

# WHIR



## Proximity testing for Reed–Solomon+

Gal Arnon



Giacomo Fenzi

**EPFL**

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**EPFL**

Eylon Yogev



# Motivation

# **SNARKs**

**Succinct Non-interactive Arguments of Knowledge**

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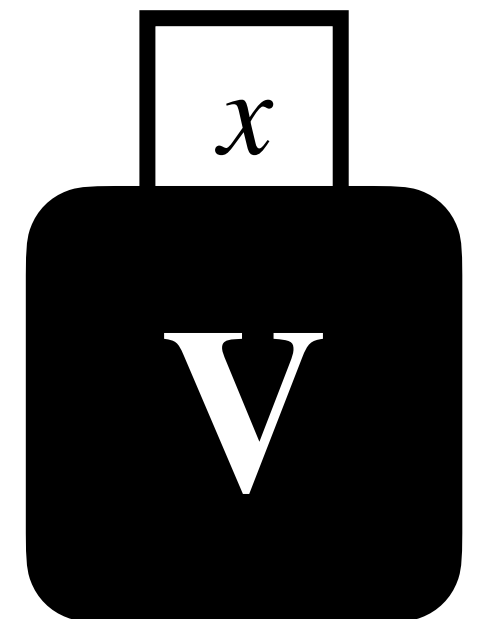
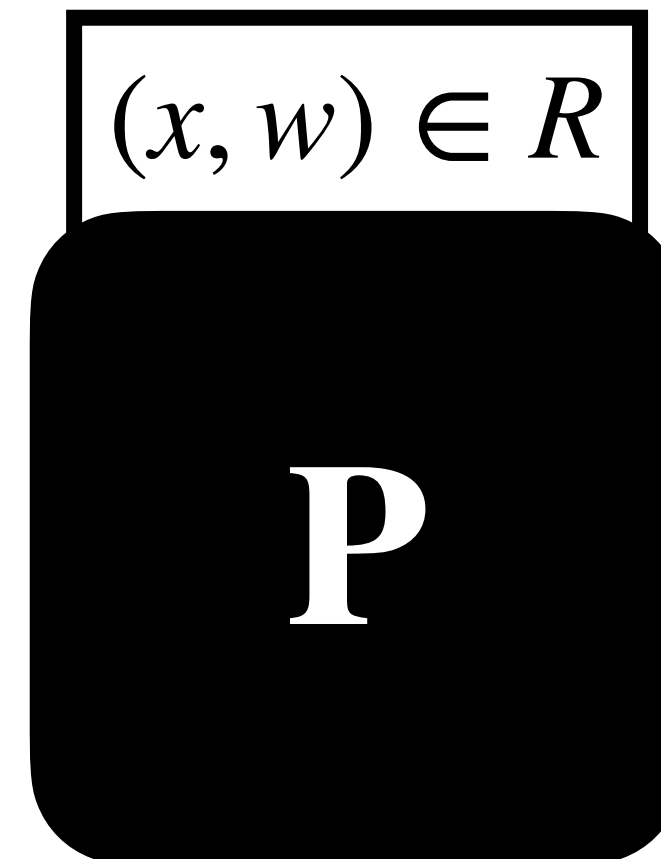
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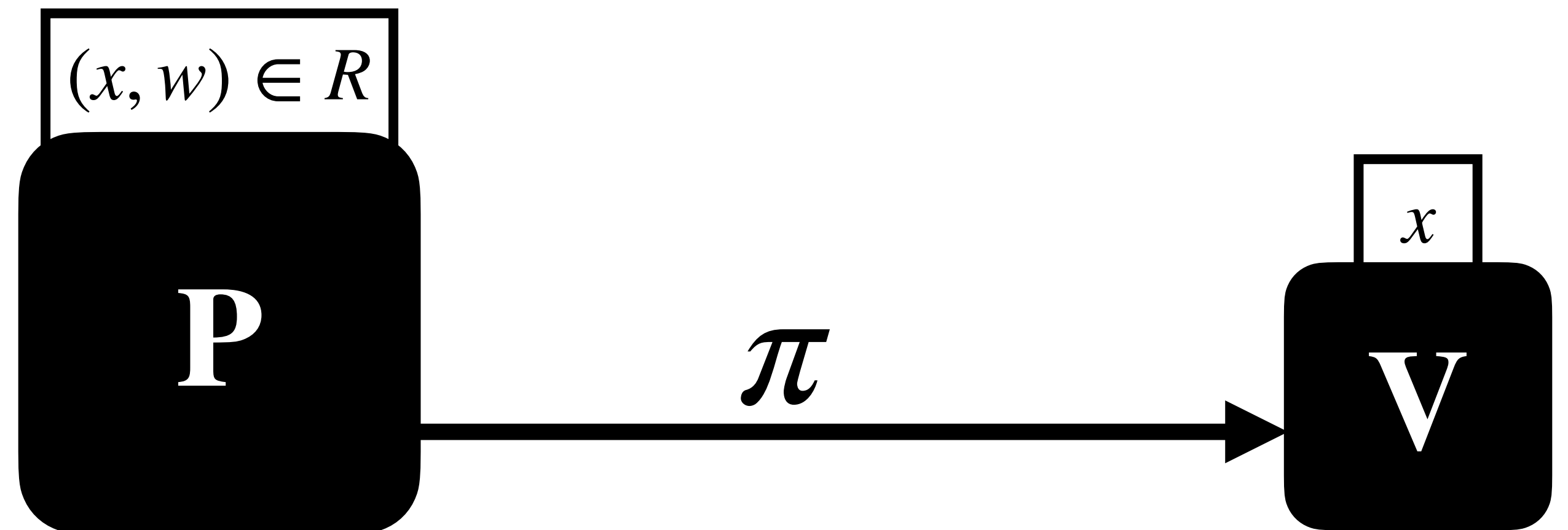
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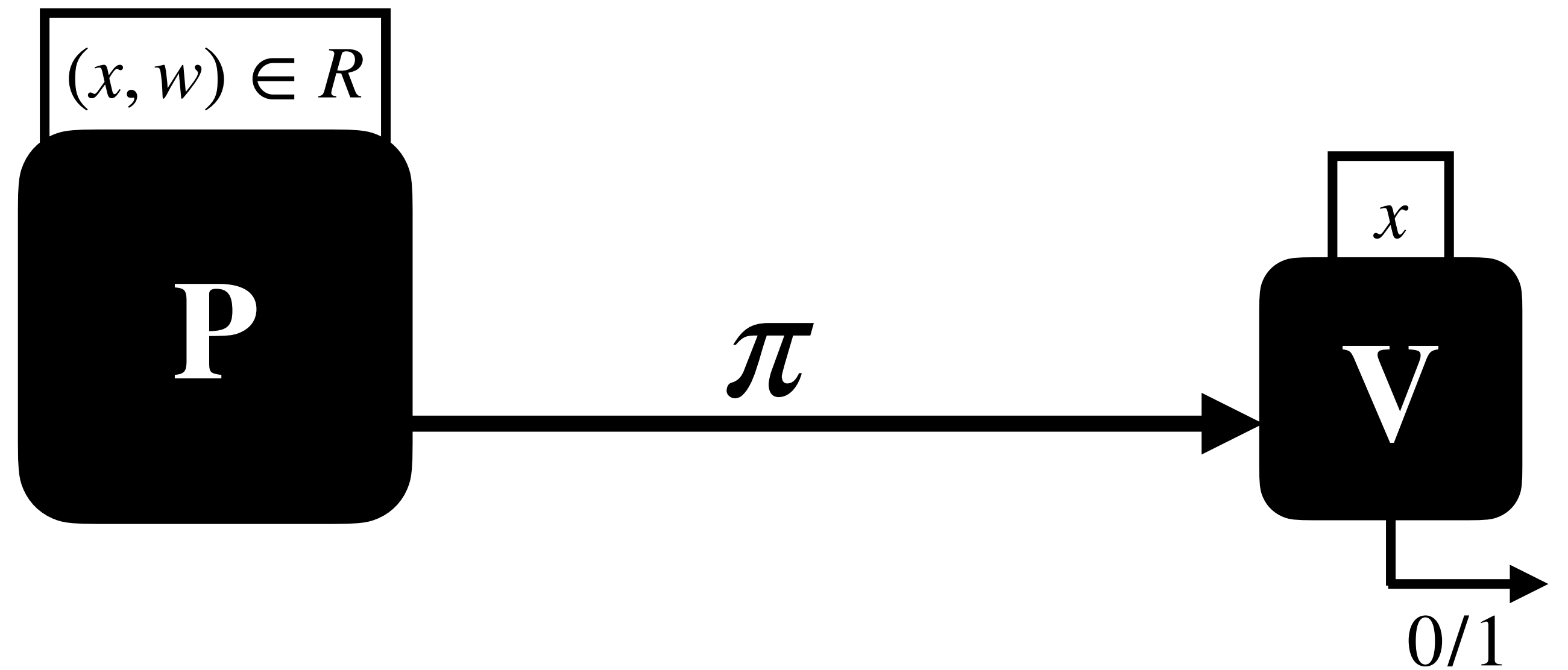
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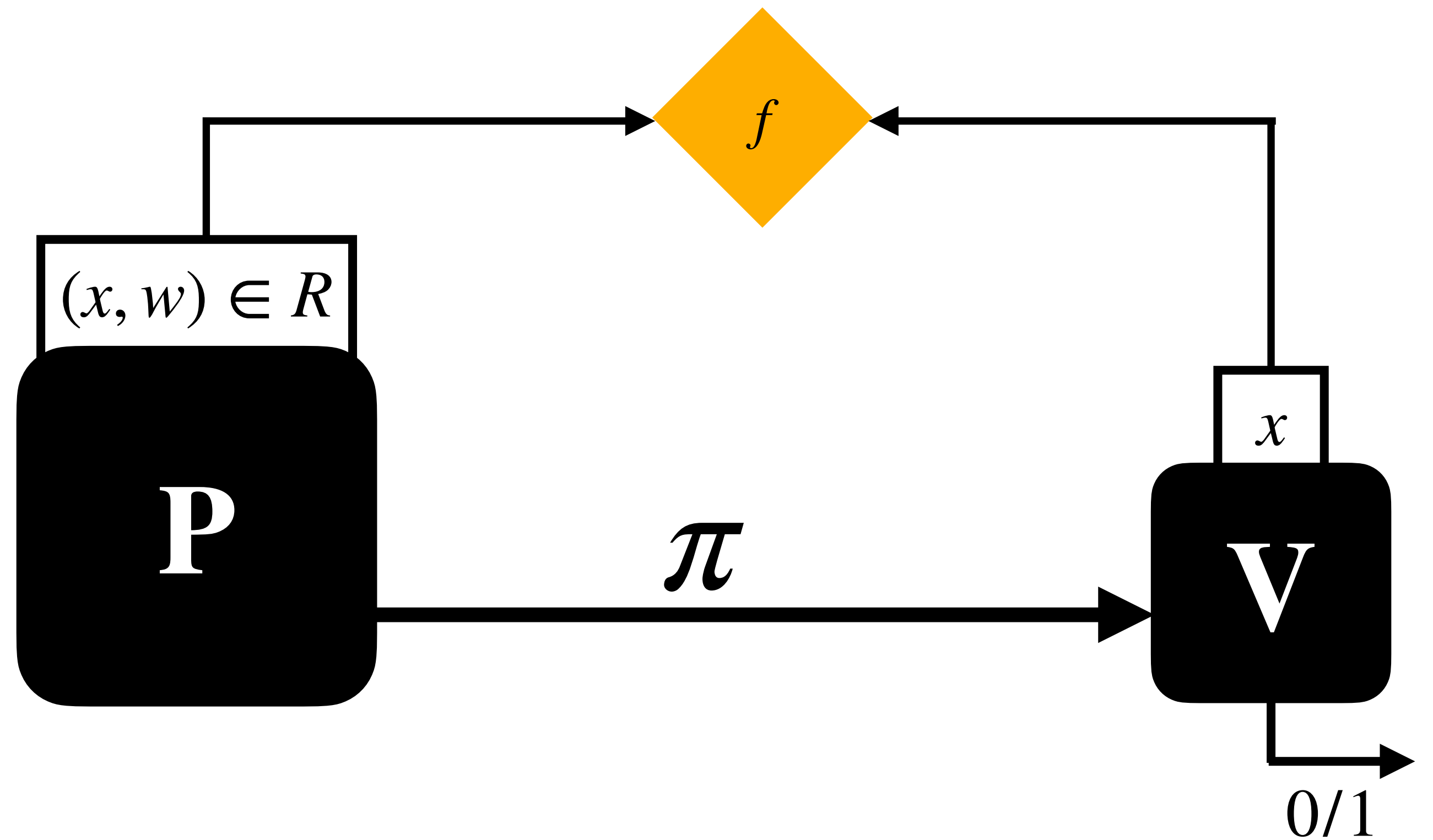




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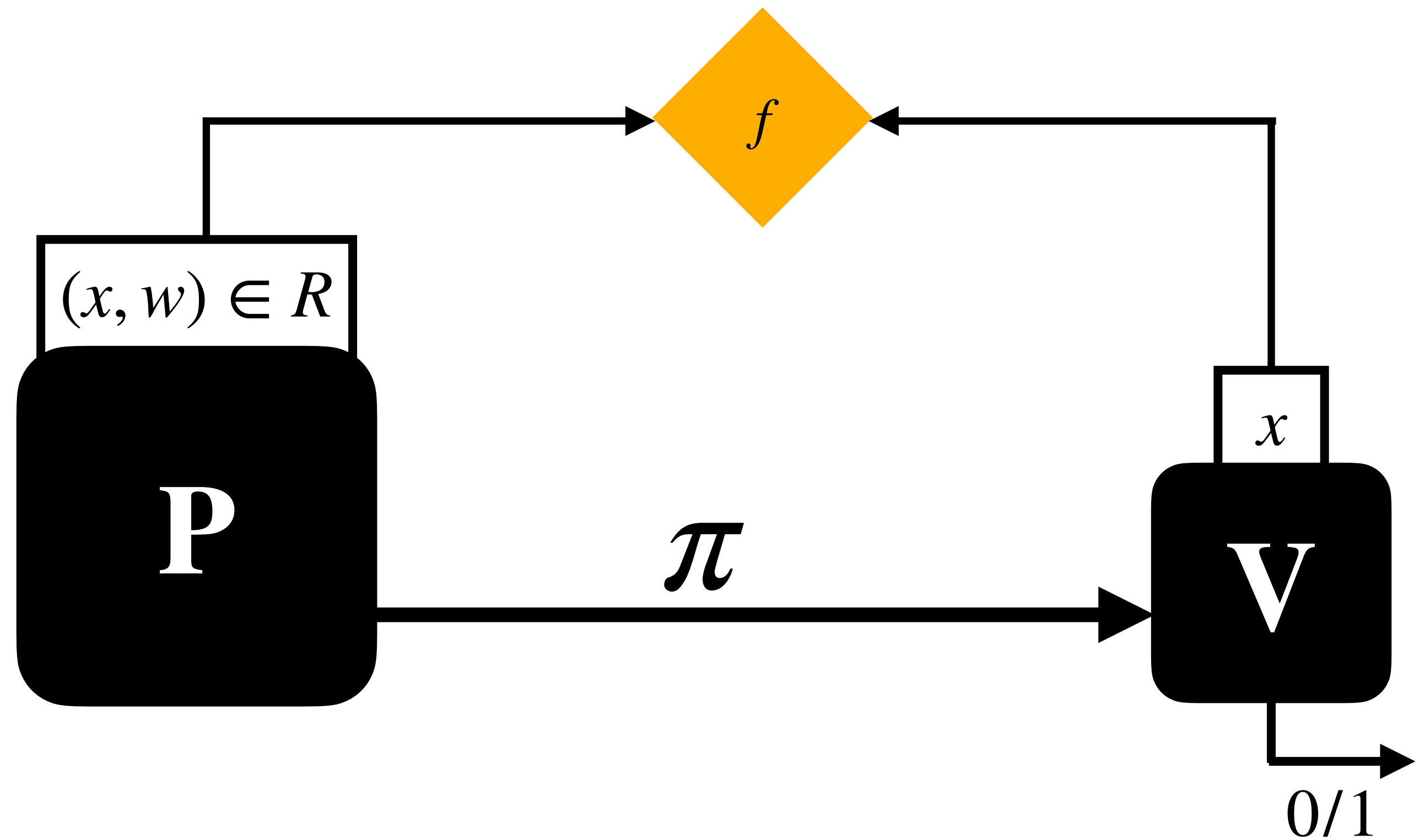


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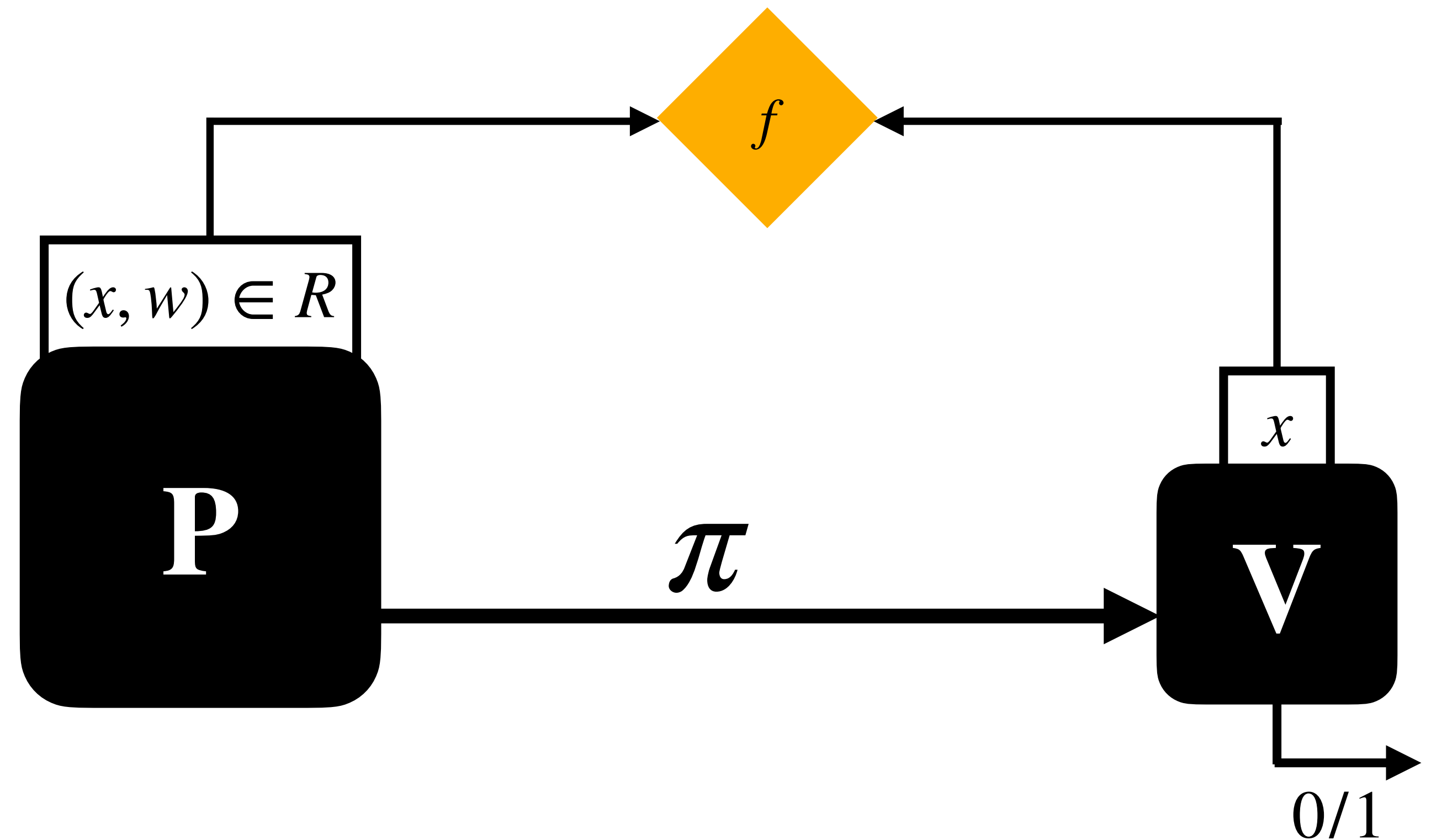
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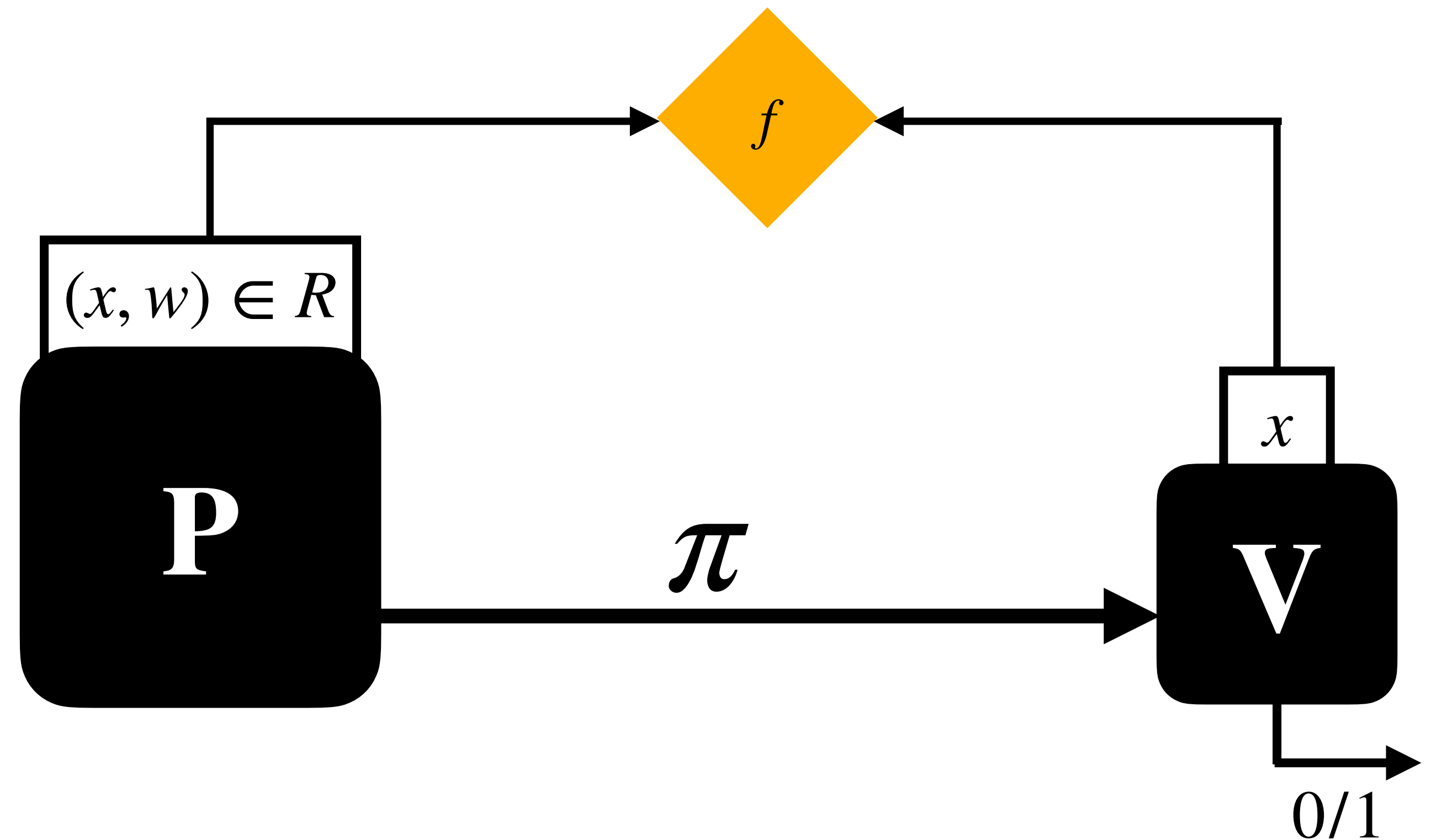
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- **Need\*** to add a **random oracle**.

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- **Today:** we limit ourselves to **pure ROM SNARKs**

- Will call these **hash-based SNARKs**.



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And many more...

# Constructing SNARKs

[BCS16] Construction

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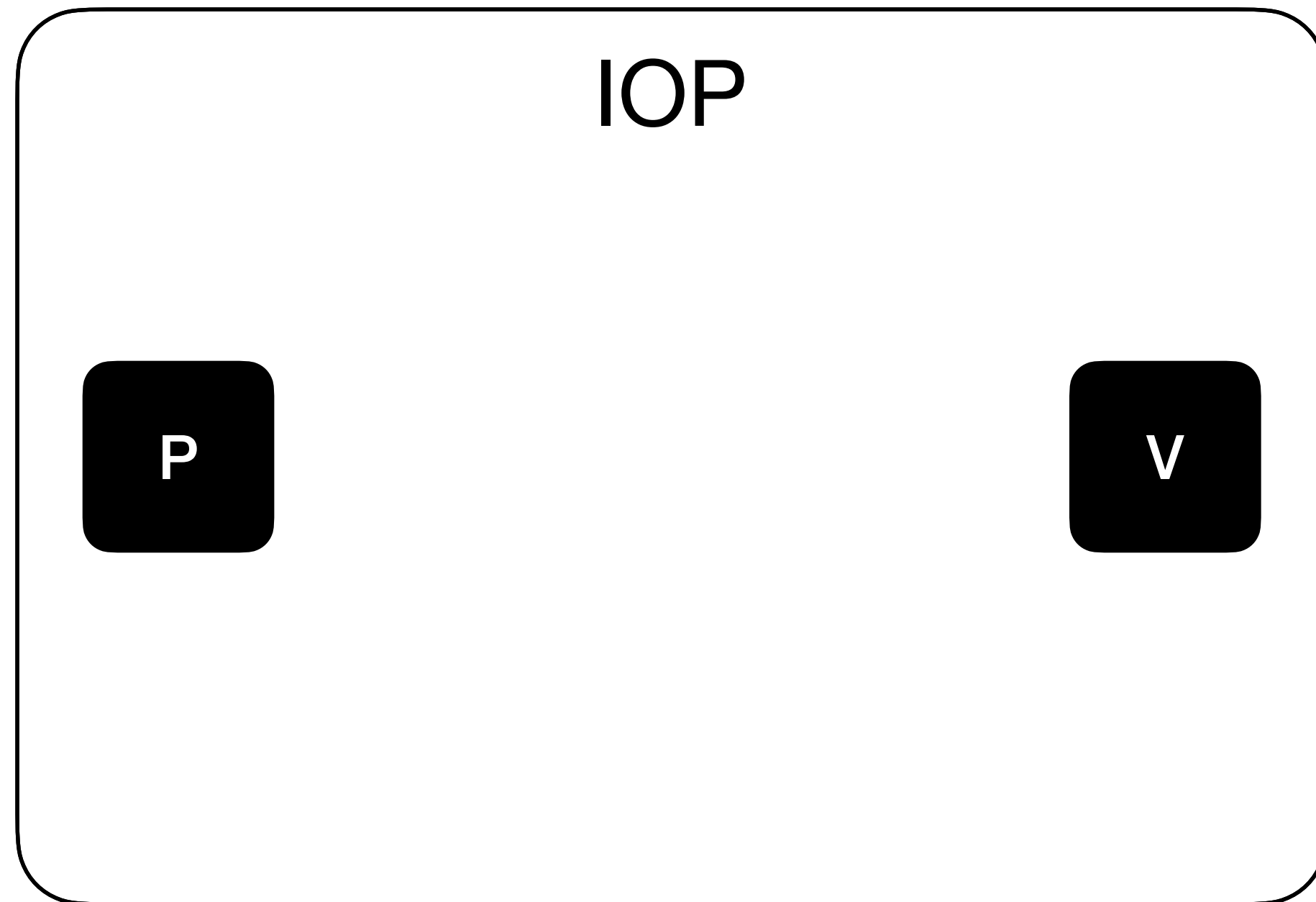
[BCS16] Construction



IOP

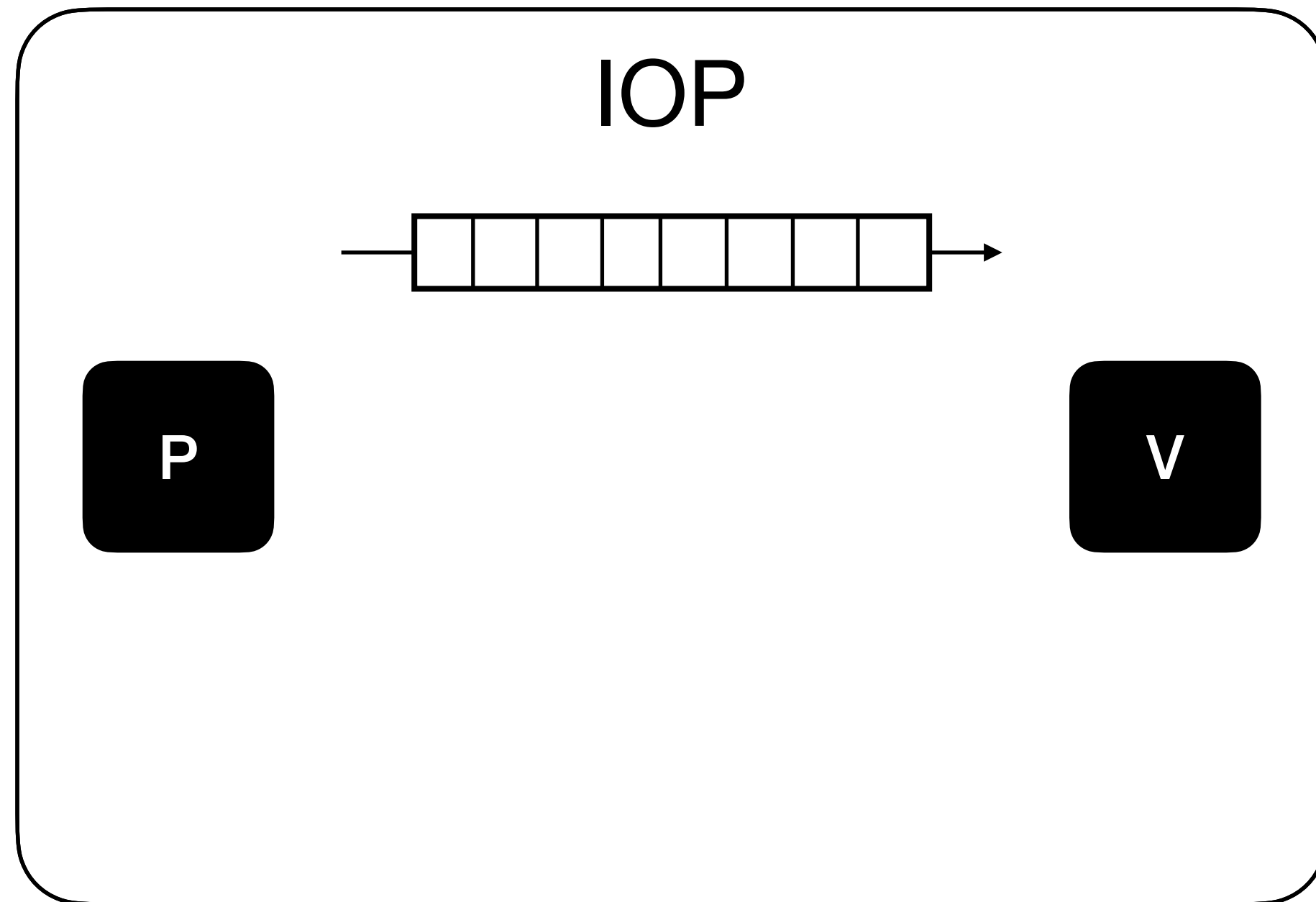
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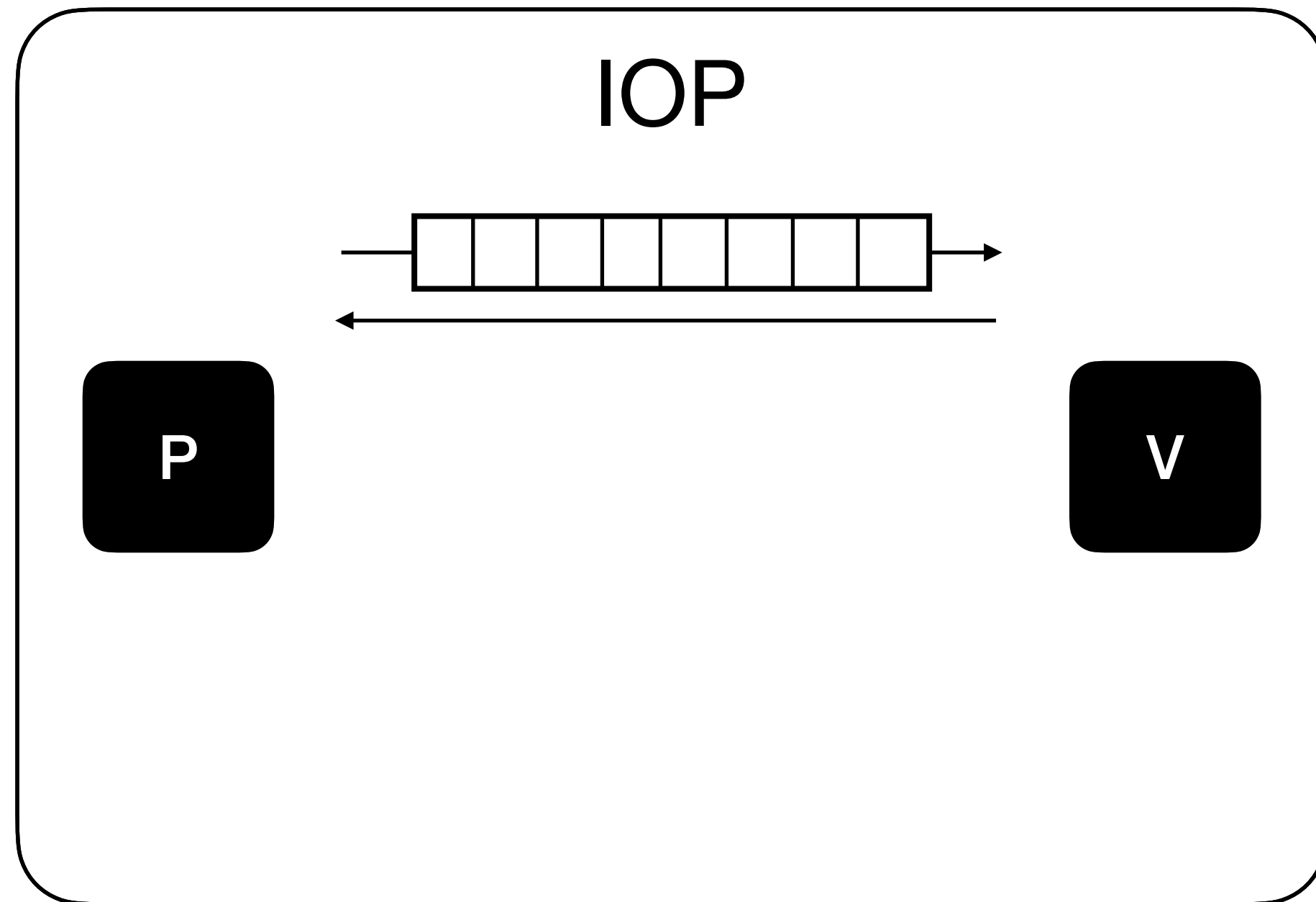
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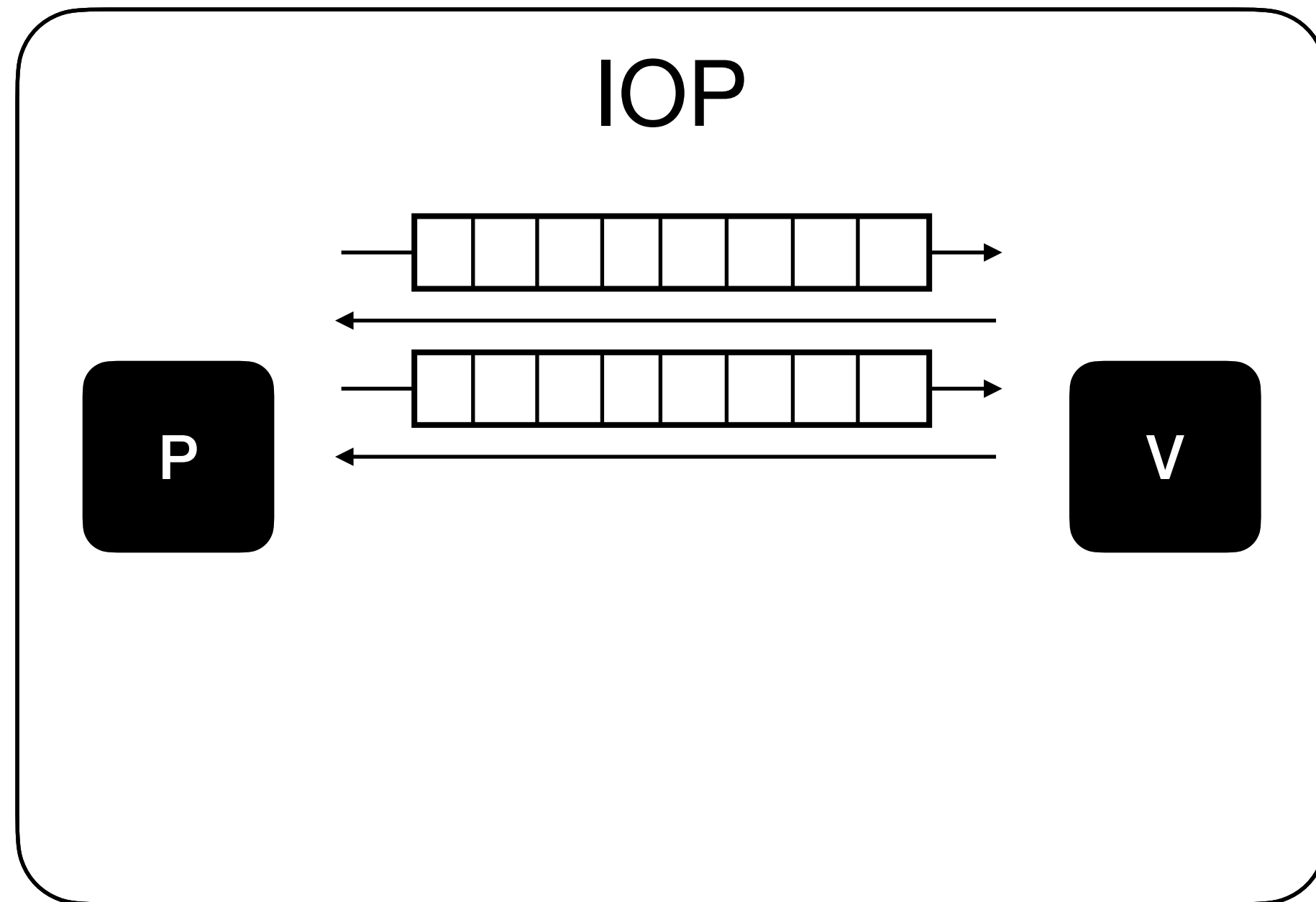
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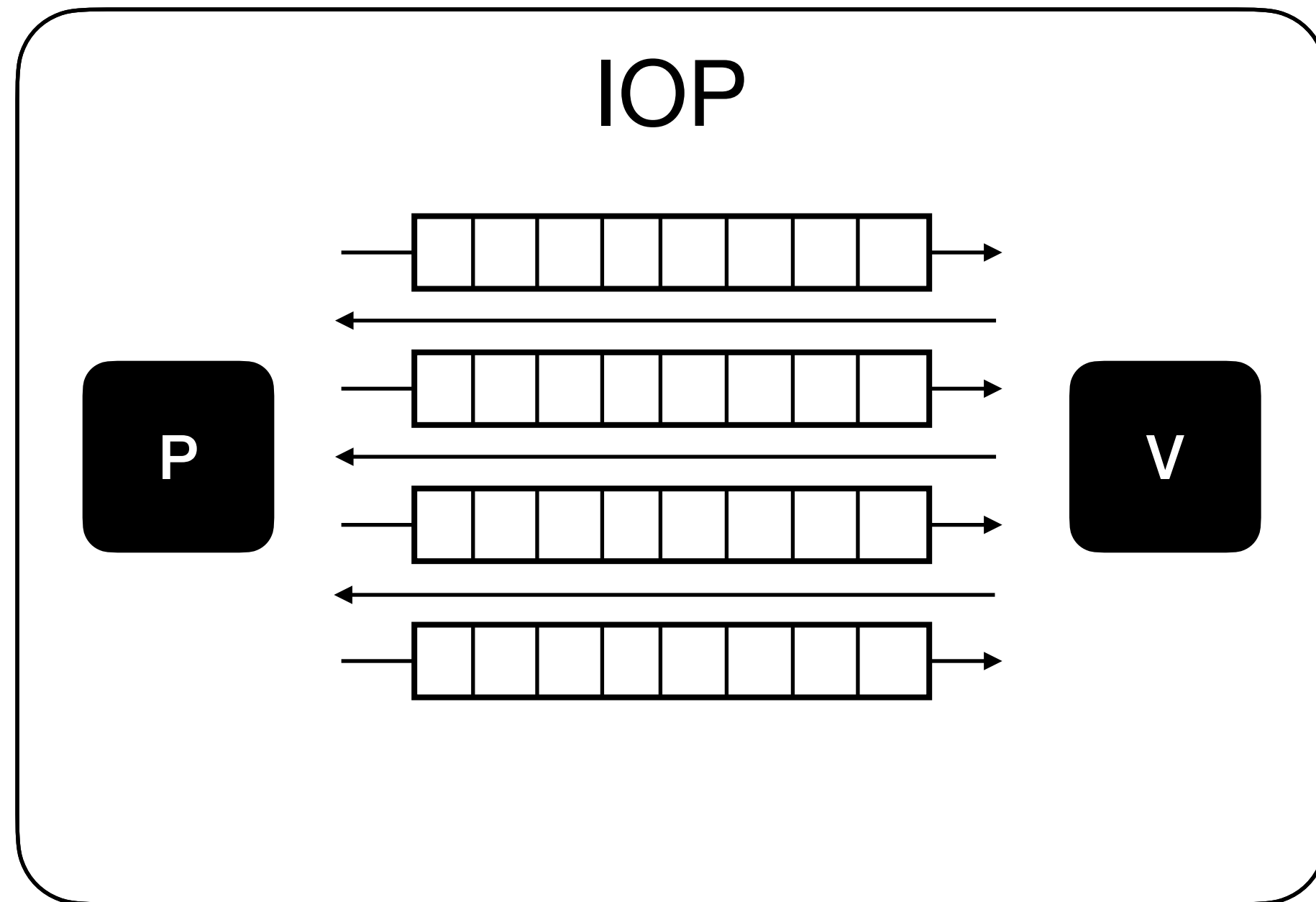
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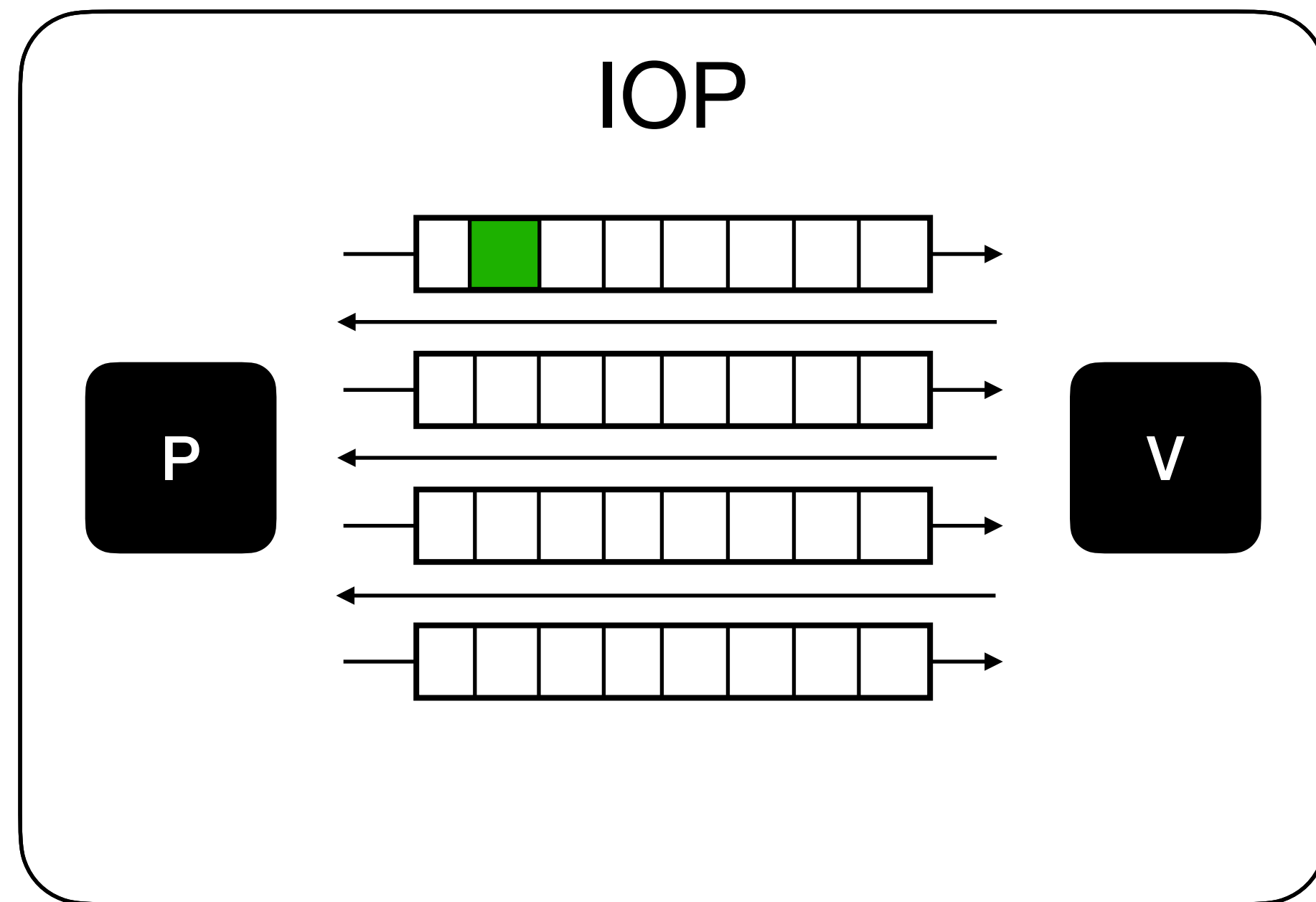
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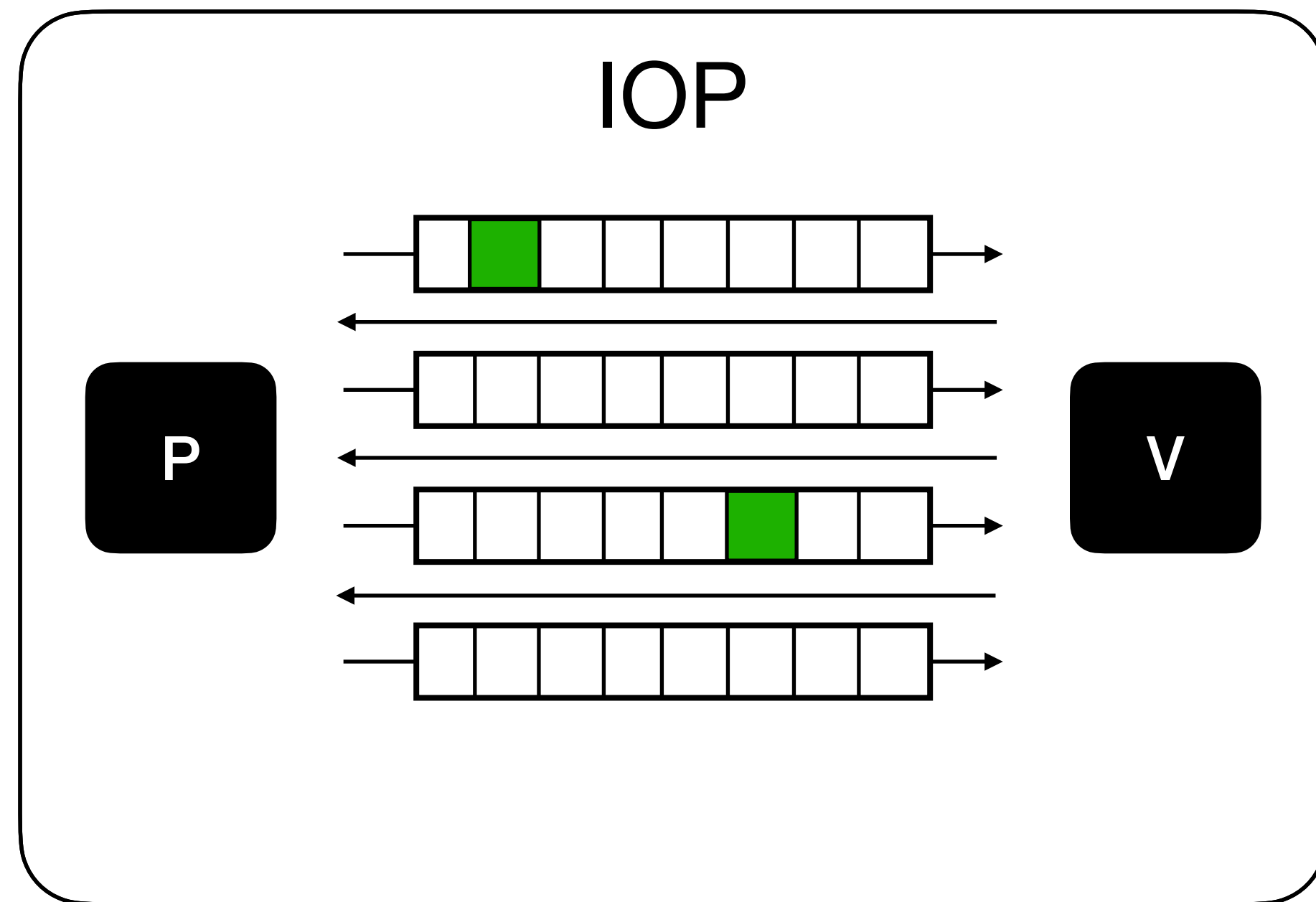
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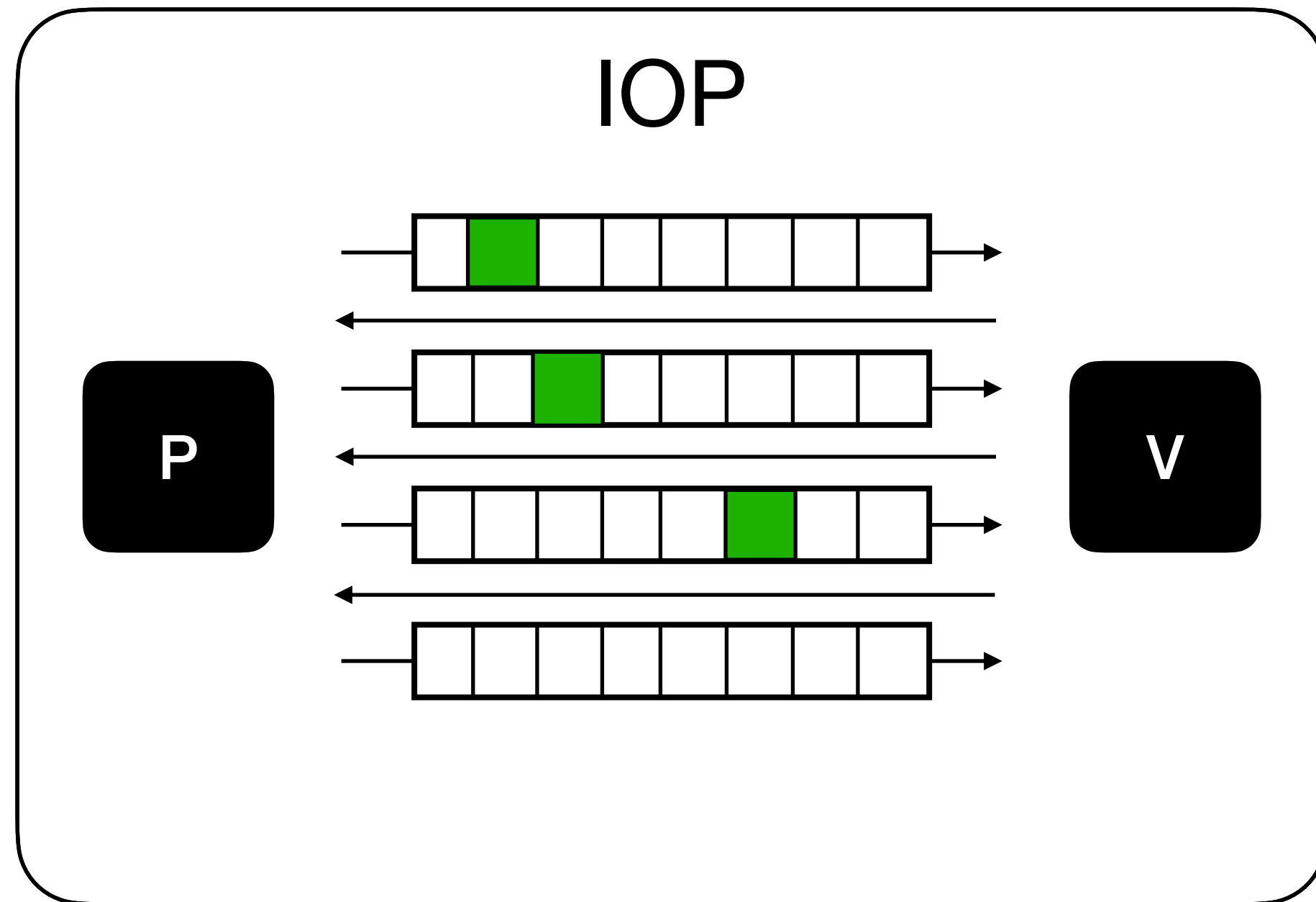
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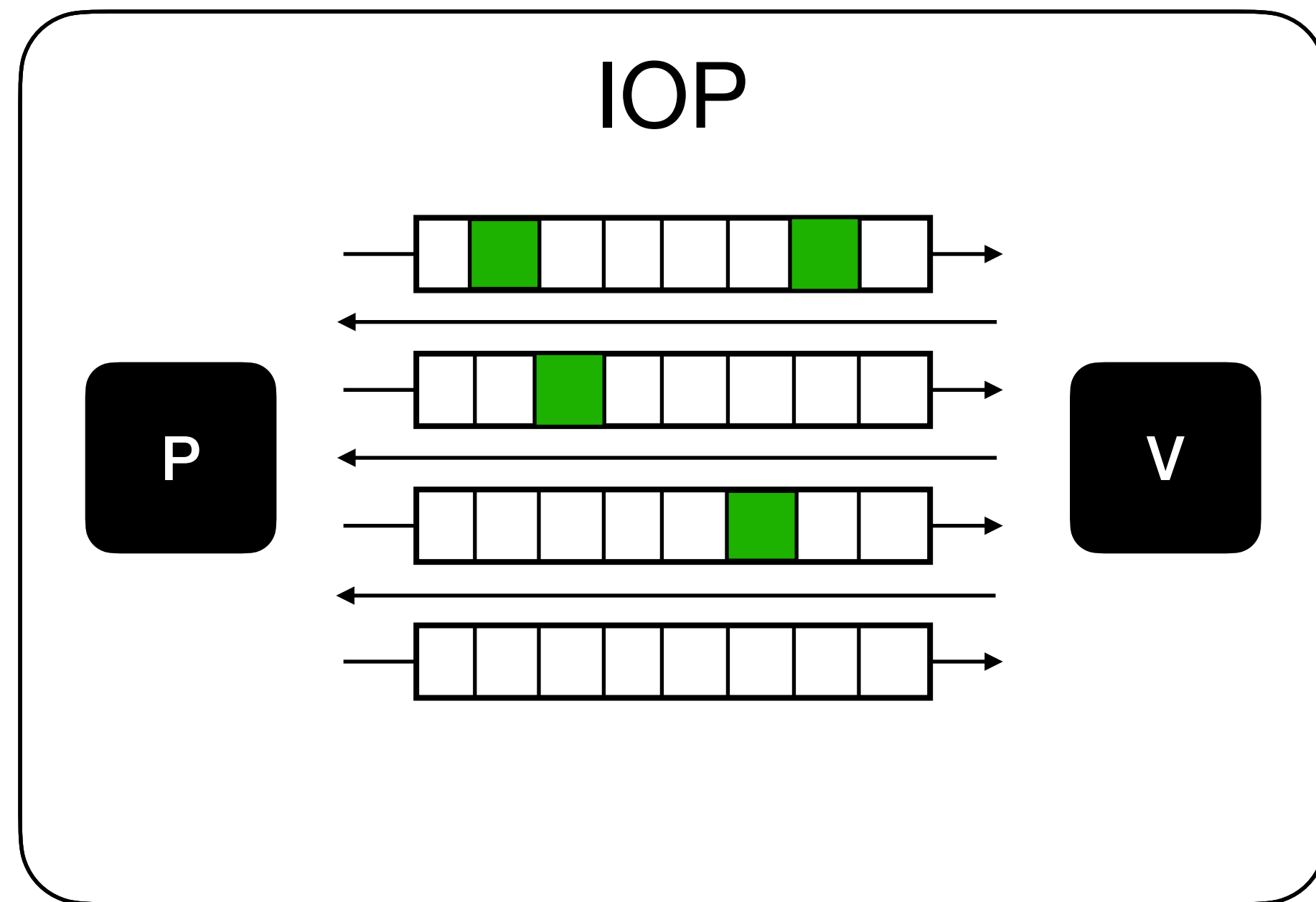
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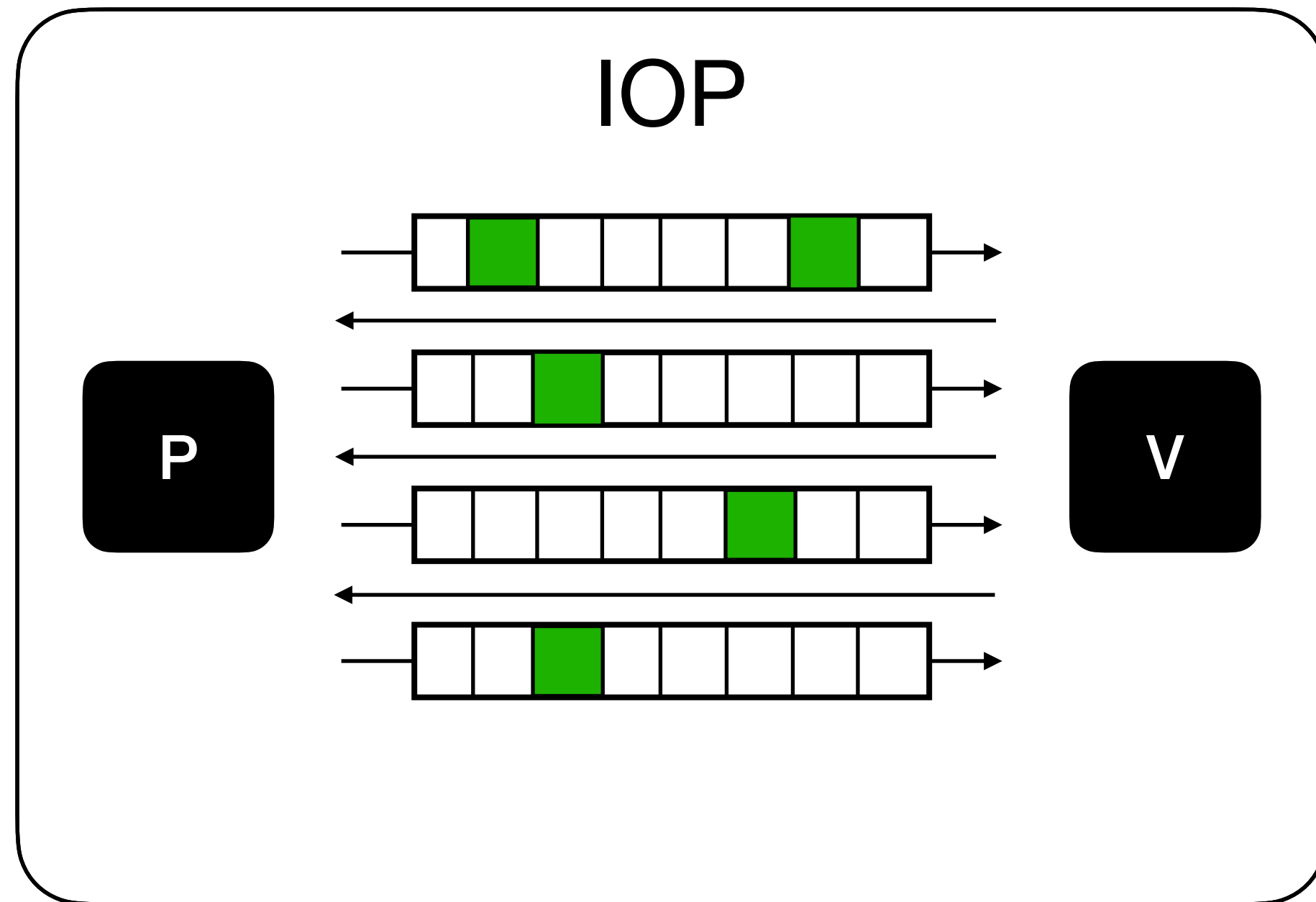
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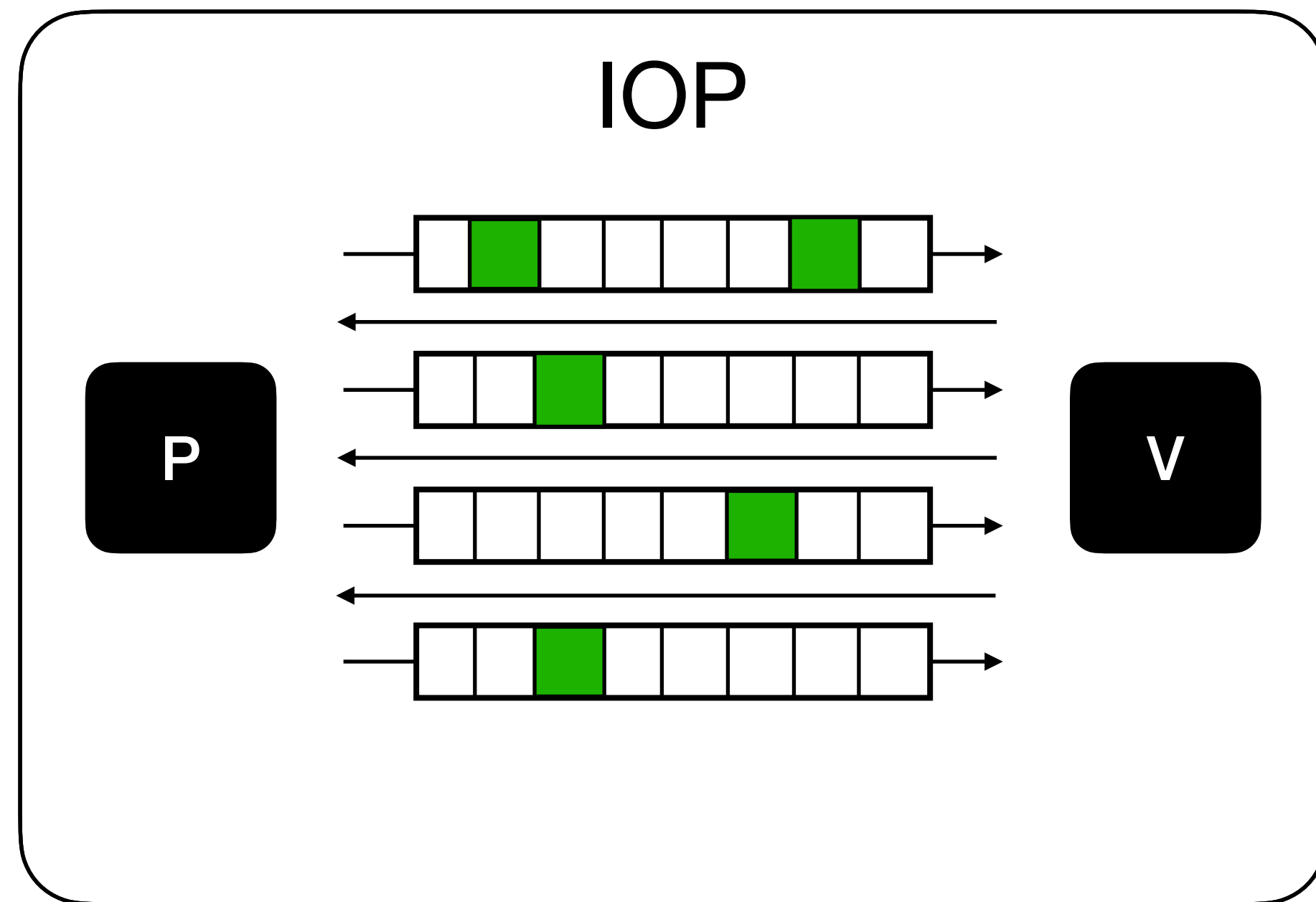
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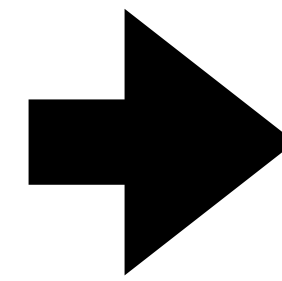


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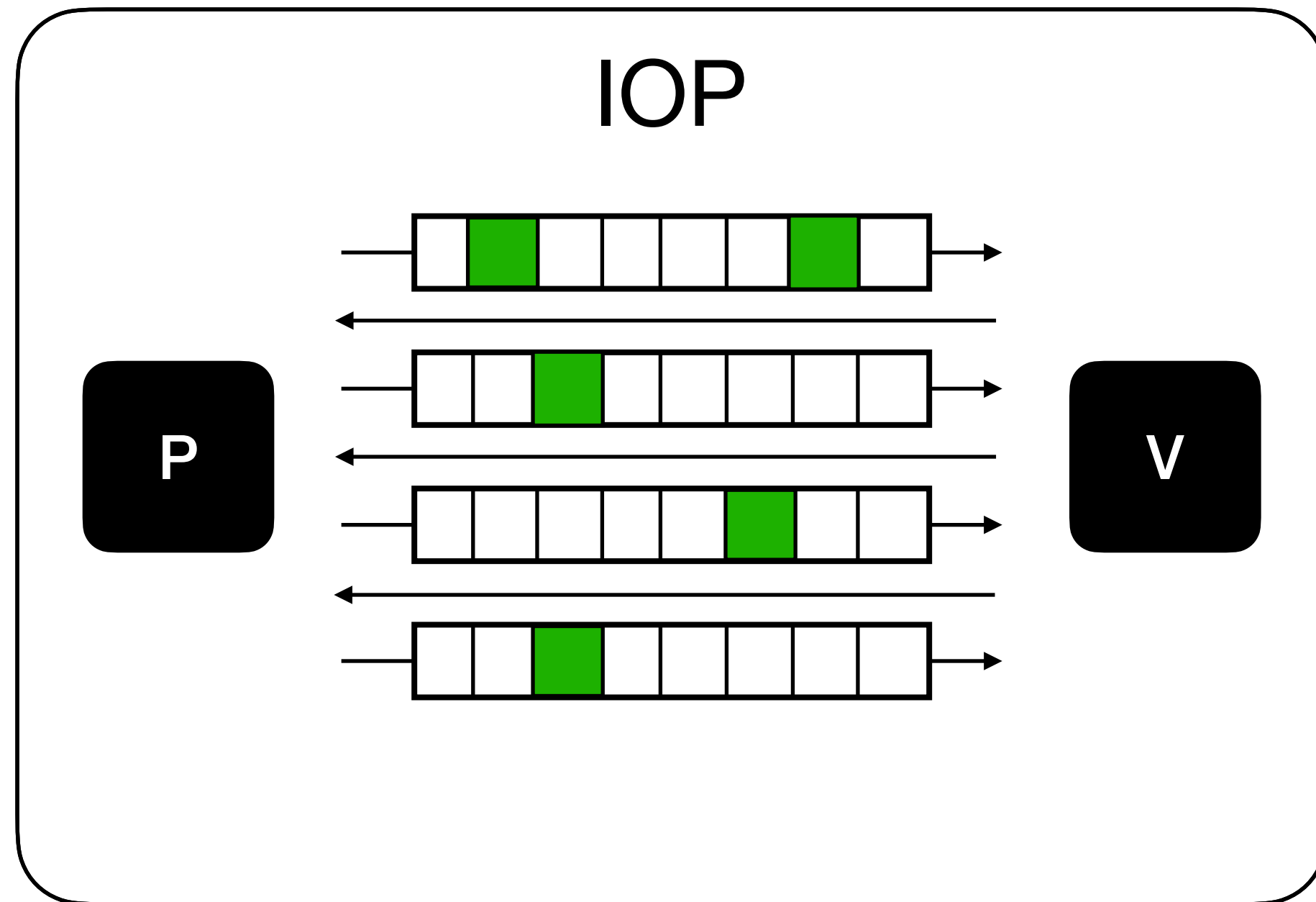
BCS



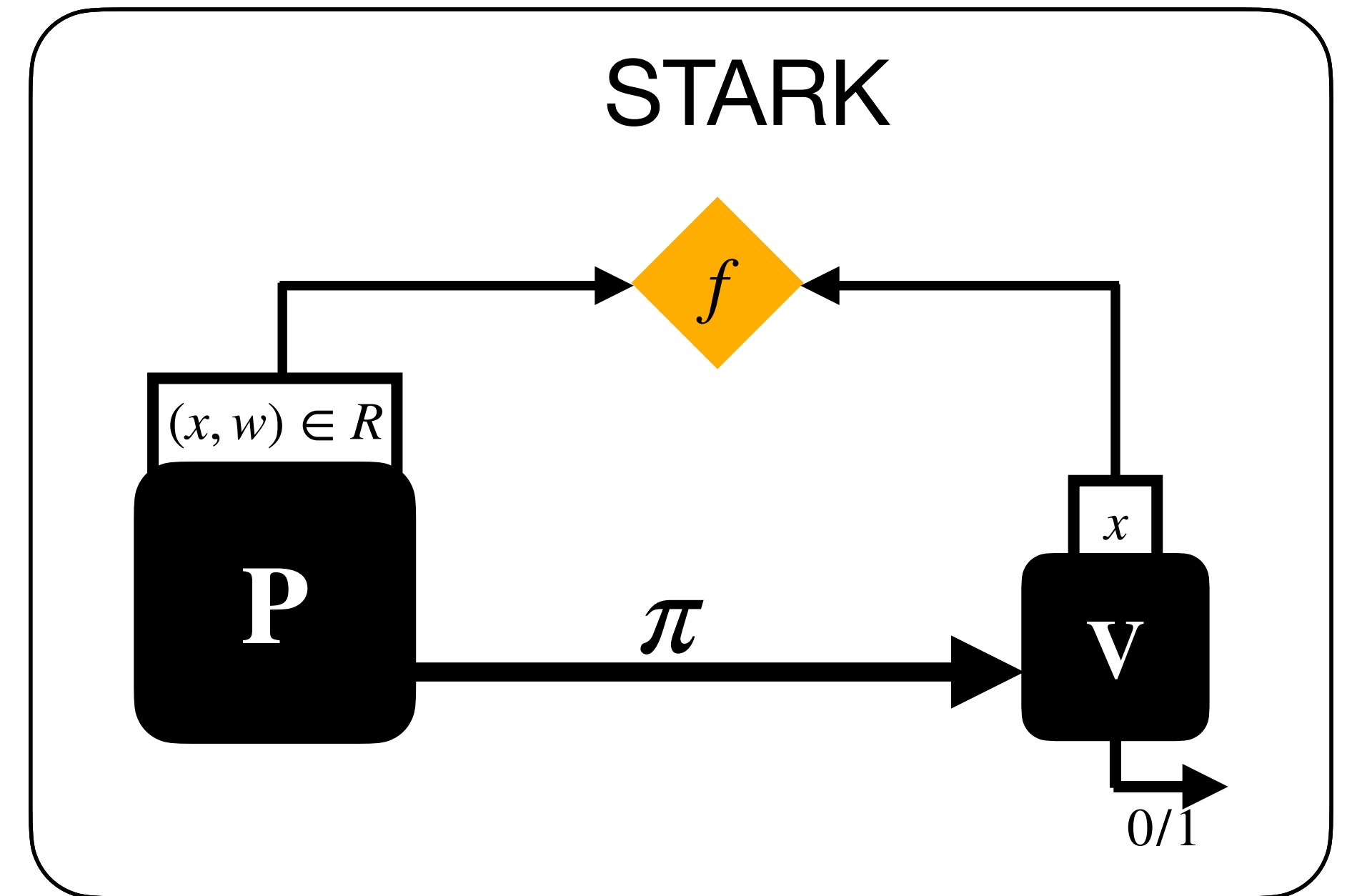
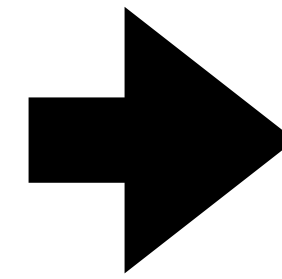


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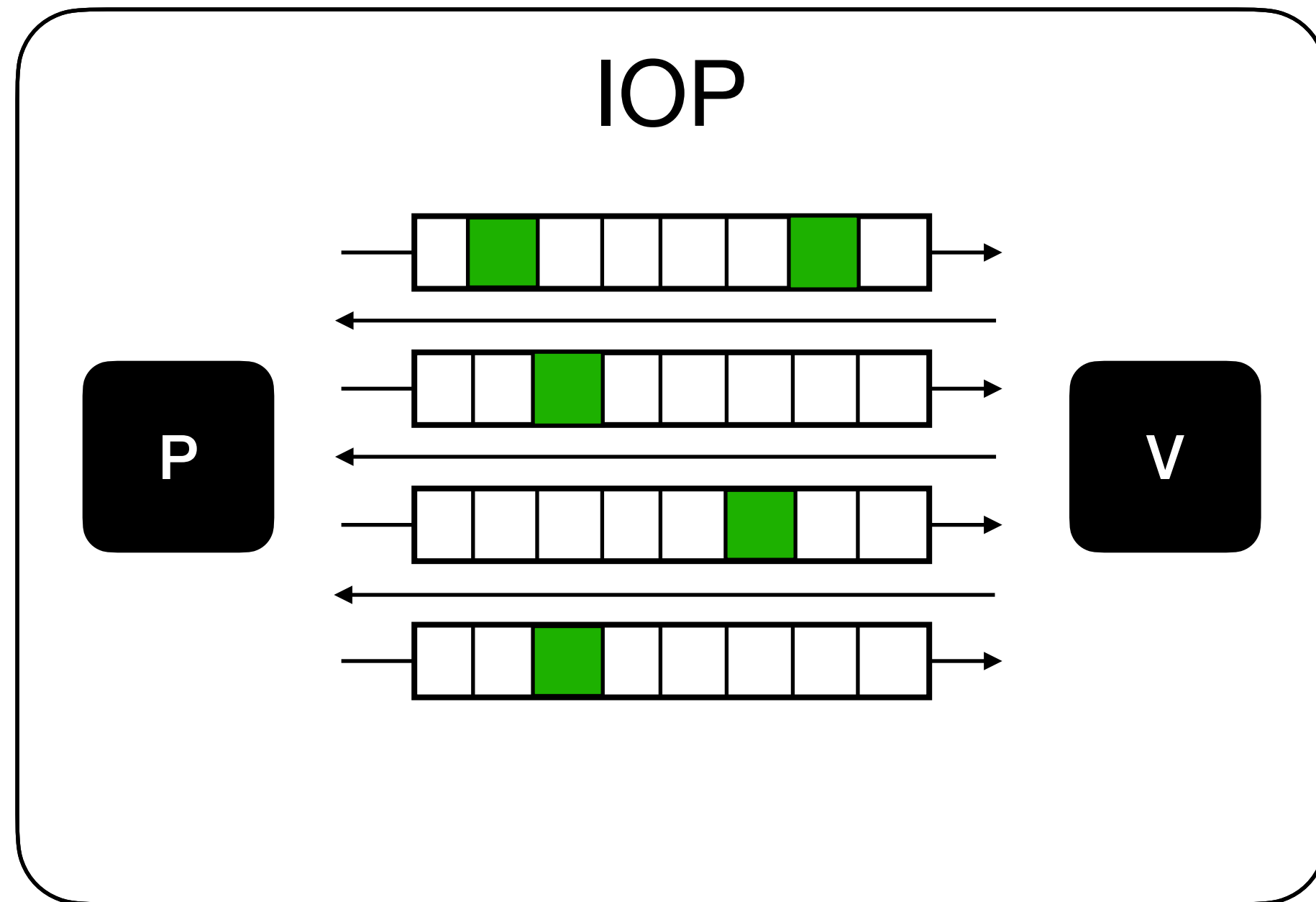


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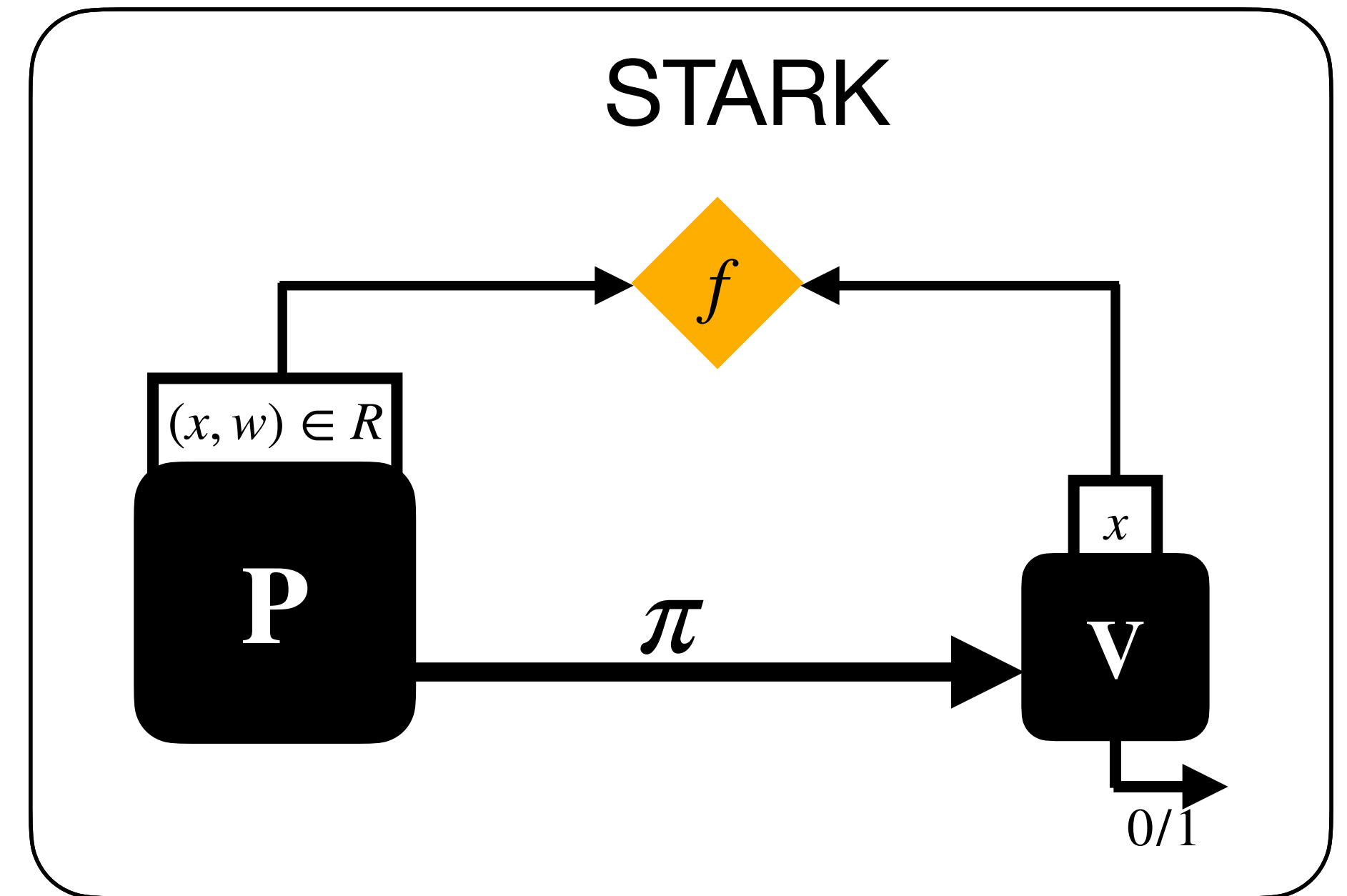
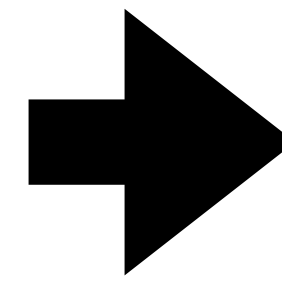


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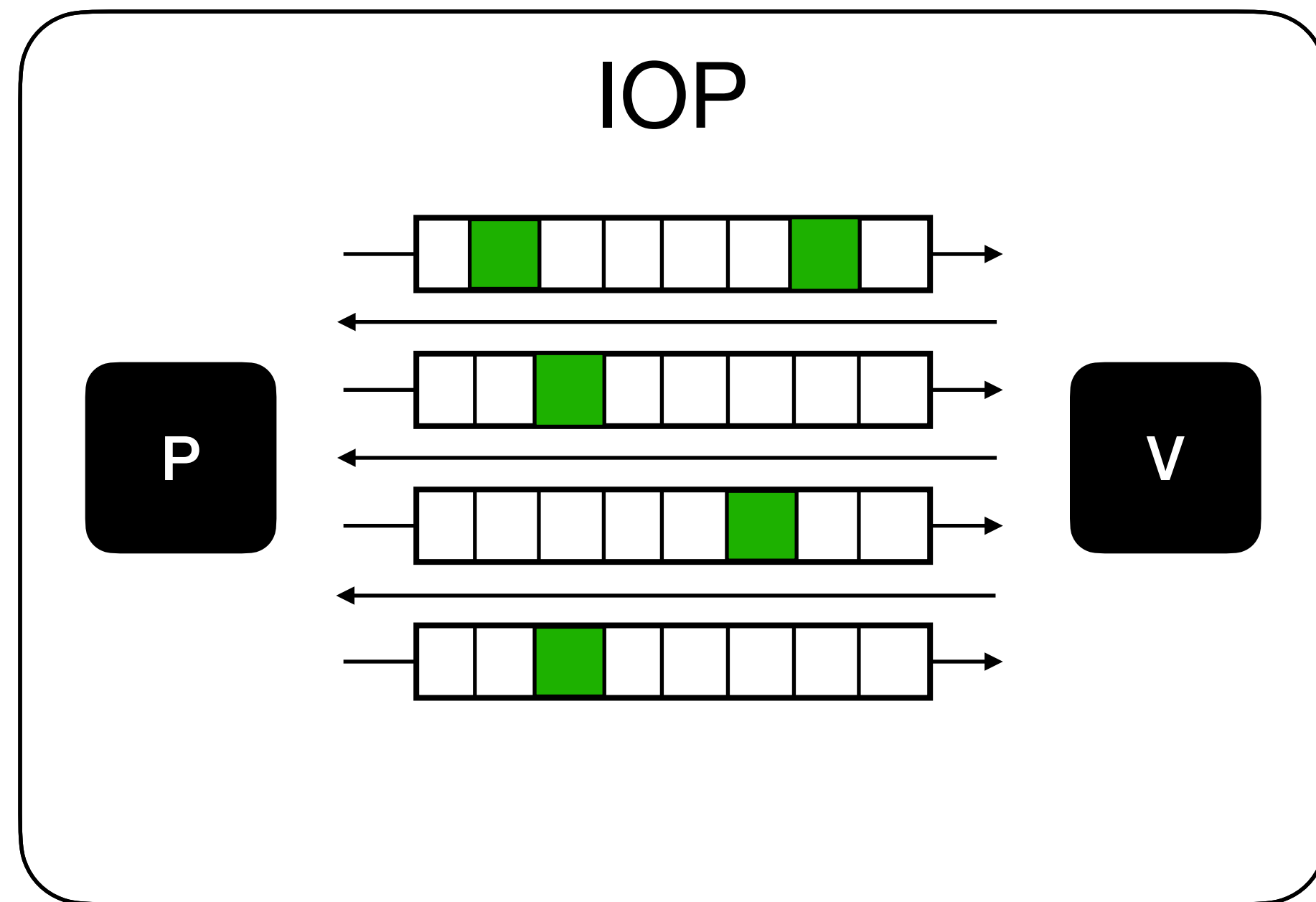


Proof length  $l \approx O(n)$

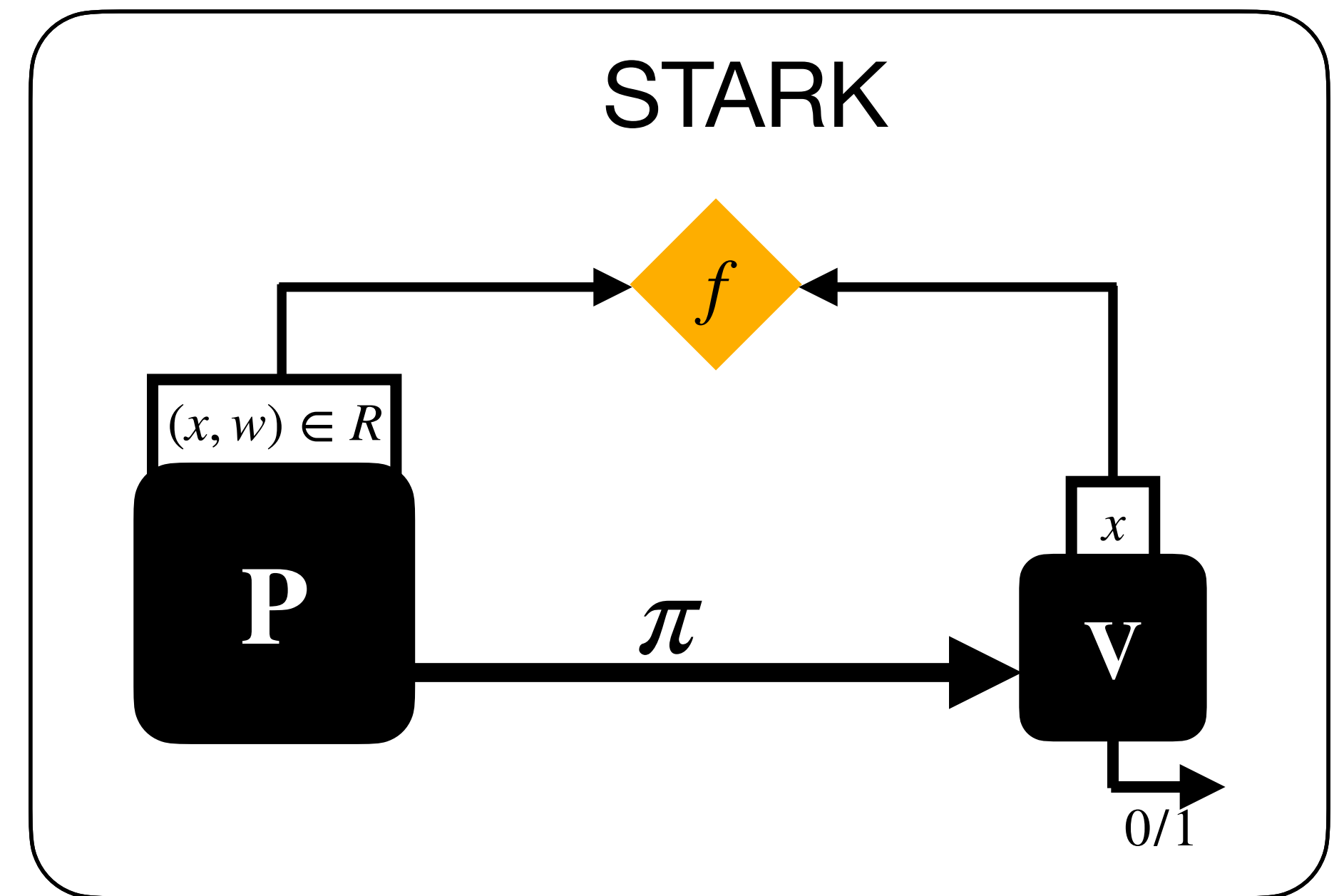
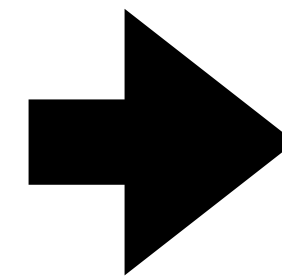
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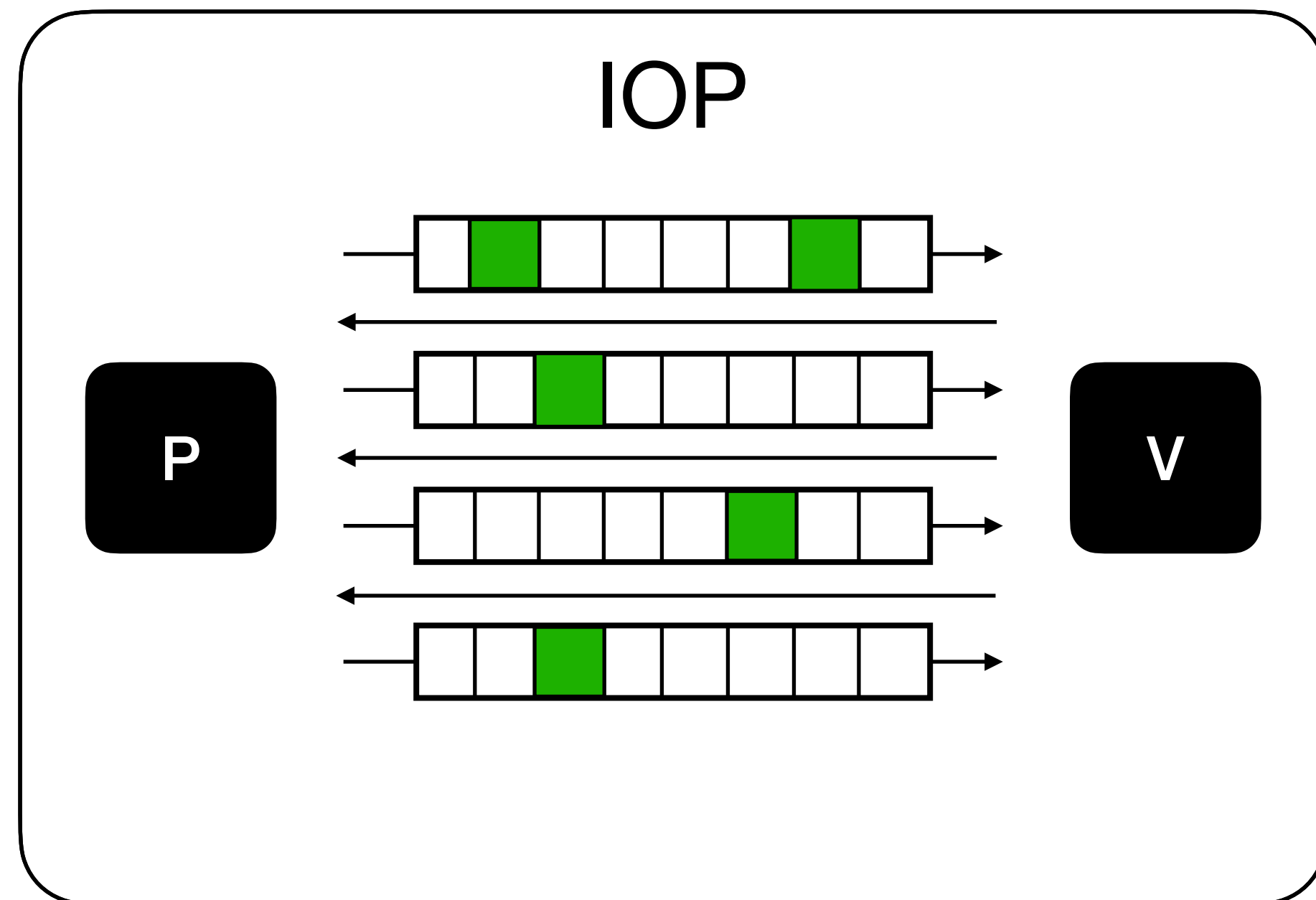


Proof length  $l \approx O(n)$  **Large, think  $2^{24}$**

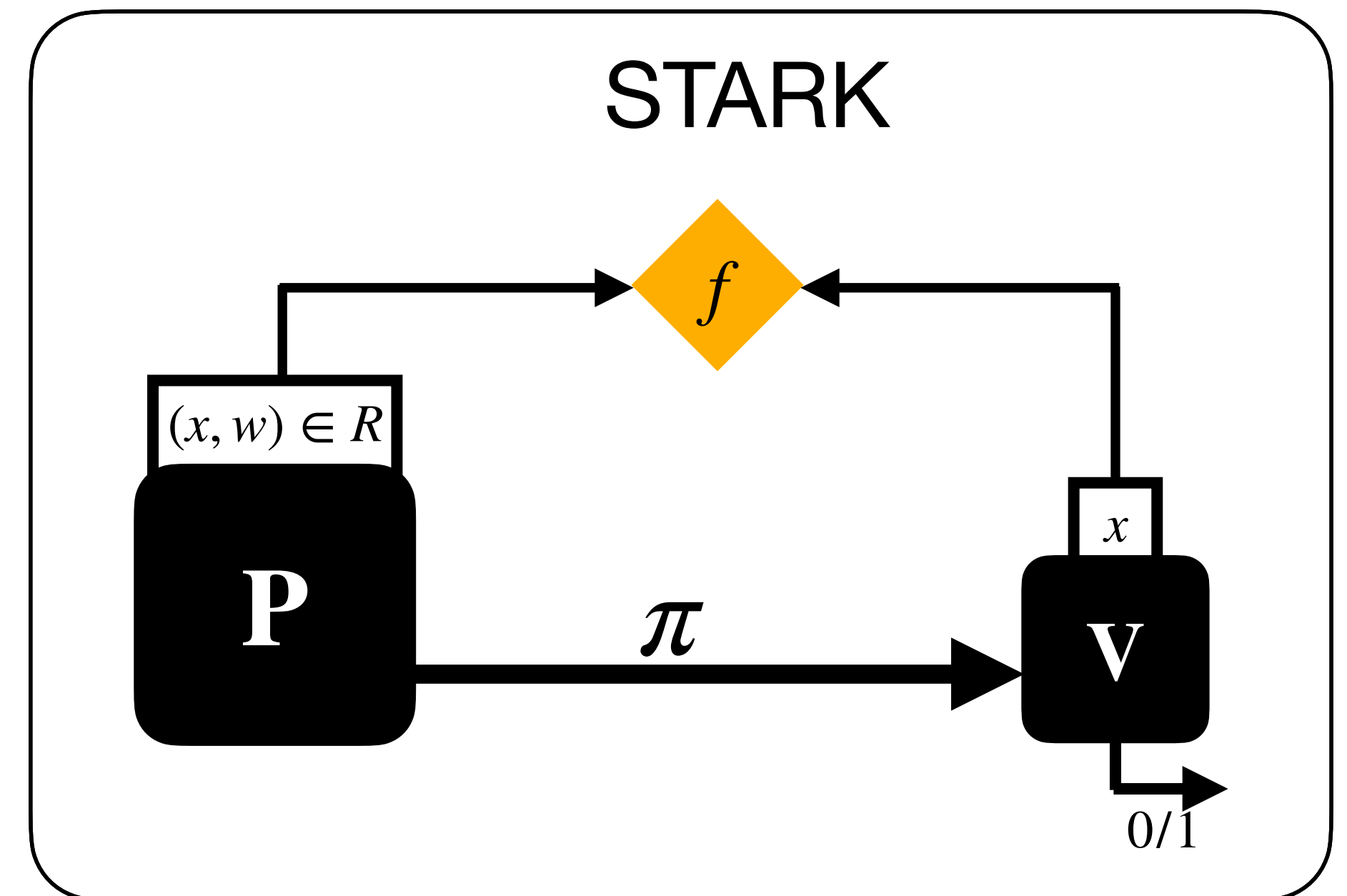
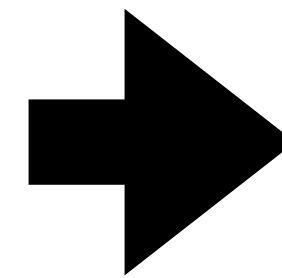
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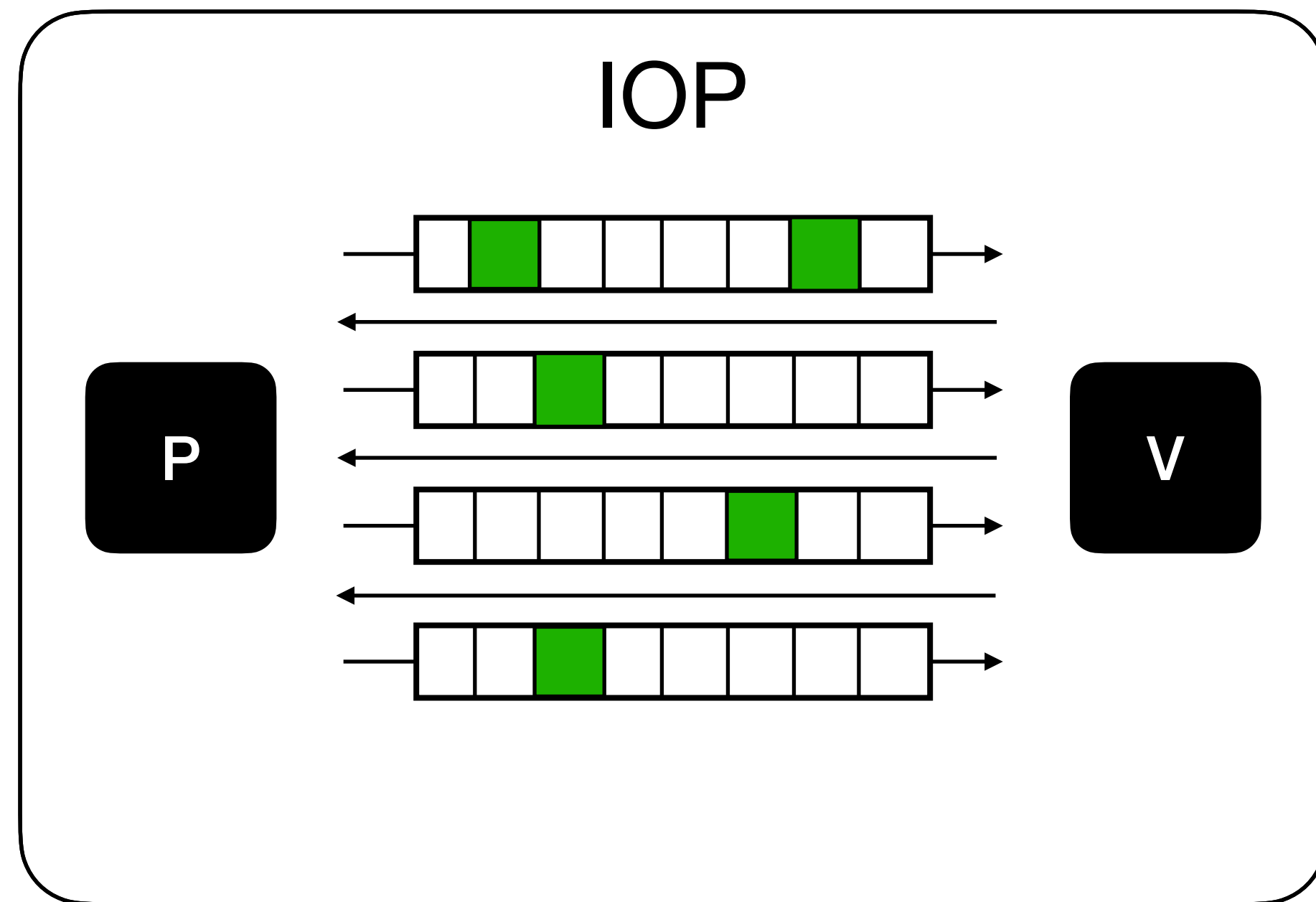
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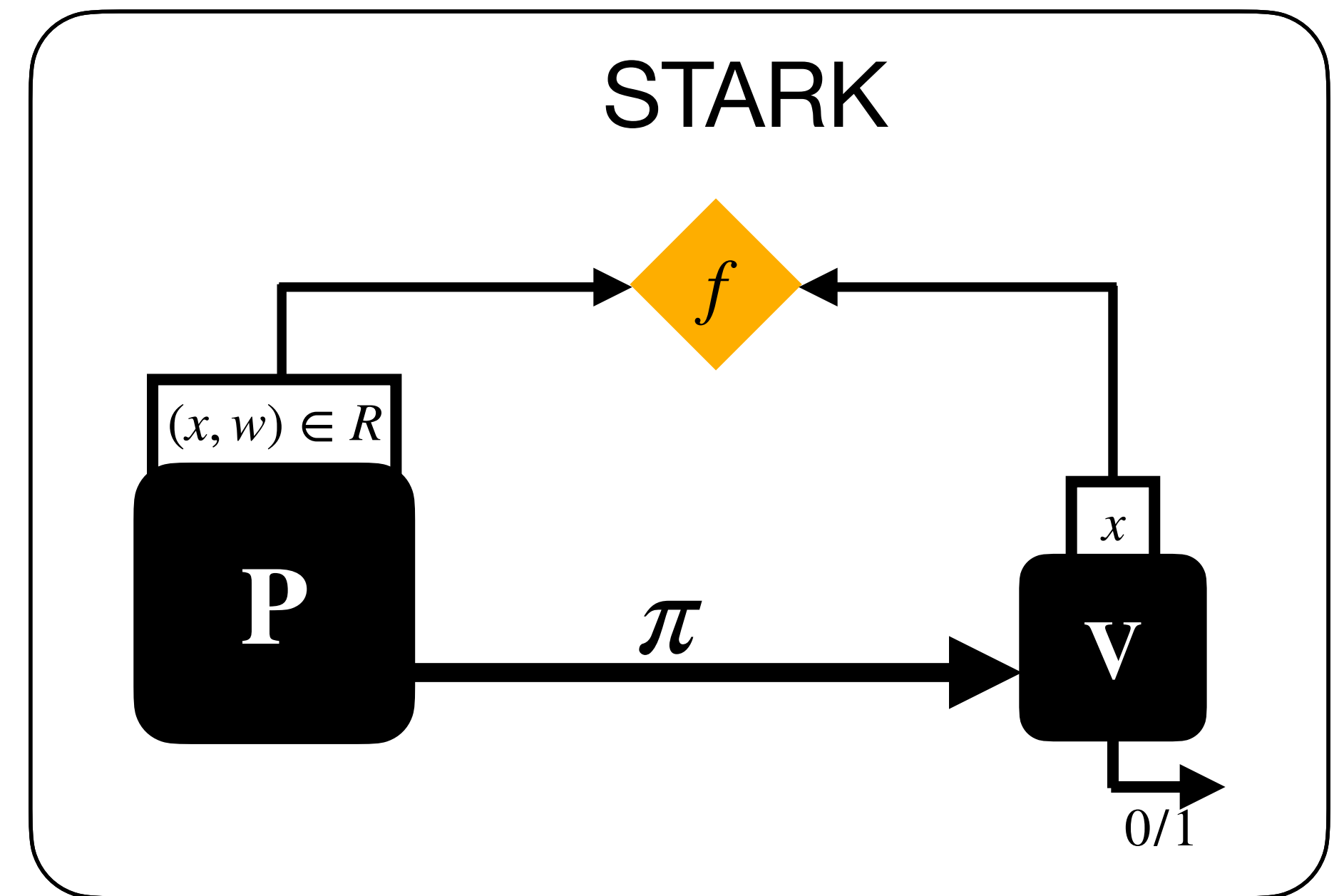
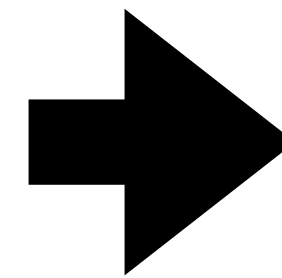
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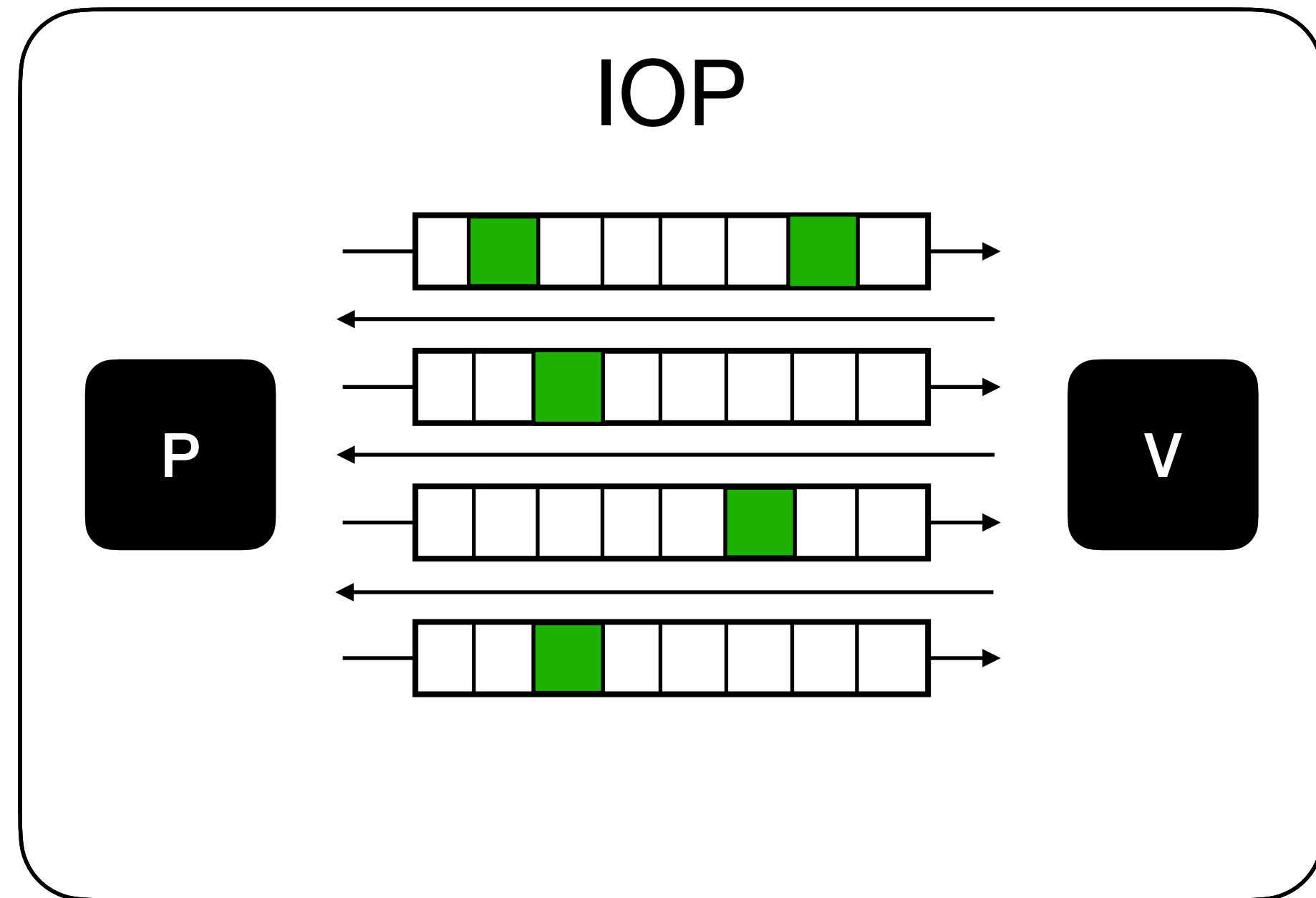
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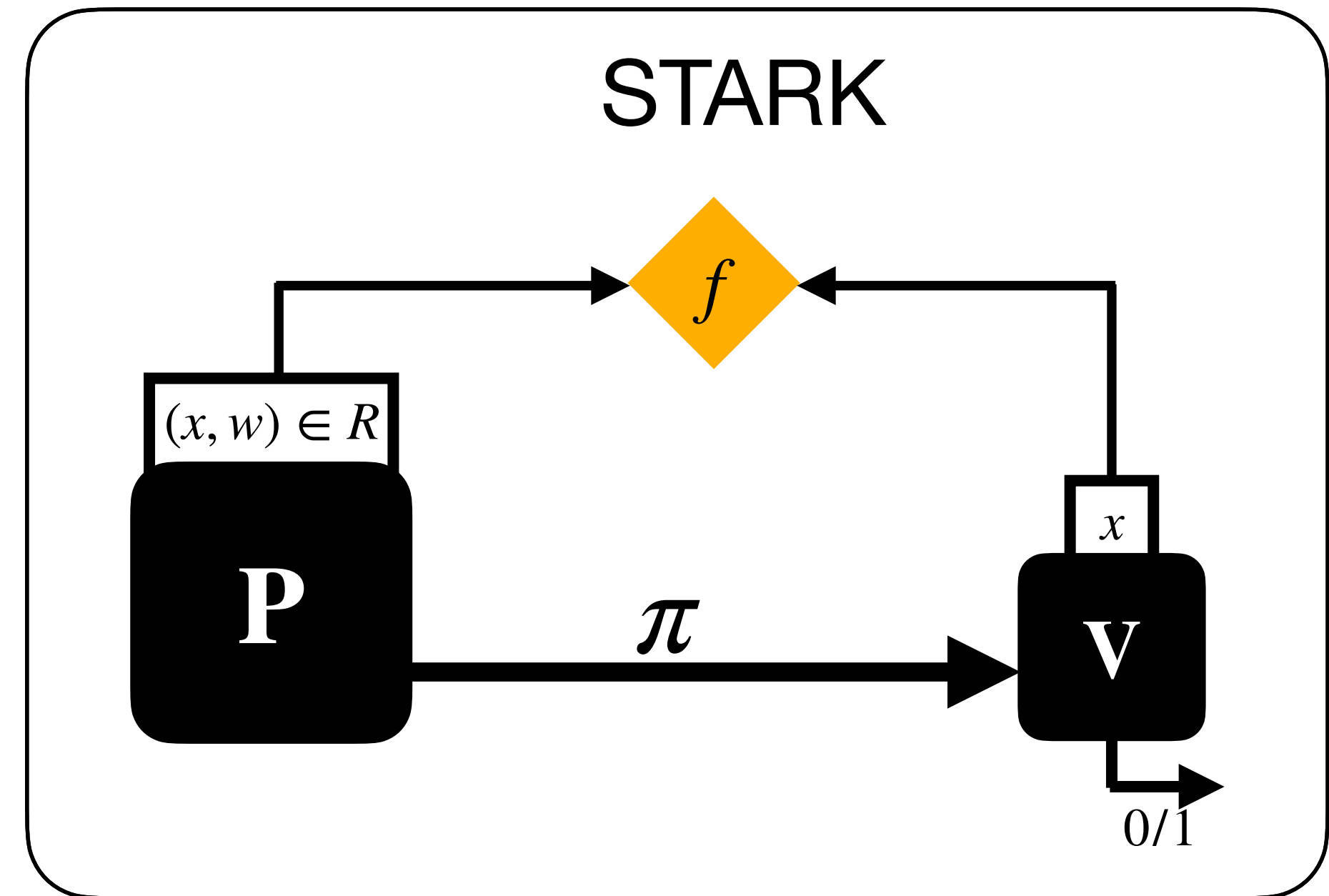
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In this talk, we focus on the IOP!



BCS  
➔



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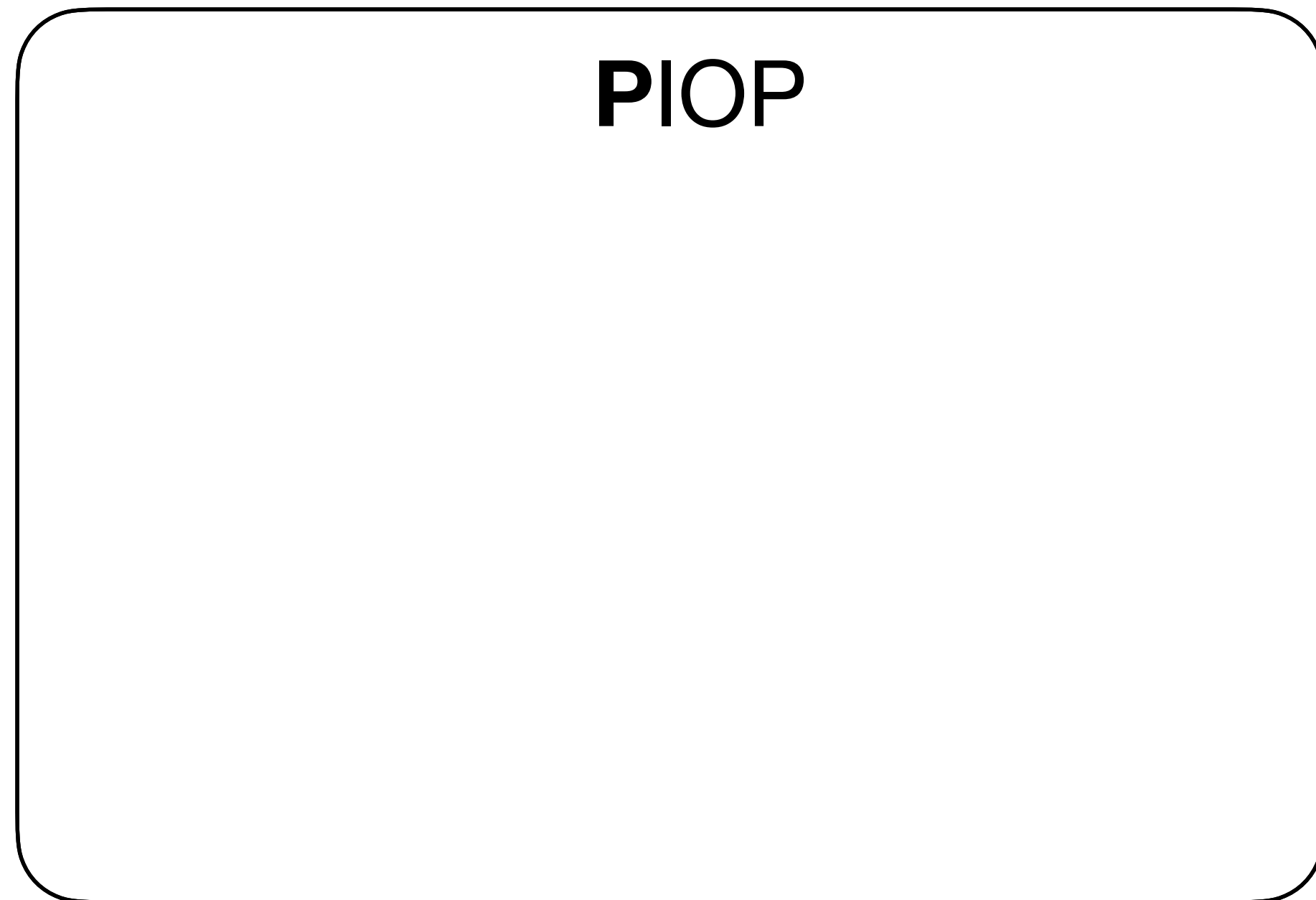


PIOP



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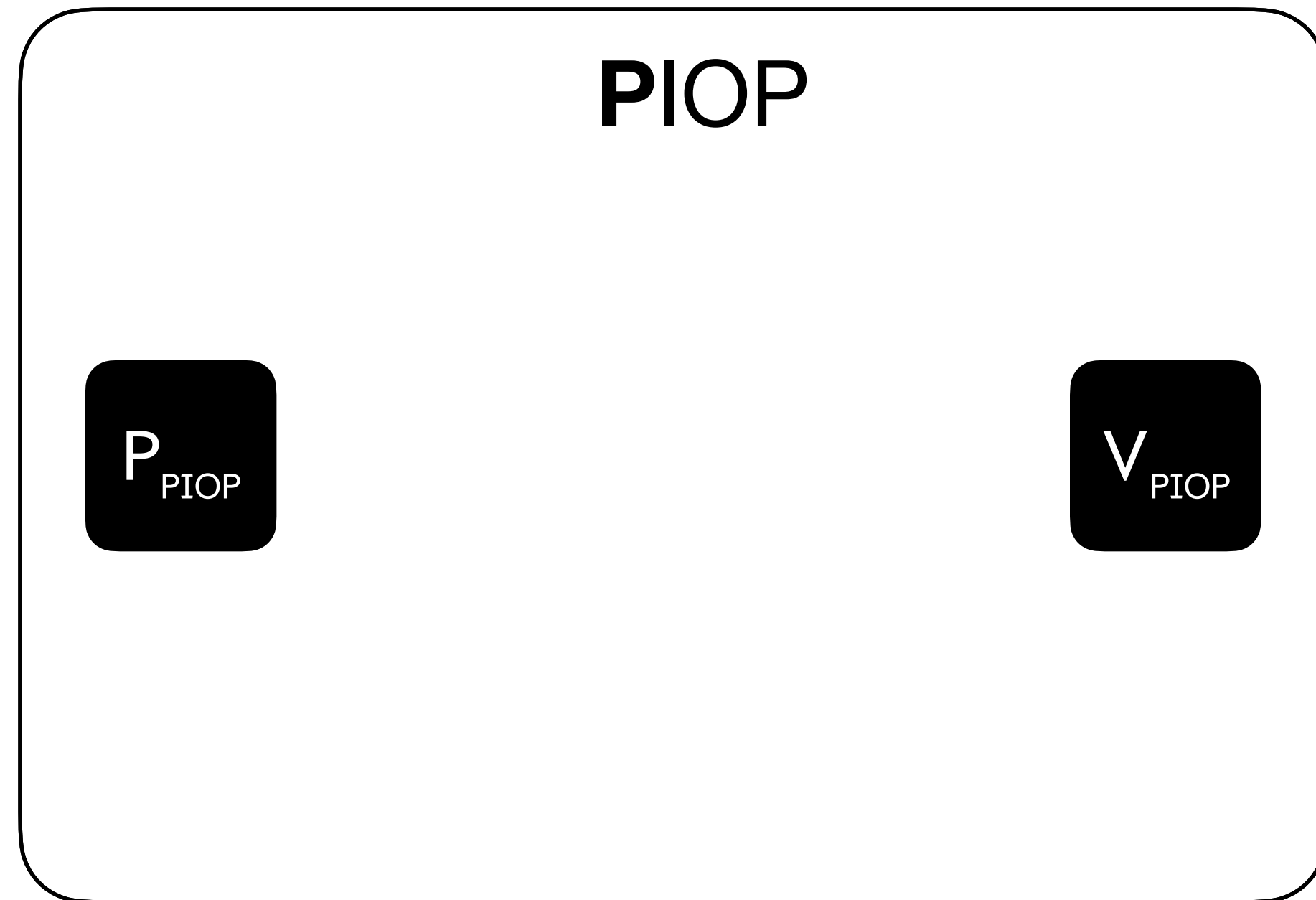


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E.g. Aurora, STARK PIOP etc.

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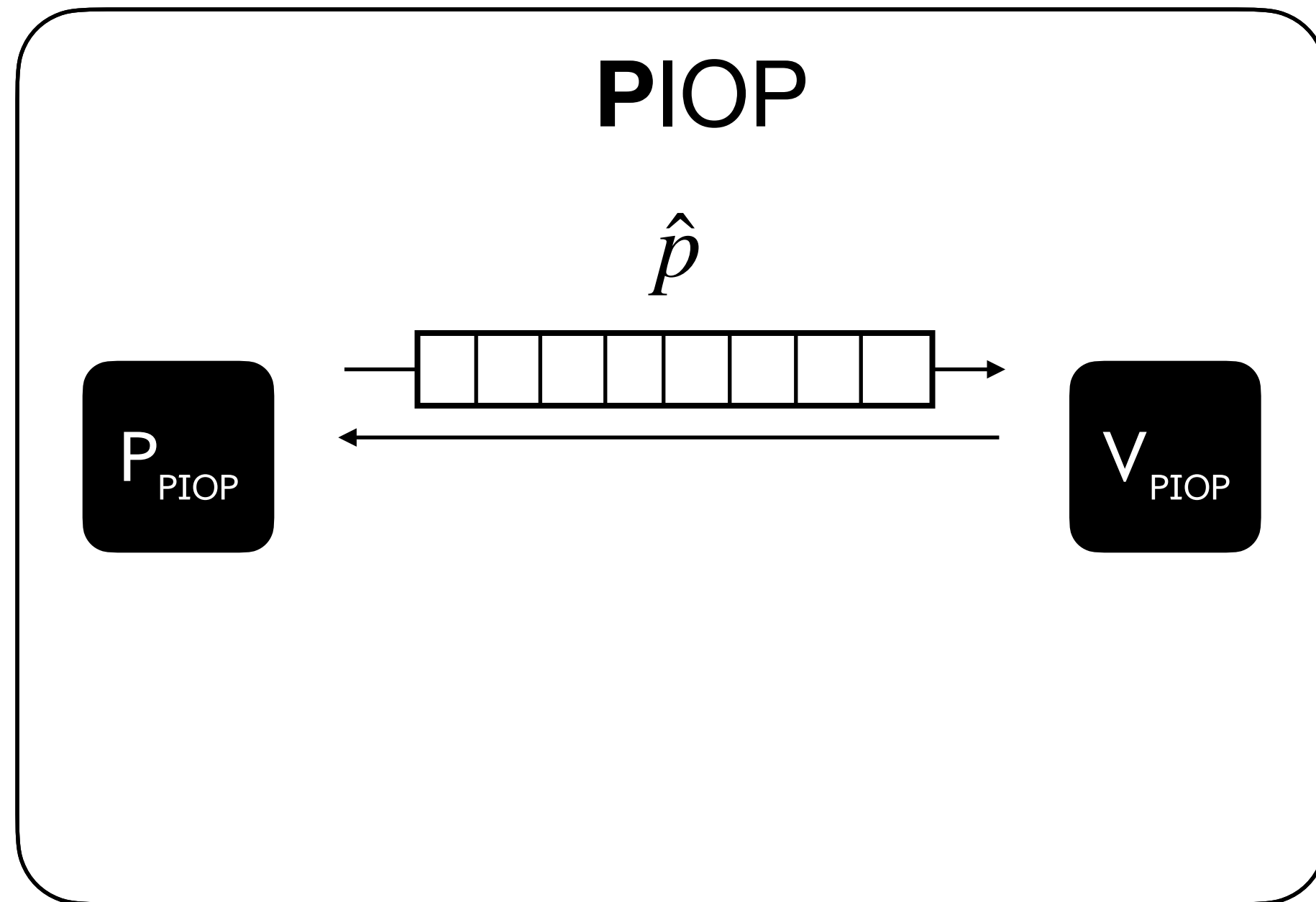


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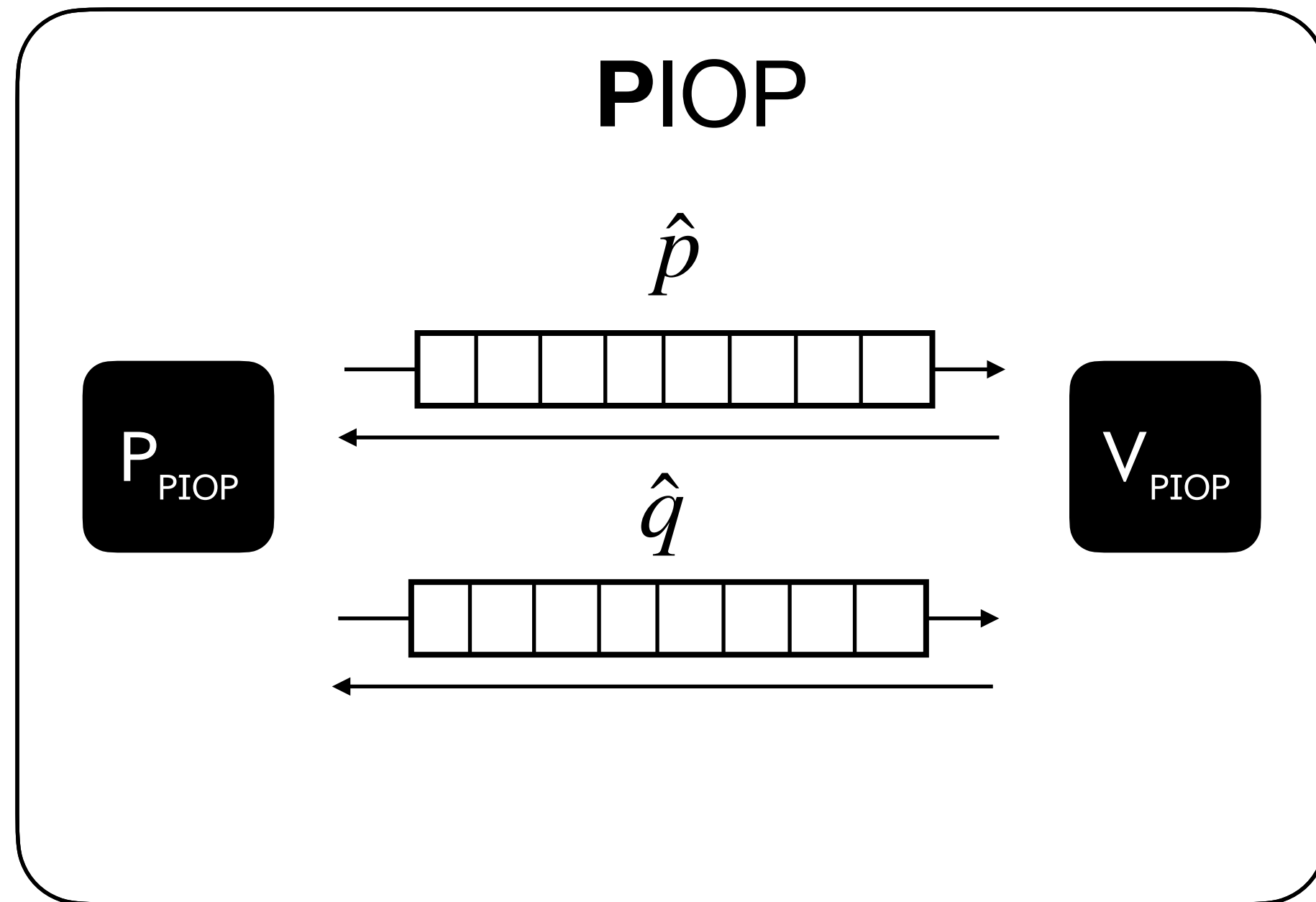


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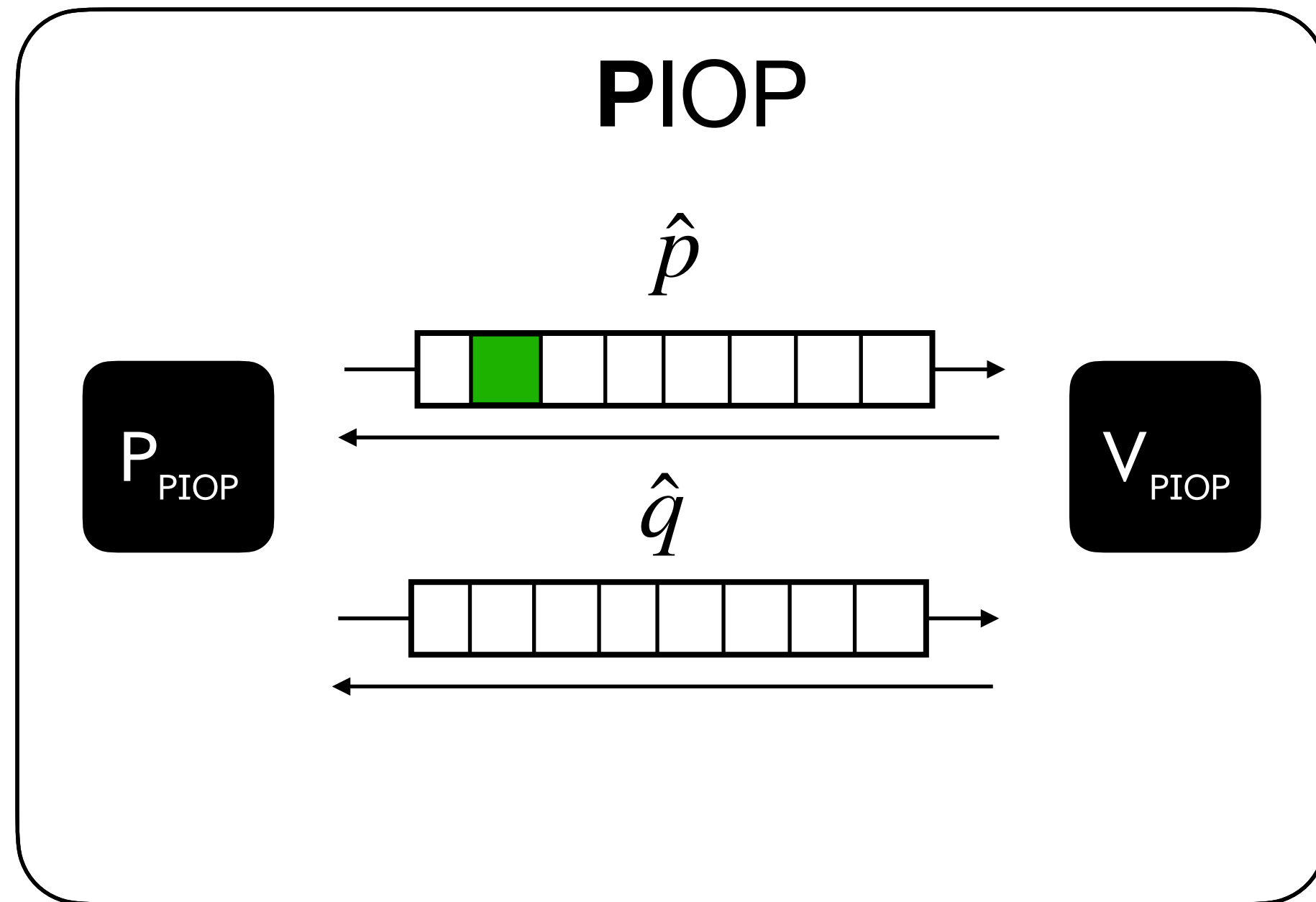


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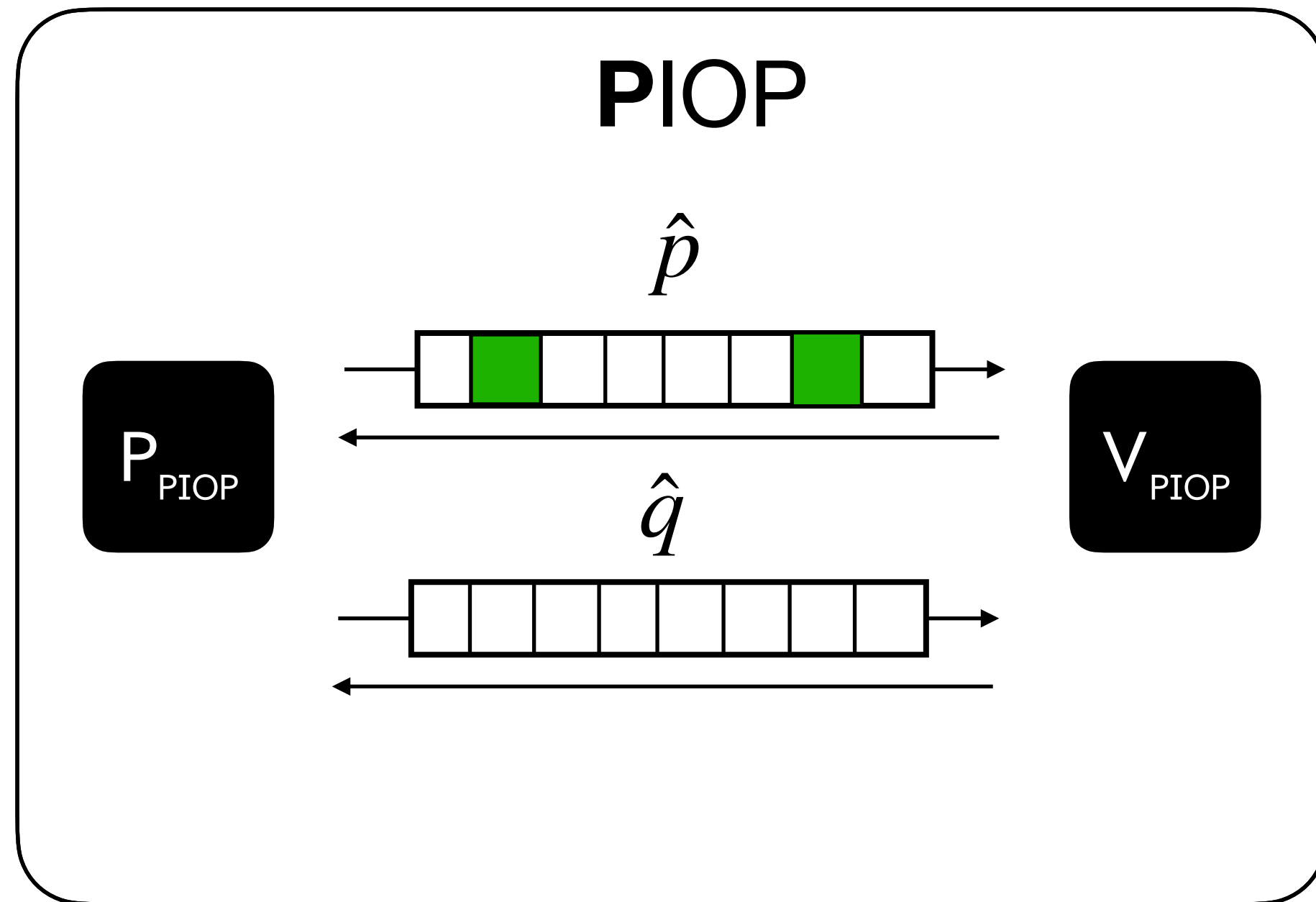


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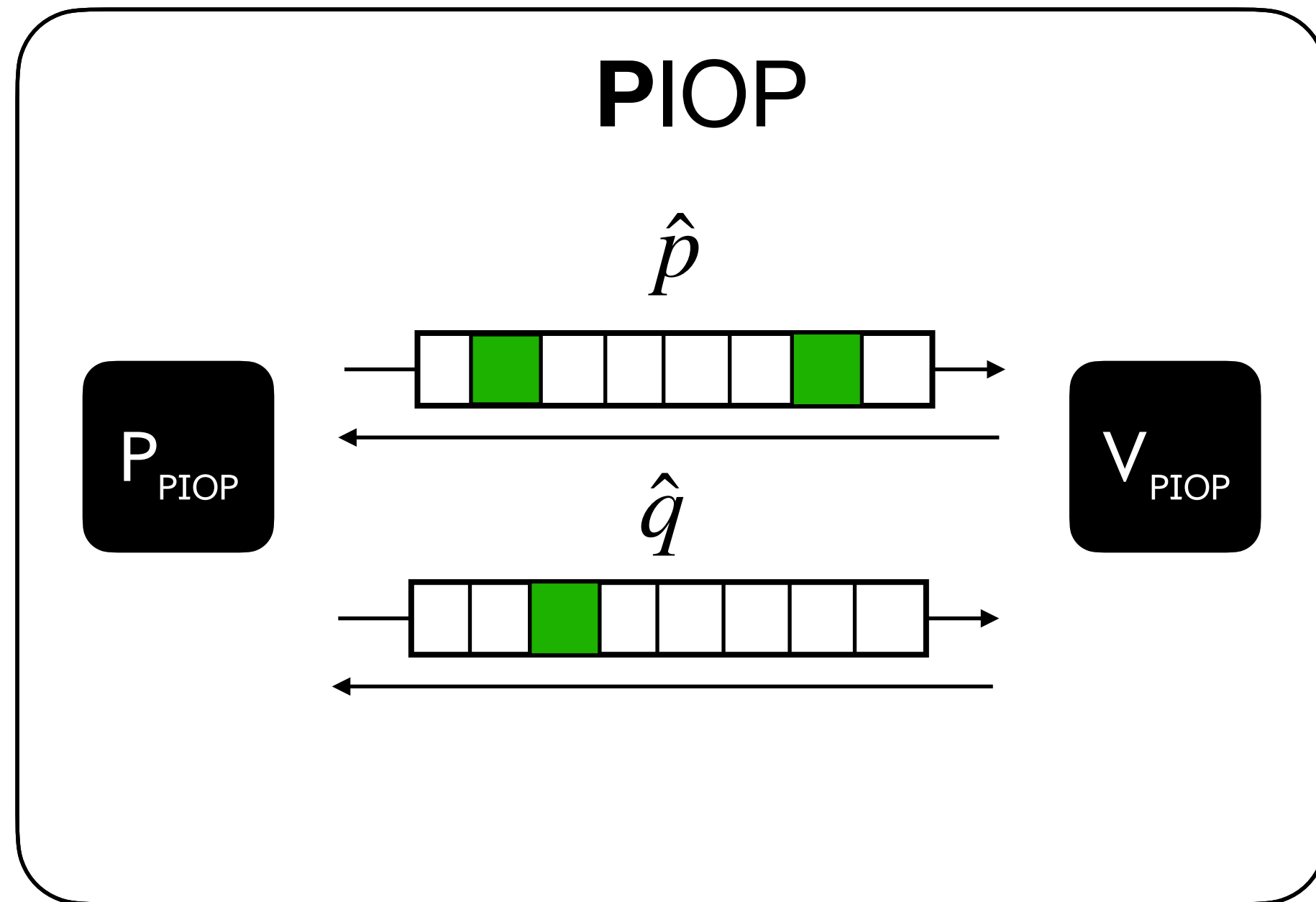


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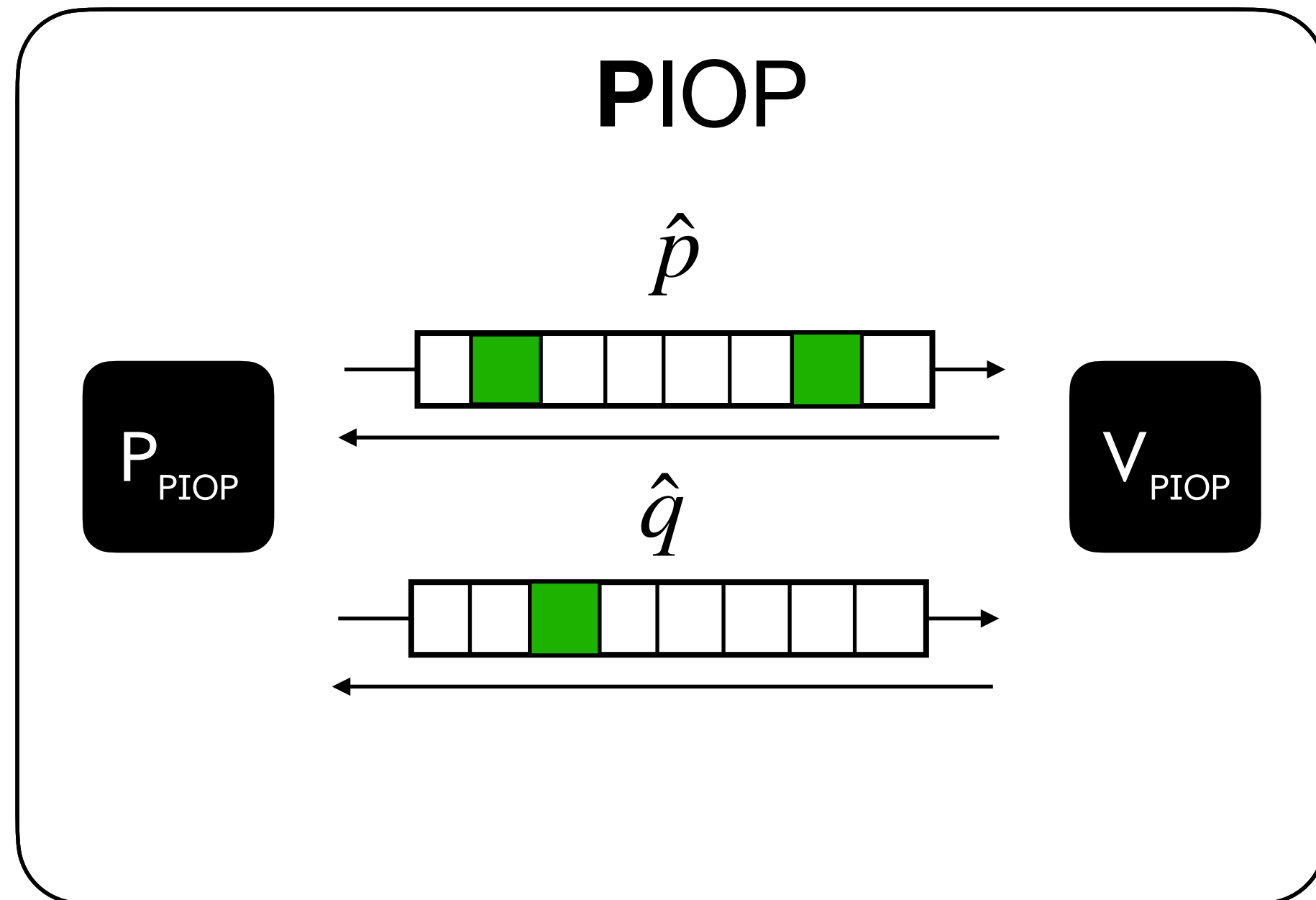
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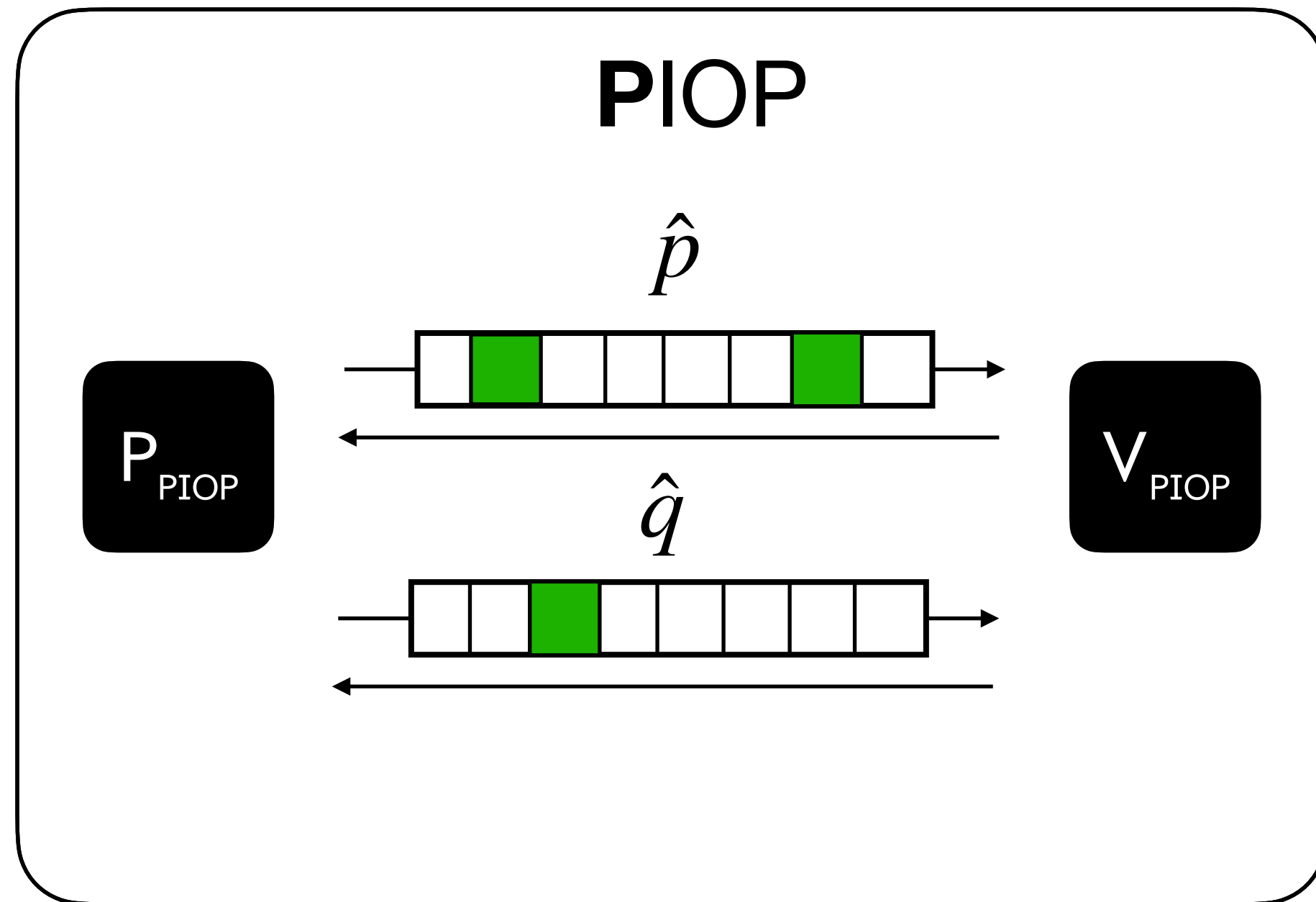
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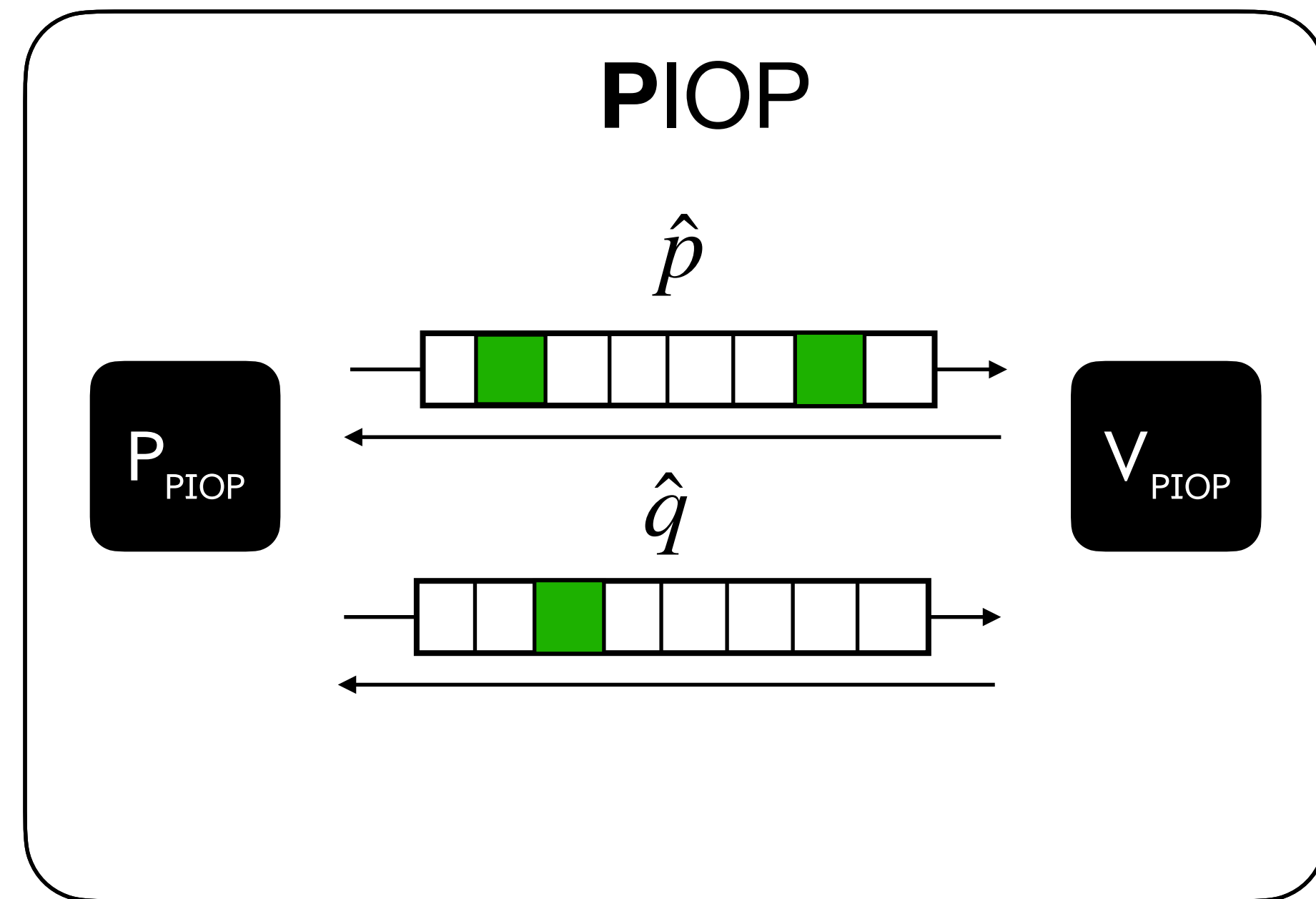


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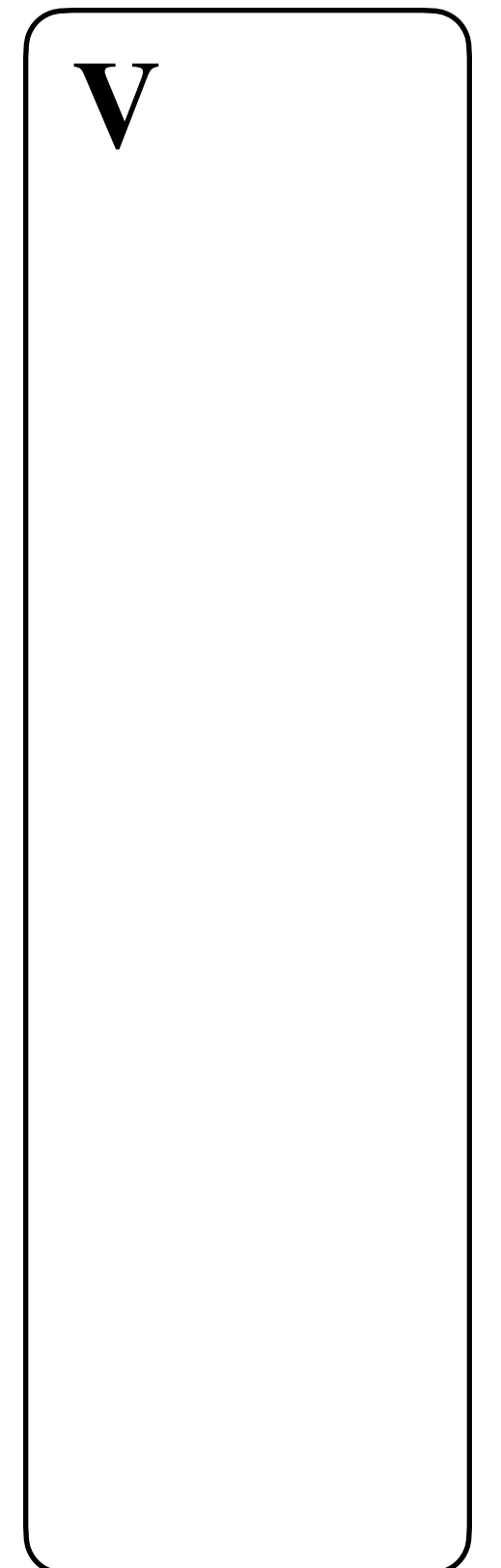
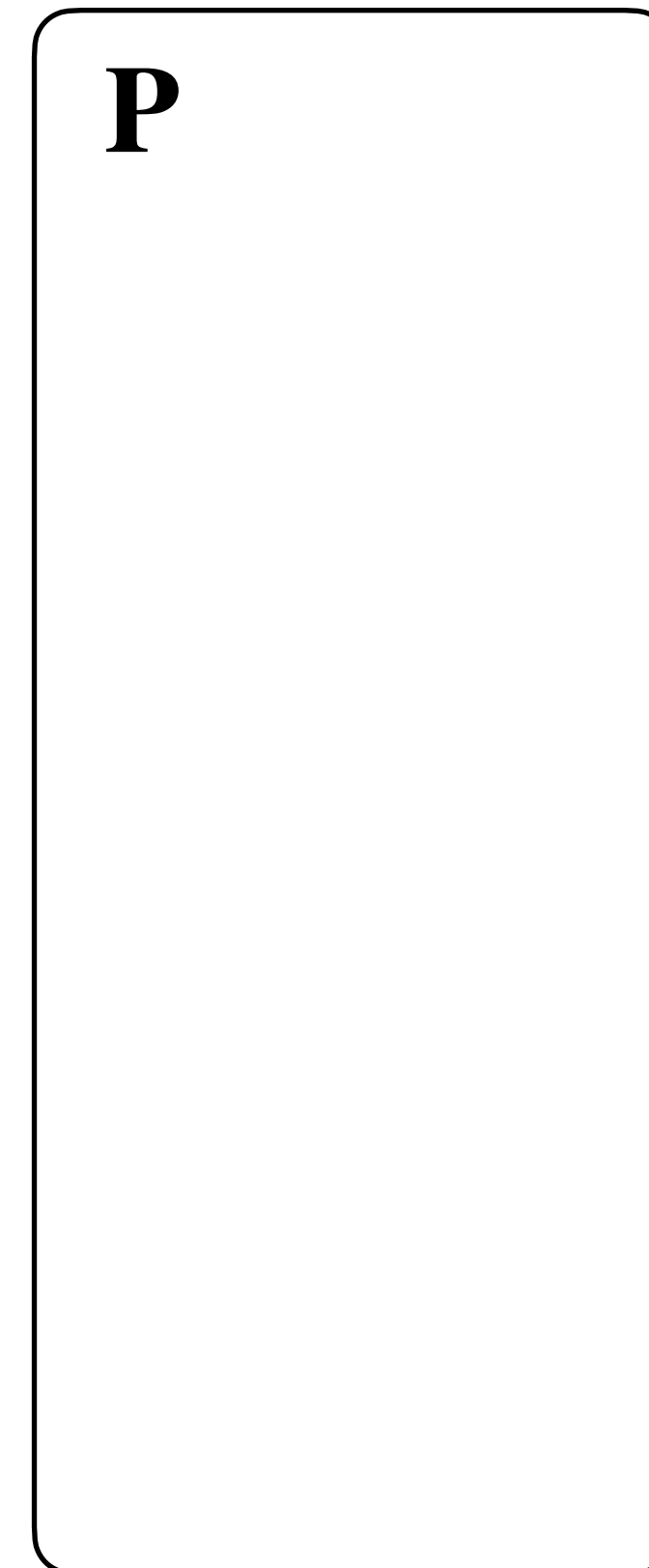
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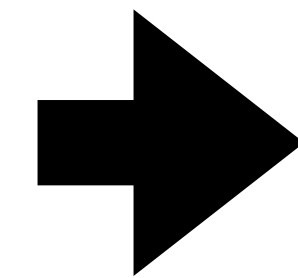
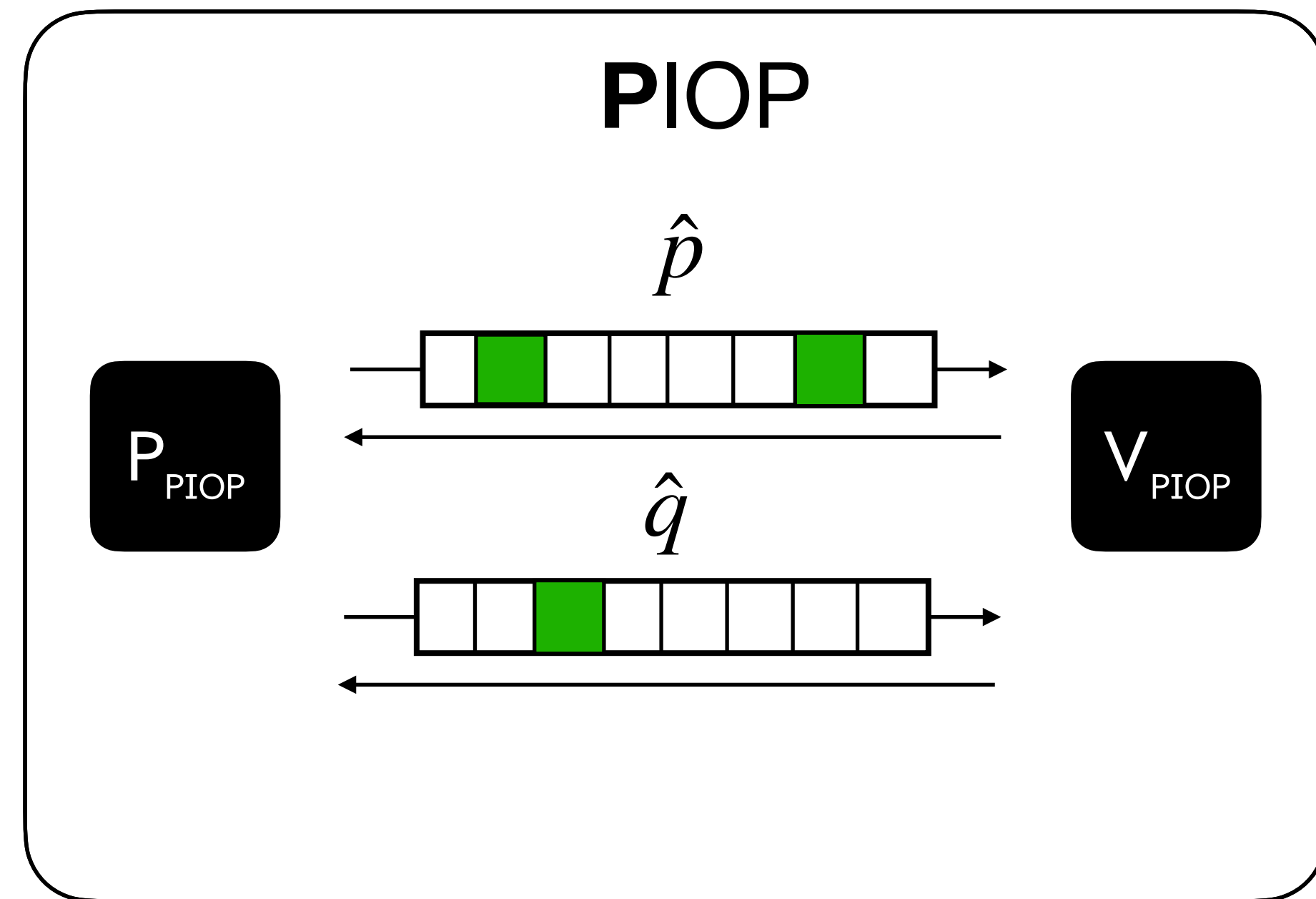
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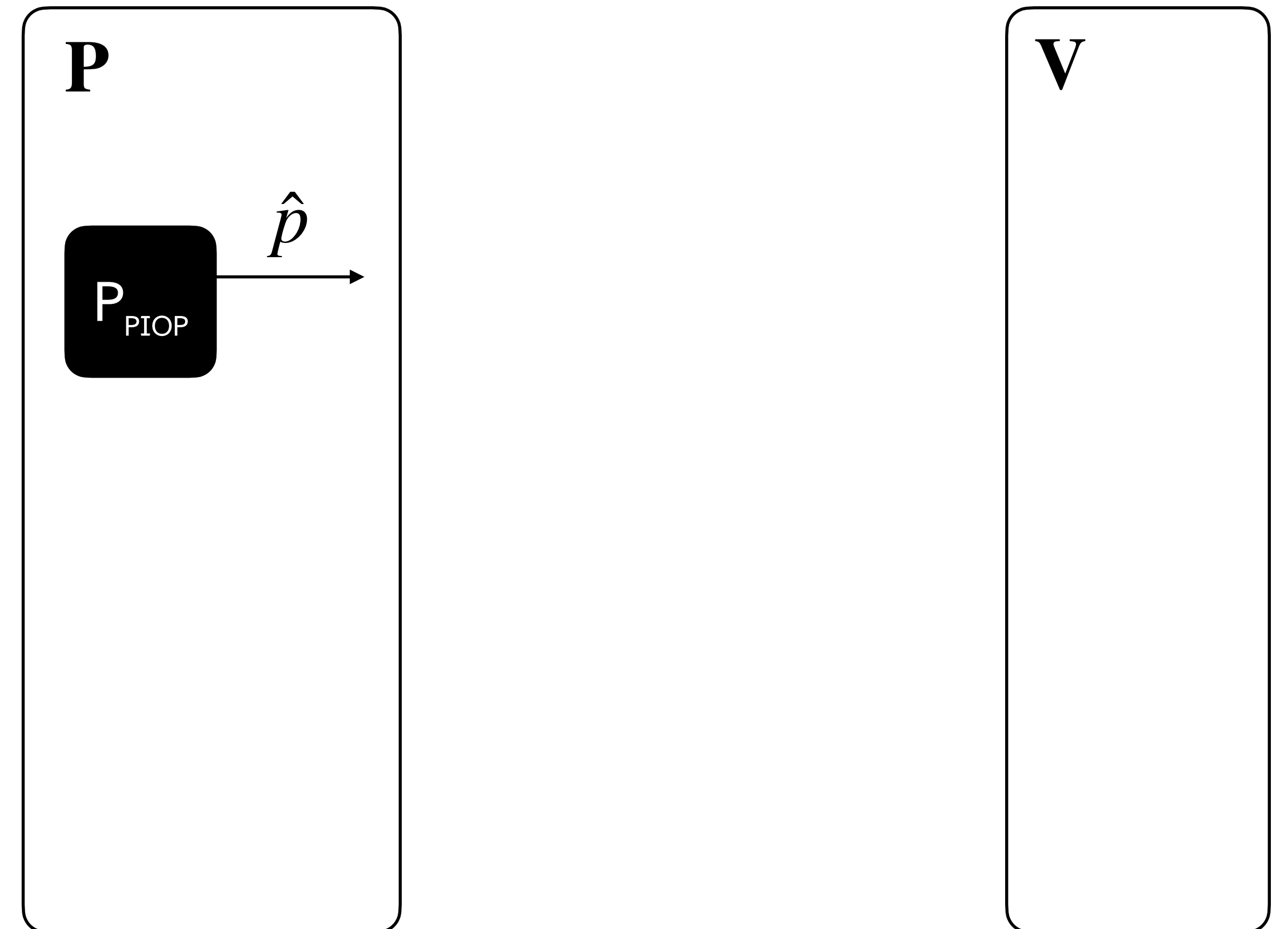
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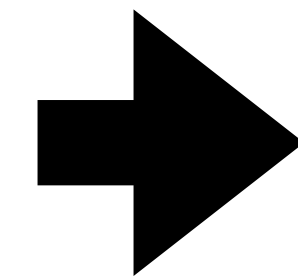
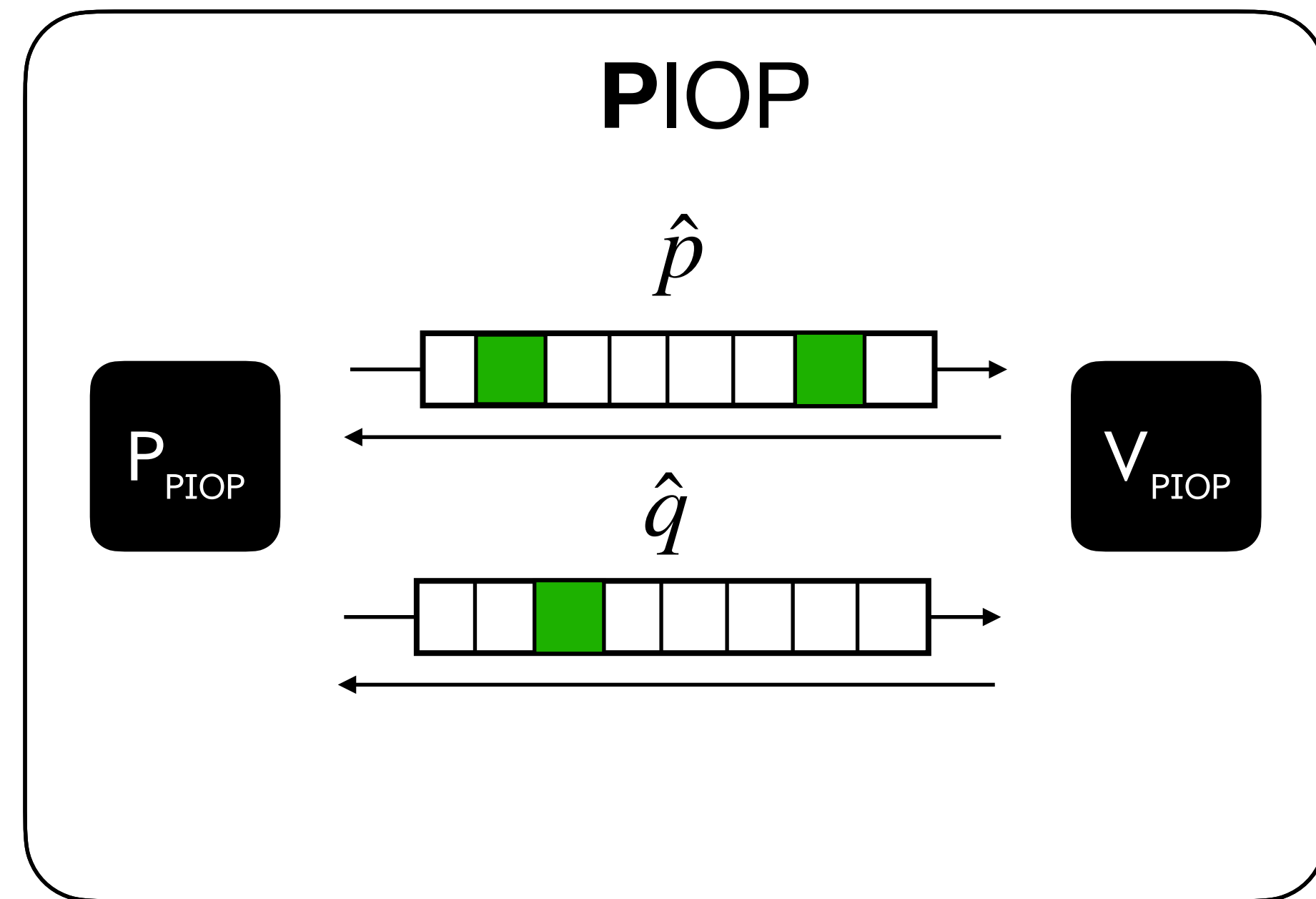


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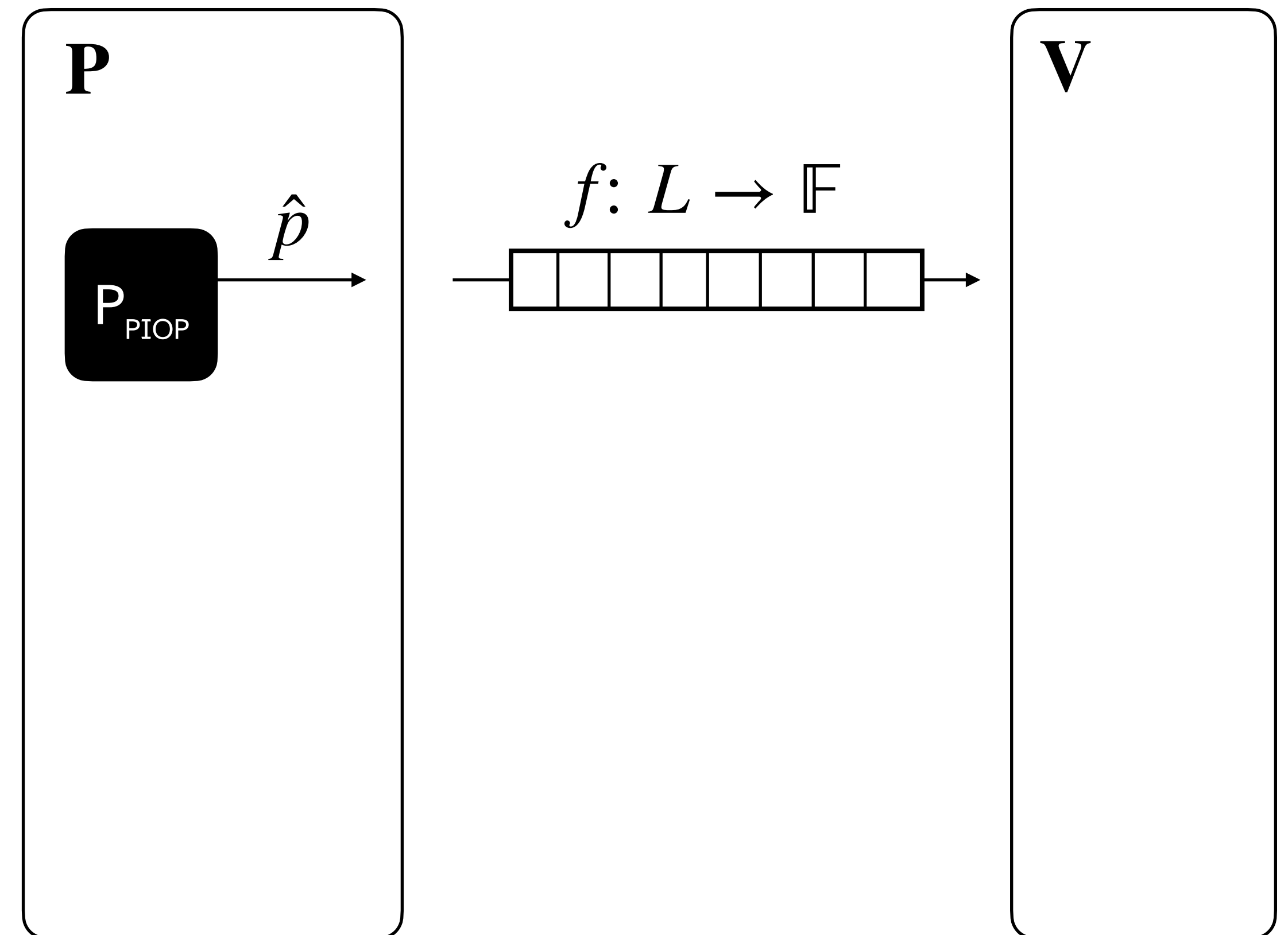
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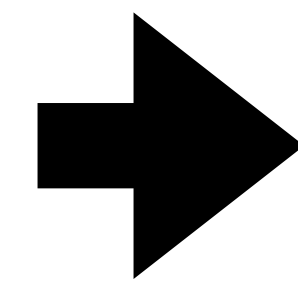
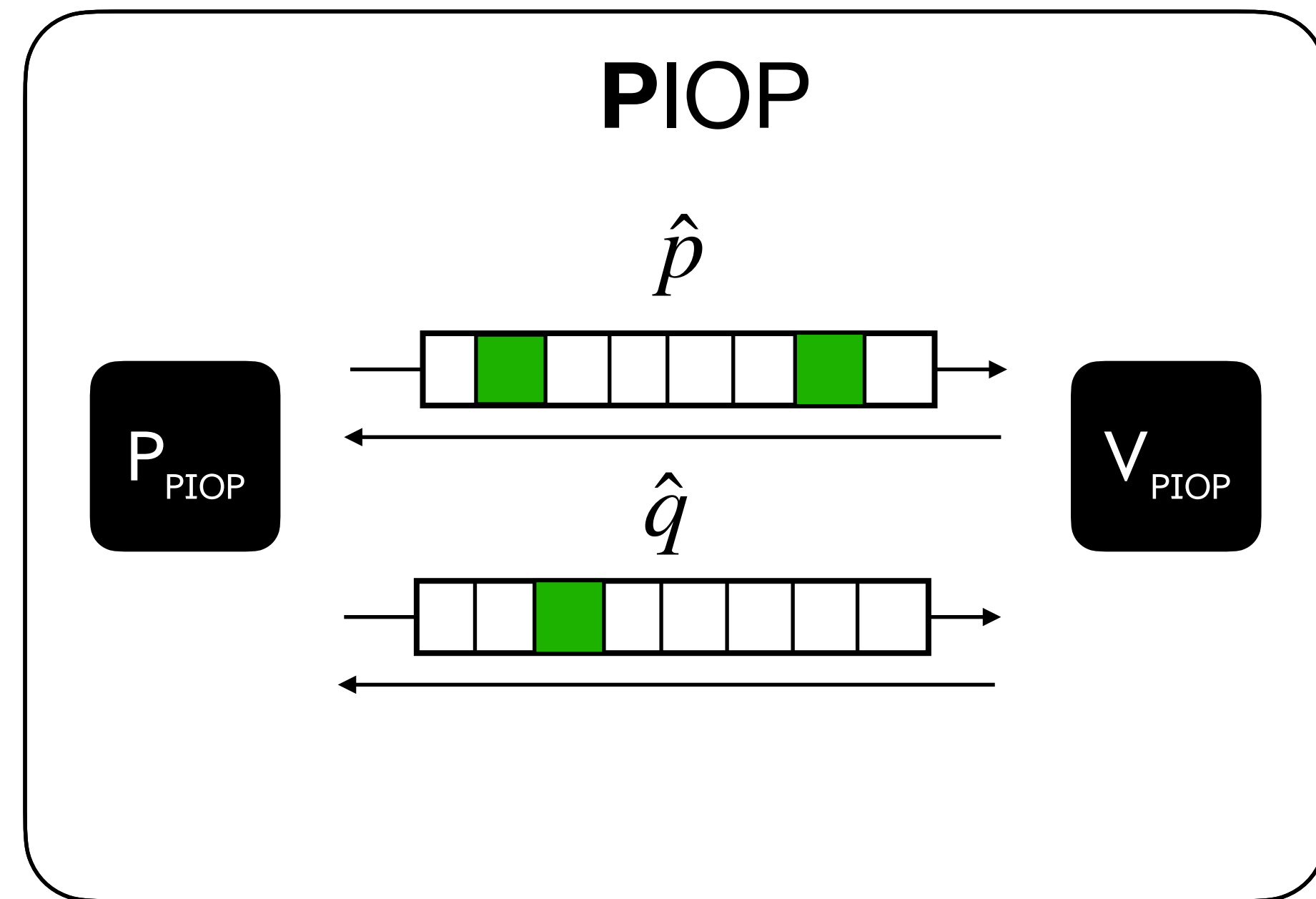


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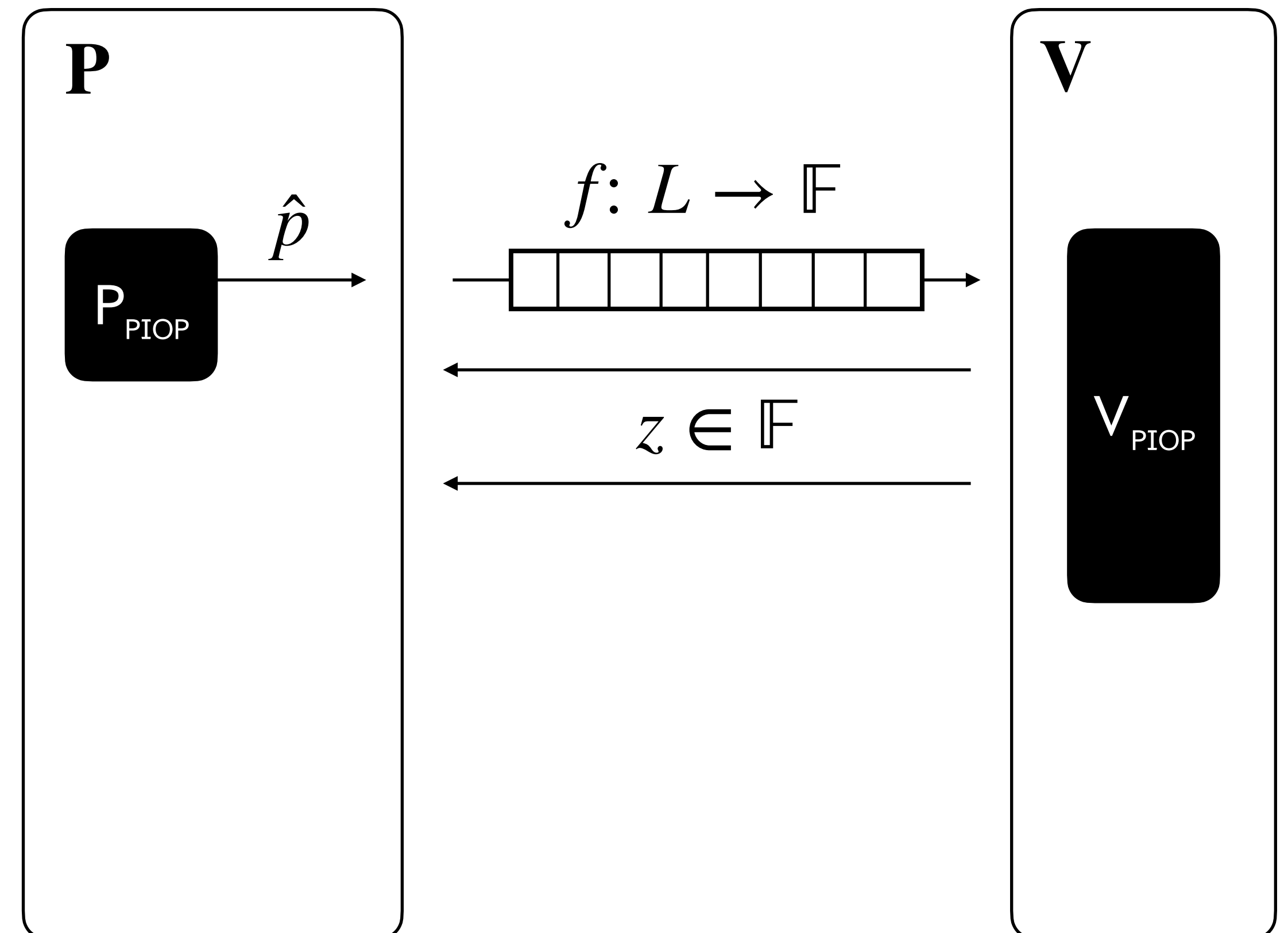
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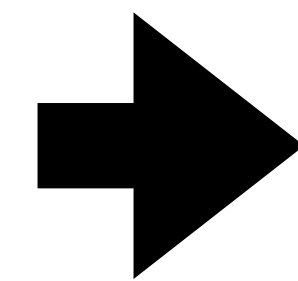
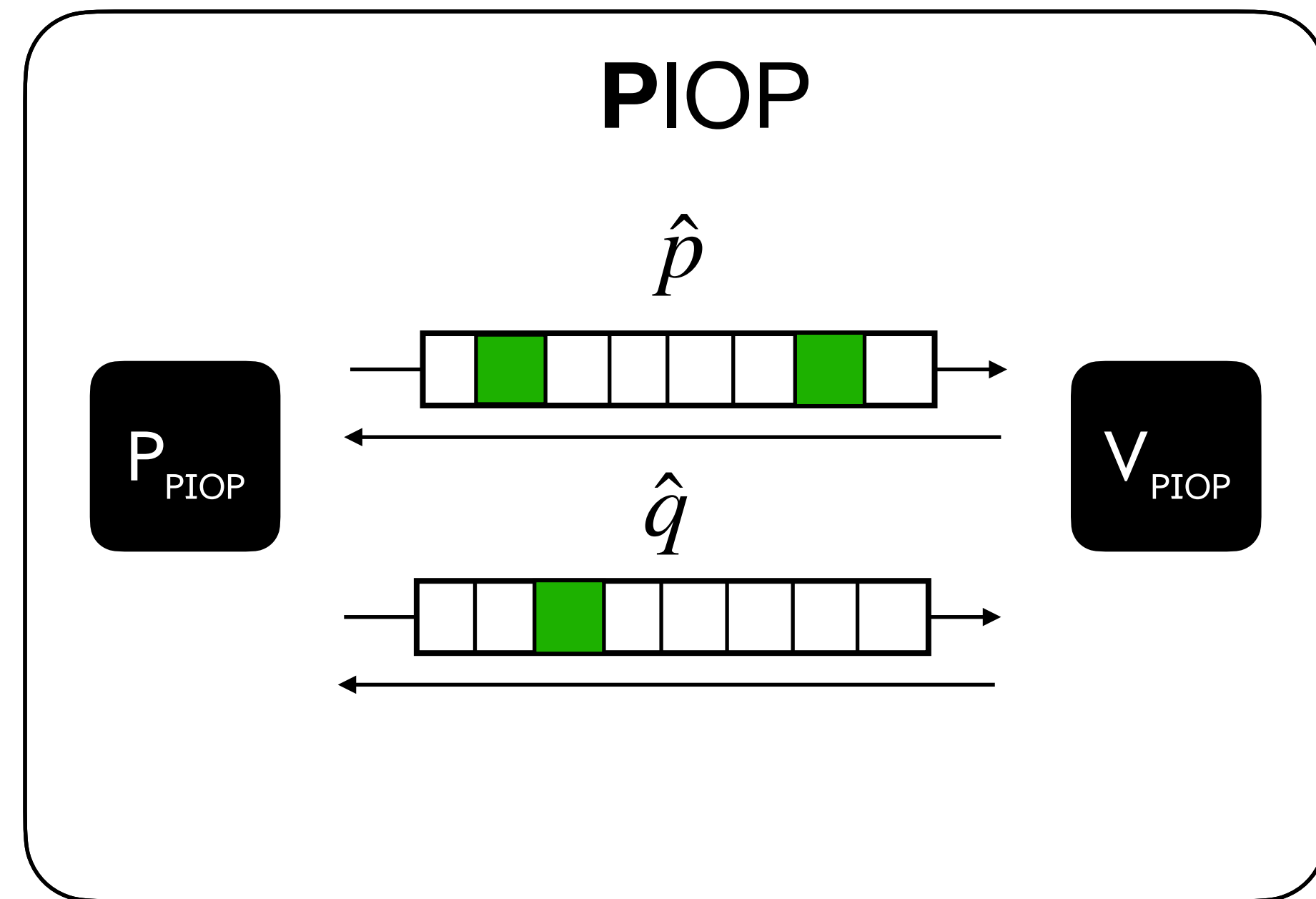


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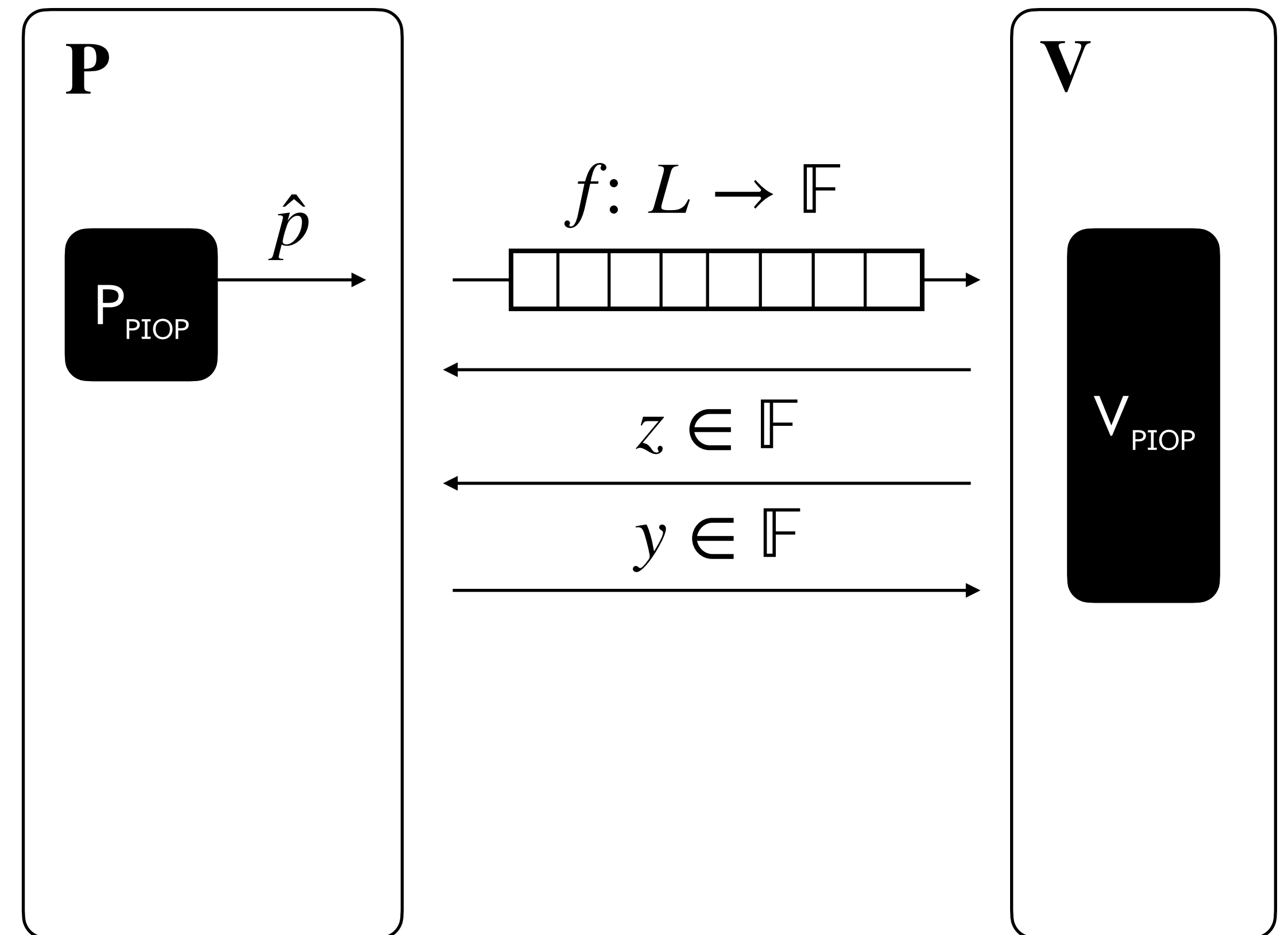
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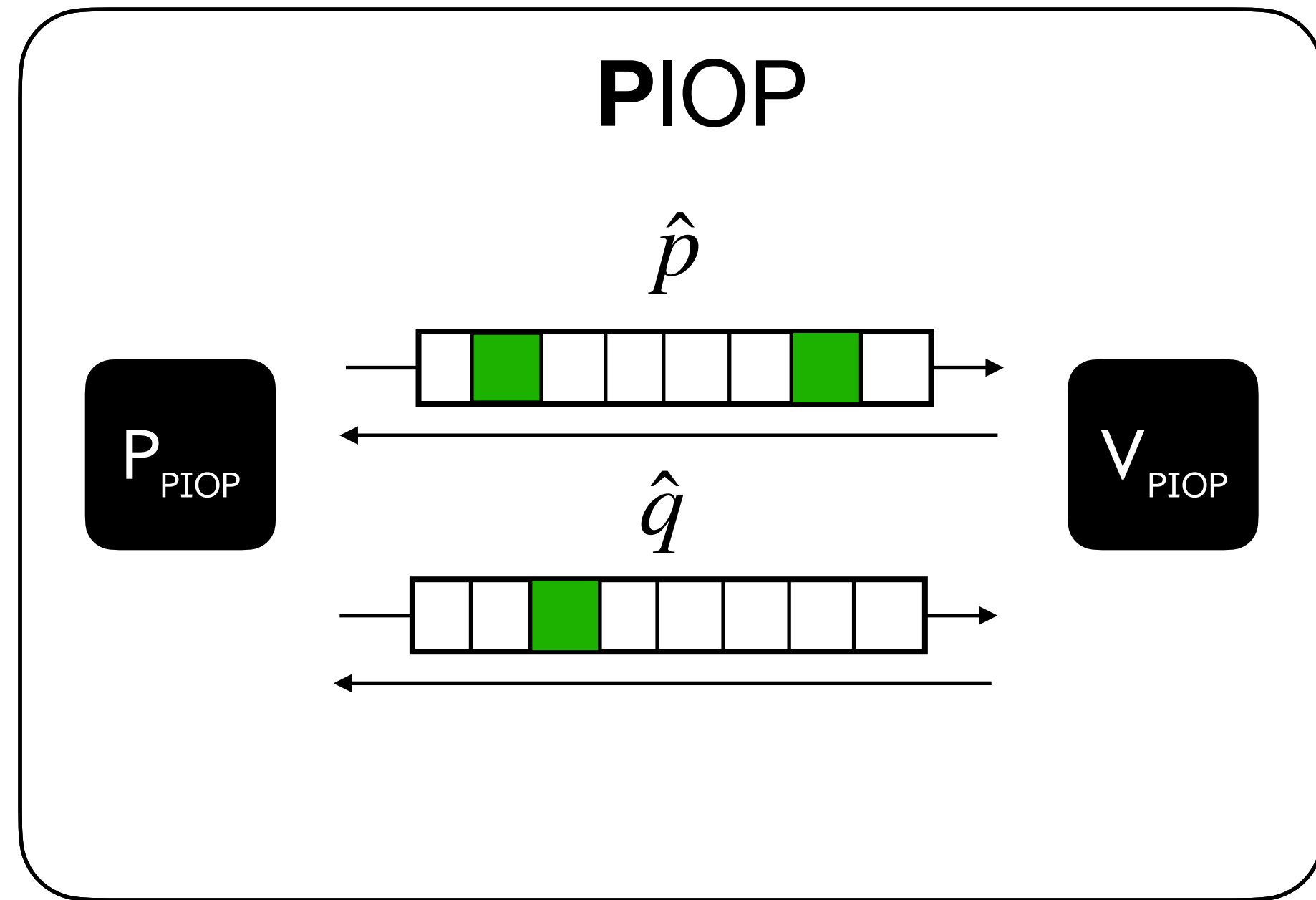
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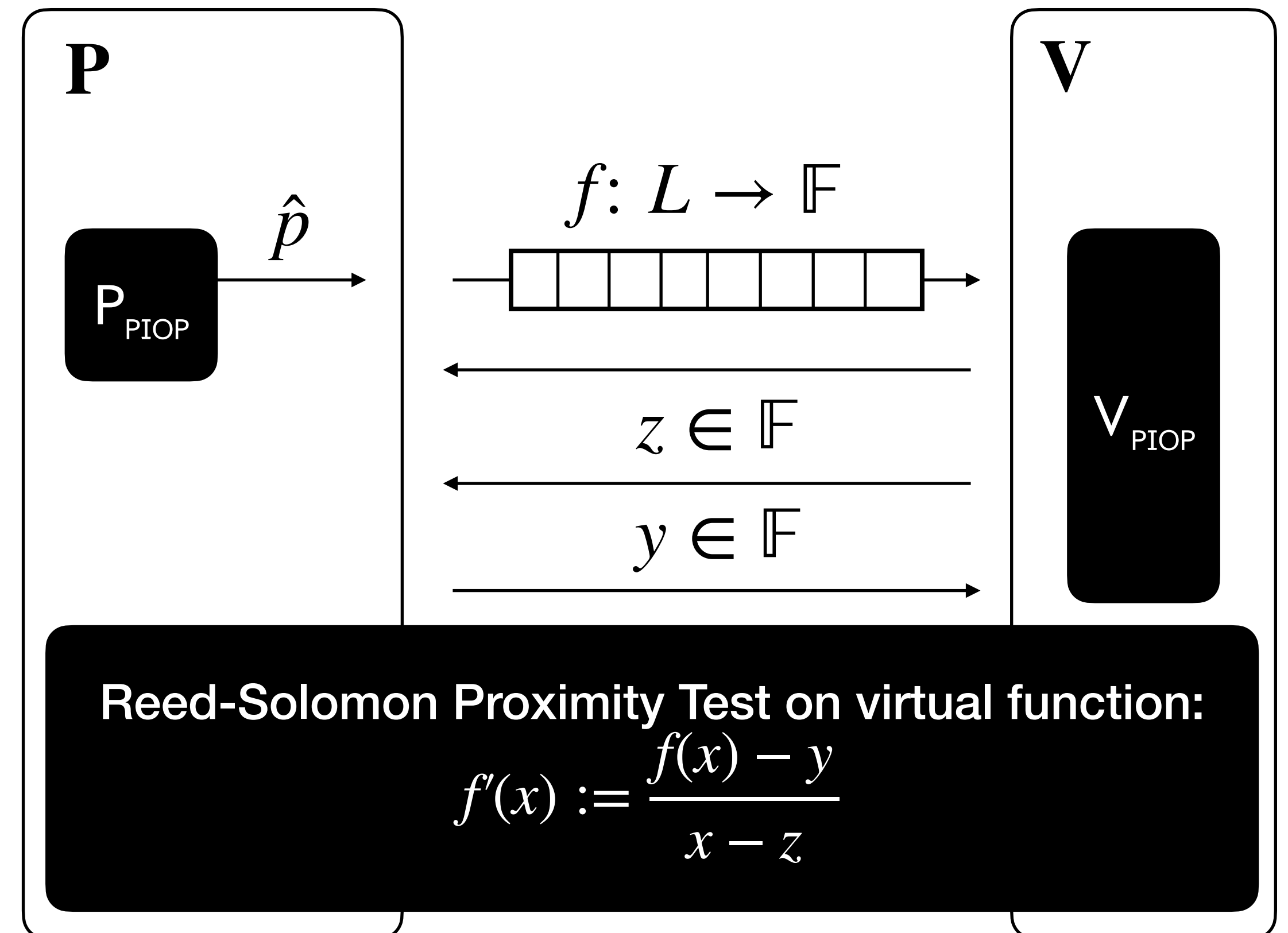
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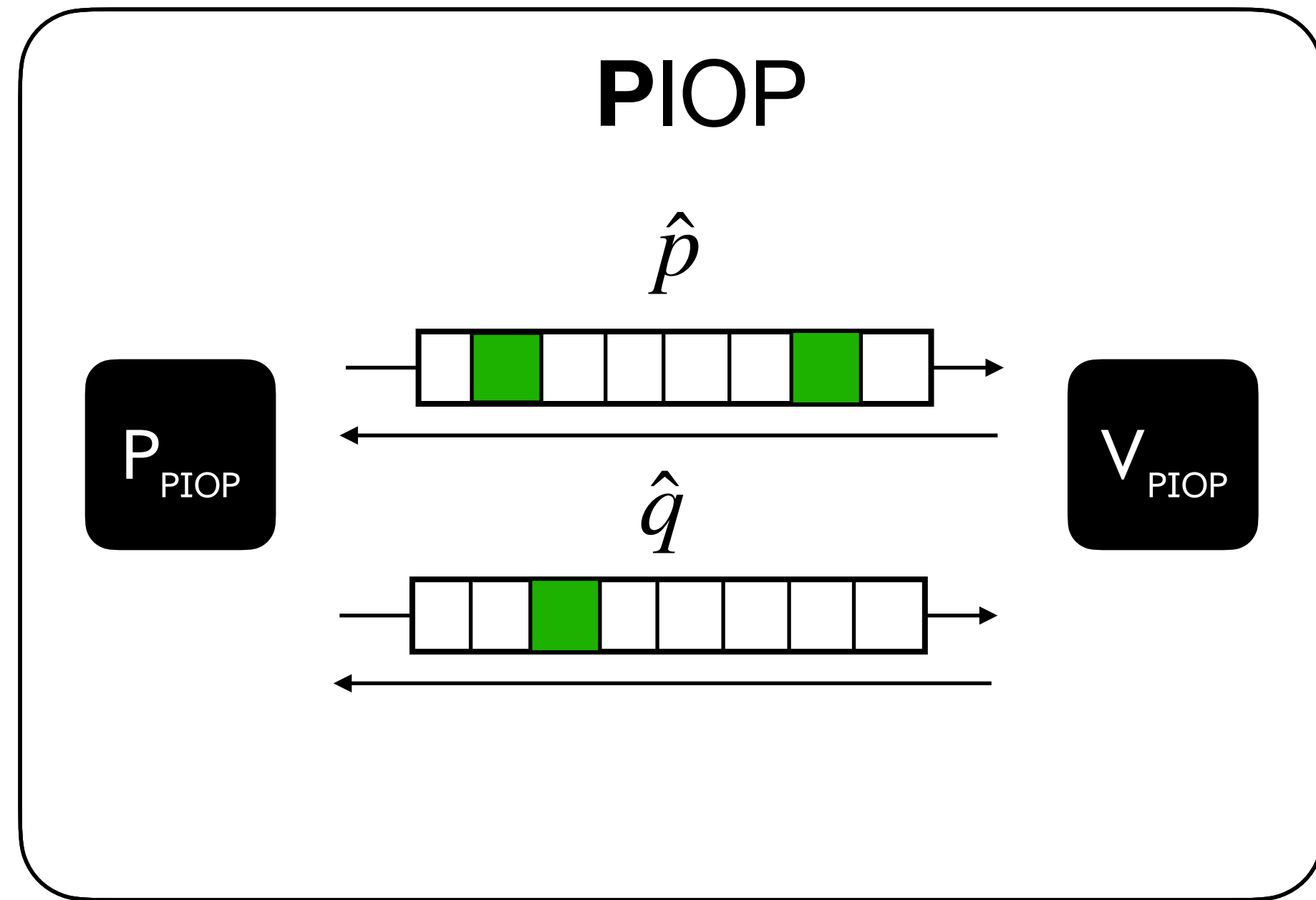
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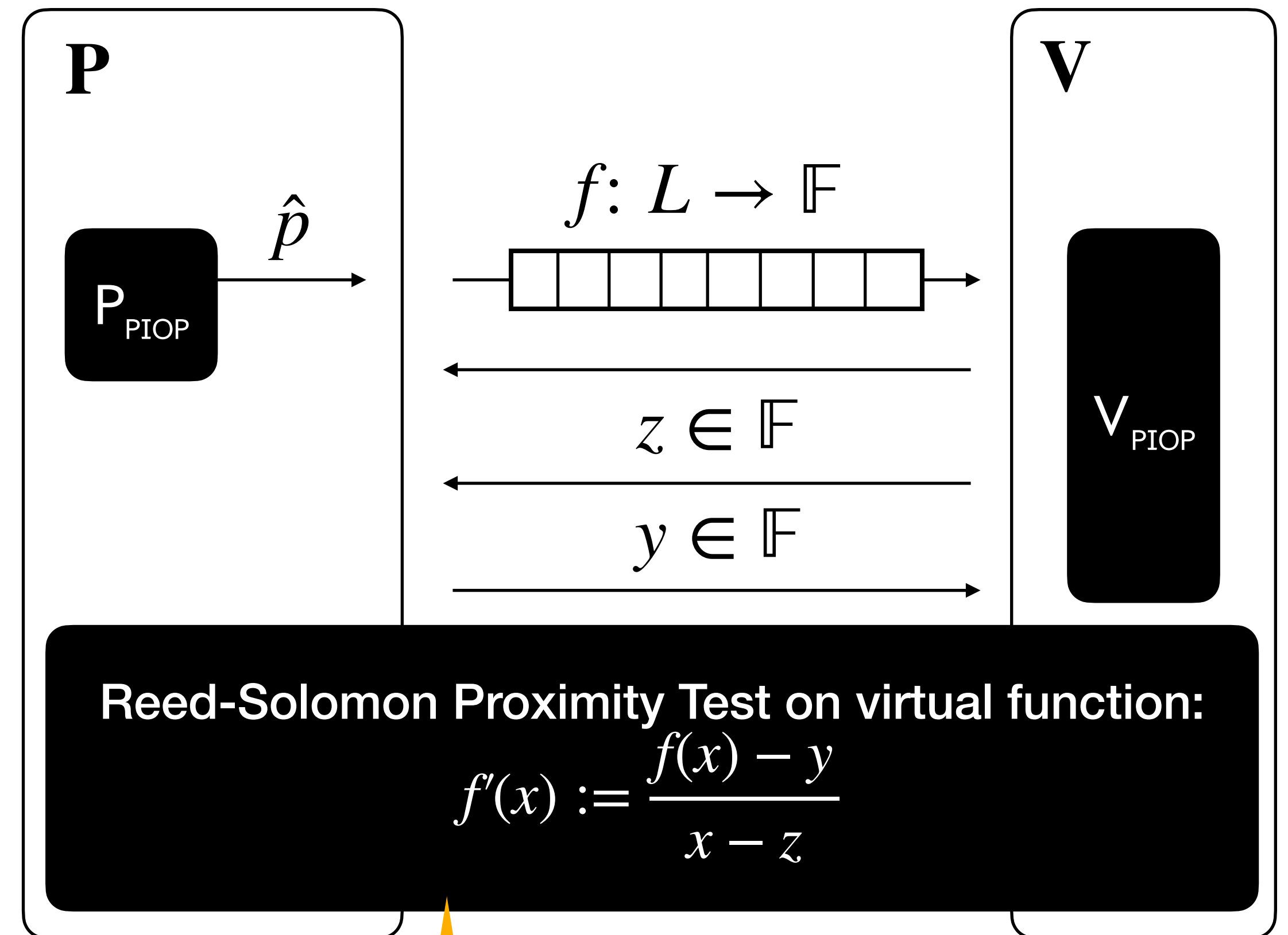
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> 80% of argument size from proximity test!



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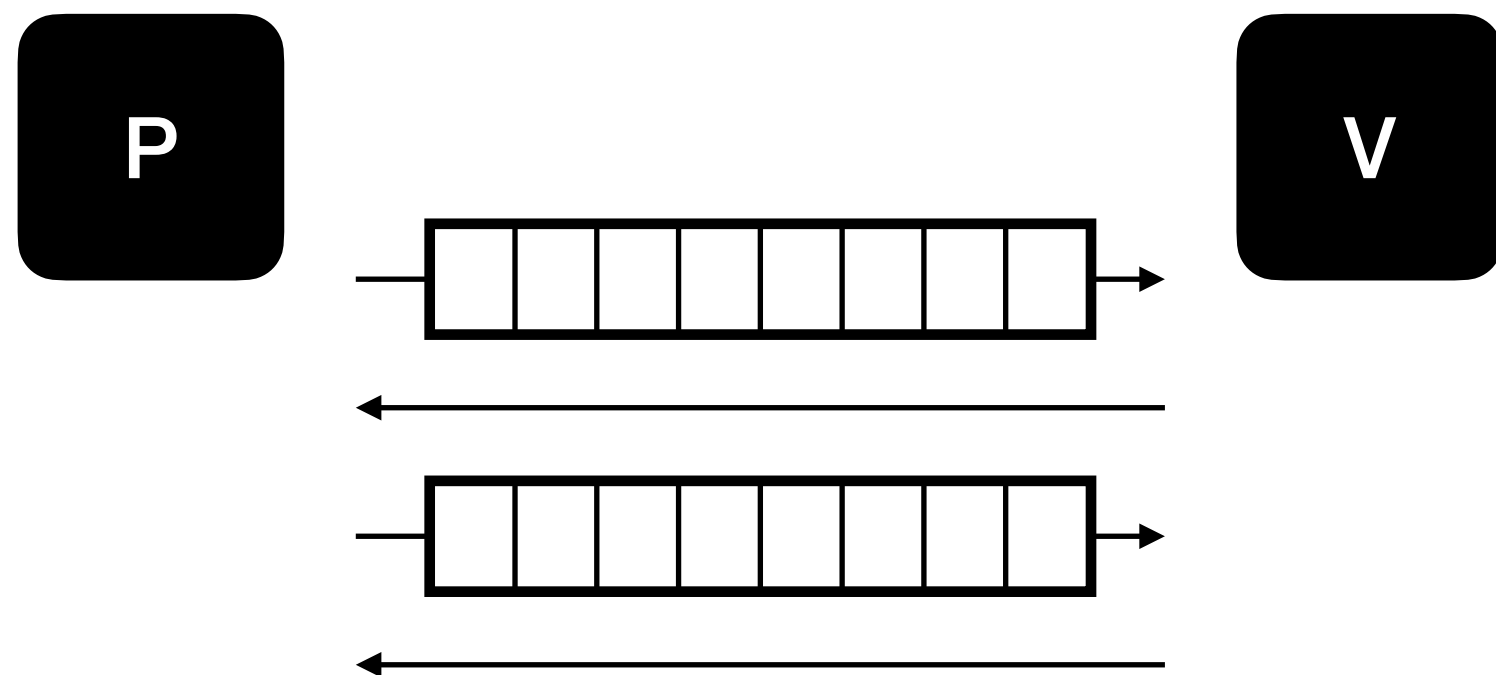
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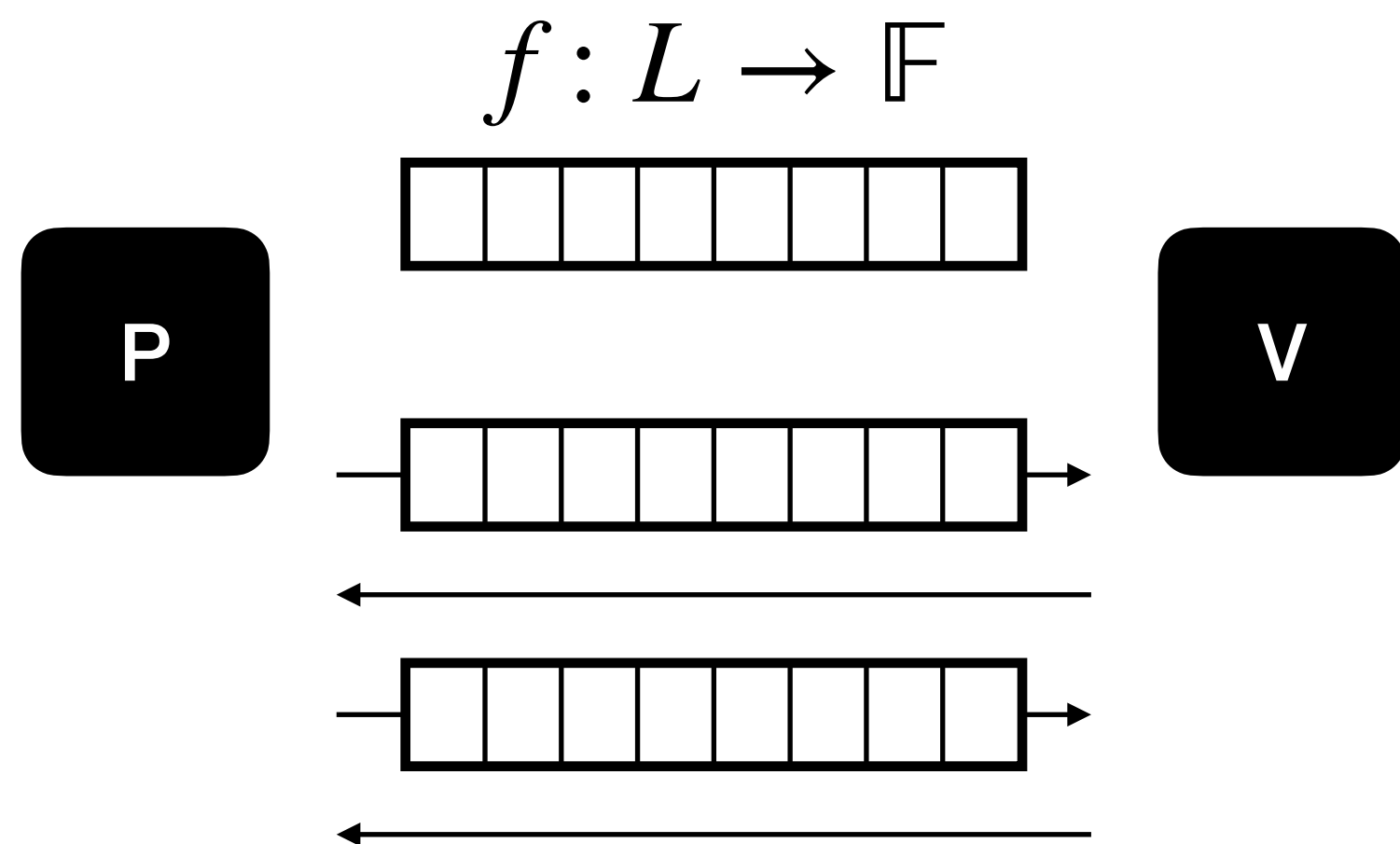
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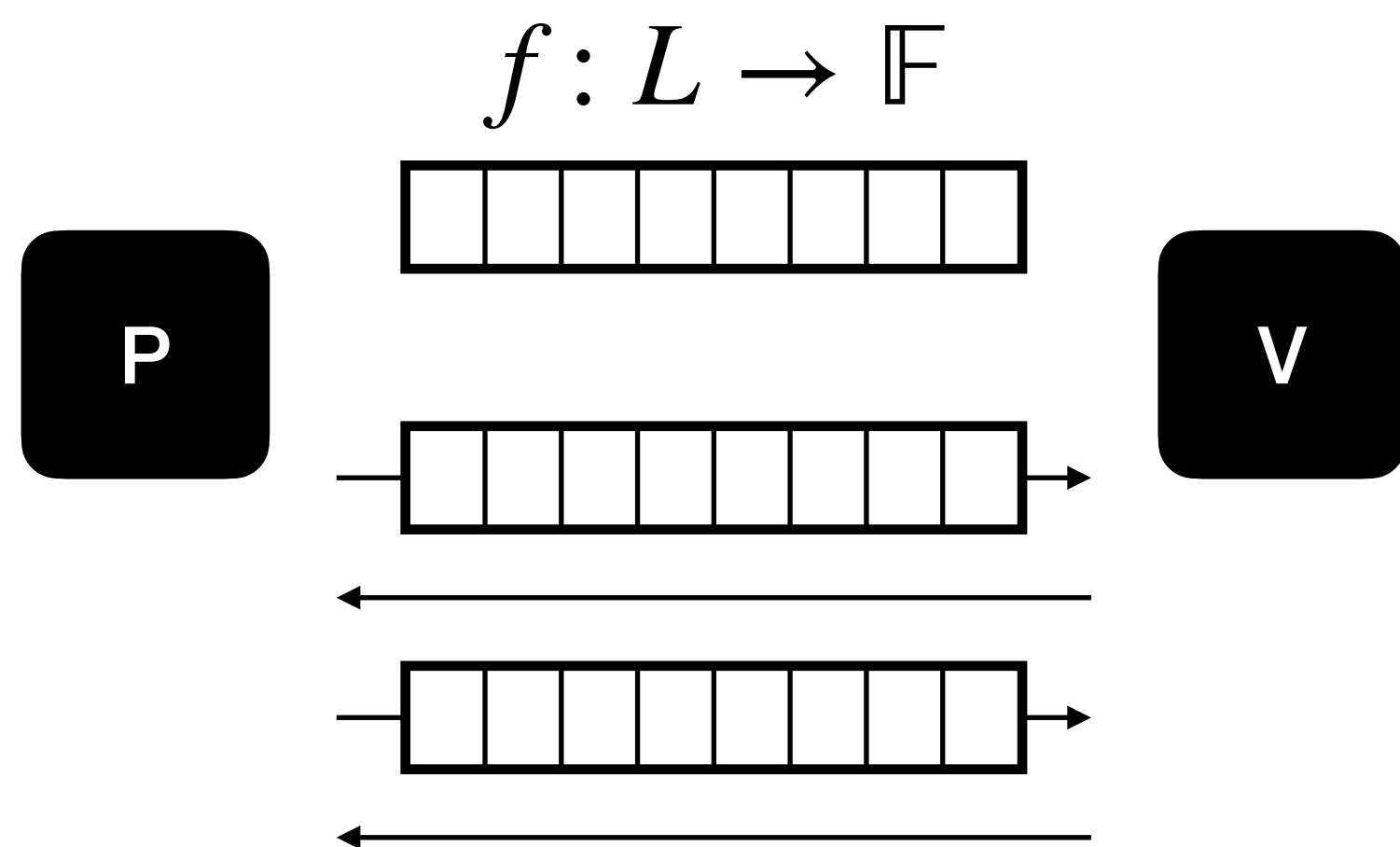
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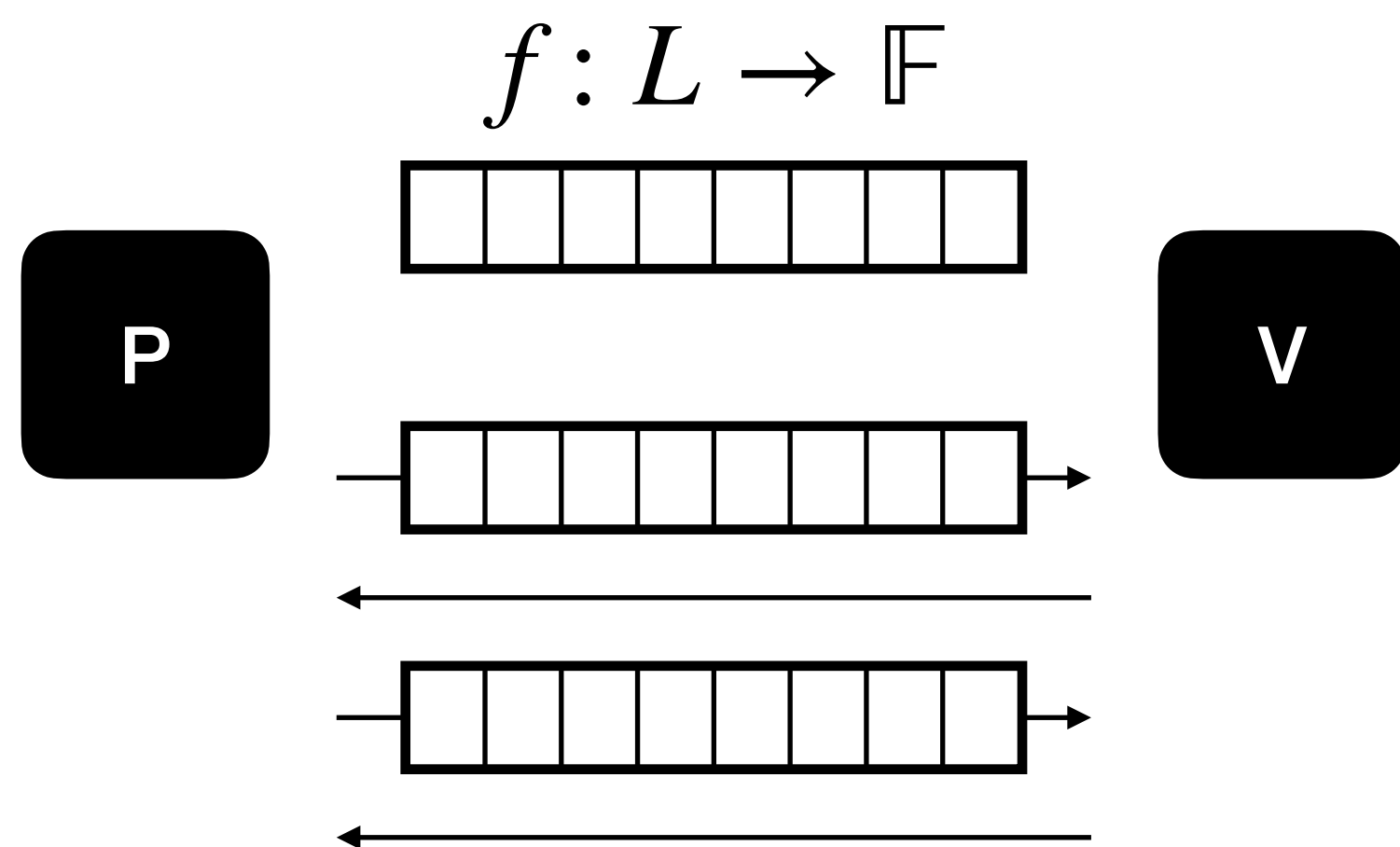
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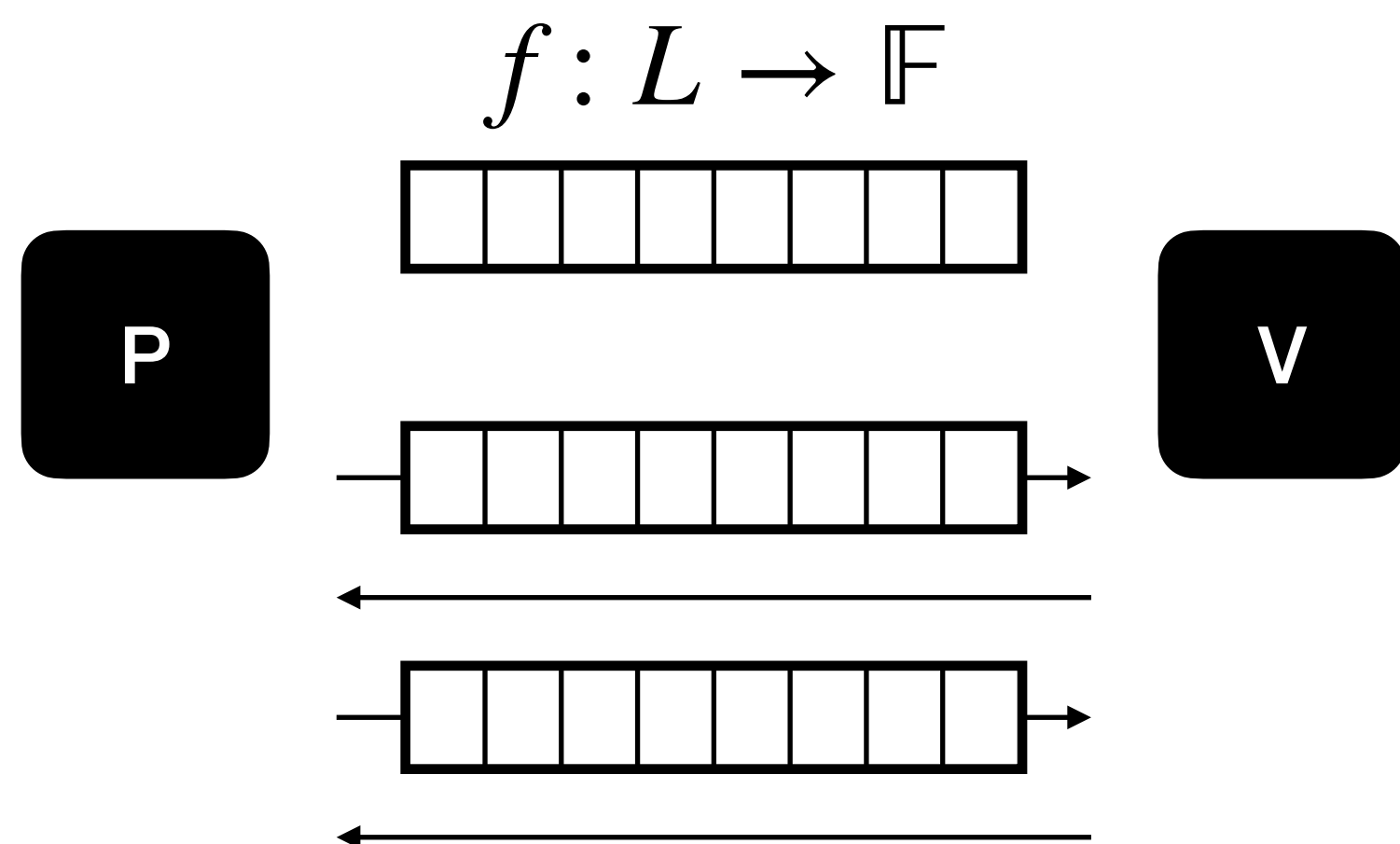
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Round by round, required by BCS transform.

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Can we move the constraint directly into the IOPP?

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$\text{RS}[n, m, \rho] = \text{CRS}[n, m, \rho, 0, 0]$

If  $\hat{w} = Z \cdot \text{eq}(\mathbf{X}, \mathbf{r})$  we recover multilinear polynomial evaluation

**Our results**

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## A constrained Reed-Solomon proximity test



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$k \approx \log m$

$\lambda \gg m$

# Comparison with prior work

	Queries	Verifier Time	Alphabet
<b>BaseFold</b>	$q_{\text{BF}} = O(\lambda \cdot m)$	$O(q_{\text{BF}})$	$\mathbb{F}^2$
<b>FRI</b>	$q_{\text{FRI}} = O\left(\frac{\lambda}{k} \cdot m\right)$	$O(q_{\text{FRI}} \cdot 2^k)$	$\mathbb{F}^{2^k}$
<b>STIR</b>	$q_{\text{STIR}} = O\left(\frac{\lambda}{k} \cdot \log m\right)$	$O(q_{\text{STIR}} \cdot 2^k + \lambda^2 \cdot 2^k)$	$\mathbb{F}^{2^k}$
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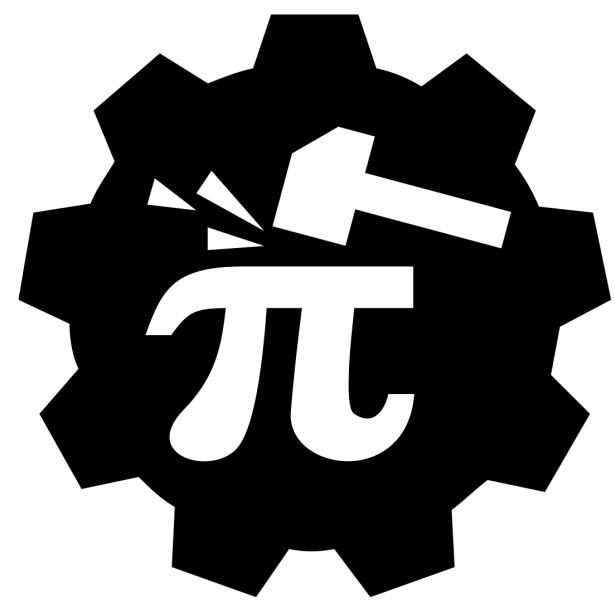
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- **Further**, super-fast verification (next)

# Implementation



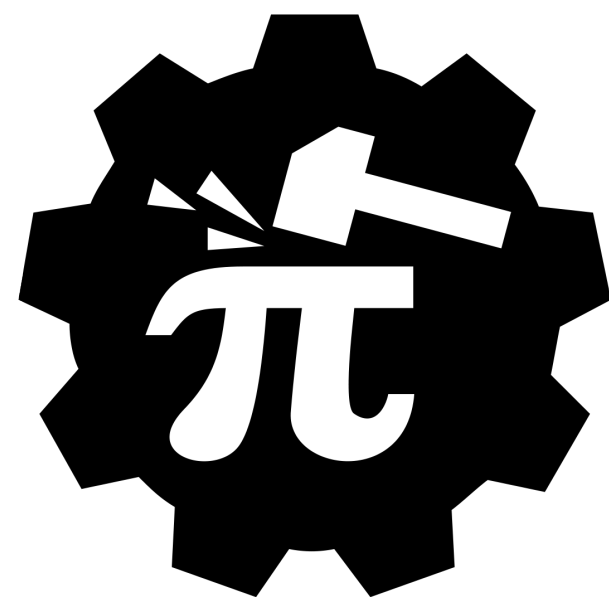
```
=====
Whir (PCS)
Field: Goldilocks2 and MT: Blake3
Number of variables: 20, folding factor: 4
Security level: 100 bits using ConjectureList security and 19 bits of PoW
initial_folding_pow_bits: 0
Num_queries: 41, rate: 2^-2, pow_bits: 18, ood_samples: 2, folding_pow: 0
Num_queries: 17, rate: 2^-5, pow_bits: 15, ood_samples: 2, folding_pow: 2
Num_queries: 11, rate: 2^-8, pow_bits: 12, ood_samples: 2, folding_pow: 4
Num_queries: 8, rate: 2^-11, pow_bits: 12, ood_samples: 2, folding_pow: 6
final_queries: 6, final_rate: 2^-14, final_pow_bits: 16, final_folding_pow_bits: 0
-----
Round by round soundness analysis:
-----
167.0 bits -- OOD commitment
102.0 bits -- (x4) prox gaps: 103.0, sumcheck: 102.0, pow: 0.0
171.0 bits -- OOD sample
100.0 bits -- query error: 82.0, combination: 94.6, pow: 18.0
100.0 bits -- (x4) prox gaps: 101.0, sumcheck: 100.0, pow: 0.0
175.0 bits -- OOD sample
100.0 bits -- query error: 85.0, combination: 93.8, pow: 15.0
100.0 bits -- (x4) prox gaps: 99.0, sumcheck: 98.0, pow: 2.0
179.0 bits -- OOD sample
100.0 bits -- query error: 88.0, combination: 92.3, pow: 12.0
100.0 bits -- (x4) prox gaps: 97.0, sumcheck: 96.0, pow: 4.0
183.0 bits -- OOD sample
100.0 bits -- query error: 88.0, combination: 90.7, pow: 12.0
100.0 bits -- (x4) prox gaps: 95.0, sumcheck: 94.0, pow: 6.0
100.0 bits -- query error: 84.0, pow: 16.0

Prover time: 356.9ms
Proof size: 58.7 KiB
Verifier time: 342.8µs
Average hashes: 1.1k
```

# Implementation



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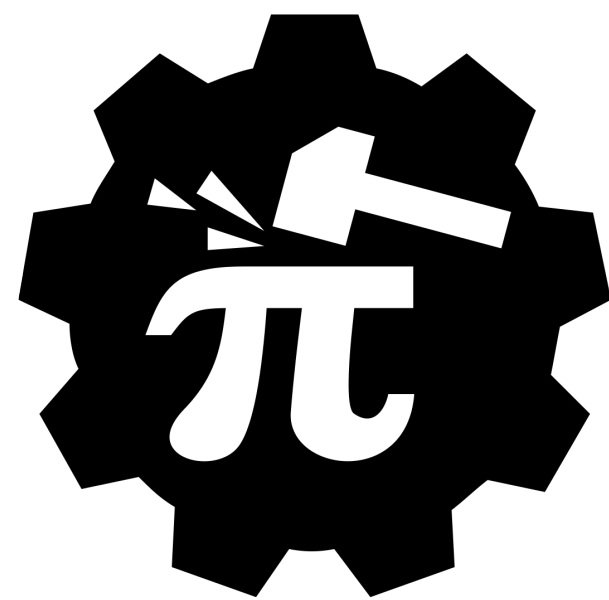
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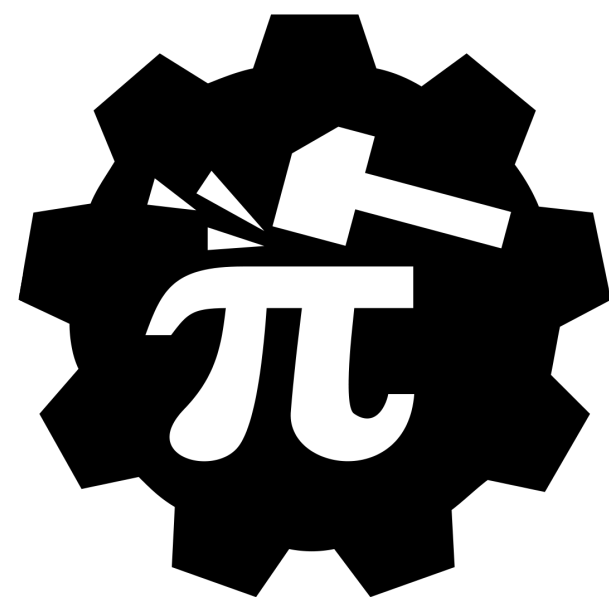
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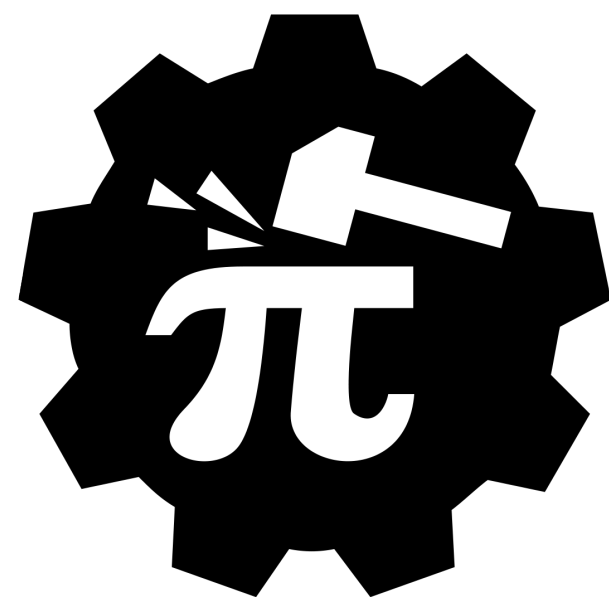
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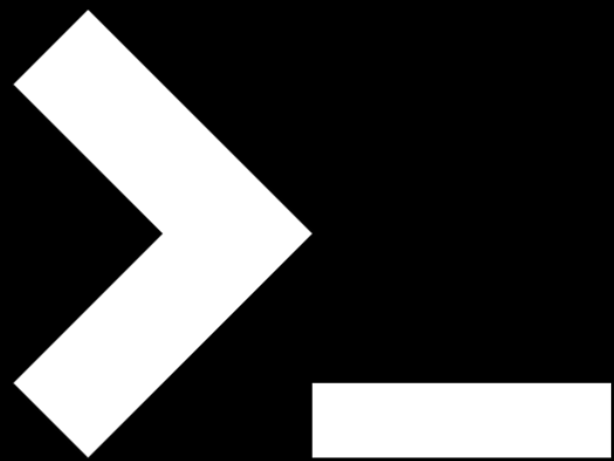
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```
Prover time: ~1s (MacBook Air)
Commit & open: 63 KiB
Verifier time: 270 μs (0.27 ms)
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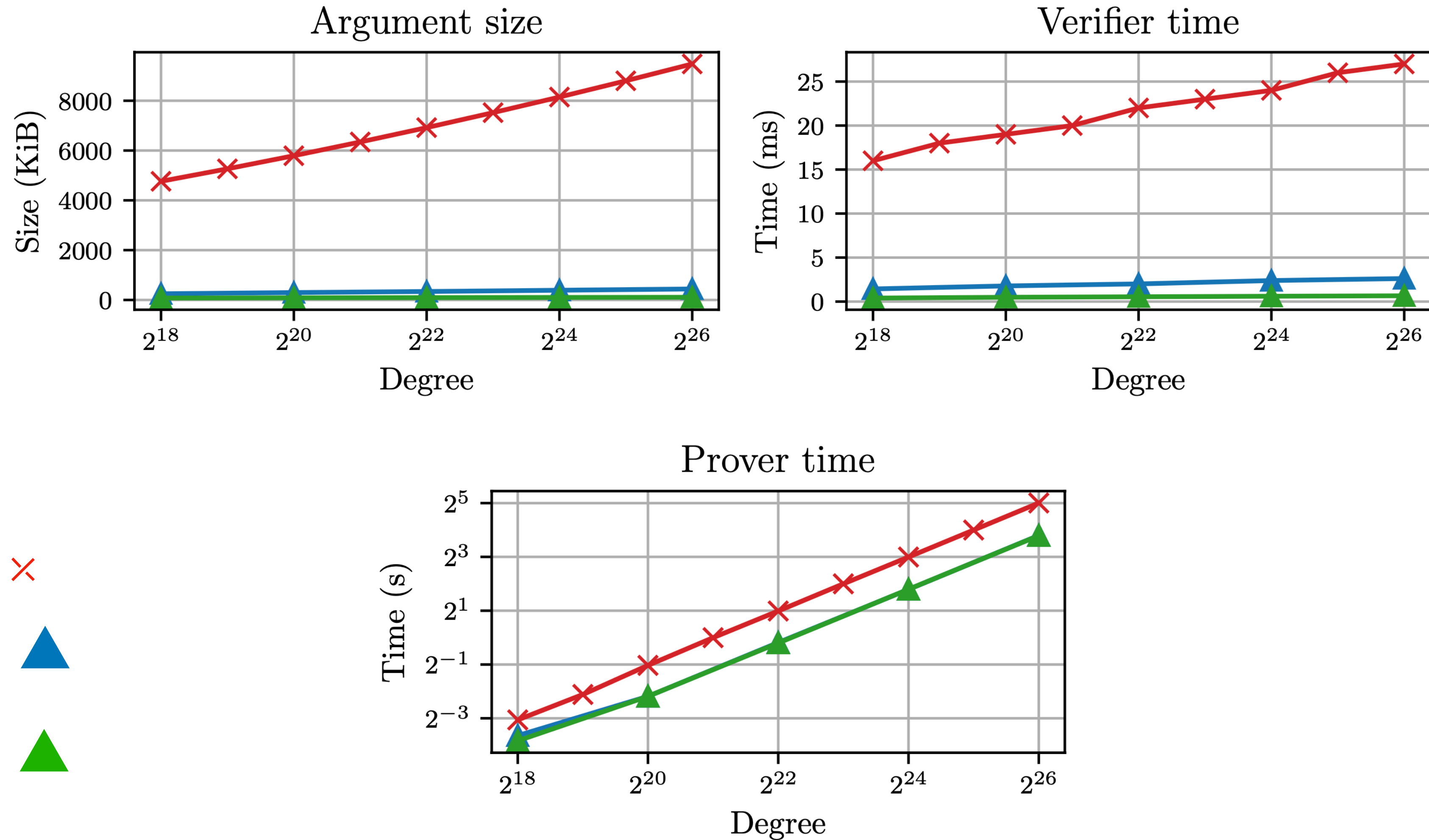
Schemes with trusted setup using pairings!

Verifier time (ms)	Brakedown	Ligero	Greyhound	Hyrax	PST	KZG	WHIR- <sup>1</sup> / <sub>2</sub>	WHIR- <sup>1</sup> / <sub>16</sub>
$\lambda = 100$	3500	733	-	100	7.81	2.42	0.61	0.29
$\lambda = 128$	3680	750	130	151	9.92	3.66	1.4	0.6

**Table 4:** Comparison of WHIR-CB's verifier time versus other polynomial commitment schemes, on 24 variables. For the KZG degree  $2^{24}$  is used instead.

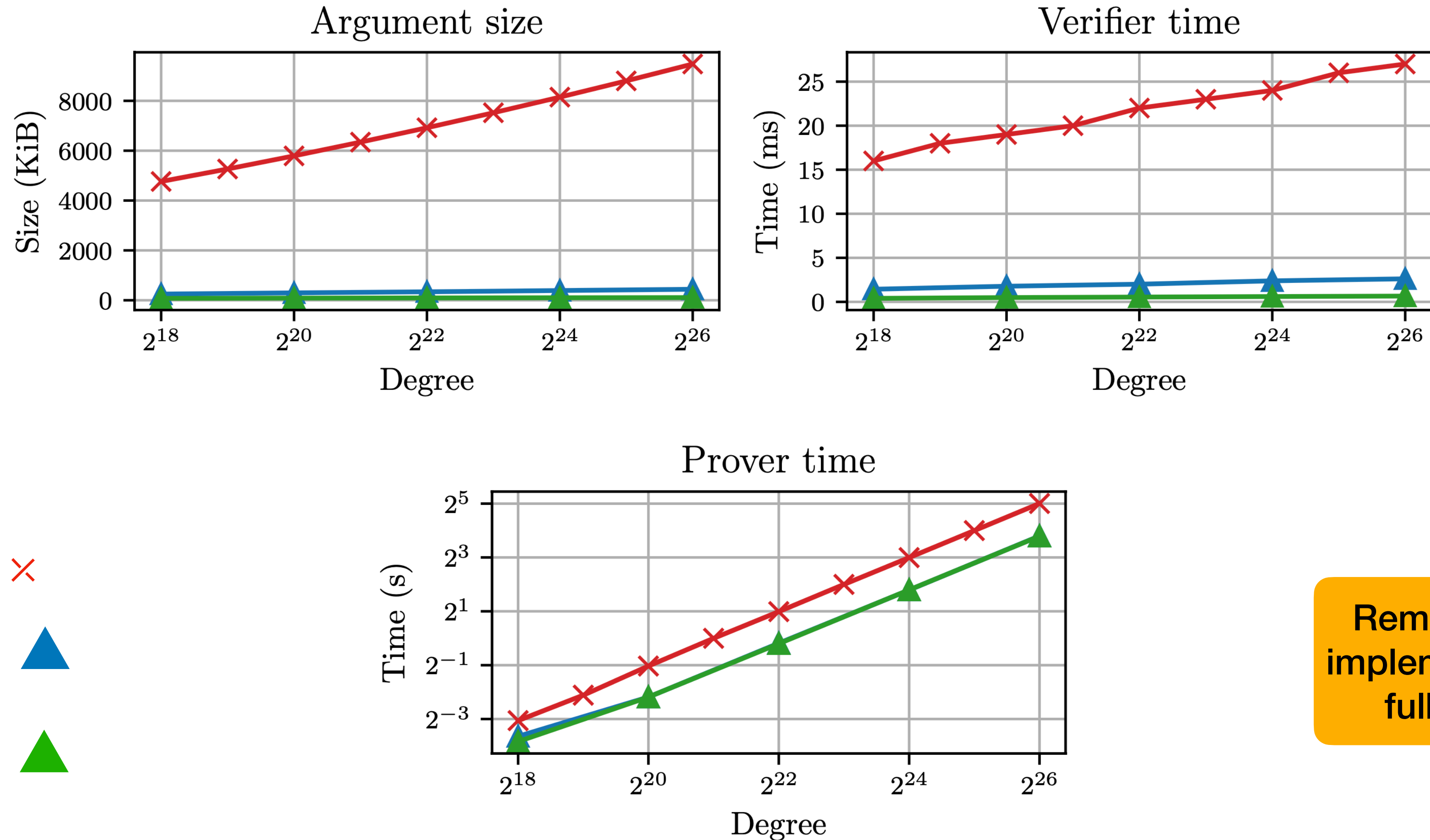


# Comparison with BaseFold



BaseFold: x  
WHIR-UD: ▲  
WHIR-CB: ▲

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Remark: BaseFold implementation is not fully optimised

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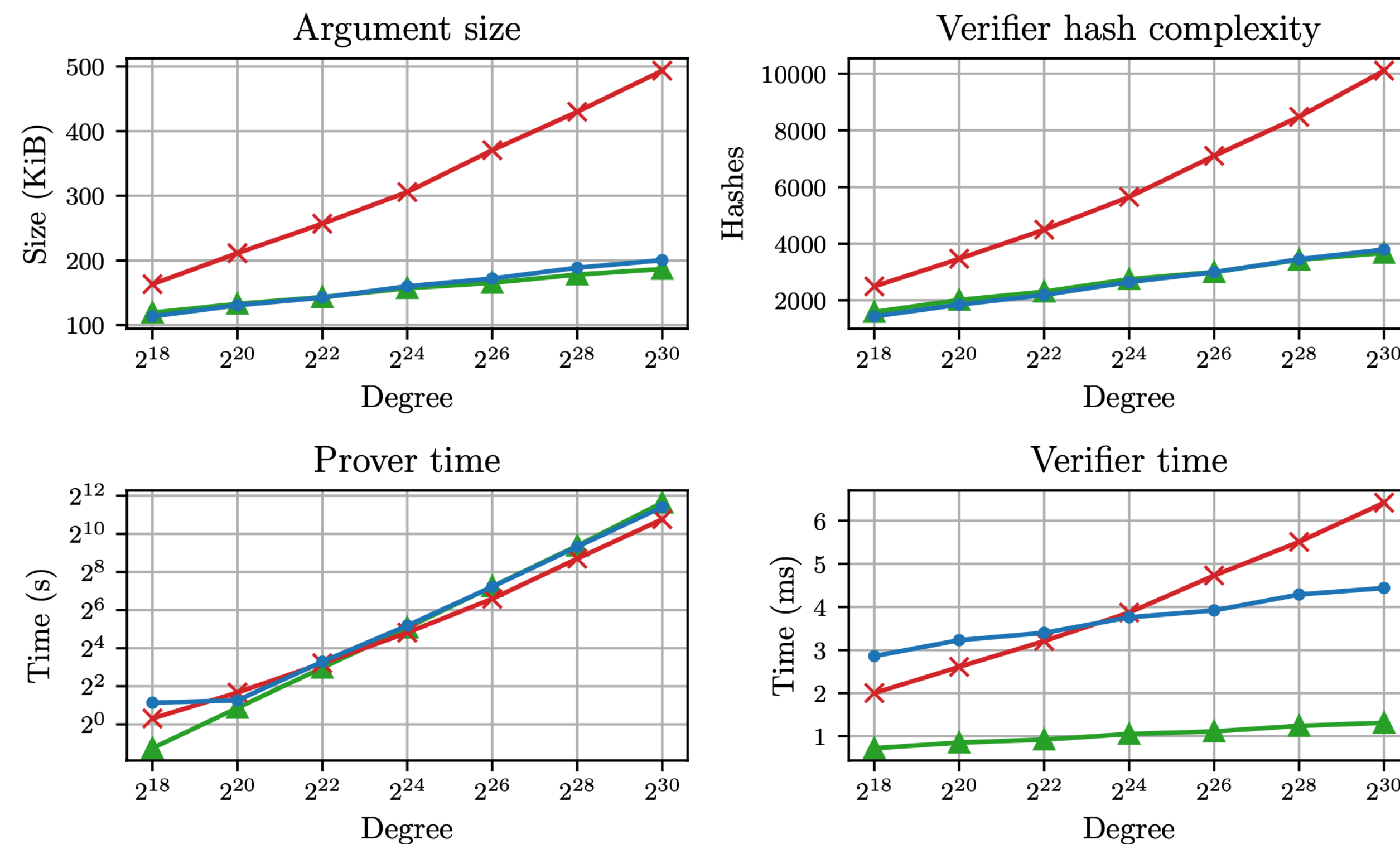
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**Figure 2:** Comparison of FRI, STIR and WHIR for  $\rho = 1/2$ . FRI:  $\times$ , STIR:  $\bullet$ , WHIR-CB:  $\blacktriangle$ . Prover time is displayed with logarithmic scaling.

**Conclusion**

# Summary



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**Extra slides**

# Techniques

# FRI & STIR Folding

**Reduce**  $RS[n, m, \rho]$  to  $RS[n/2^k, m - k, \rho]$

(Think  $k = 4$ )



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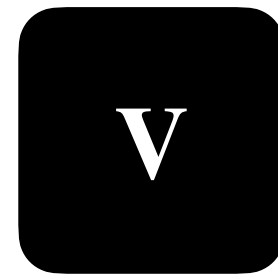
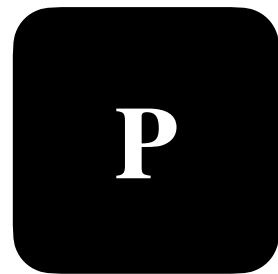
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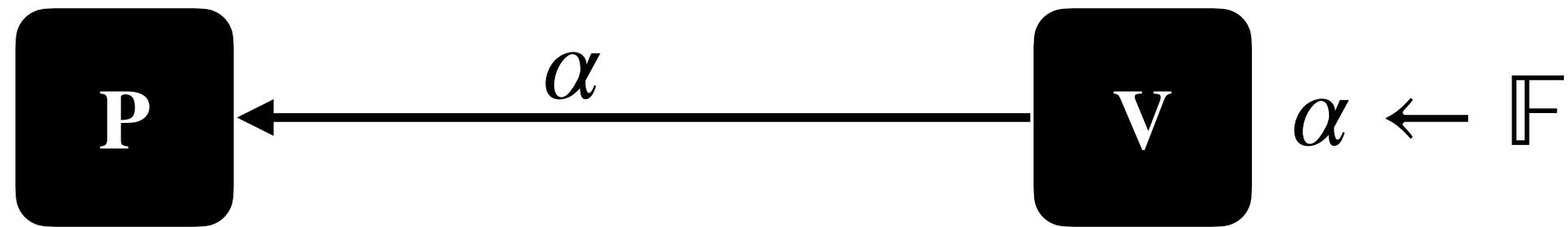
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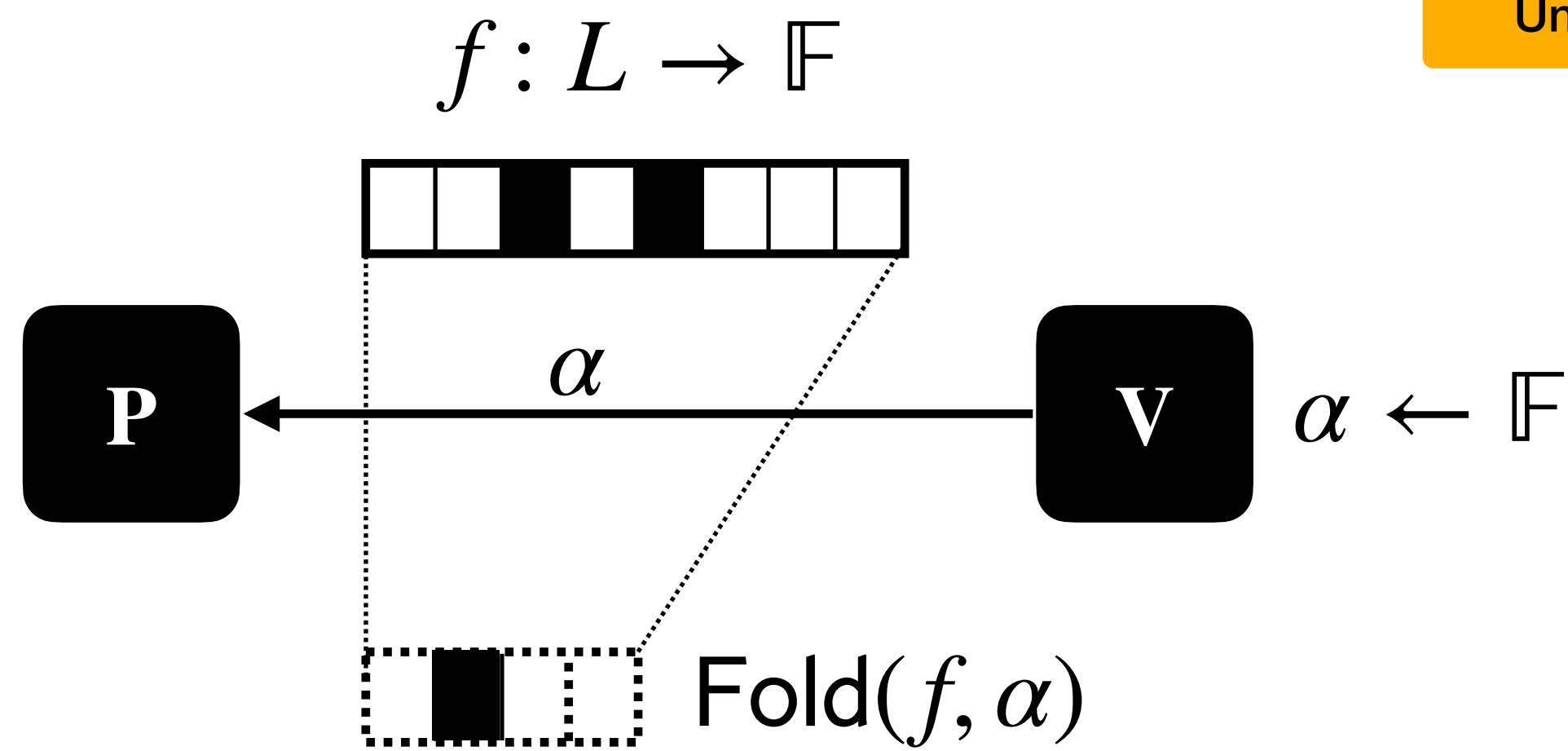


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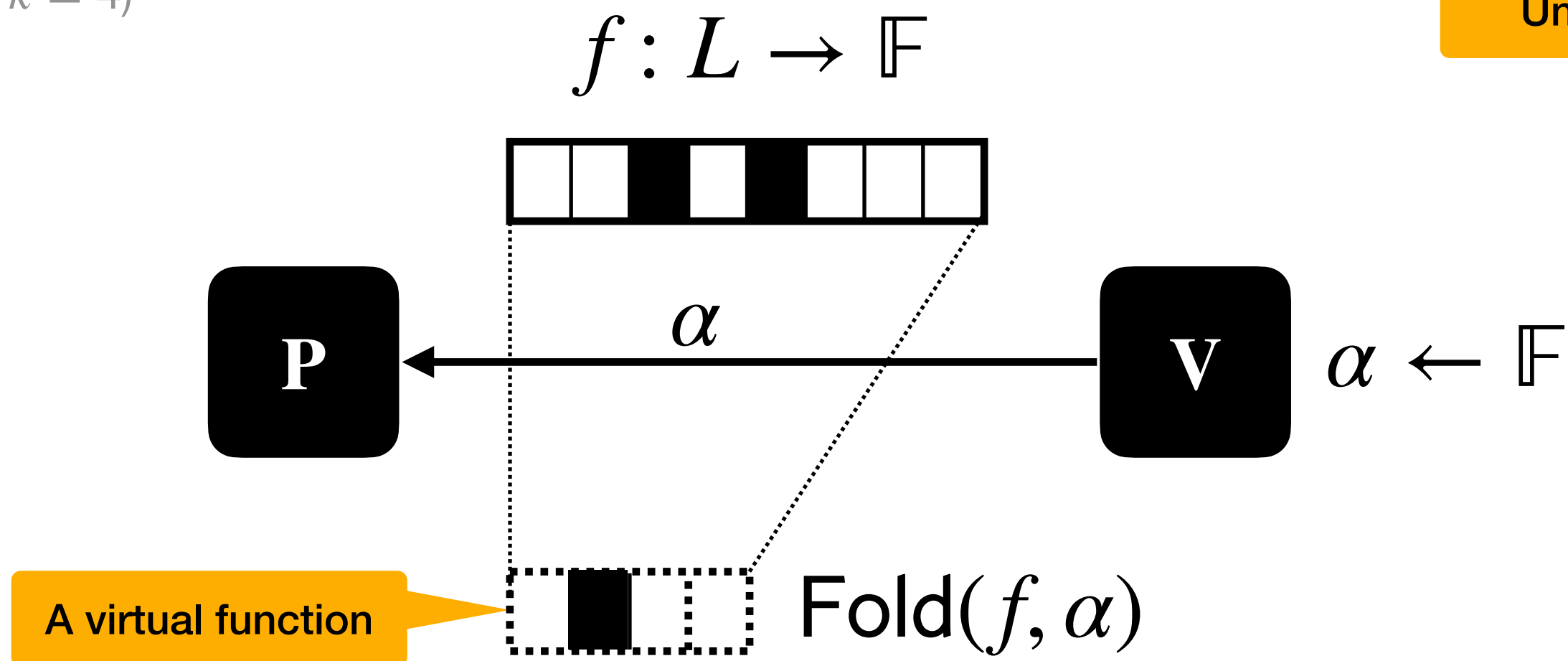
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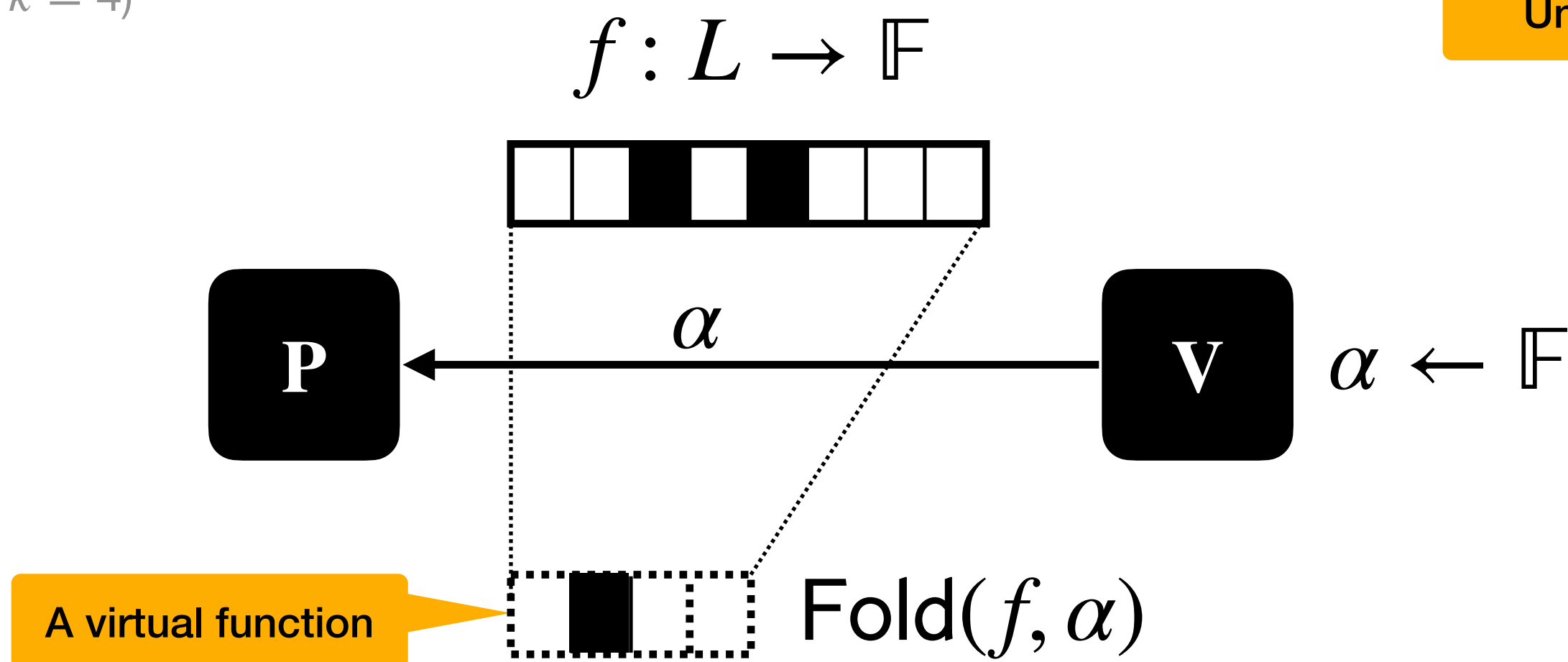


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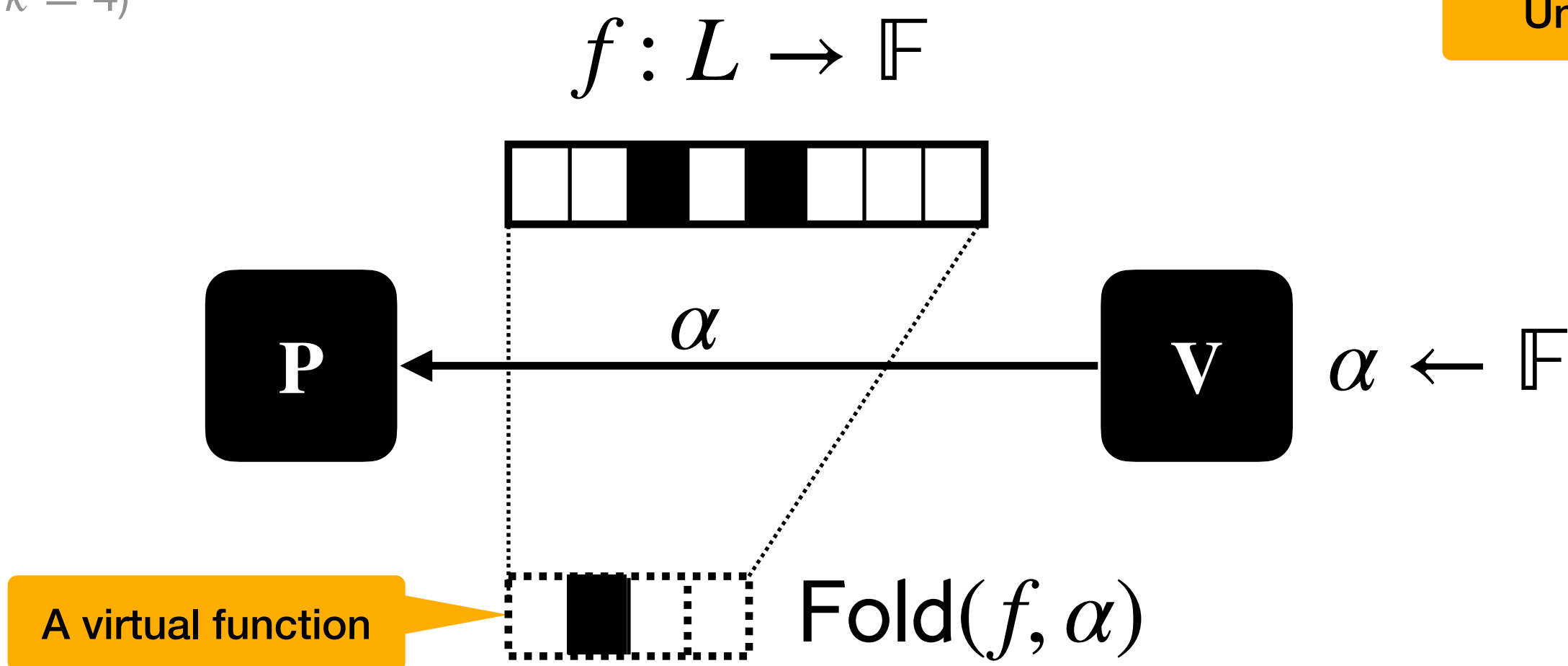
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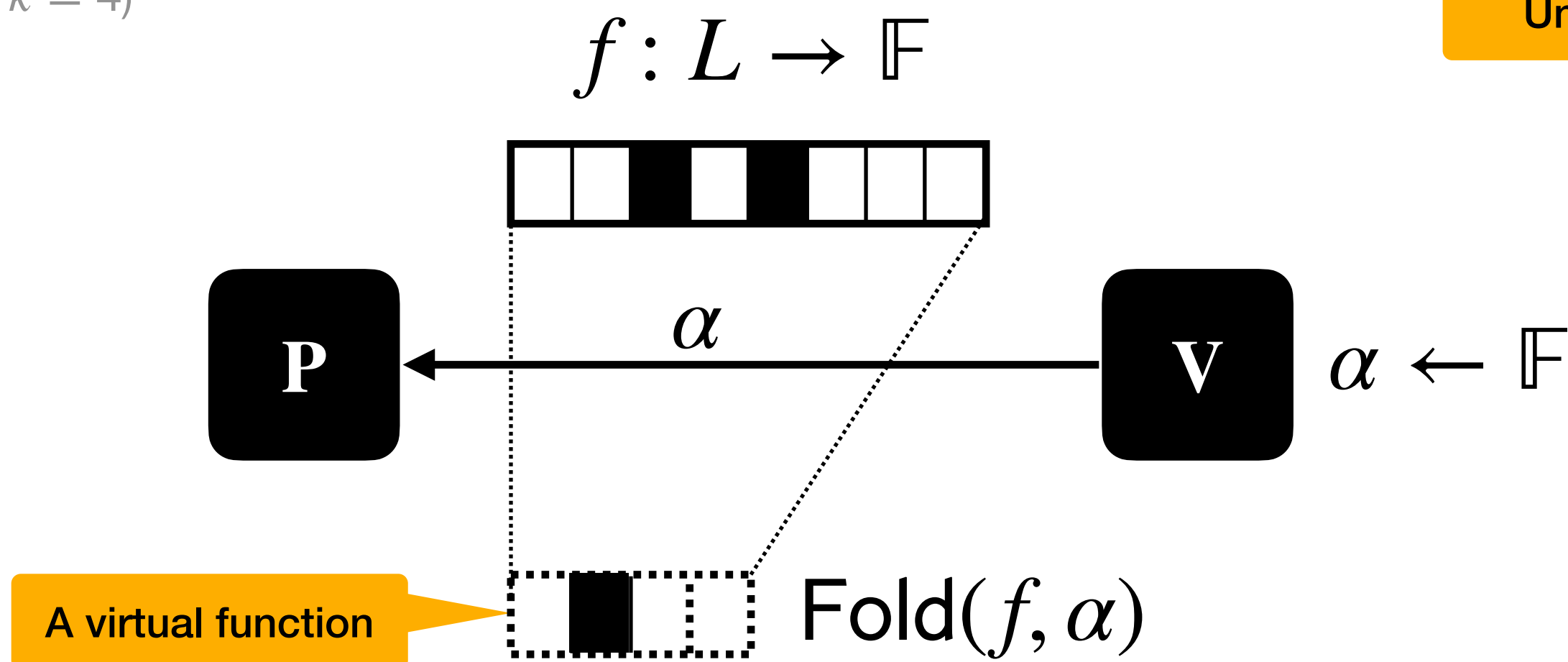
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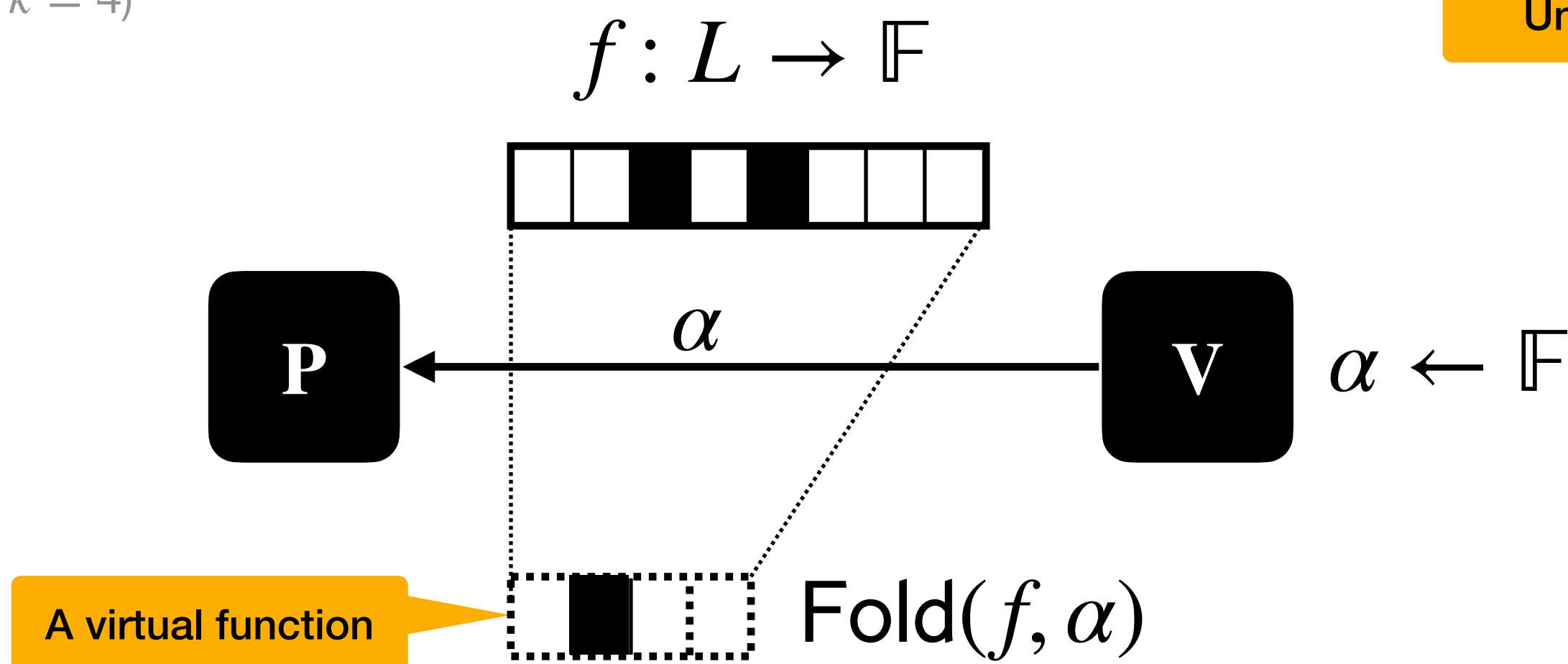
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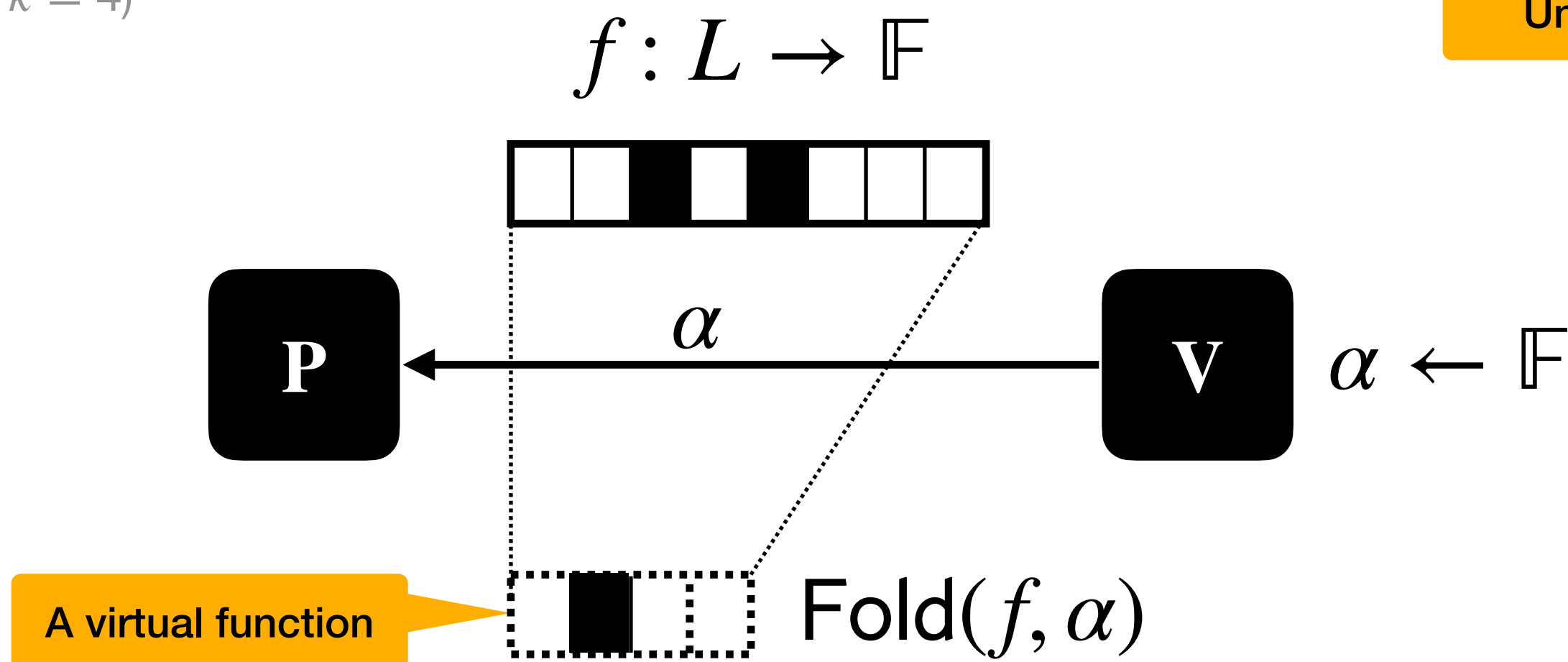
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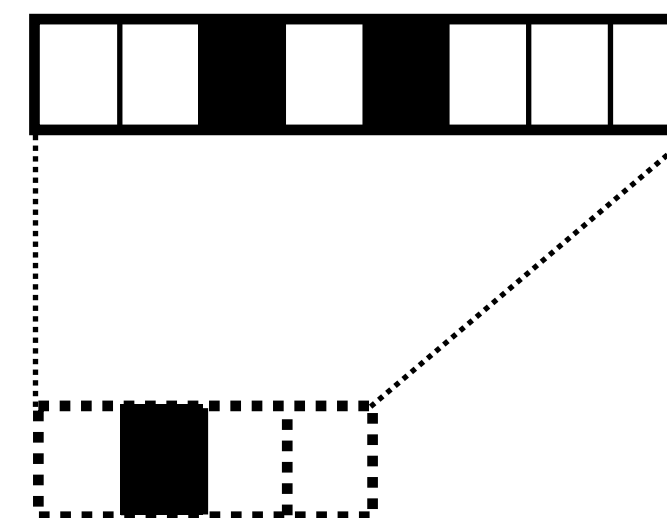
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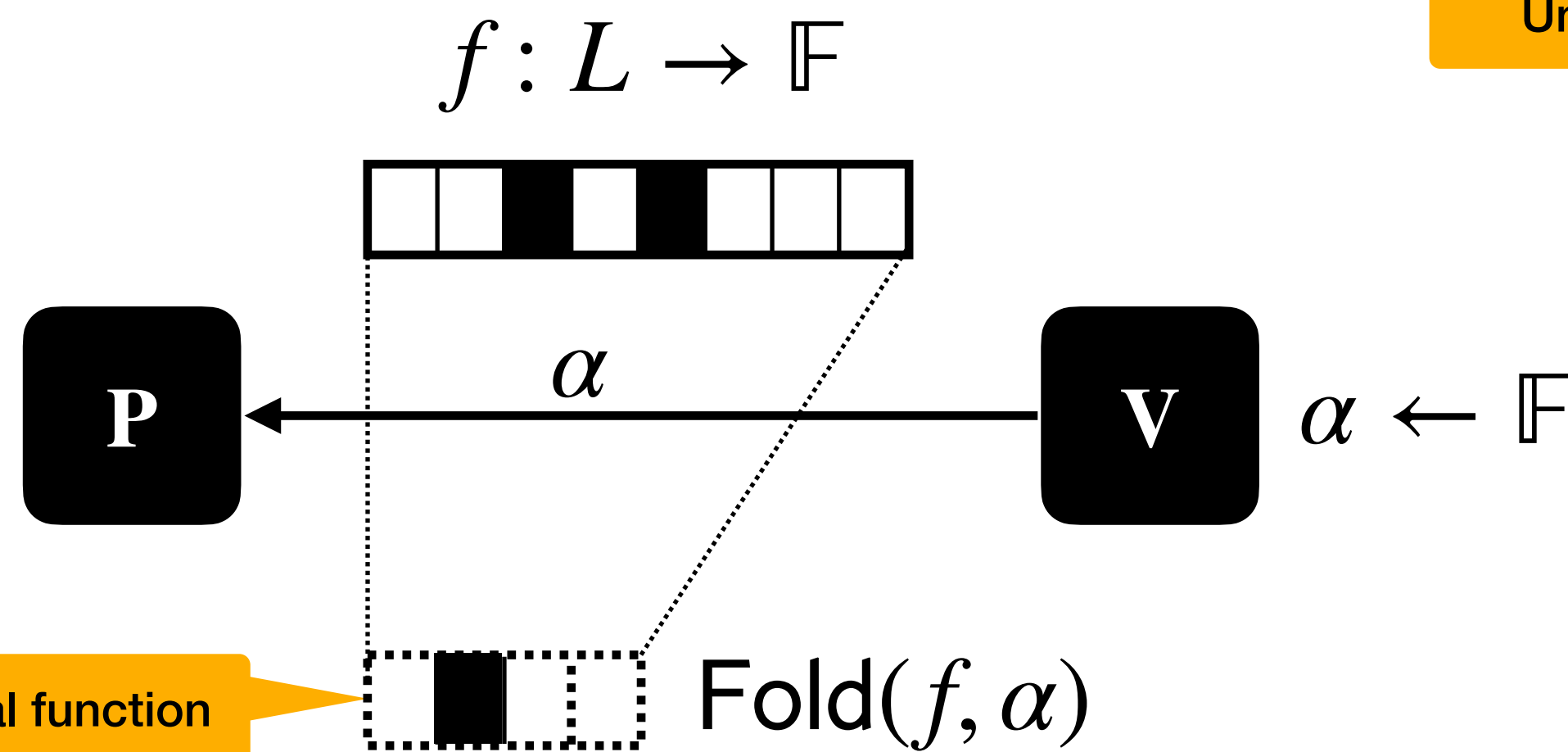


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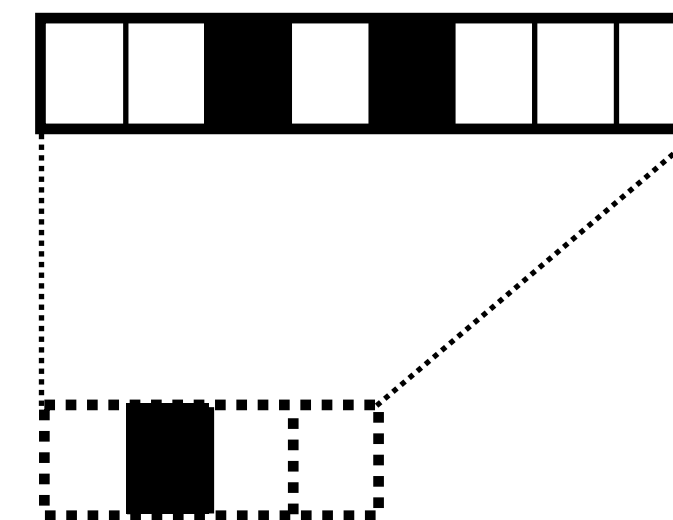
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Proximity Gaps for Reed–Solomon Codes

Eli Ben-Sasson\* Dan Carmon\* Yuval Ishai† Swastik Kopparty ‡

Shubhangi Saraf§

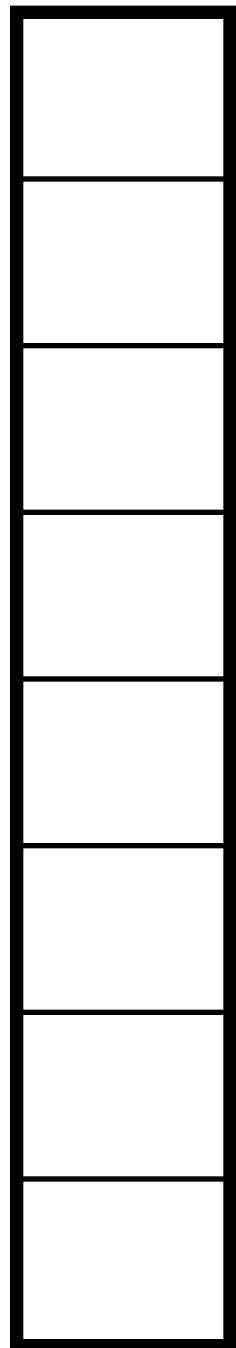
July 3, 2021

# Mutual correlated agreement

Test a random linear combination

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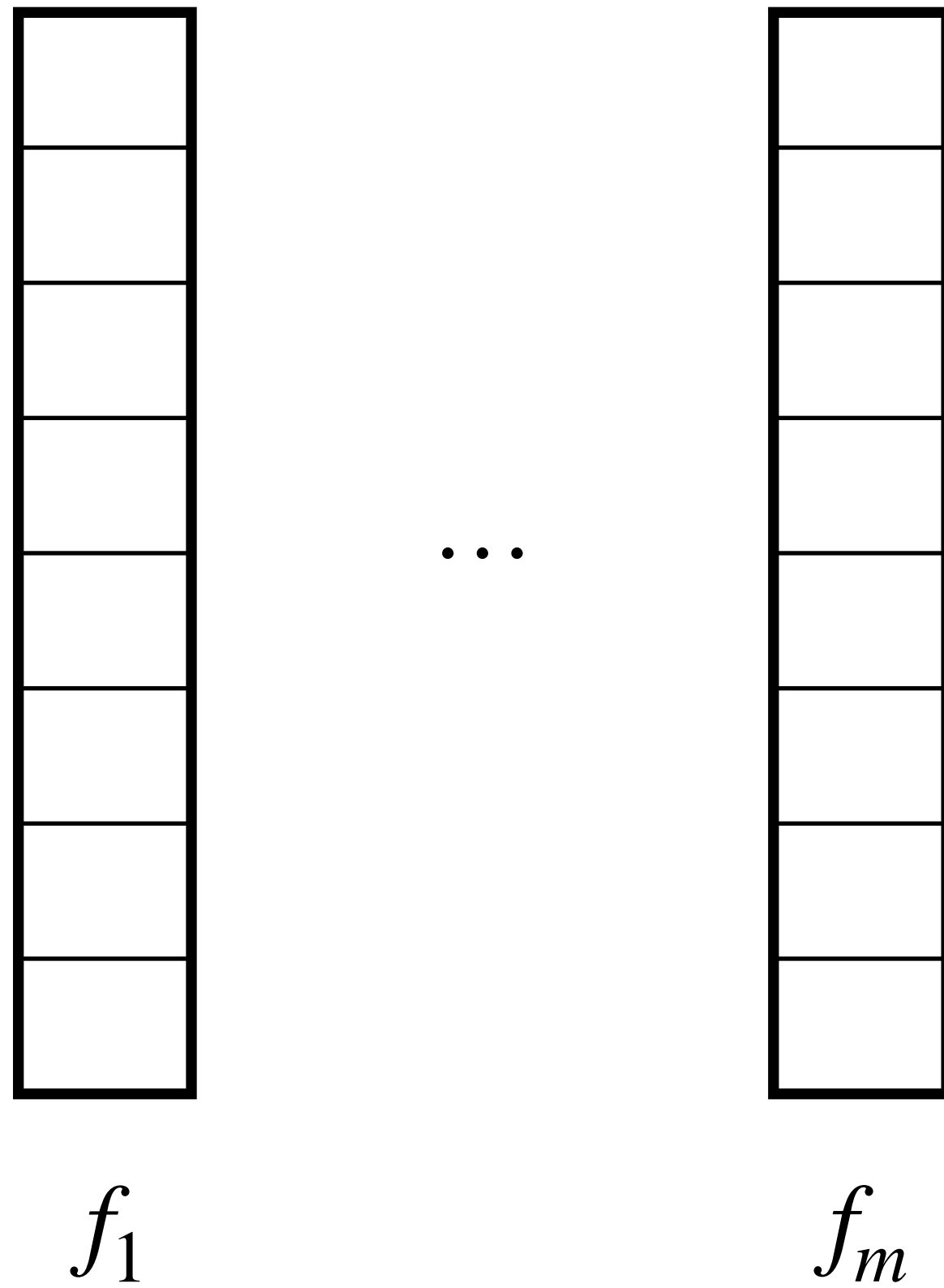
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$f_1$

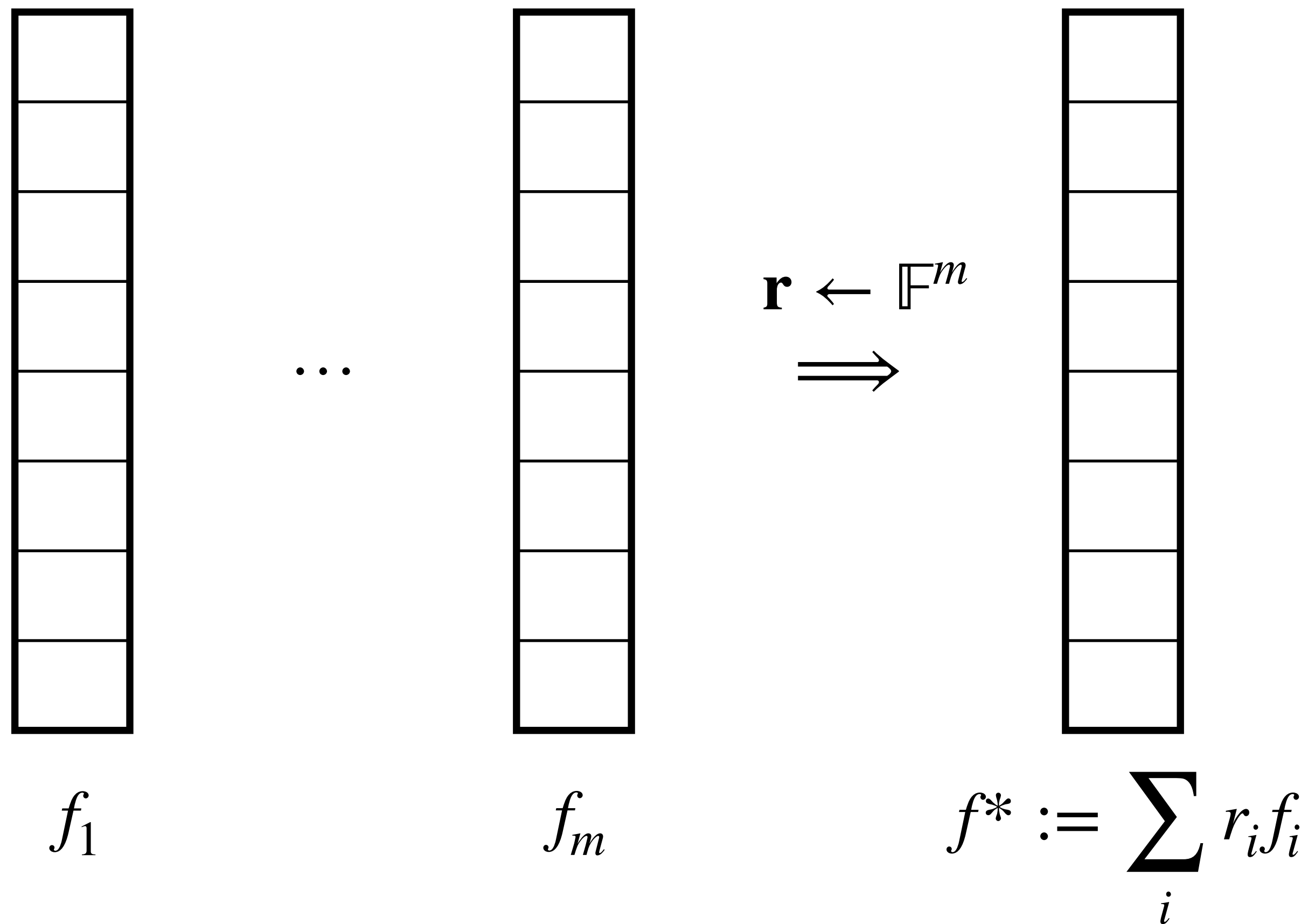
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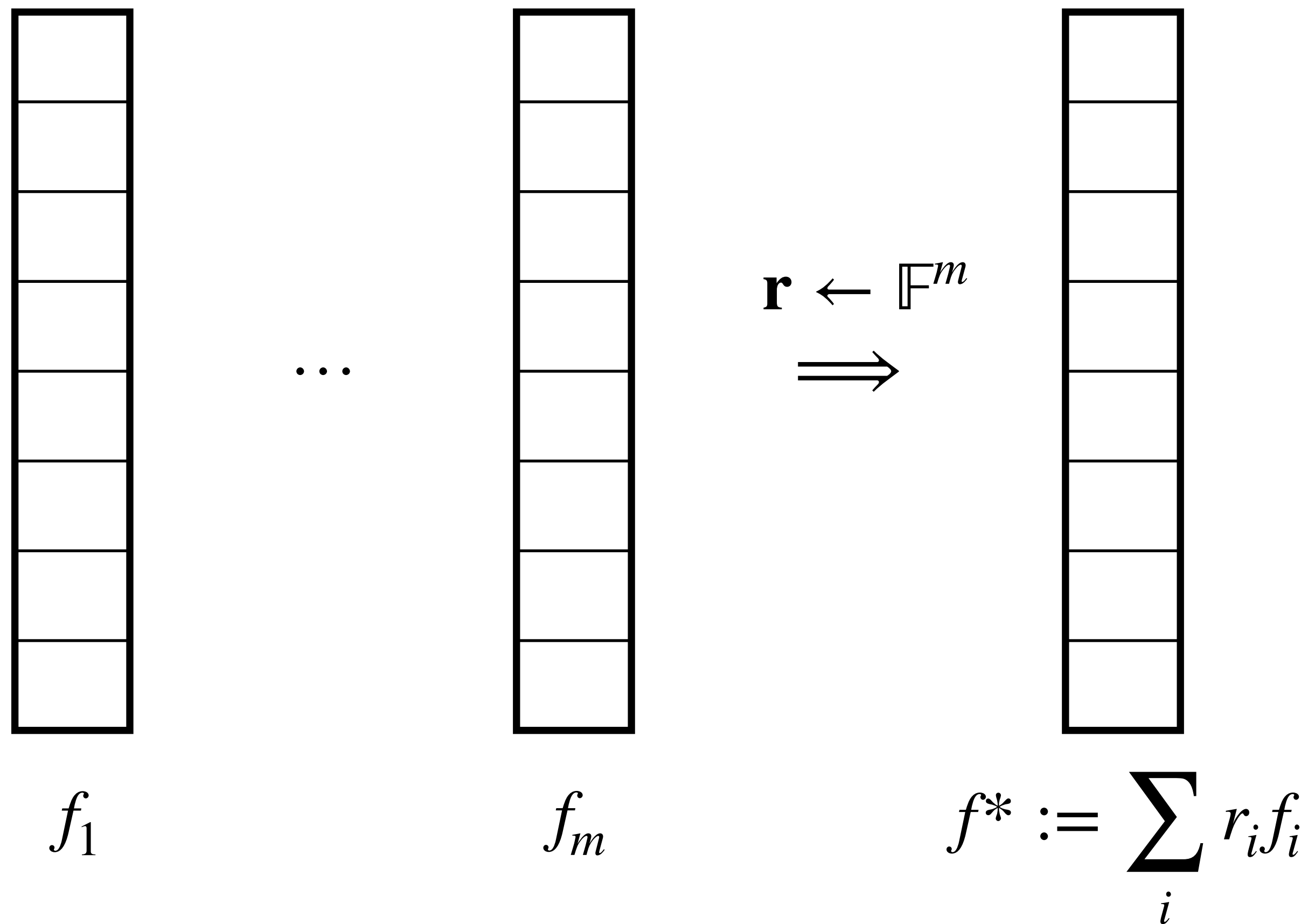
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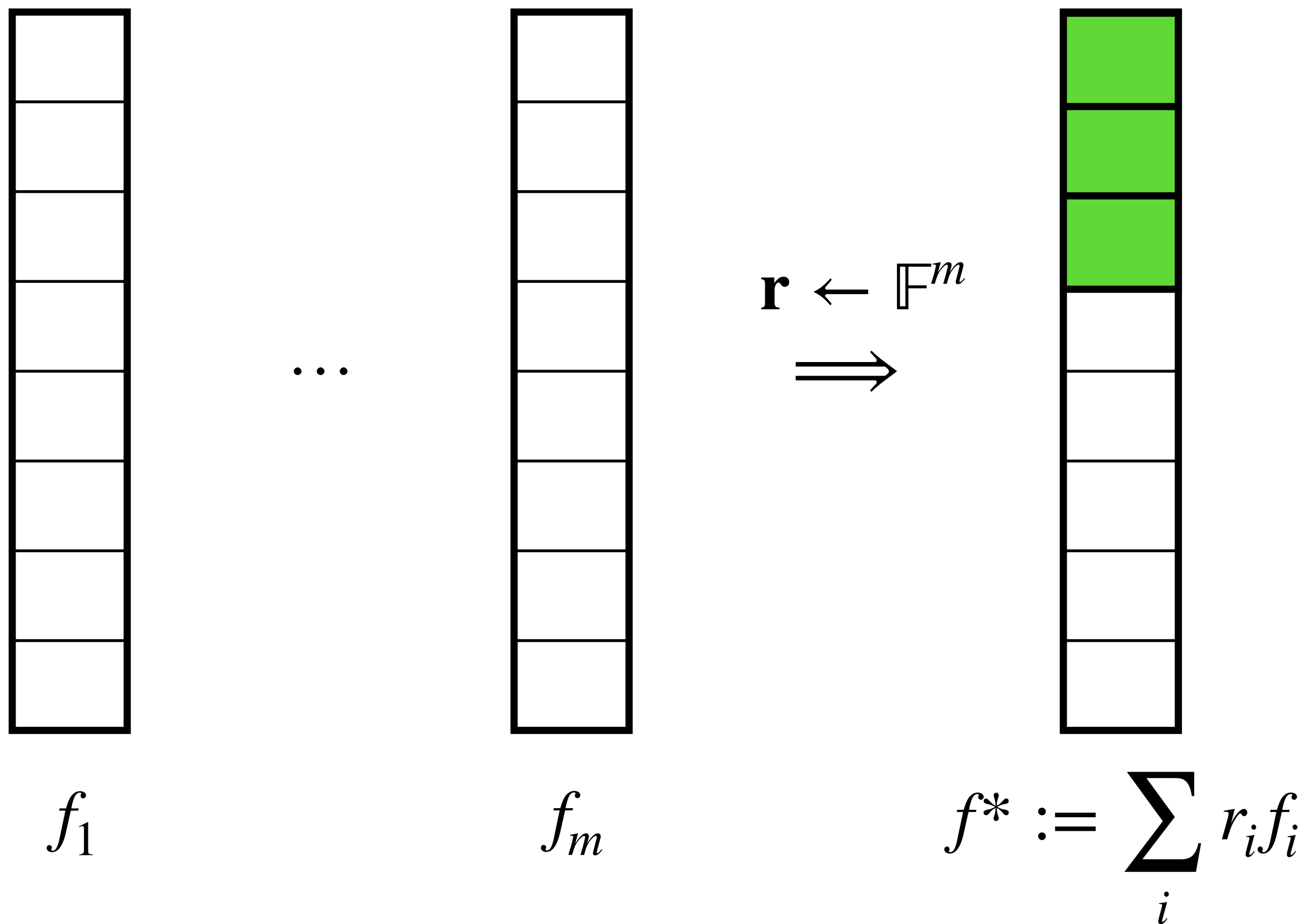




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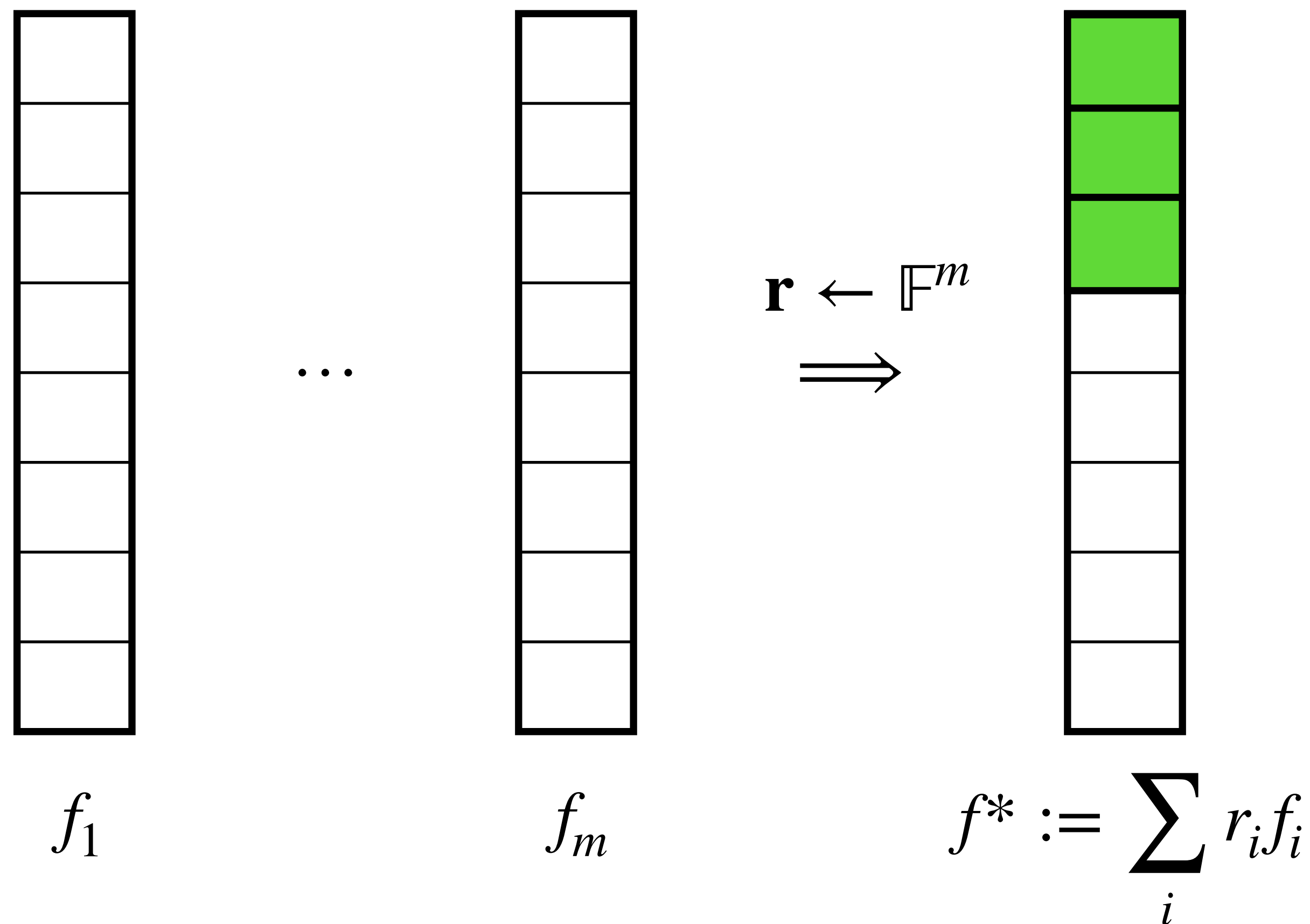
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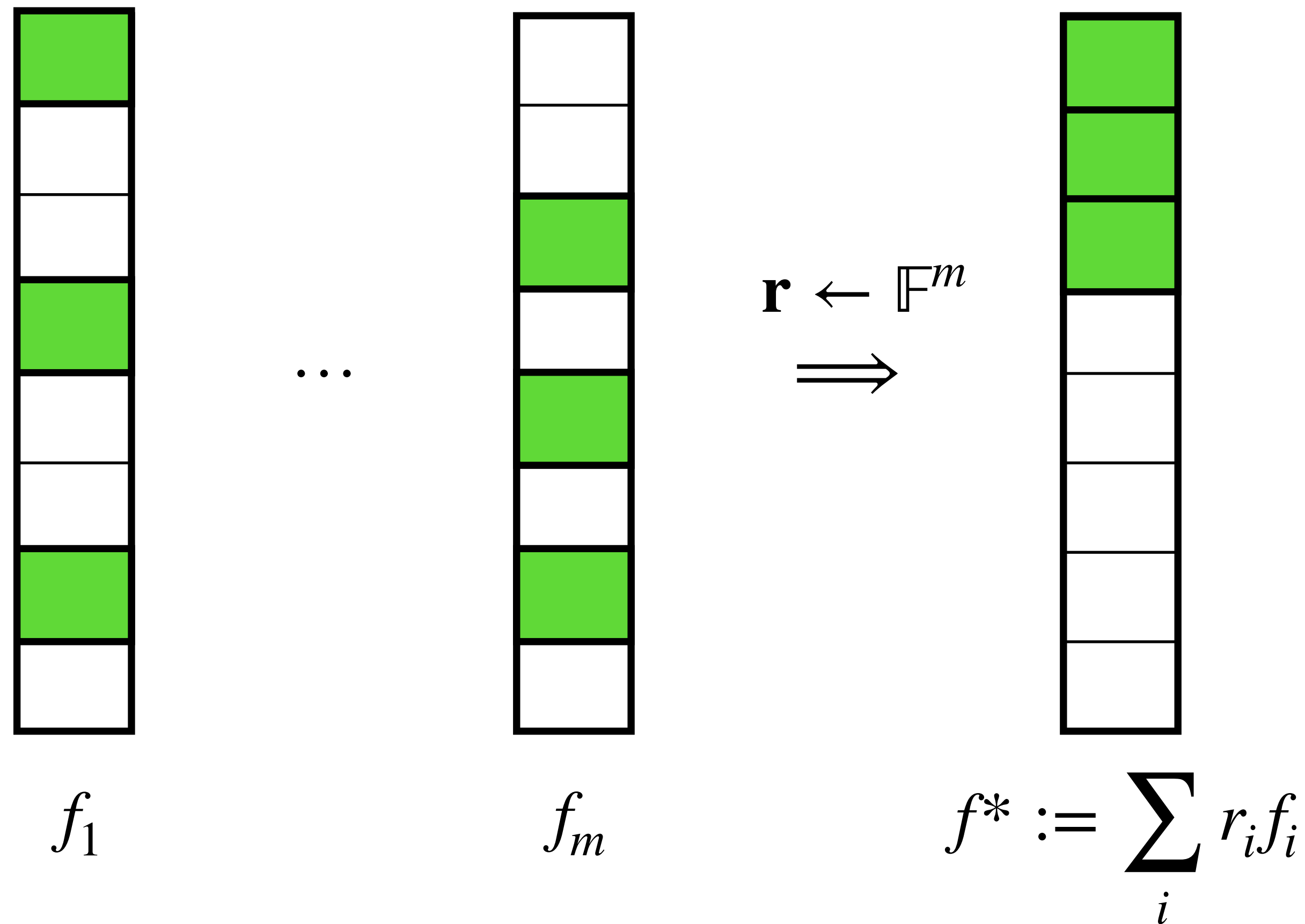


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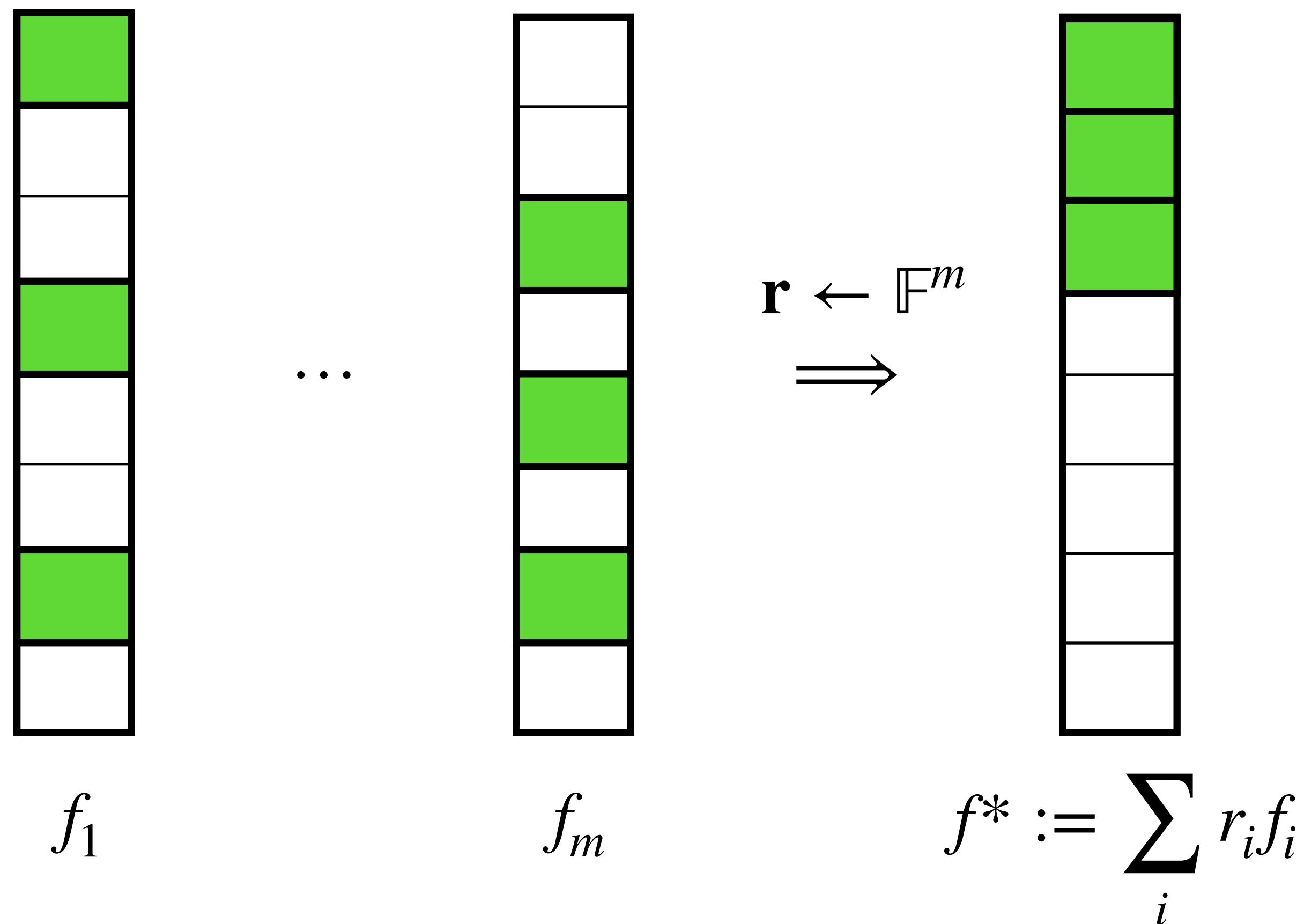


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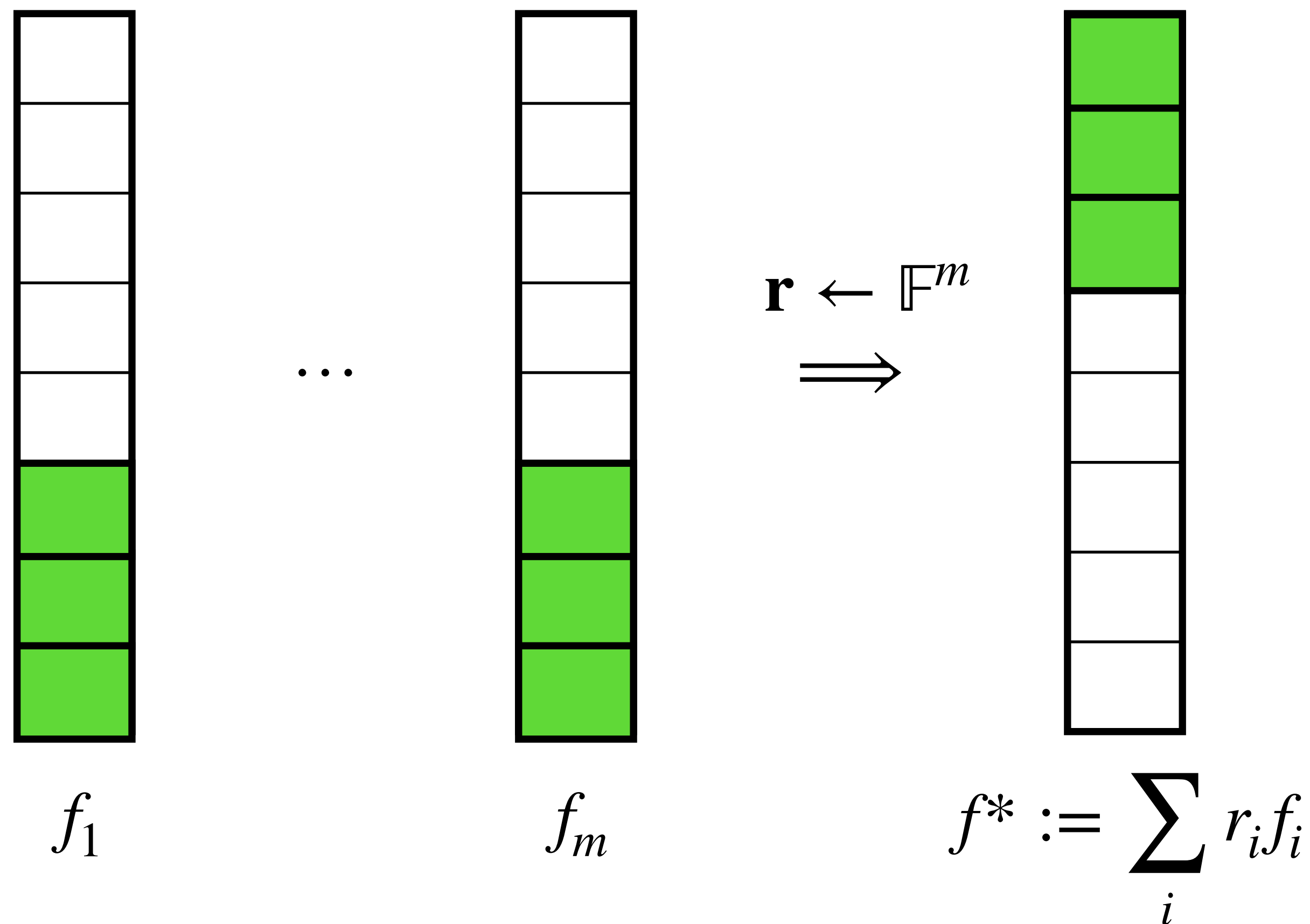
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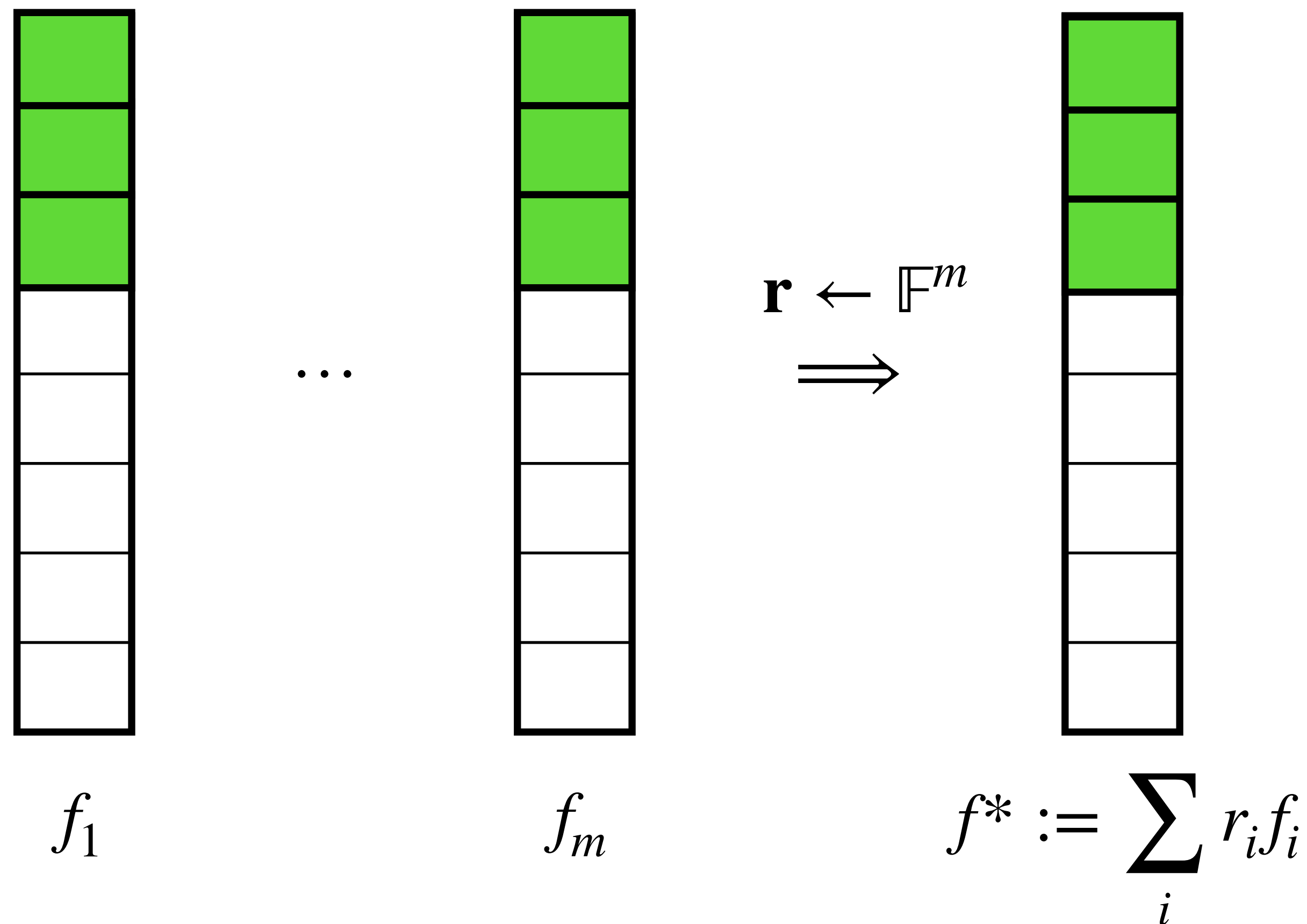
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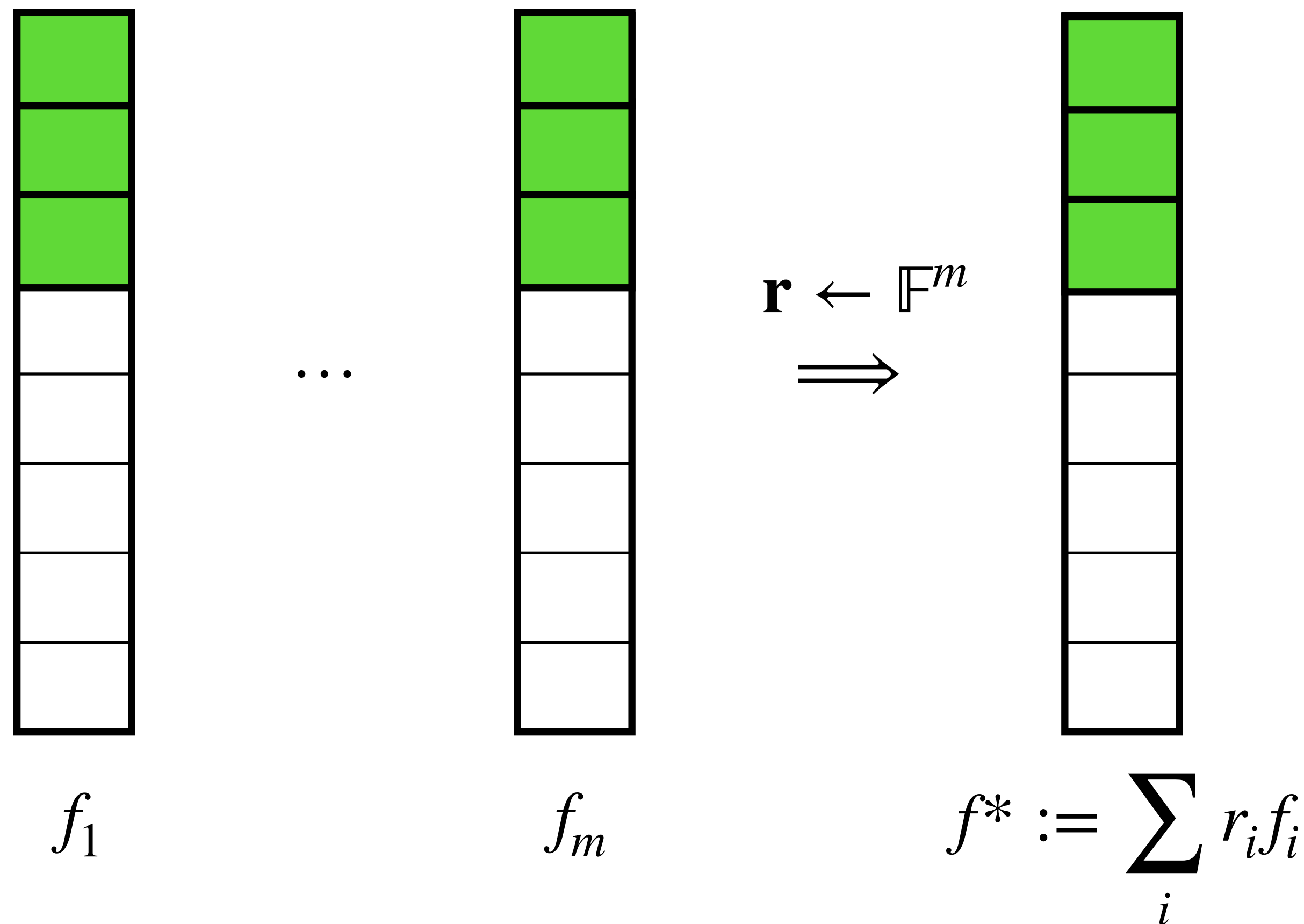
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**Mutual correlated agreement:** the stripe in which  $f_1, \dots, f_m$  agree with  $\mathcal{C}$  is the same on which  $f^*$  does:

*“No new correlated domains appear”*

# List-RLC lemma and List-Fold

Implied by mutual correlated agreement

$\Lambda(\mathcal{C}, f, \delta)$  is the list of  
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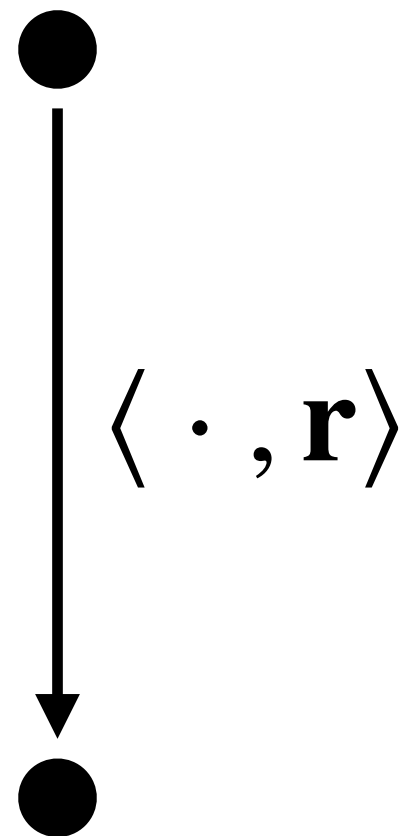
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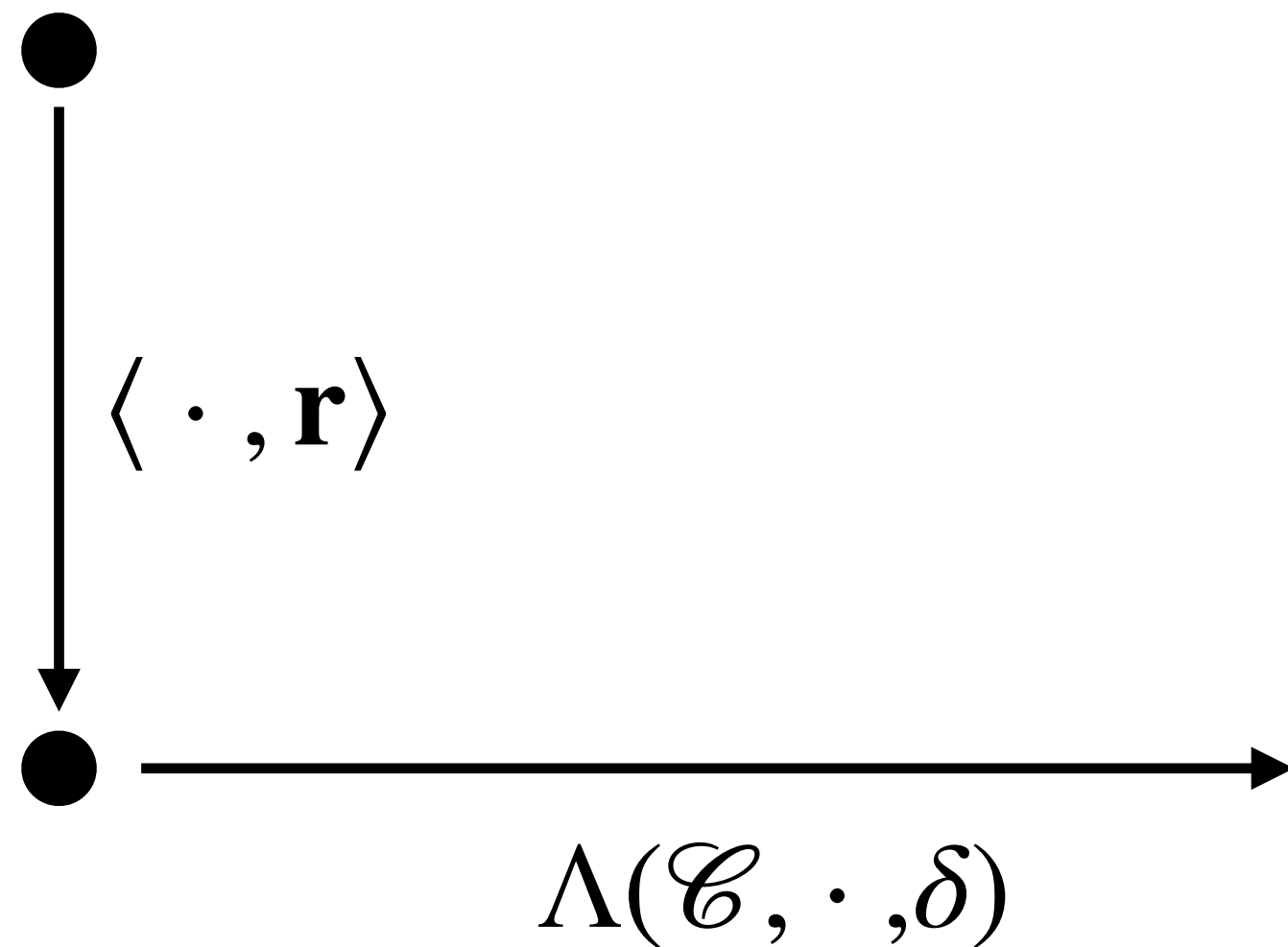
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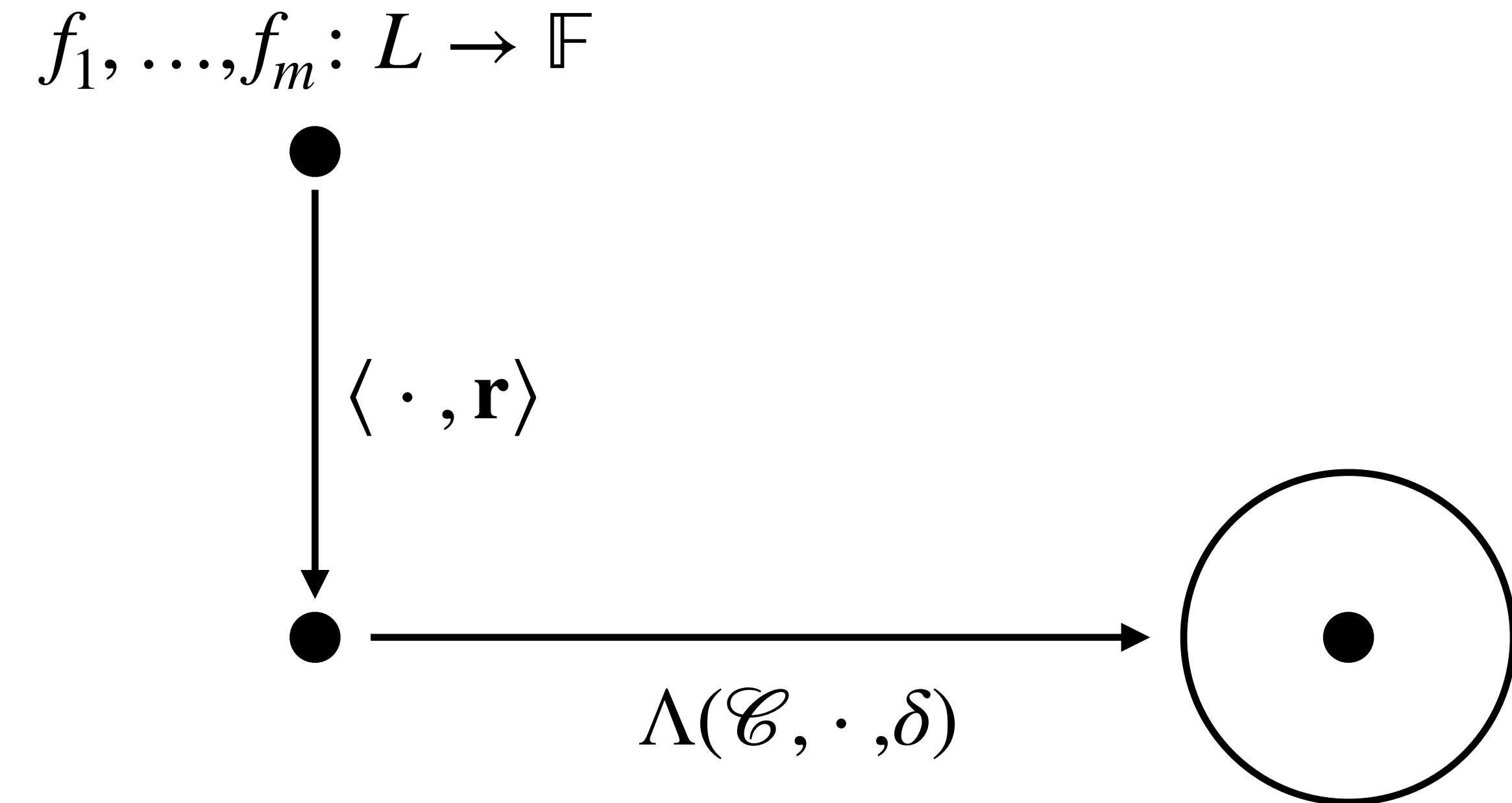


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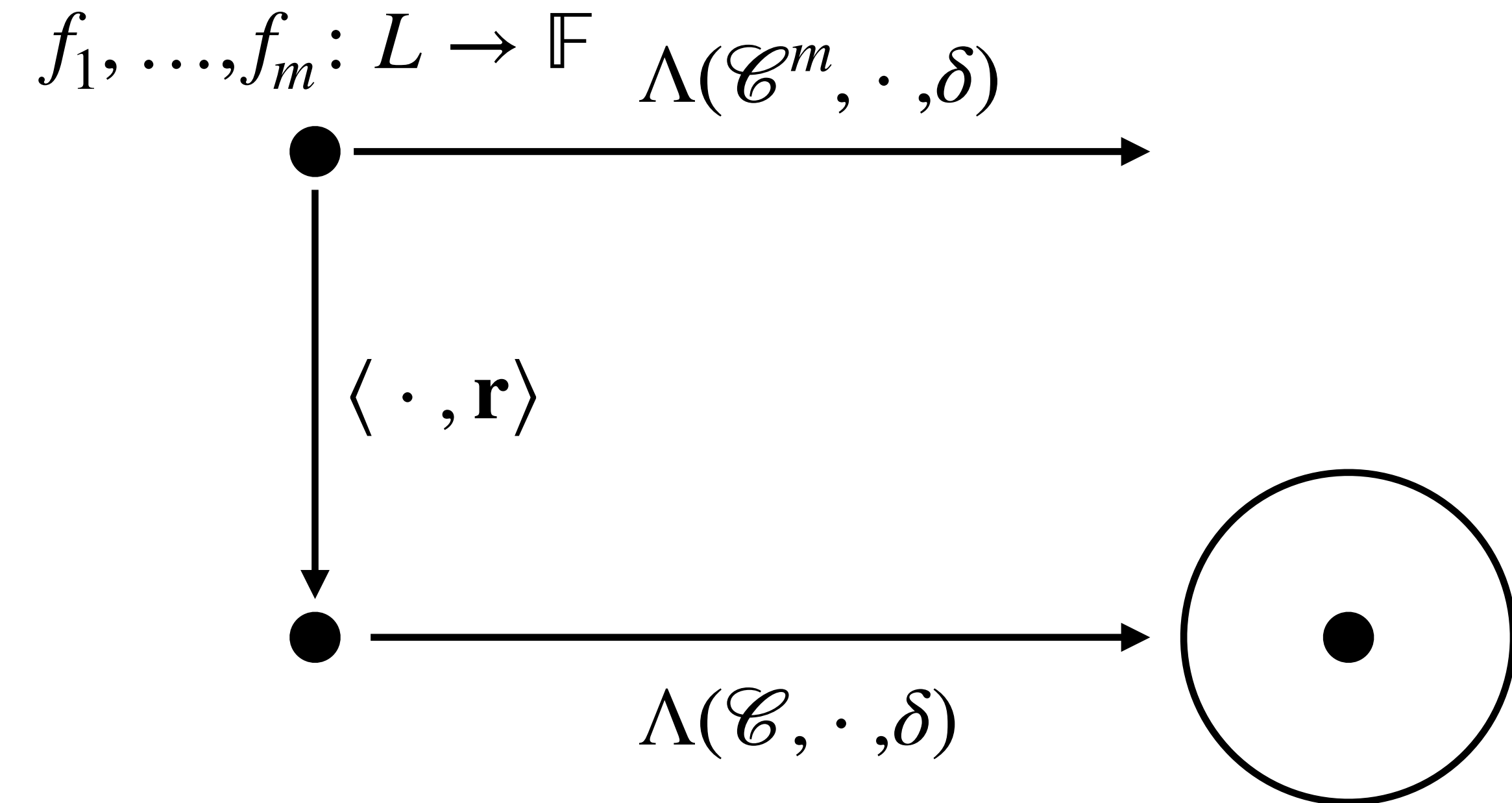


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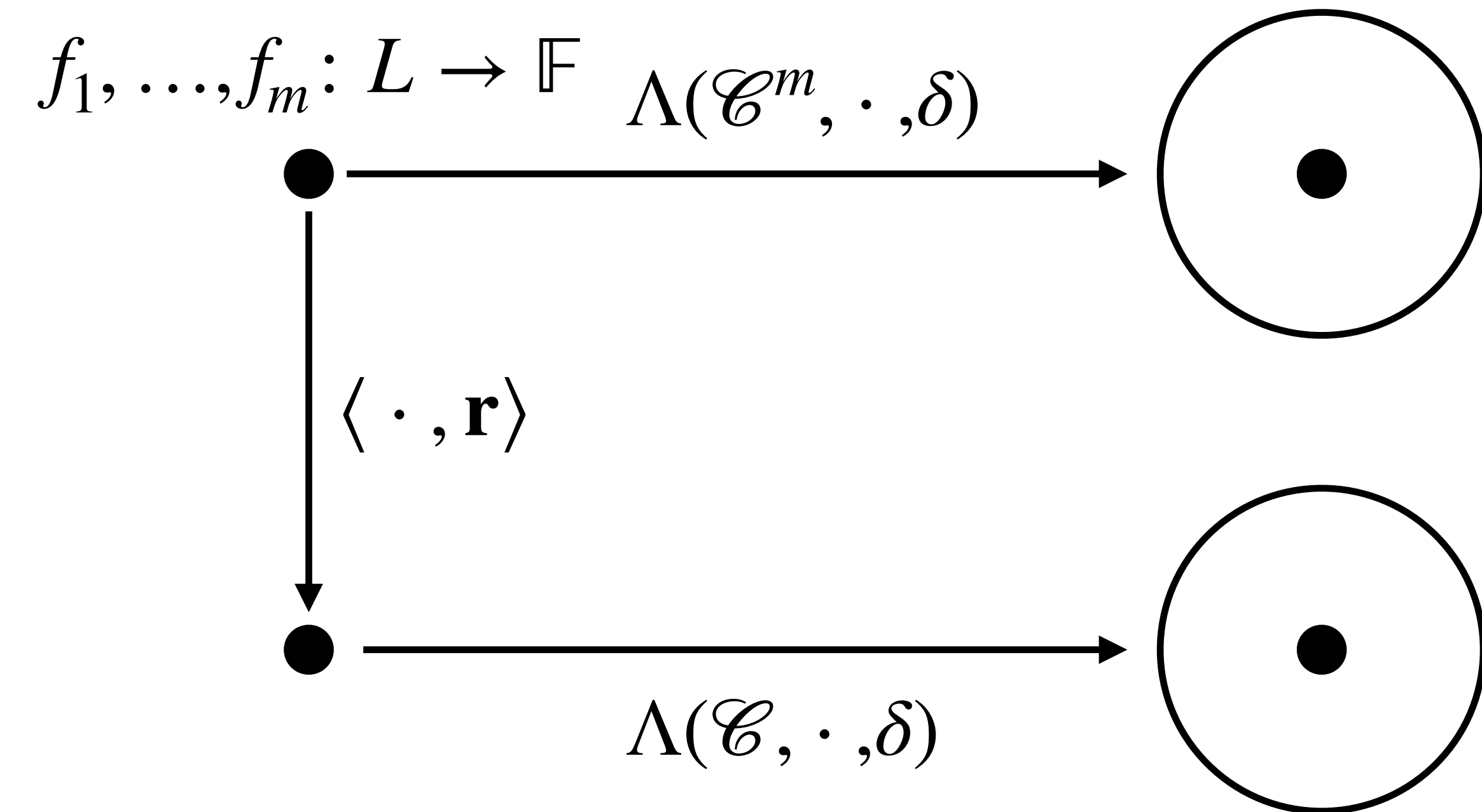


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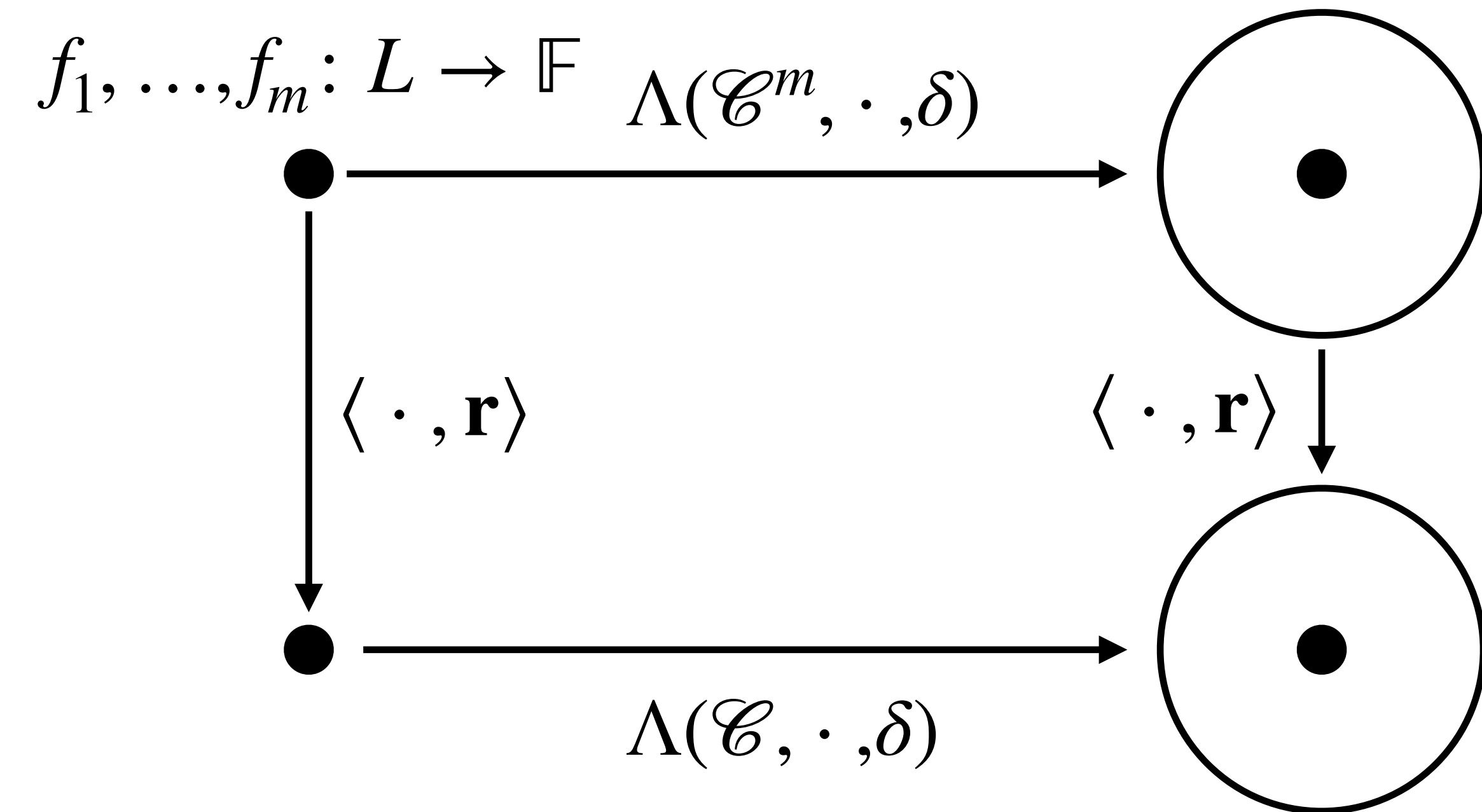


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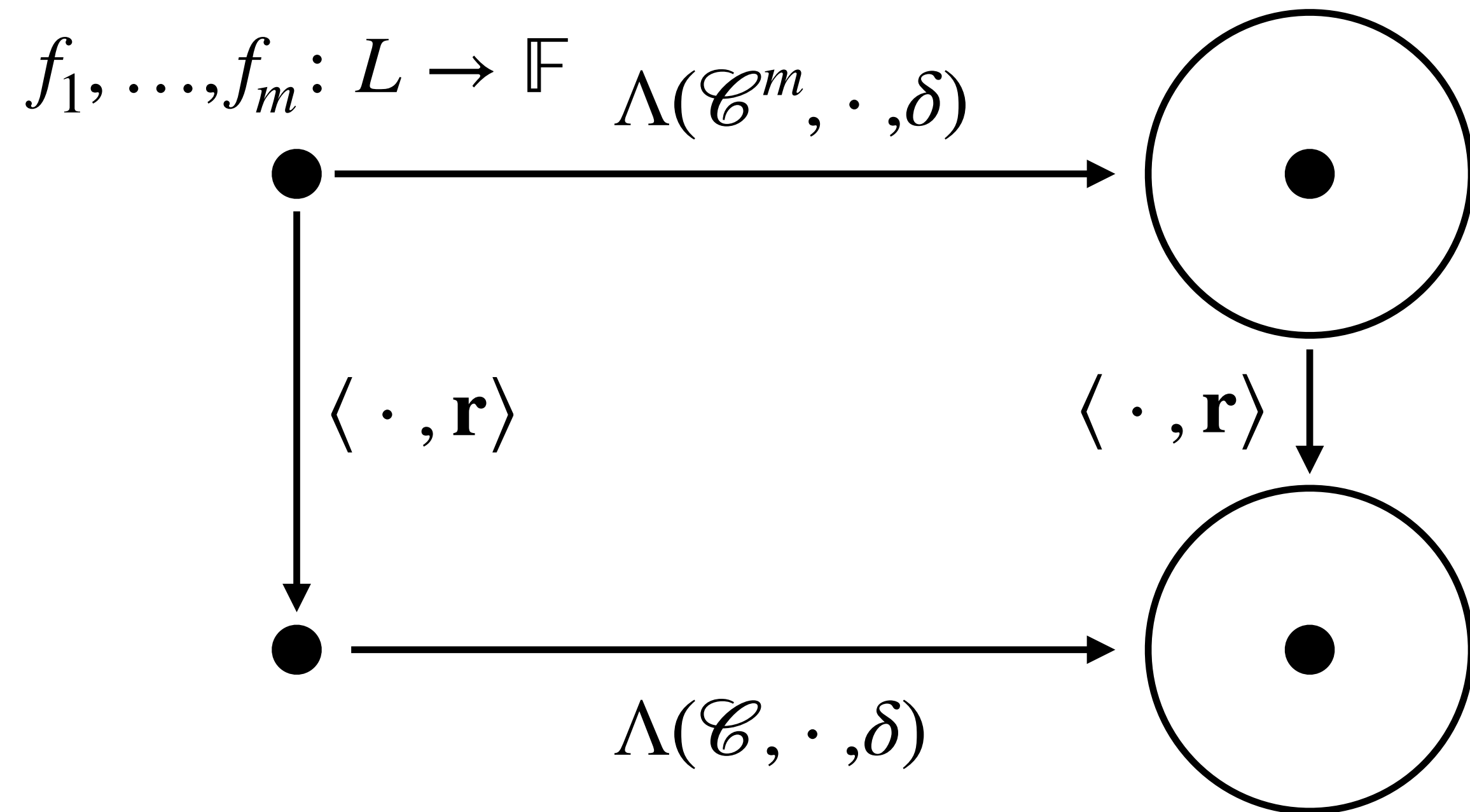




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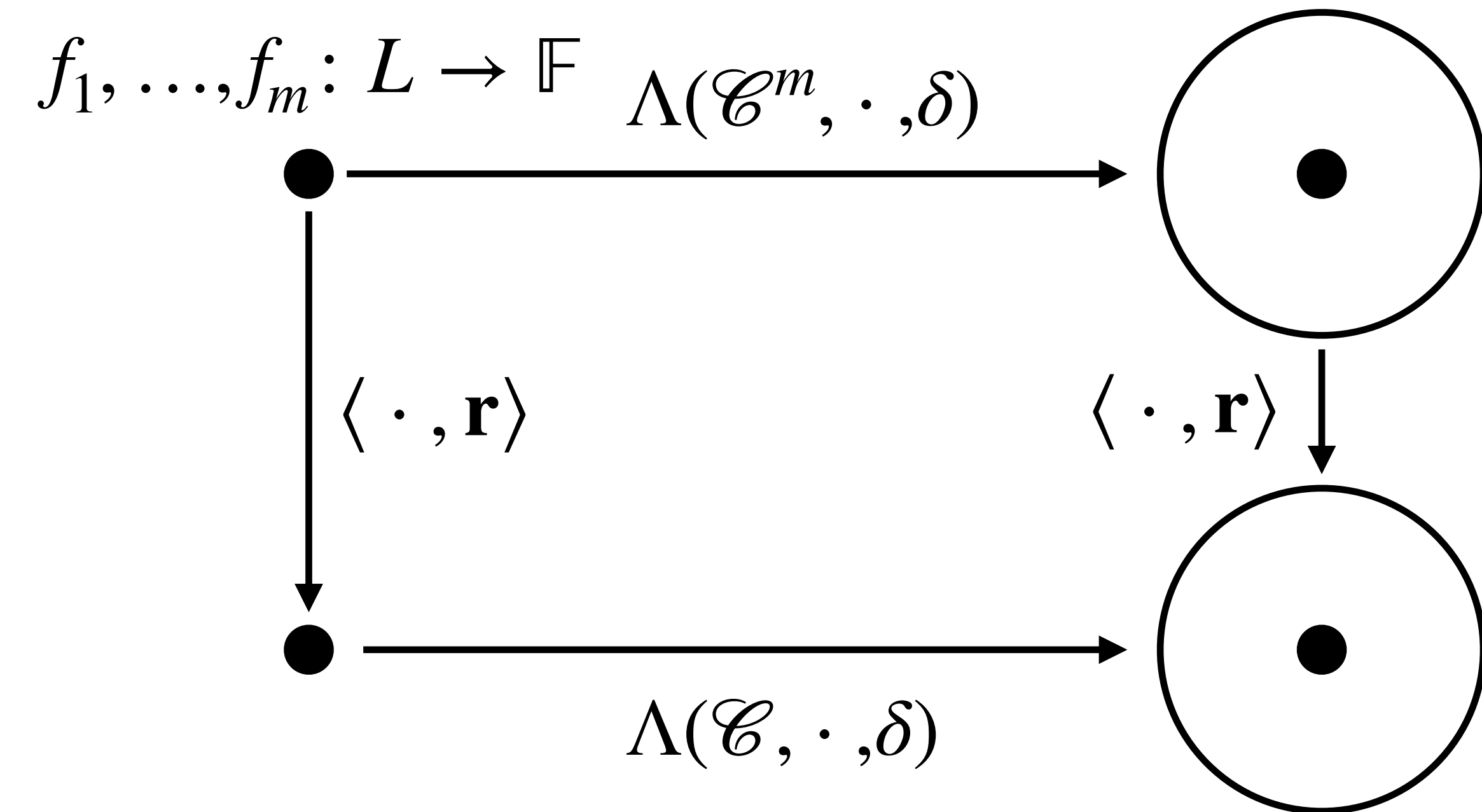


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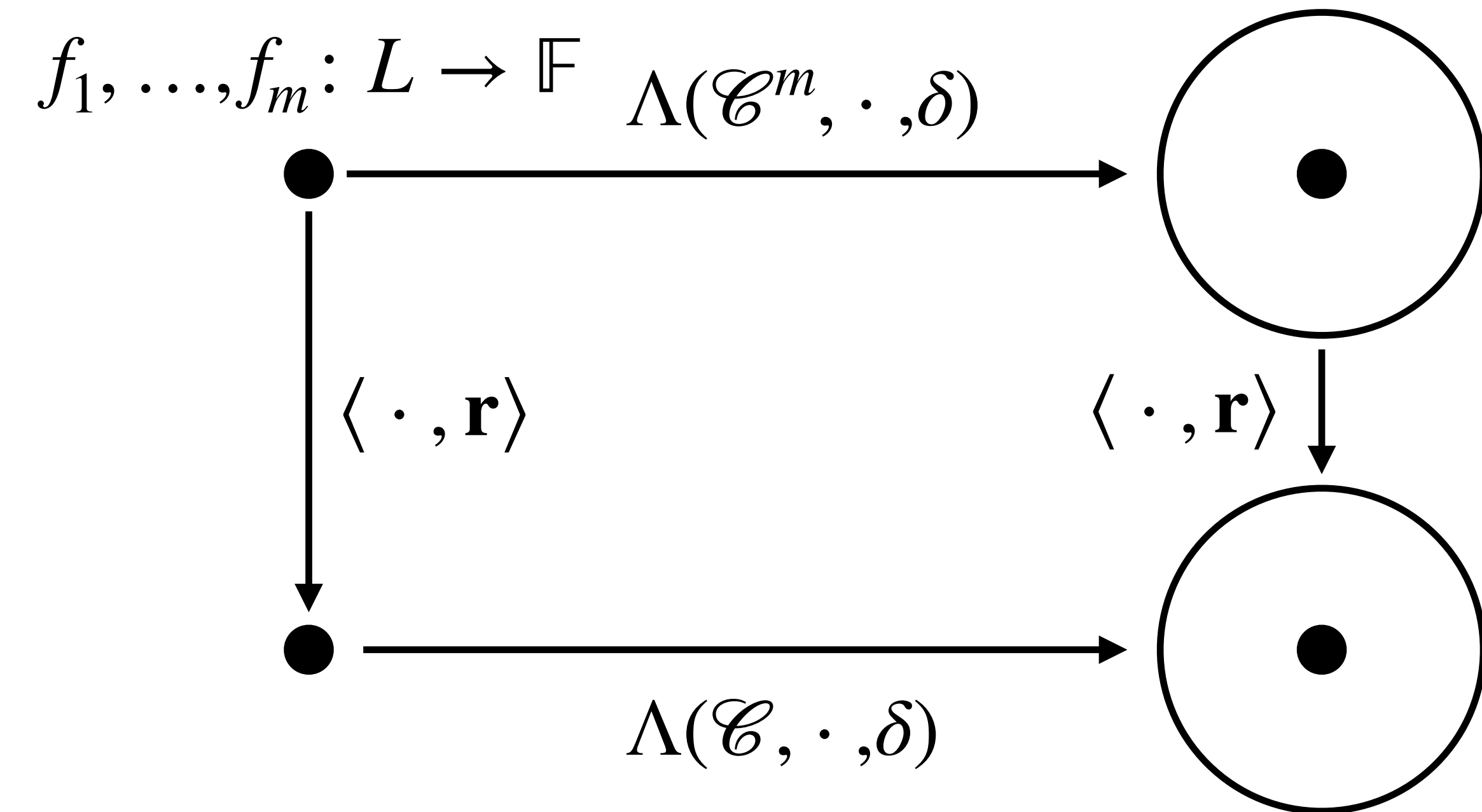


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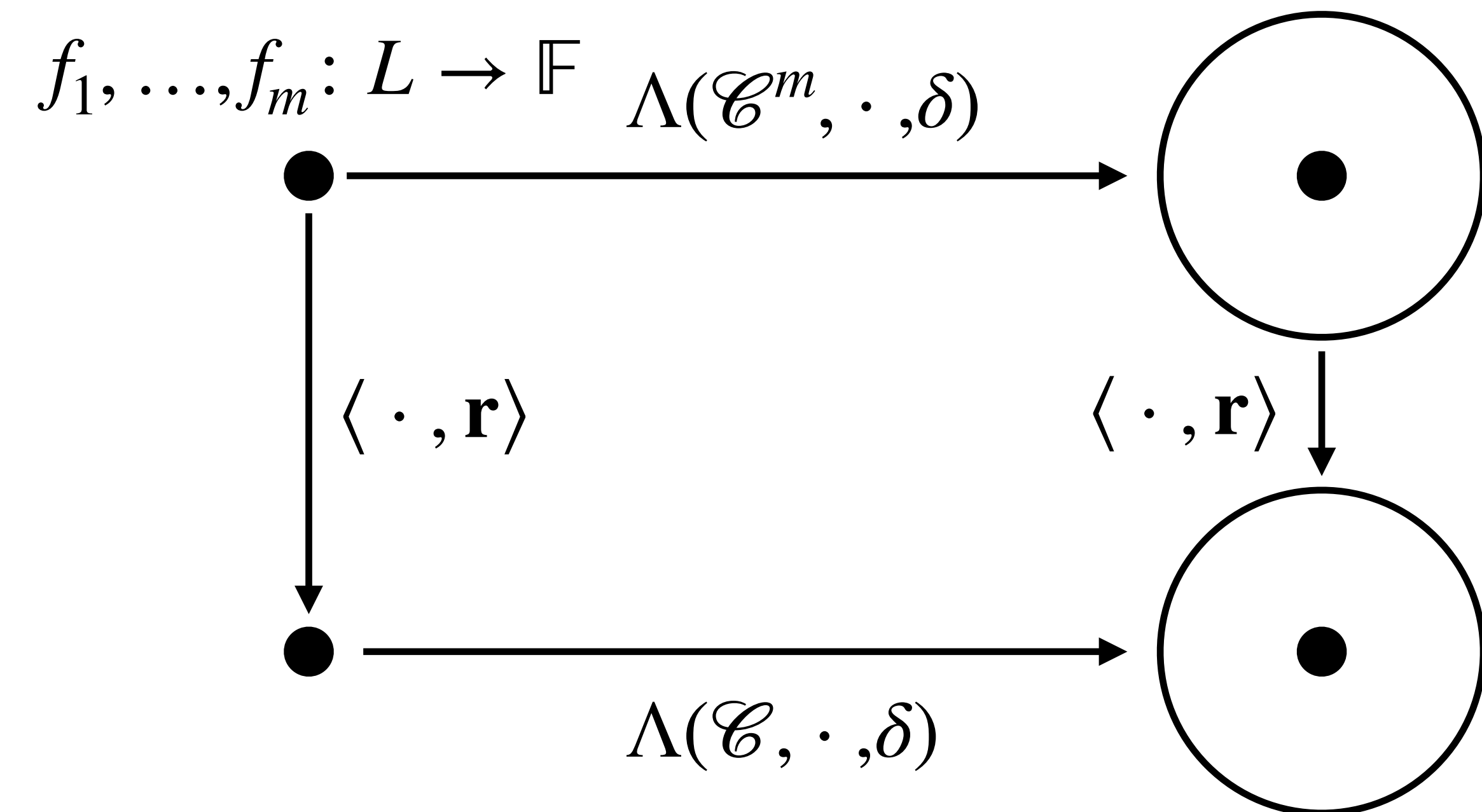


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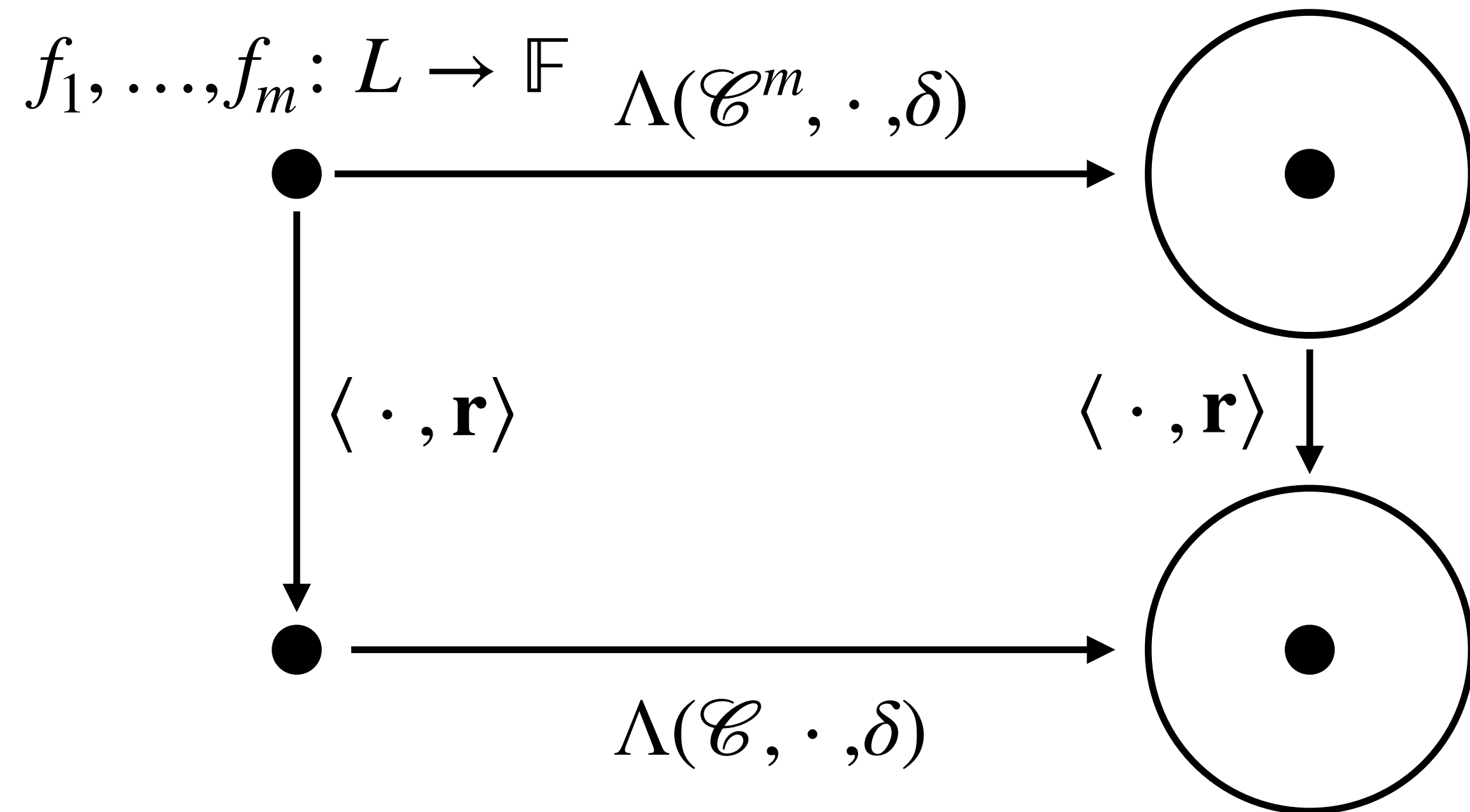
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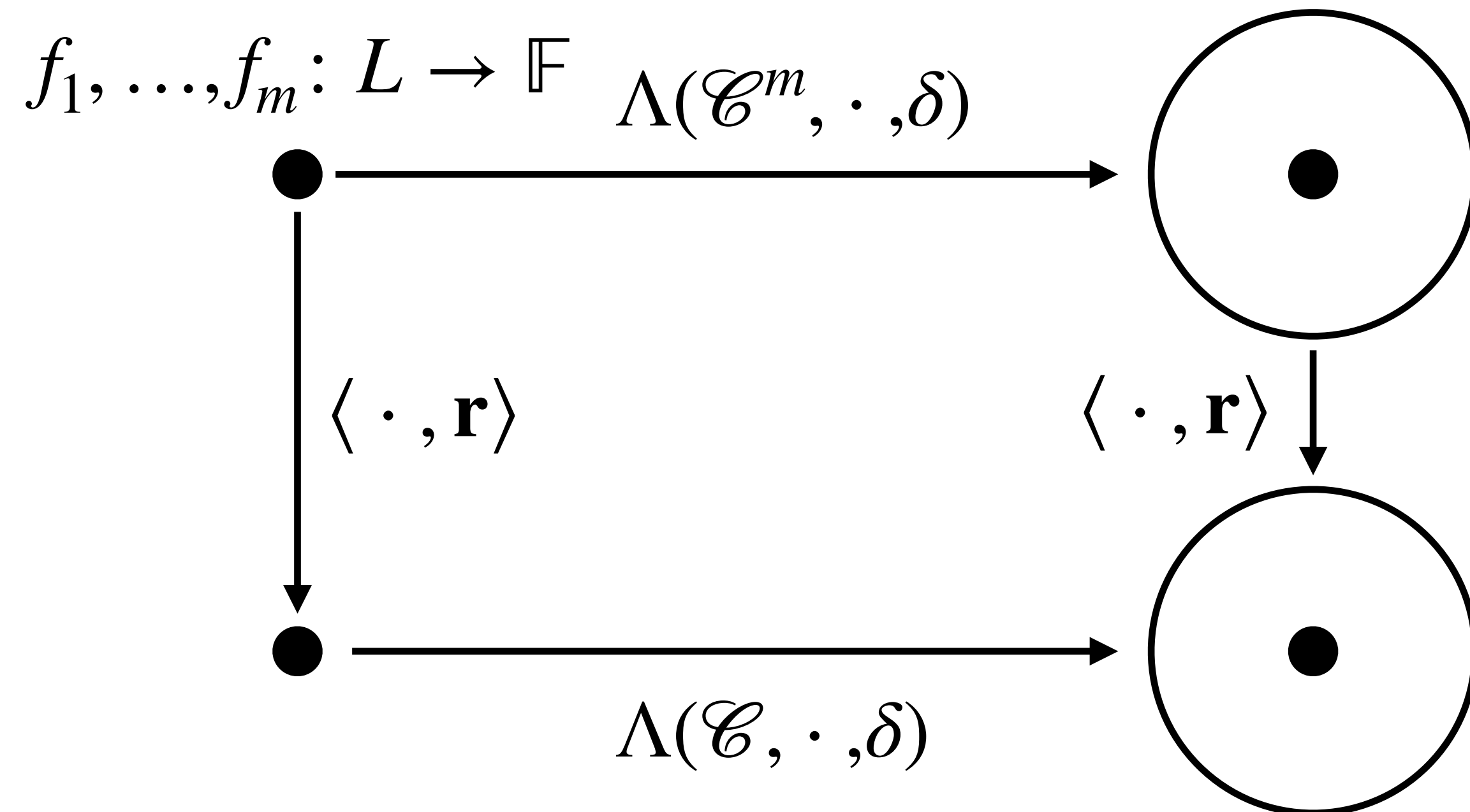
Stronger than what is required for STIR's soundness

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Recent results show it holds up to 1.5 Johnson for general linear codes!

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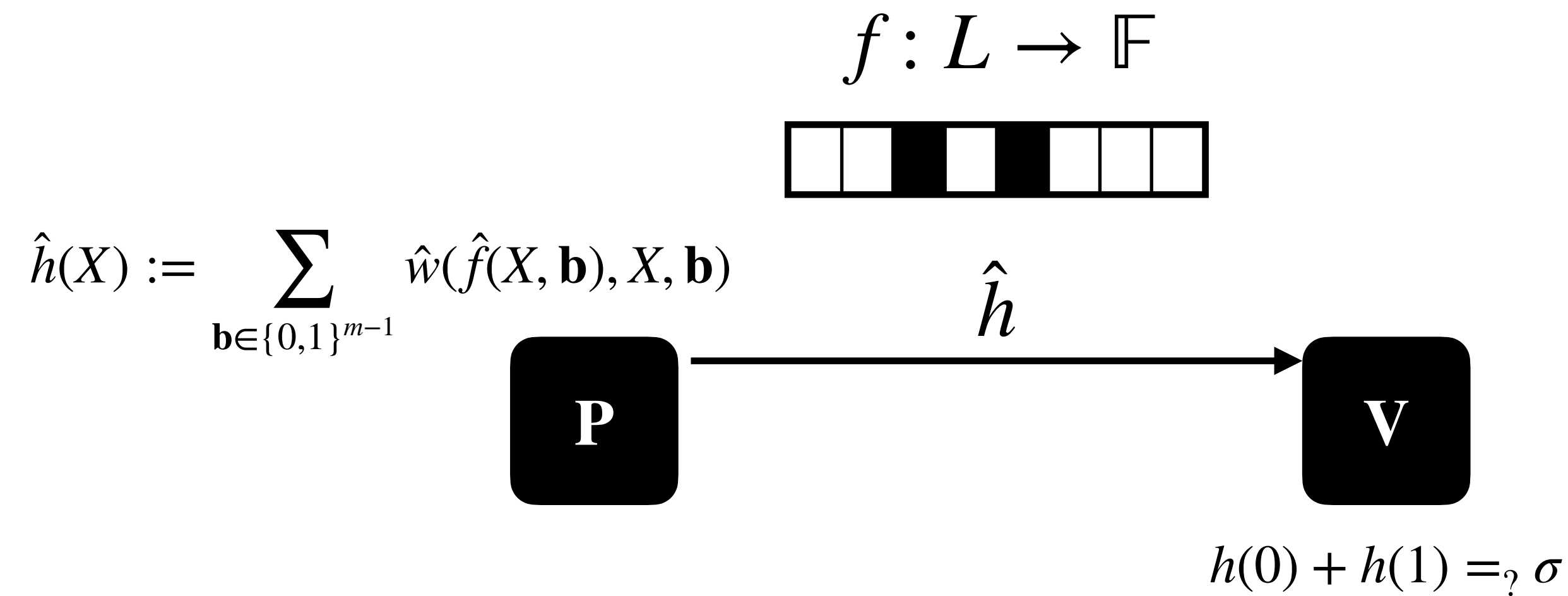
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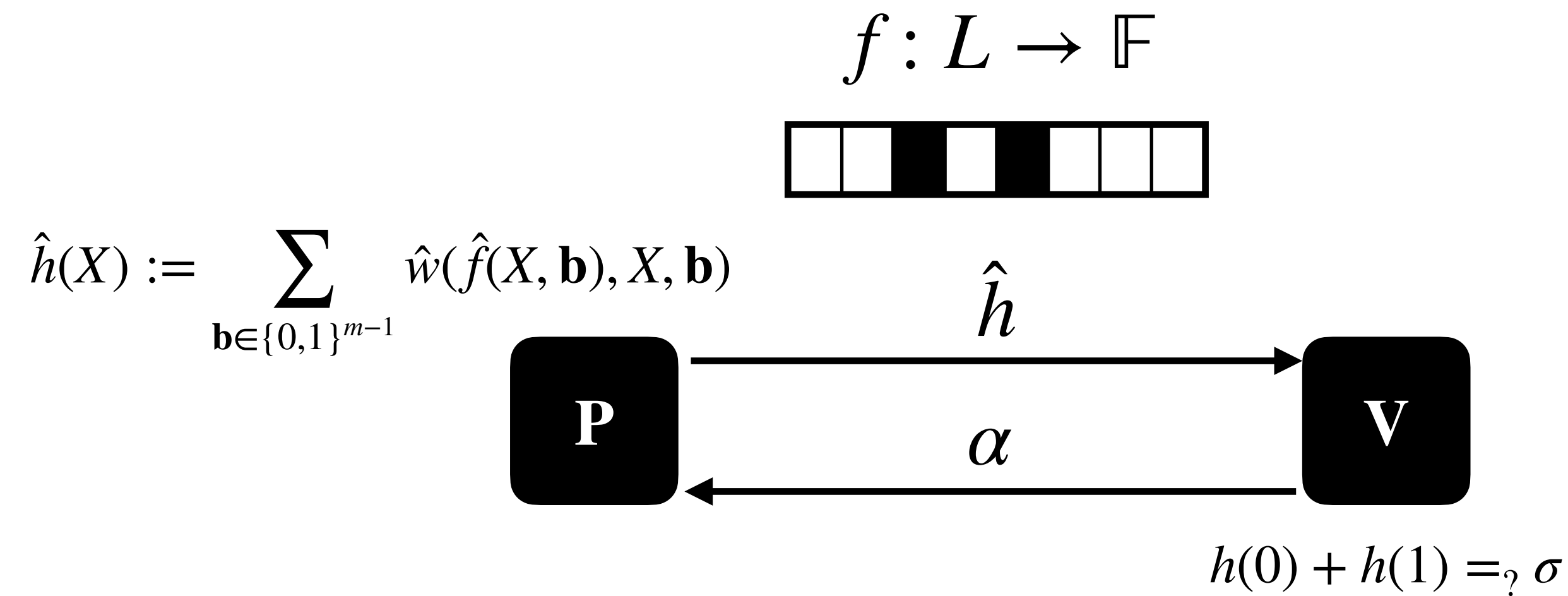
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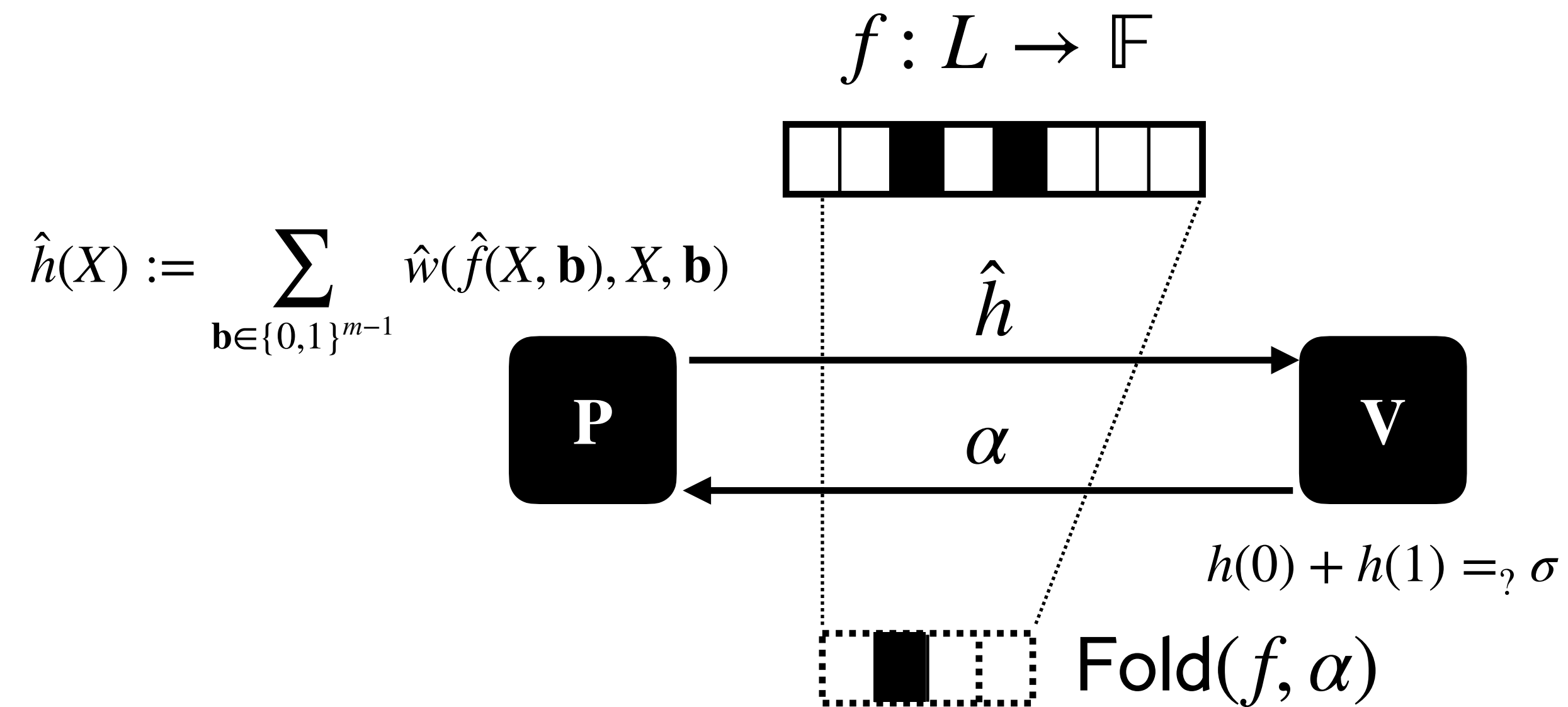
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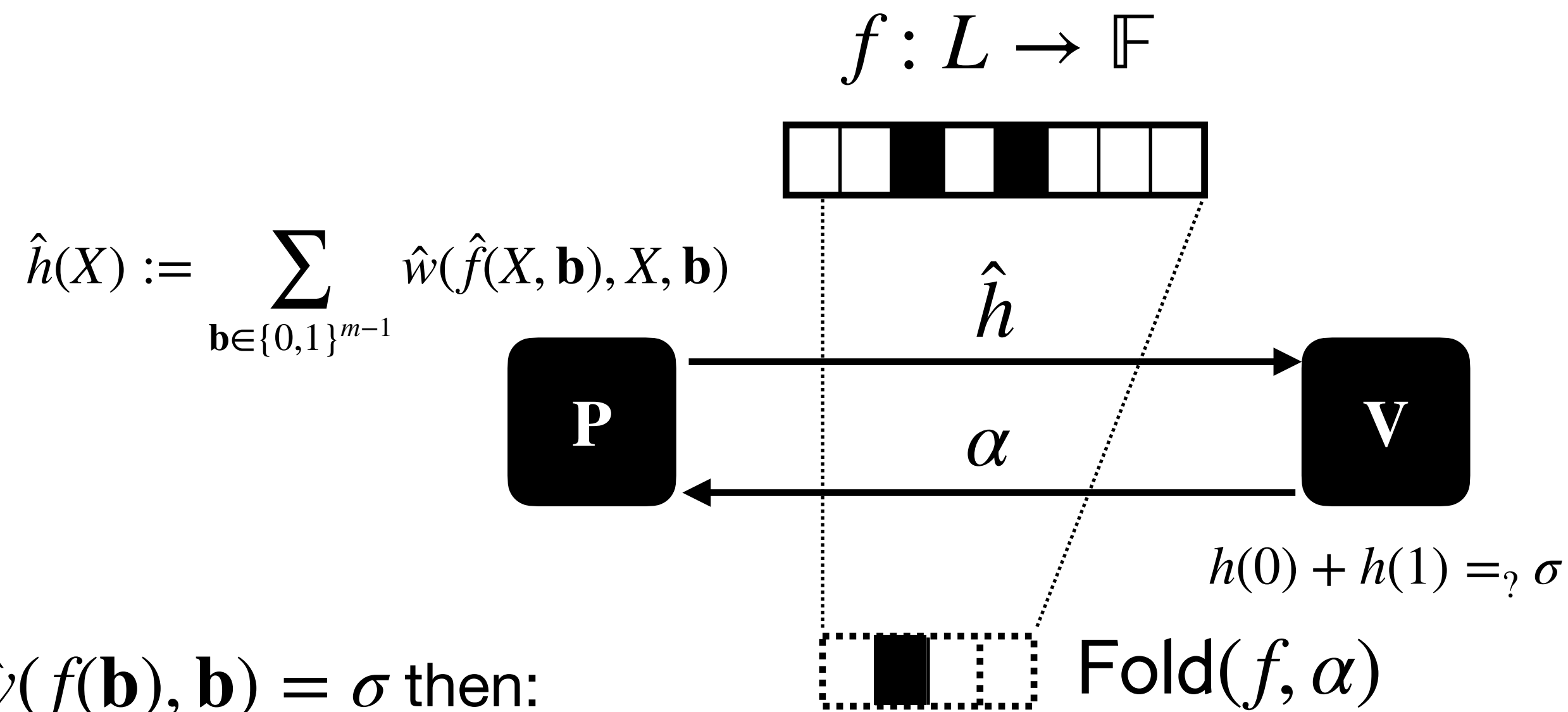
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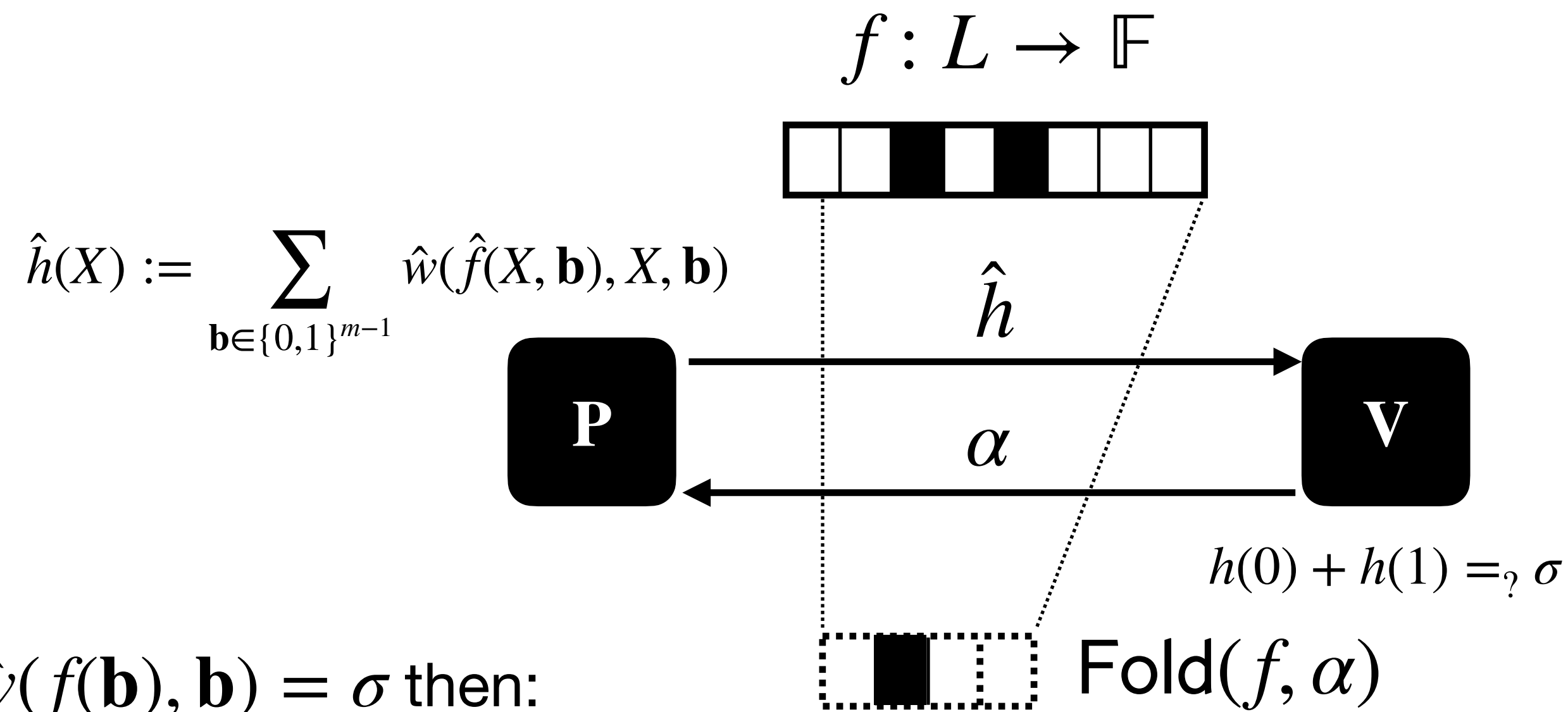


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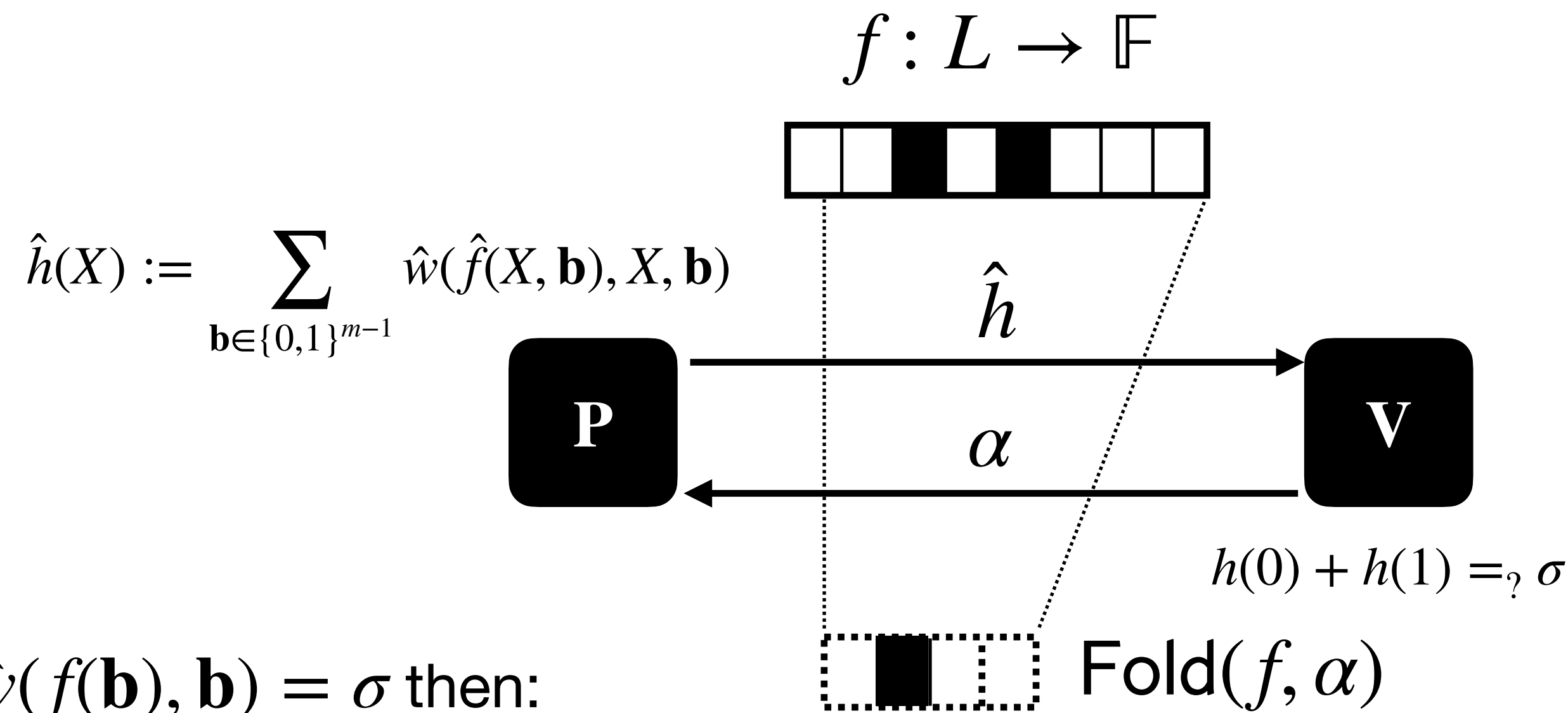
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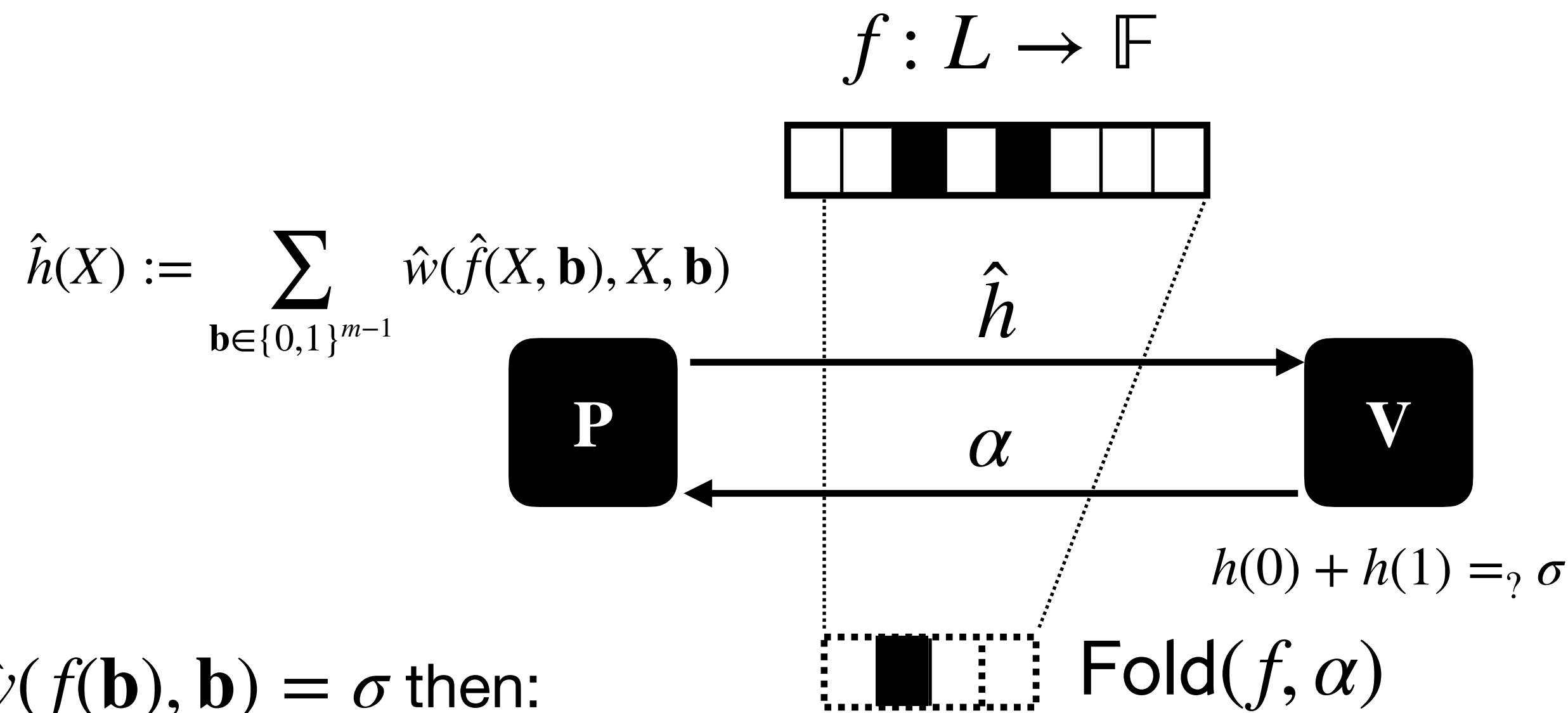
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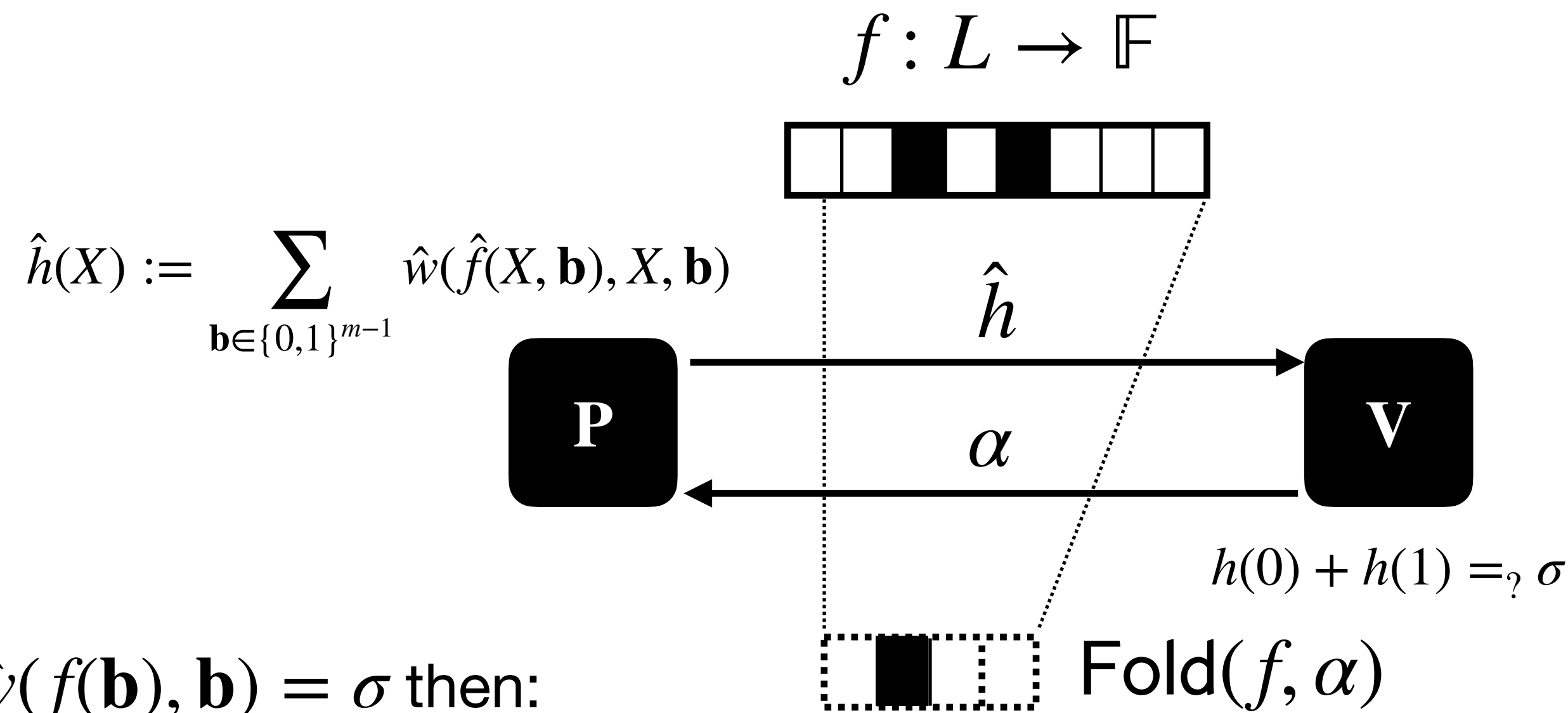
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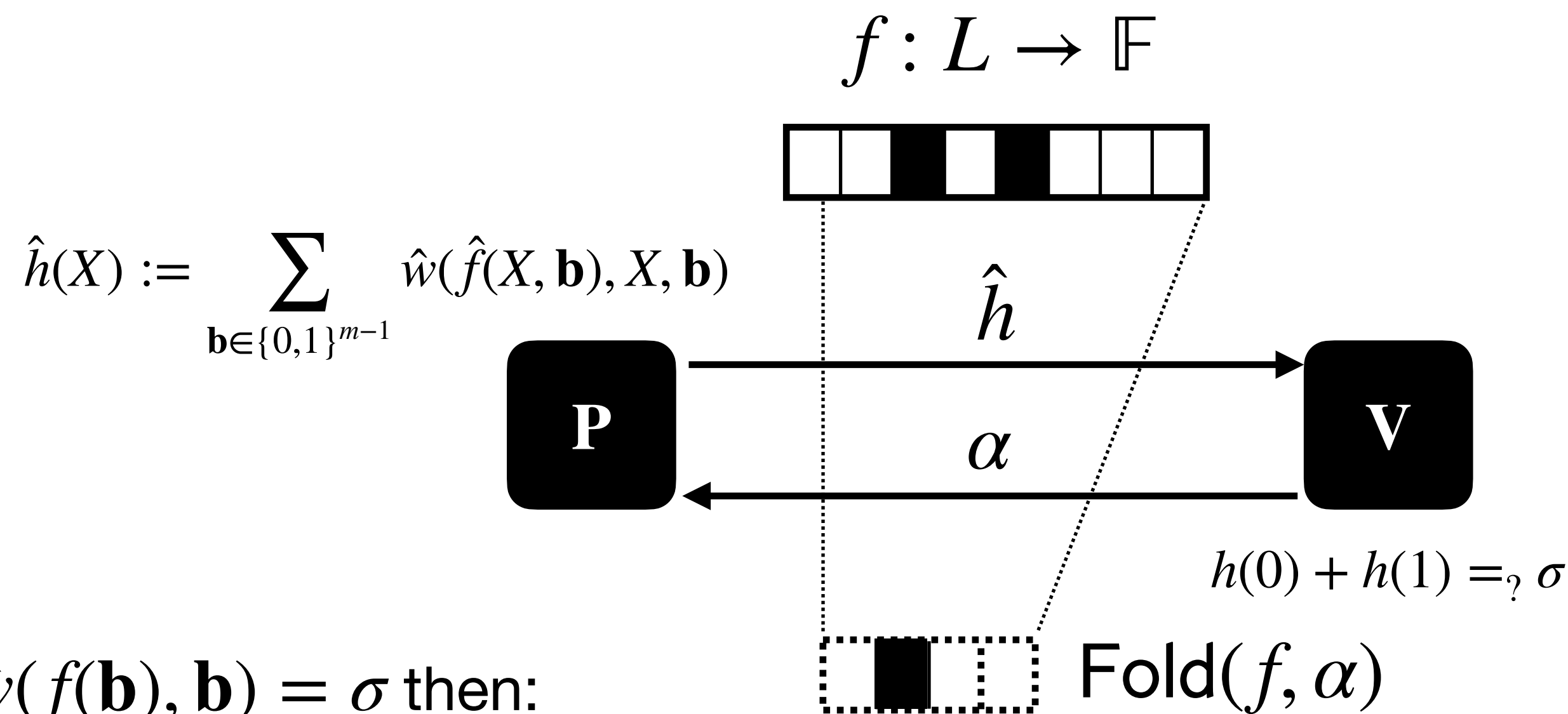
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**Soundness:** by mutual correlated agreement, w.h.p. if  $\Delta(f, \text{CRS}[n, m, \rho, \hat{w}, \sigma]) > \delta$  then  $\Delta(\text{Fold}(f, \alpha), \text{CRS}[n/2, m - 1, \rho, \hat{w}_\alpha, \hat{h}(\alpha)]) > \delta$

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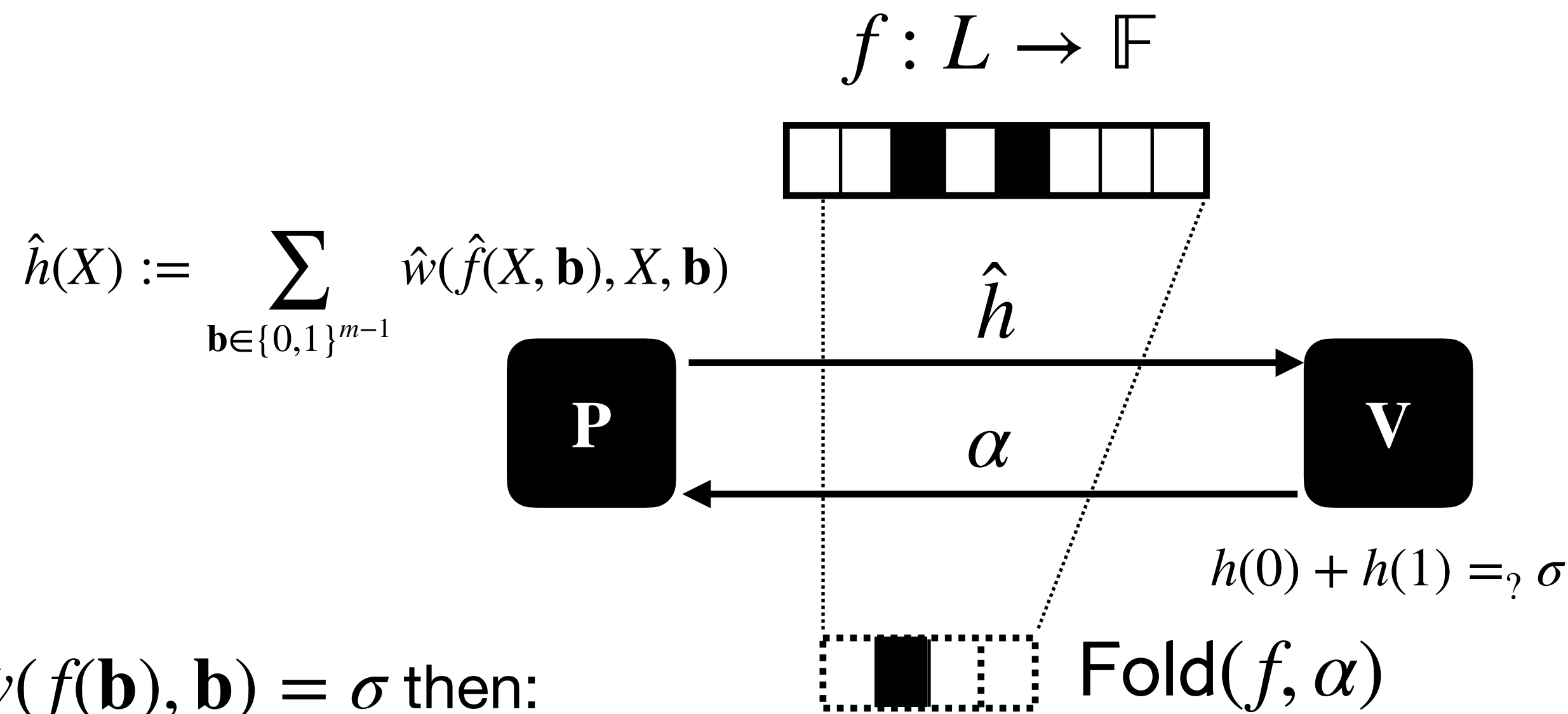
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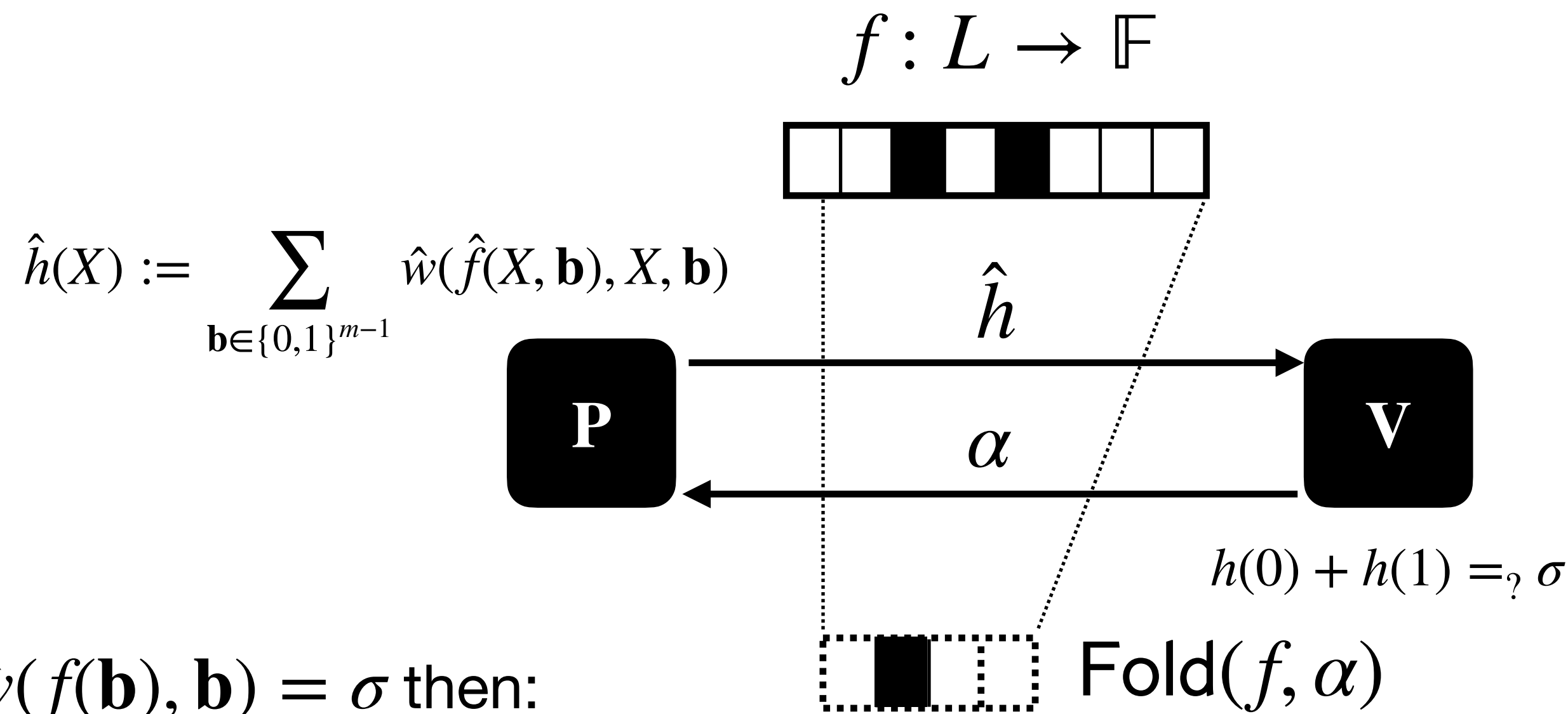
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Unchanged!

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In the full protocol, we fold by 2-by-2  $k$  times. Can also fold by  $2^k$  at a time (nice for first round!)

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- $h(0) + h(1) = \sigma,$
- $\sum_{\mathbf{b}} \hat{w}(f(\alpha, \mathbf{b}), \alpha, \mathbf{b}) = \hat{h}(\alpha)$
- $\widehat{\text{Fold}(f, \alpha)} = \hat{f}(\alpha, \cdot)$

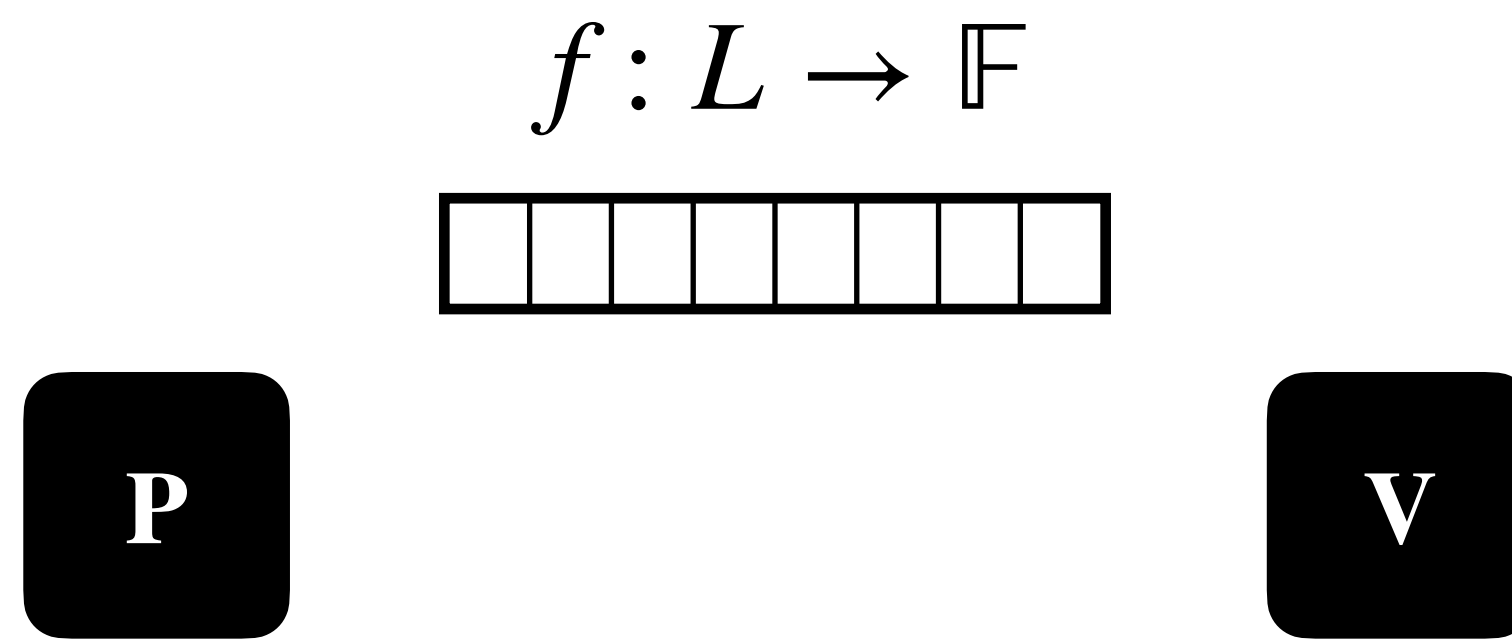
**Soundness:** by mutual correlated agreement, w.h.p. if  $\Delta(f, \text{CRS}[n, m, \rho, \hat{w}, \sigma]) > \delta$  then  $\Delta(\text{Fold}(f, \alpha), \text{CRS}[n/2, m - 1, \rho, \hat{w}_\alpha, \hat{h}(\alpha)]) > \delta$

$\hat{w}_\alpha(Z, \mathbf{X}) = \hat{w}(Z, \alpha, \mathbf{X})$

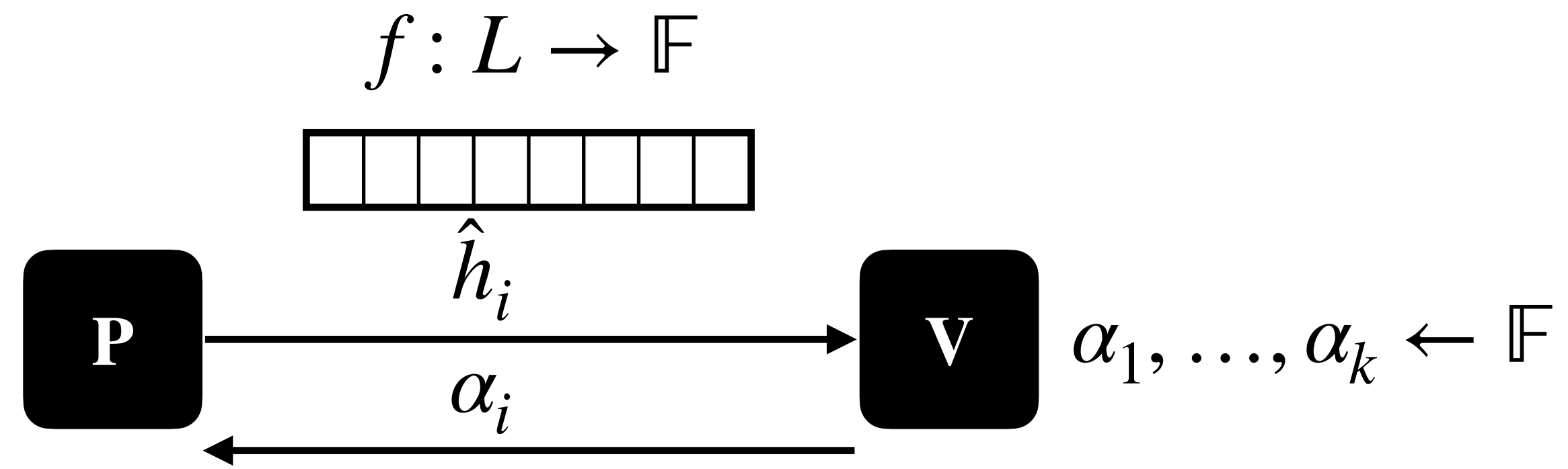
Unchanged!

# WHIR iteration

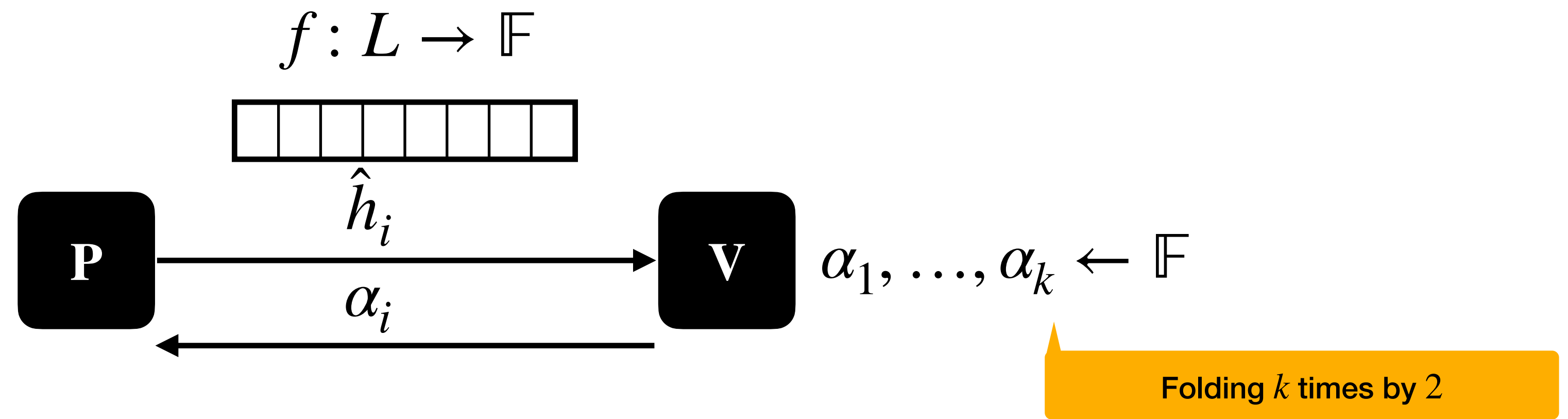
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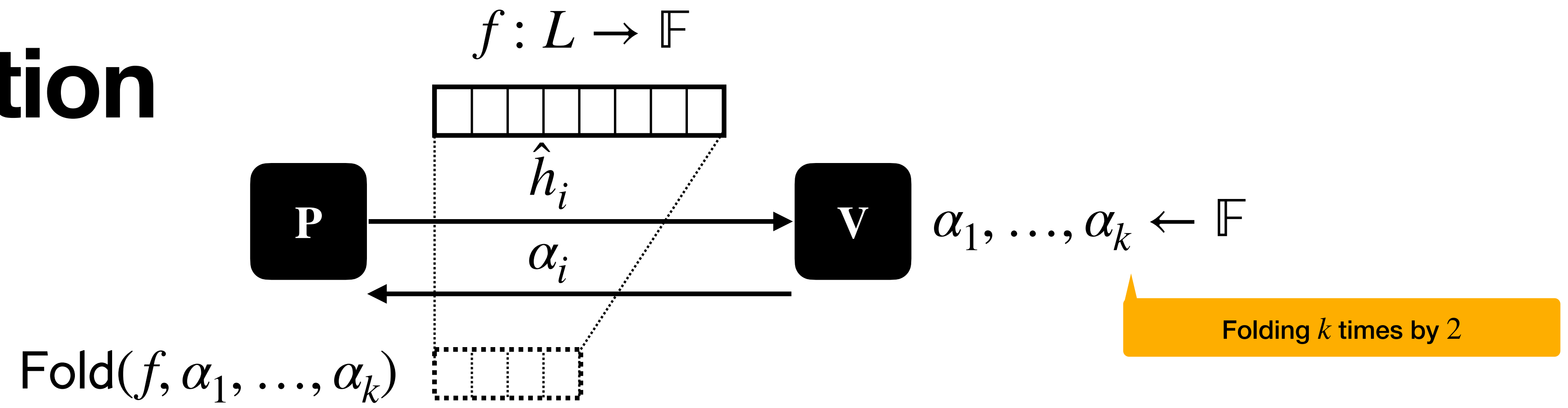


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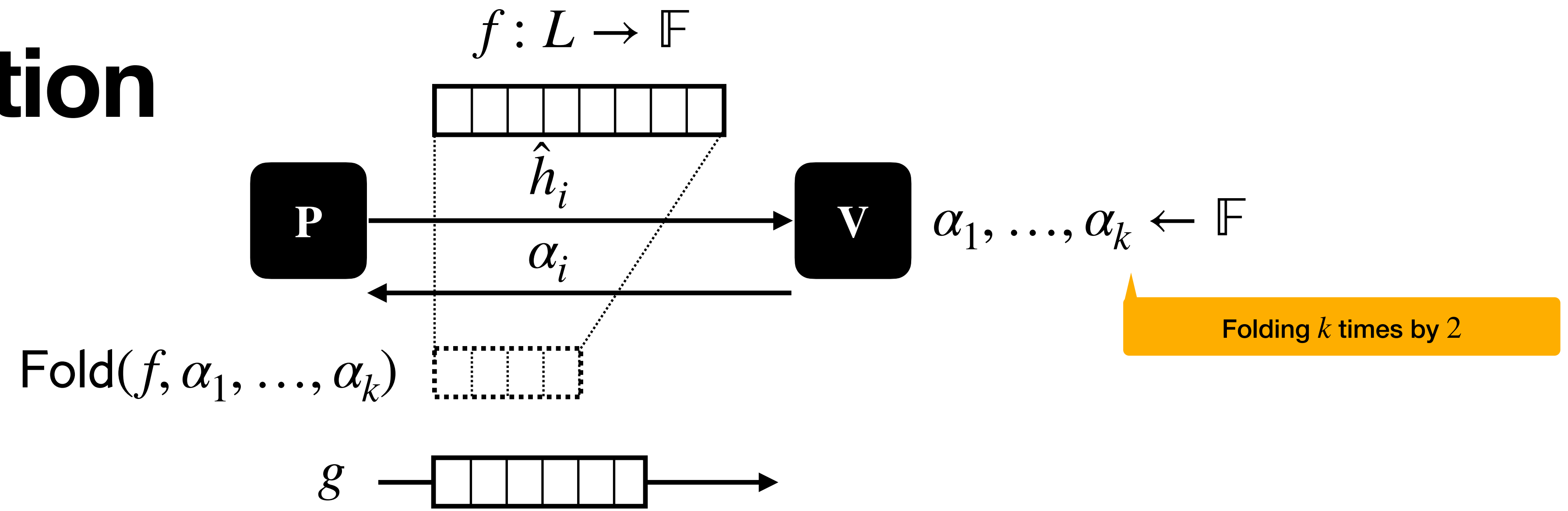




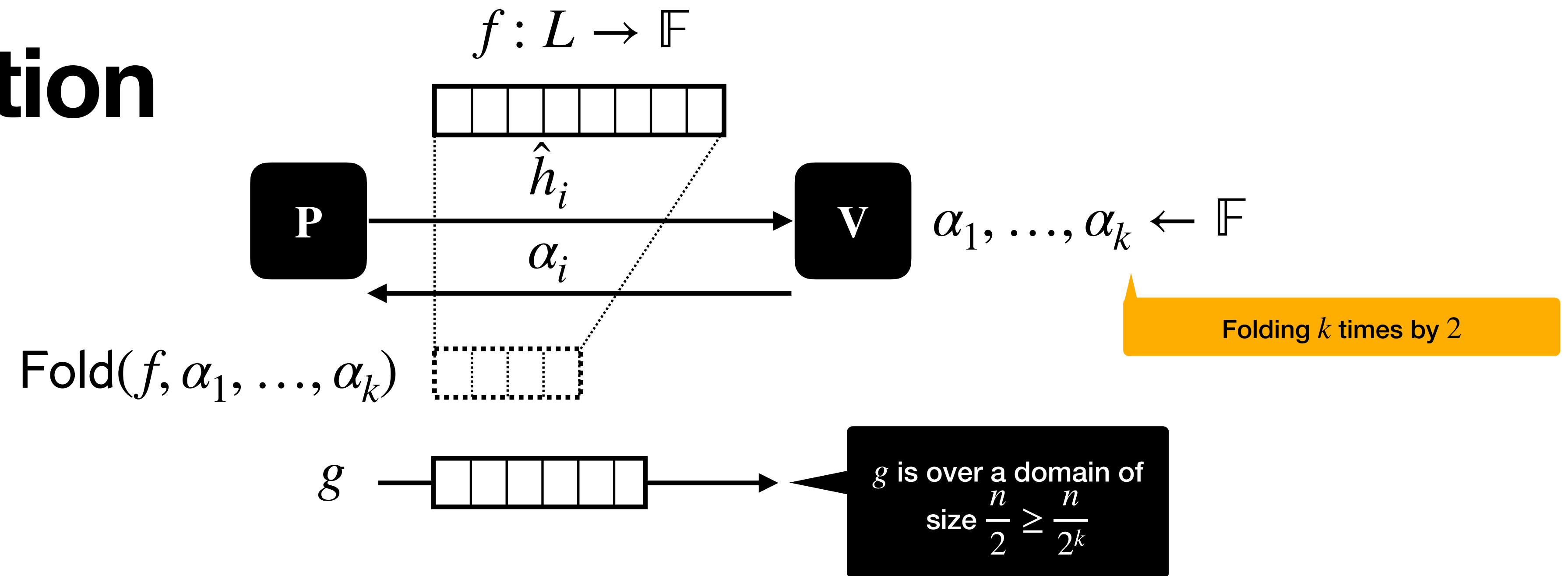
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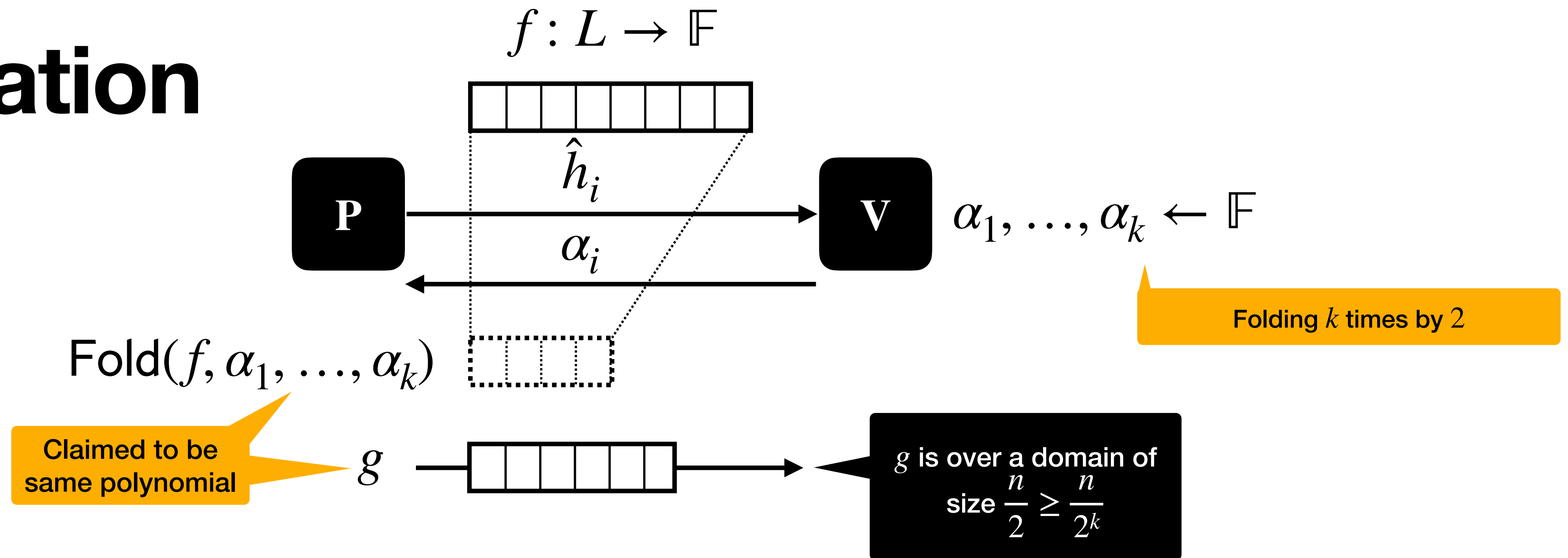
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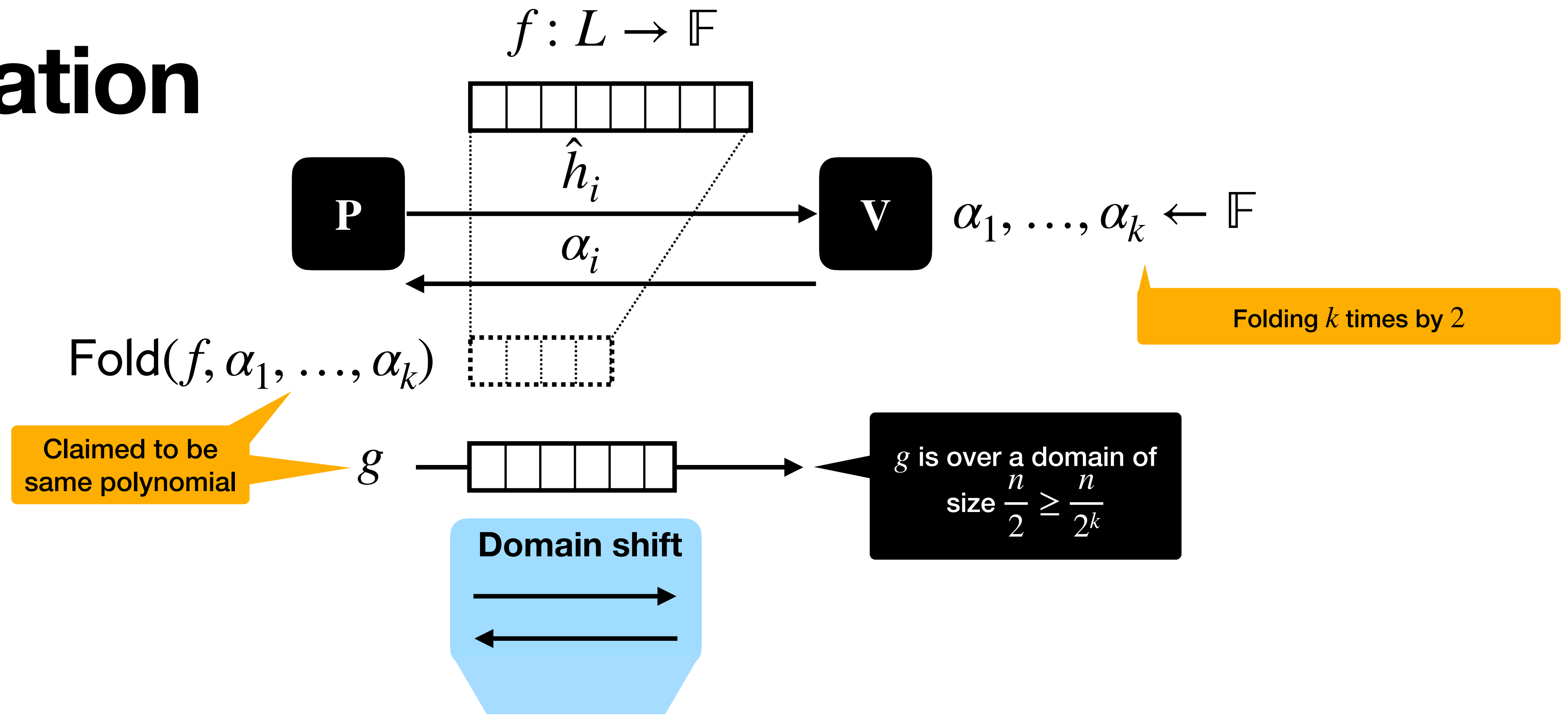
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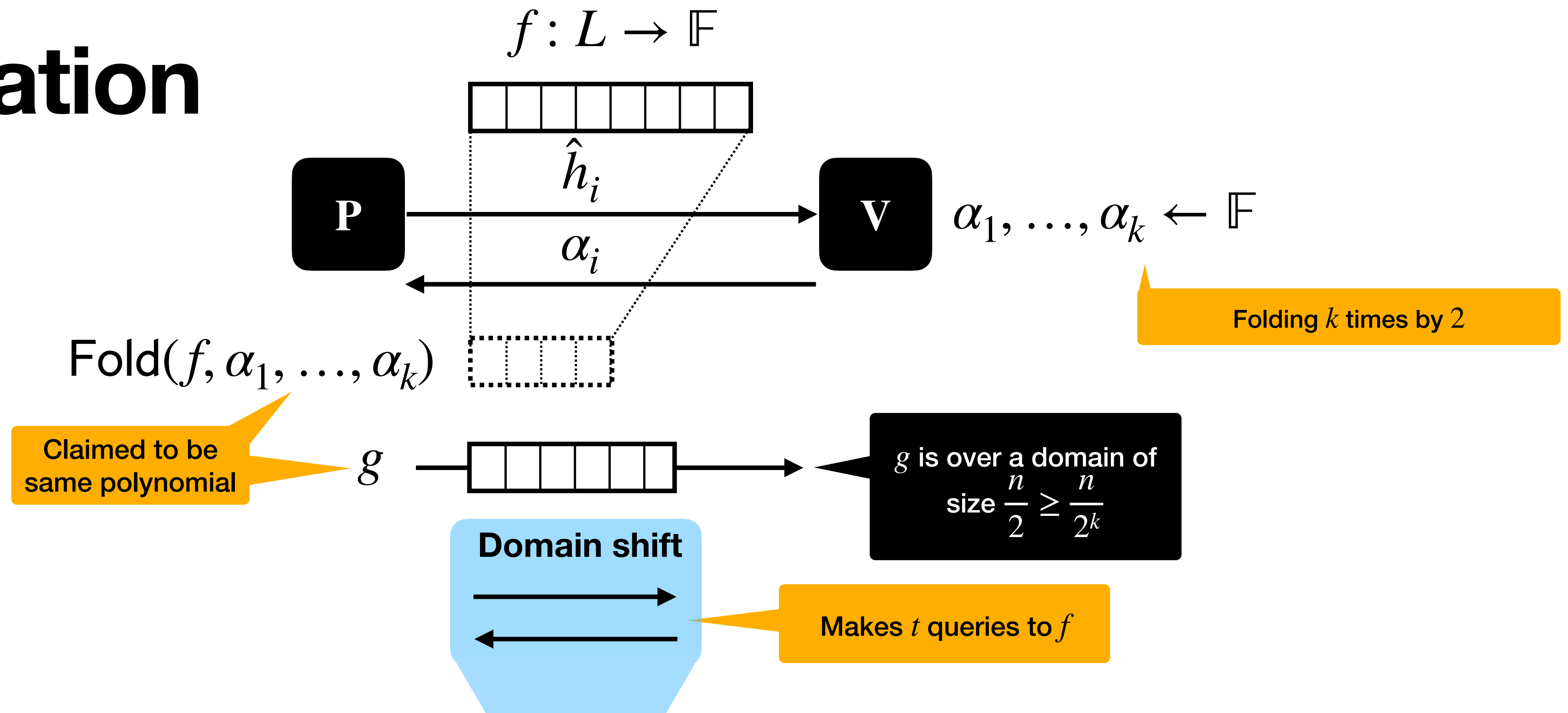
# WHIR iteration



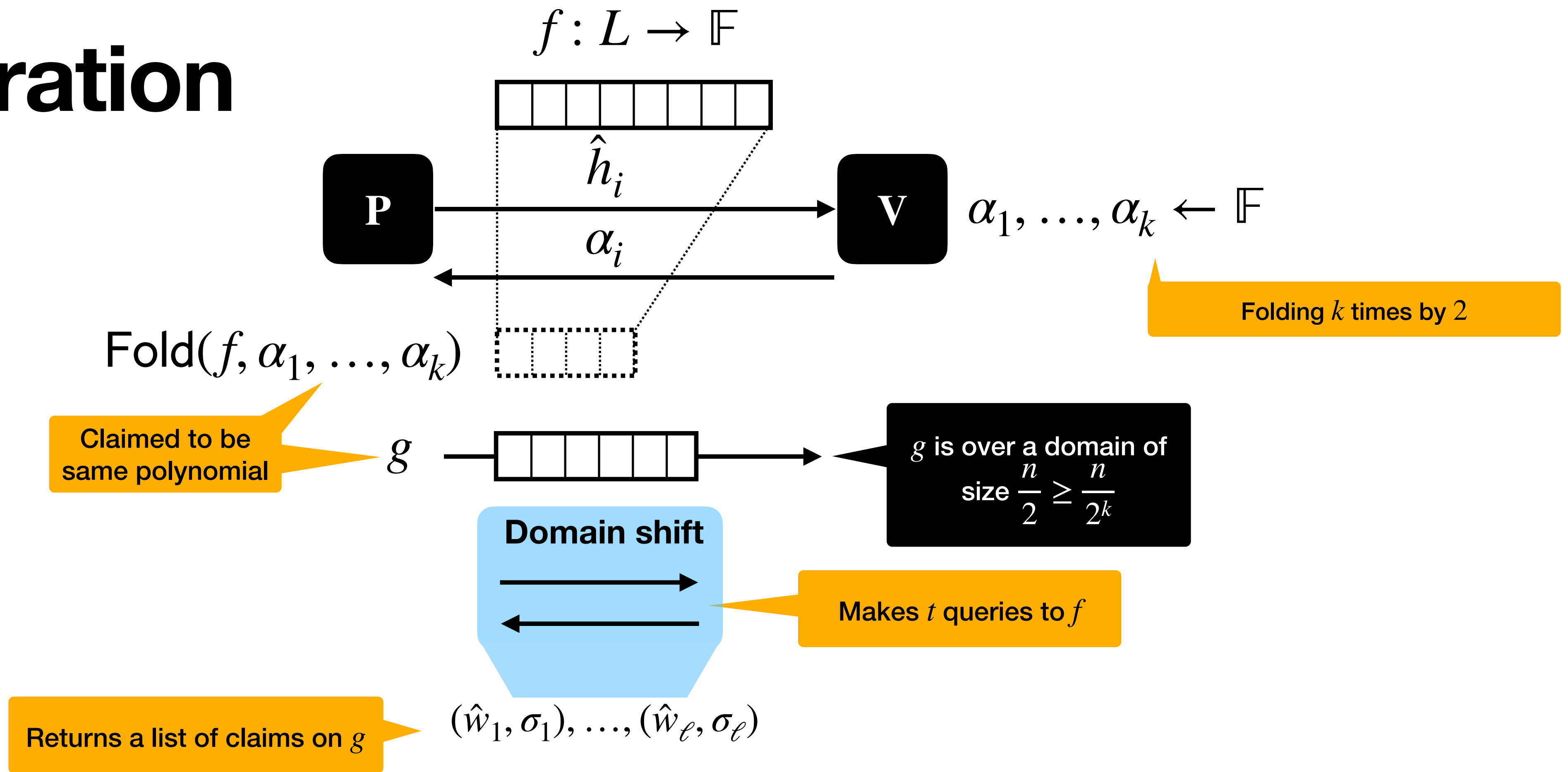
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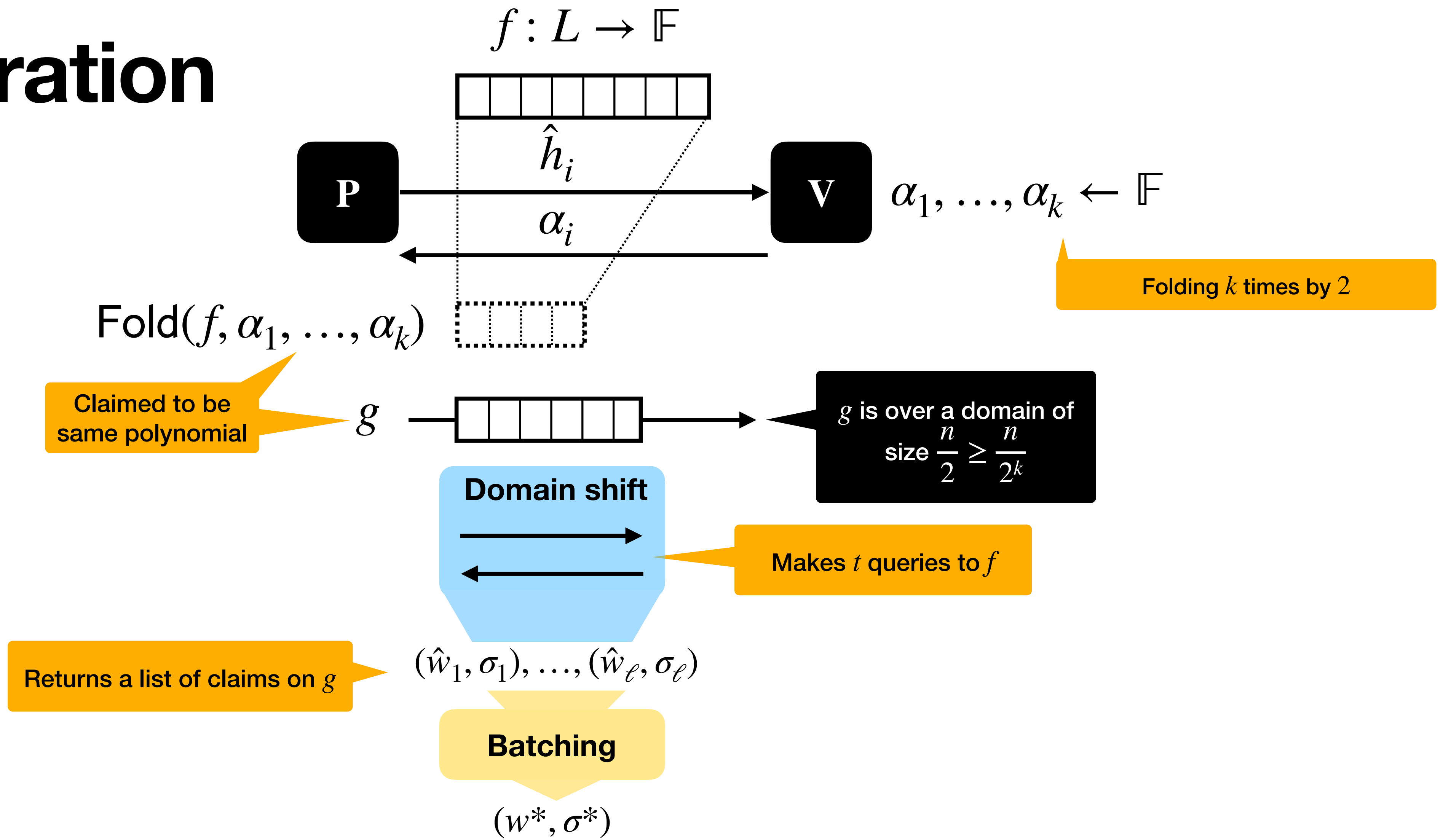
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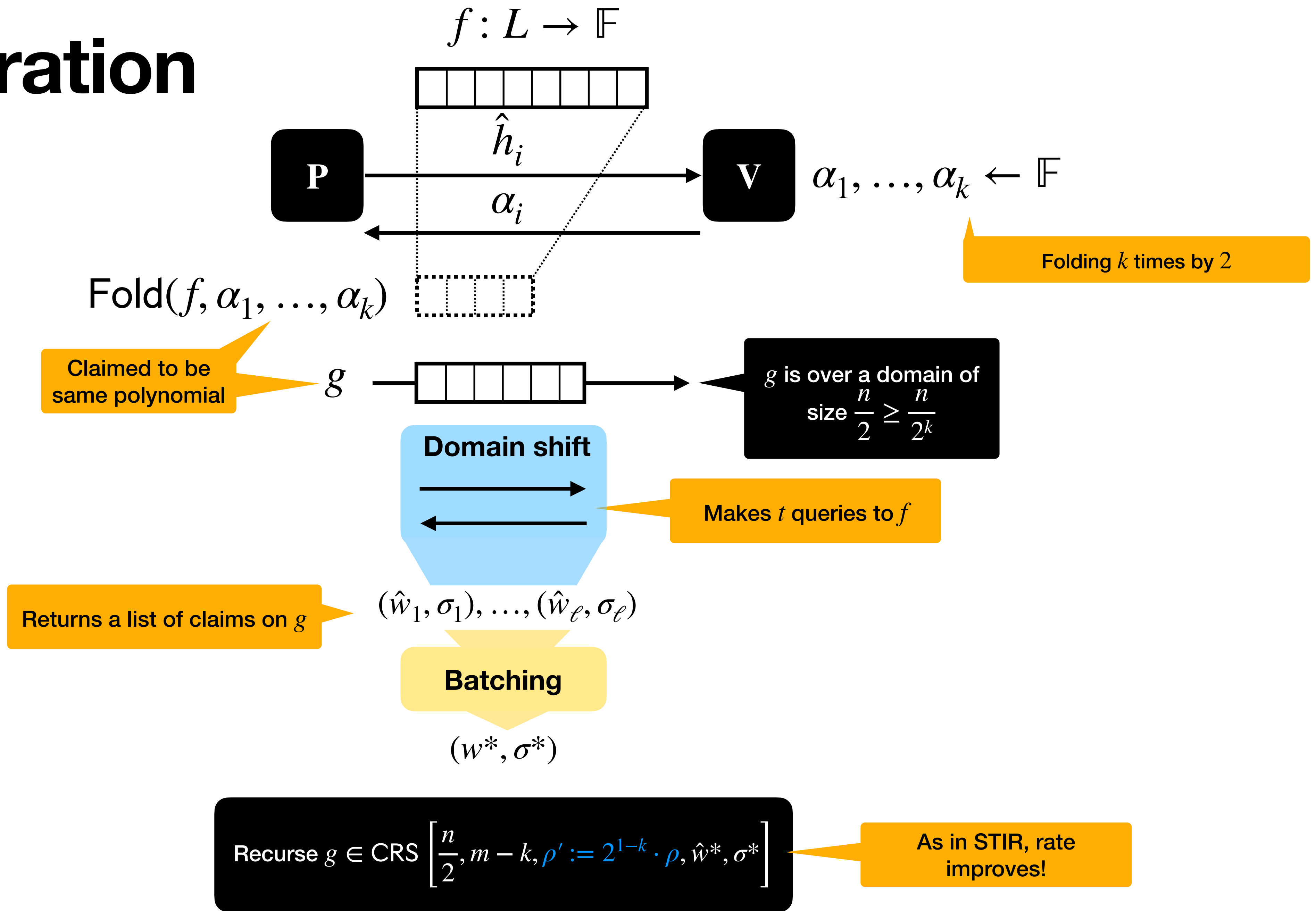


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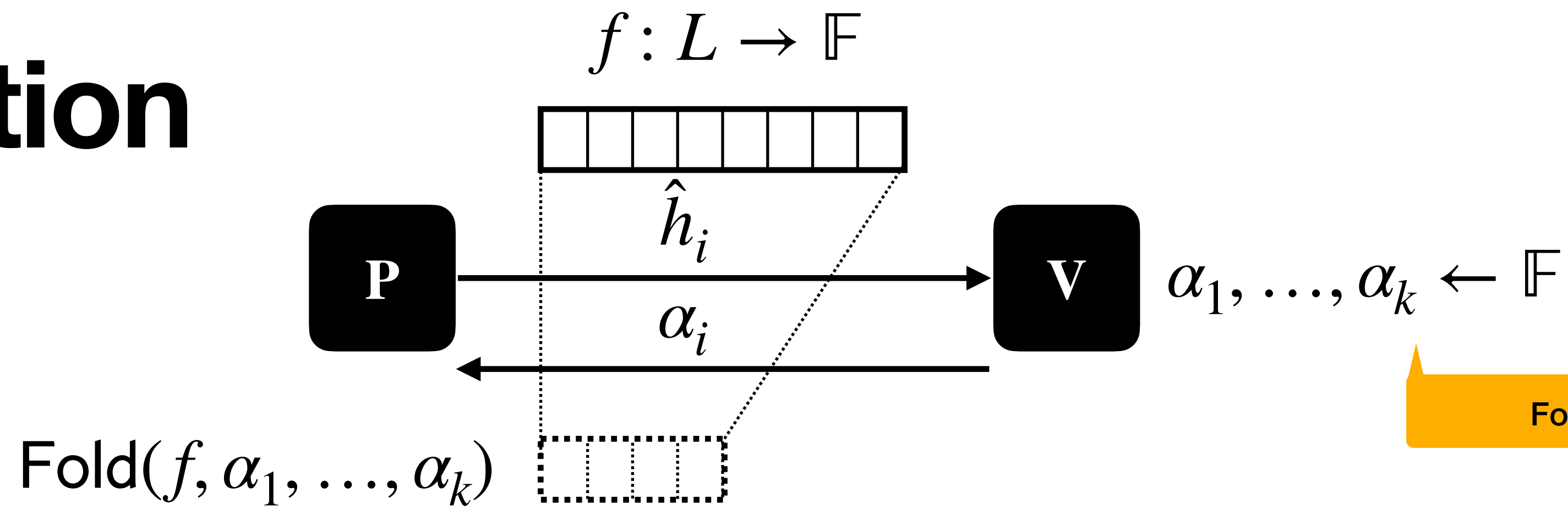




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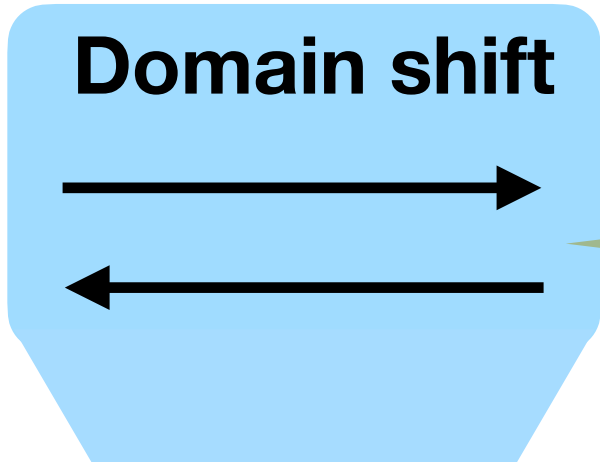
Folding  $k$  times by 2

Claimed to be same polynomial



$g$  is over a domain of size  $\frac{n}{2} \geq \frac{n}{2^k}$

Similar structure to STIR!  
Multilinear structure forbids using quotients: we need new ideas to domain shift!



Makes  $t$  queries to  $f$

Returns a list of claims on  $g$

$(\hat{w}_1, \sigma_1), \dots, (\hat{w}_\ell, \sigma_\ell)$



$(w^*, \sigma^*)$

Recurse  $g \in \text{CRS} \left[ \frac{n}{2}, m - k, \rho' := 2^{1-k} \cdot \rho, \hat{w}^*, \sigma^* \right]$

As in STIR, rate improves!

# Domain shifting

# Domain shifting

**Claim on  $f$ :**  $(\hat{w}, \sigma)$

$$f: L \rightarrow \mathbb{F}$$



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**Output claims on  $g$ :**  
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$f$  and  $g$  claimed to be evaluations of same polynomial. Want to output **claims** on  $g$ .

**Goal:** If  $f$  is  $(1 - \sqrt{\rho})$ -far from  $\text{CRS}[|L|, m, \rho, \hat{w}, \sigma]$ , w.h.p.  $g$  is  $(1 - \sqrt{\rho'})$ -far from  $\text{CRS}[|L^*|, m, \rho', \hat{w}_i, \sigma_i]$  for at least one  $i \in [\ell]$

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OOD subprotocol (next)



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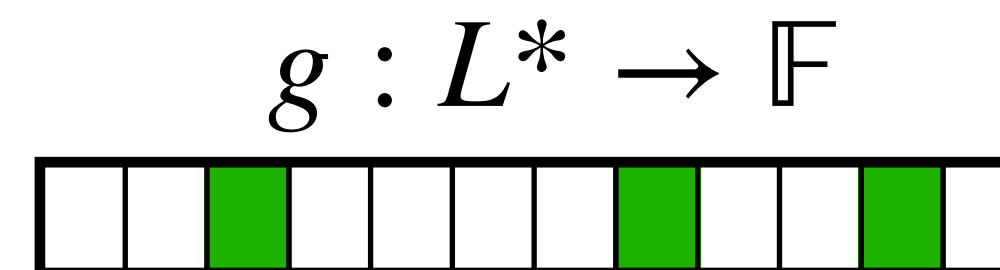
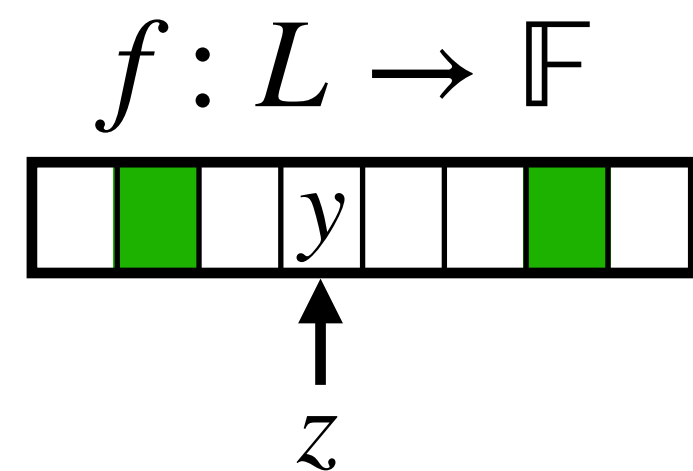
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Then, if  $\hat{p}$  satisfies the  $(\hat{w}, \sigma)$ -constraint  $f$  must be  $(1 - \sqrt{\rho})$ -far from it.

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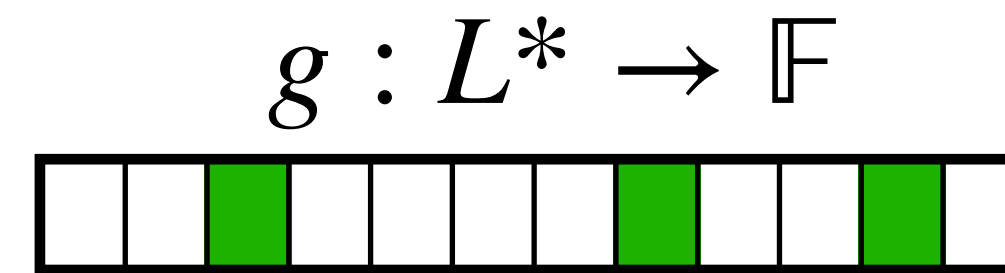
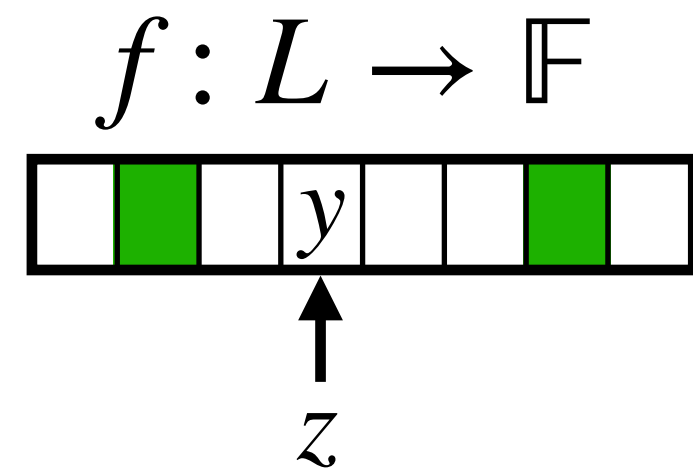
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OOD subprotocol (next)

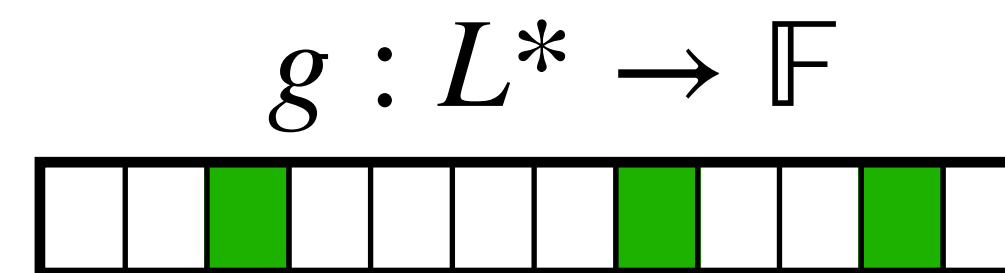
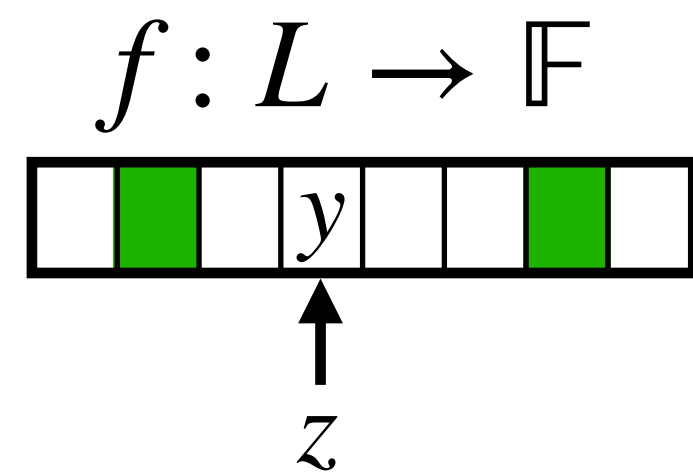
Then, if  $\hat{p}$  satisfies the  $(\hat{w}, \sigma)$ -constraint  $f$  must be  $(1 - \sqrt{\rho})$ -far from it.

**New constraints:** (i) original constraint  $(\hat{w}, \sigma)$  (ii)  $\hat{p}(z) = y$  for some random point  $z$ .

Just an evaluation constraint which we know how to handle!

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**New constraints:** (i) original constraint  $(\hat{w}, \sigma)$  (ii)  $\hat{p}(z) = y$  for some random point  $z$ .

Just an evaluation constraint which we know how to handle!

So, except with probability  $\sqrt{\rho}$ ,  $g$  is  $(1 - \sqrt{\rho'})$ -far from  $\text{CRS}[|L^*|, m, \rho', (\hat{w}_1, \sigma_1), \dots, (\hat{w}_\ell, \sigma_\ell)]$ .

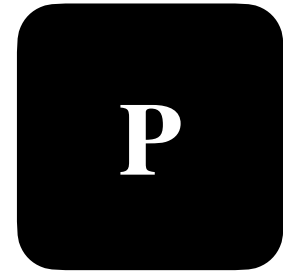
Can amplify to  $\sqrt{\rho}^t$

# **Out Of Domain**

## **Subprotocol to force unique**

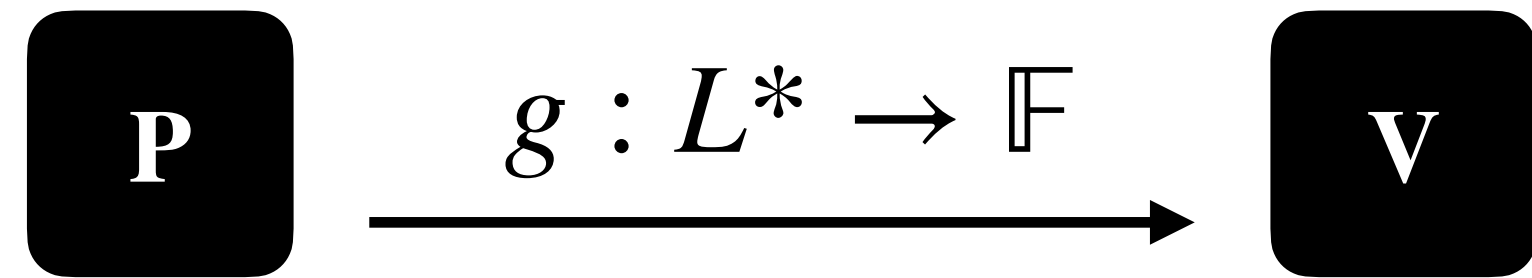
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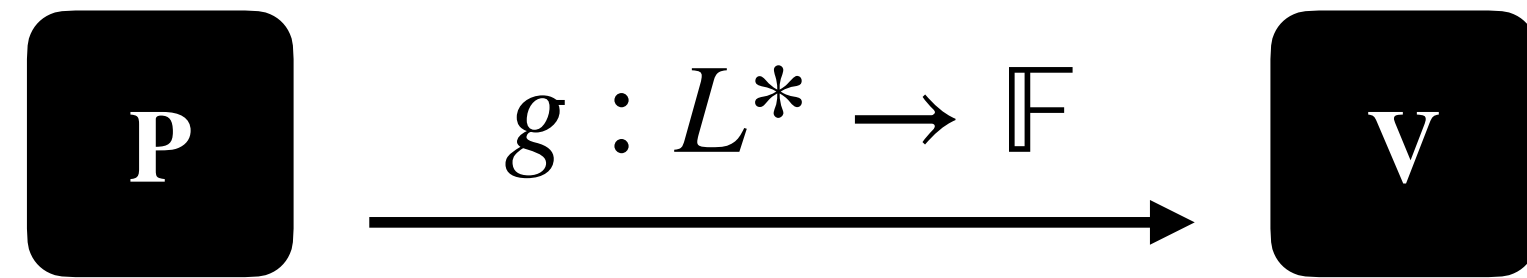
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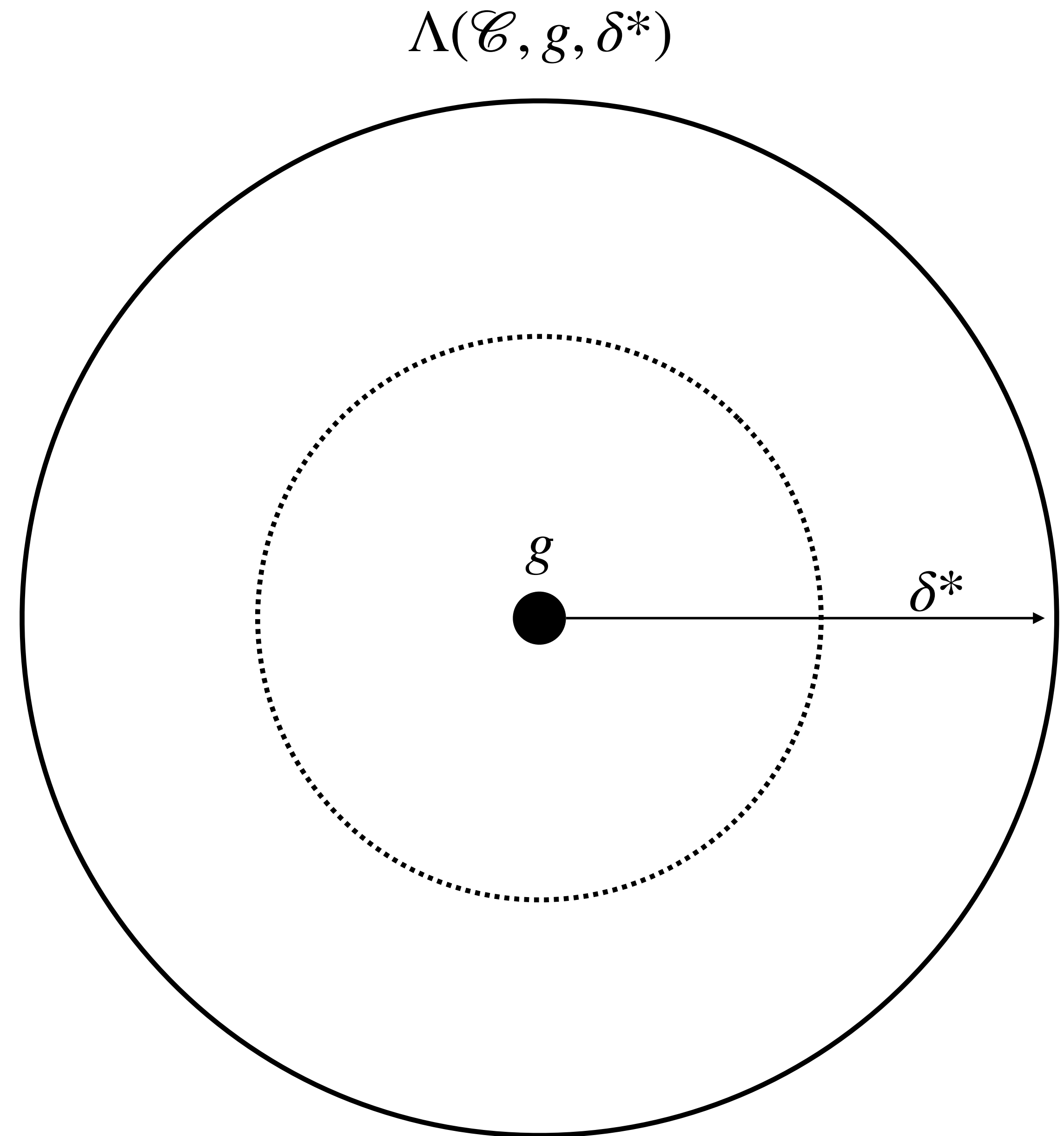
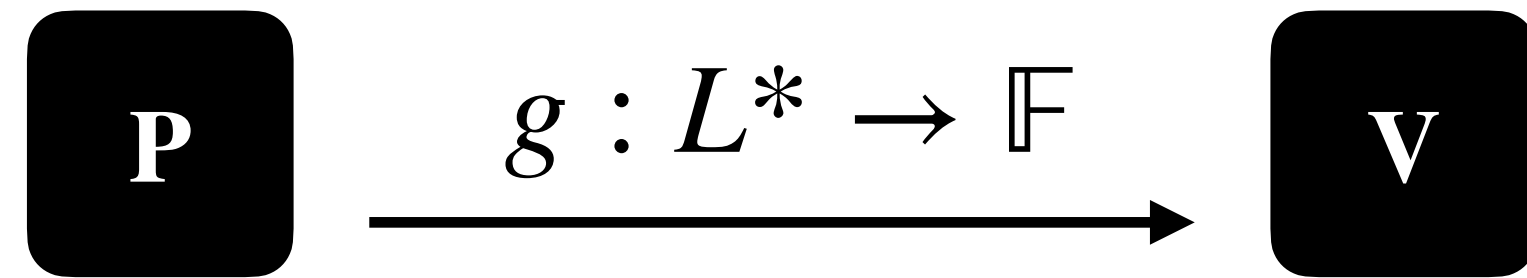
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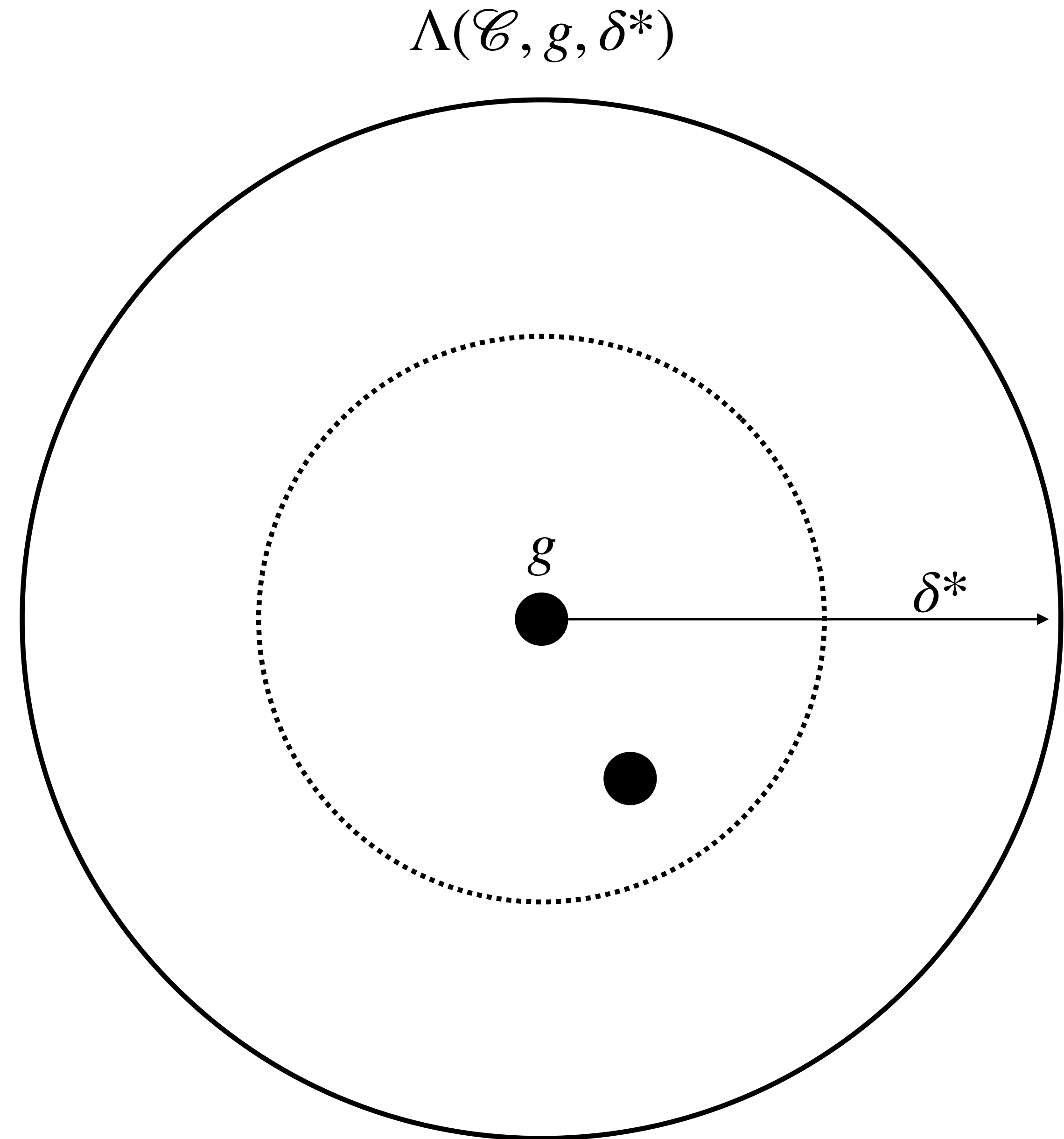
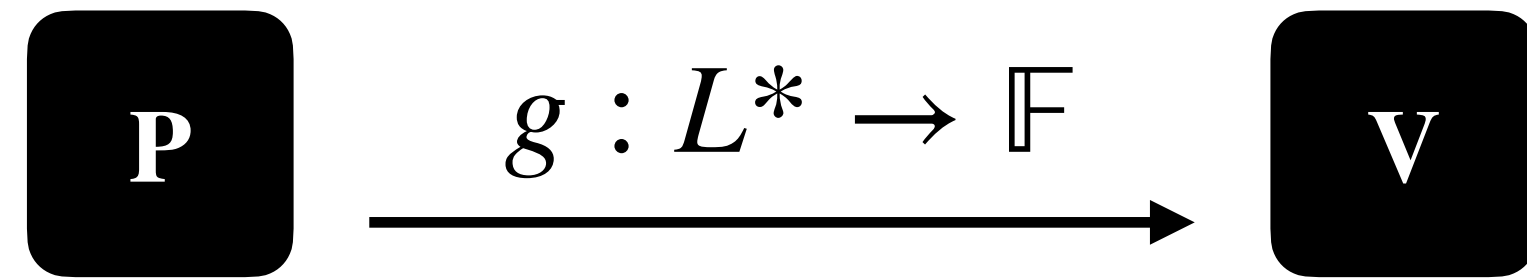
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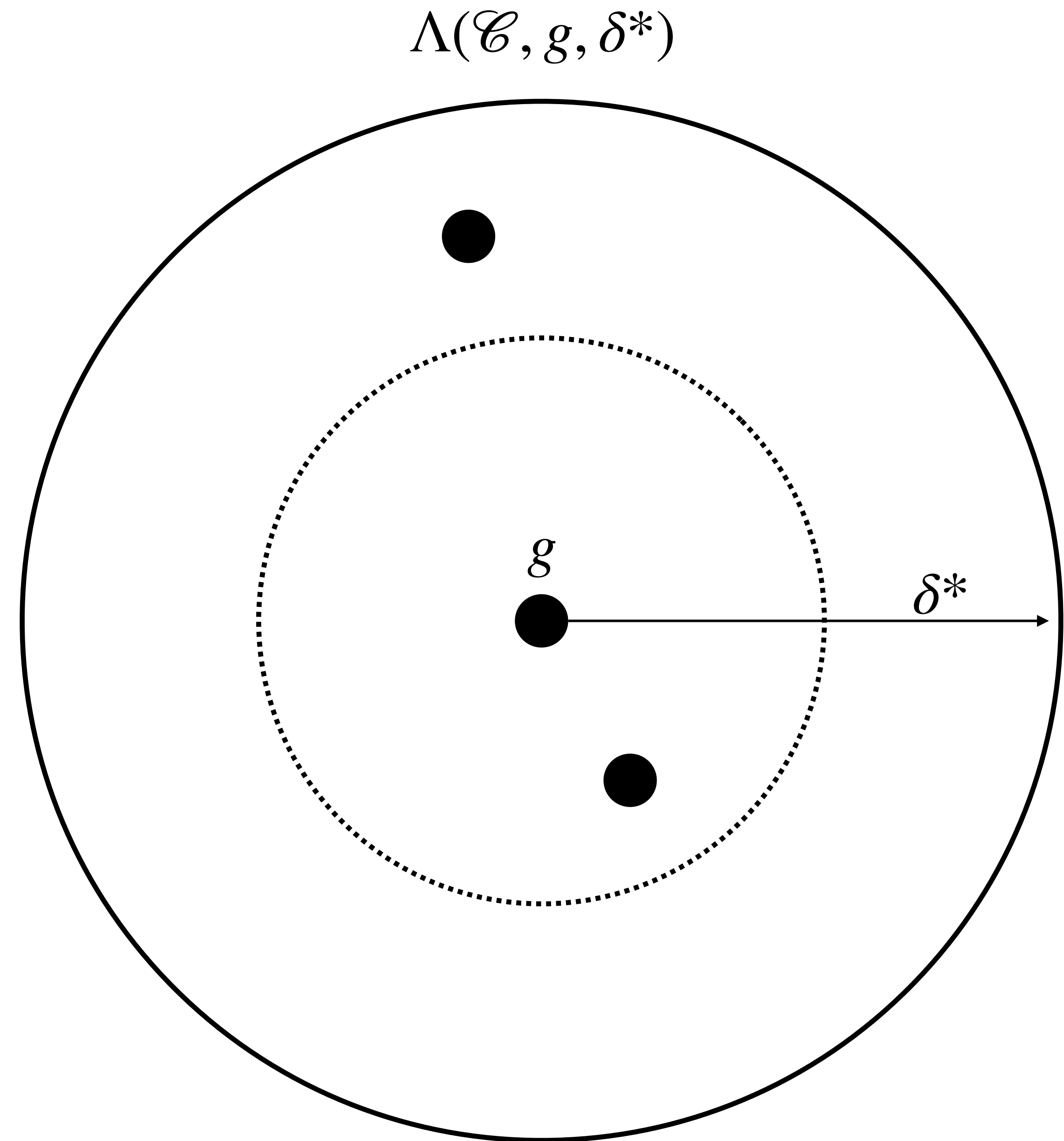
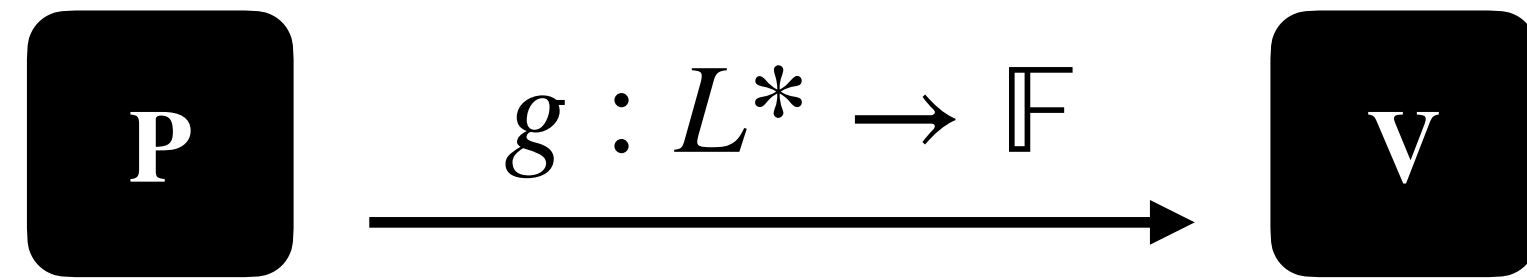
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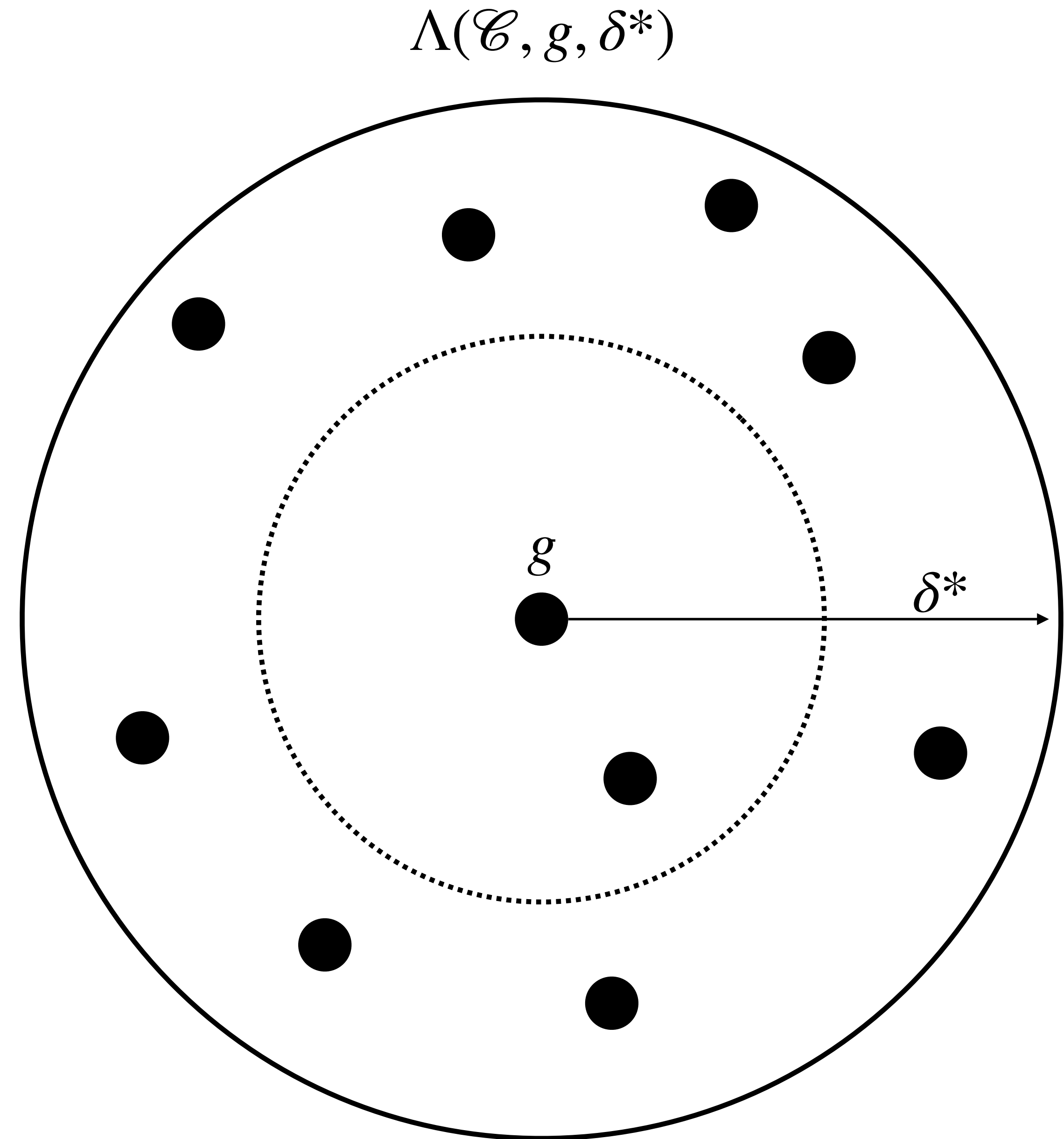
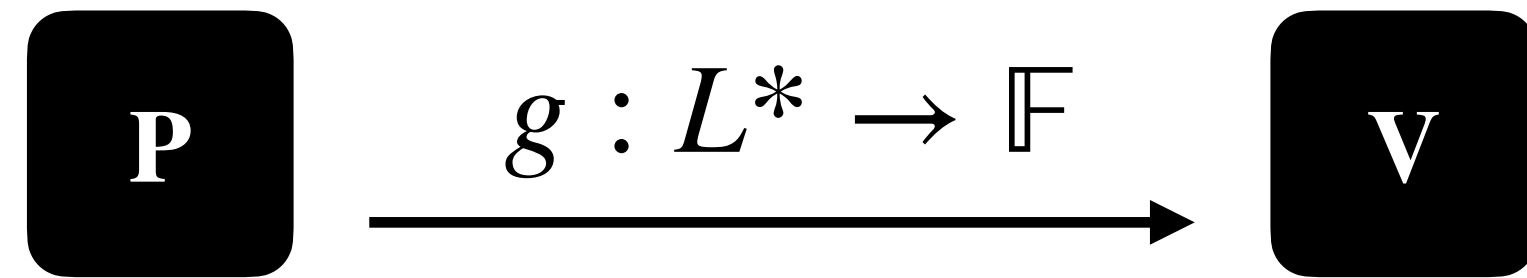
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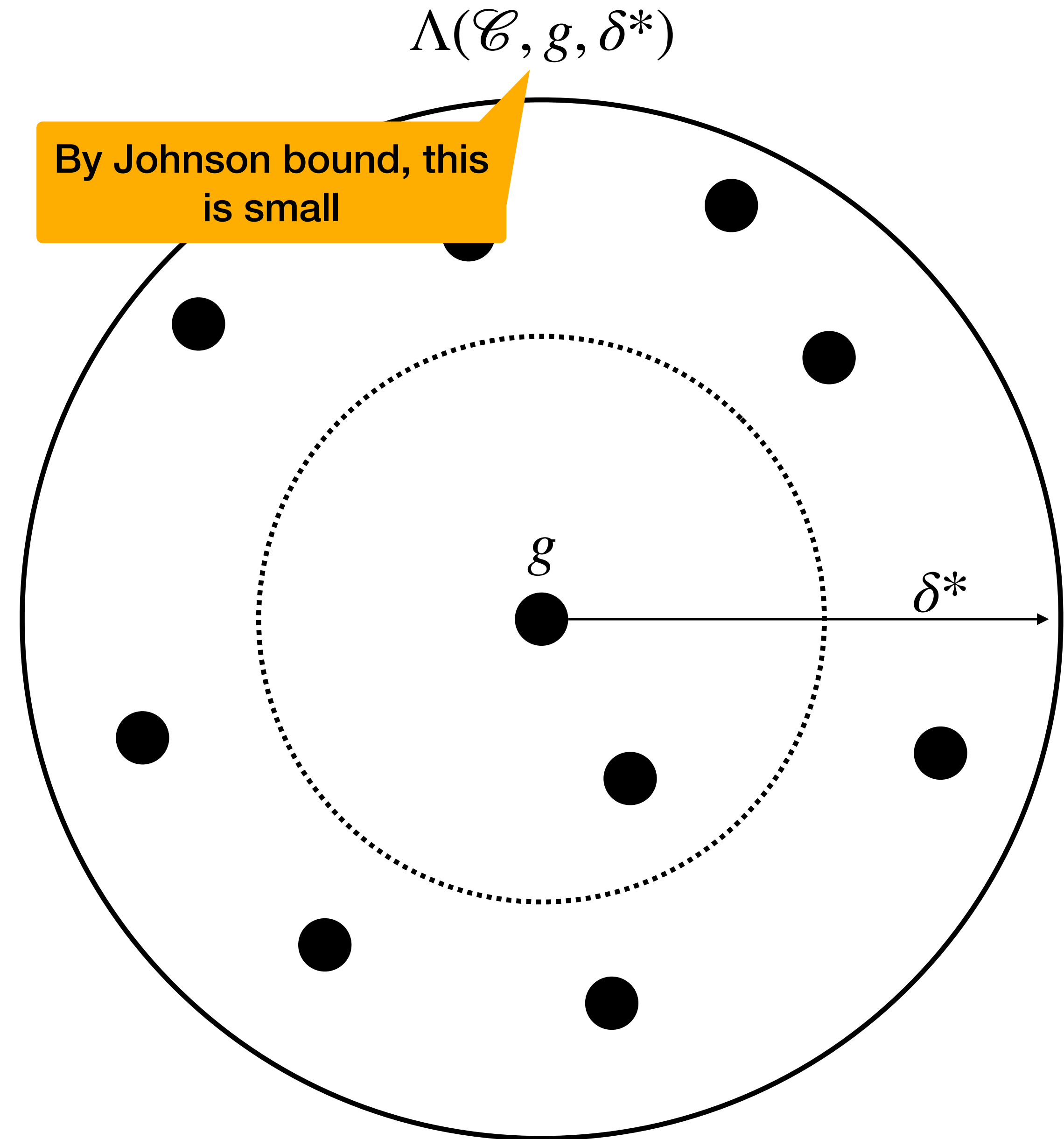
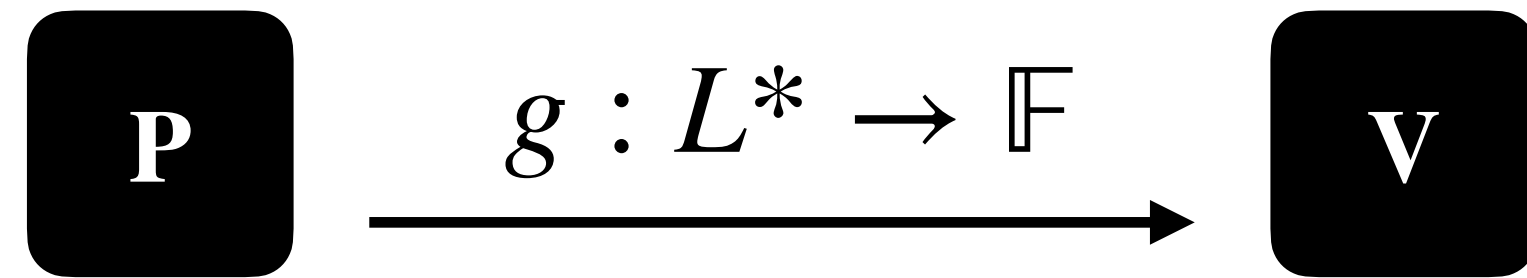
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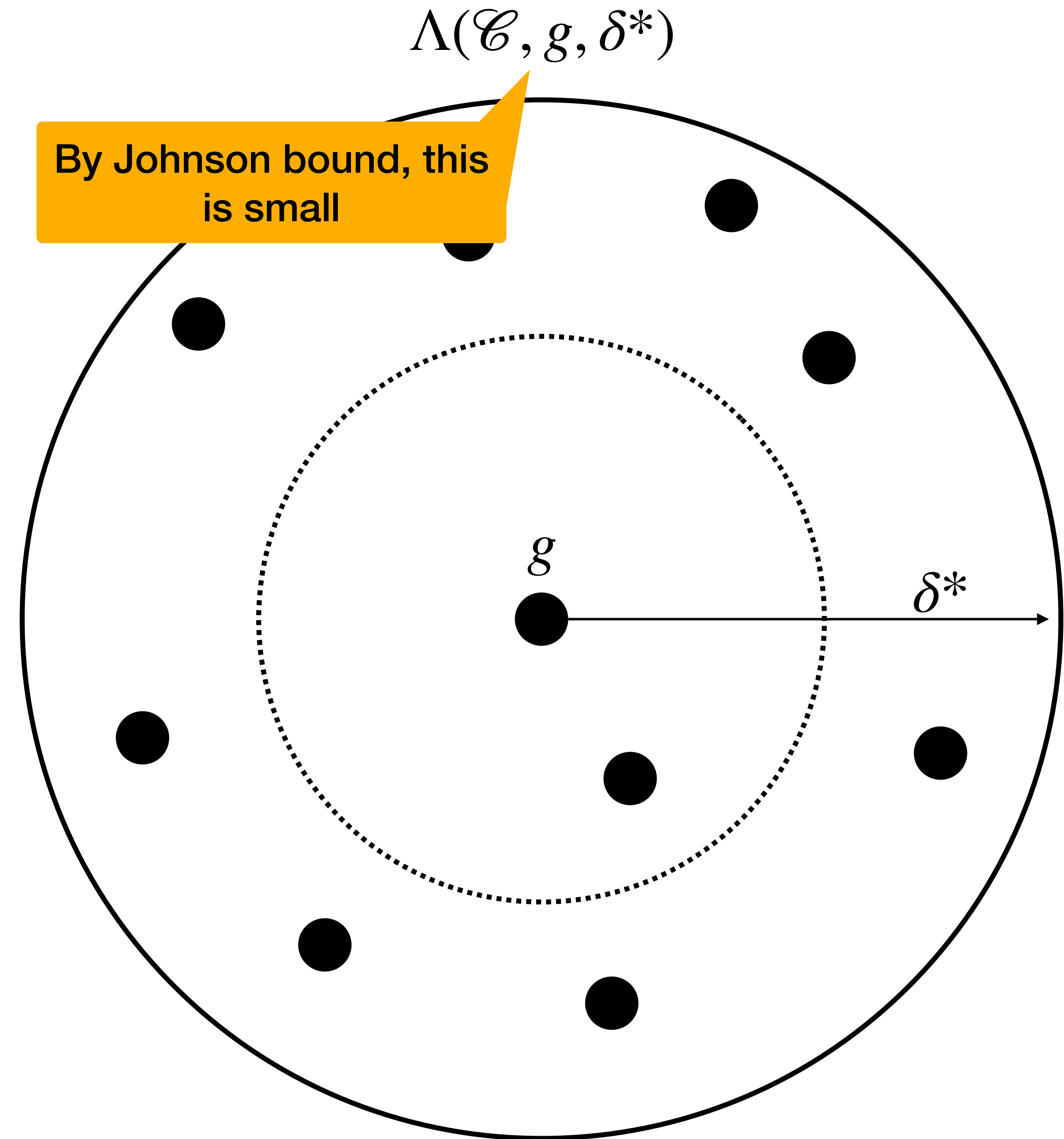
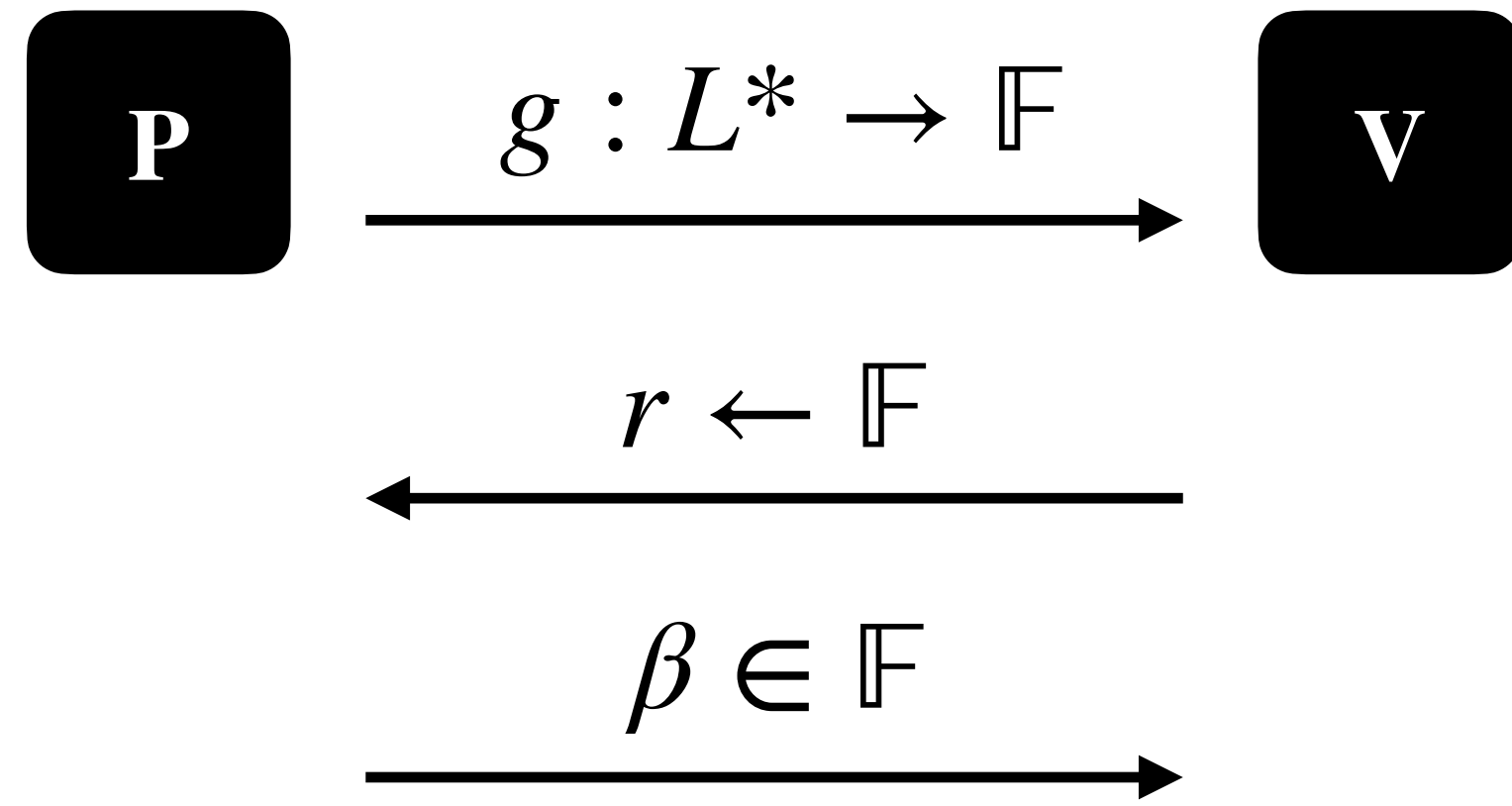
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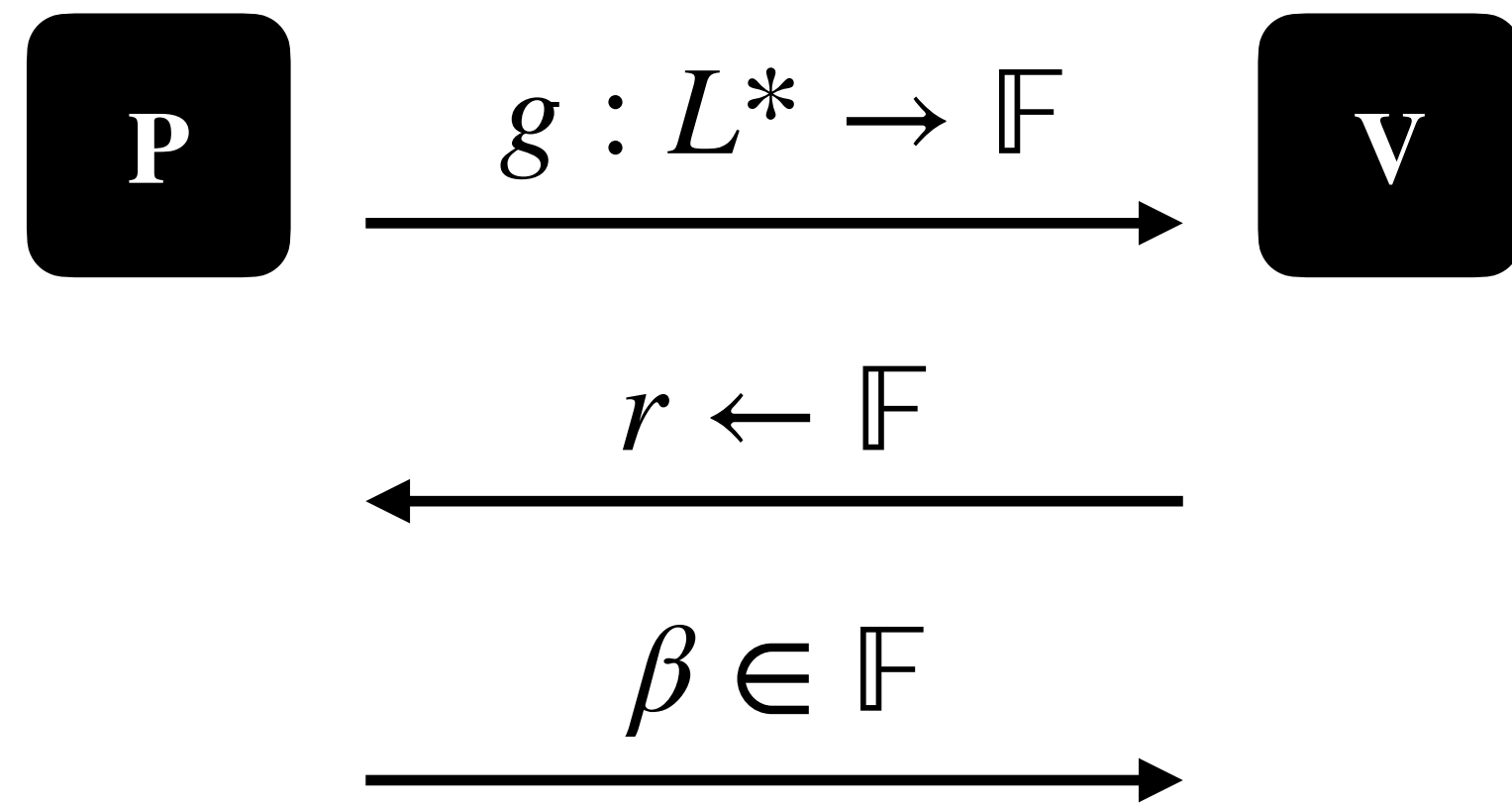
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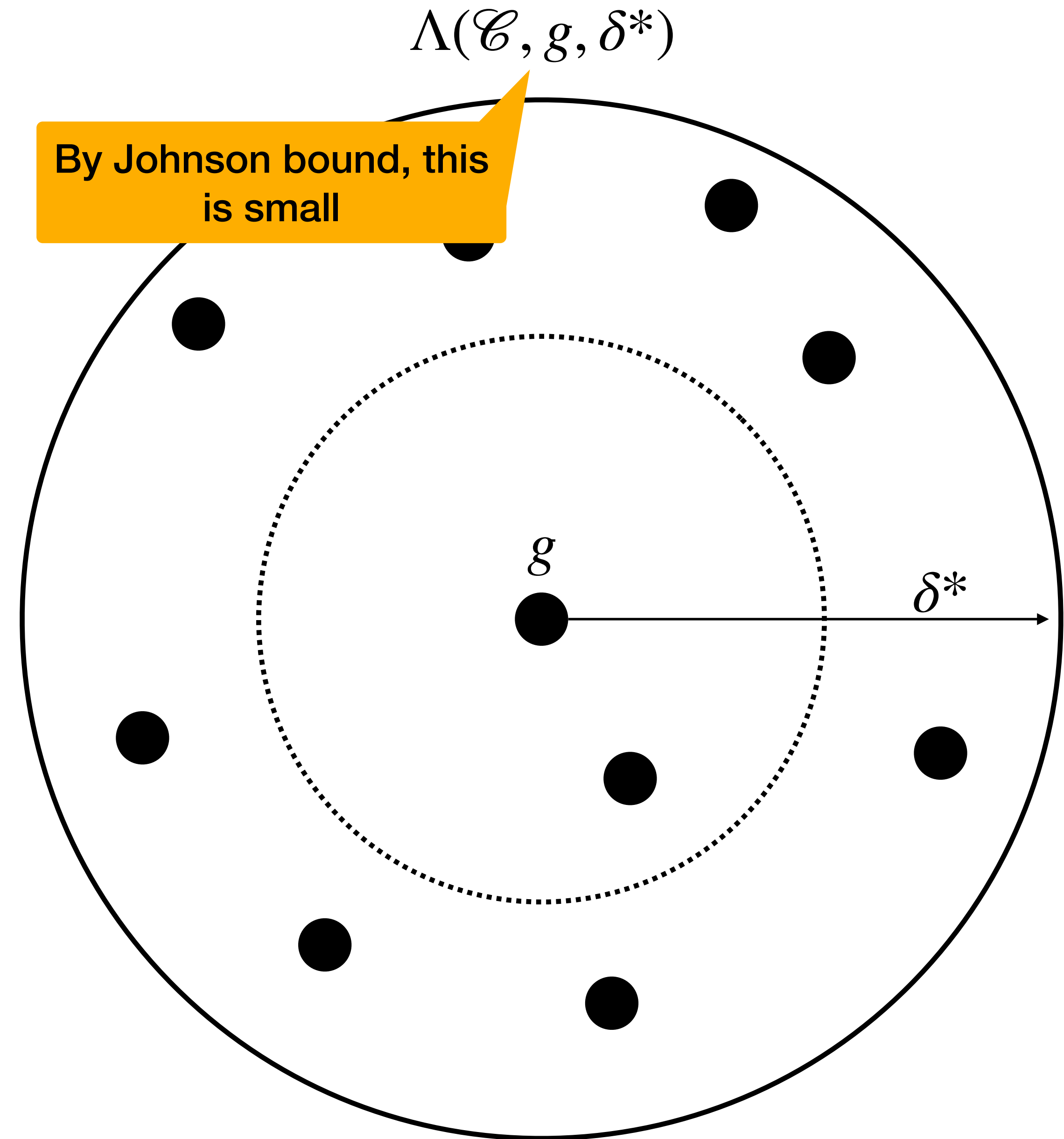


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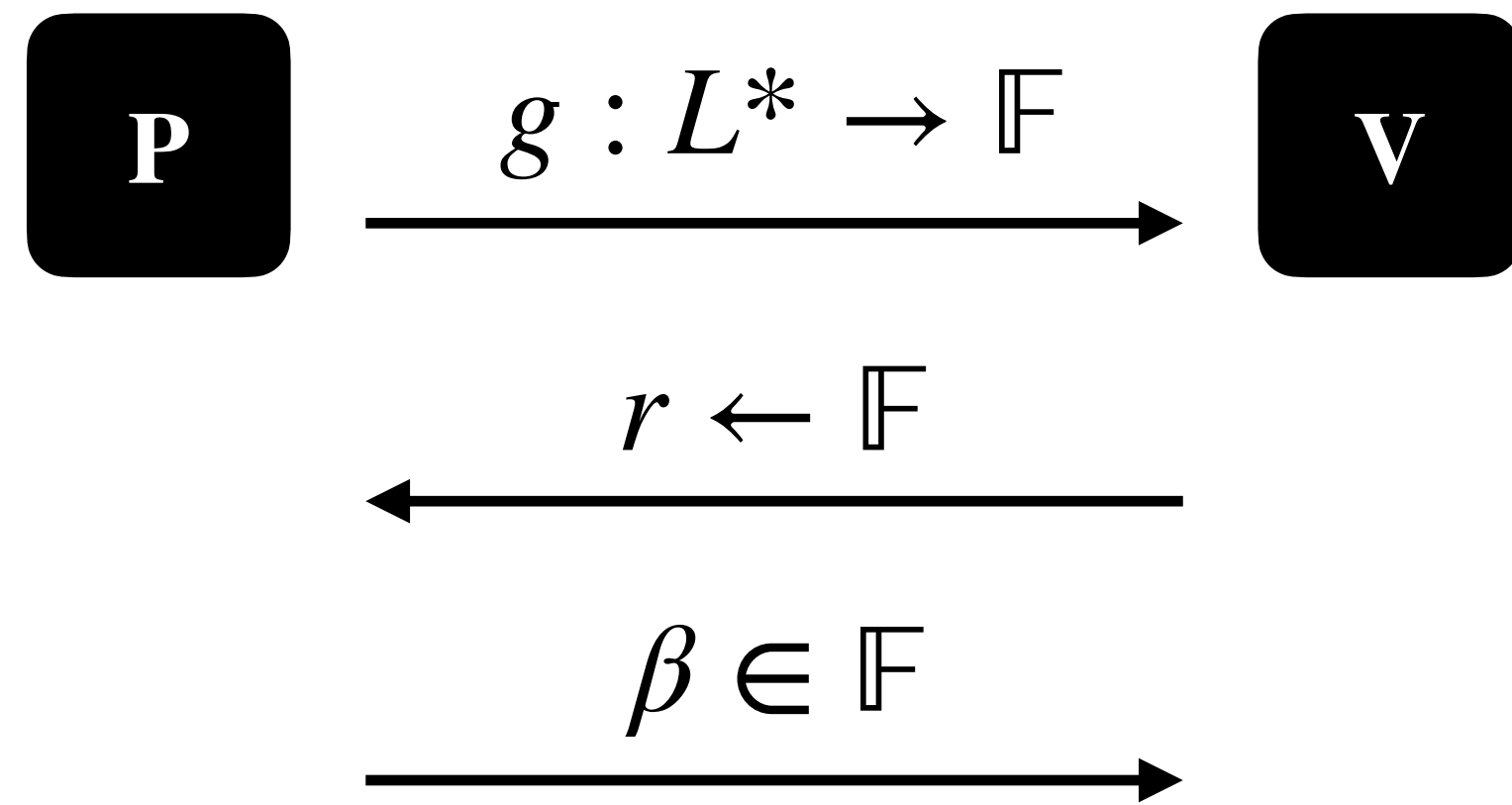


- By fundamental theorem of algebra of w.h.p. no pair  $\hat{u}, \hat{v}$  with  $\hat{u}(r) = \hat{v}(r)$
- Prover "chooses" which codeword  $\hat{u}$  it "commits" to

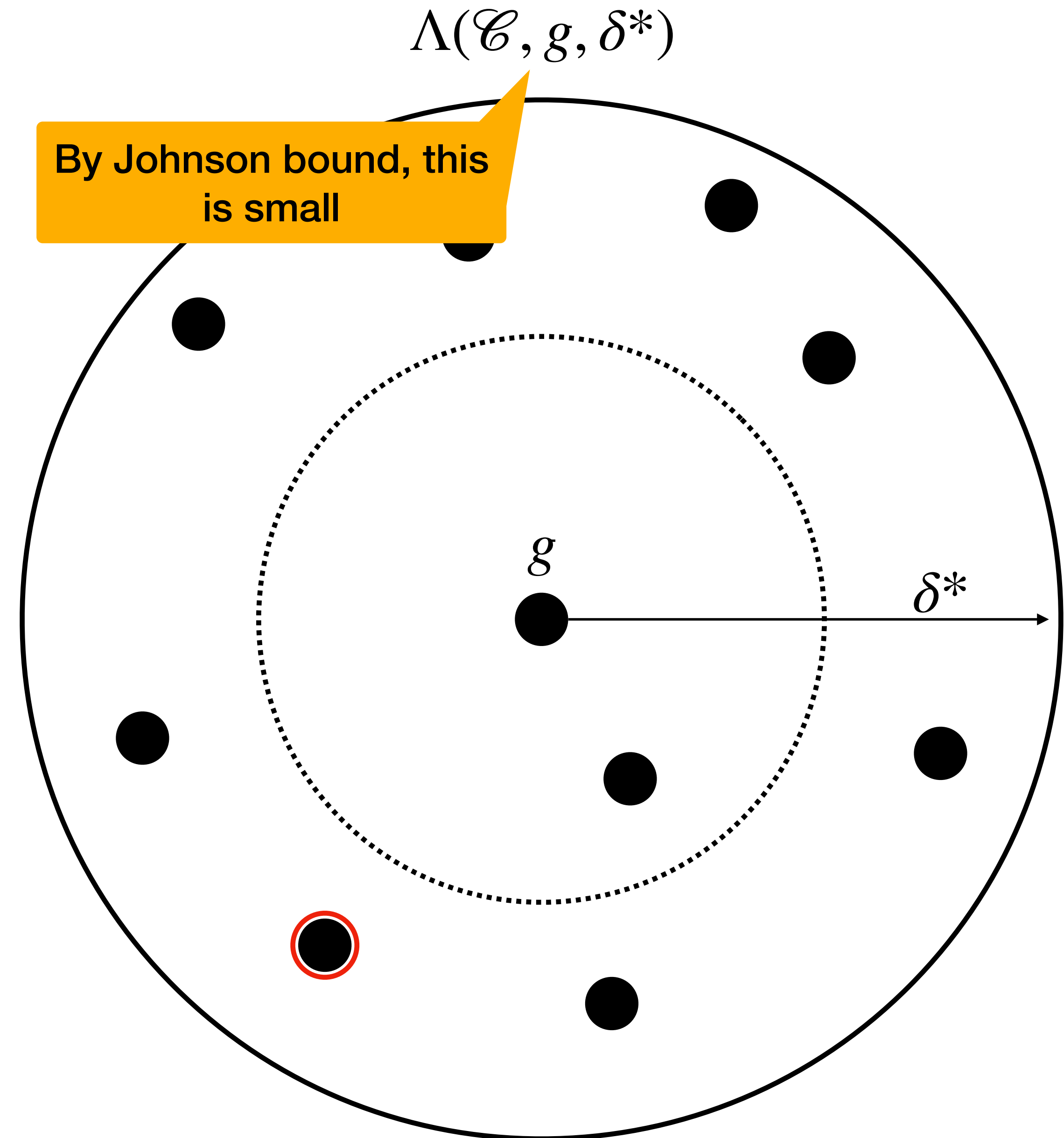


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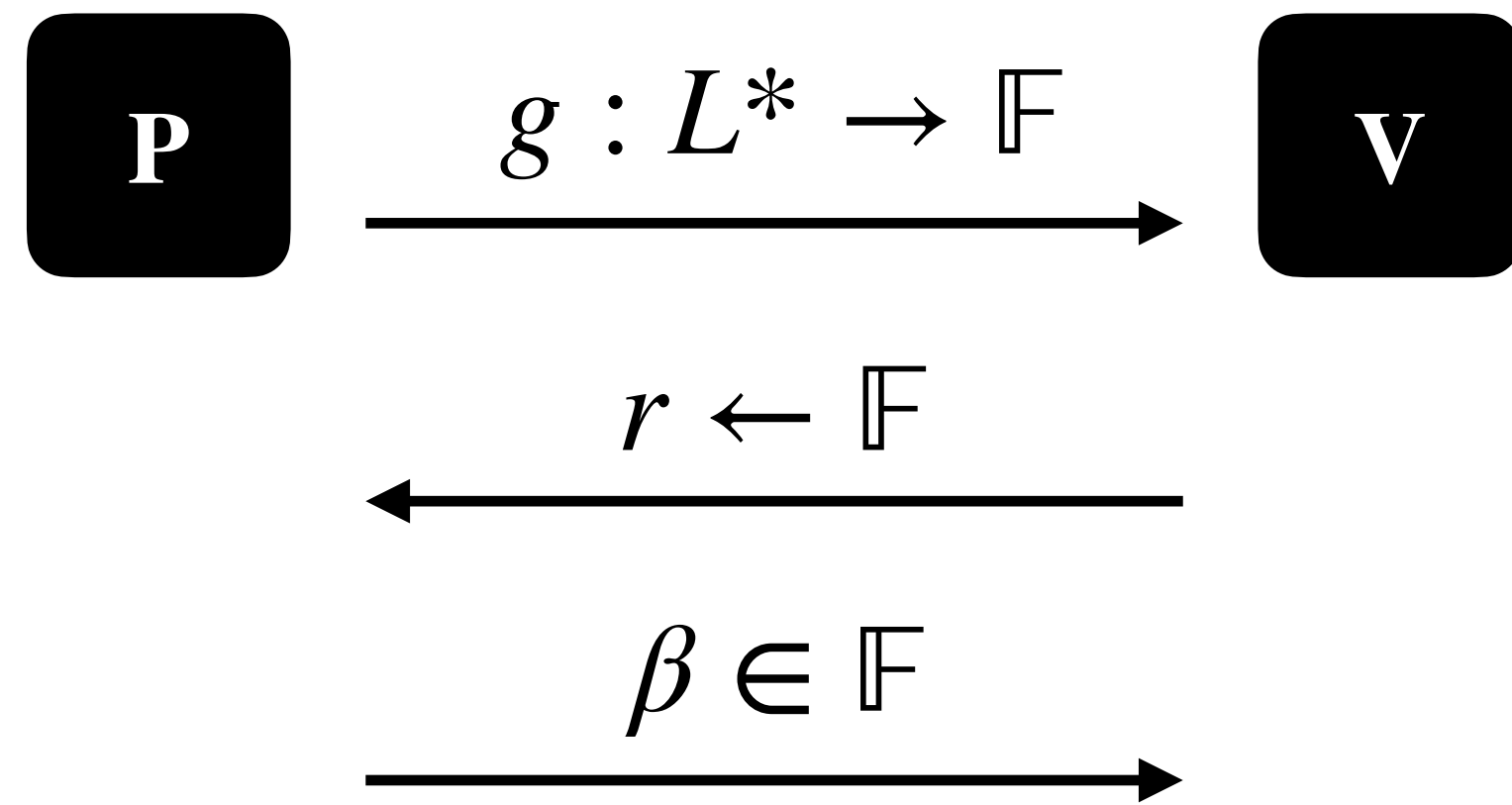
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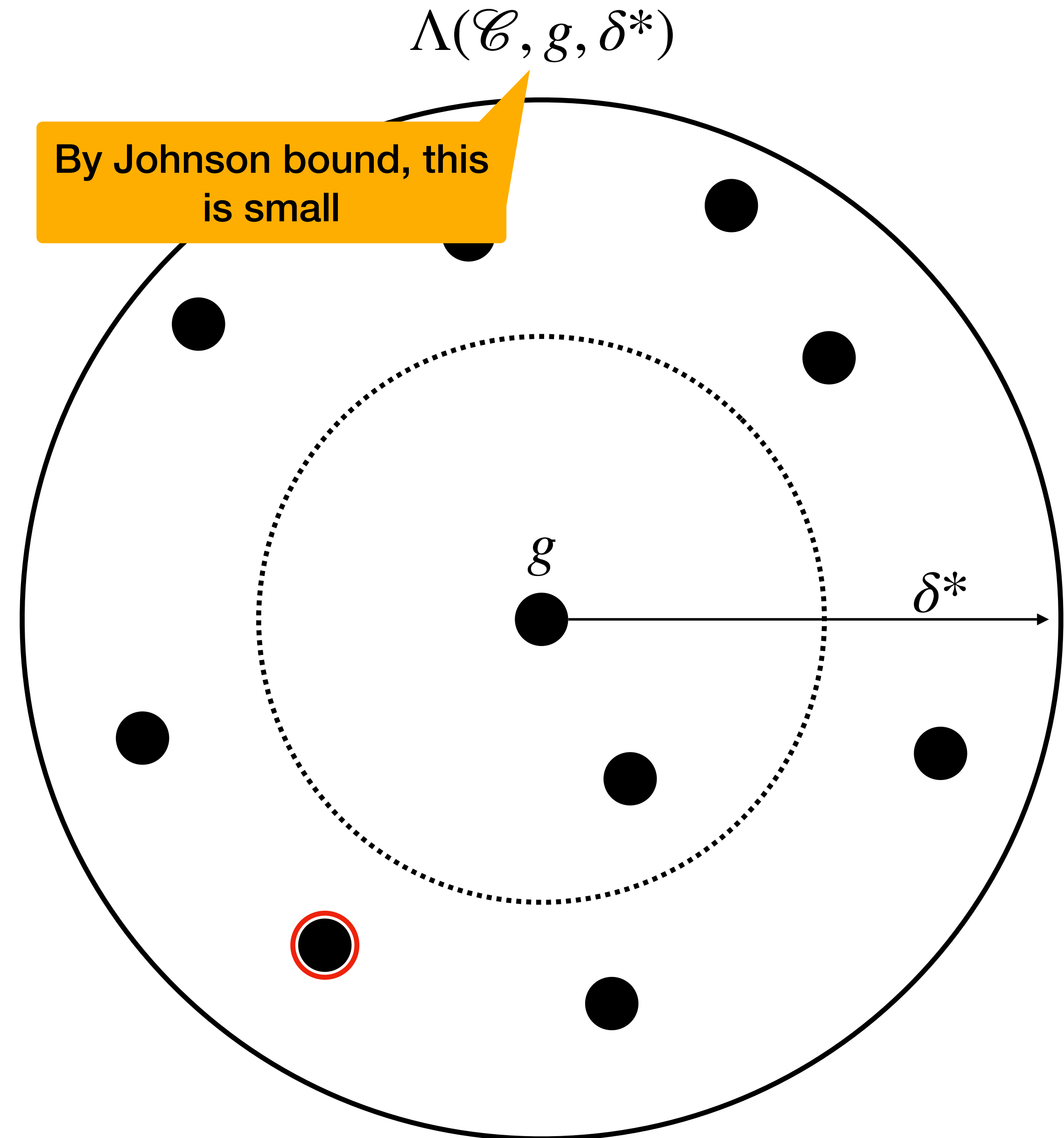
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Add to list of constraints to **enforce!**



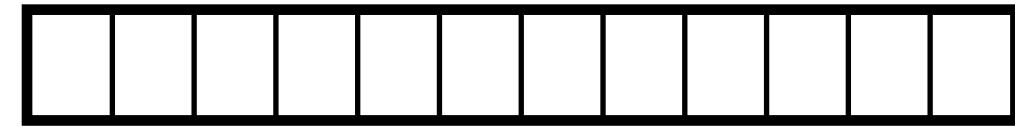
# Batching

**Pick your favourite sumcheck batching**

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$$g : L \rightarrow \mathbb{F}$$



**Sumcheck claims on  $g$ :**

$$(\hat{w}_1, \sigma_1), \dots, (\hat{w}_\ell, \sigma_\ell)$$

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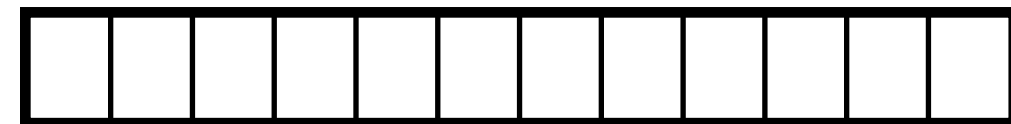
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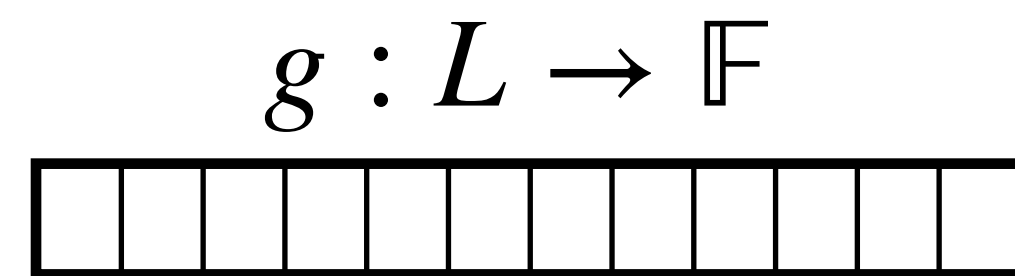
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Sumcheck claim on  $g$ :  $(\hat{w}^*, \sigma^*)$

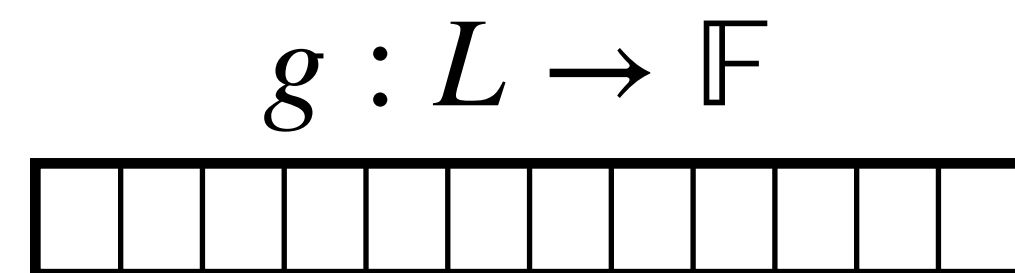
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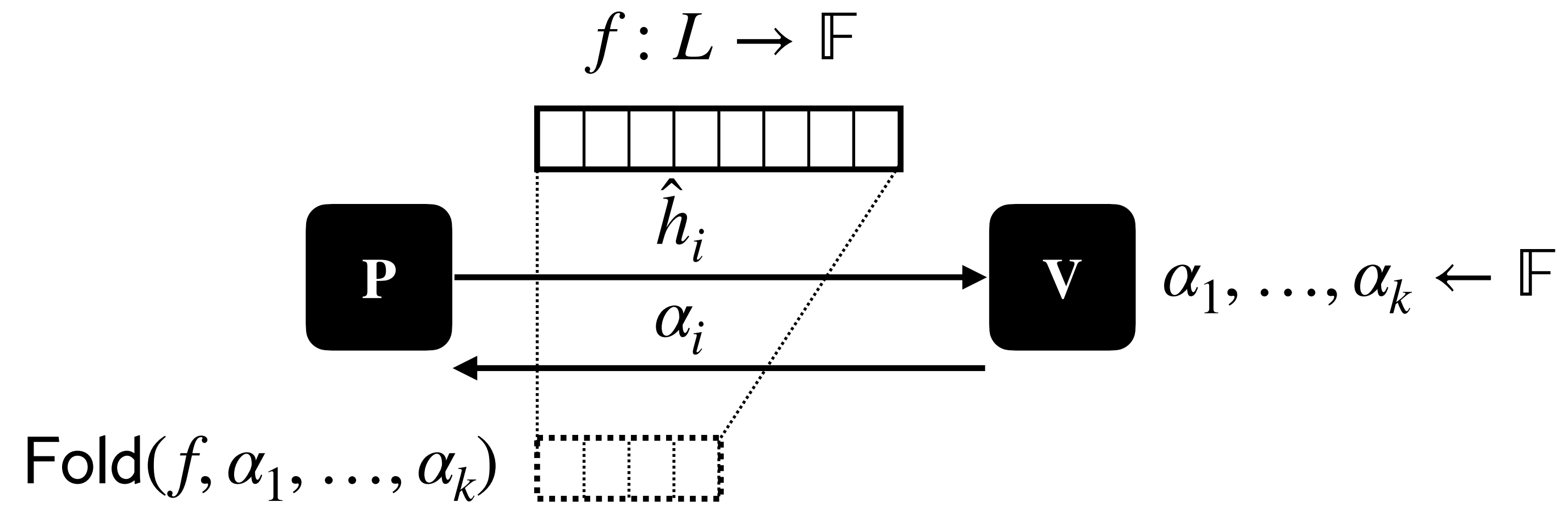


Sumcheck claim on  $g$ :  $(\hat{w}^*, \sigma^*)$

Many ways this can be done: **we chose random linear combination.**

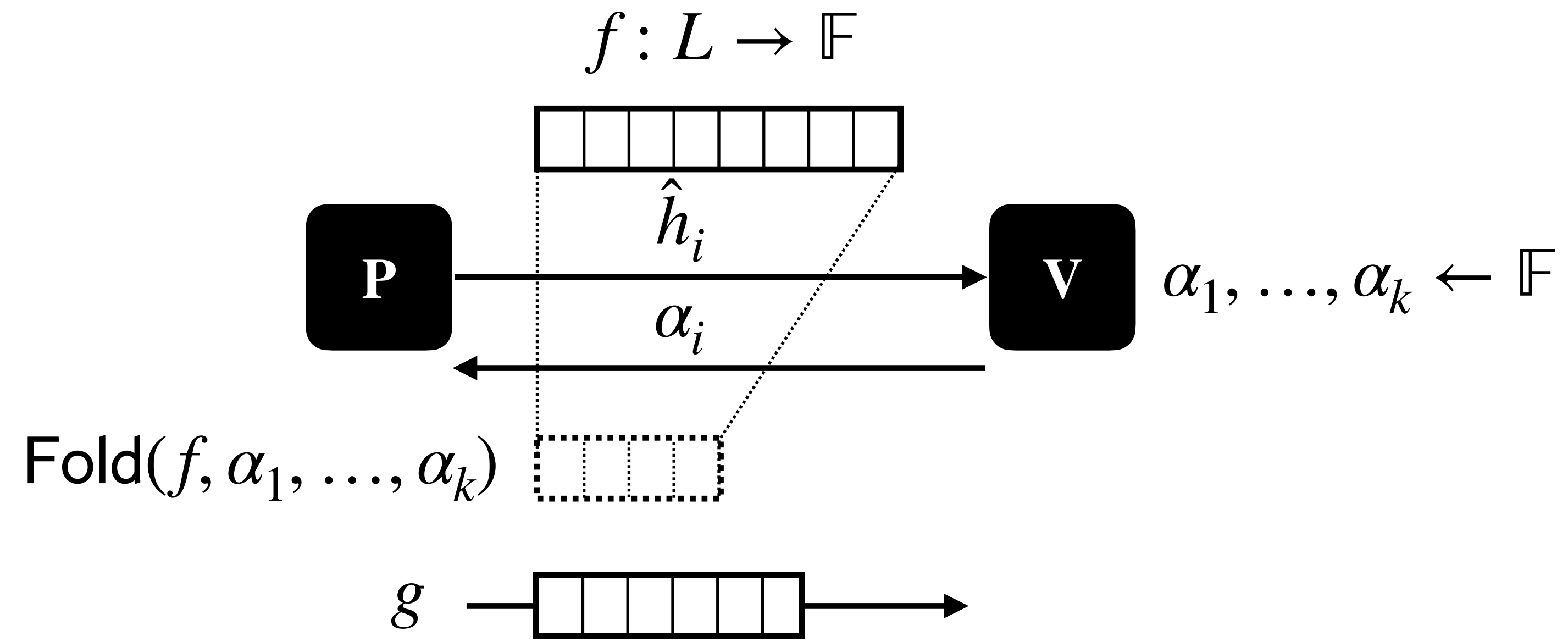
**WHIR** 

# WHIR

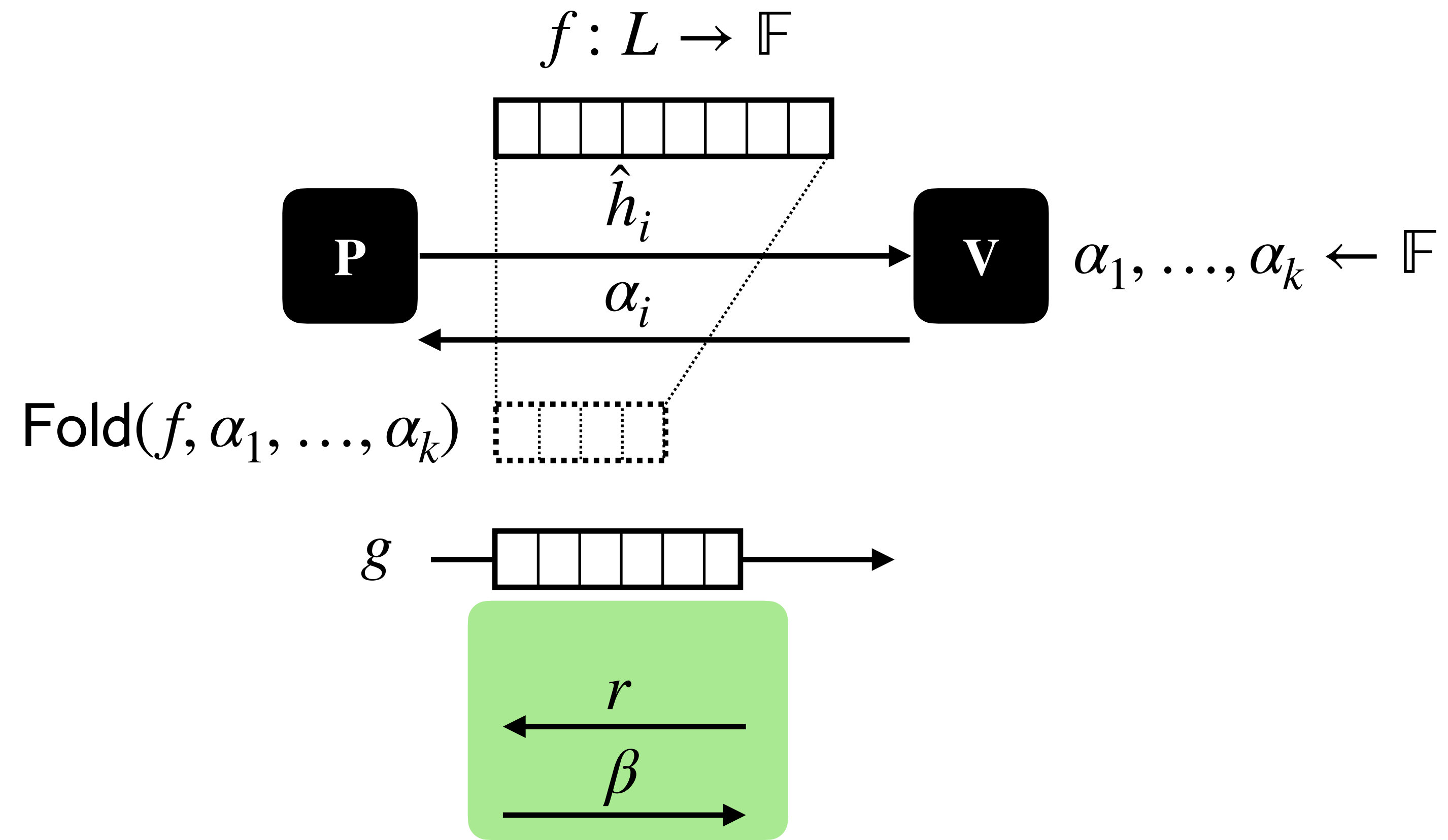




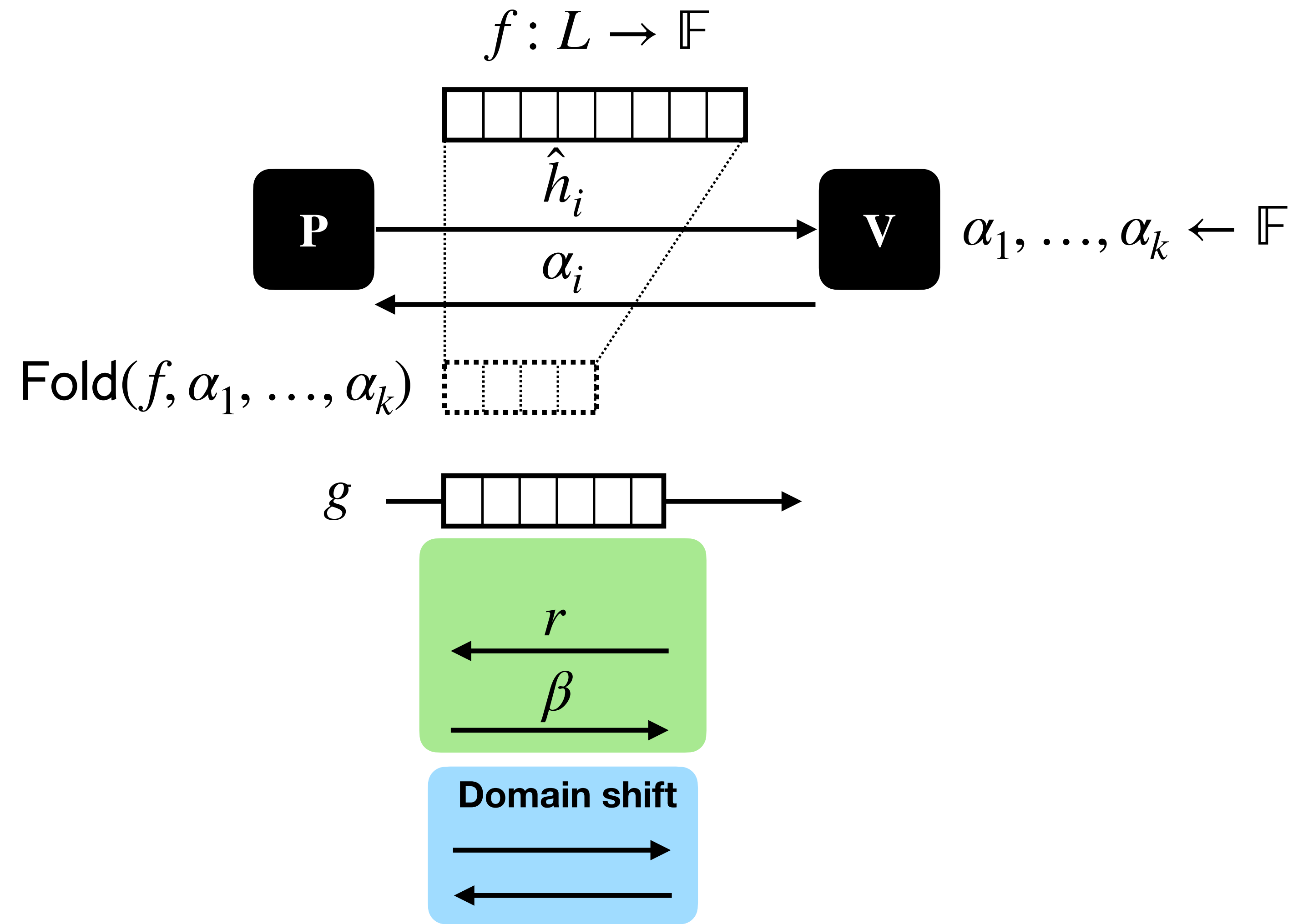
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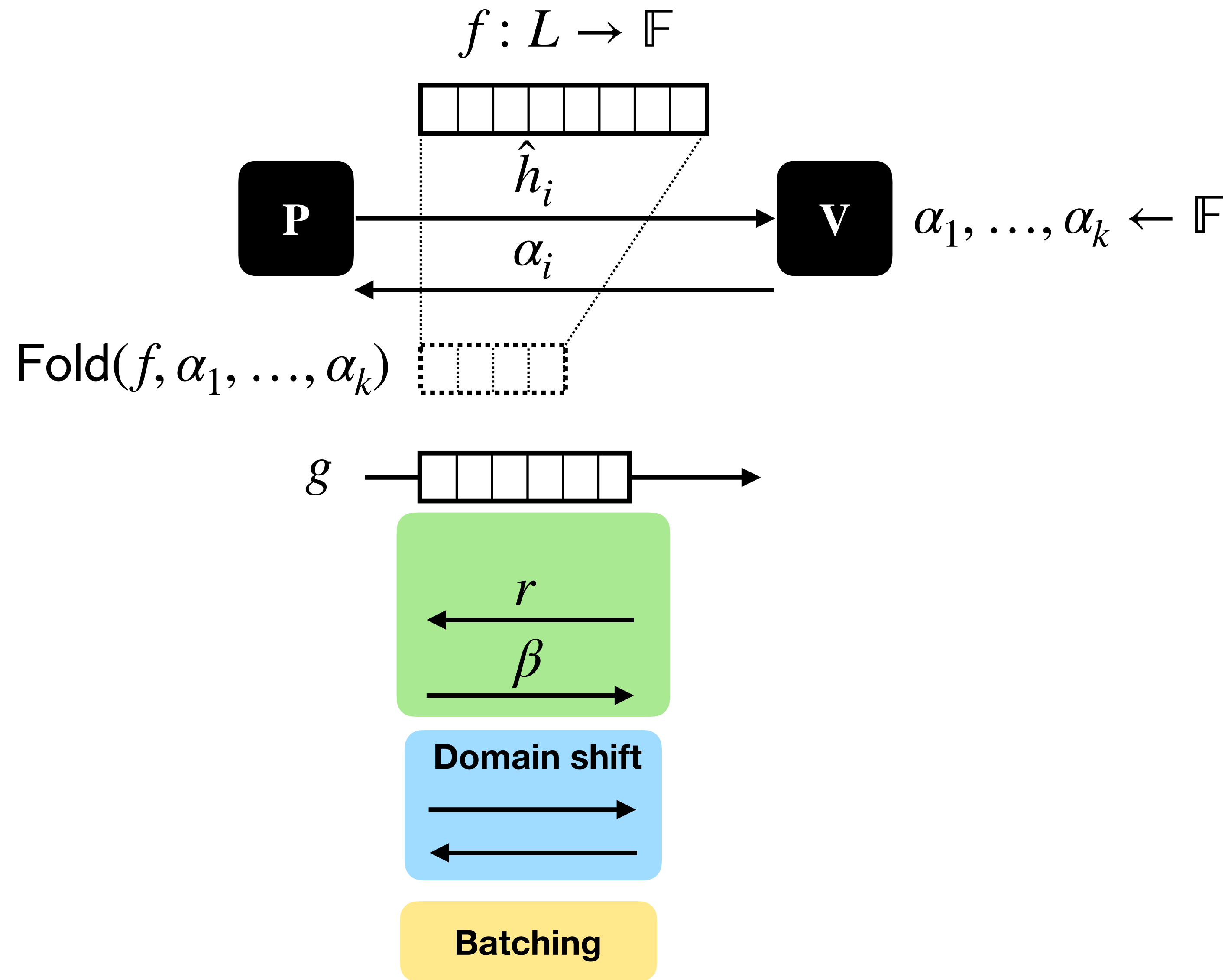
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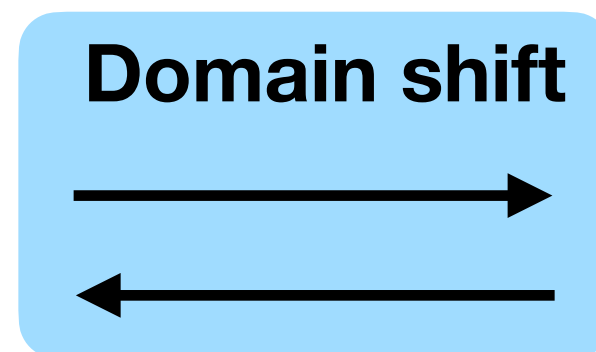
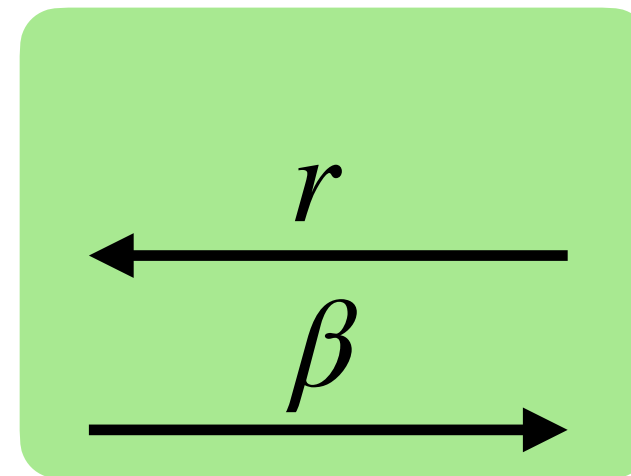
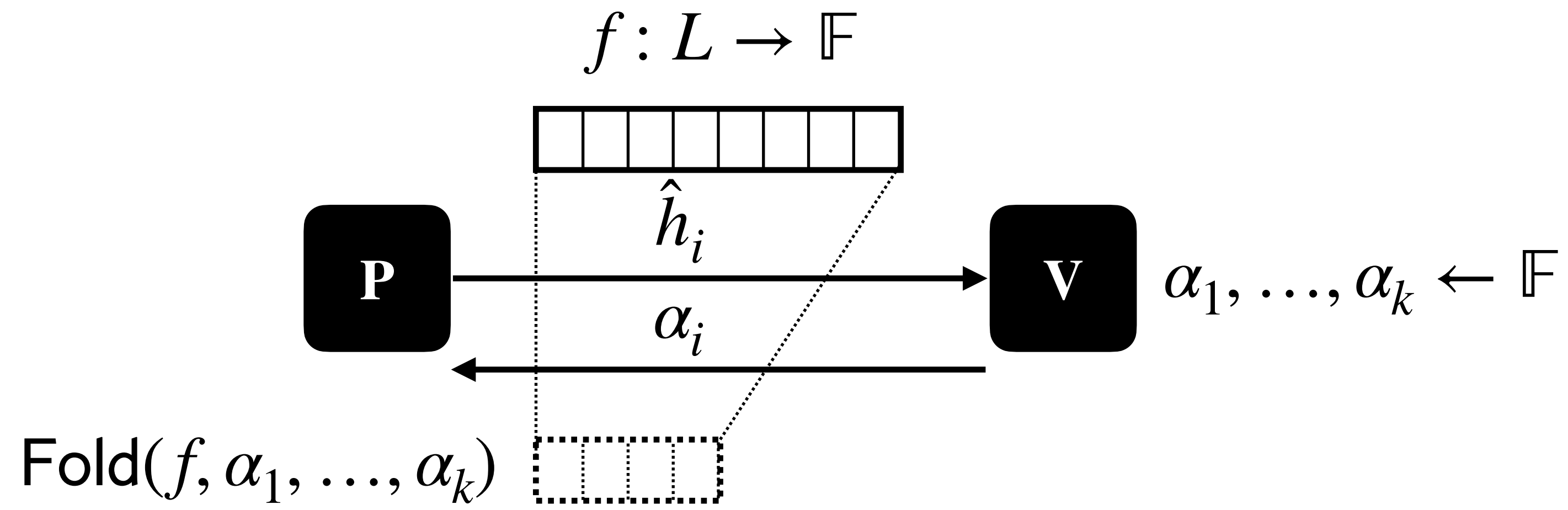
# WHIR



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# Application: $\Sigma$ -IOP

High soundness compilation using constrained codes

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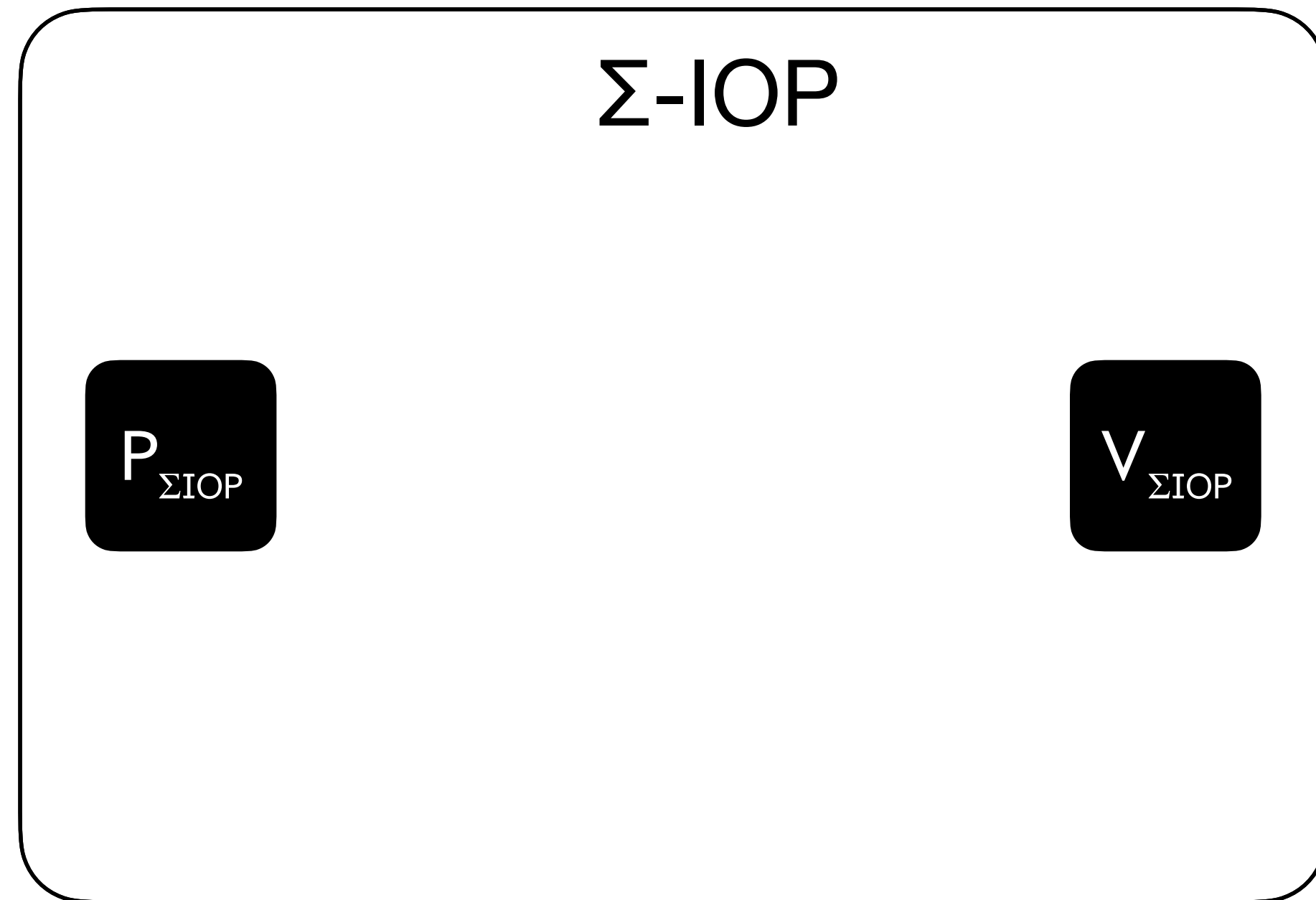
High soundness compilation using constrained codes



$\Sigma$ -IOP

# Application: $\Sigma$ -IOP

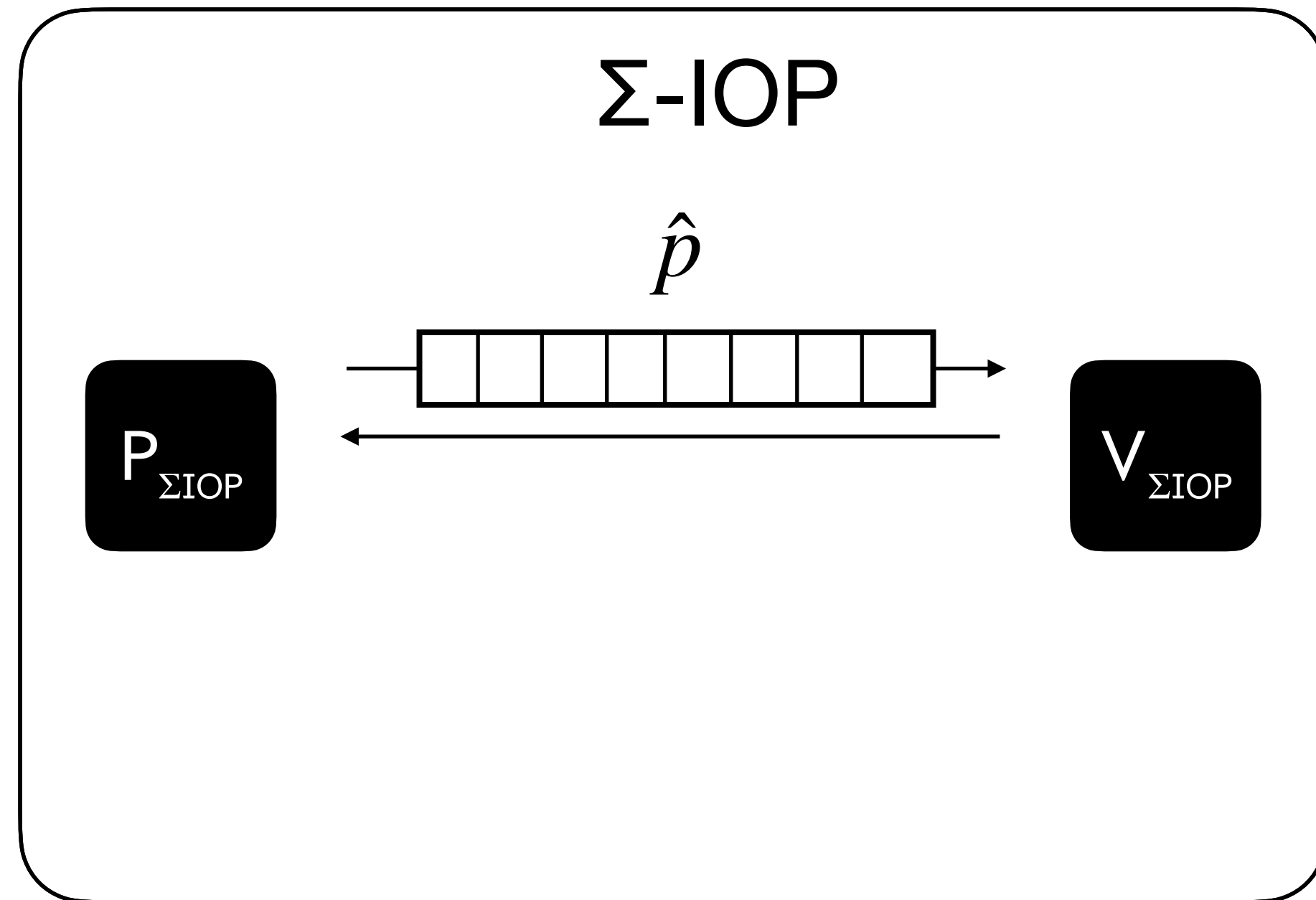
High soundness compilation using constrained codes





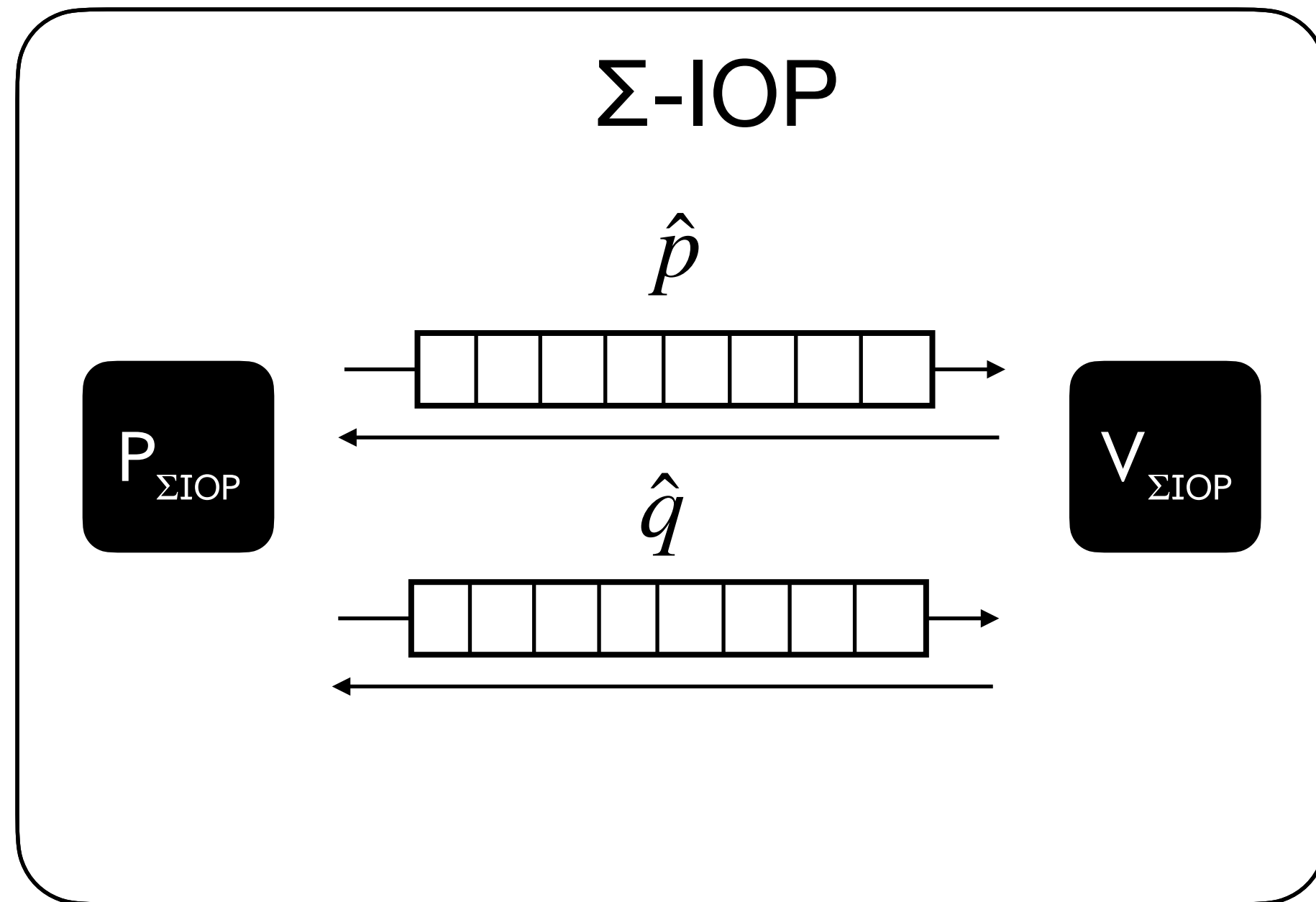
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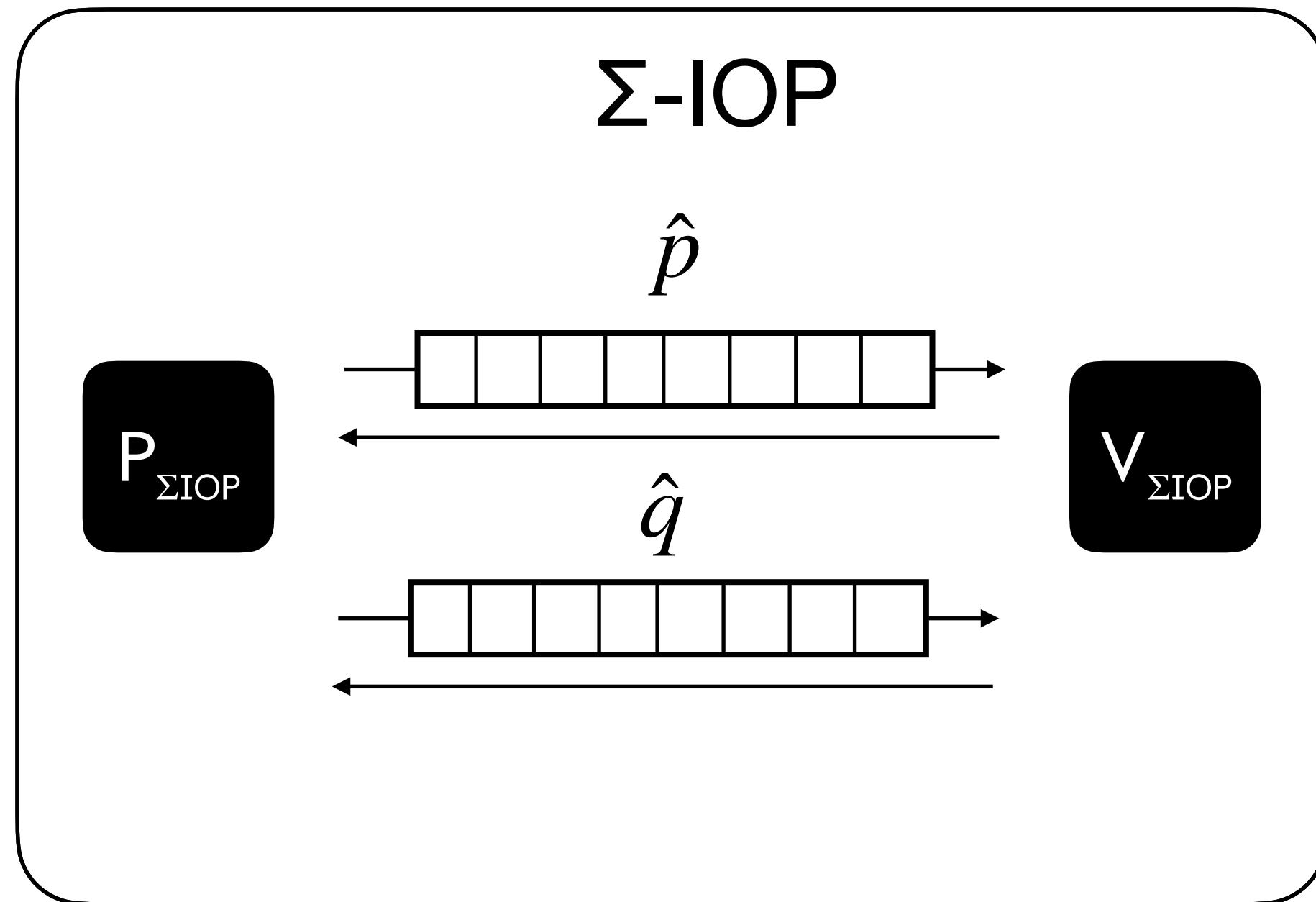
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Generalizes univariate and multilinear PIOPs at no extra cost!

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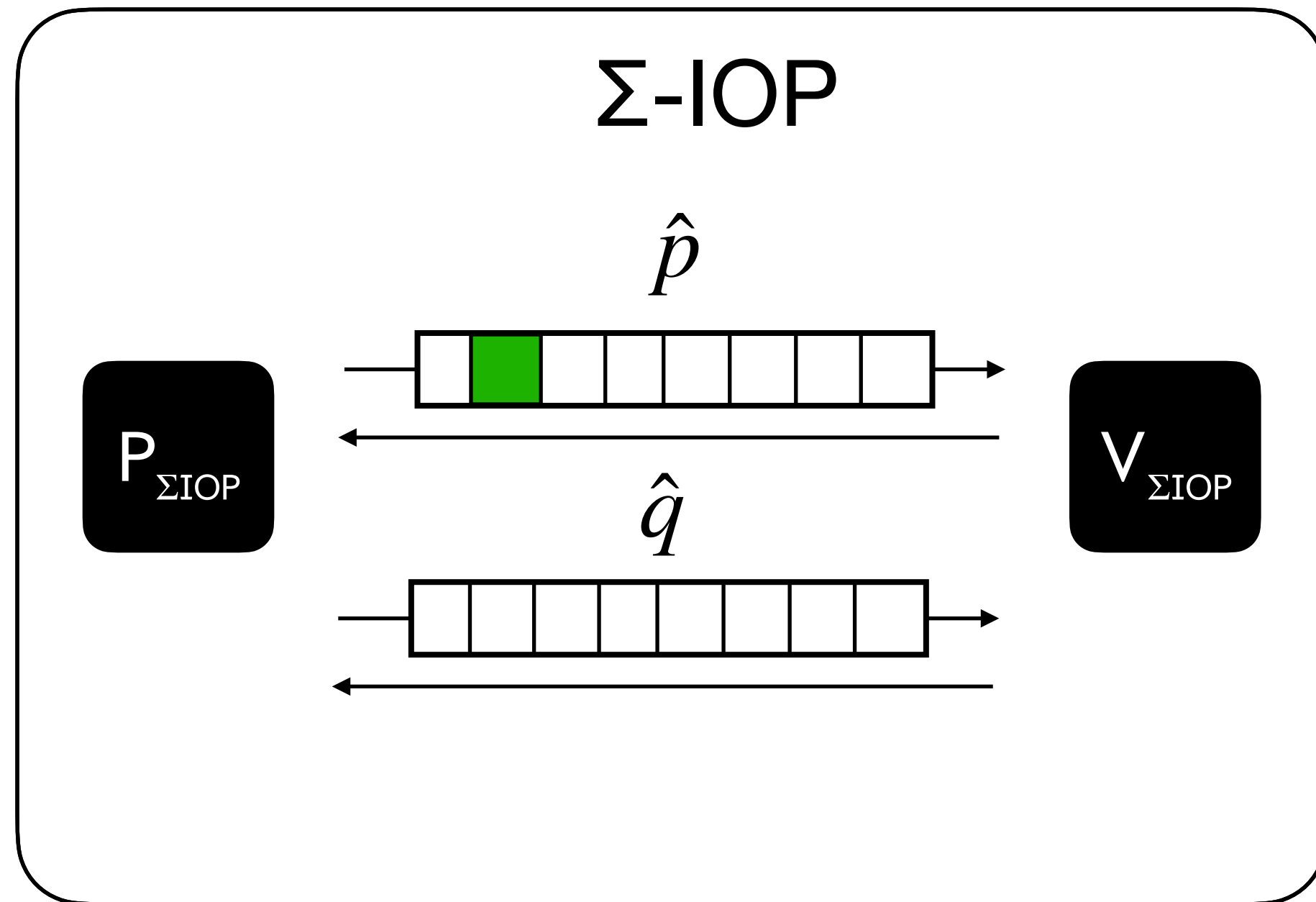
Verifier can ask **sumcheck queries**

i.e. send  $\hat{w}$  and receive  $\sum_{\mathbf{b}} \hat{w}(\hat{f}(\mathbf{b}), \mathbf{b})$

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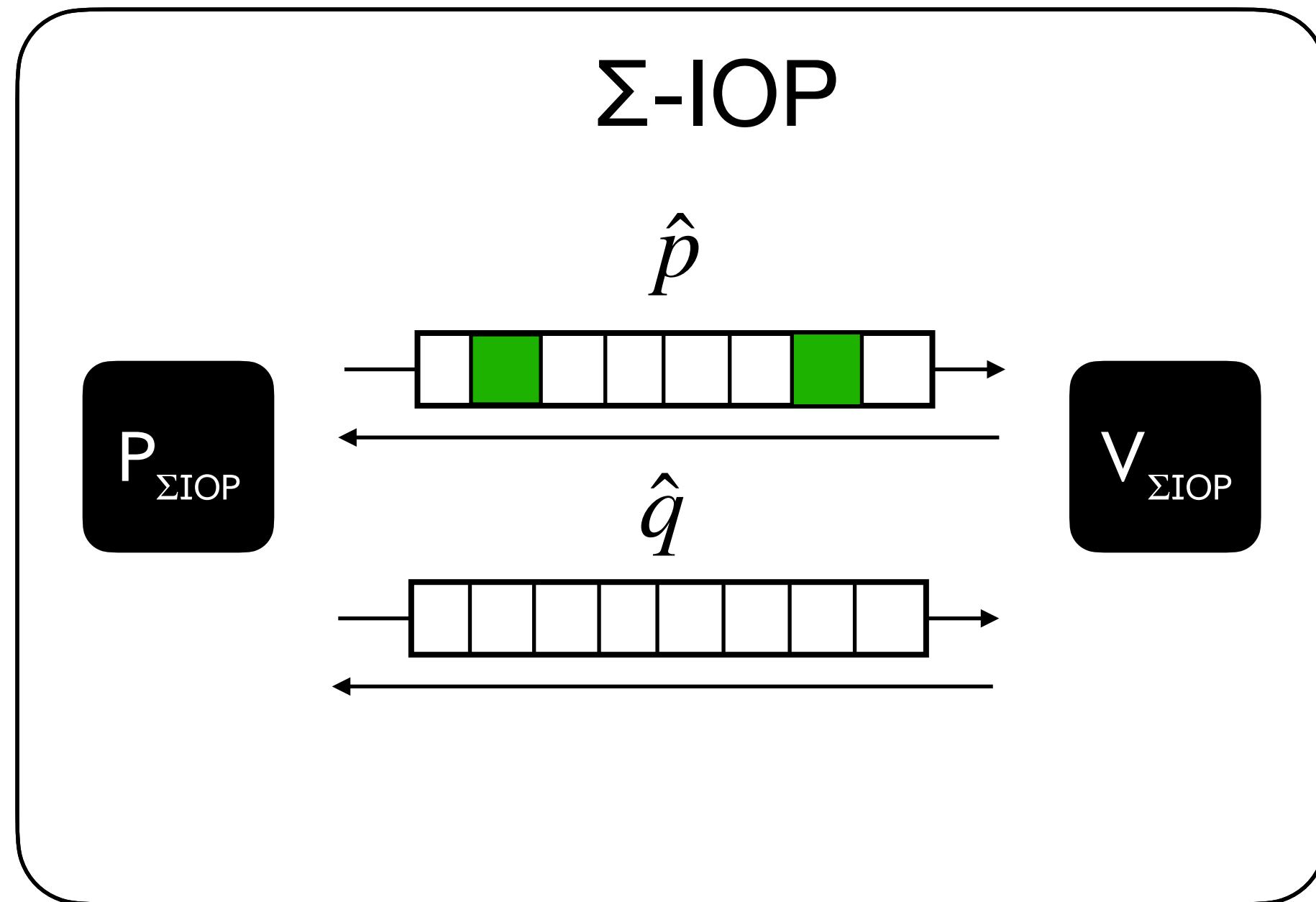
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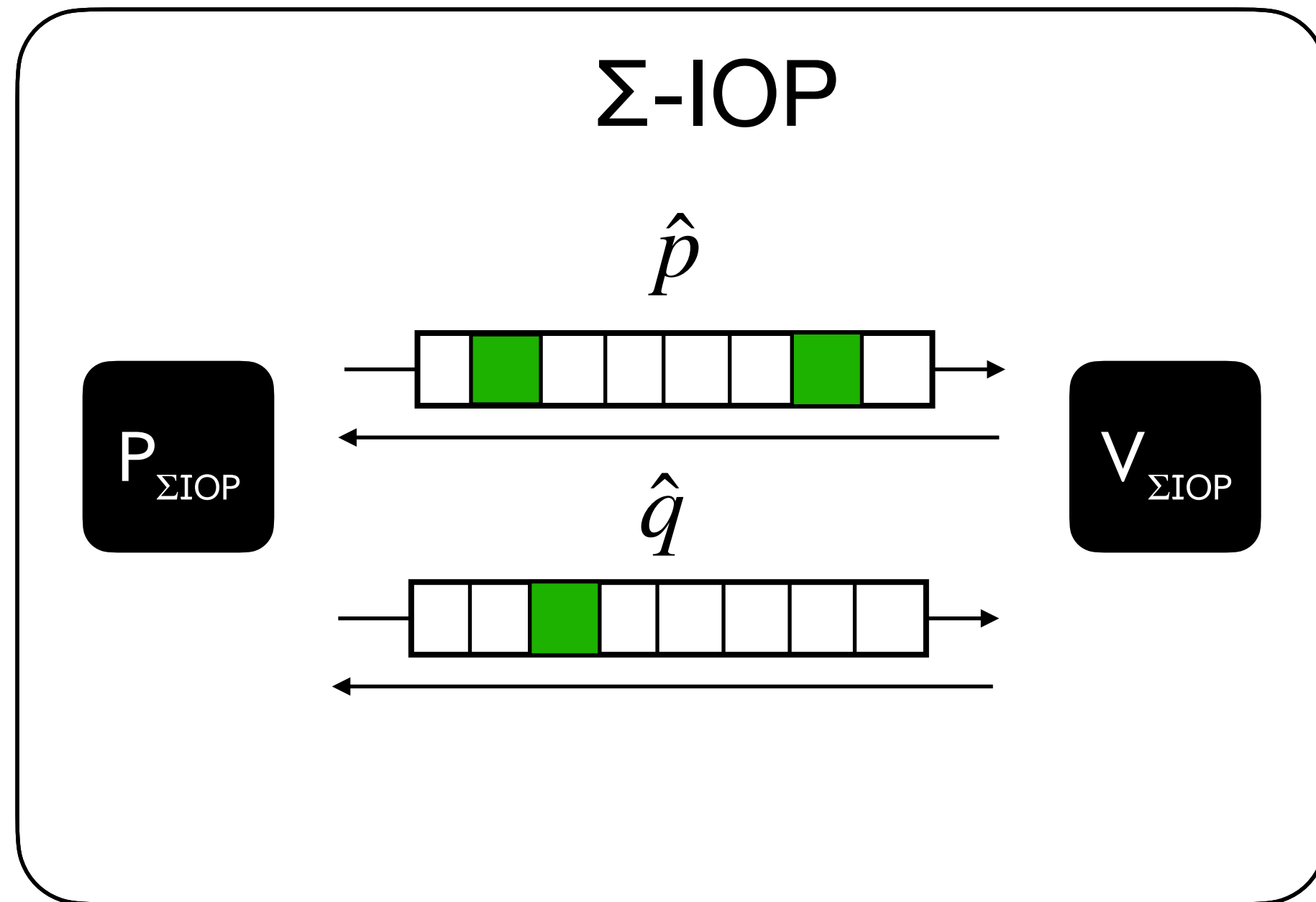
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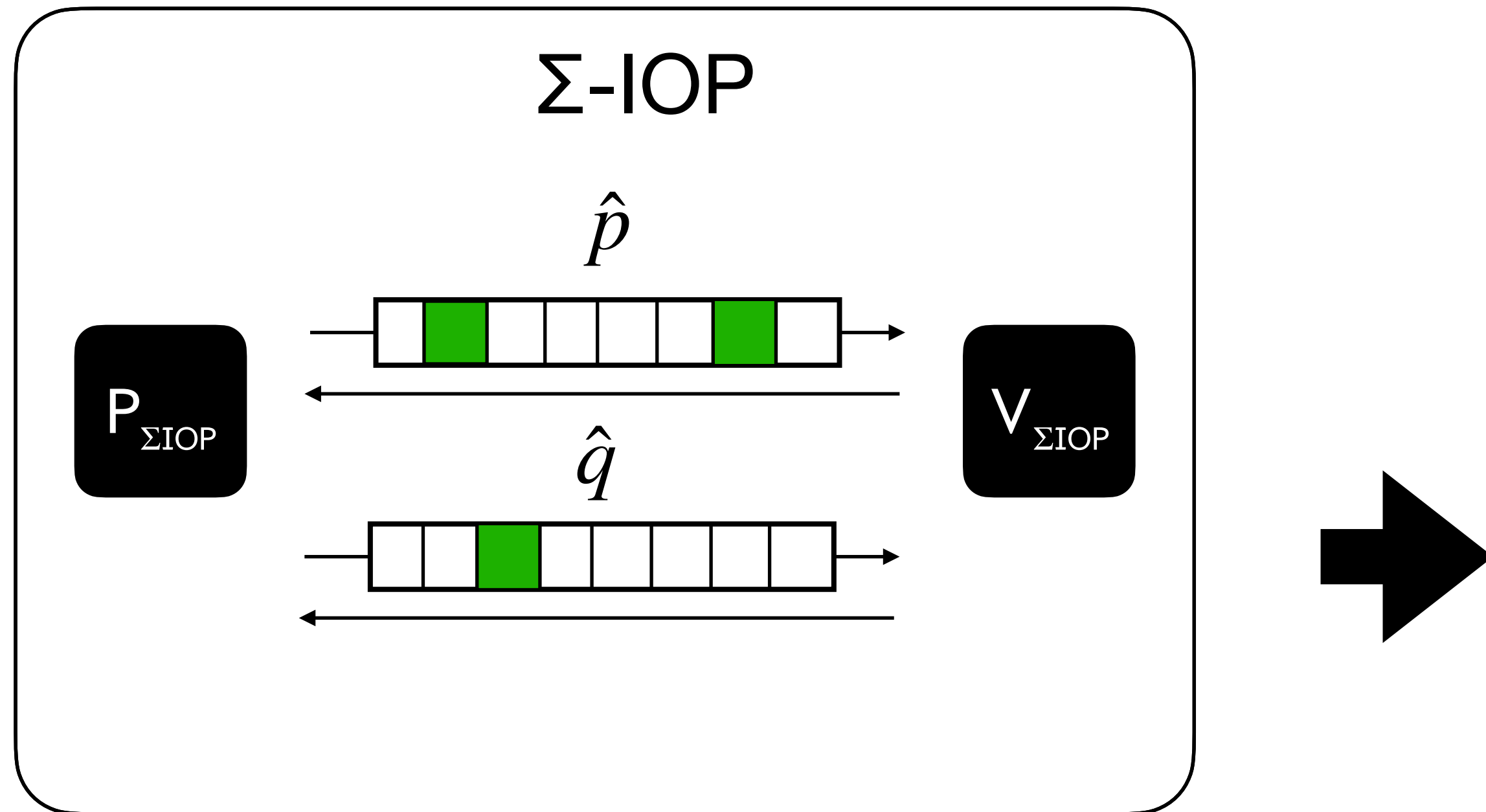
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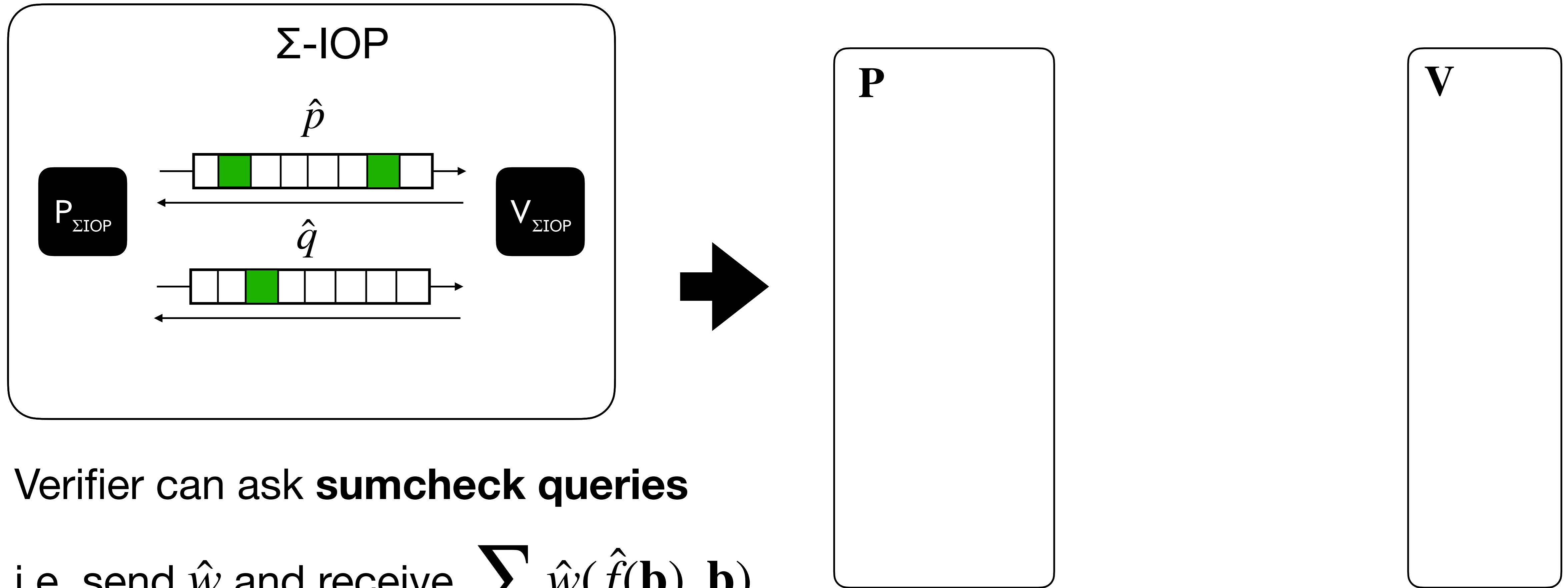
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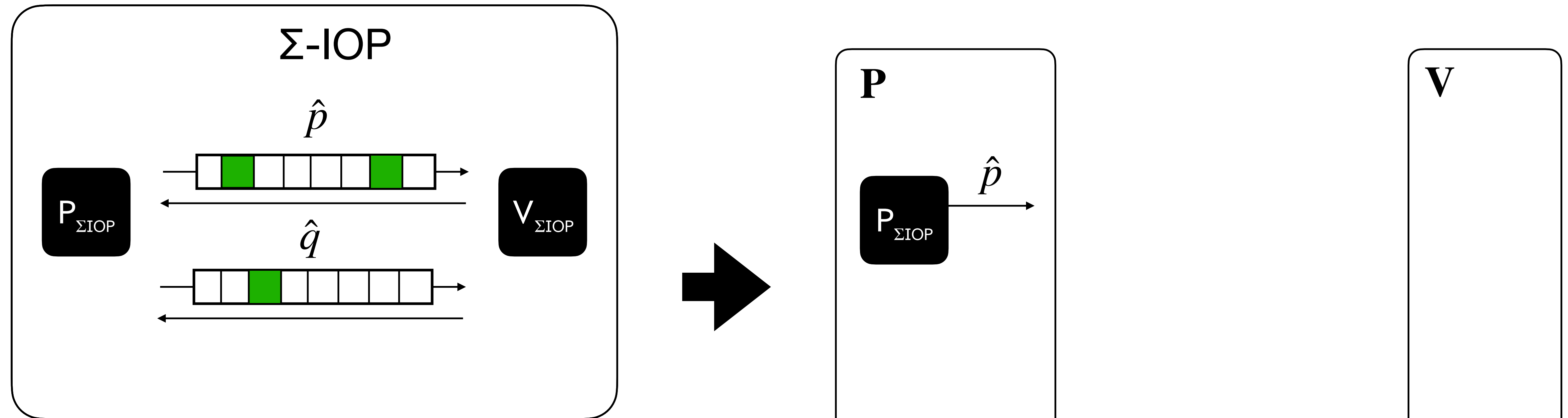
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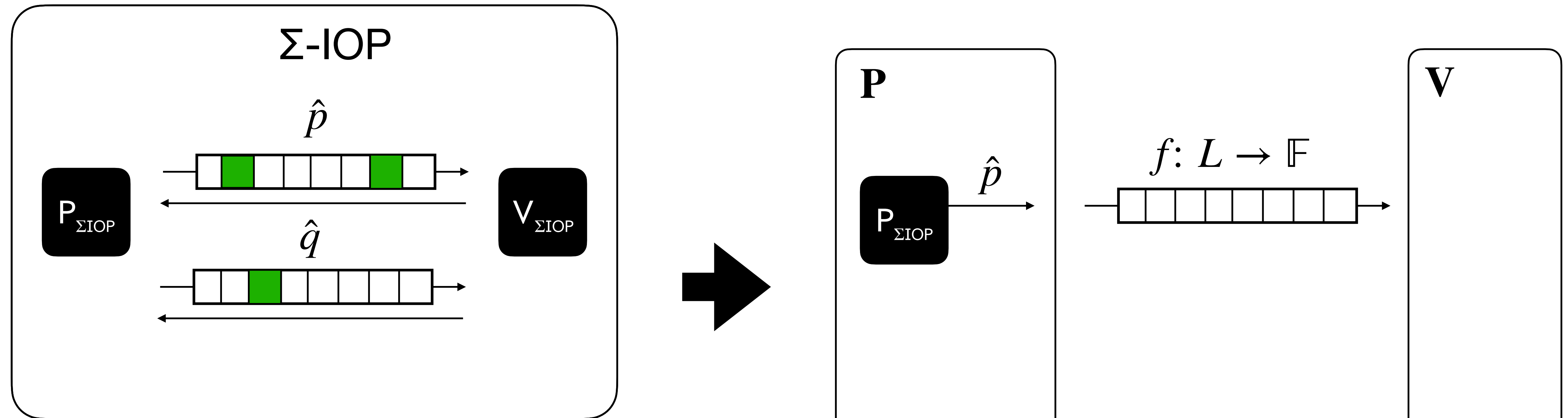
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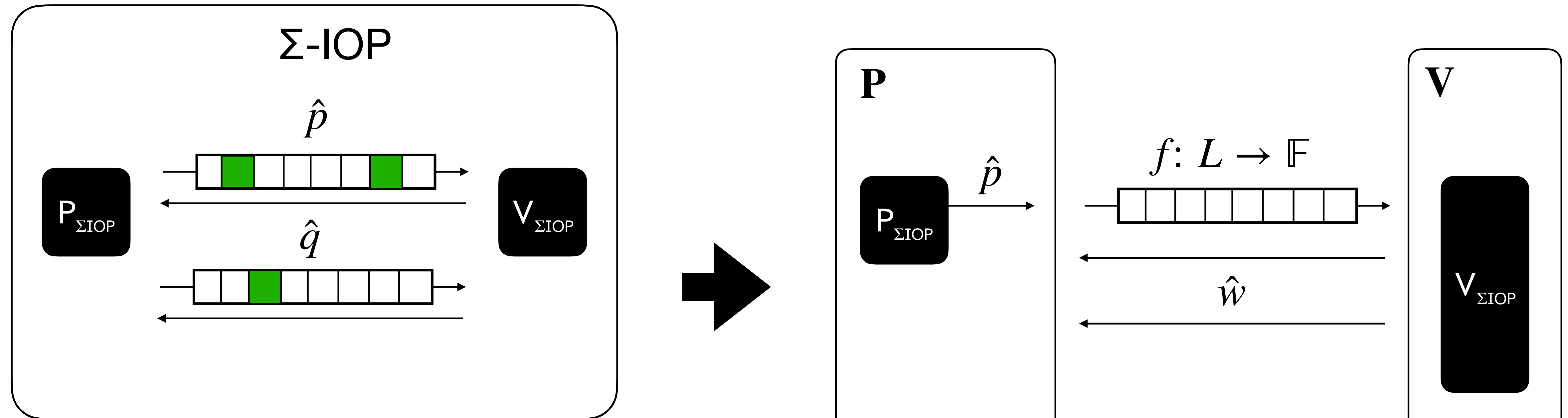
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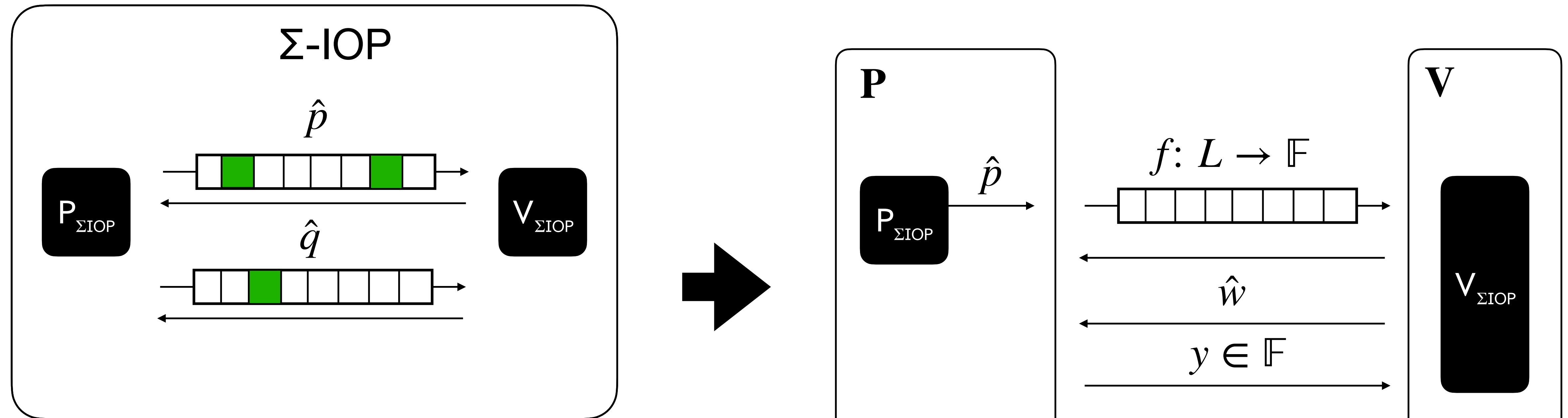
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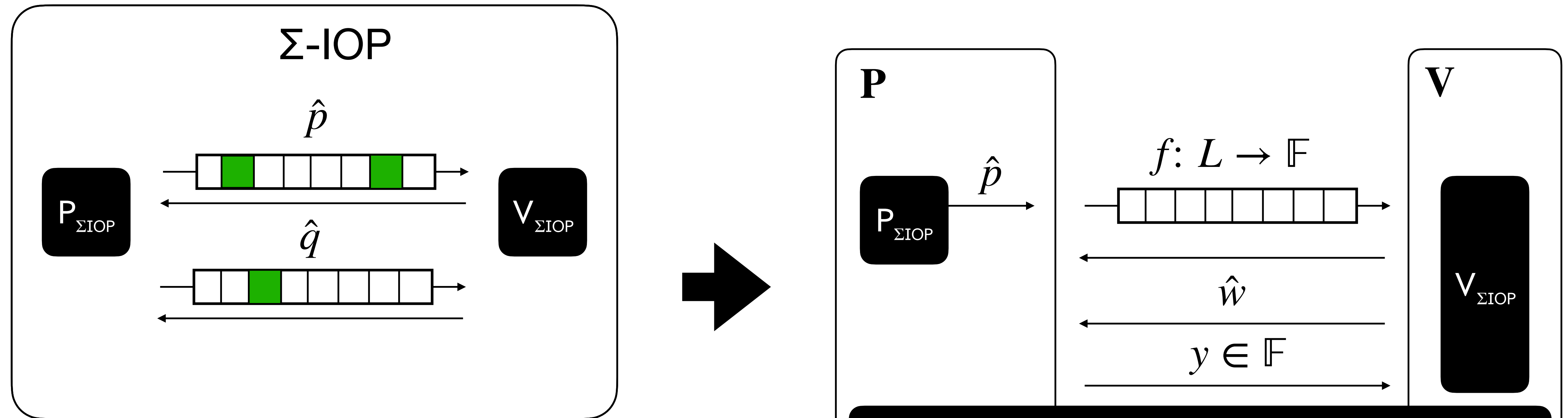
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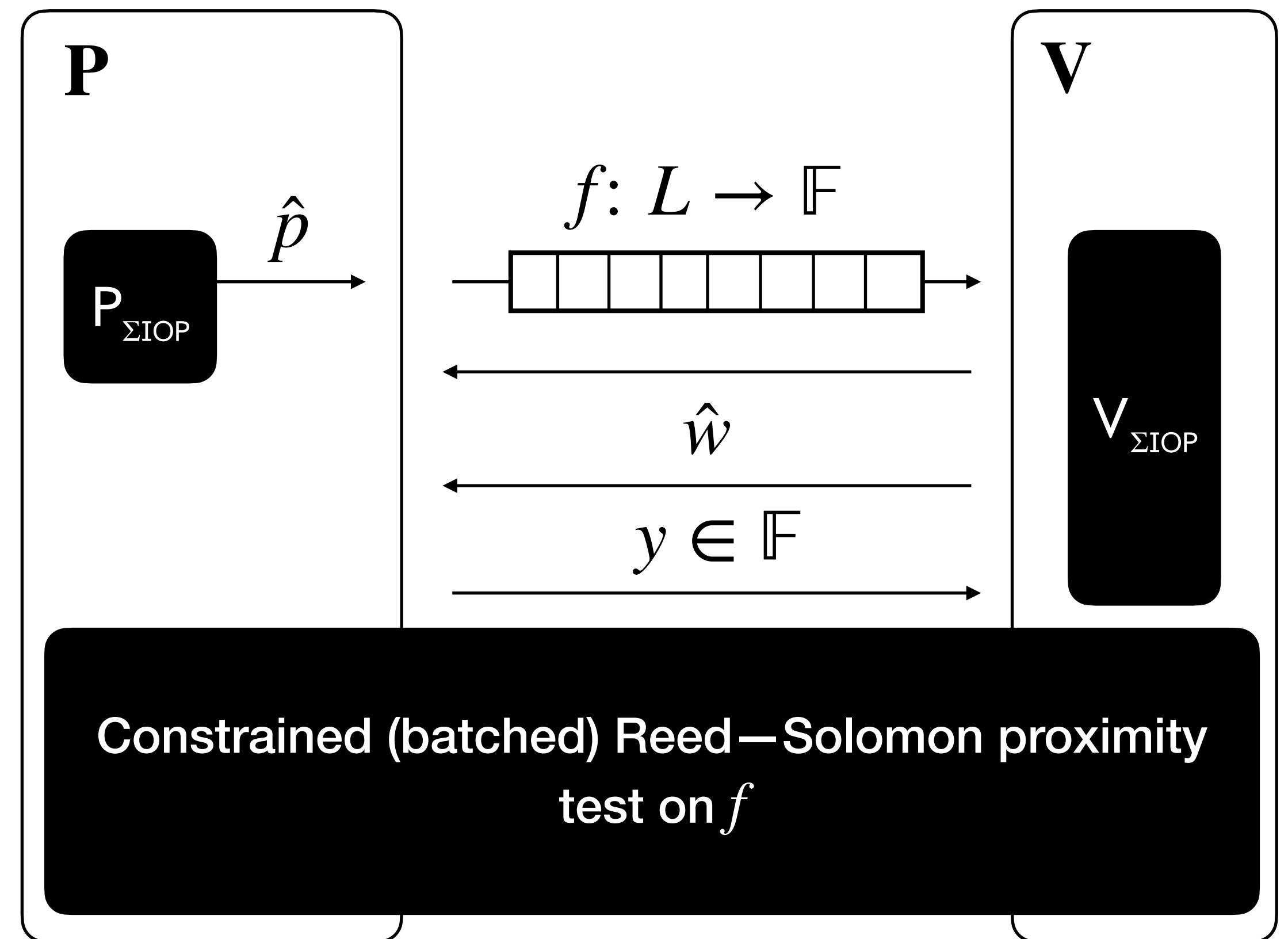
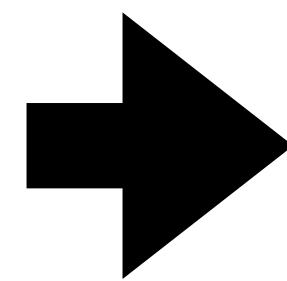
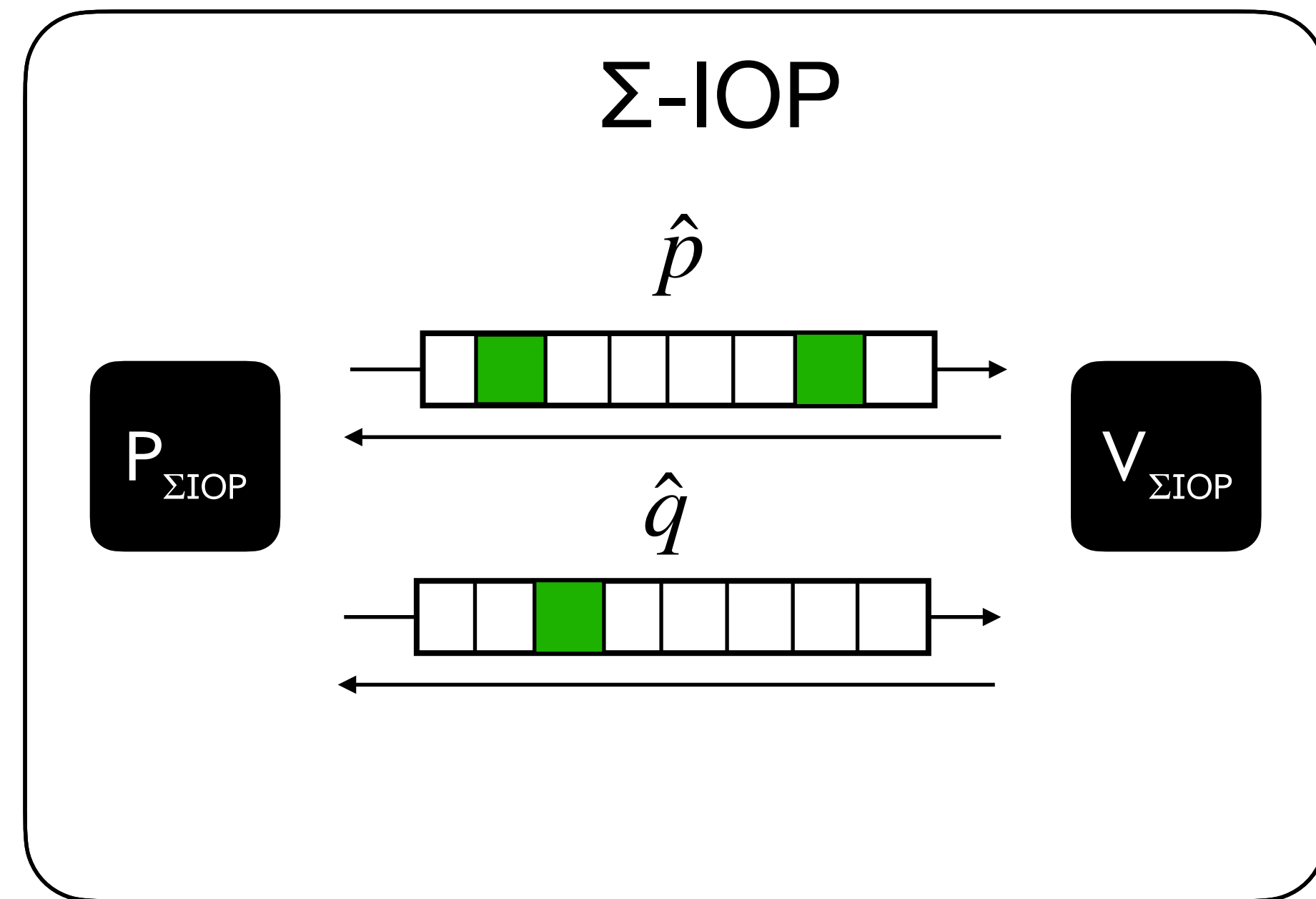
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Q: Can we use this to do more efficient arithmetizations?

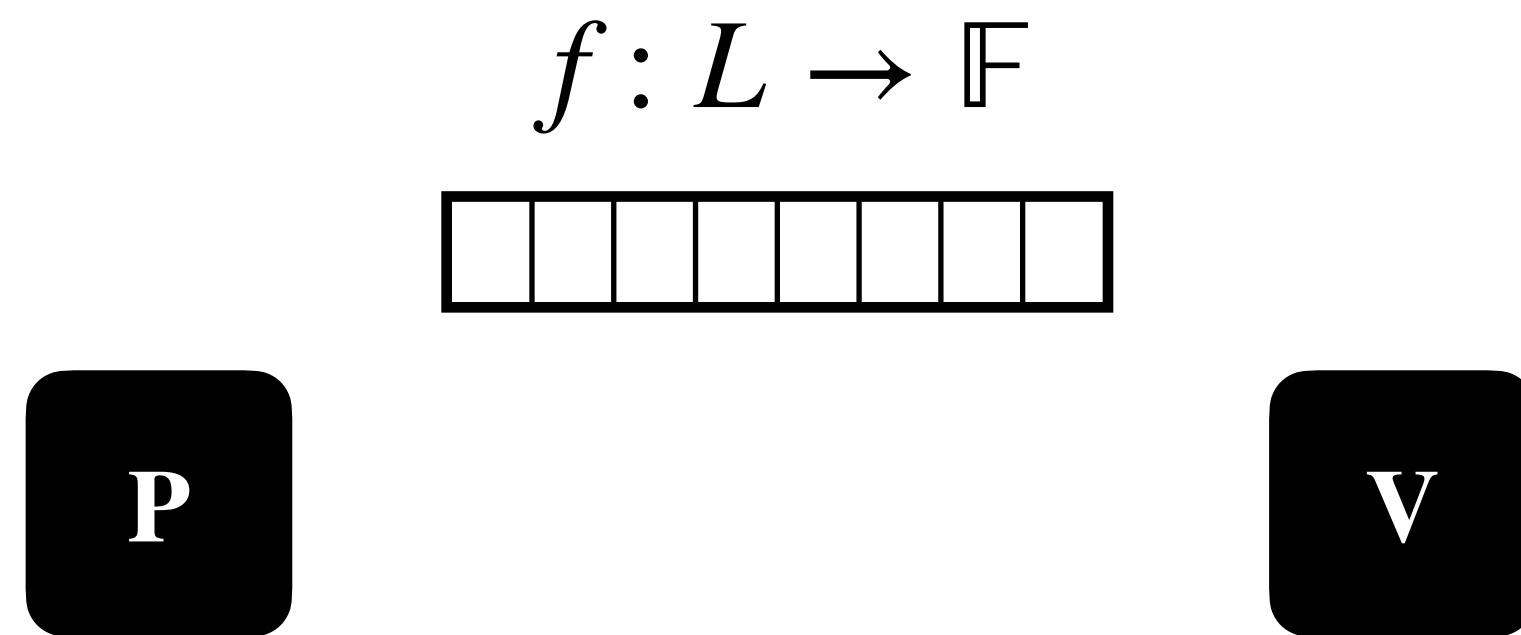


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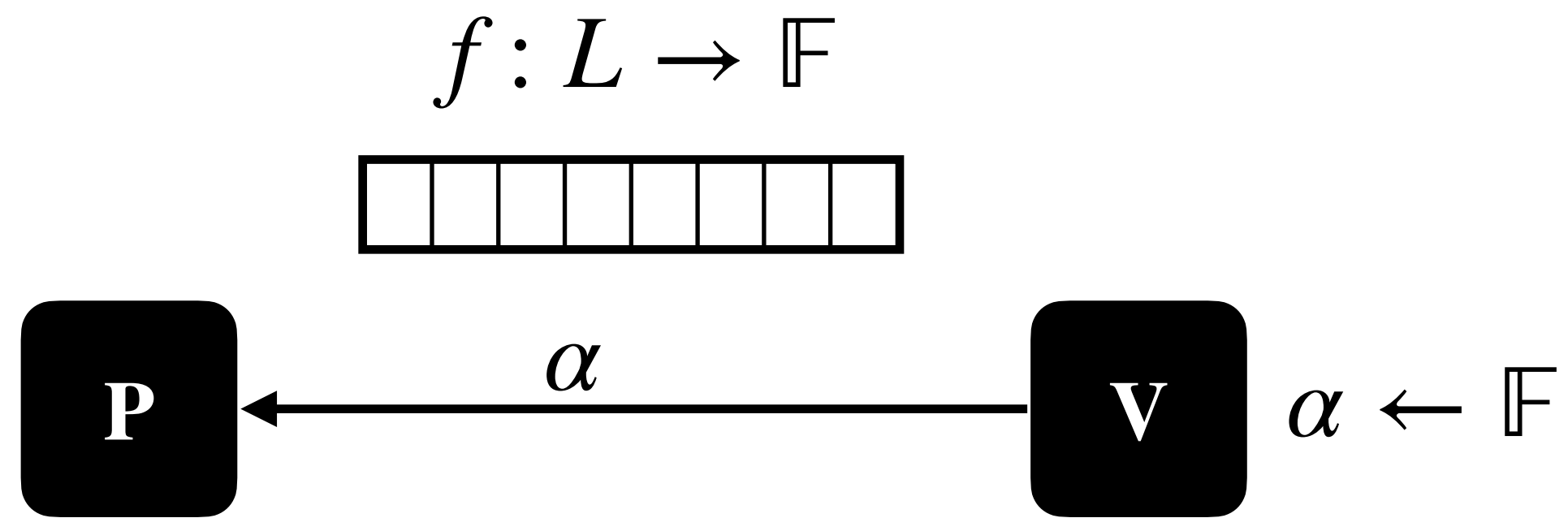
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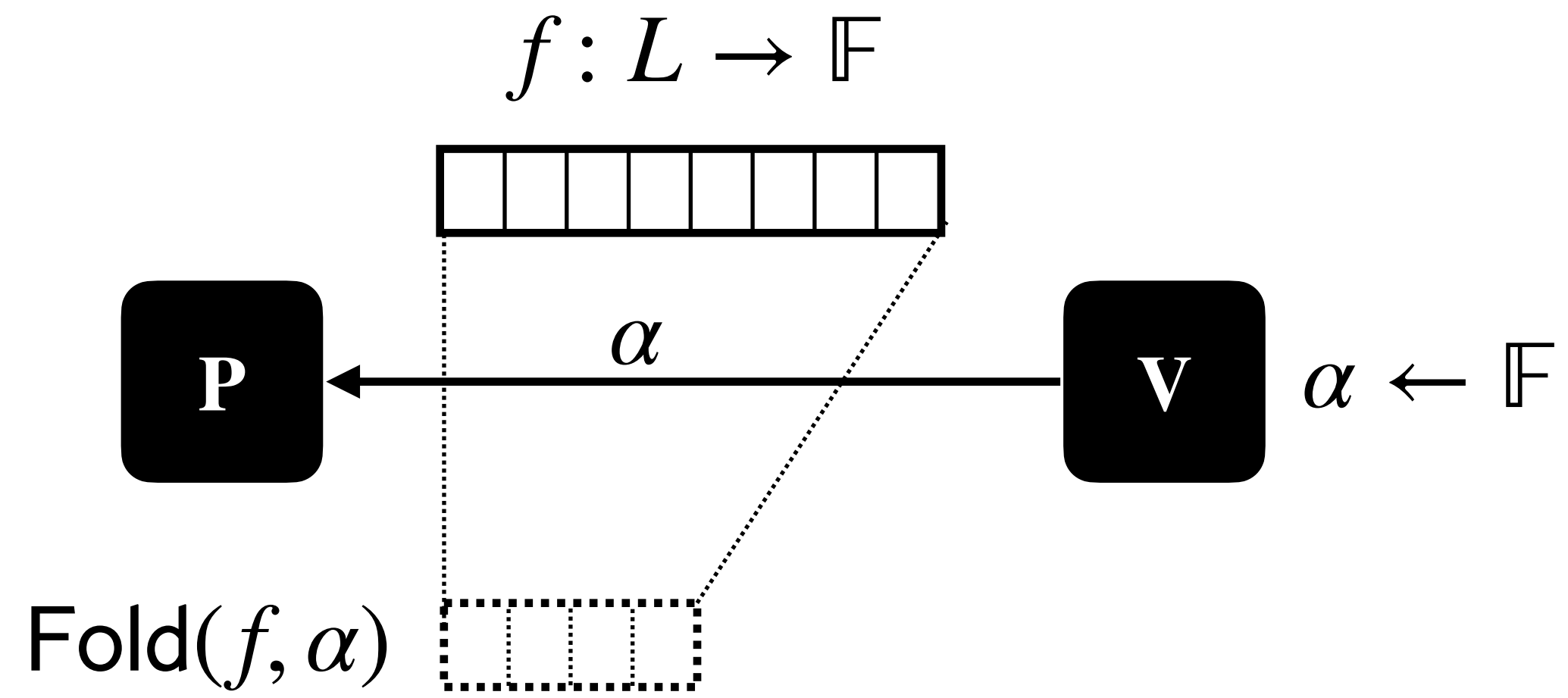




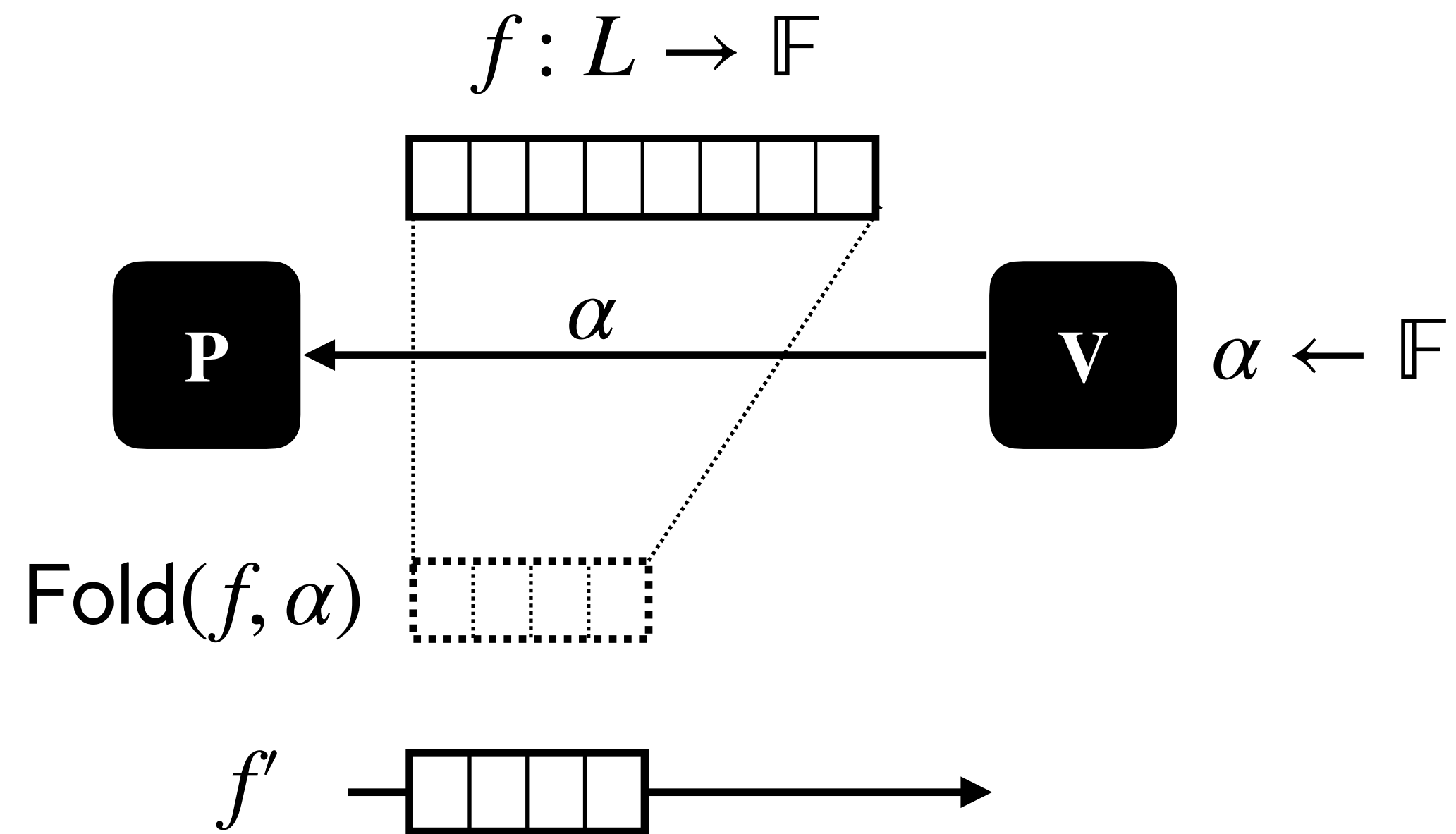
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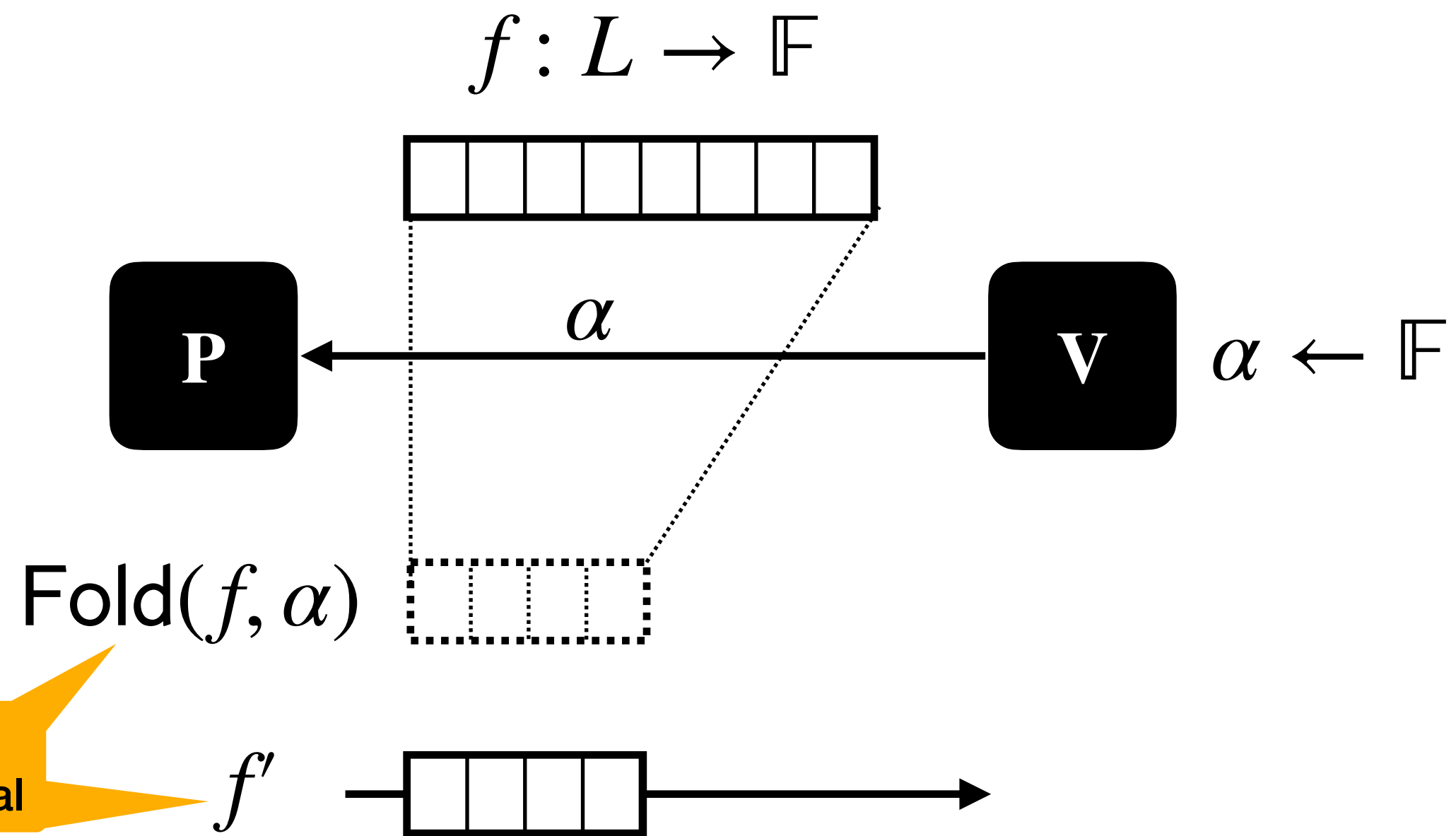
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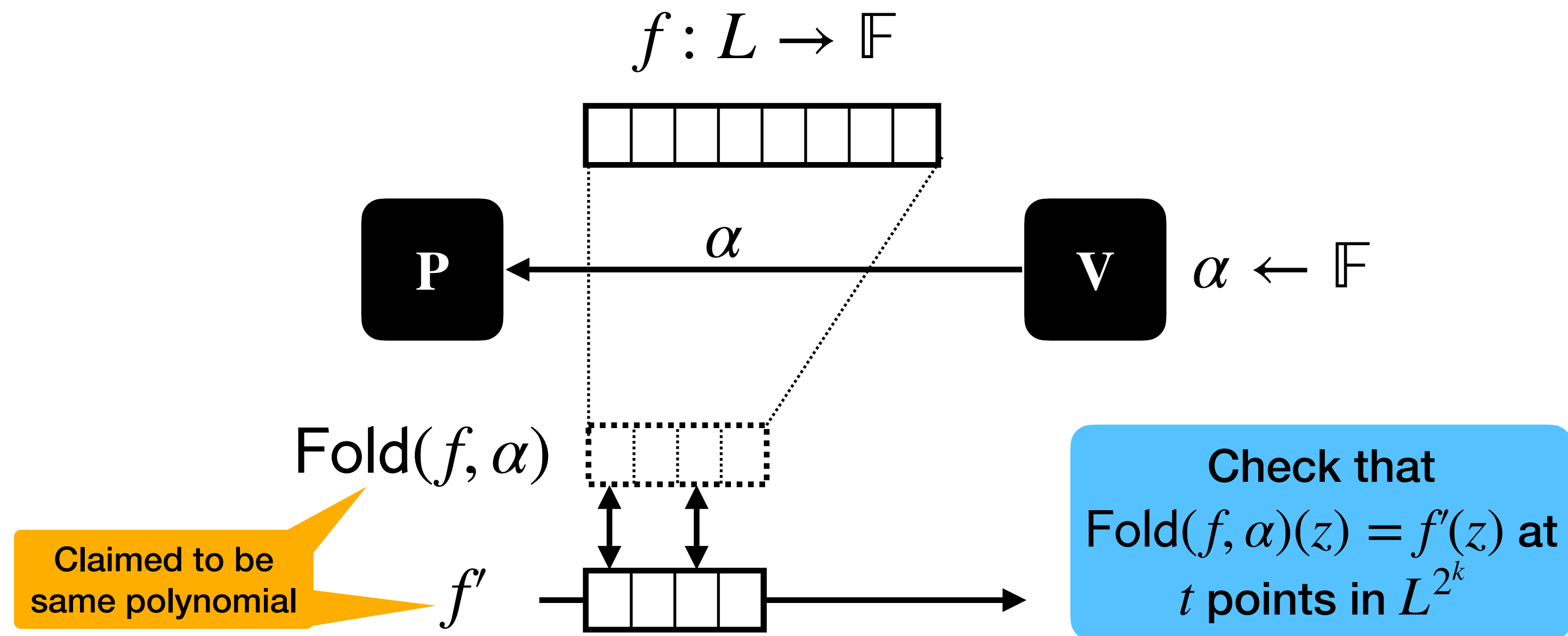
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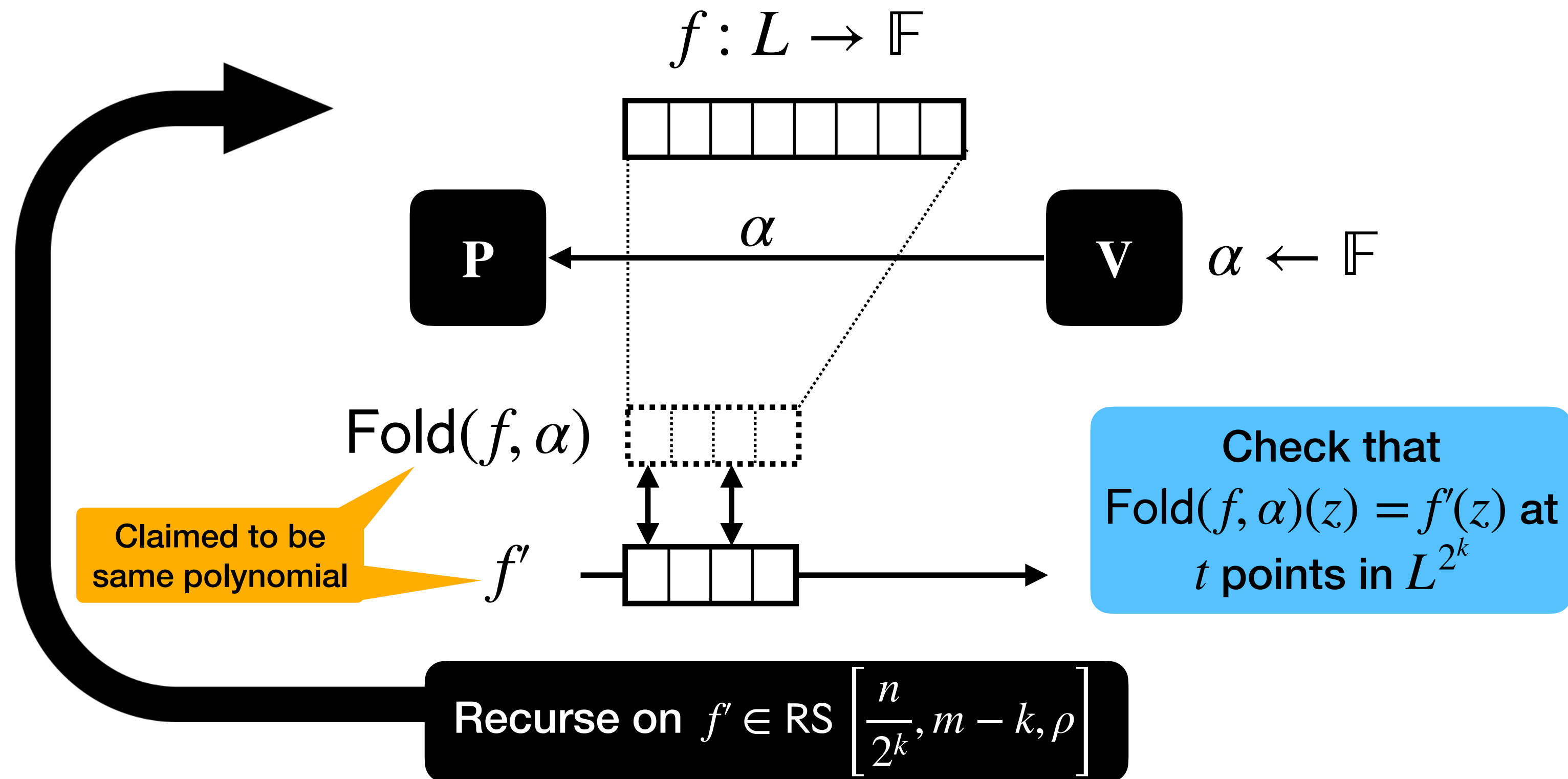
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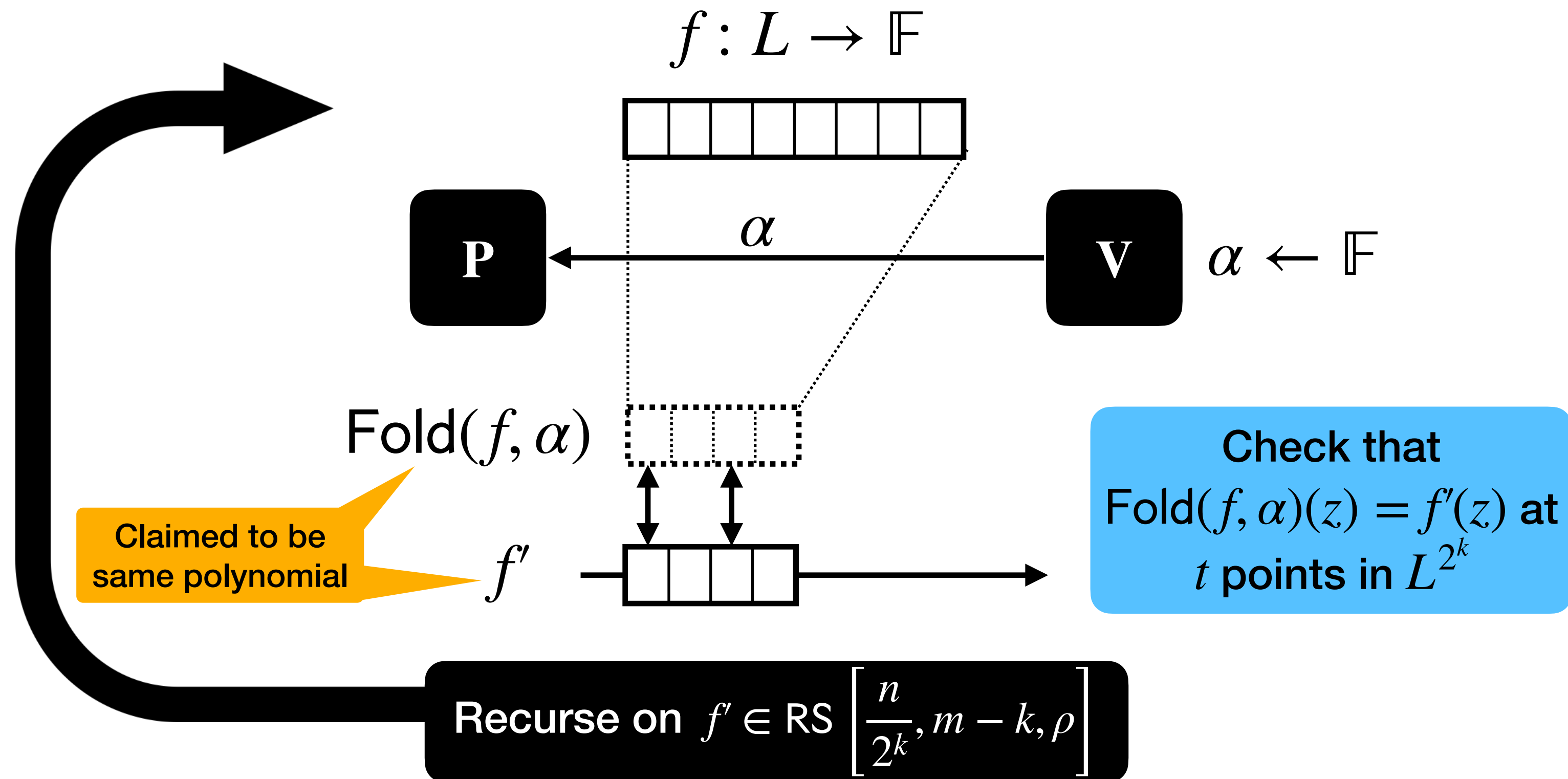
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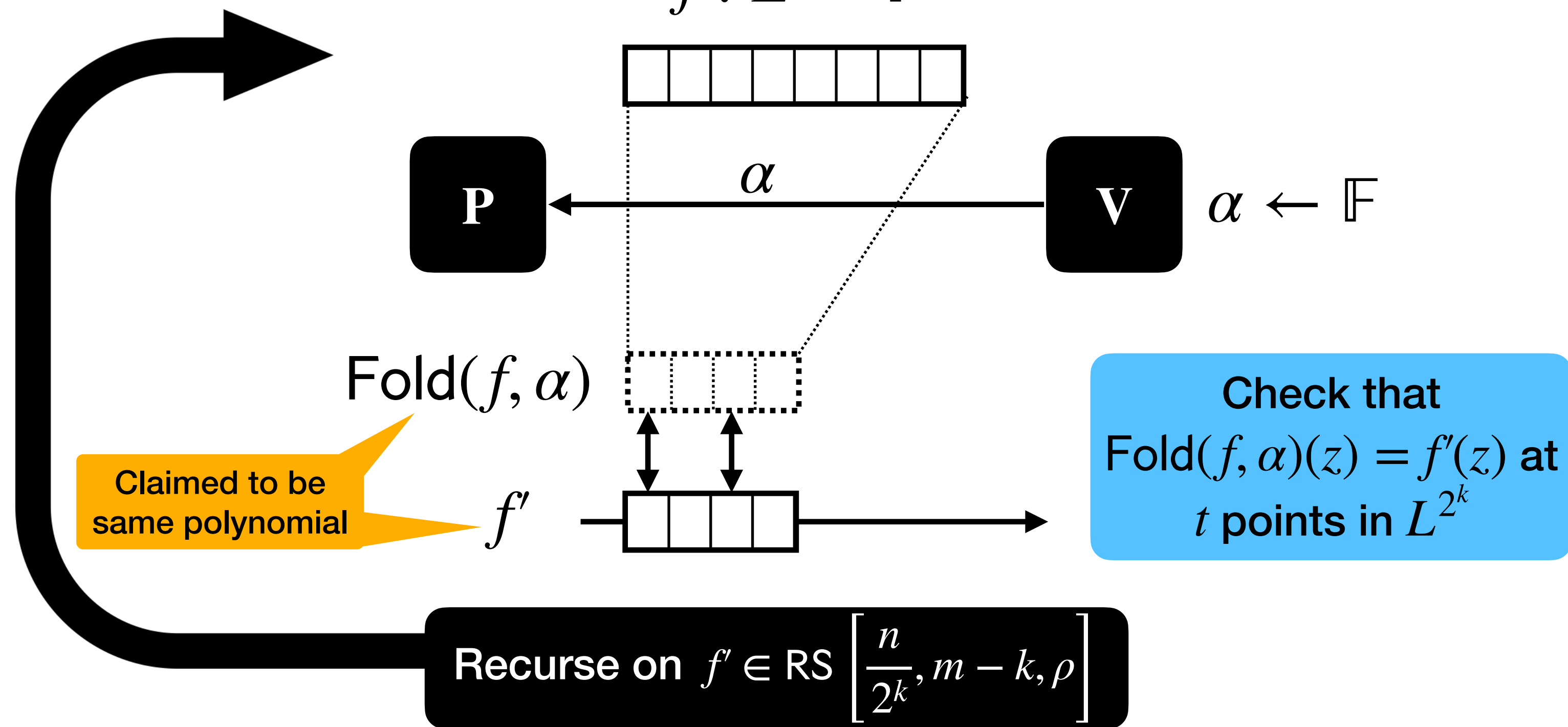


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**Disclaimer:** in full FRI consistency checks are correlated between rounds.

# Review: FRI iteration



## Soundness:

Suppose that  $f' \in \text{RS}[n/2^k, m - k, \rho]$ .

If  $f$  is  $\delta$ -far from  $\text{RS}[n, m, \rho]$ ,

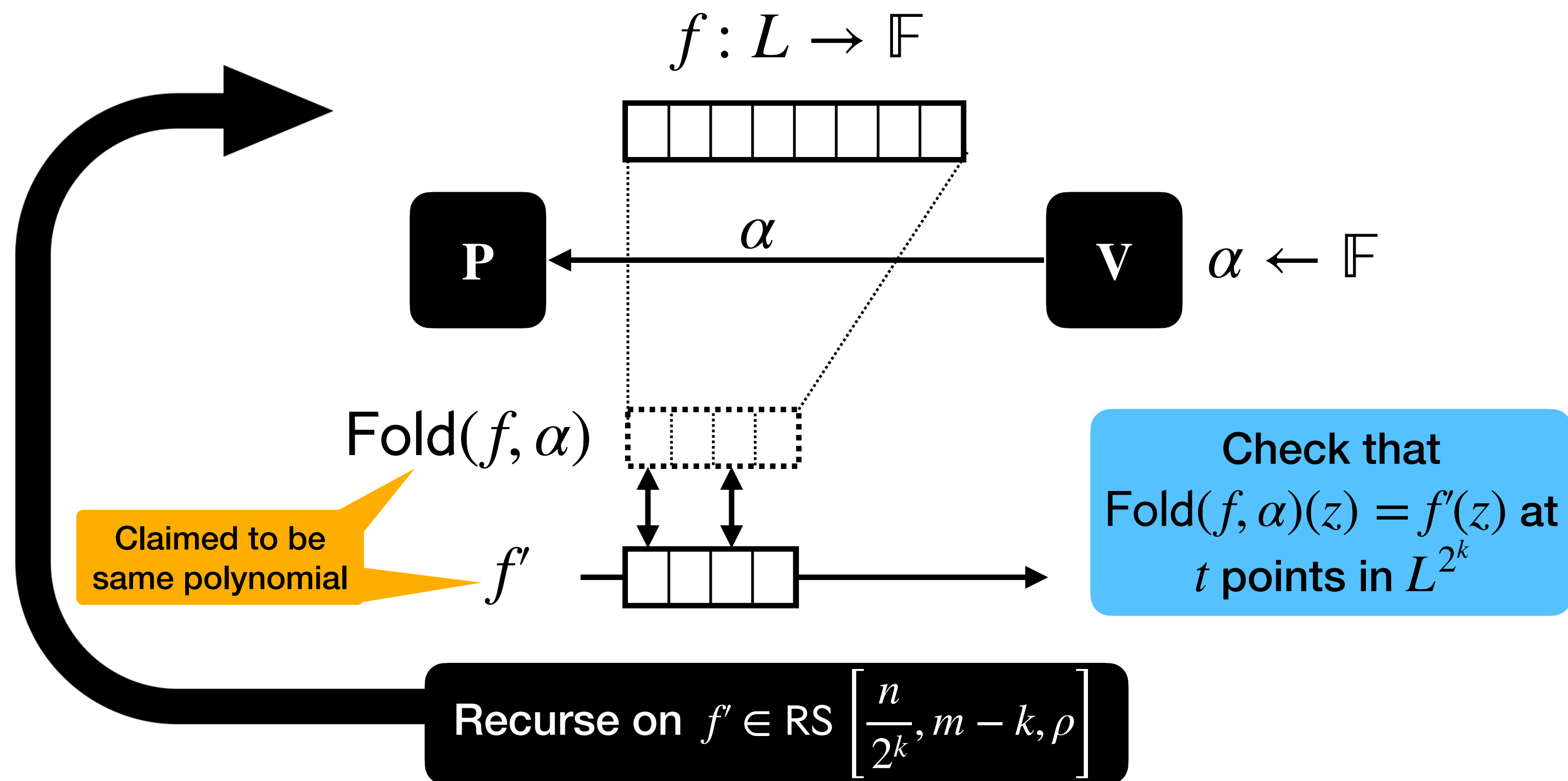
$\text{Fold}(f, \alpha)$  must be  $\delta$ -far from  $\text{RS}[n/2^k, m - k, \rho]$

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Suppose that  $f' \in \text{RS}[n/2^k, m - k, \rho]$ .

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$\text{Fold}(f, \alpha)$  must be  $\delta$ -far from  $\text{RS}[n/2^k, m - k, \rho]$

Then,  $f'$  and  $\text{Fold}(f, \alpha)$  differ on a  $\delta$ -fraction.

**Soundness error** is  $(1 - \delta)^t$

To get soundness error  $\epsilon_{\text{RBR}} \leq 2^{-\lambda}$ :  
 set  $\delta := 1 - \sqrt{\rho}$  and  $t := \frac{\lambda}{-\log \sqrt{\rho}}$

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