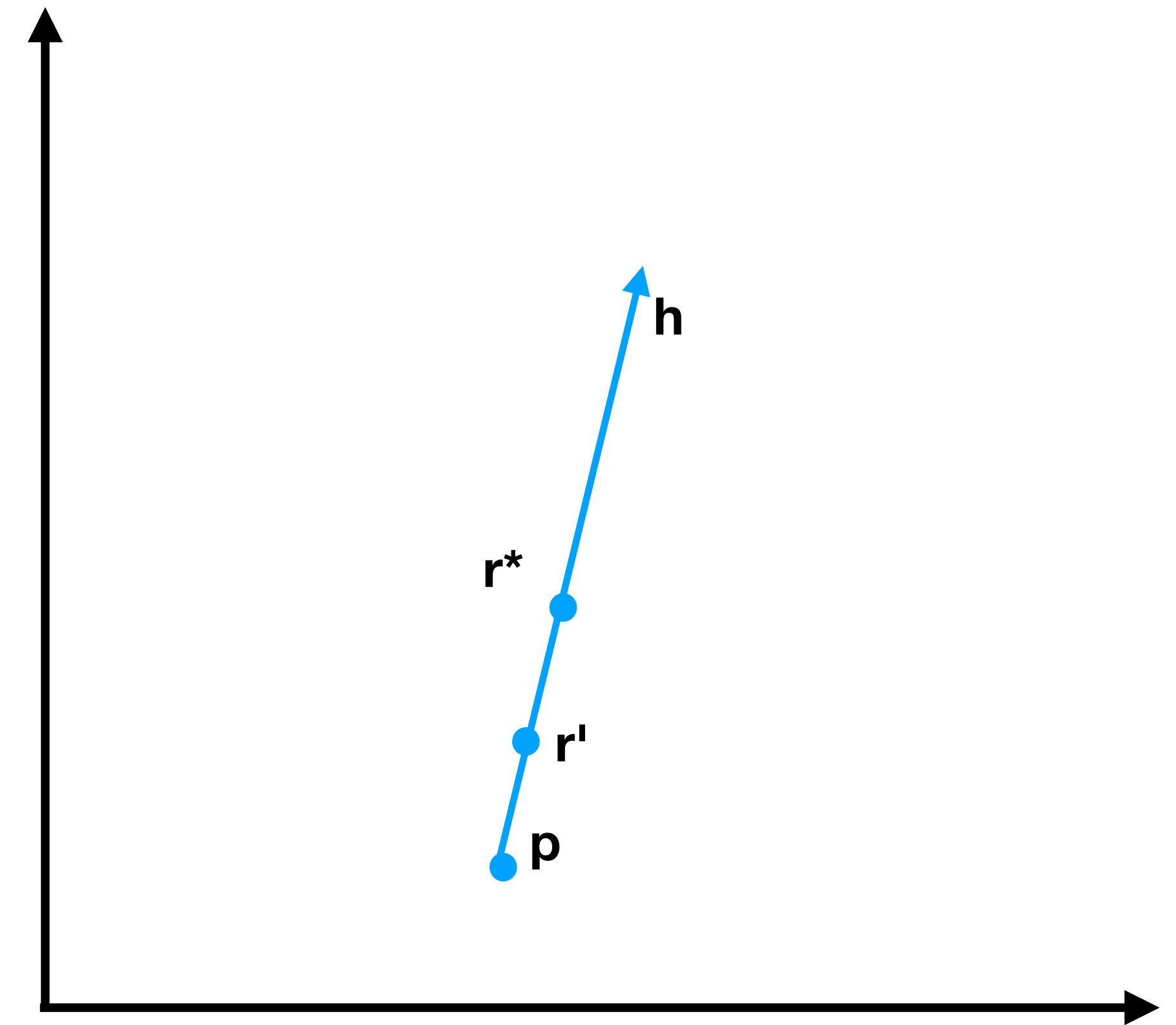
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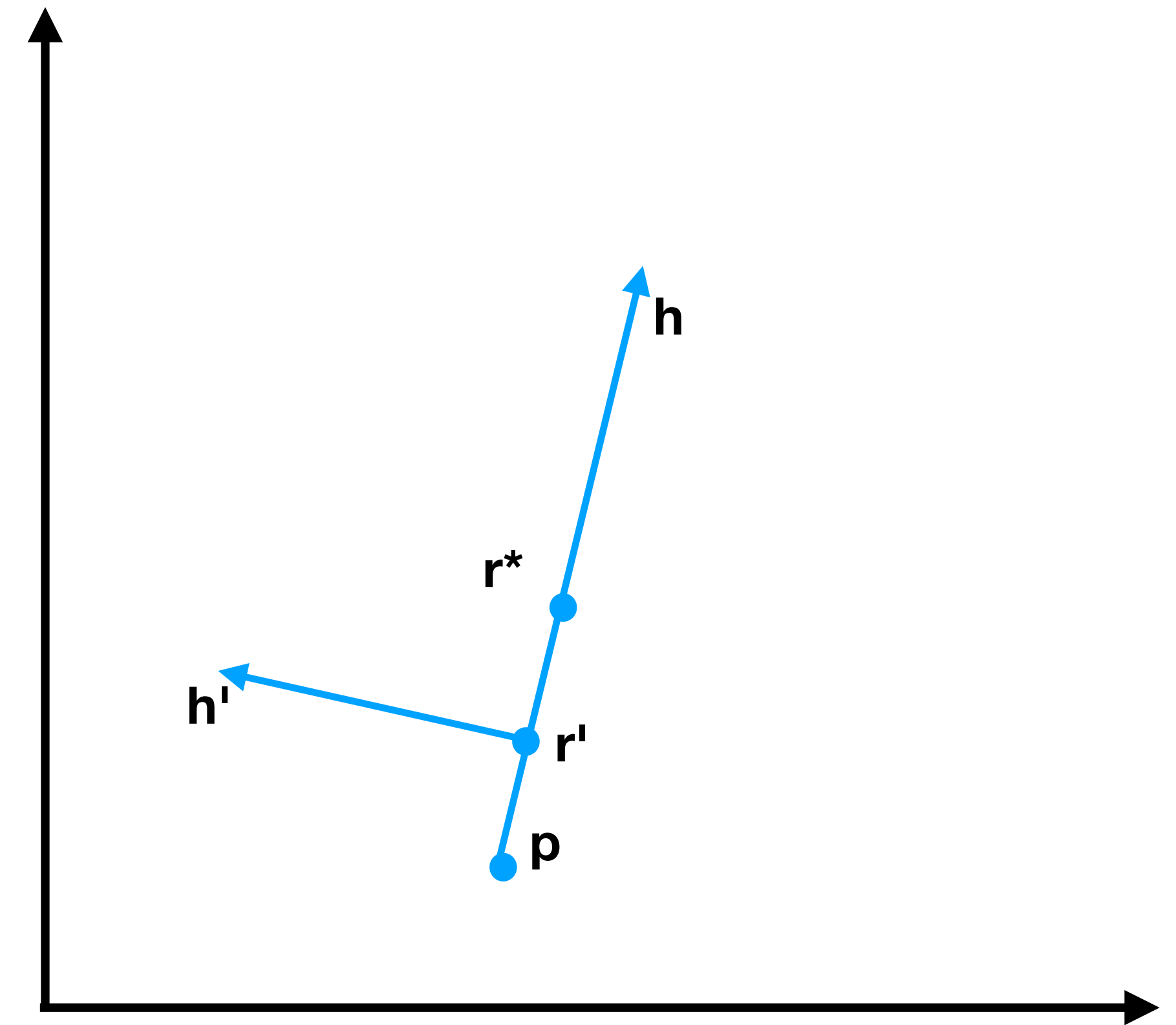
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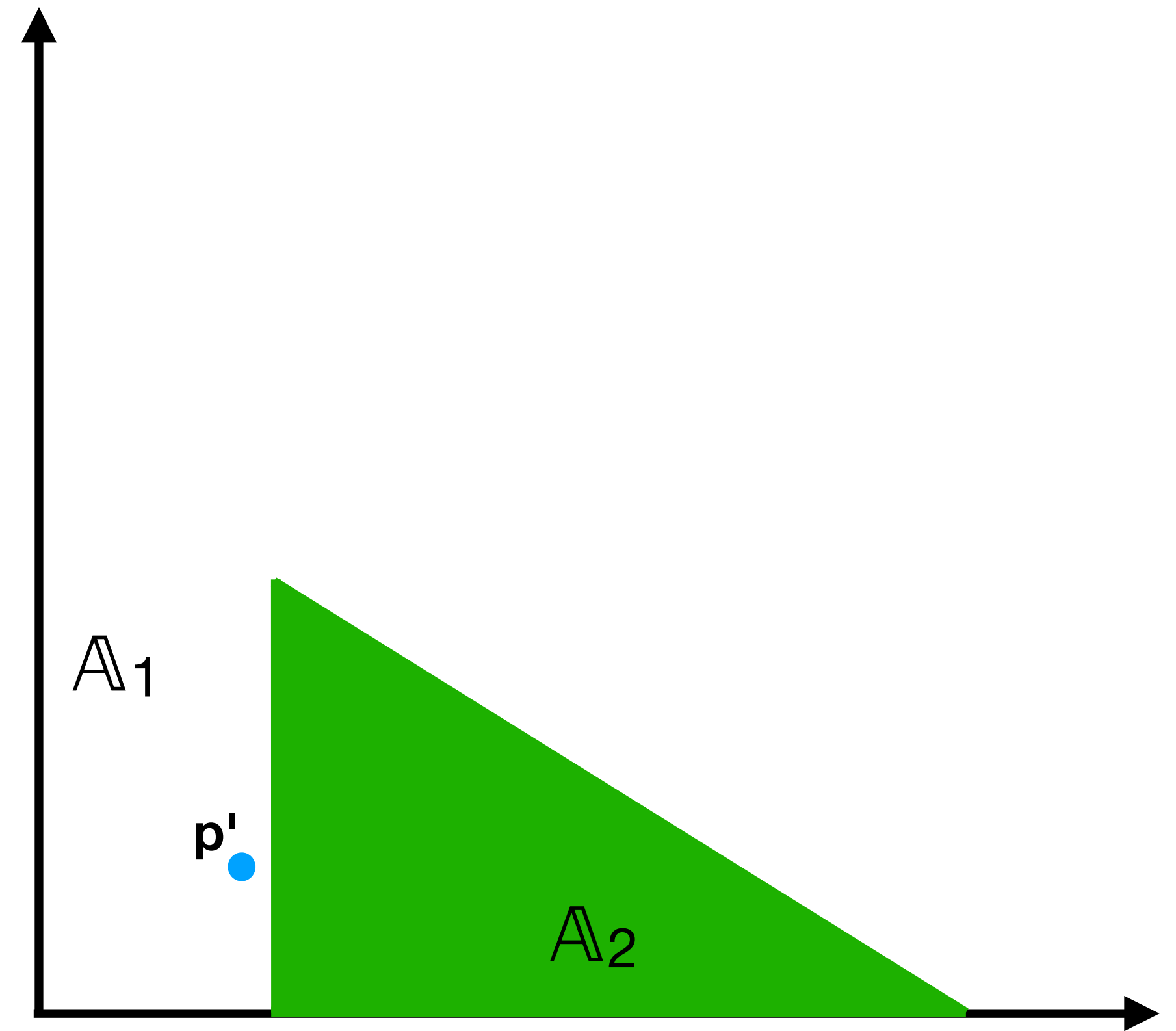
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A/A^* is either optimal for some distance along h' , or some alignment that is optimal for some distance is returned.



Completing the decomposition

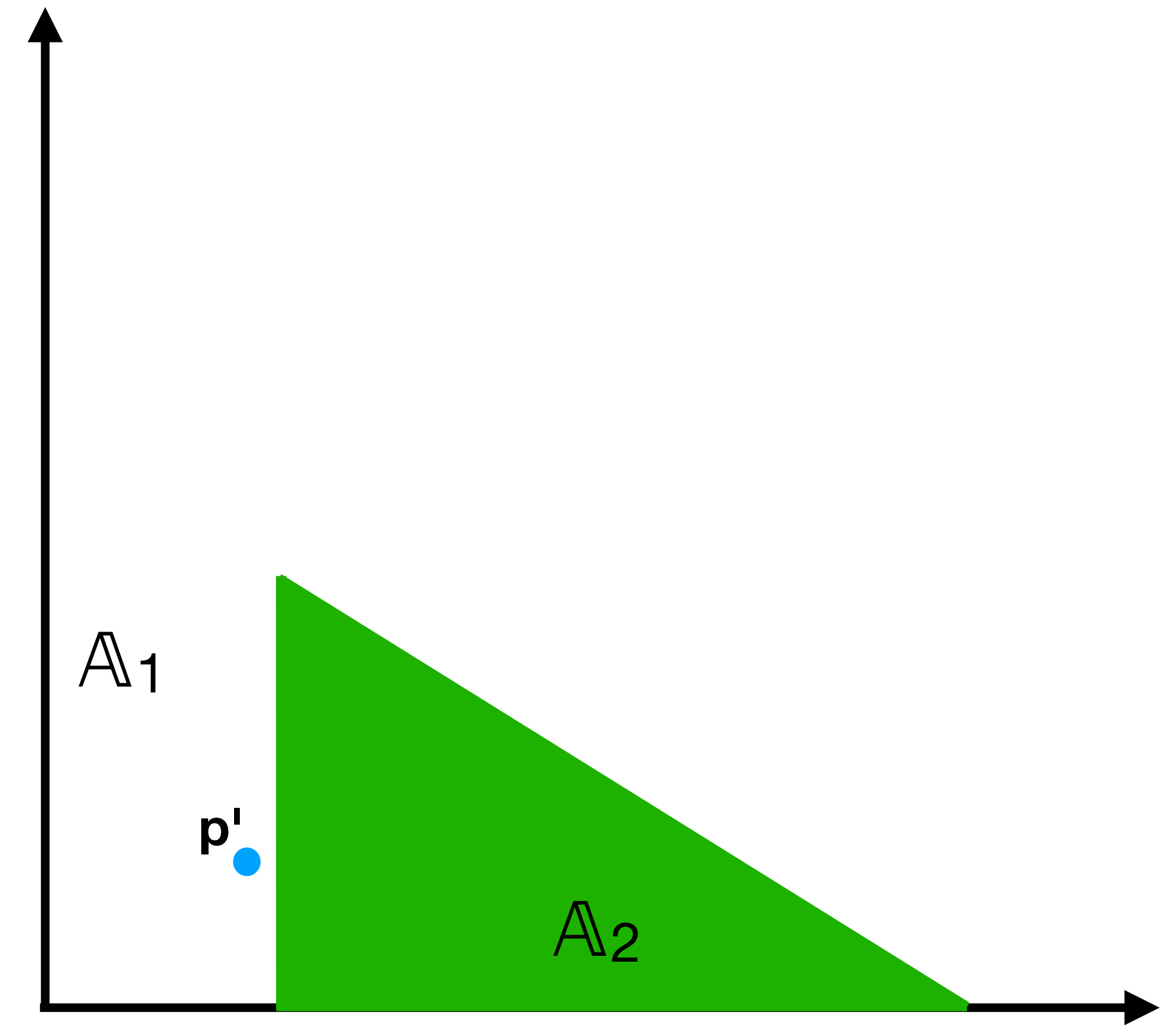
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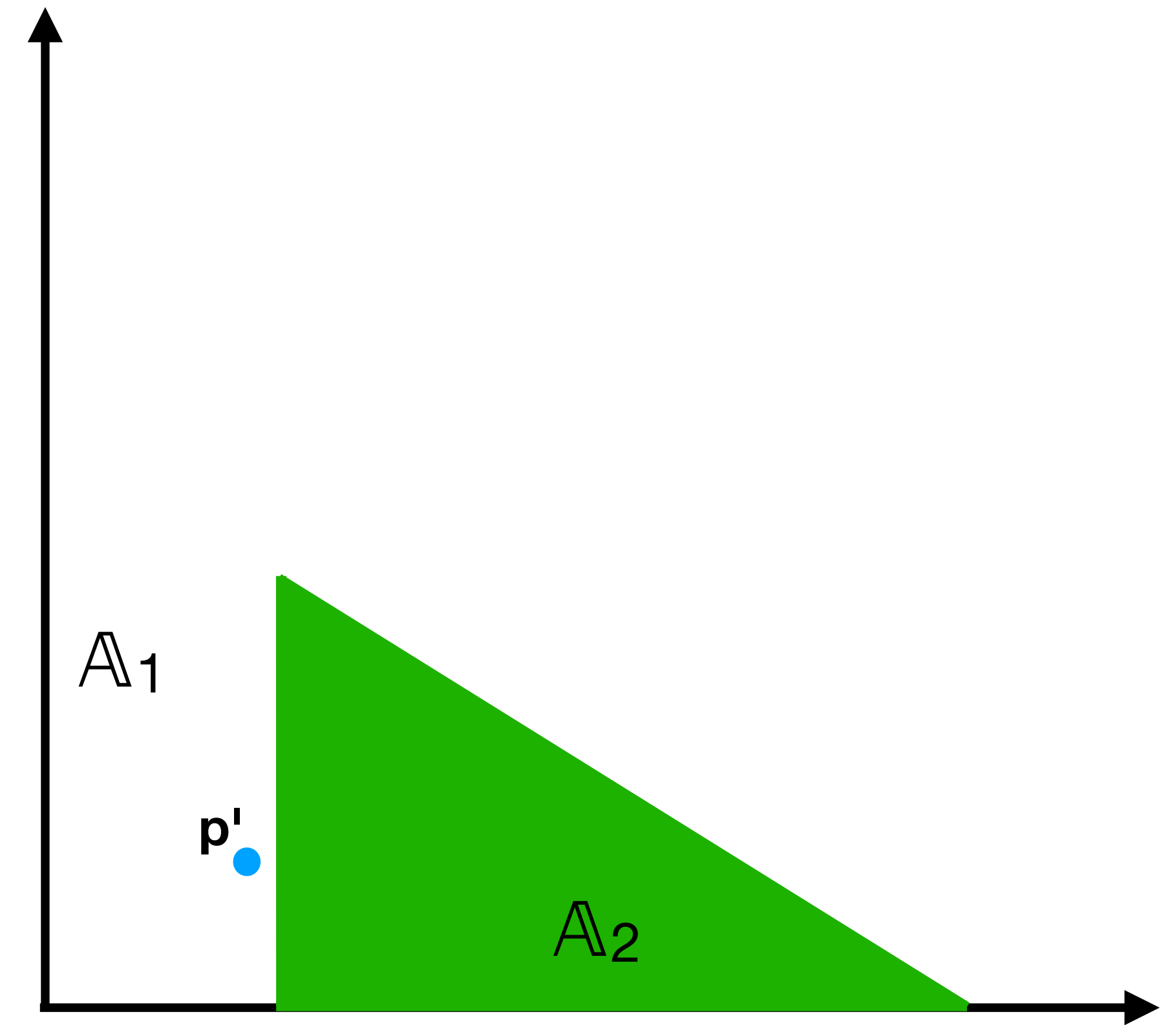


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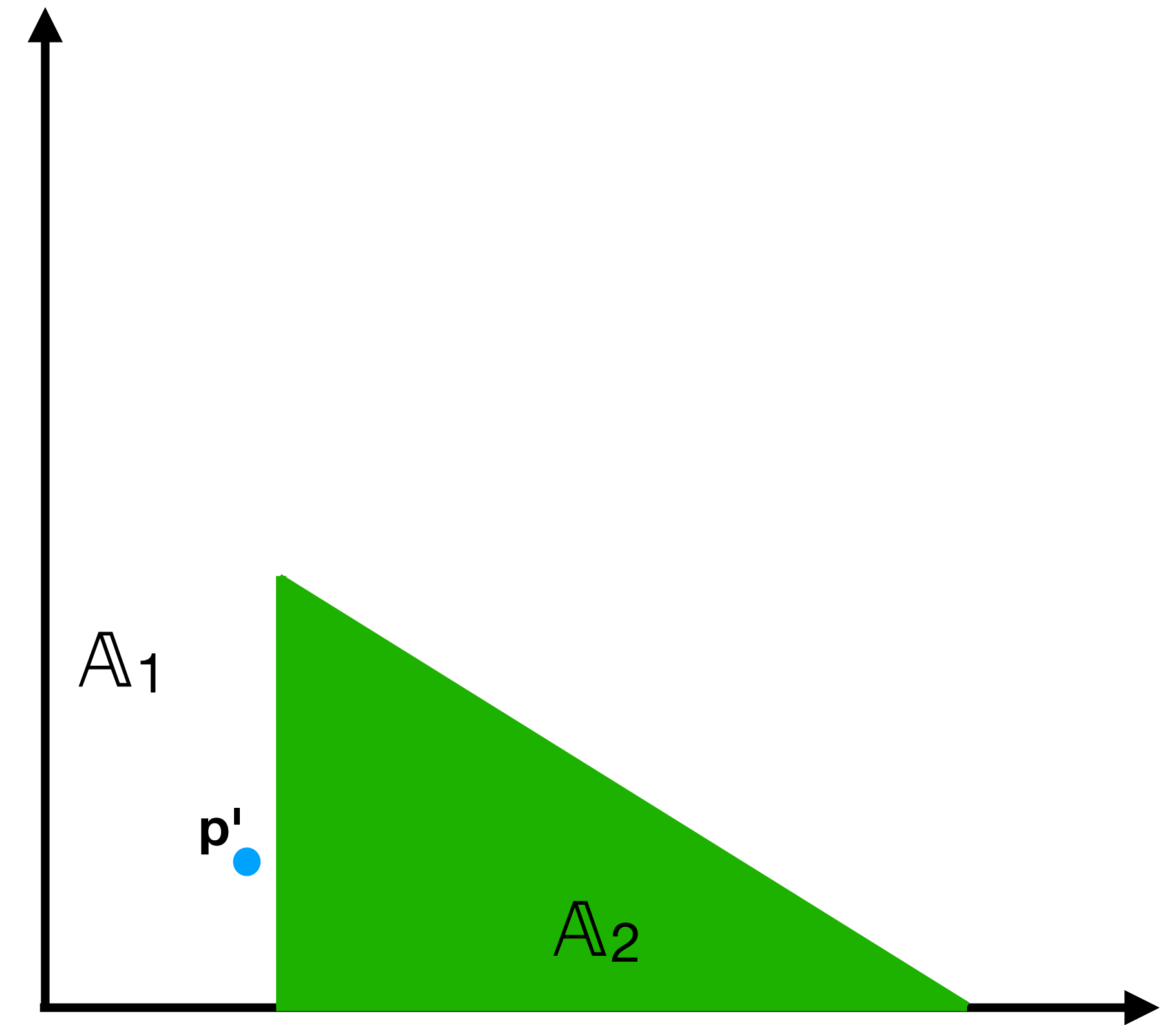
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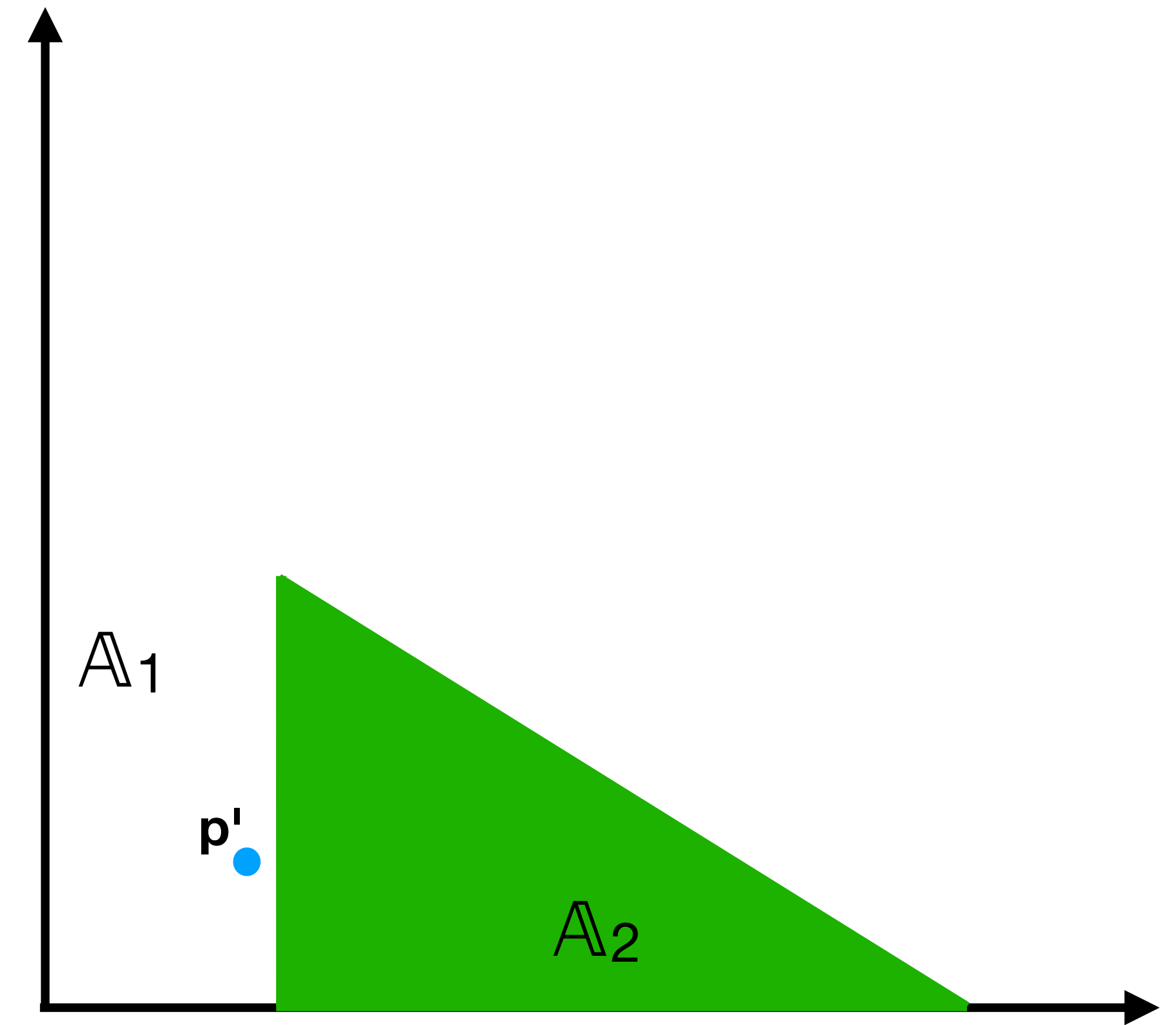
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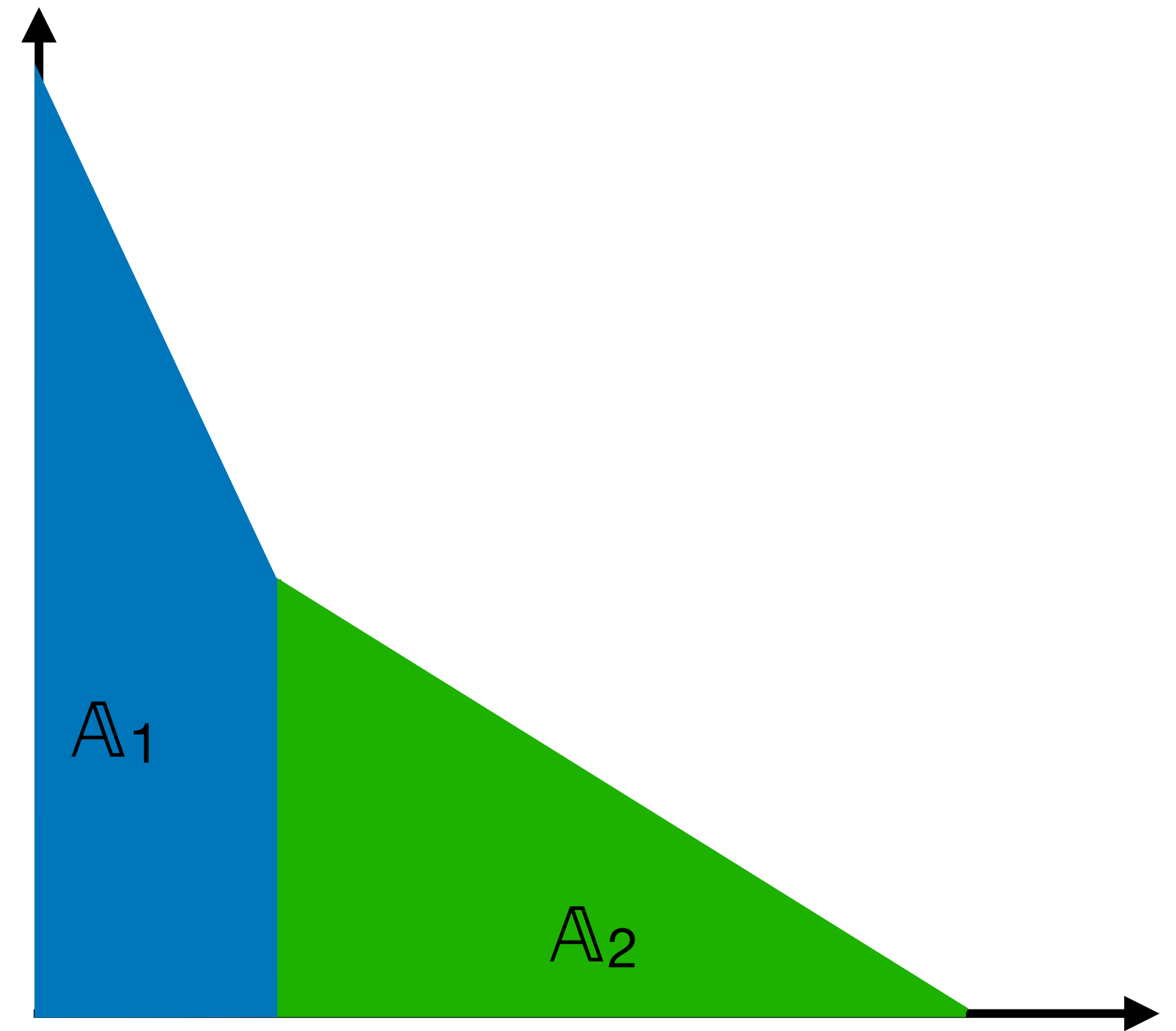
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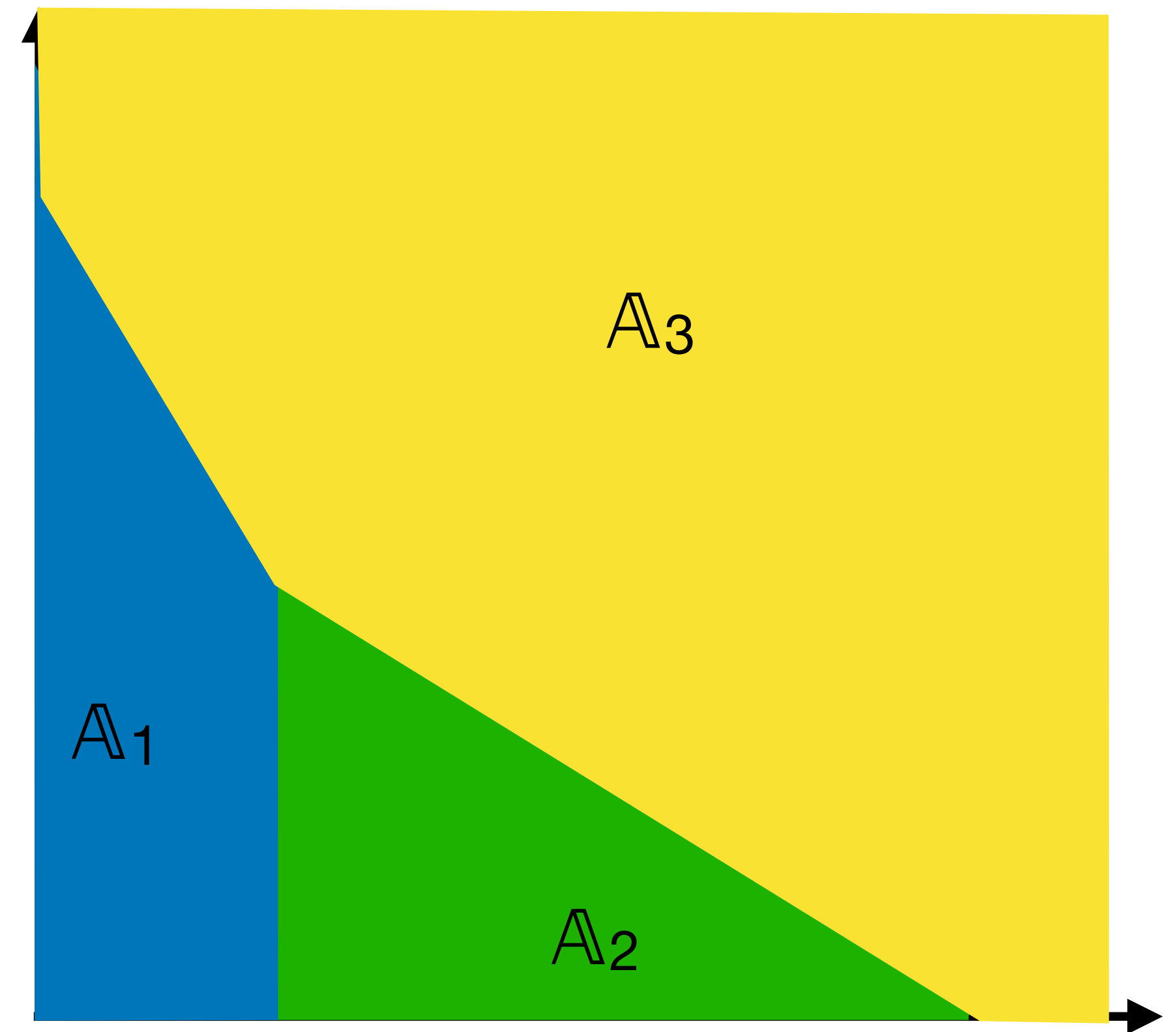
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When entire list is marked, the decomposition is complete.



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Any two alignments with the same triple are optimal at exactly the same points.

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Can be $O(m^4)$!

Decomposition Speed

Keep a list L of optimal alignments found along the way (the values of **mt**, **ms**, **id**, & **gp**)

Any time a new h is chosen, find the place where the line intersecting the optimal alignment and every alignment in L intersect h

Start the ray search at the closest to p rather than the boundary.

The size of L is bounded by sum of the number of vertices (V), edges (E), and polygons (R) in the decomposition.

$O(E)$ ray searches to find all edges, therefore $O(E(V+E+R)) = O(m^4)$ extra work to use L .

When doing a ray search, we only compute an alignment for points not in L , then they are added to L , so the number of alignments is bounded by the same size as L , therefore the alignment running time is $O((V+E+R)m^2) = O(m^4)$

Parametric sequence alignment

For a fixed input:

- there are $O(m^2)$ optimal alignments when two parameters are free
- the regions can be found by repeated ray-search

More free parameters

$$f_{\alpha,\beta,\gamma,\delta}(\mathbb{A}) = \alpha \cdot \mathbf{mt}_{\mathbb{A}} - \beta \cdot \mathbf{ms}_{\mathbb{A}} - \gamma \cdot \mathbf{id}_{\mathbb{A}} - \delta \cdot \mathbf{gp}_{\mathbb{A}}$$

- $\mathbf{mt}_{\mathbb{A}}$ -- number of columns where both characters match
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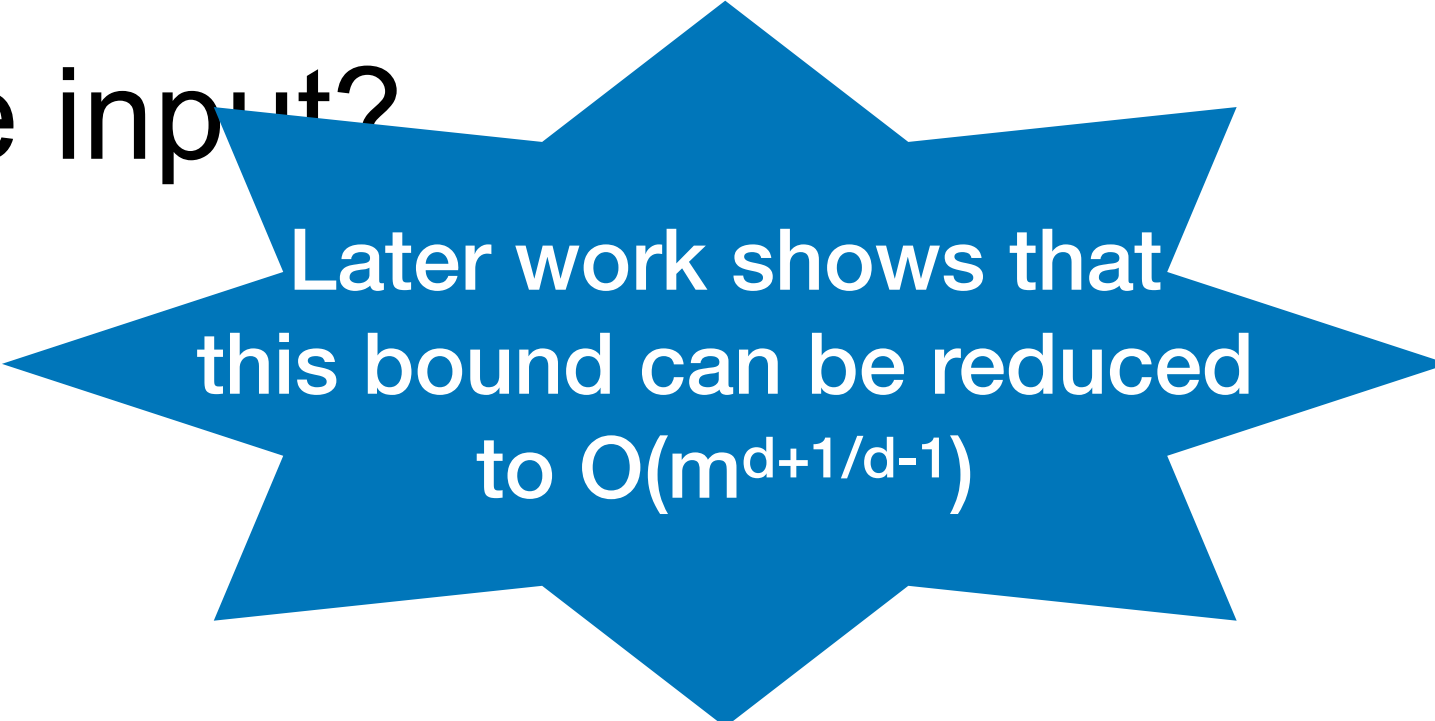
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Later work shows that
this bound can be reduced
to $O(m^{d+1/d-1})$