

Isotonic subgroup selection

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12th April 2024

Collaborators



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University of Bristol



Timothy I. Cannings
University of Edinburgh



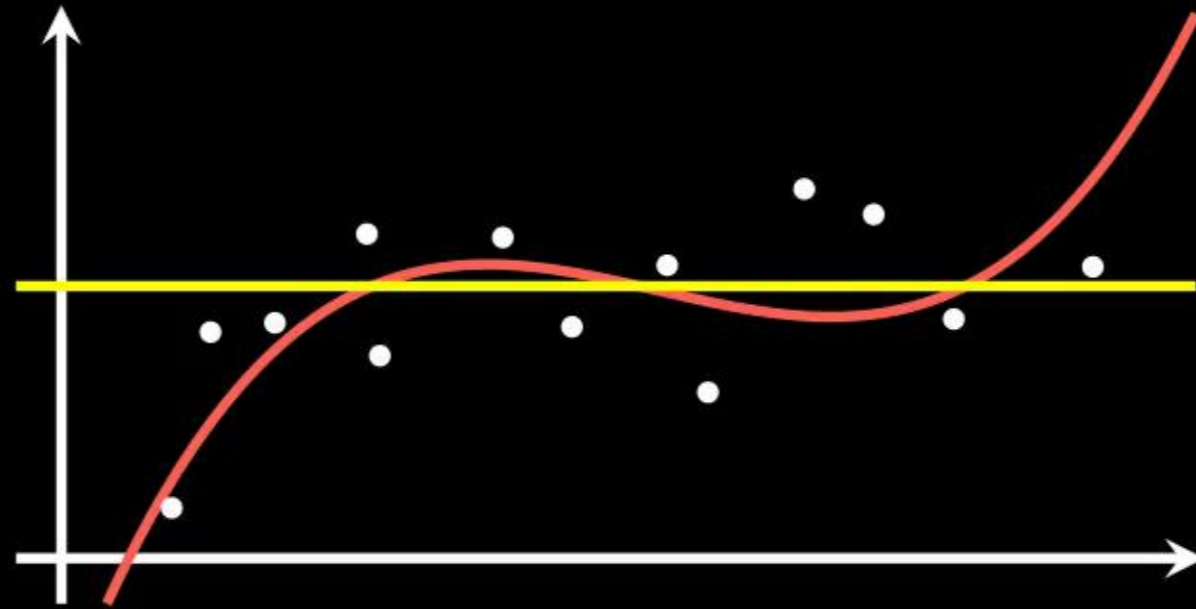
Richard J. Samworth
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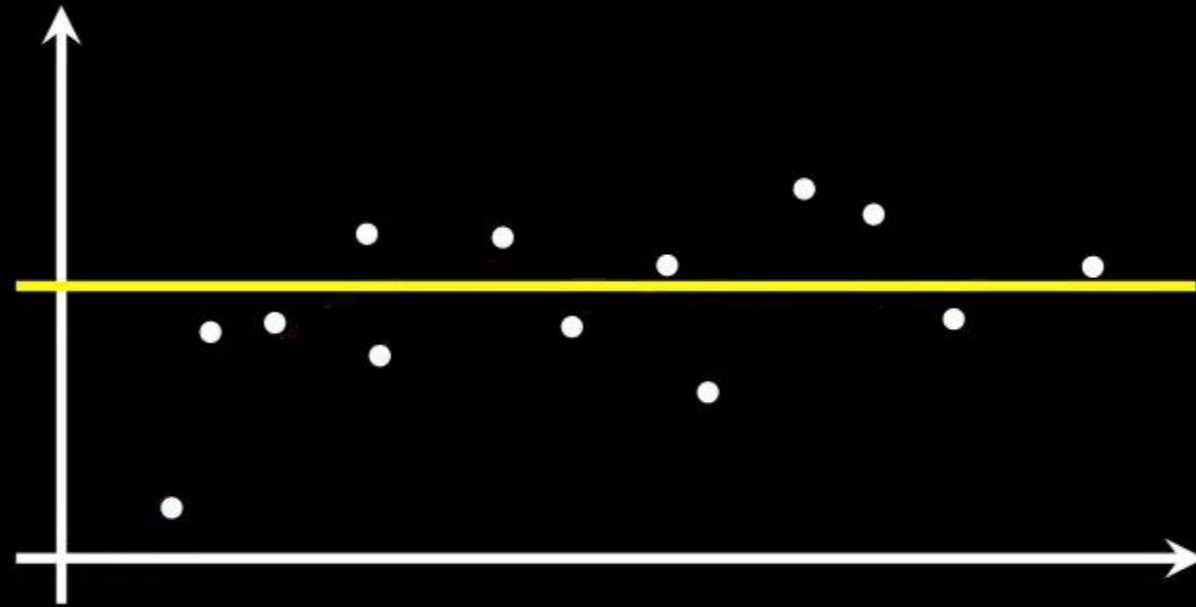
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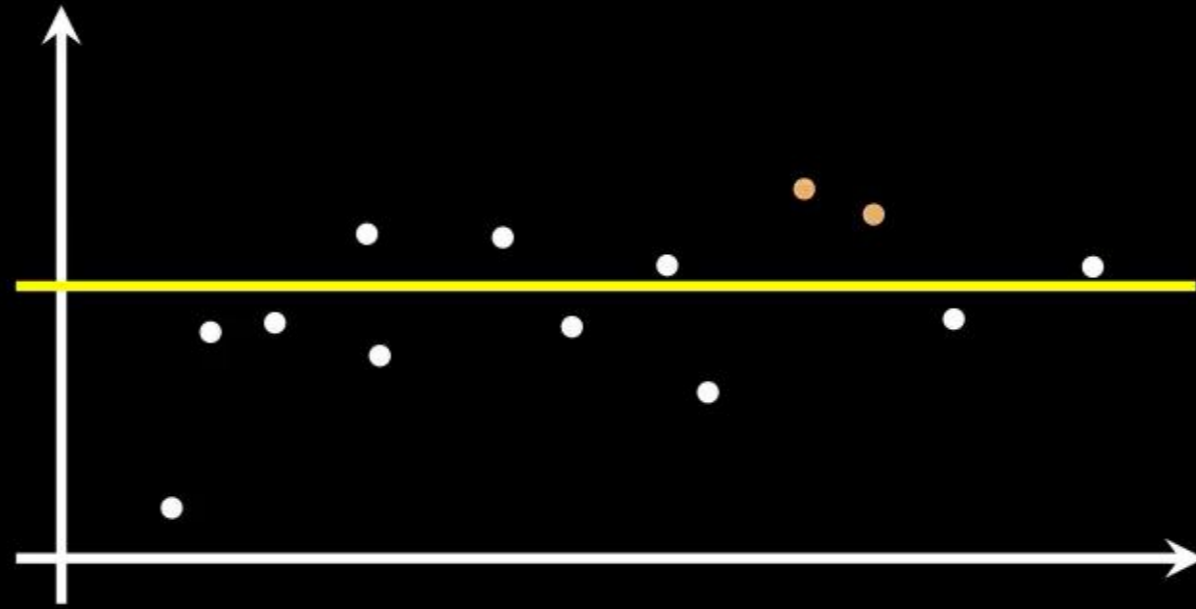
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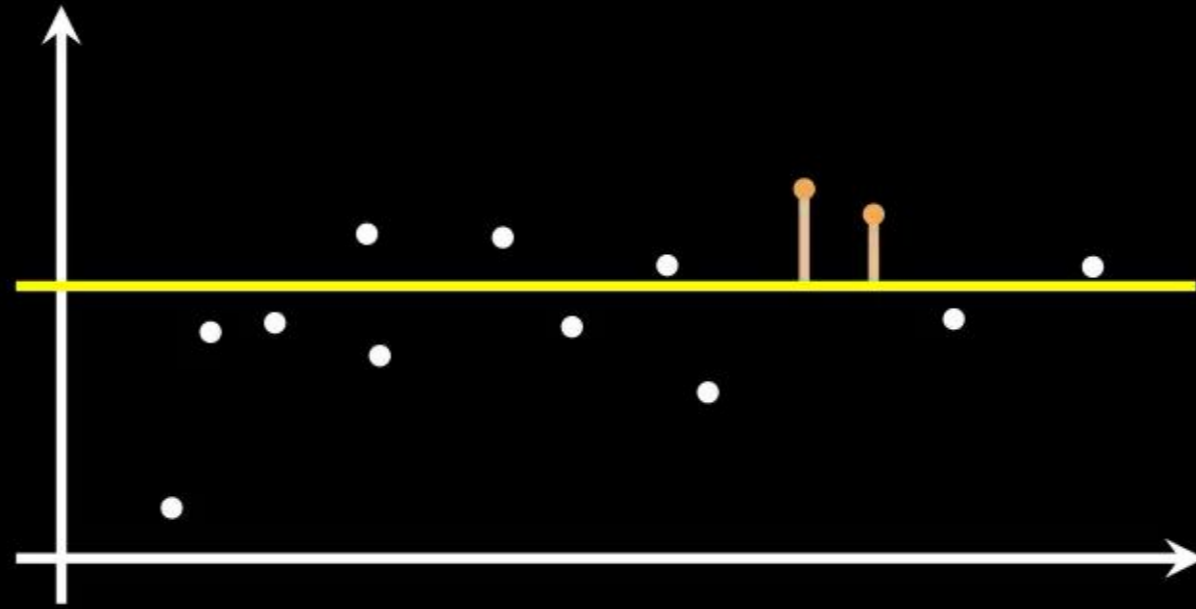
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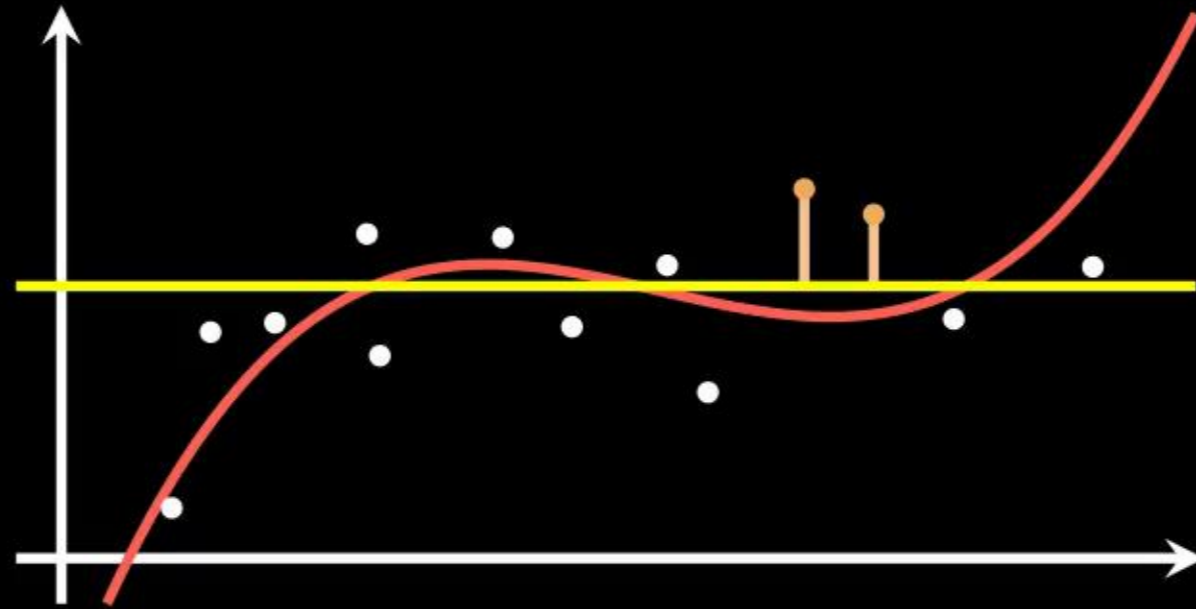
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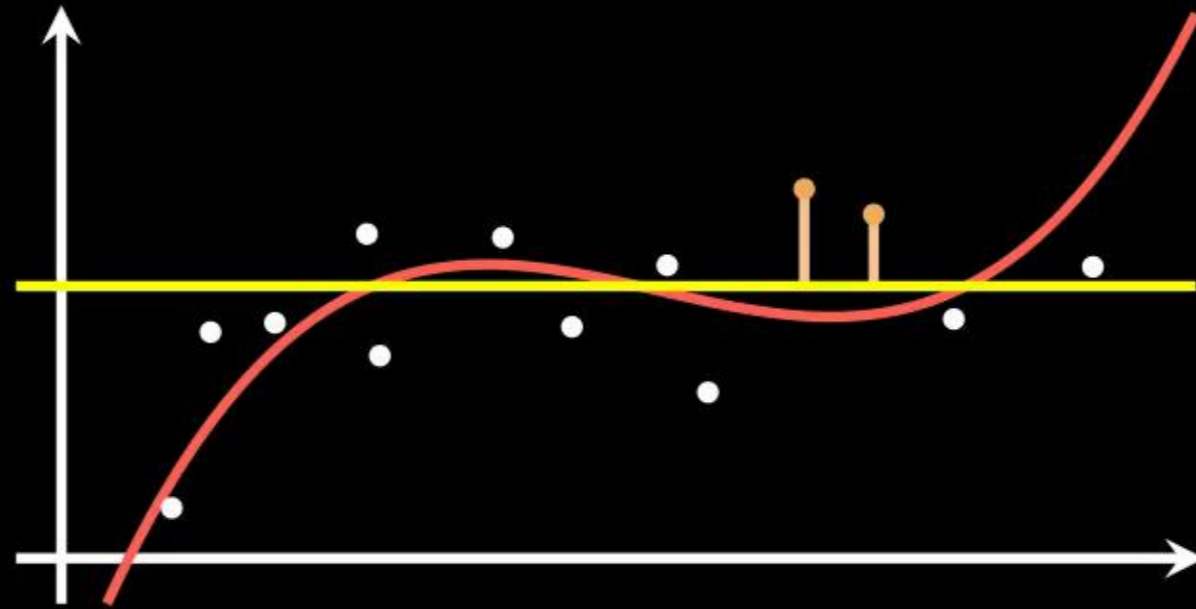
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Example. Efficacy of a new vaccine.

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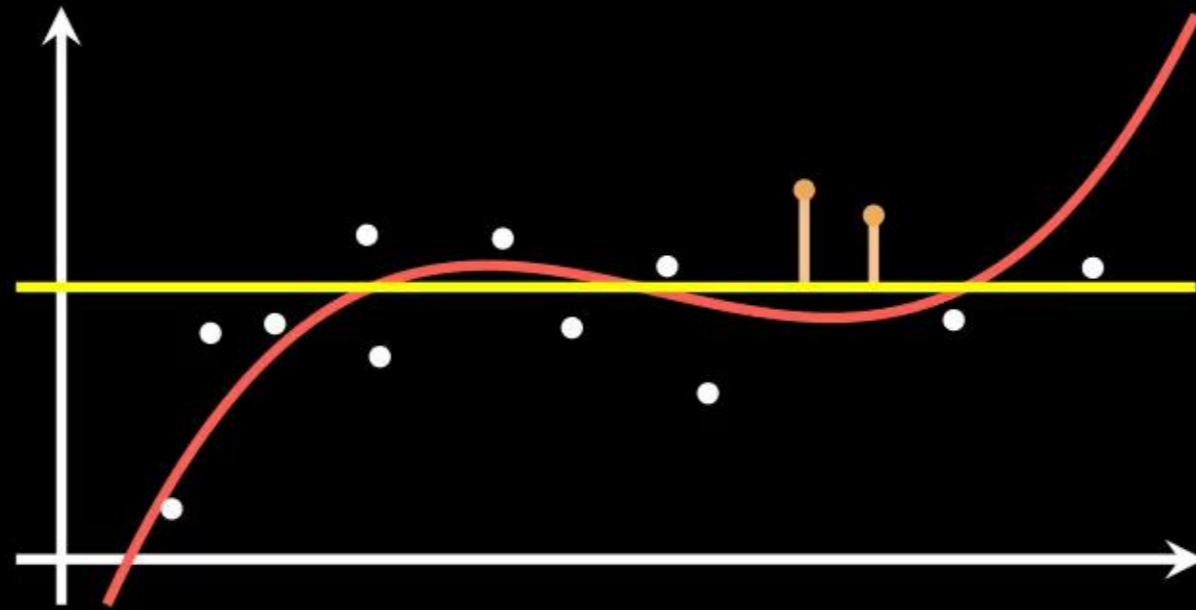
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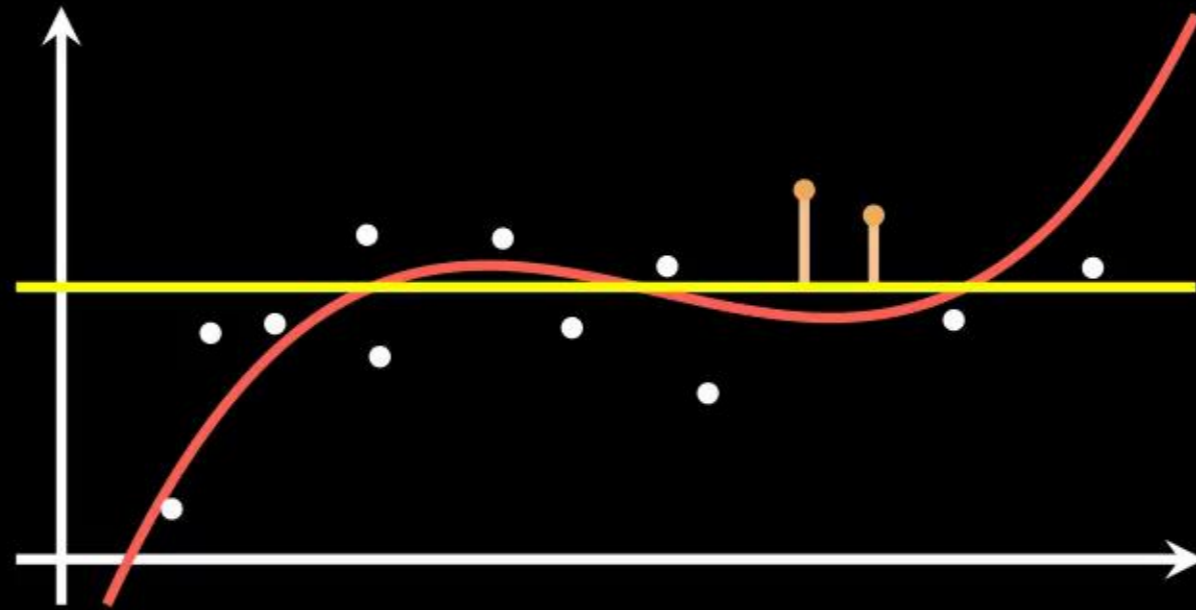


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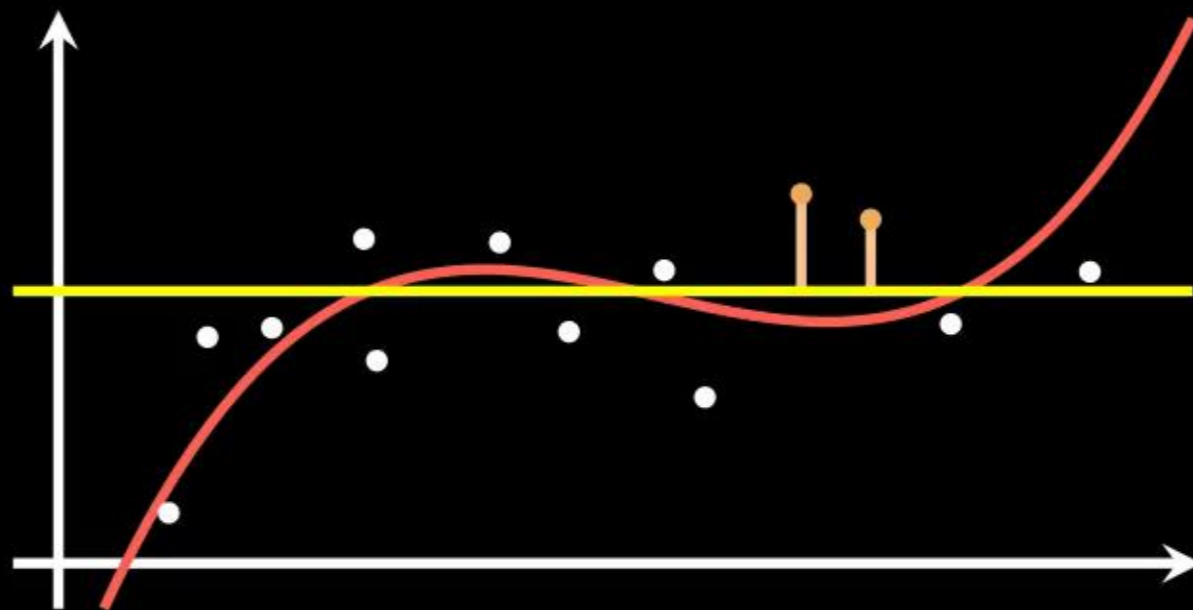
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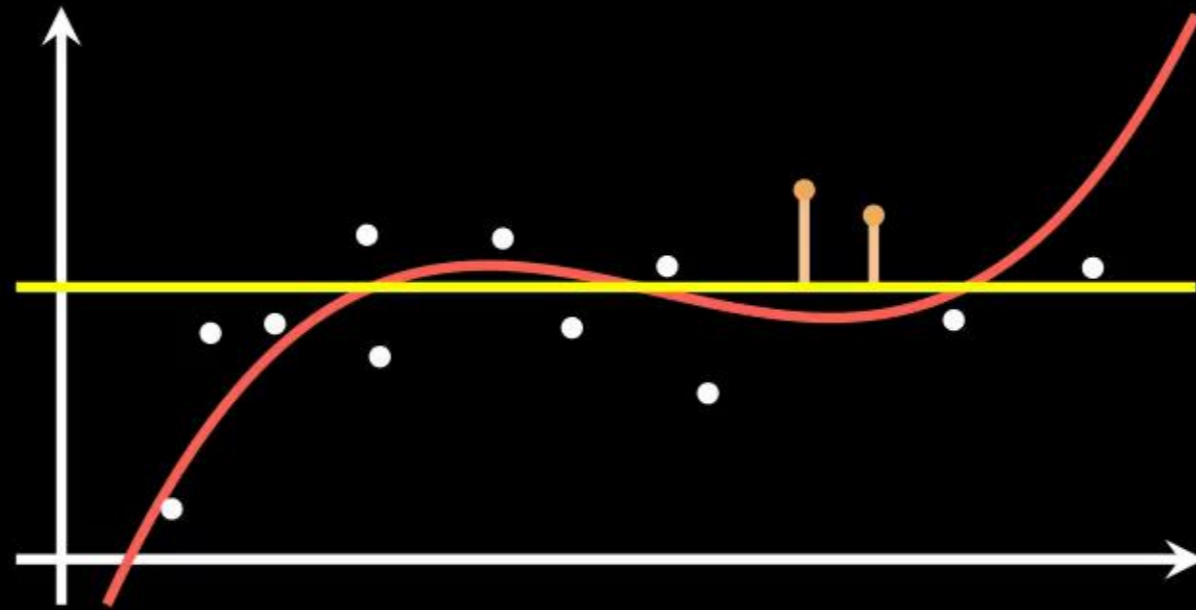
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→ Asymmetry of errors

Statistical setting

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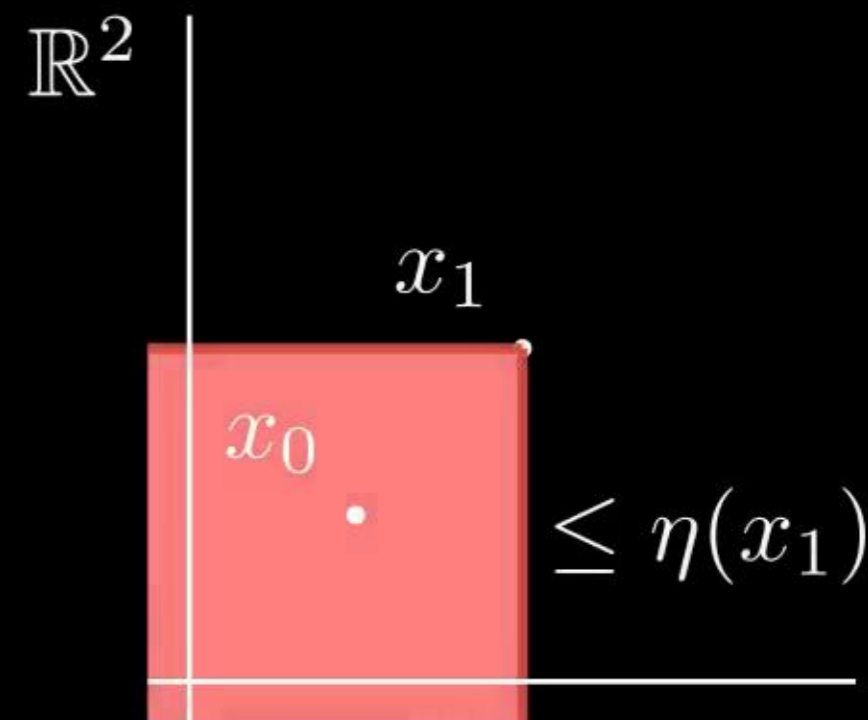
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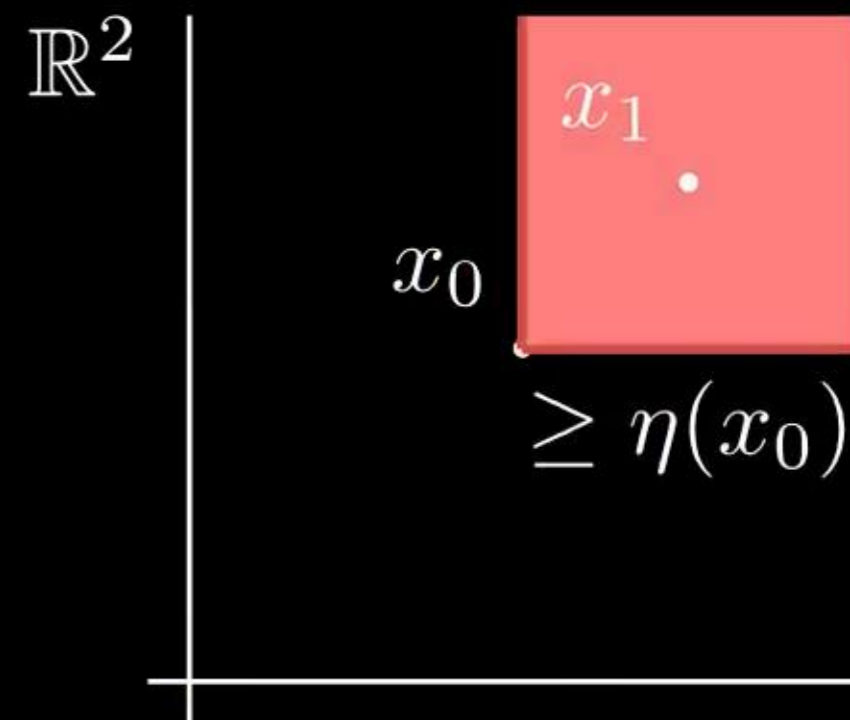


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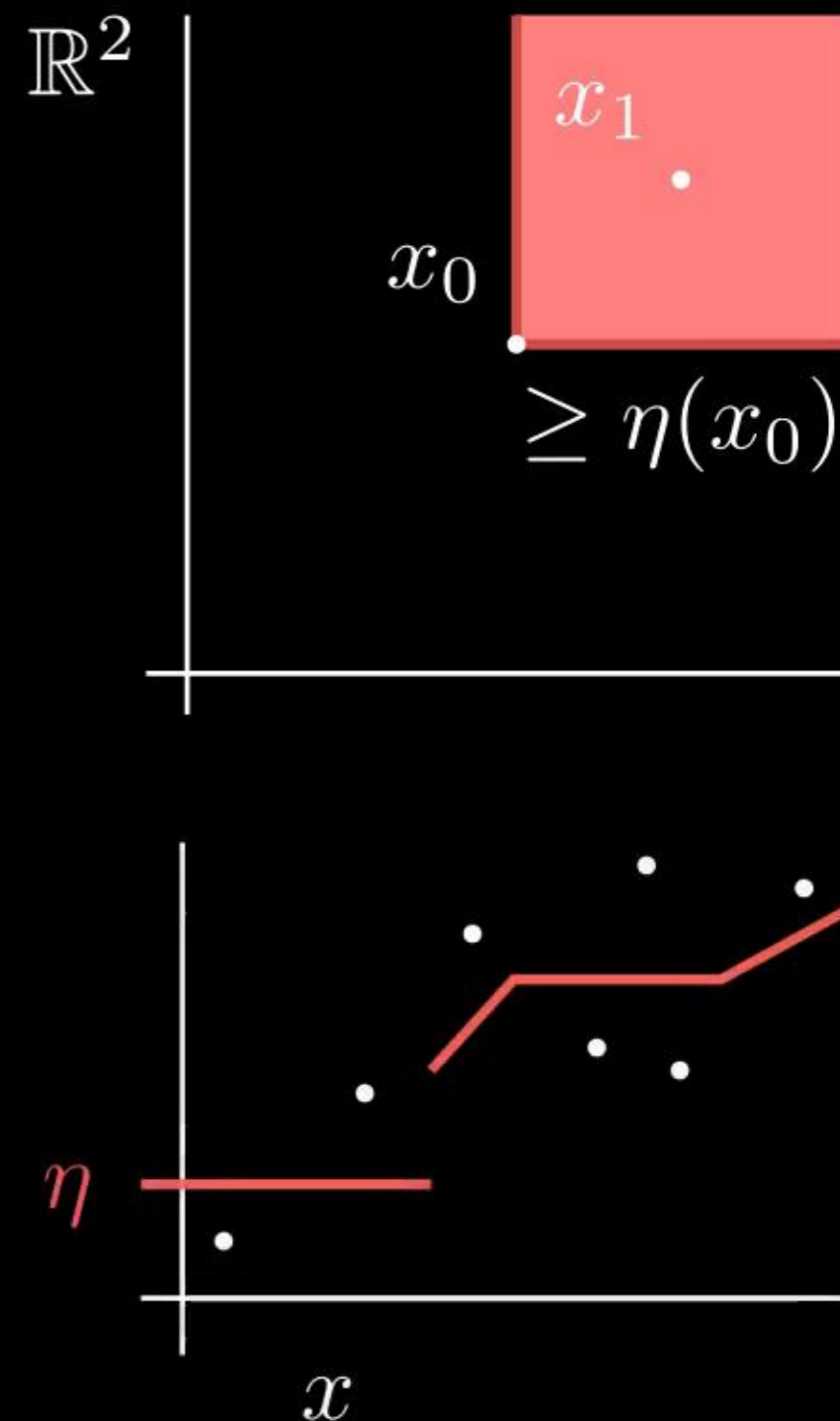


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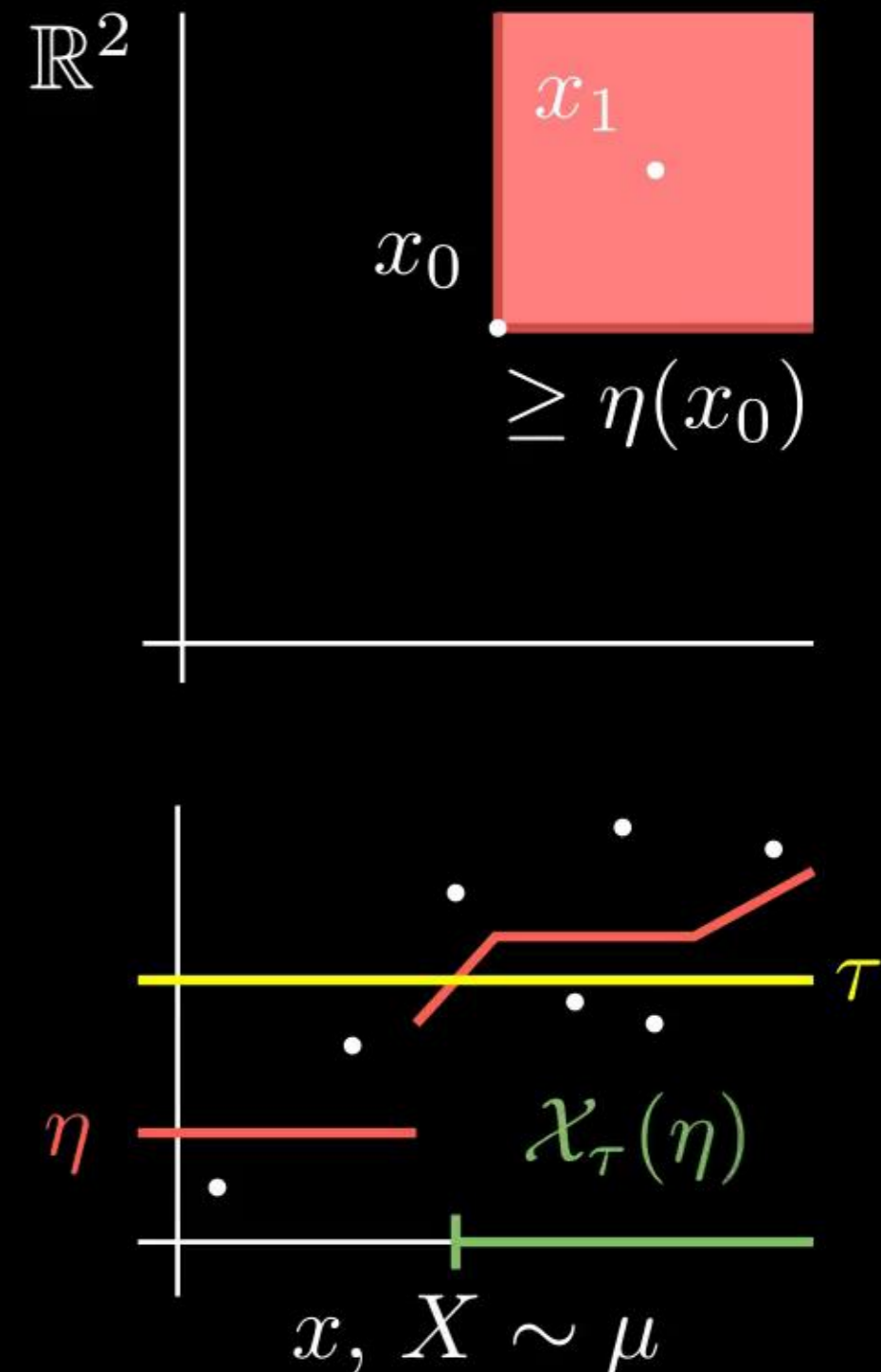
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Notation:

- Fix $\tau \in \mathbb{R}$. Define τ -superlevel set by

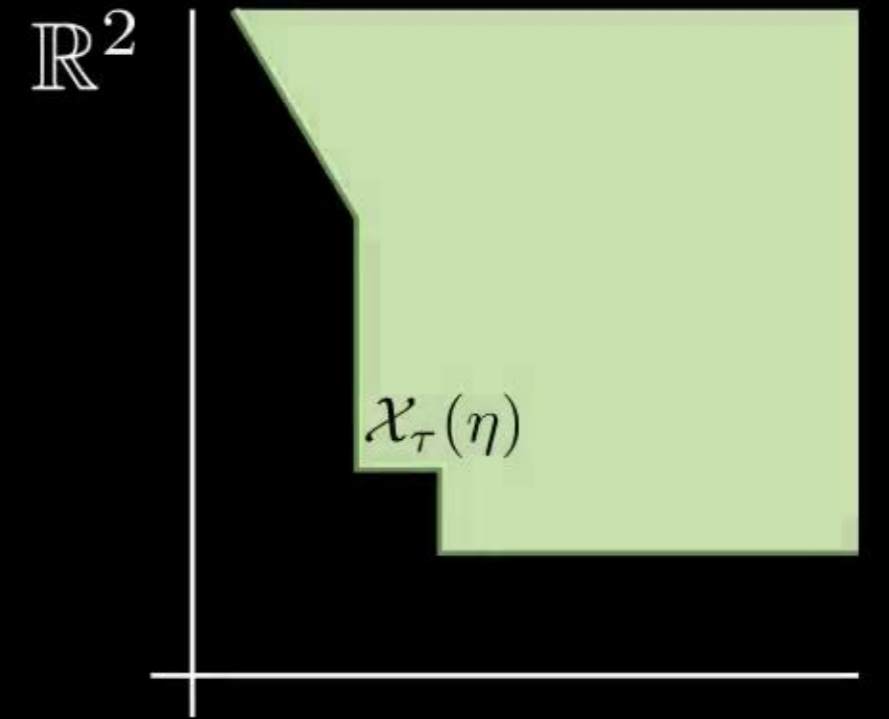
$$\mathcal{X}_\tau(\eta) := \{x \in \mathbb{R}^d : \eta(x) \geq \tau\}$$

- Denote the marginal distribution of X by μ .



Goal

Writing $\mathcal{D} := ((X_1, Y_1), \dots, (X_n, Y_n)) \sim P^n$, we want $\hat{A} : \mathcal{D} \mapsto \hat{A}(\mathcal{D}) \subseteq \mathbb{R}^d$ such that:

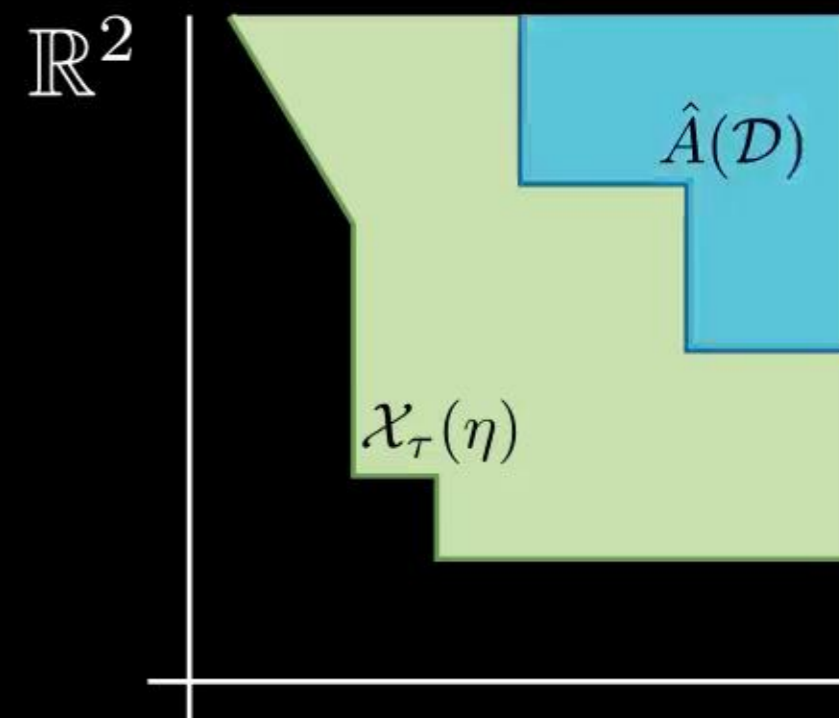


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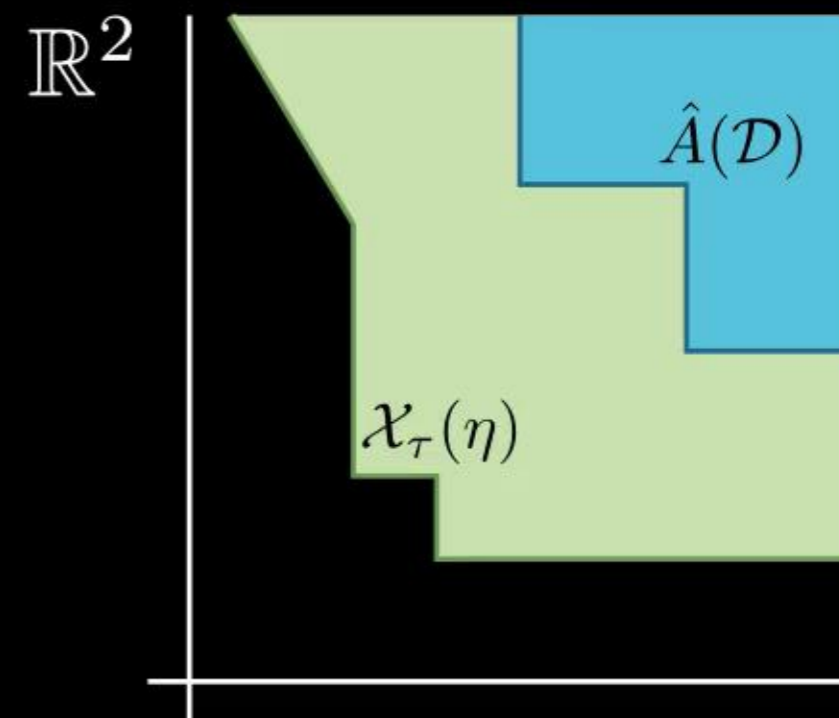
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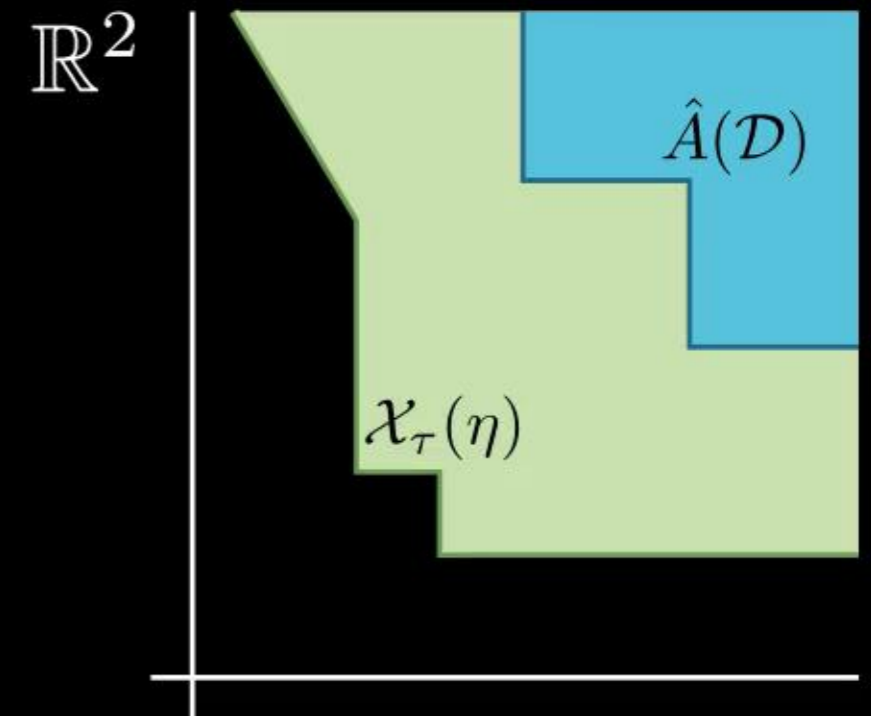
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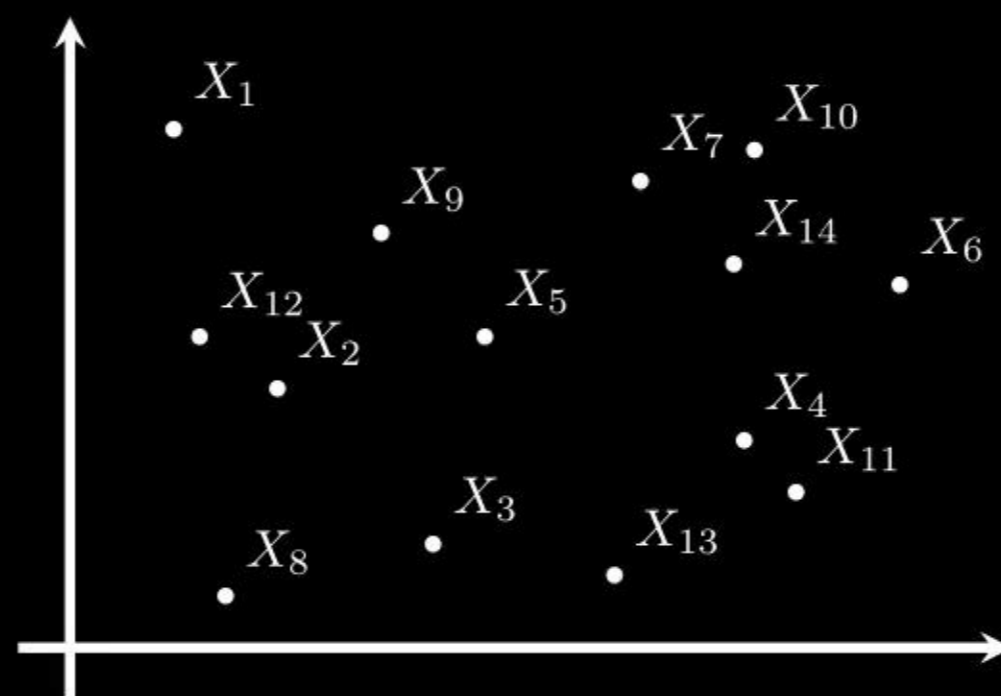
Example:

Type I error: \hat{A} should contain only those HIV-patients for whom the conditional probability of not facing a negative event within the next 5 years is at least 98%.



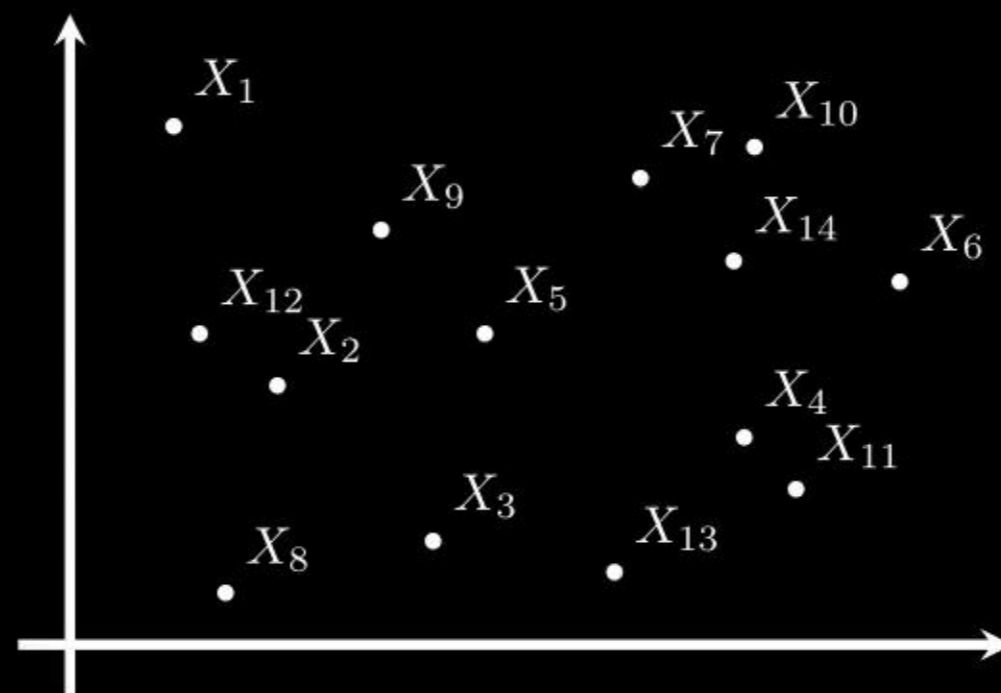
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For $x_0 \in \mathbb{R}^d$, define null hypothesis $H_0(x_0) := \{P \in \mathcal{P}_{\text{Mon},d}(\sigma) : \eta(x_0) < \tau\}$.



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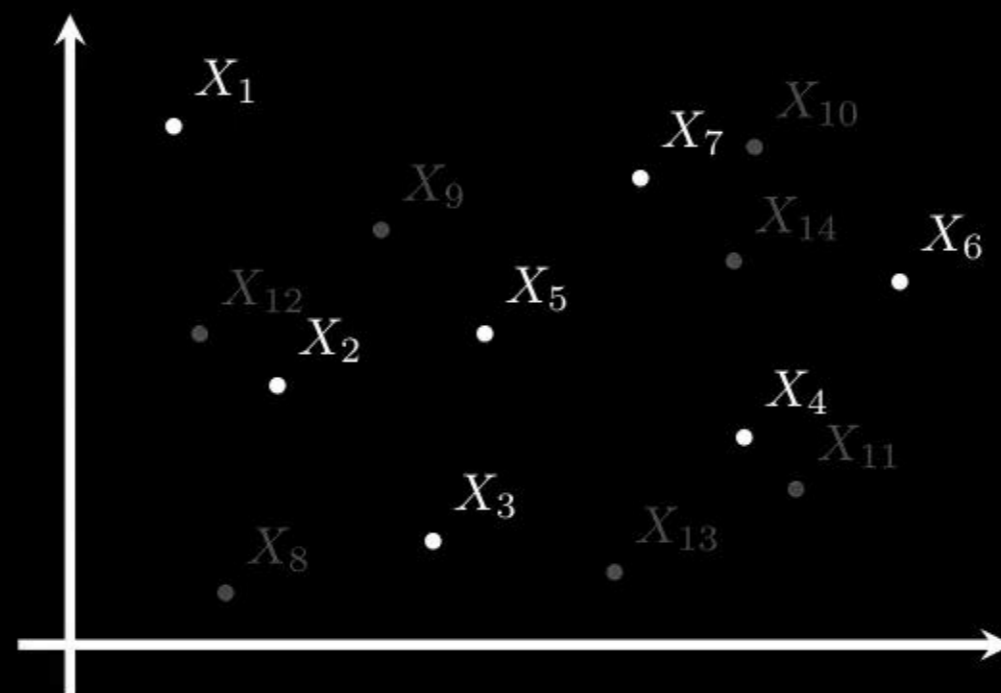
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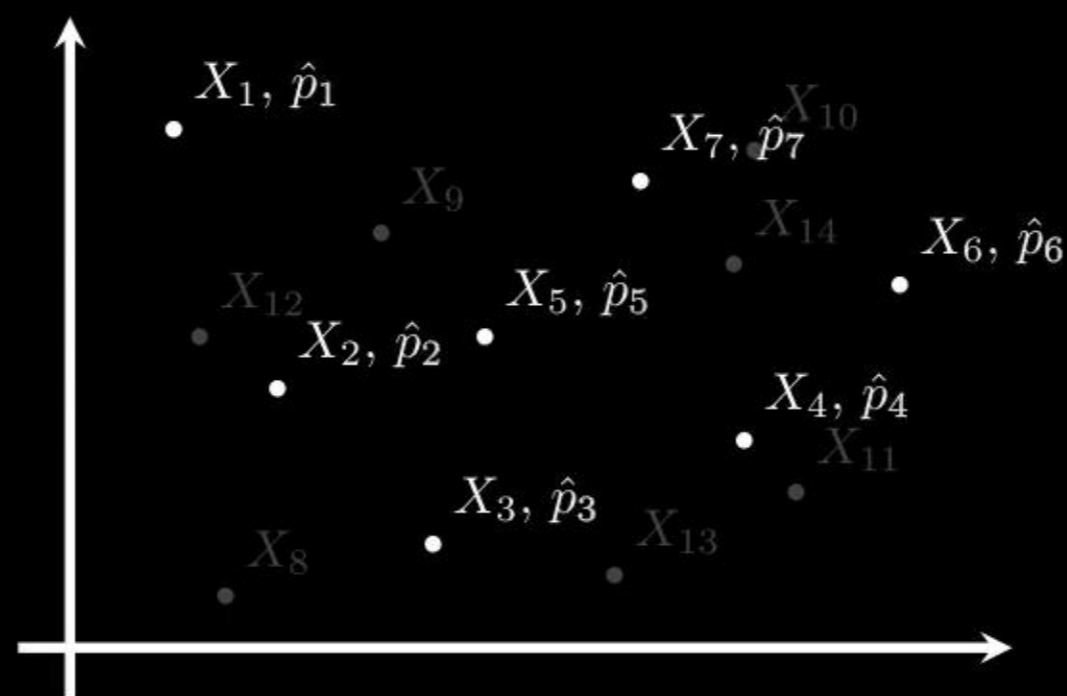


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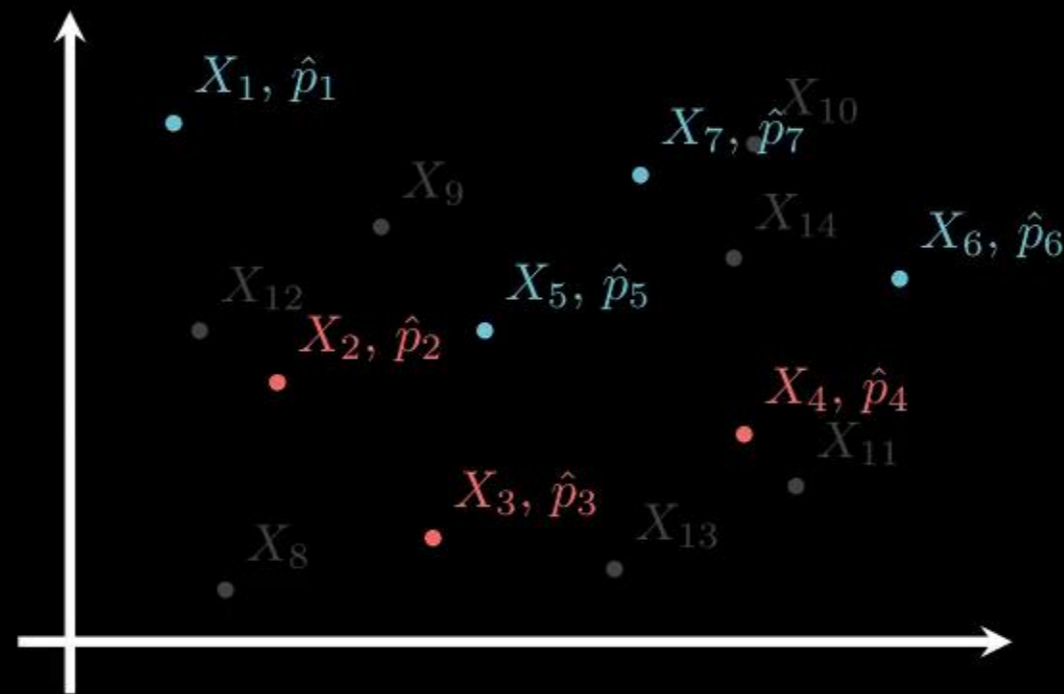


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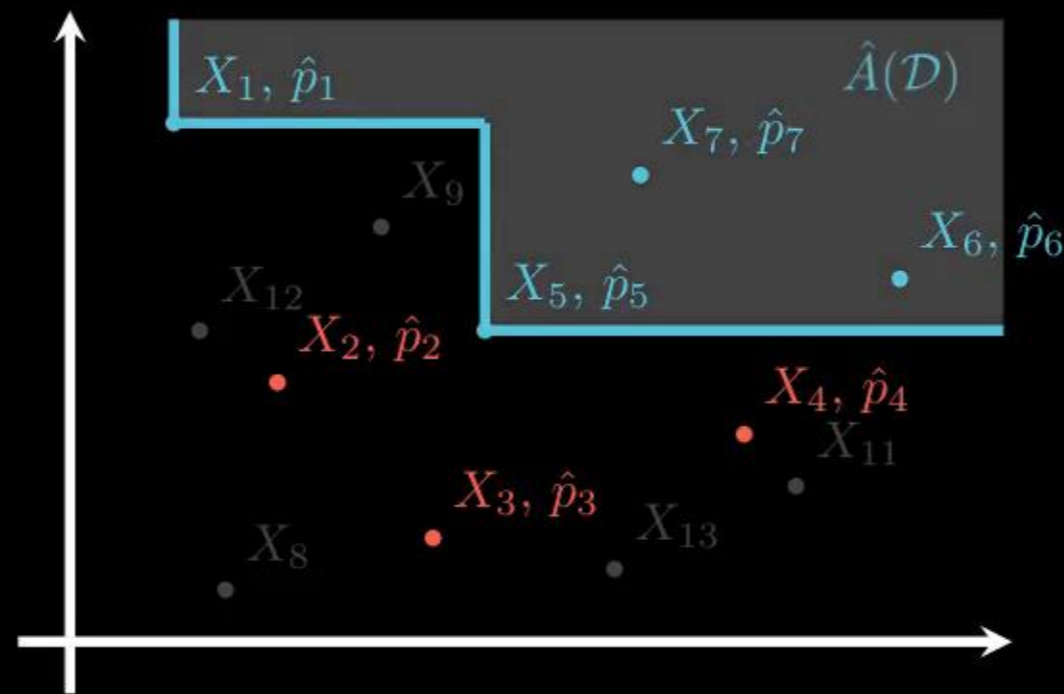


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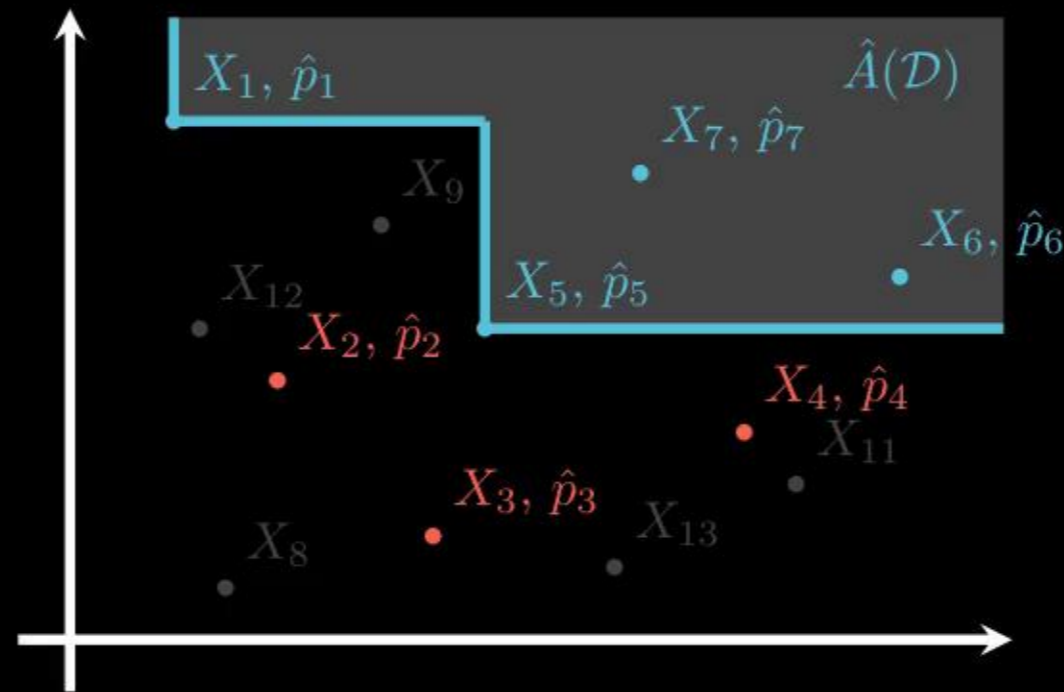


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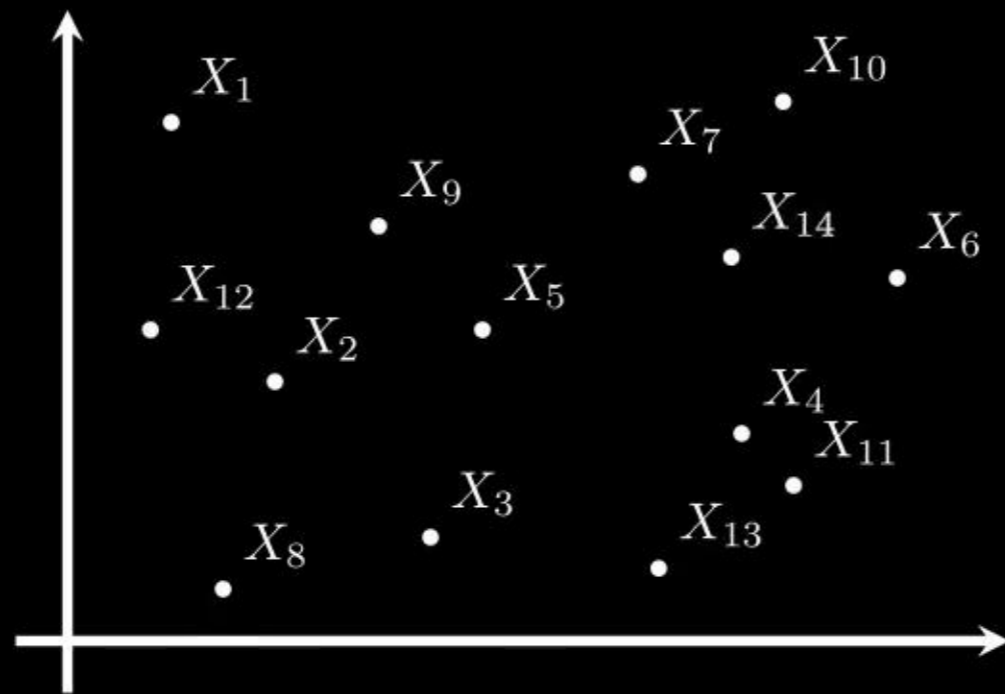
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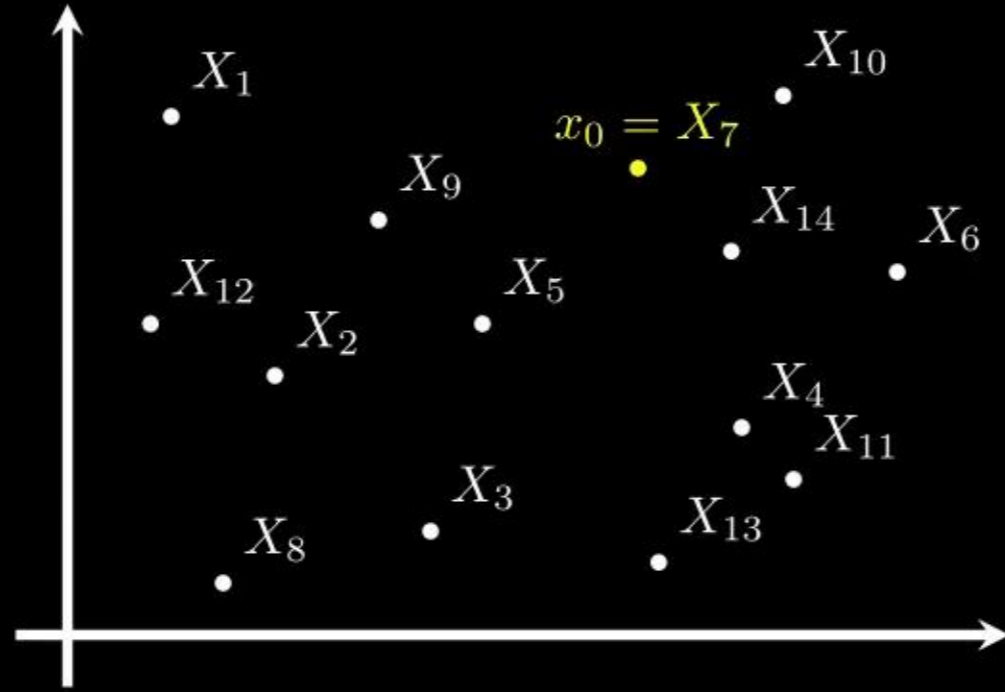
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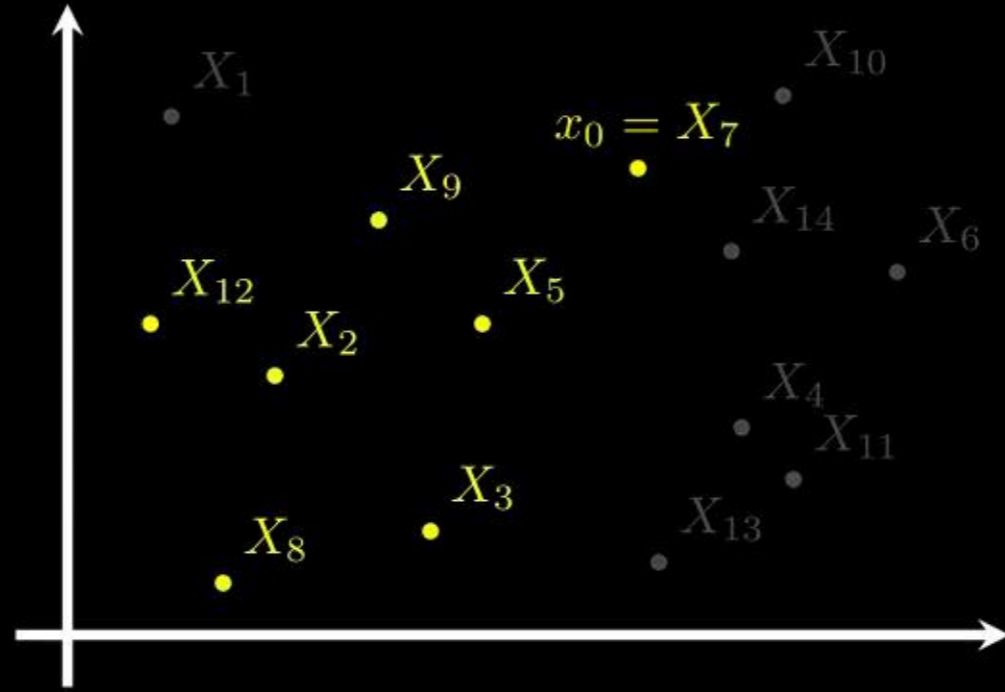
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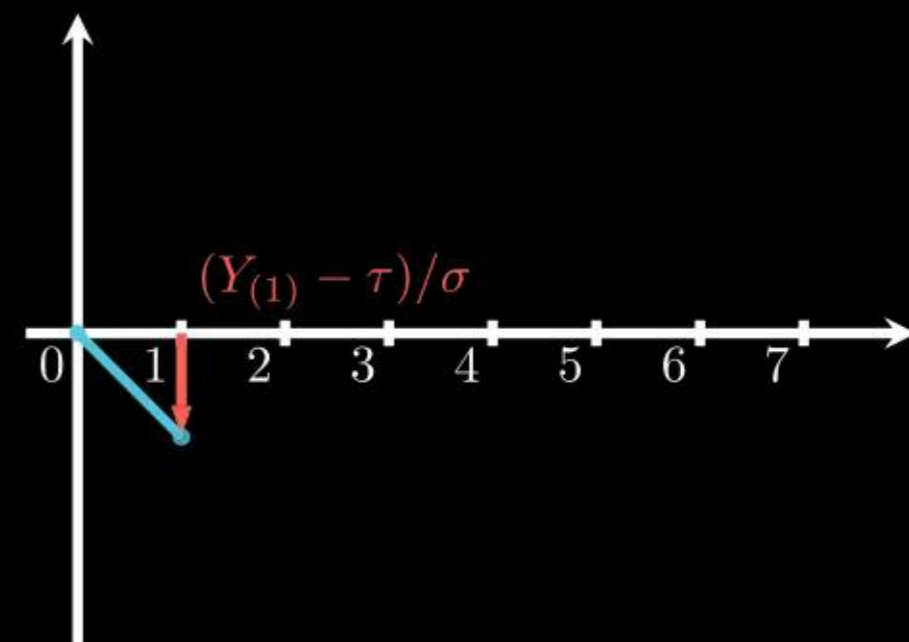
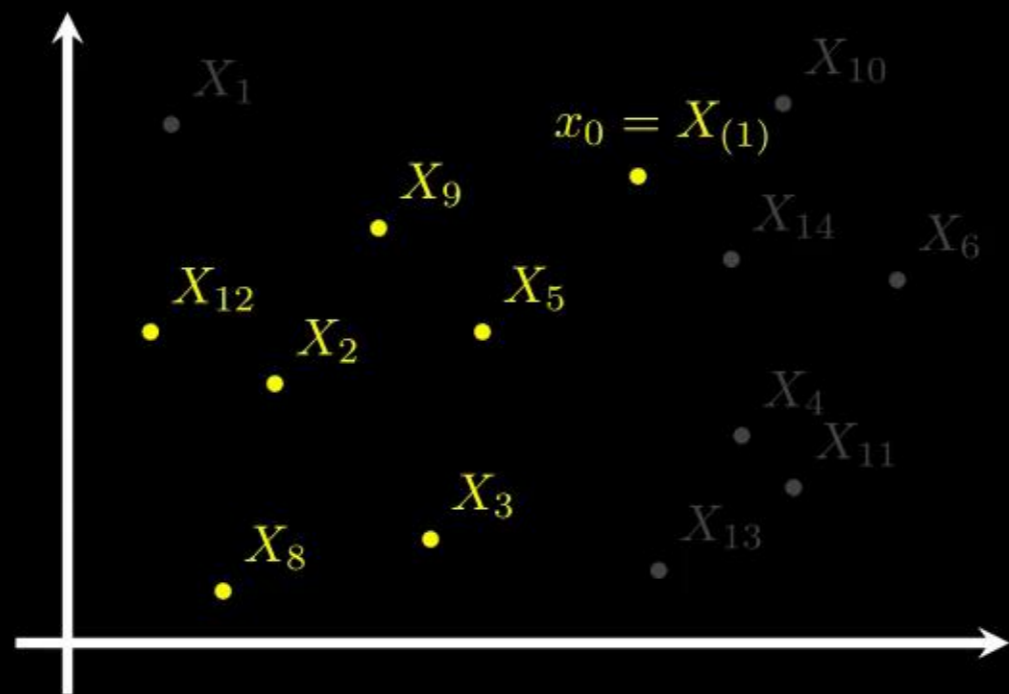
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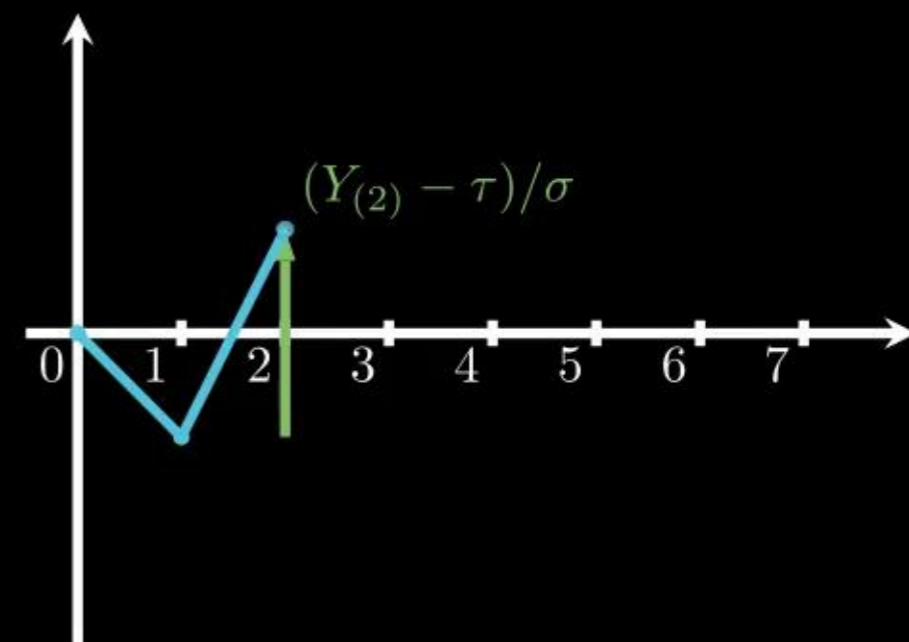
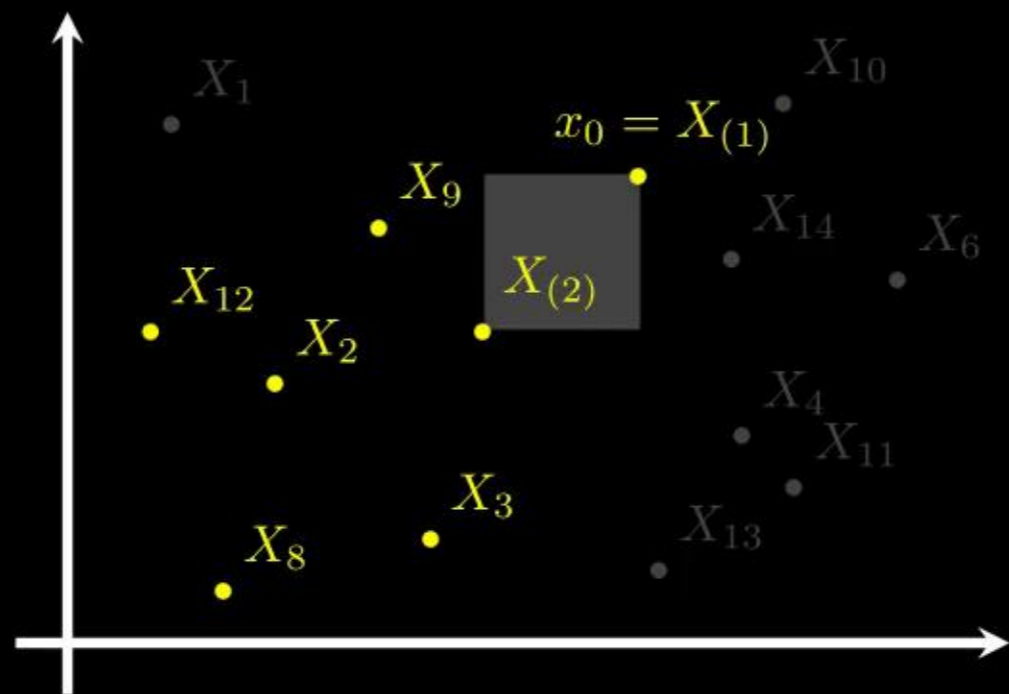


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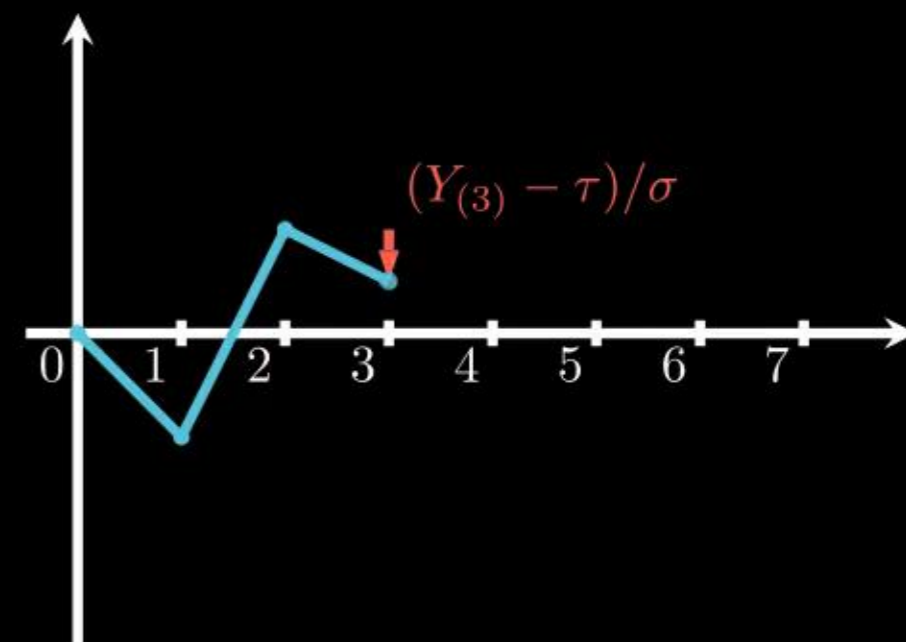
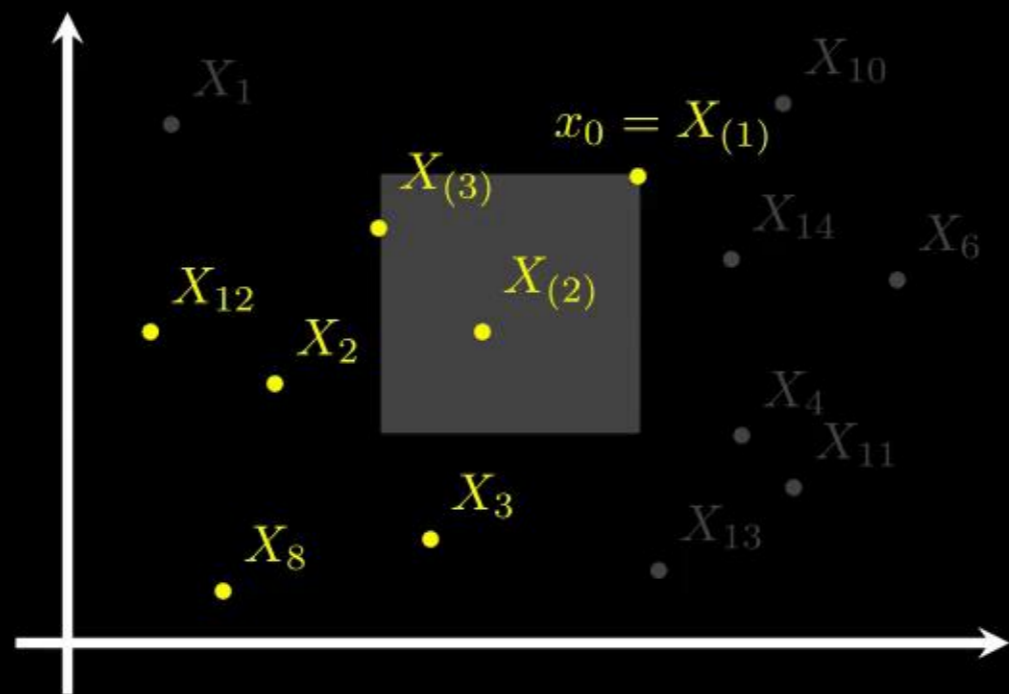


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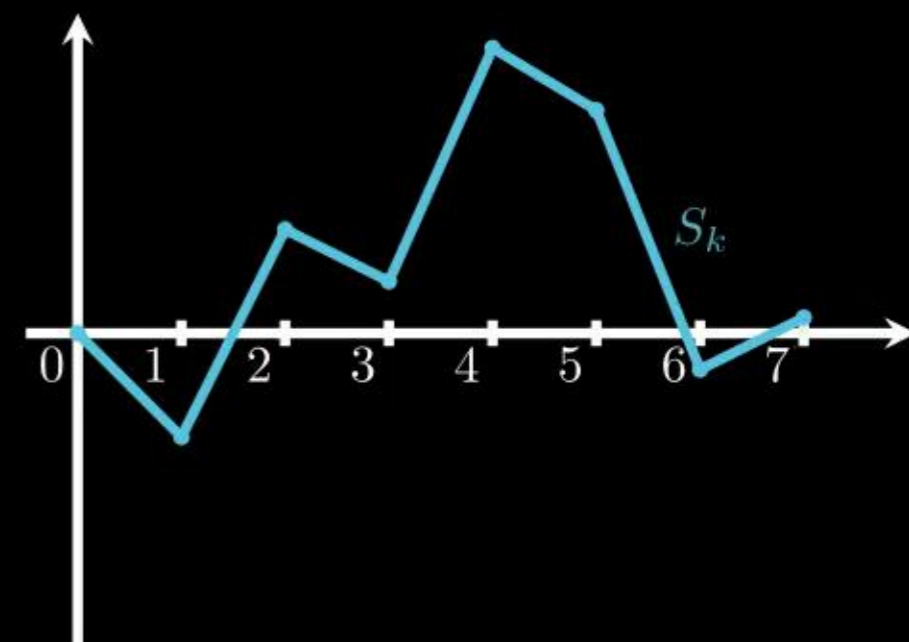
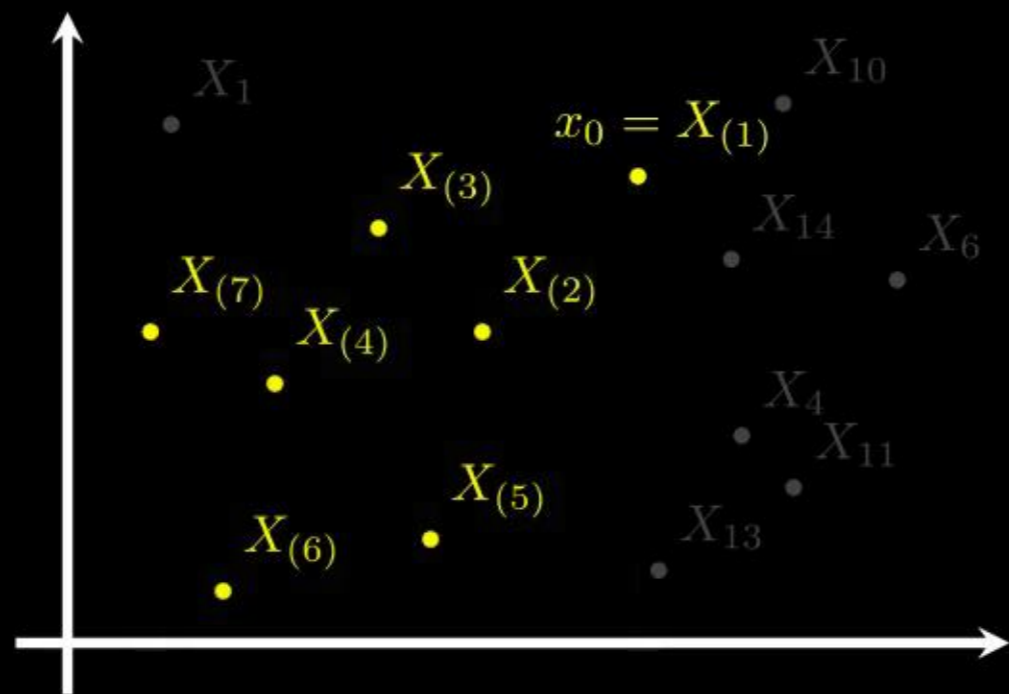


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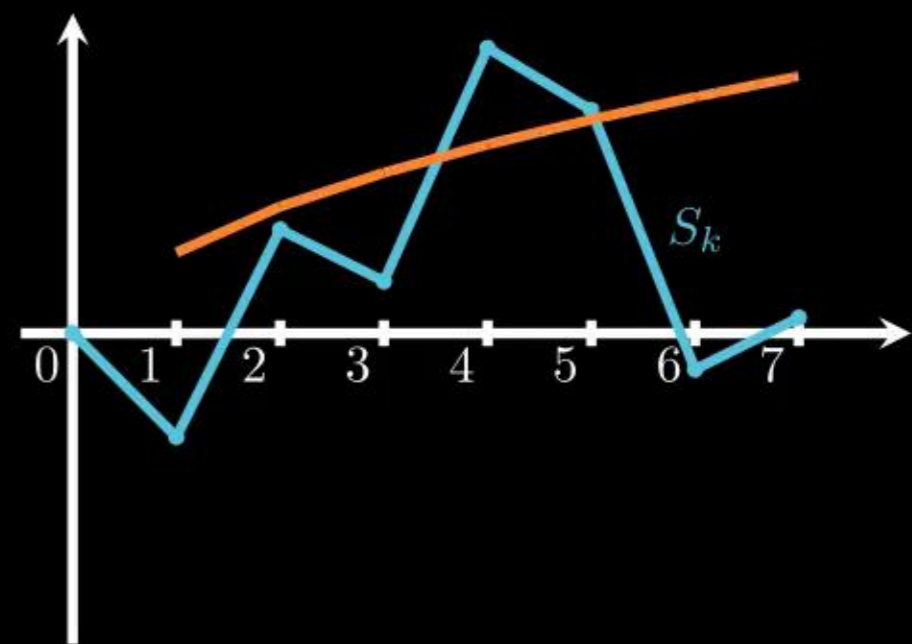
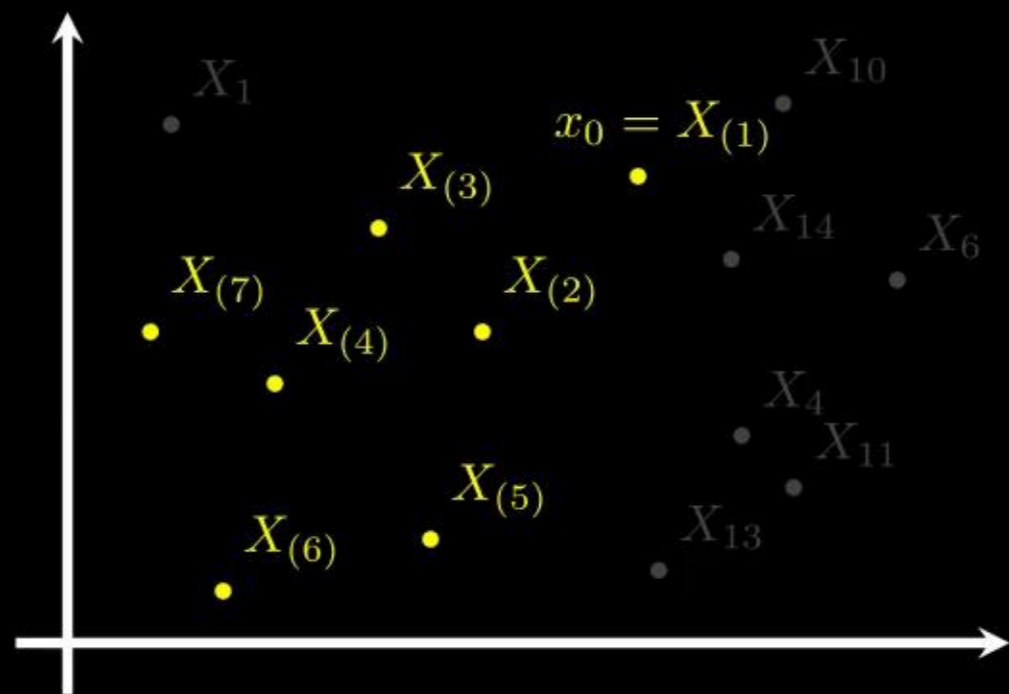


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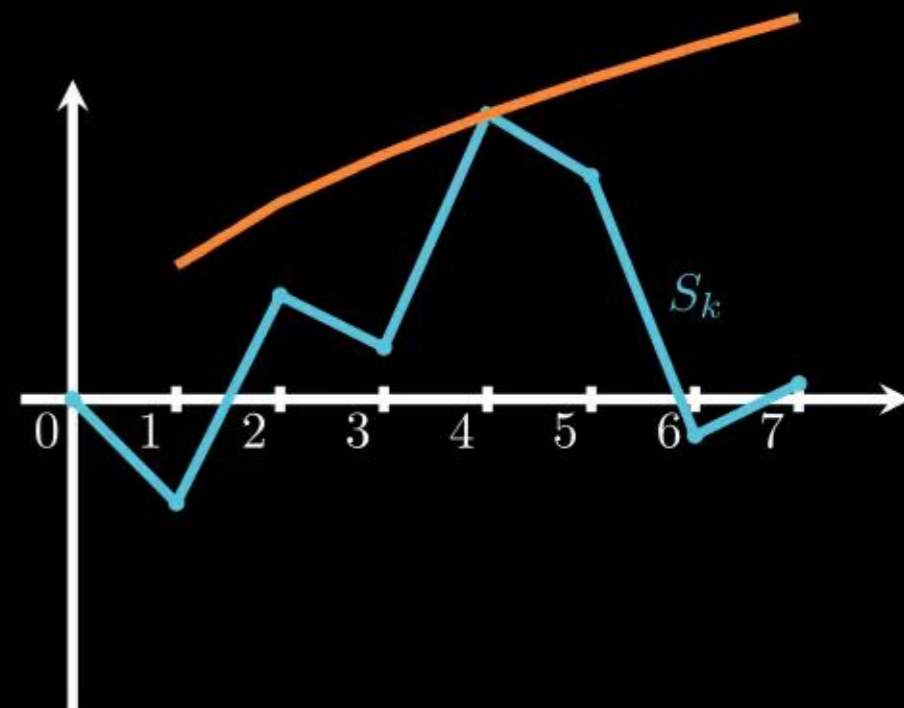
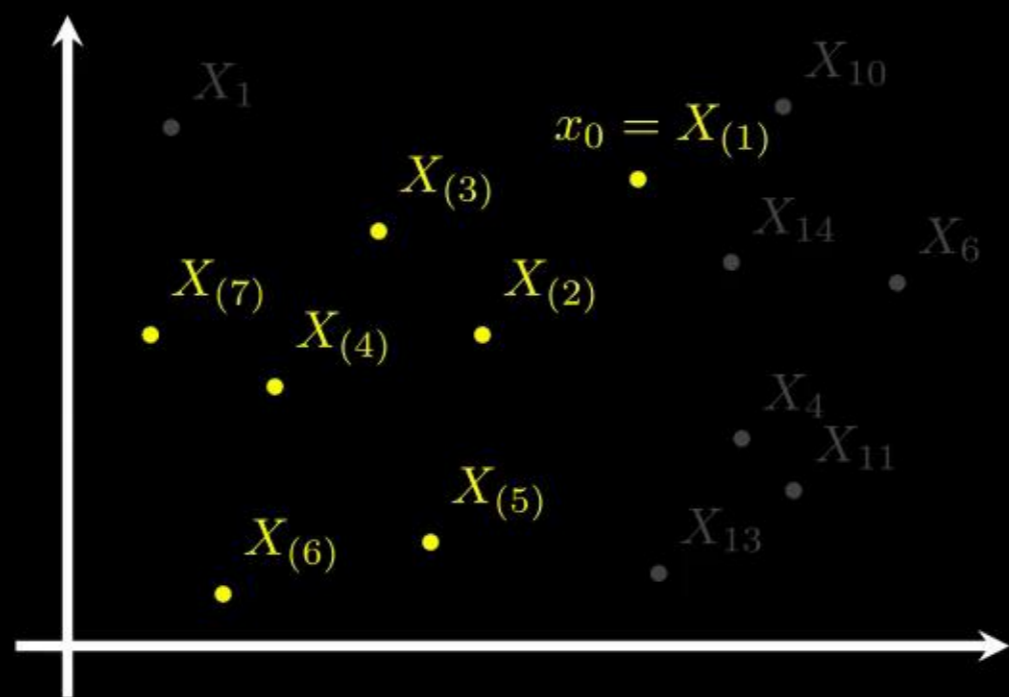
Let

$$S_k := \sum_{j=1}^k \frac{Y_{(j)} - \tau}{\sigma}.$$

Then S_k is a supermartingale under $P \in H_0(x_0)$. Combination with time-uniform bounds by Howard et al. (2021) gives p -values from this martingale test (Duan et al., 2020).

Construct p -values \hat{p}_i for $H_0(X_i)$, $i \in [m]$

Given $x_0 \in \mathbb{R}^d$, we seek a p -value for $H_0(x_0) := \{P \in \mathcal{P}_{\text{Mon},d}(\sigma) : \eta(x_0) < \tau\}$.



Denote $\mathcal{I}(x_0) := \{i \in [n] : X_i \preceq x_0\}$, $n(x_0) := |\mathcal{I}(x_0)|$.

Let $X_{(j)}$ be the j th nearest neighbour of x_0 among X_i , $i \in \mathcal{I}(x_0)$, in sup-norm and let $Y_{(j)}$ be the corresponding response.

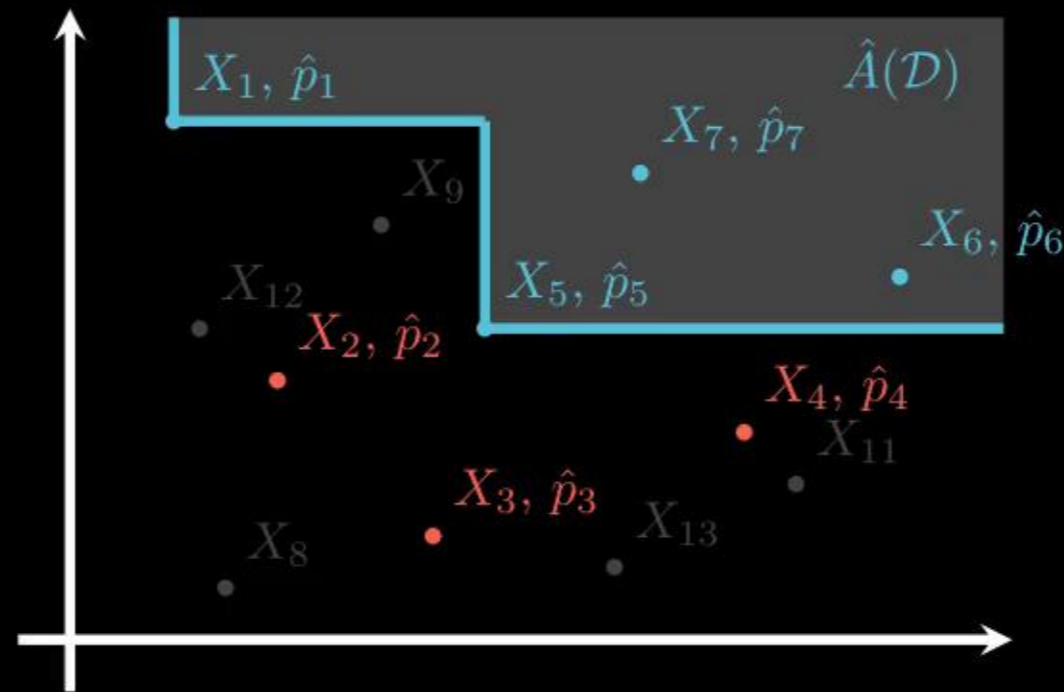
Let

$$S_k := \sum_{j=1}^k \frac{Y_{(j)} - \tau}{\sigma}.$$

Then S_k is a supermartingale under $P \in H_0(x_0)$. Combination with time-uniform bounds by Howard et al. (2021) gives p -values from this martingale test (Duan et al., 2020).

High-level strategy

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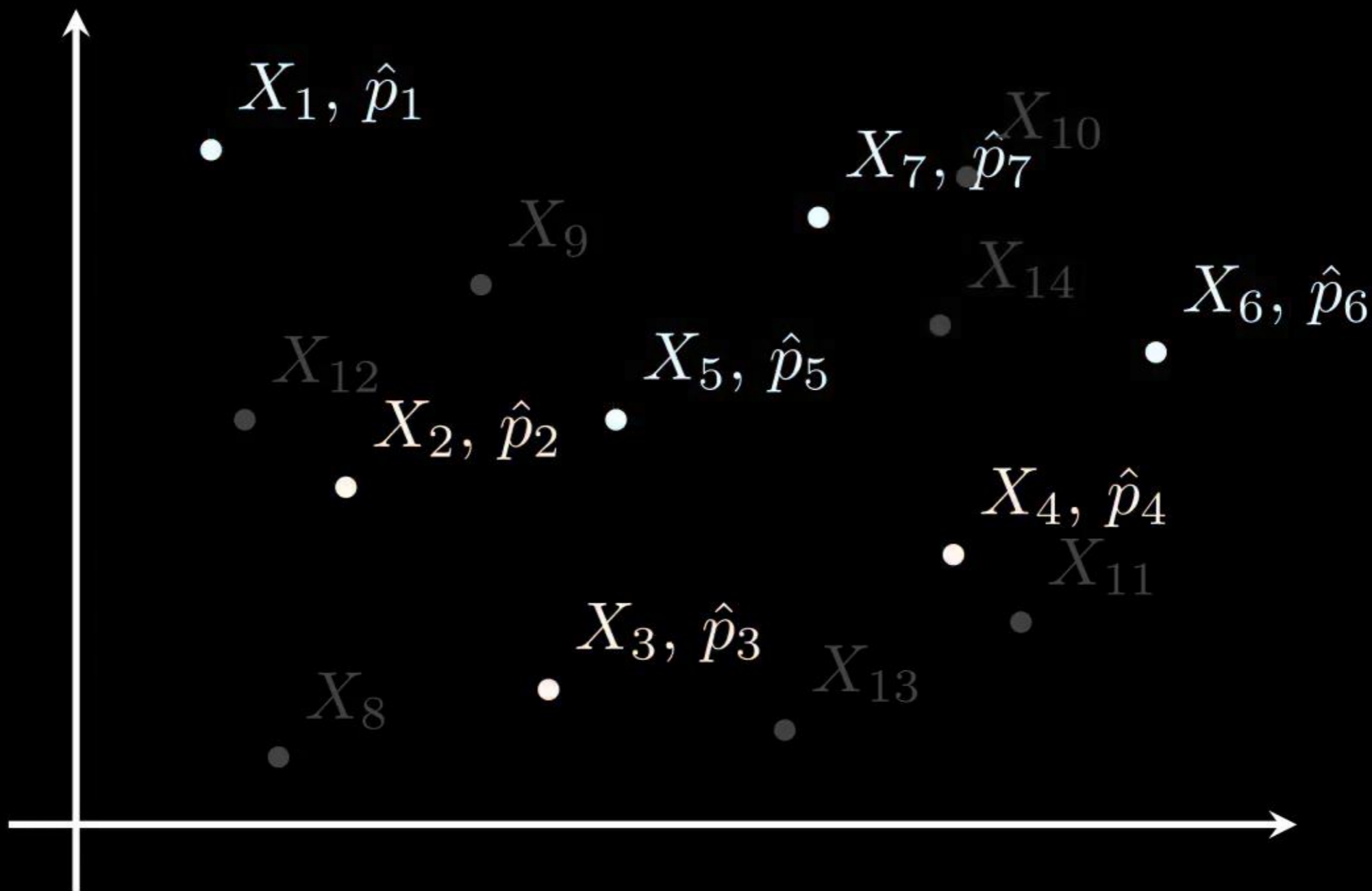


High-level strategy:

1. Subsample m covariate vectors X_1, \dots, X_m with $m \leq n$;
2. Calculate p -values \hat{p}_i for $H_0(X_i)$, $i \in [m]$;
3. Apply a *multiple testing procedure* with FWER-control at level α to reject $\mathcal{R}_\alpha \subseteq [m]$;
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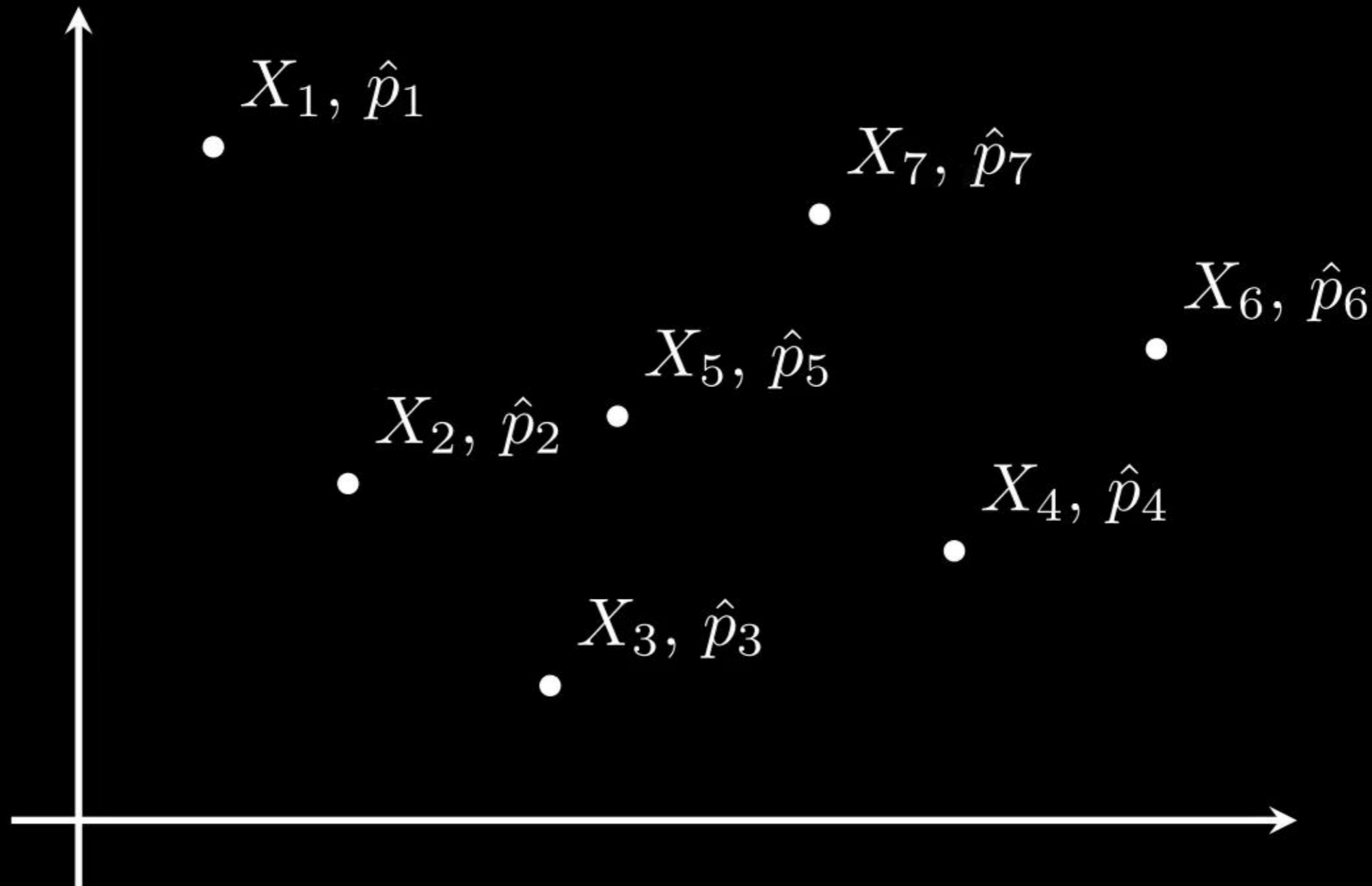
Multiple testing procedure

Key idea: logical relationships of hypotheses $H_0(X_i)$, $i \in [m]$, induce DAG with vertex set $[m]$. We combine the sequential rejection principle (Goeman and Solari, 2010) with careful α -budget allocation to construct a *DAG testing procedure*.



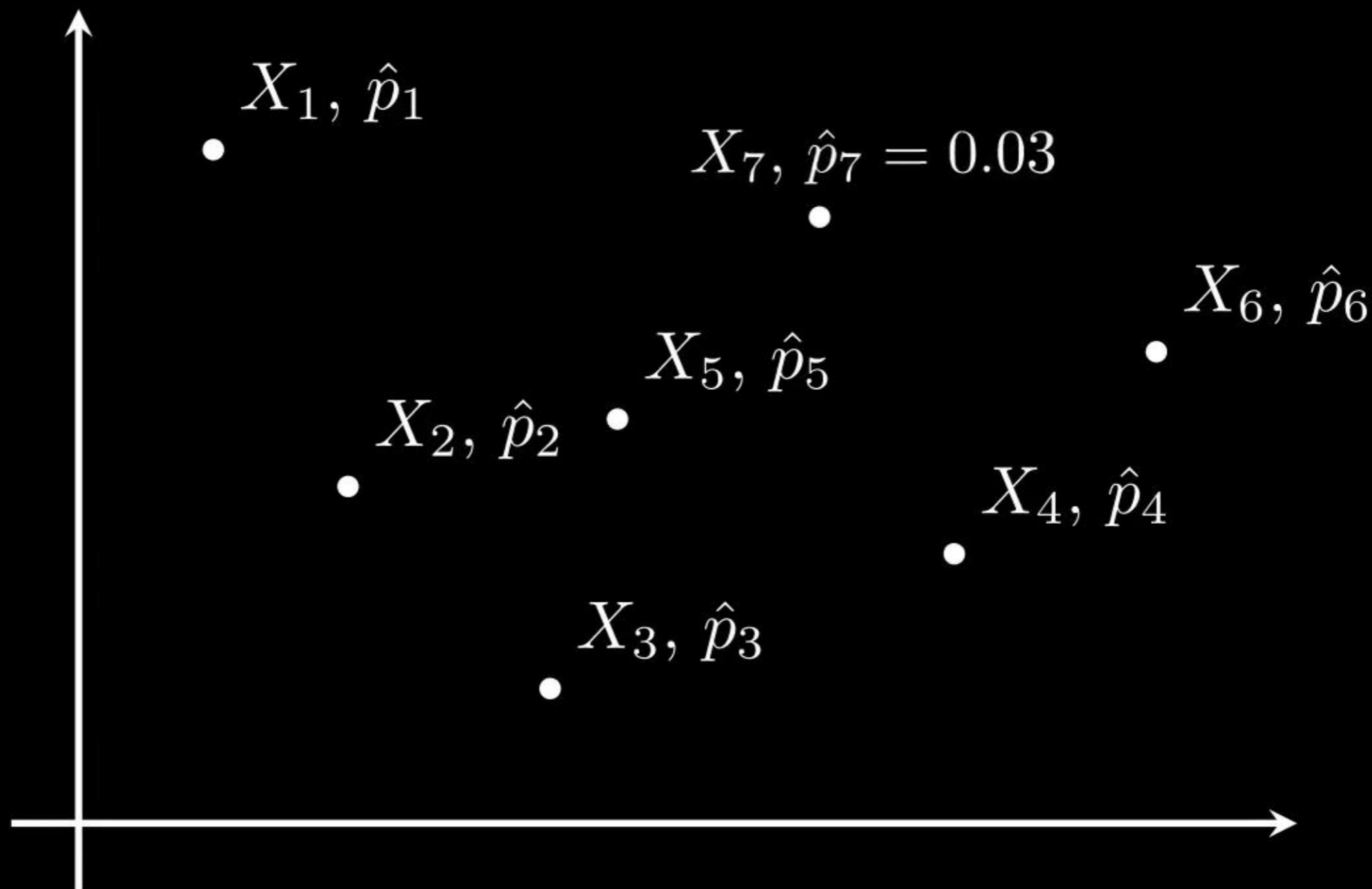
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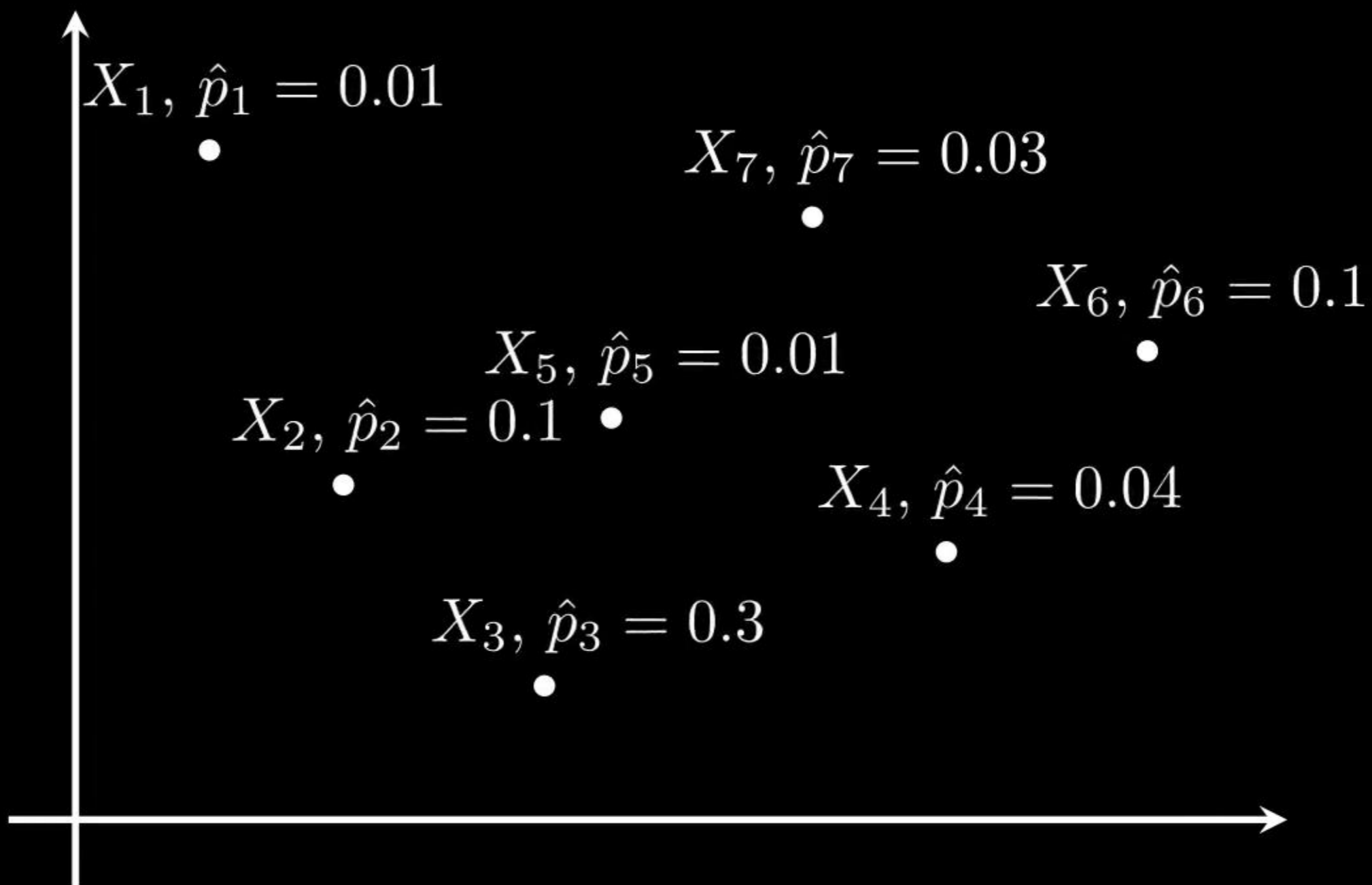
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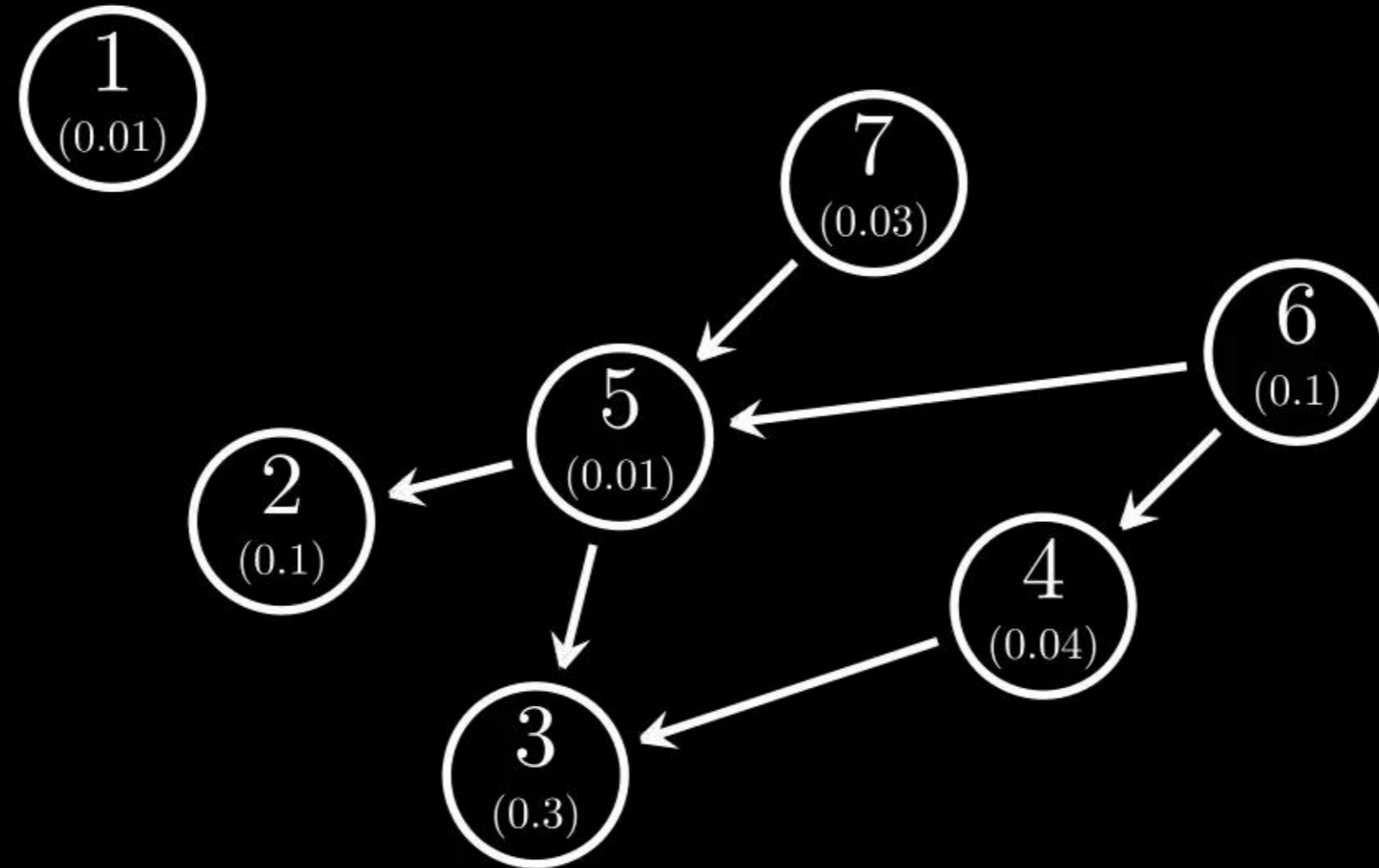
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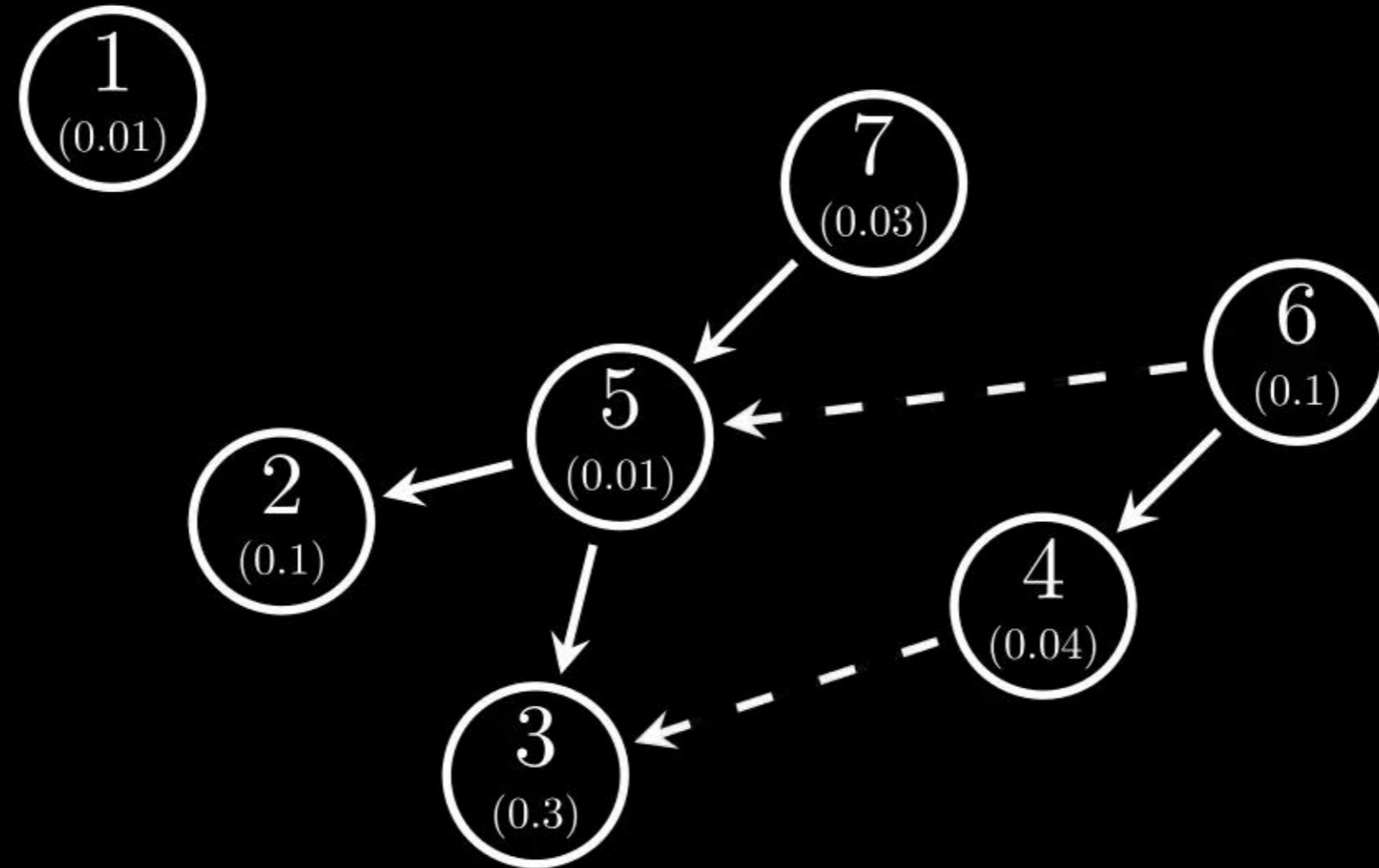
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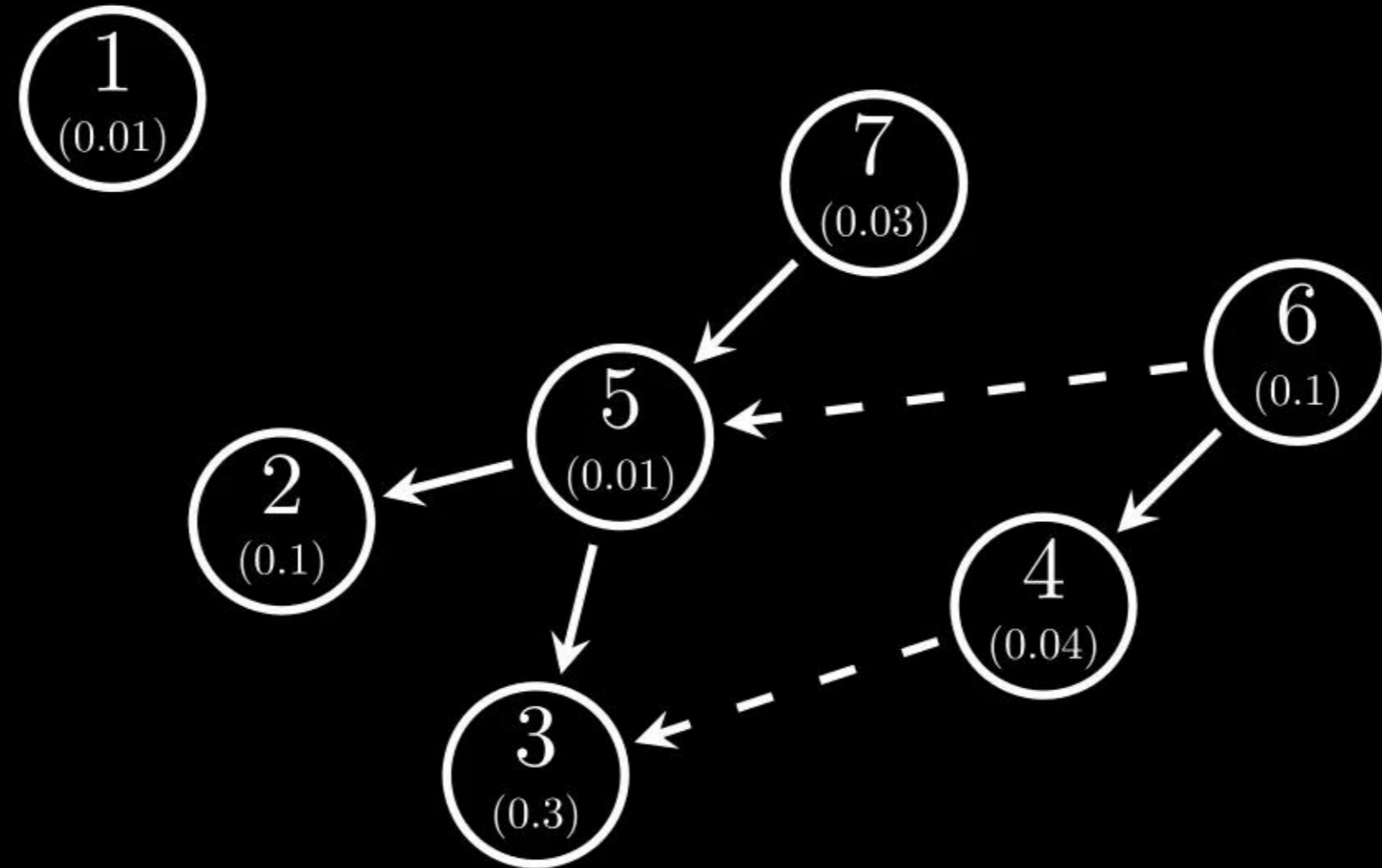
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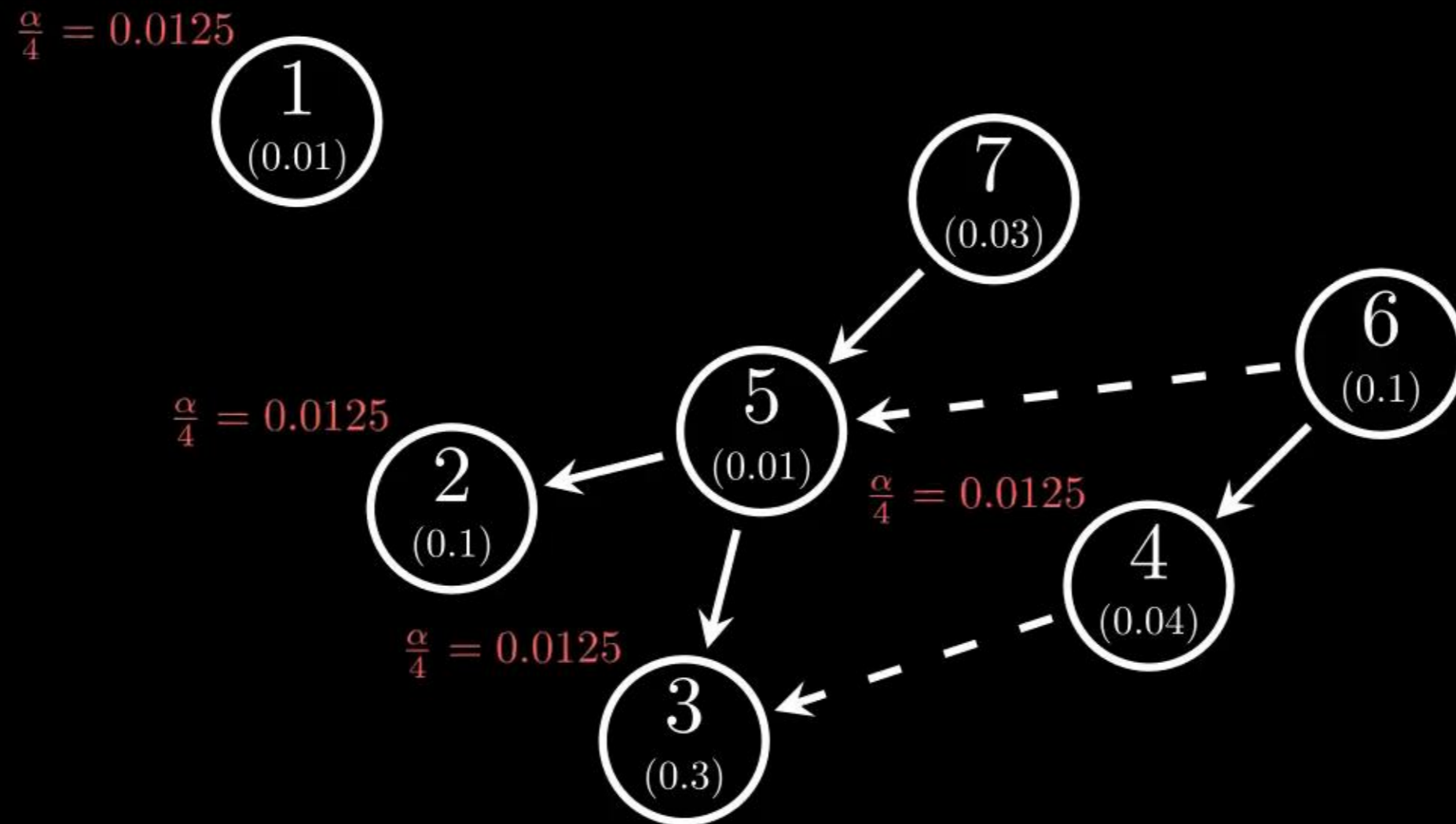
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Here: $\alpha = 0.05$.

Multiple testing procedure

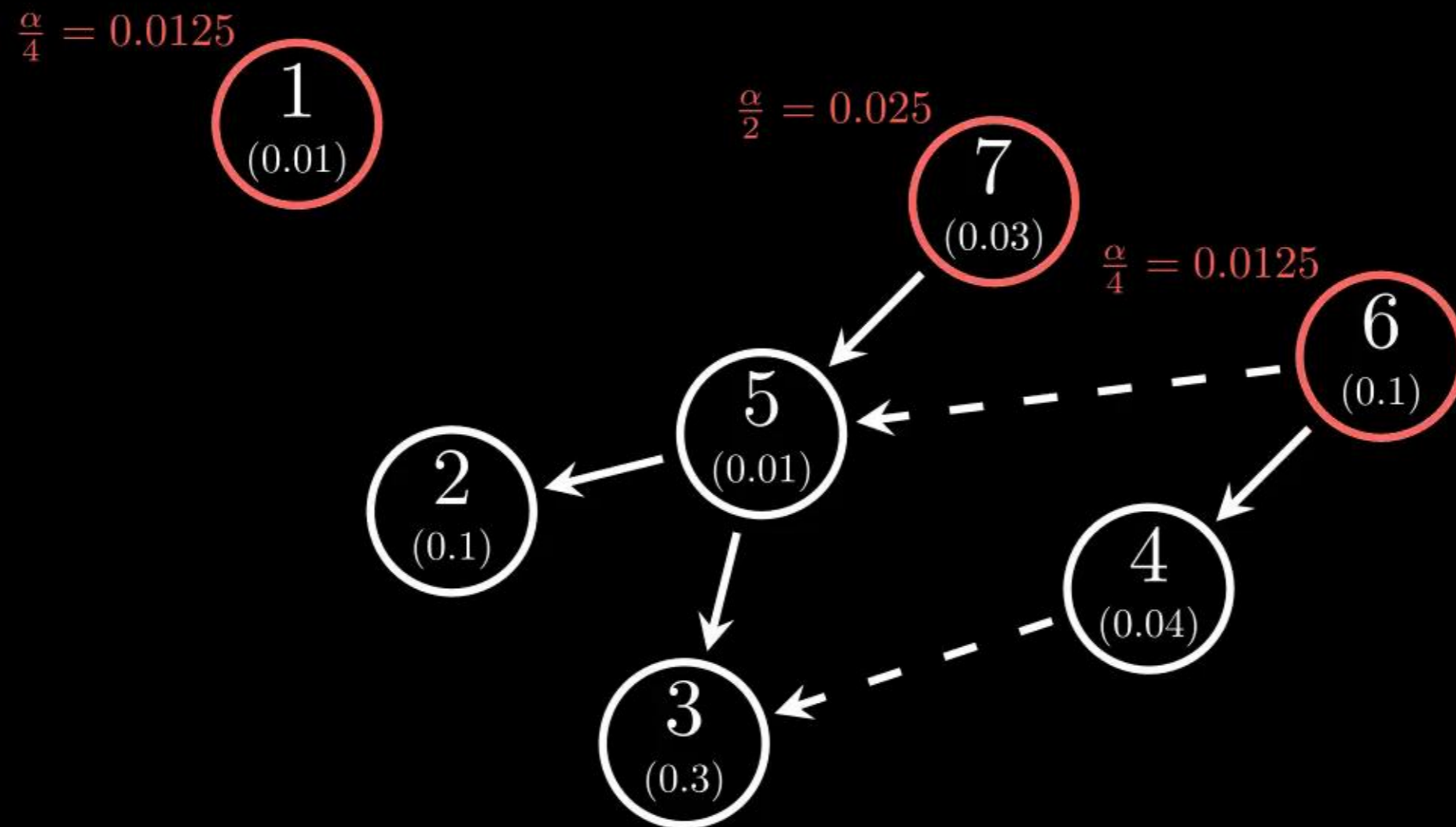
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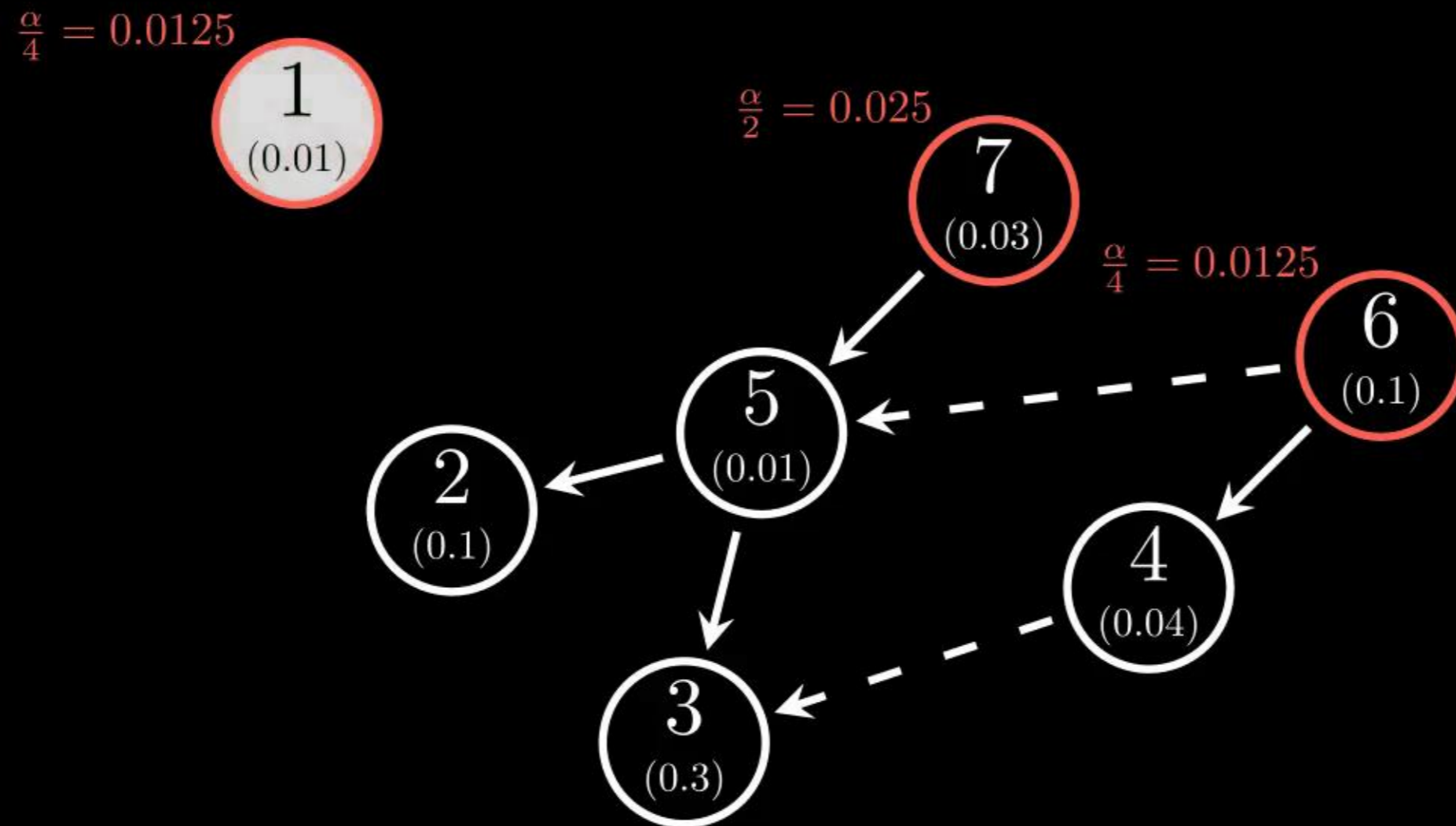
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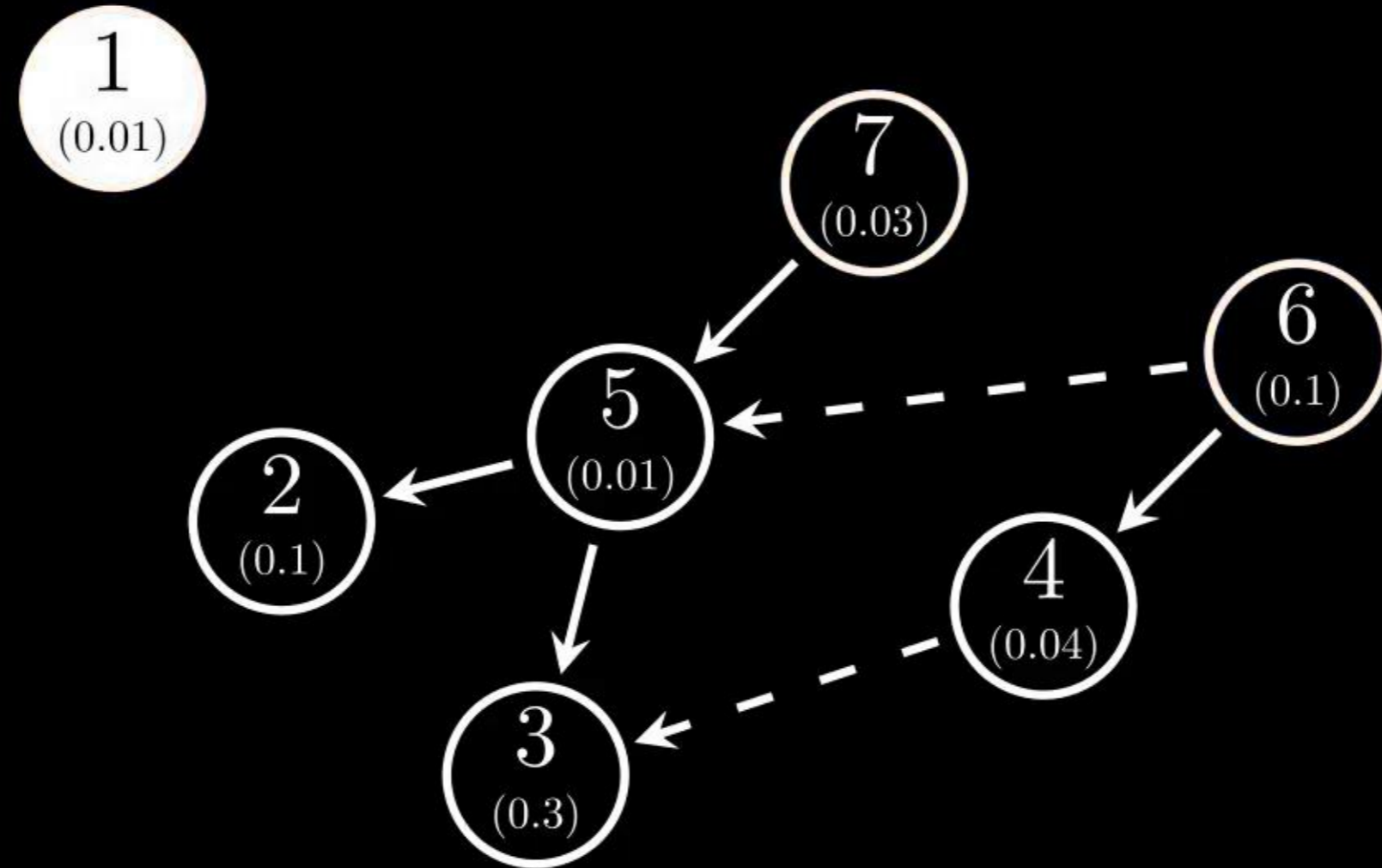
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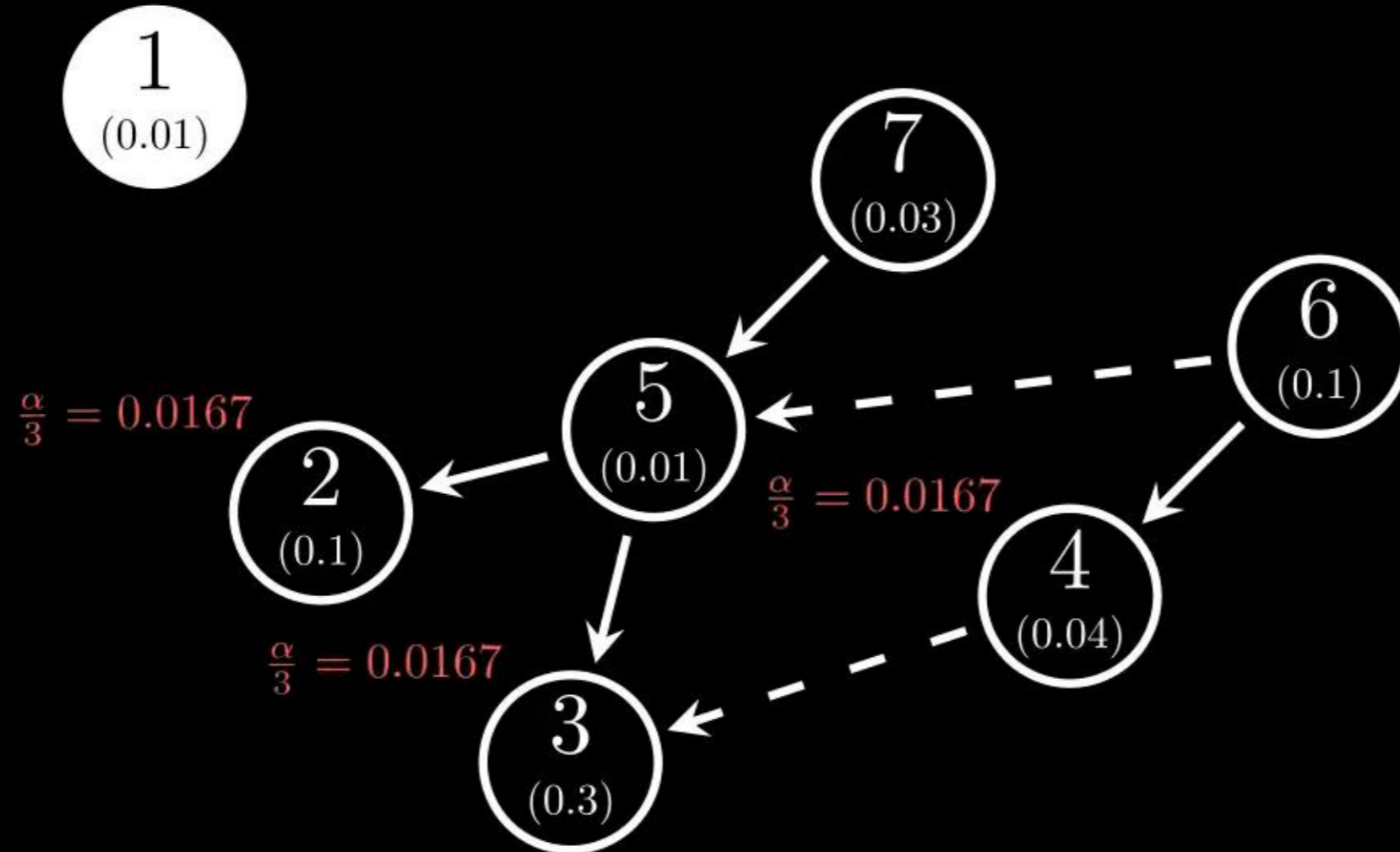
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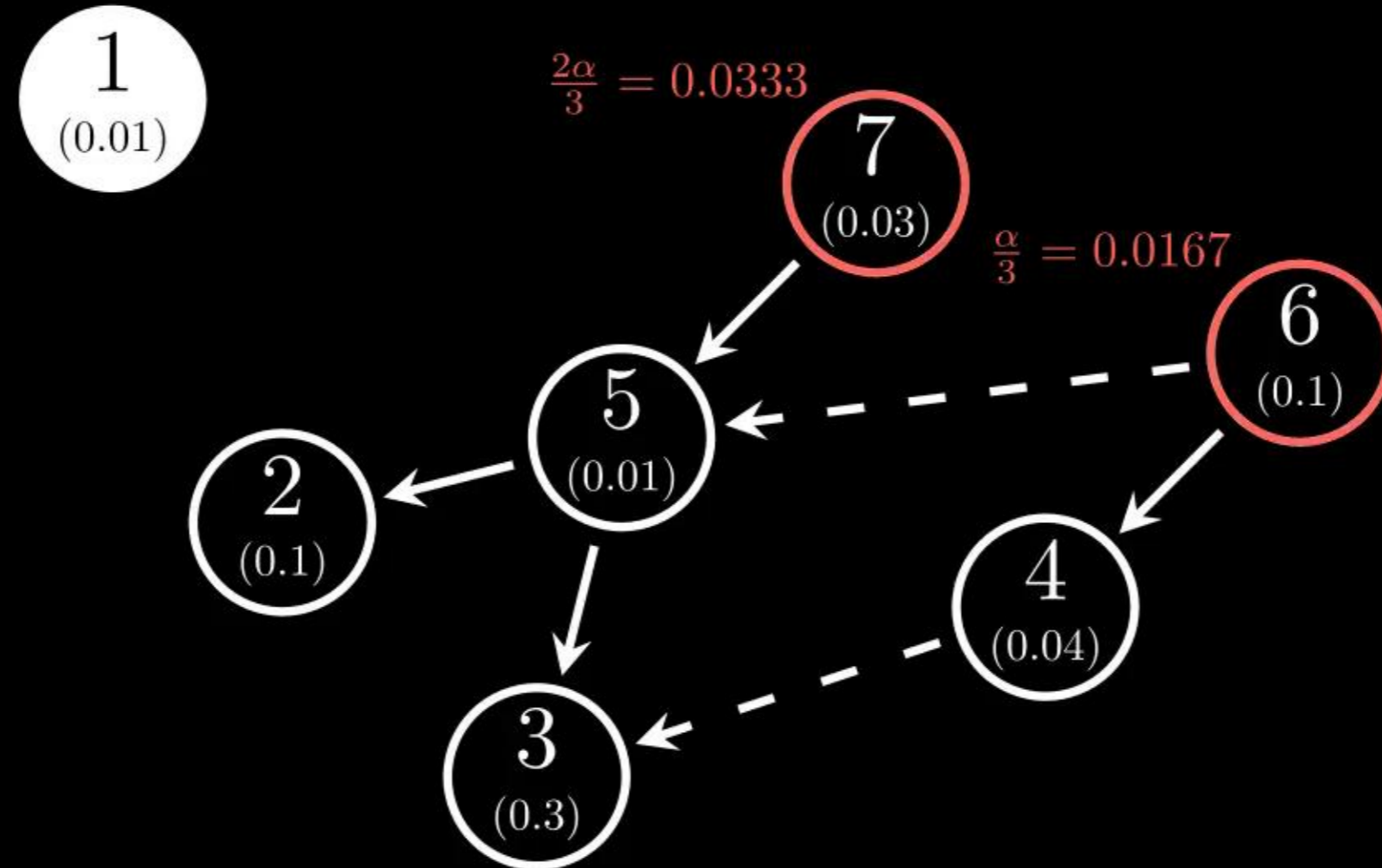
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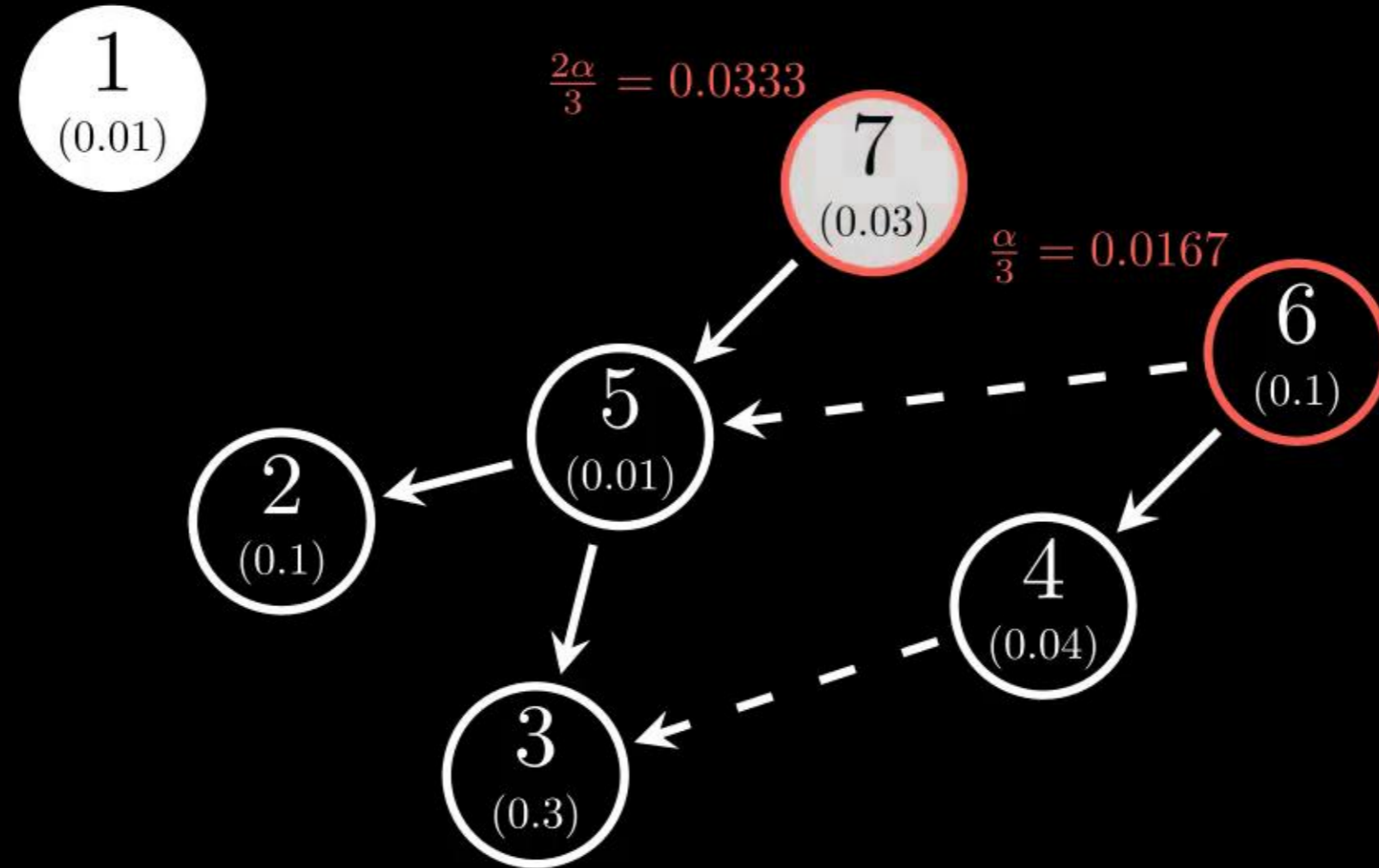
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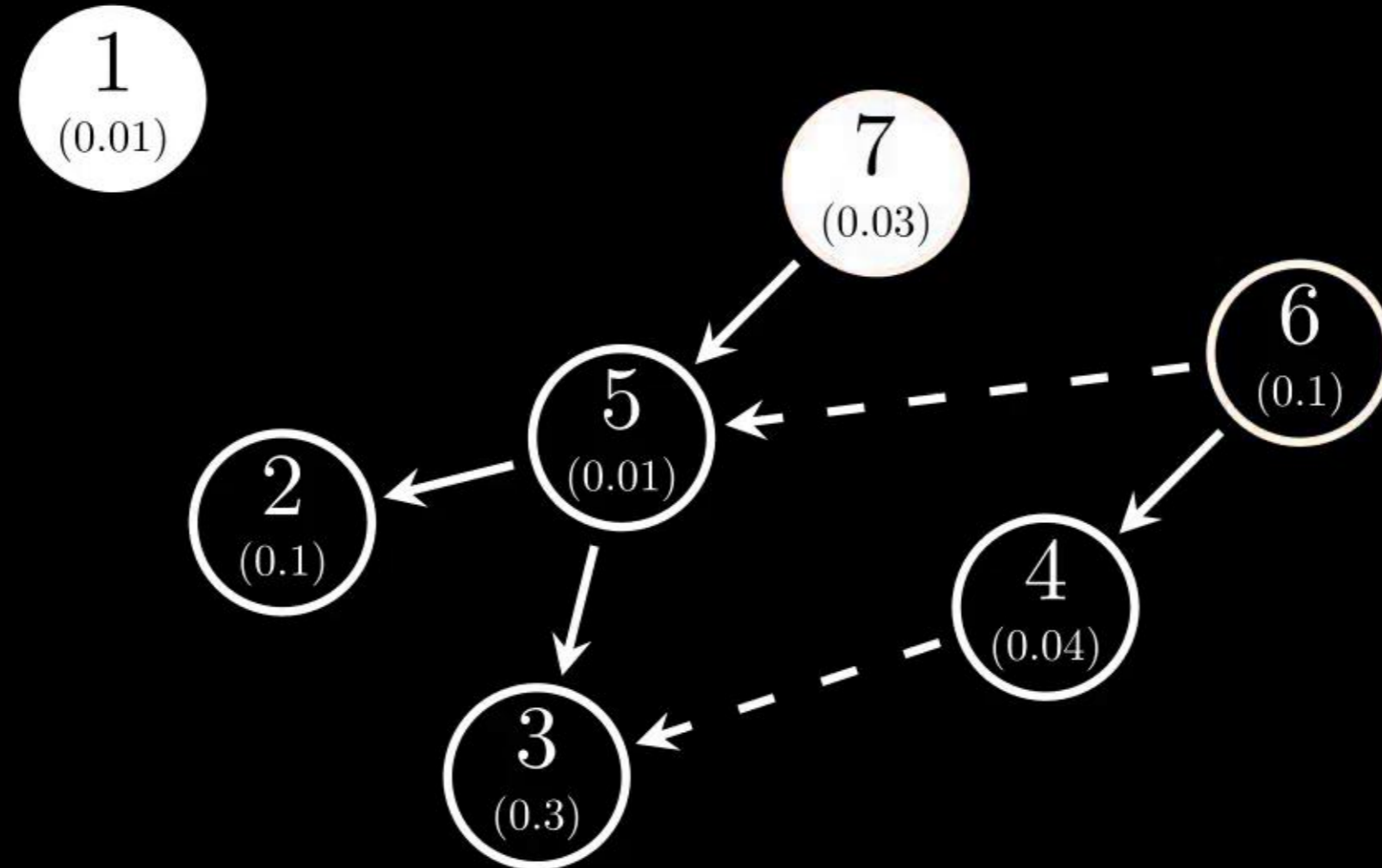
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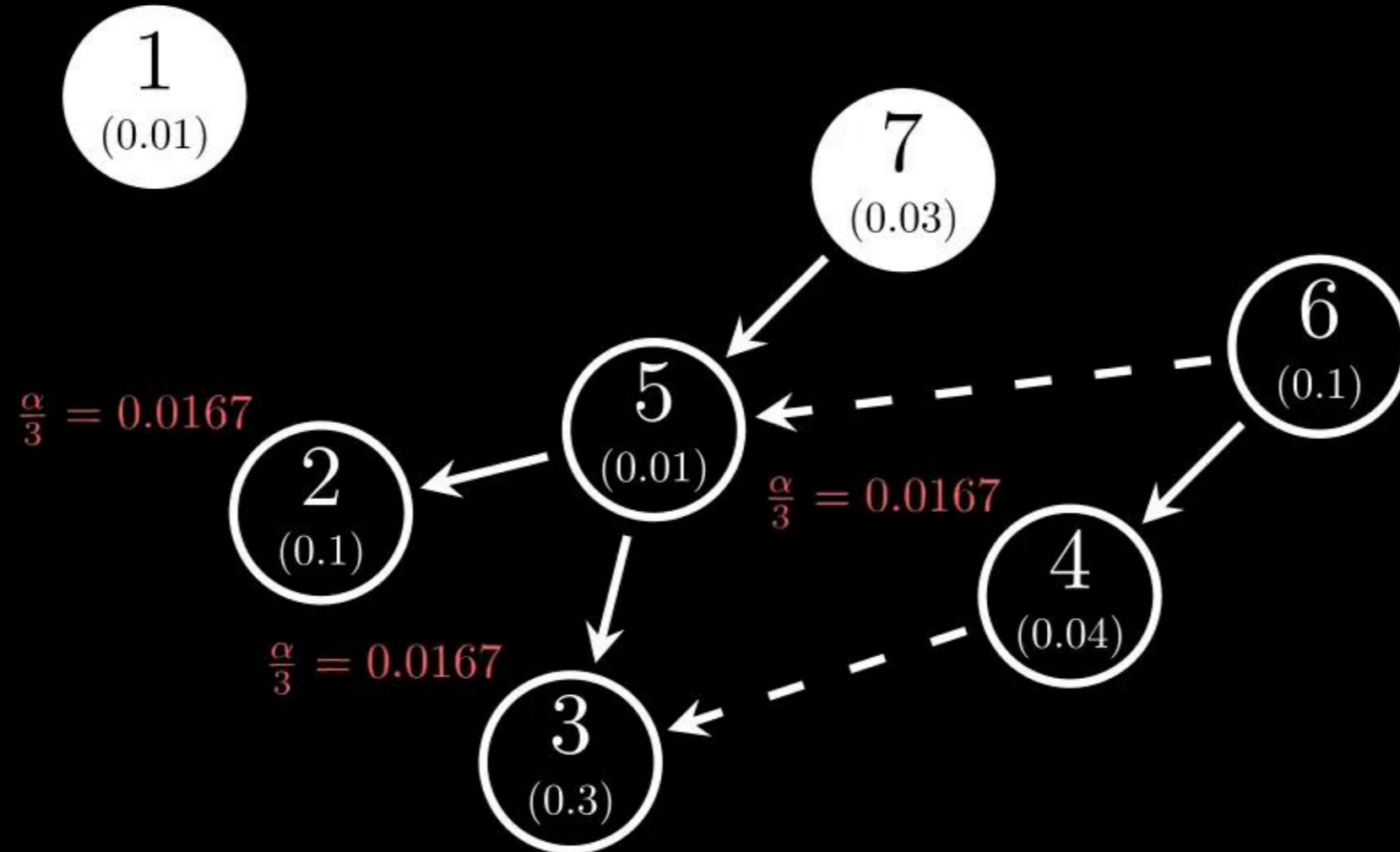
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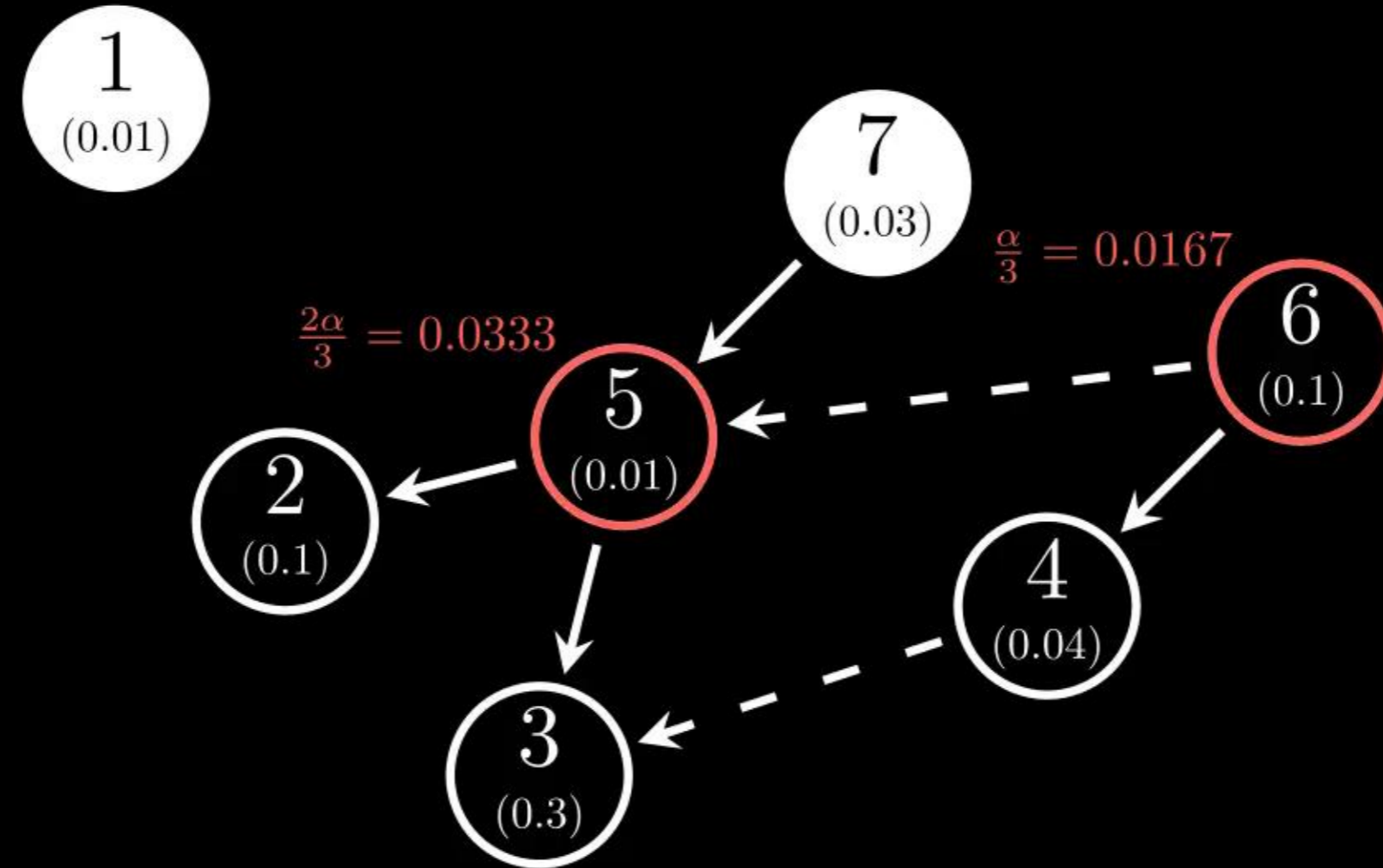
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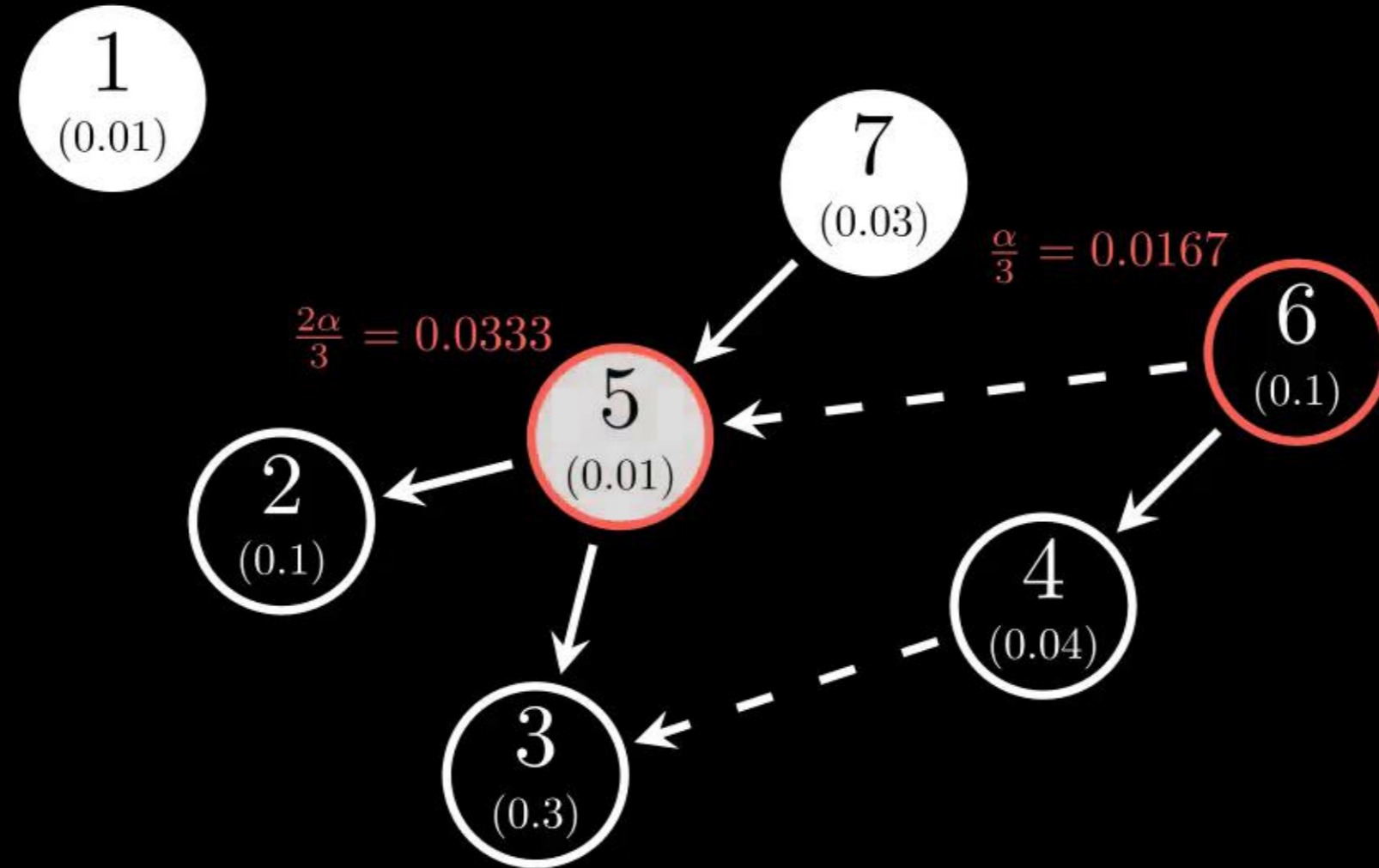
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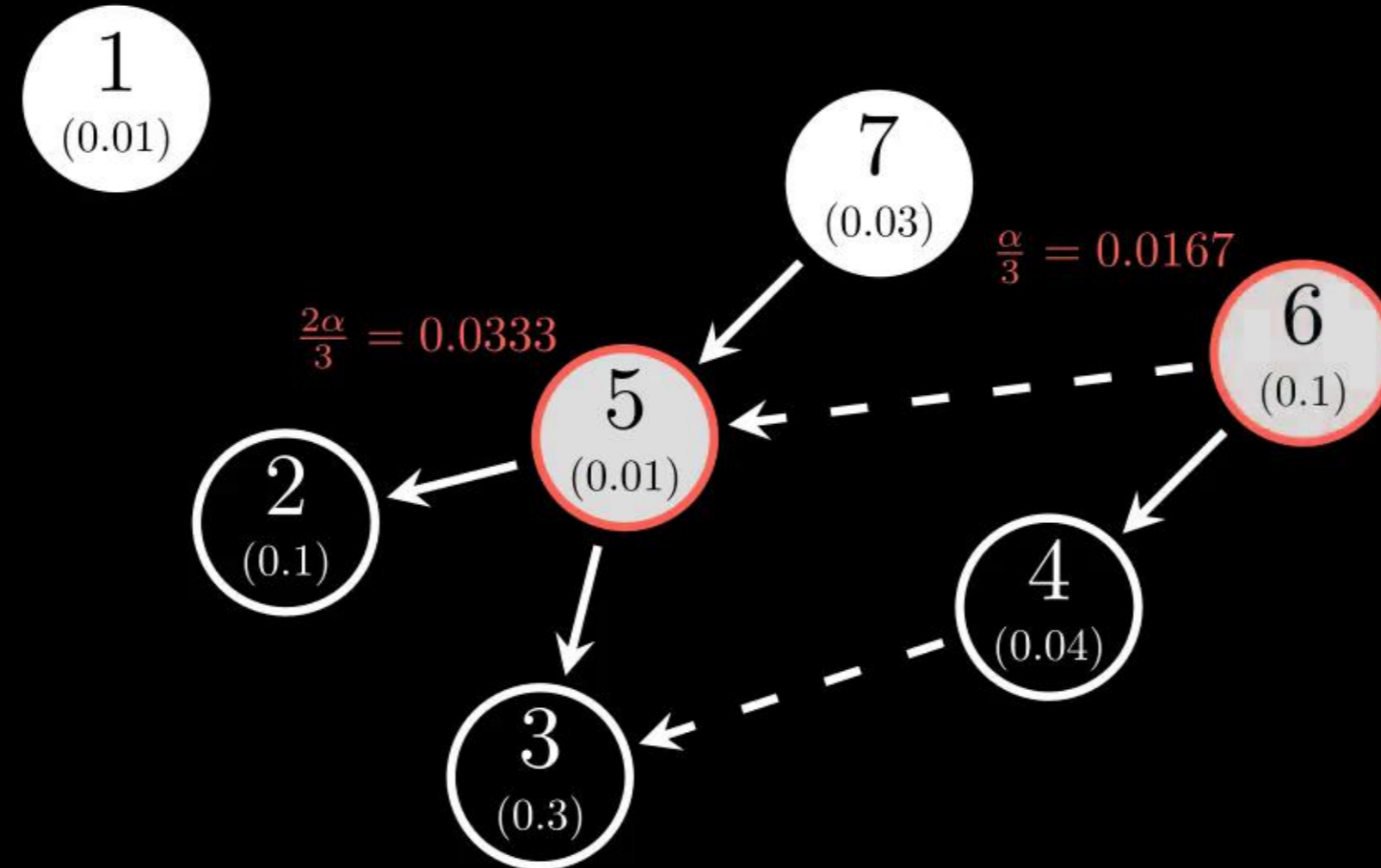
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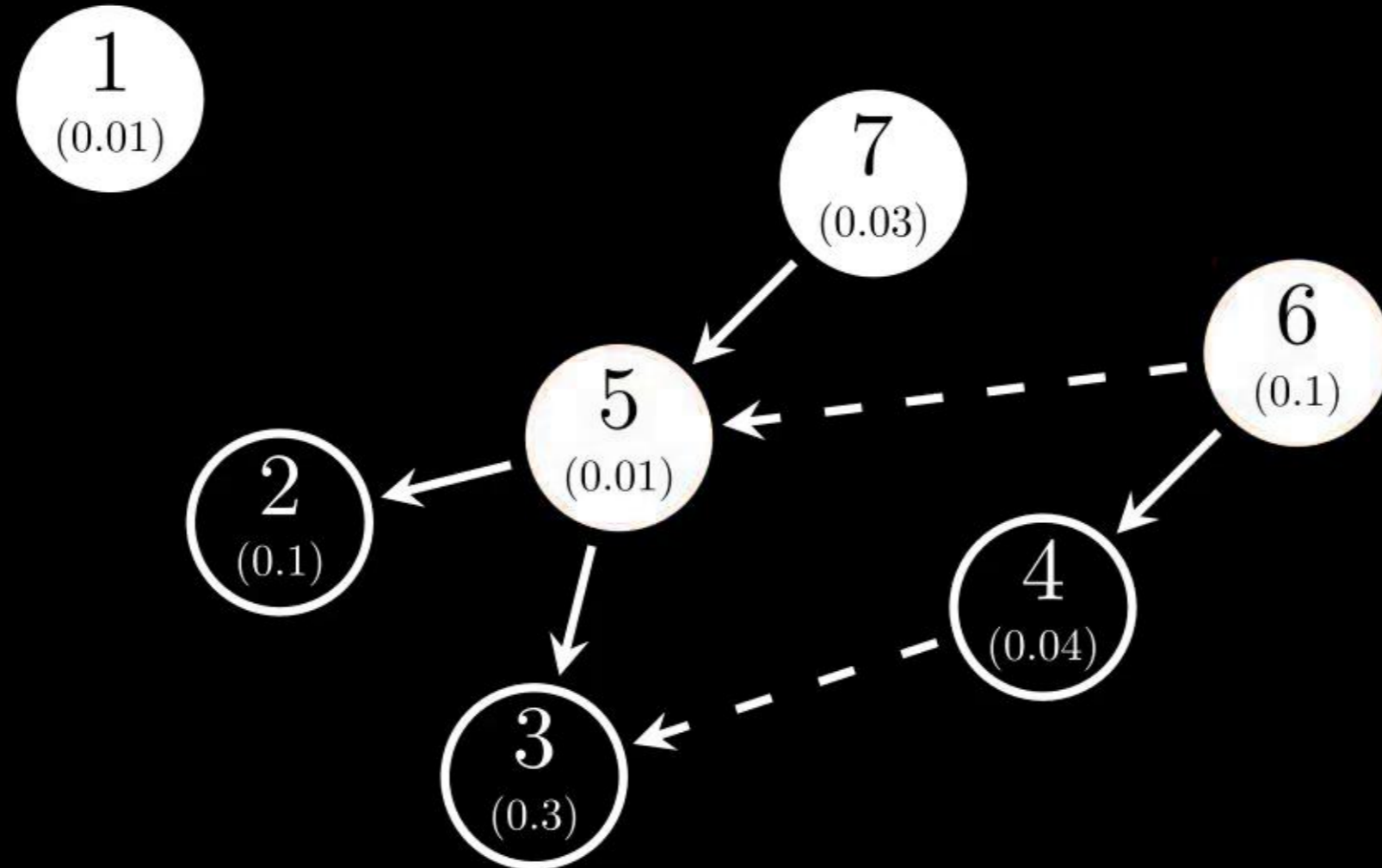
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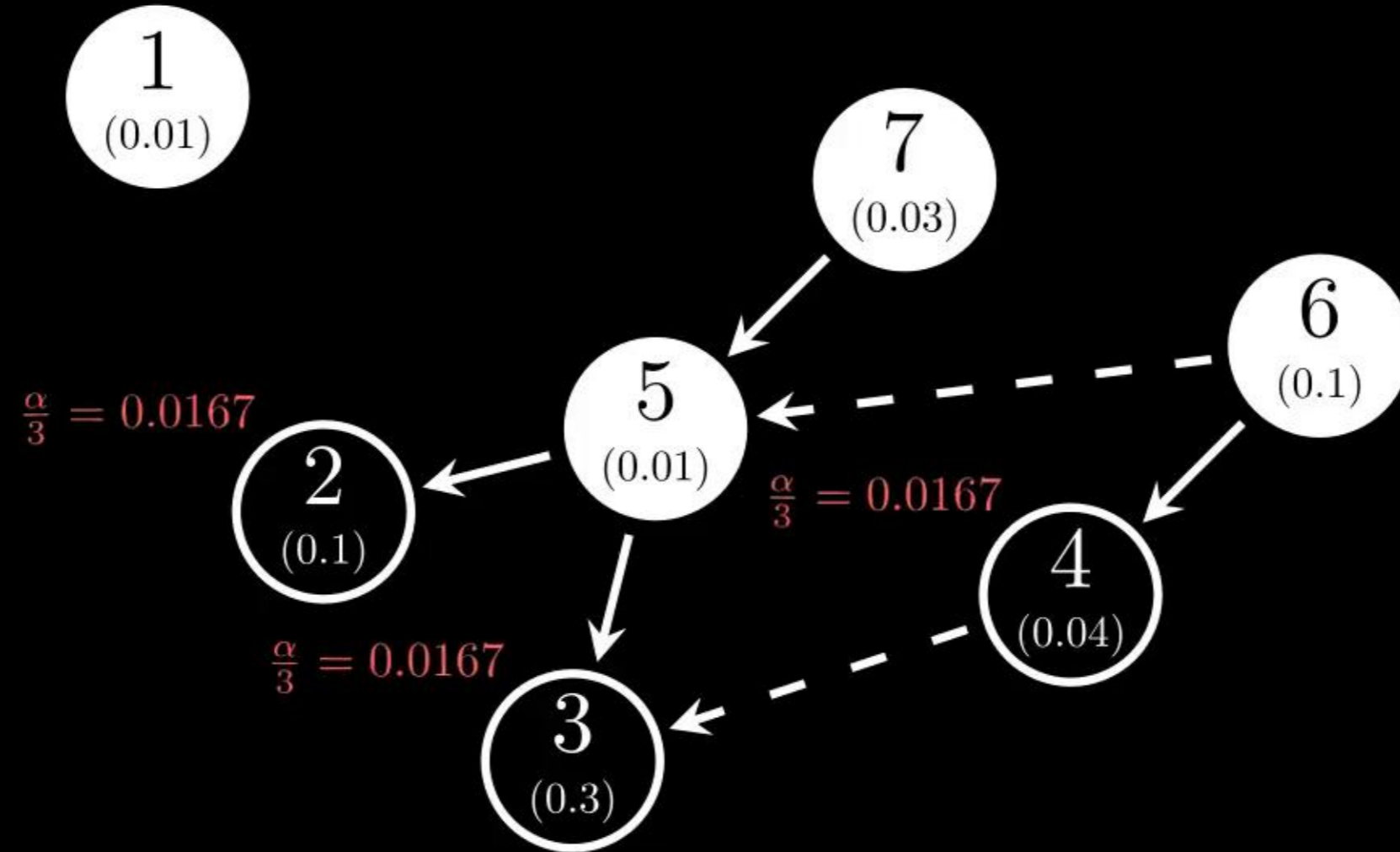
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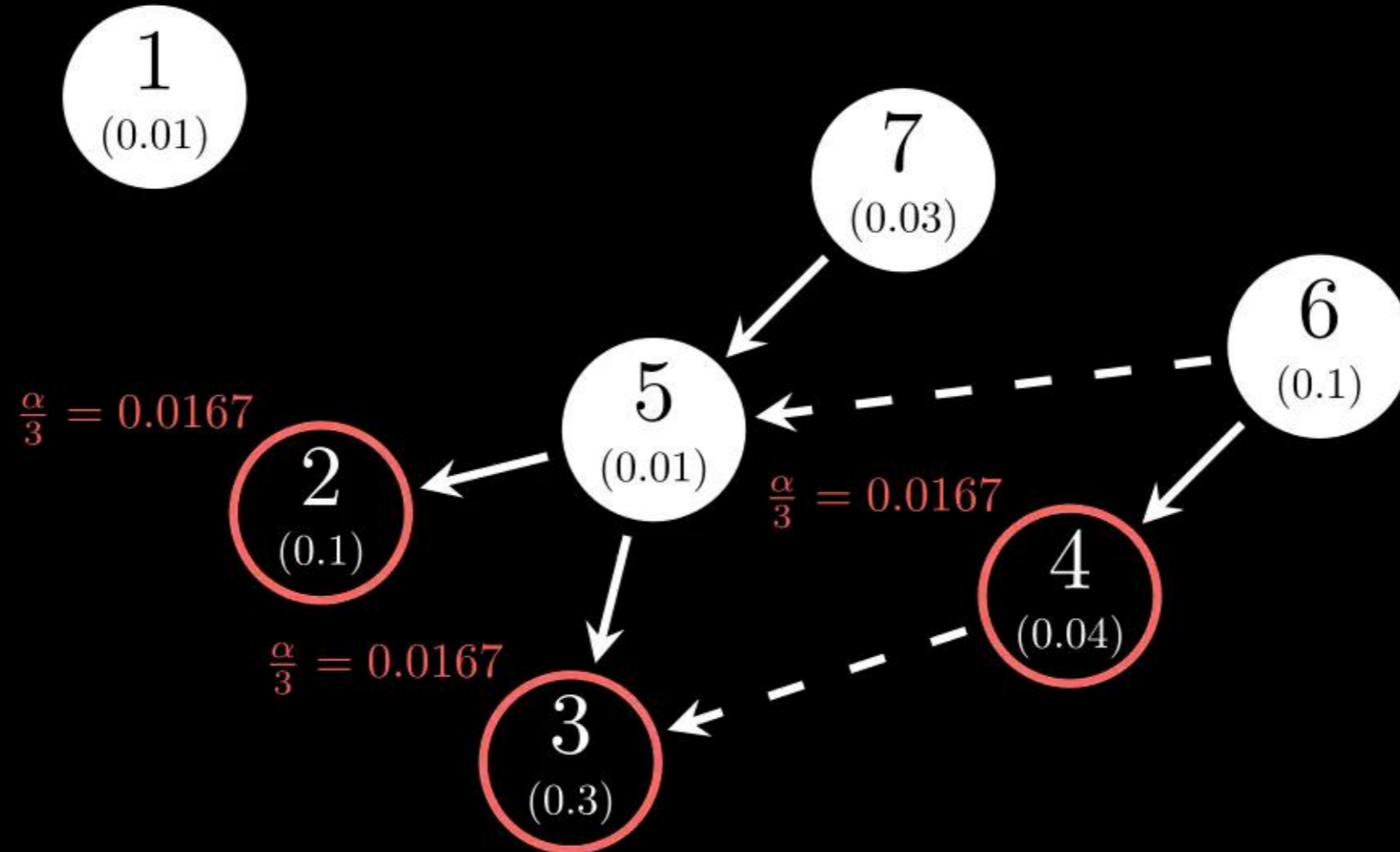
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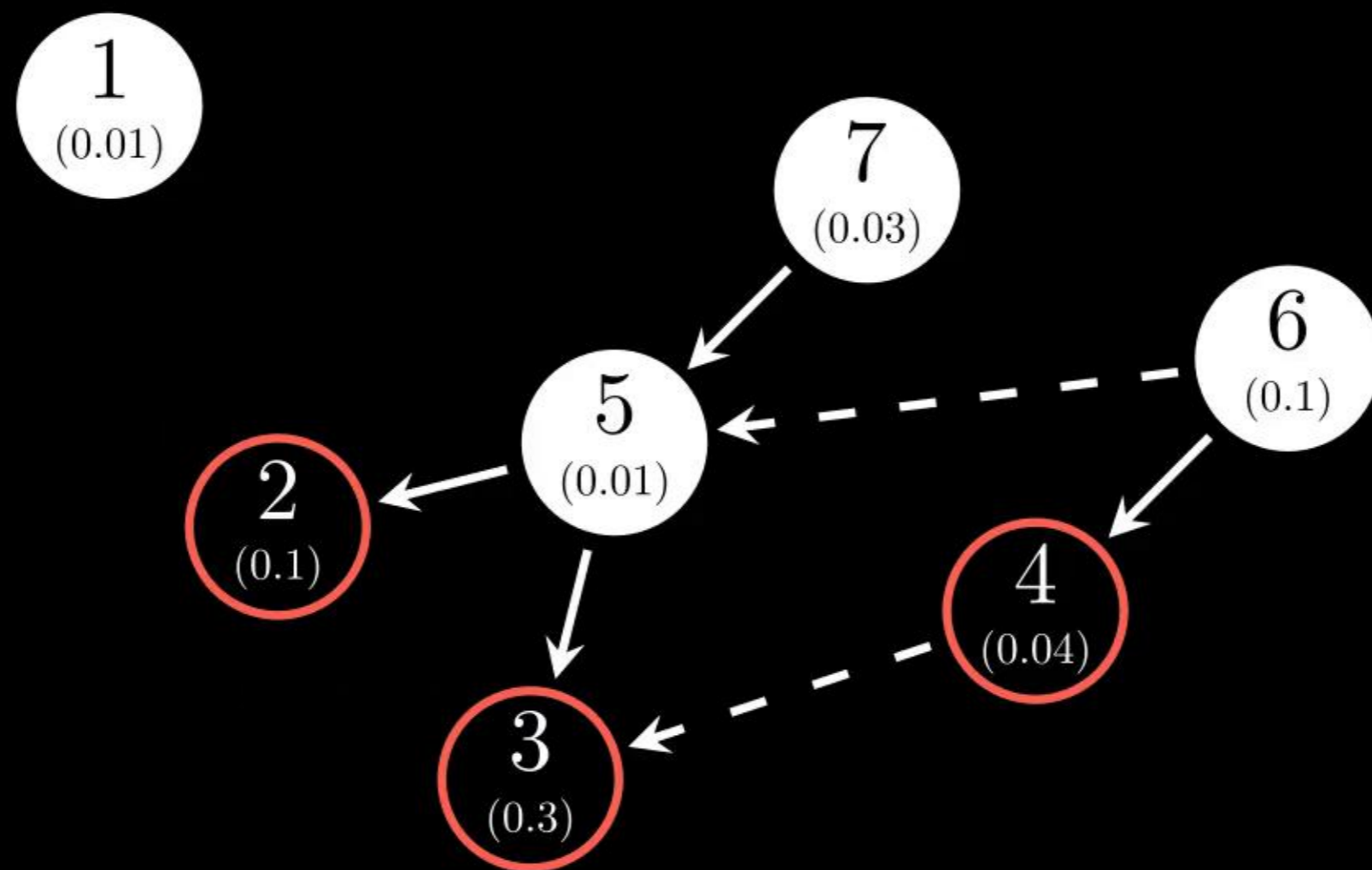
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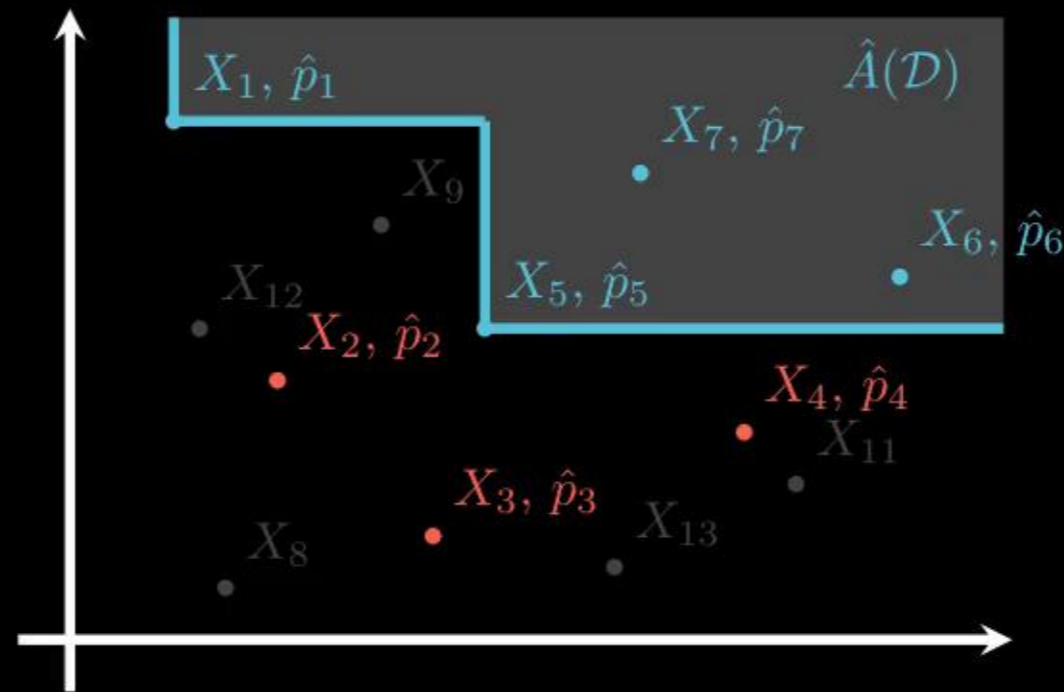
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Here: $\alpha = 0.05$. The procedure terminates with $\mathcal{R}_\alpha = \{1, 5, 6, 7\}$.

High-level strategy

For $x_0 \in \mathbb{R}^d$, define null hypothesis $H_0(x_0) := \{P \in \mathcal{P}_{\text{Mon},d}(\sigma) : \eta(x_0) < \tau\}$.

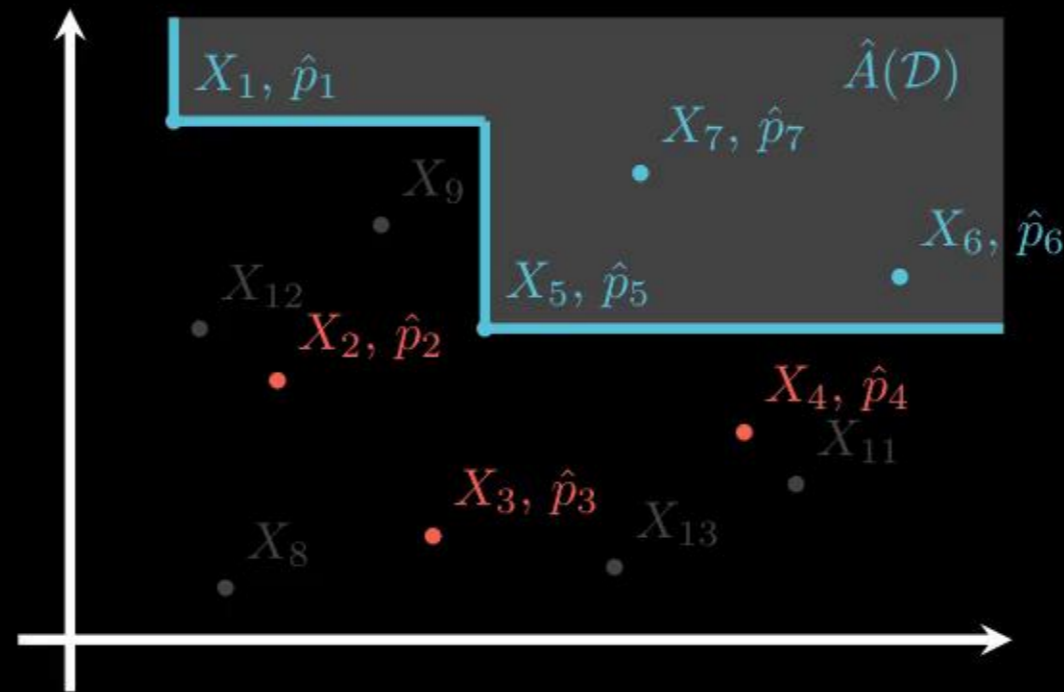


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Theorem. For any $n \geq 1$, $m \in [n]$, $\alpha \in (0, 1)$, $\sigma > 0$, we have

$$\inf_{P \in \mathcal{P}_{\text{Mon}, d}(\sigma)} \mathbb{P}(\hat{A}^{\text{ISS}} \subseteq \mathcal{X}_{\tau}(\eta) \mid X_1, \dots, X_n) \geq 1 - \alpha.$$

Further conditions are needed for power

Let $\hat{\mathcal{A}}_n(\tau, \alpha, \mathcal{P})$ denote the set of *data-dependent selection sets* controlling Type I error over \mathcal{P} . Recall $R_\tau(\hat{A}) := \mathbb{E}\{\mu(\mathcal{X}_\tau(\eta) \setminus \hat{A})\}$.

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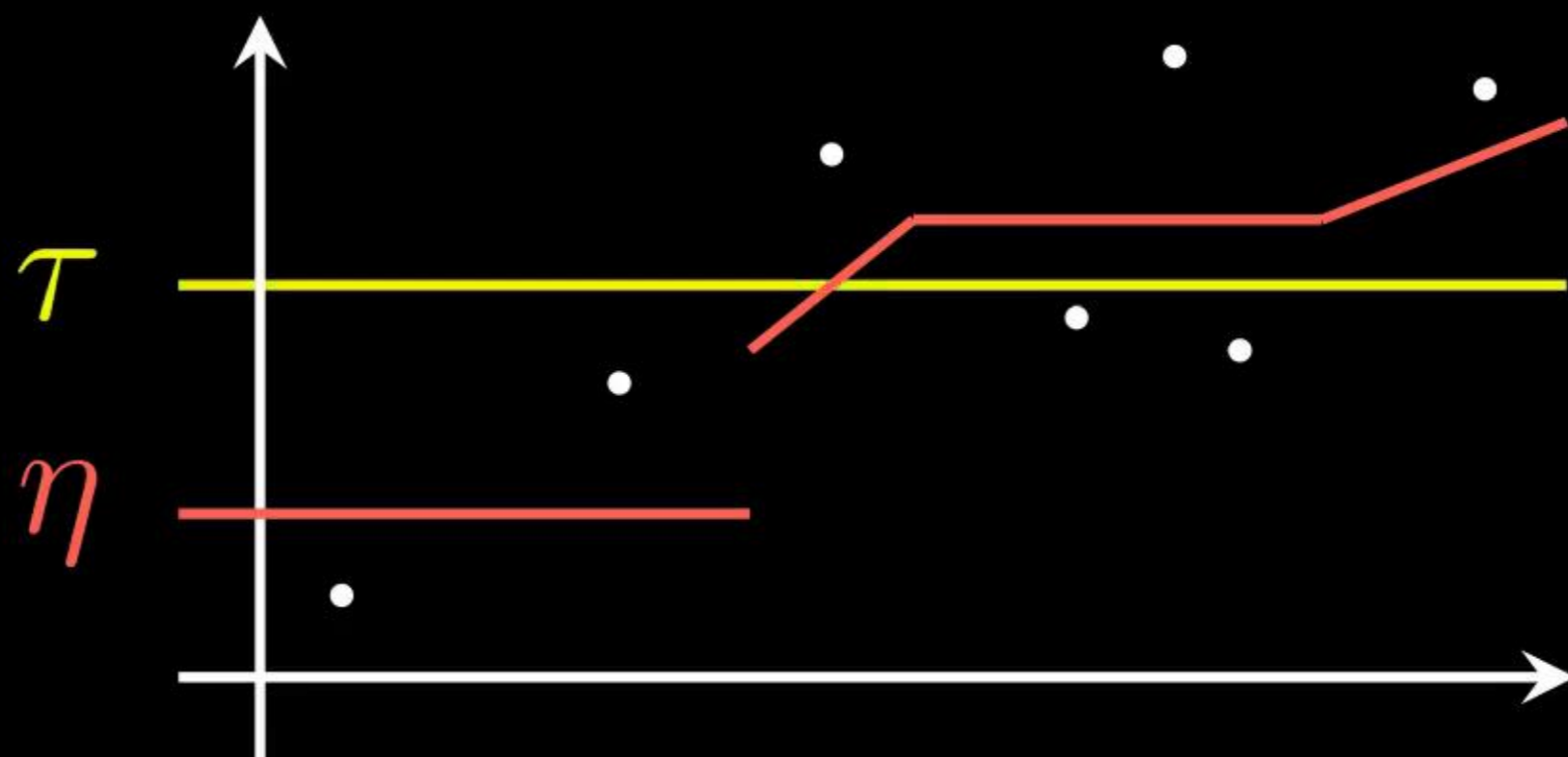
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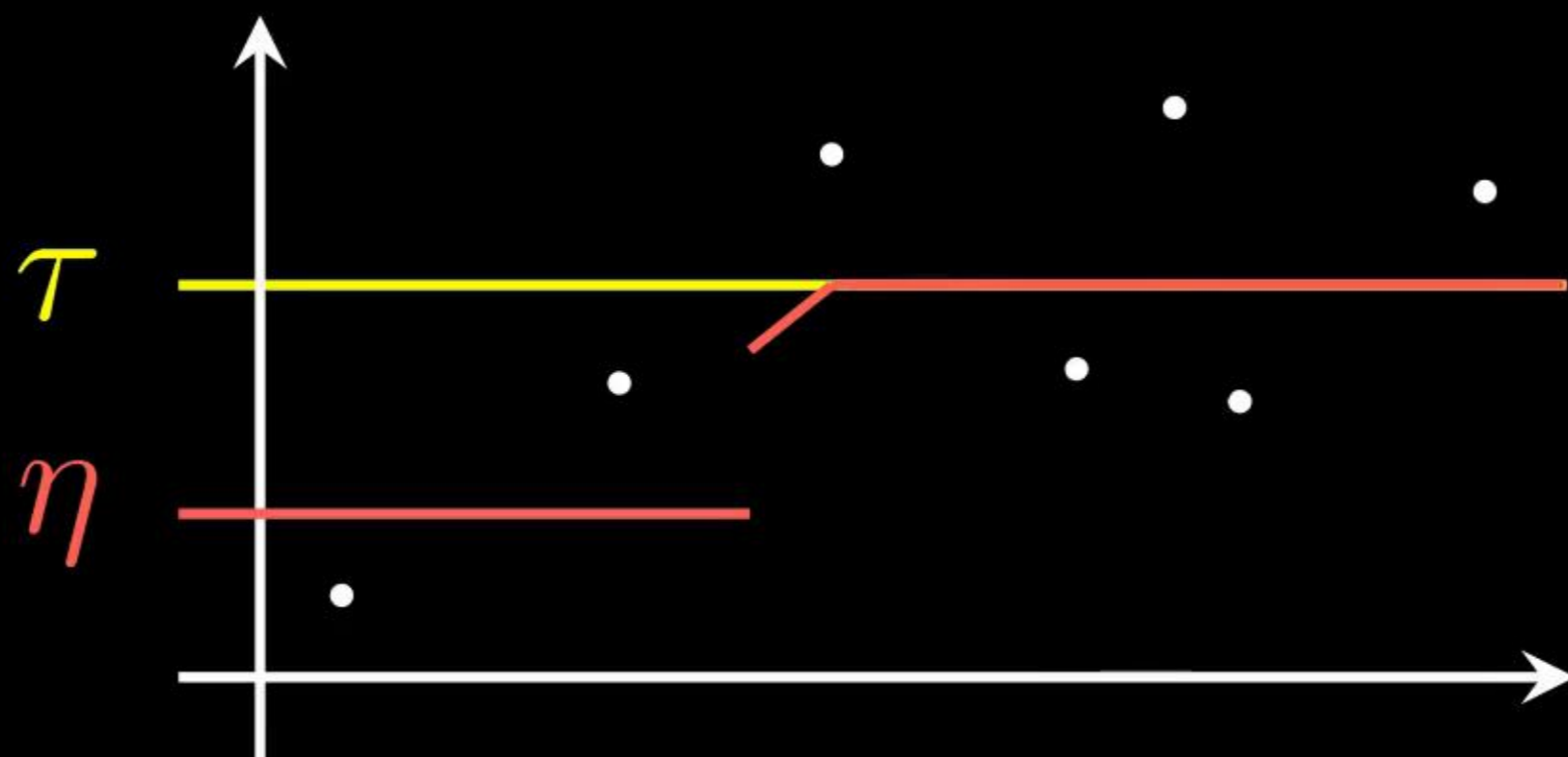


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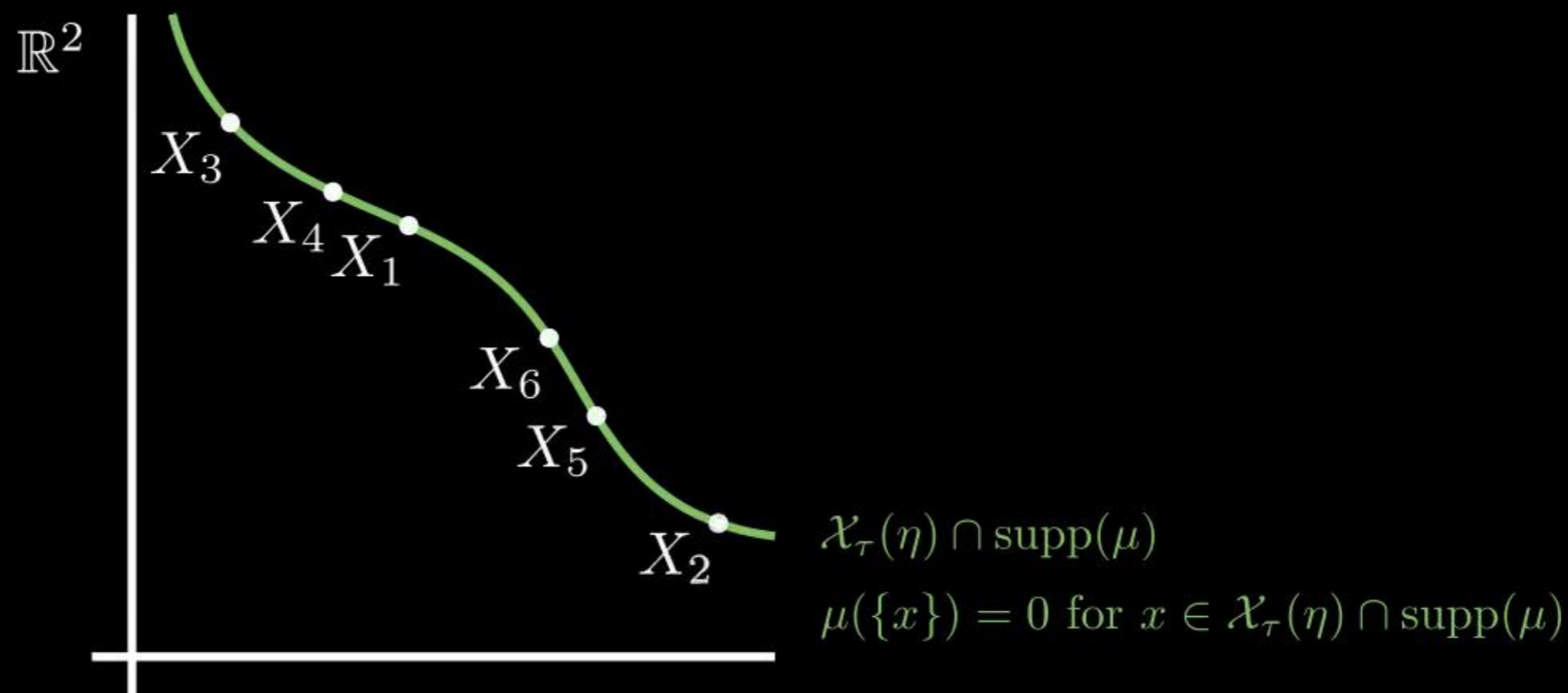
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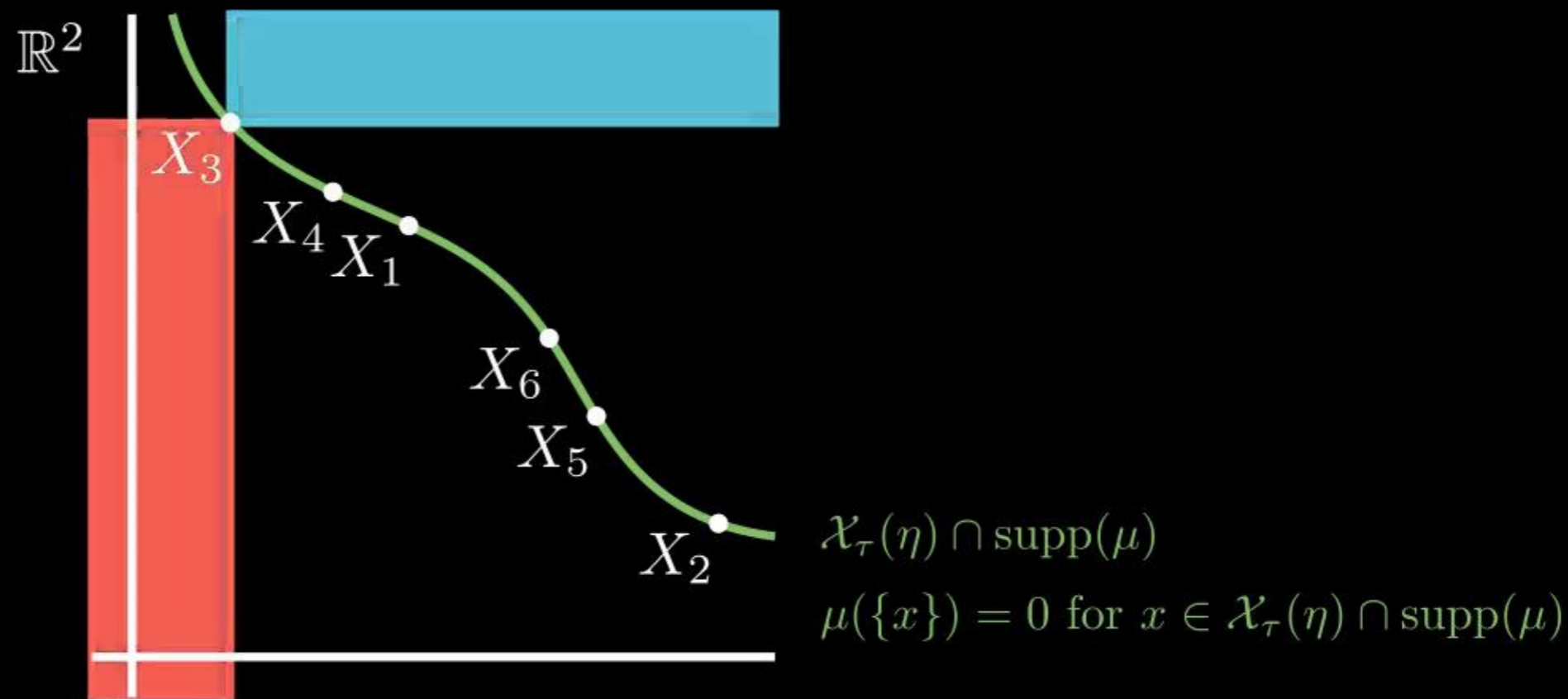


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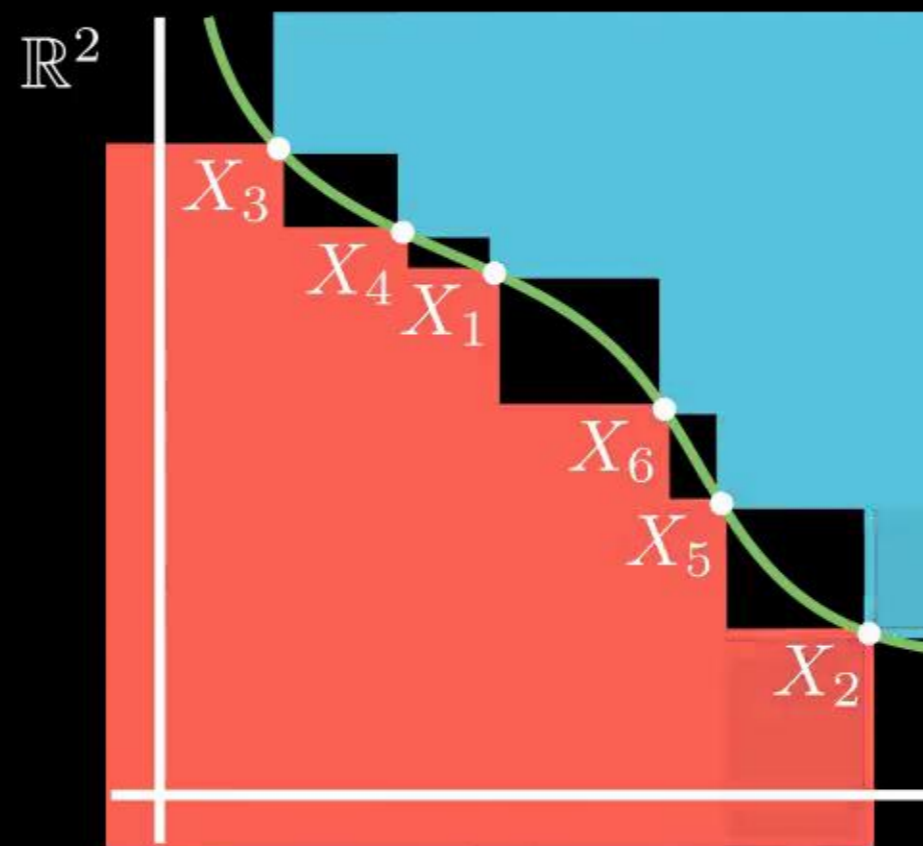


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$\mathcal{X}_\tau(\eta) \cap \text{supp}(\mu)$

$\mu(\{x\}) = 0$ for $x \in \mathcal{X}_\tau(\eta) \cap \text{supp}(\mu)$

Power

Definition. Let $\sigma, \gamma, \lambda > 0$ and $\theta > 1$. $\mathcal{P}_{\text{MonReg}} \equiv \mathcal{P}_{\text{MonReg},d}(\sigma, \tau, \gamma, \lambda, \theta)$ denotes the set of distributions $P \in \mathcal{P}_{\text{Mon},d}(\sigma)$ for which additionally

1. $\eta(x + r(1, \dots, 1)^\top) \geq \tau + \lambda \cdot r^\gamma$ for all $x \in \mathcal{X}_\tau(\eta) \cap \text{supp}(\mu)$ and $r \in (0, 1]$;
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Theorem. $\exists C(d, \theta) \geq 1$, such that for any $n \geq 1$, $m \in [n]$ and $\alpha \in (0, 1)$,

$$\sup_{P \in \mathcal{P}_{\text{MonReg}}} R_\tau(\hat{A}^{\text{ISS}}) \leq 1 \wedge C(d, \theta) \left\{ \left(\frac{\sigma^2}{n\lambda^2} \log_+ \left(\frac{m \log_+ n}{\alpha} \right) \right)^{1/(2\gamma+d)} + \left(\frac{\log_+ m}{m} \right)^{1/d} \right\},$$

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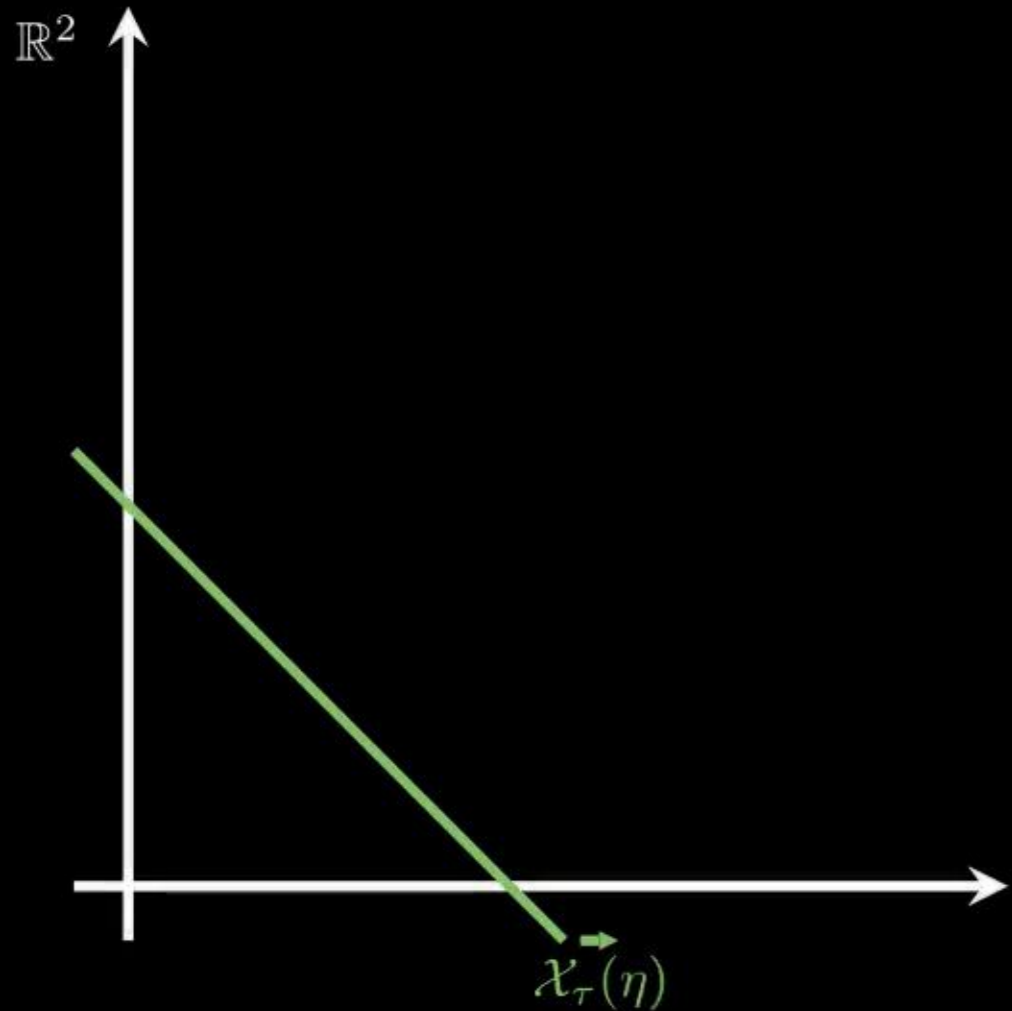
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where $\log_+ x := \log(x \vee e)$ for $x \in \mathbb{R}$.

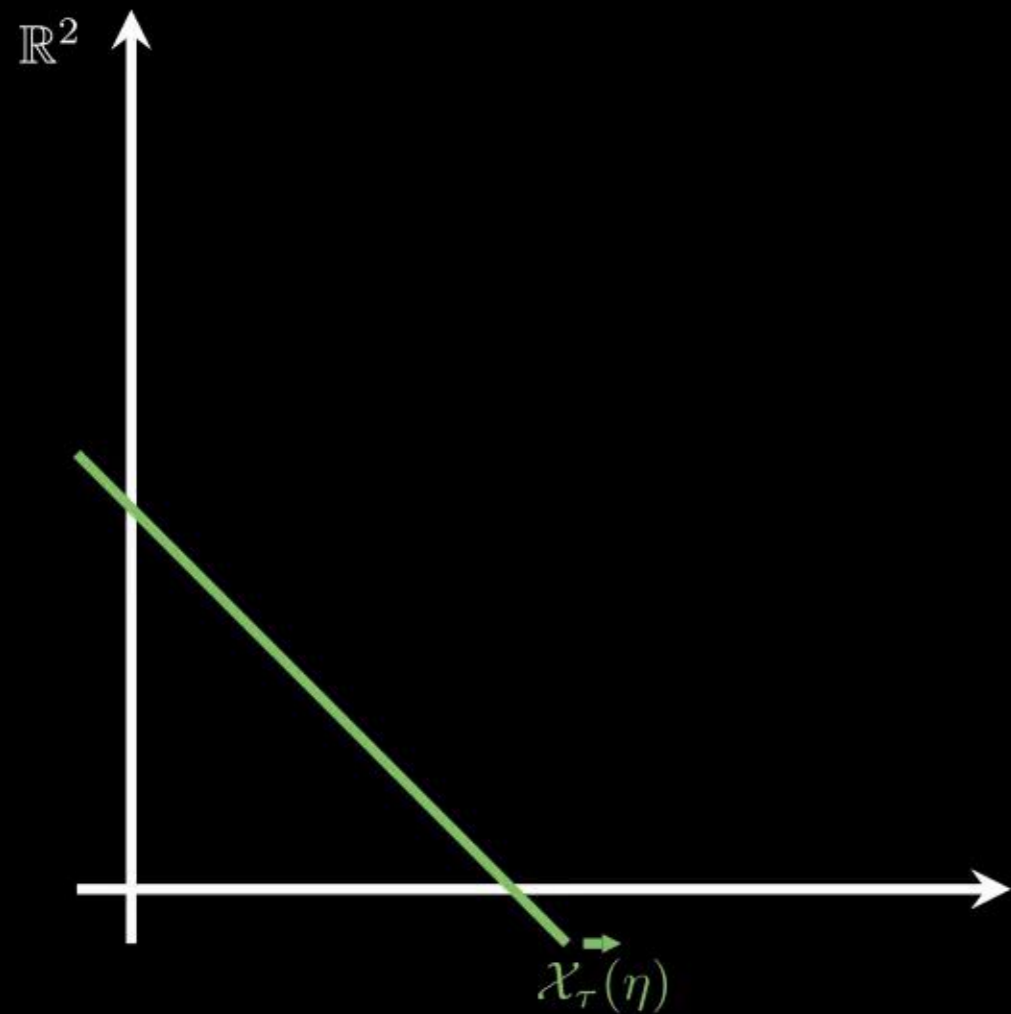


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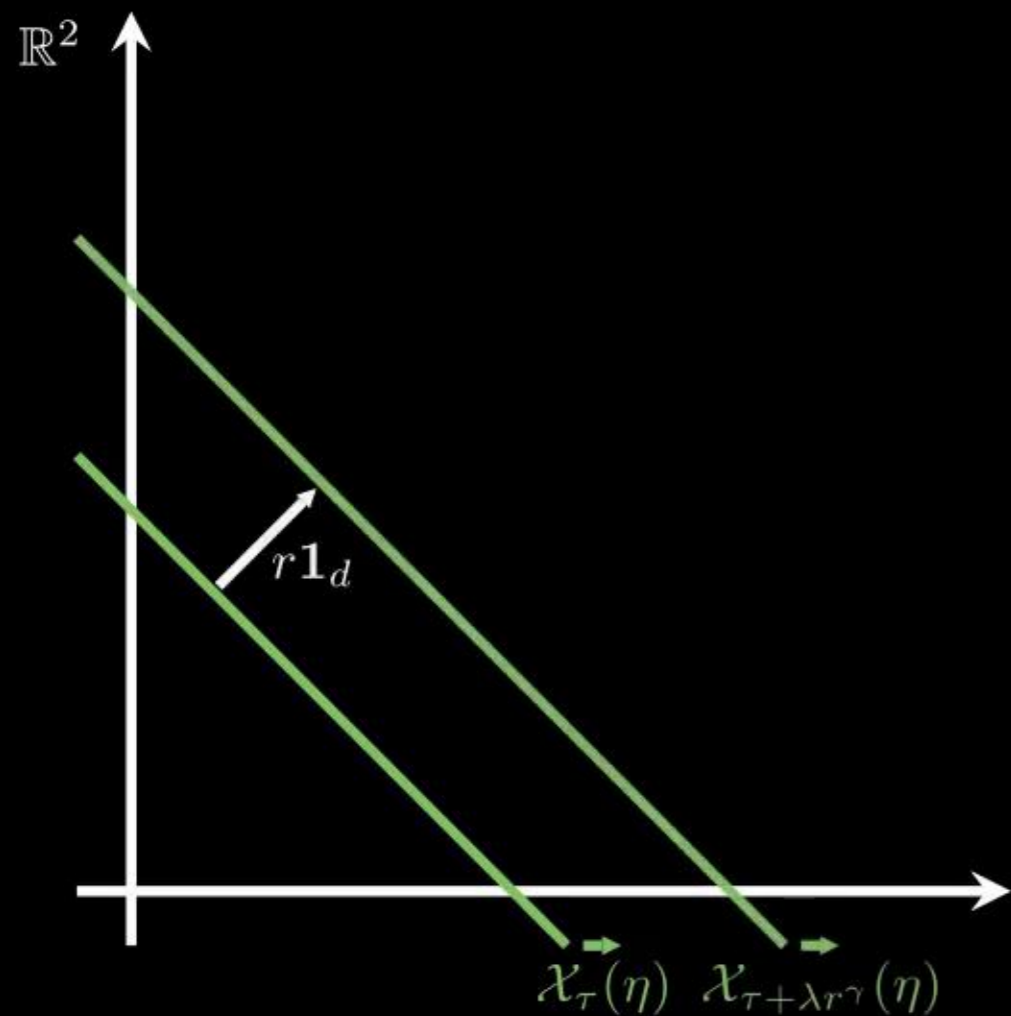
Let $r > 0$ be small.

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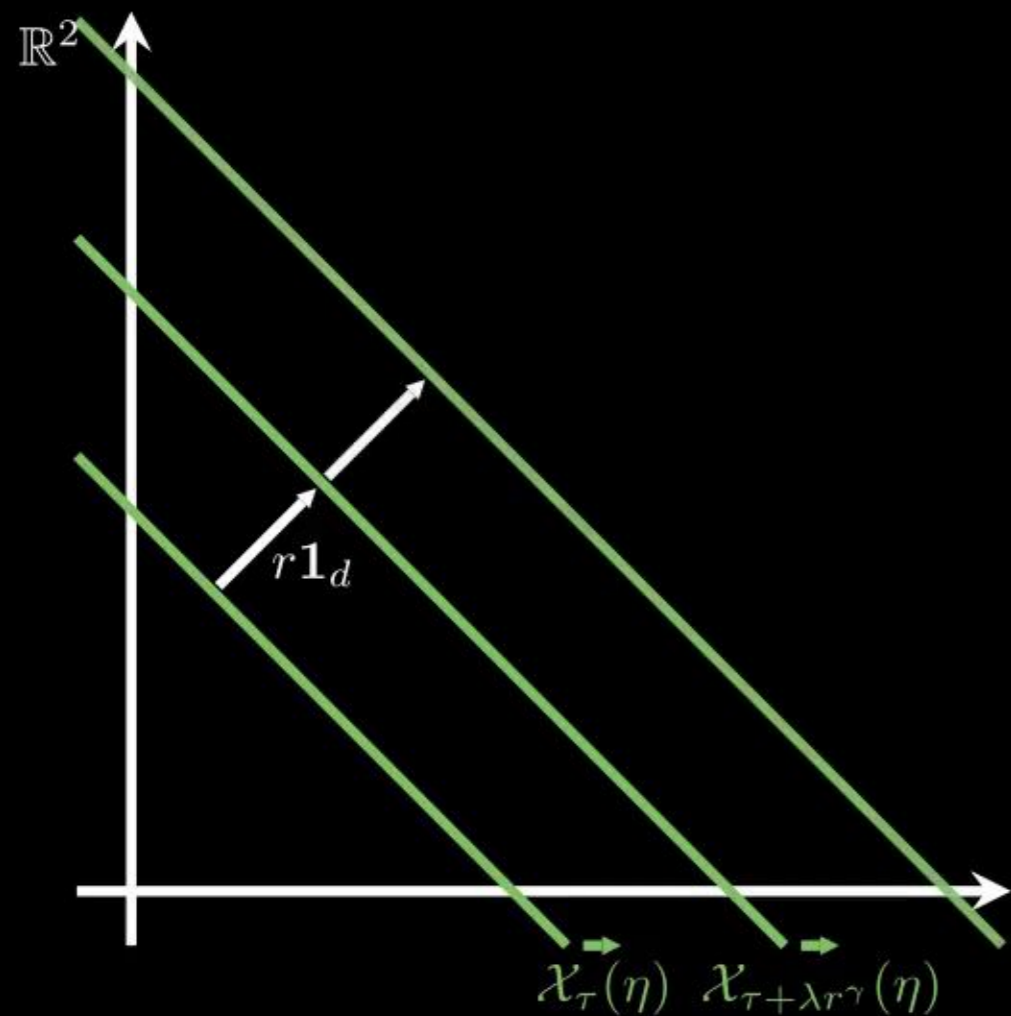
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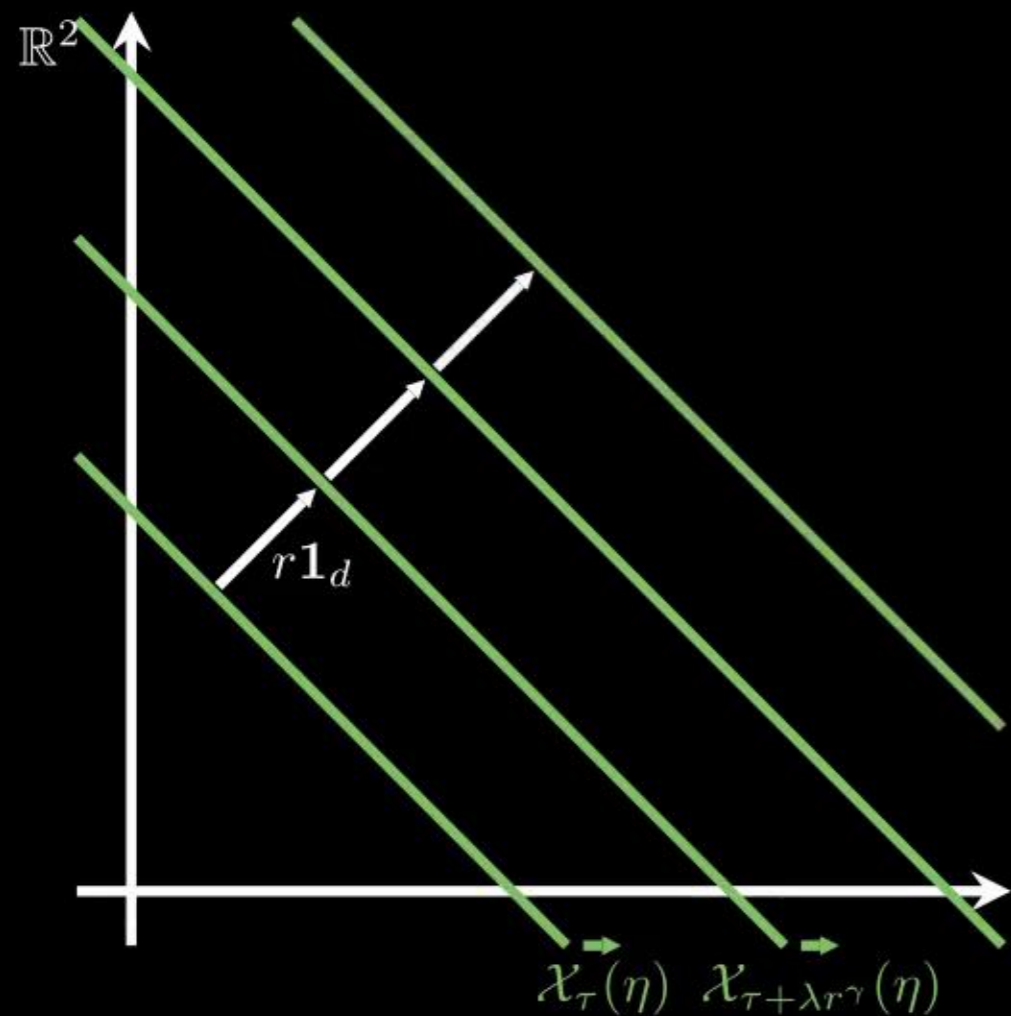
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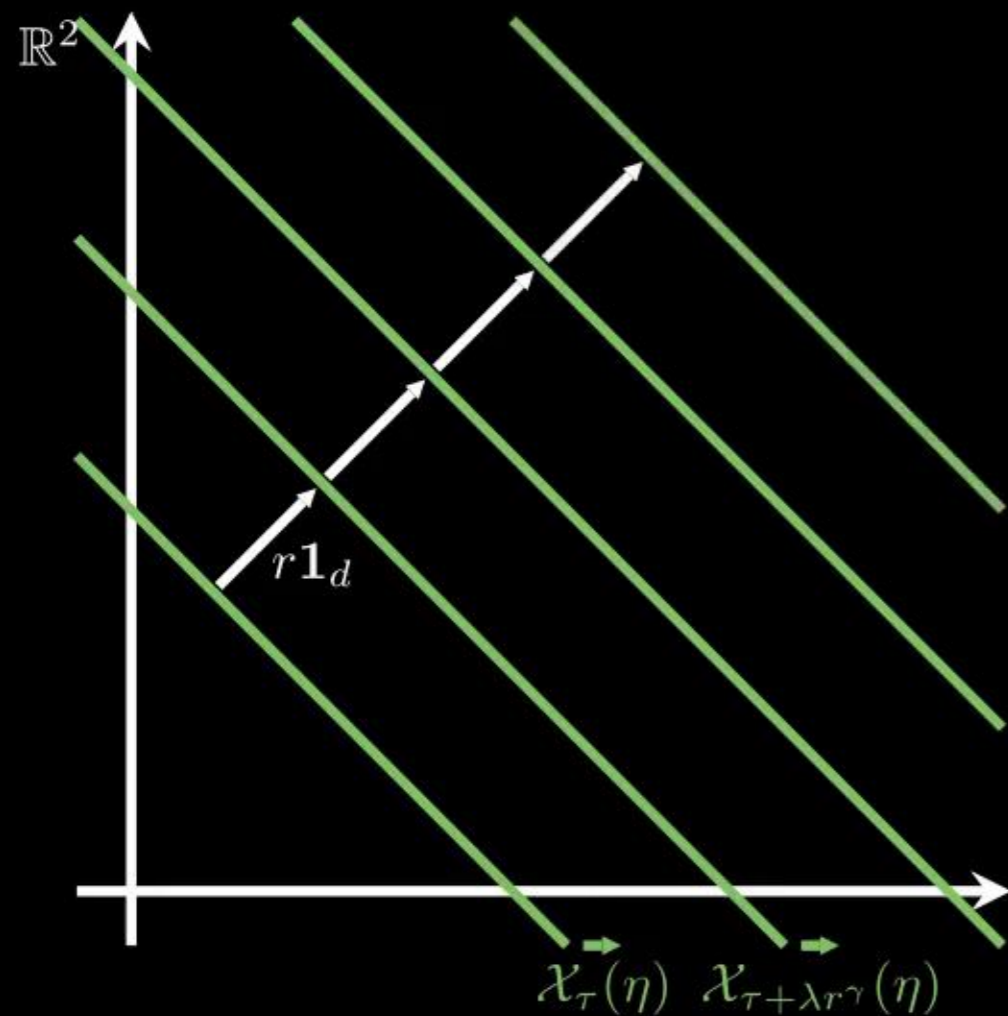
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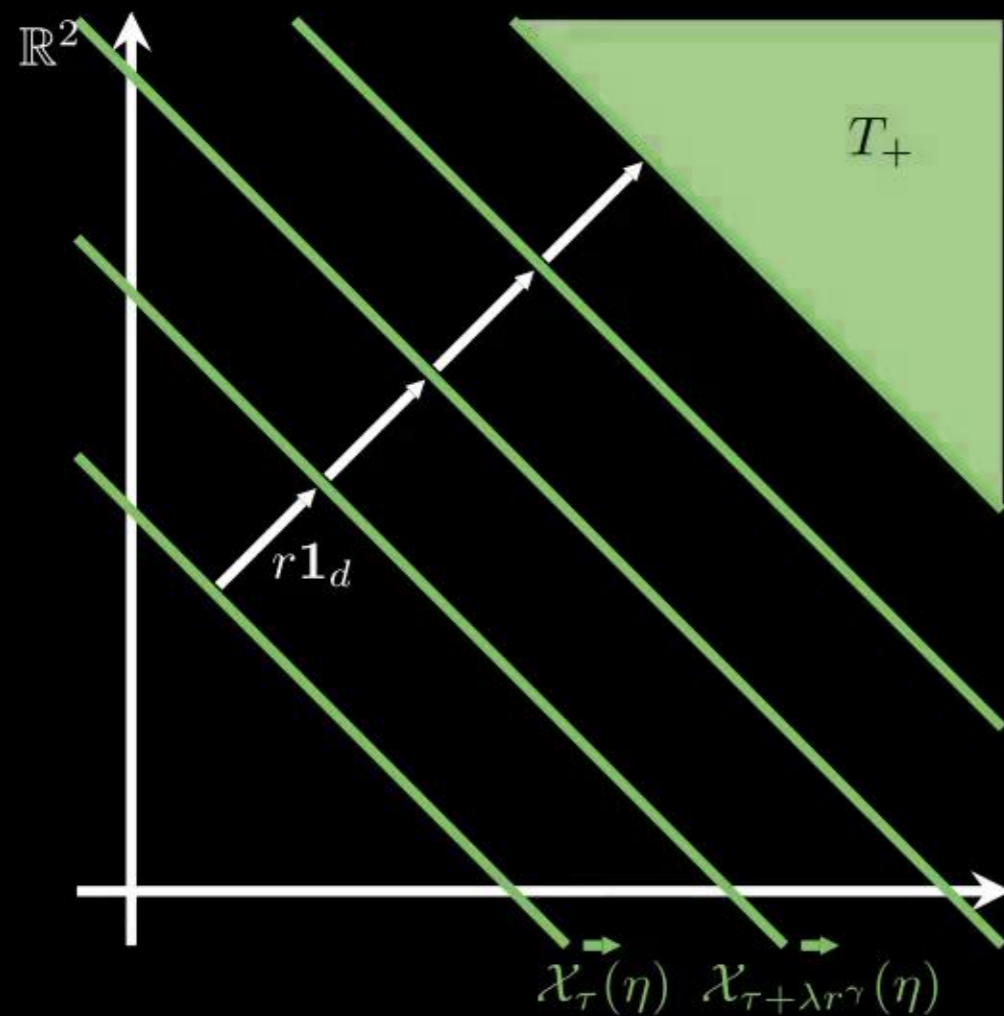
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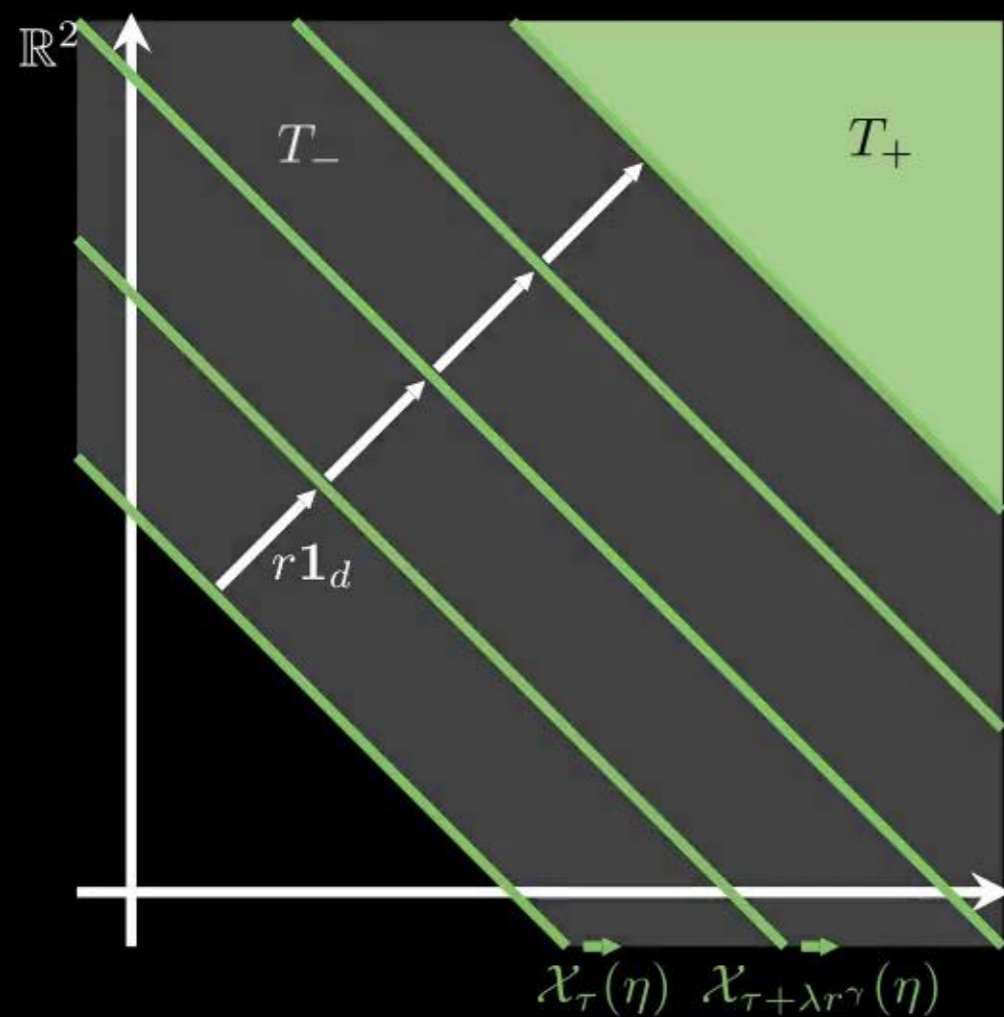
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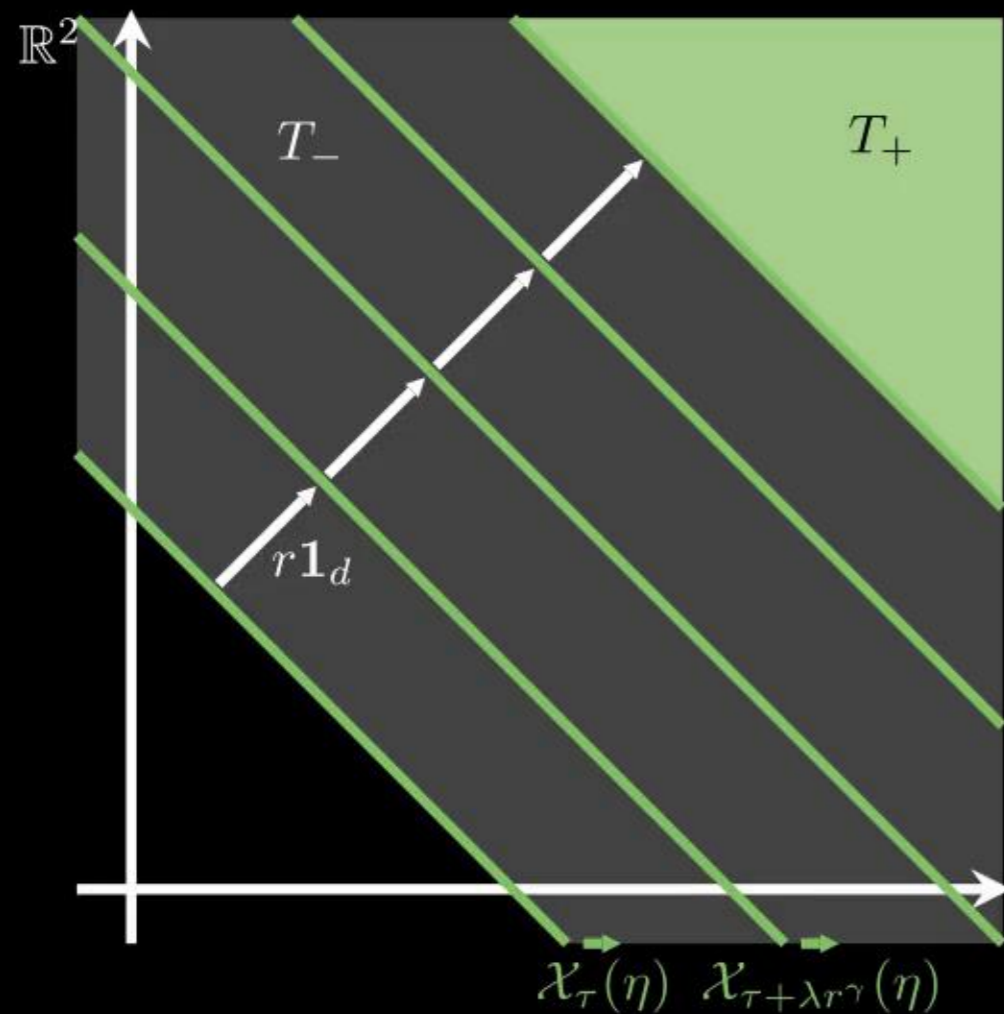
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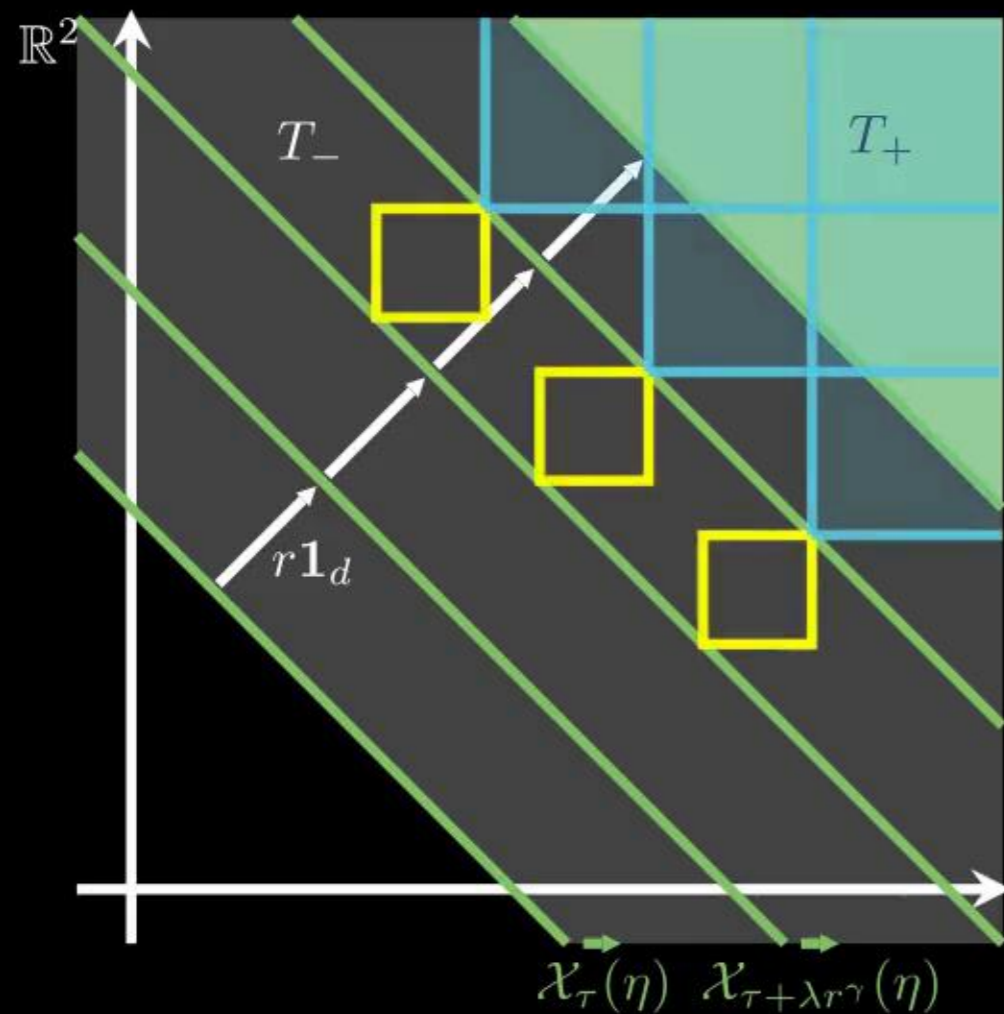
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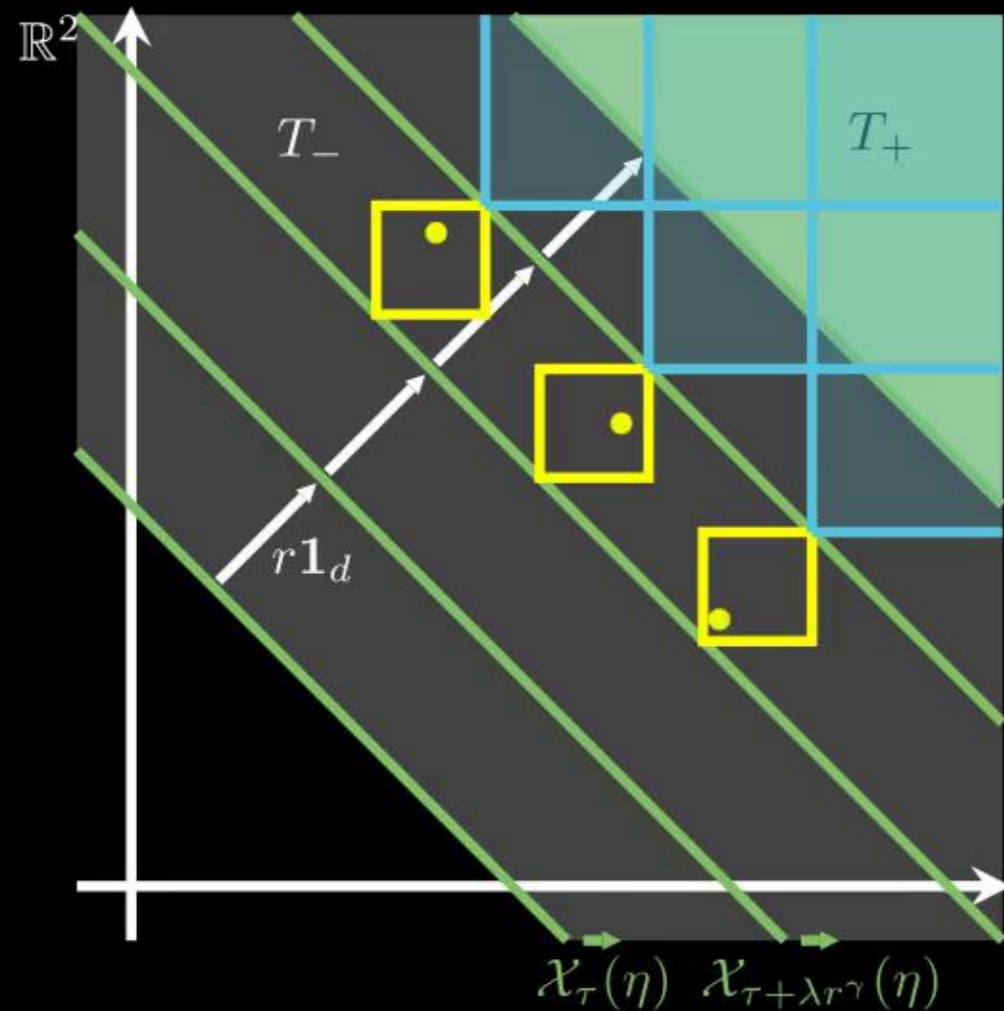
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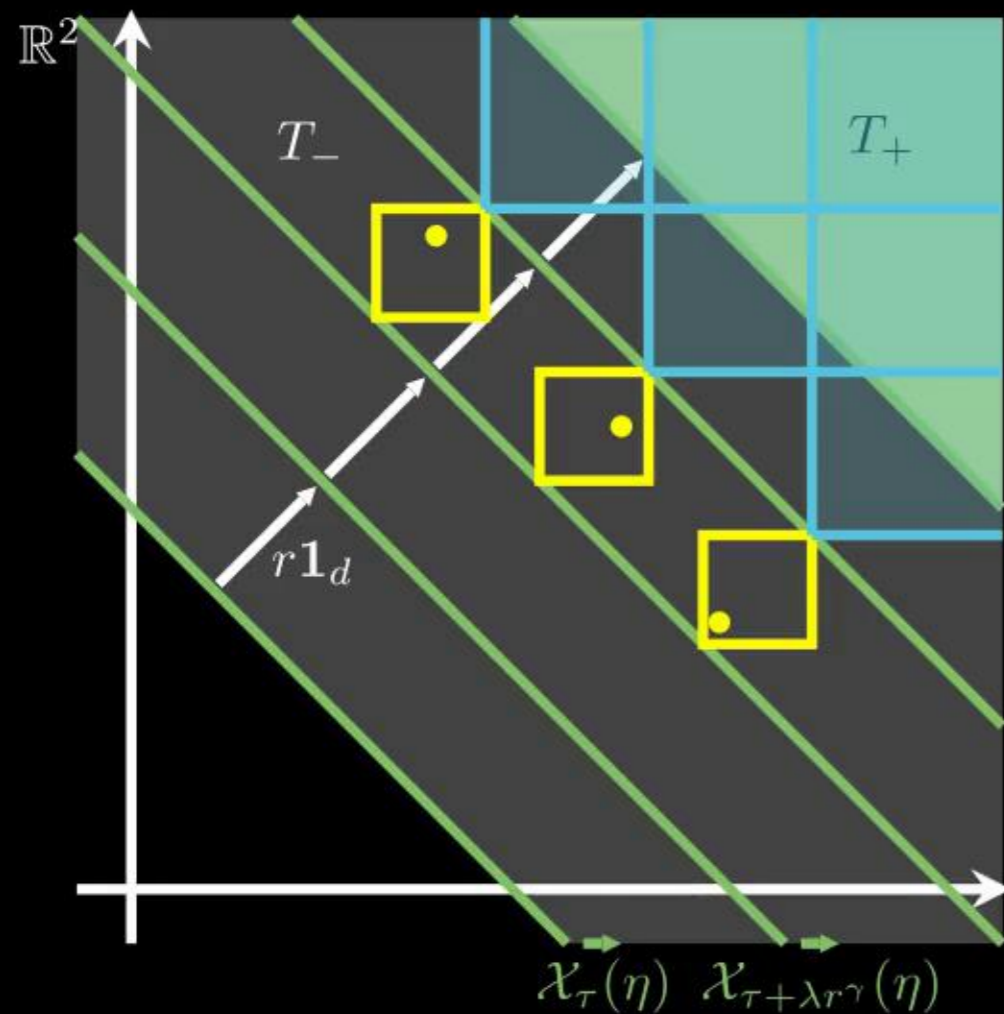
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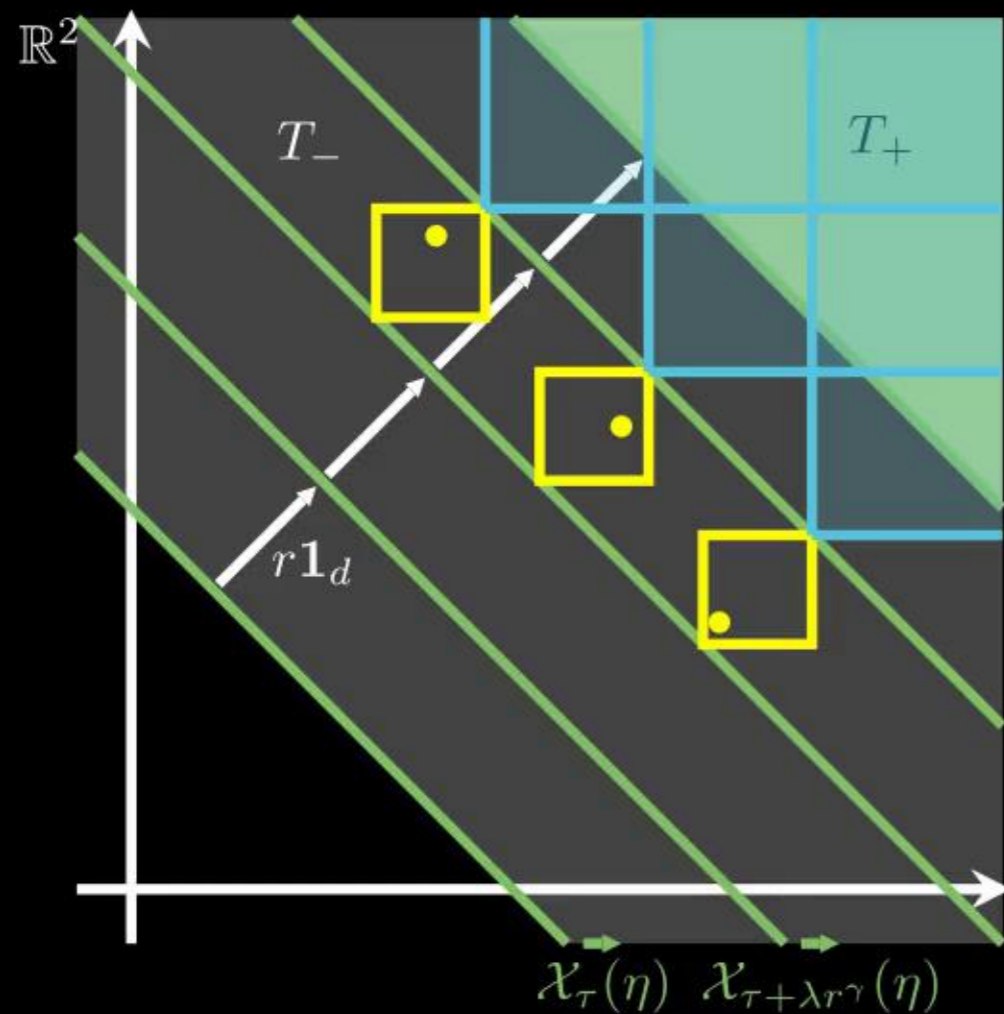
Out of $(X_i)_{i \in [m]}$ roughly mr^d in each box.

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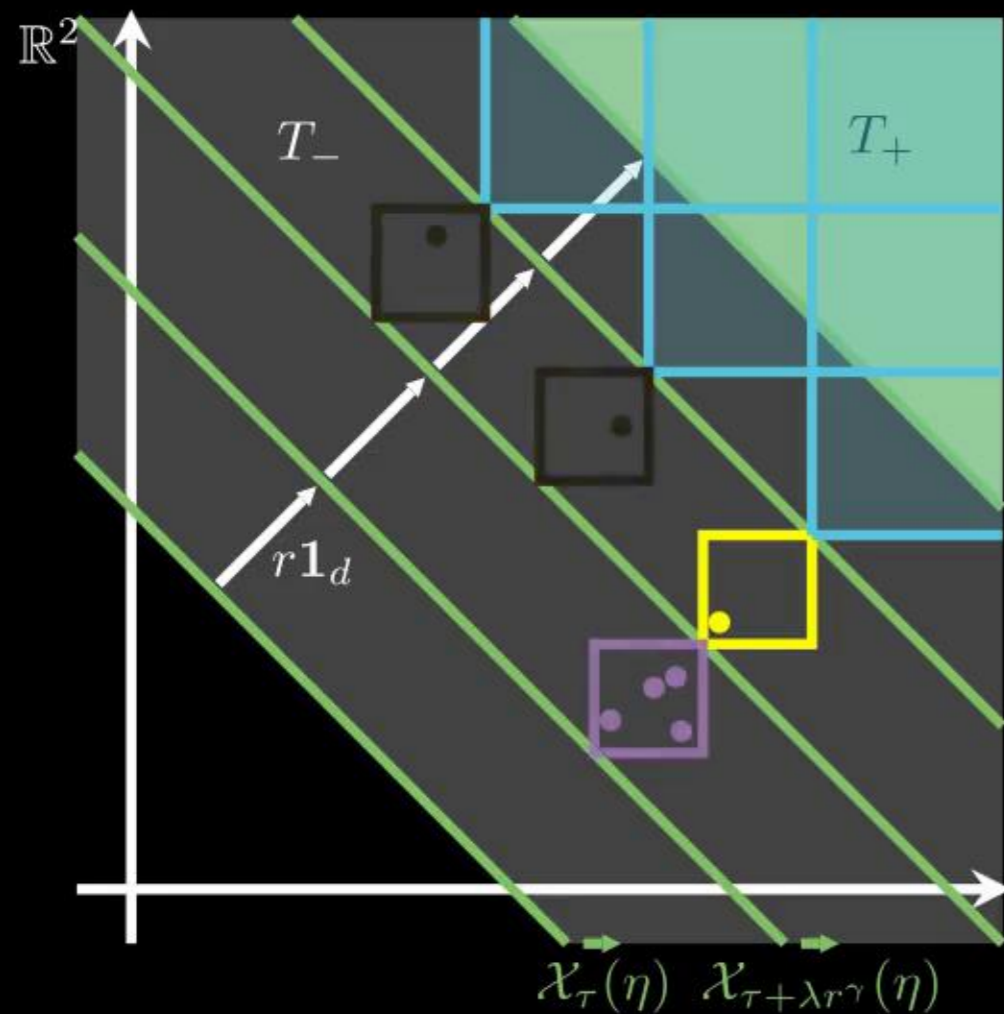
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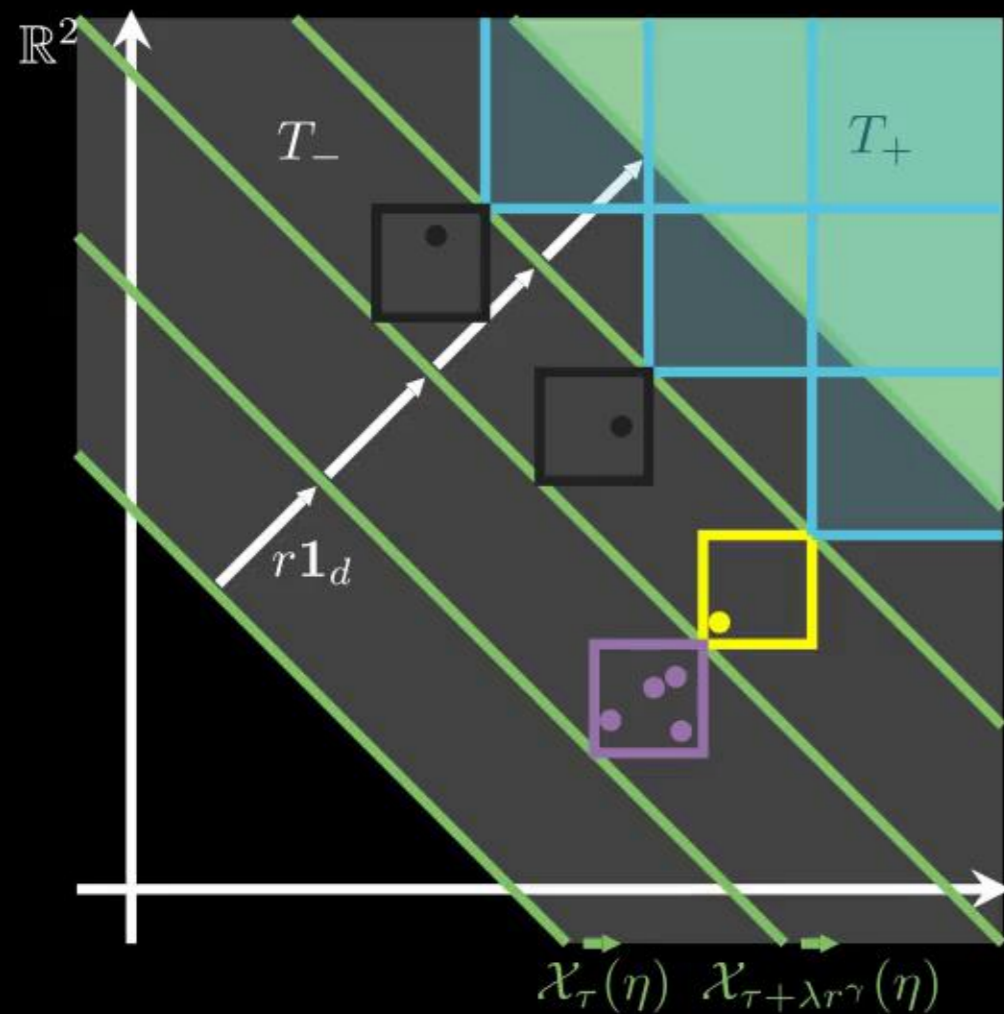
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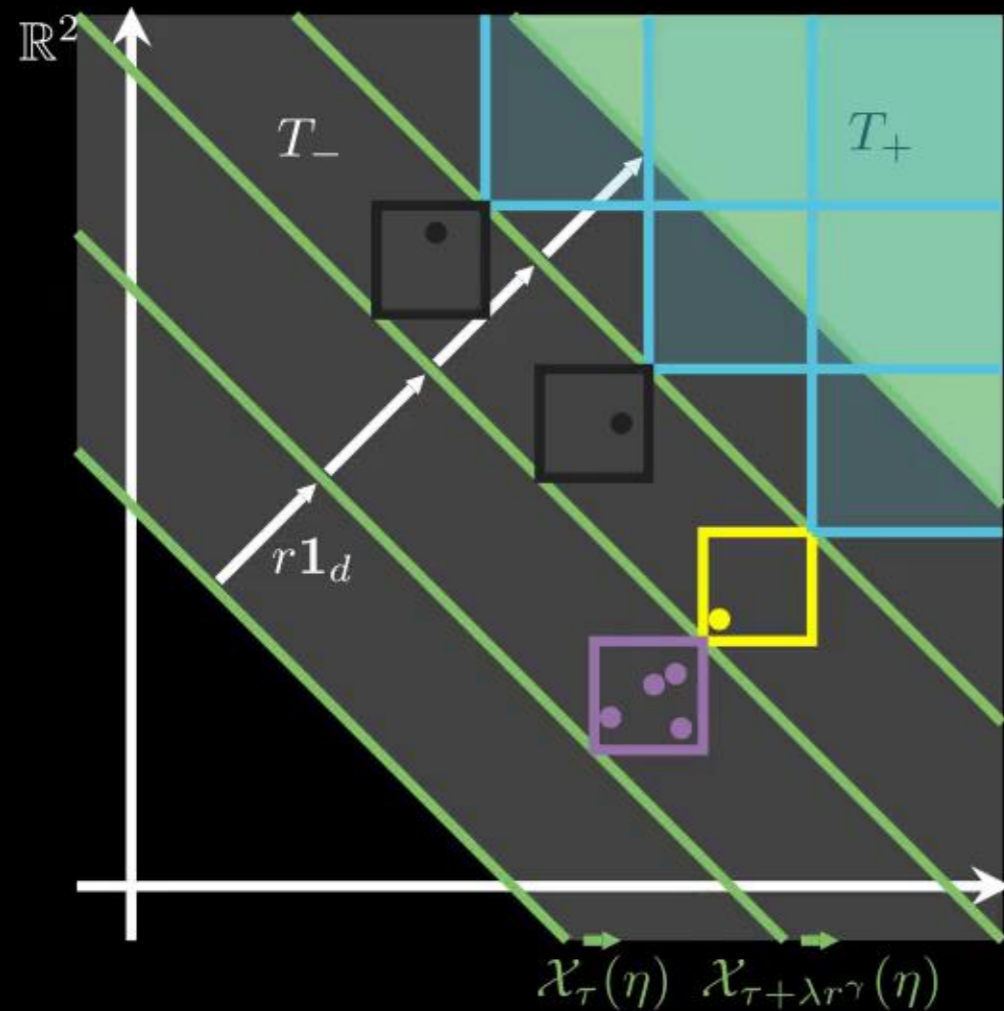
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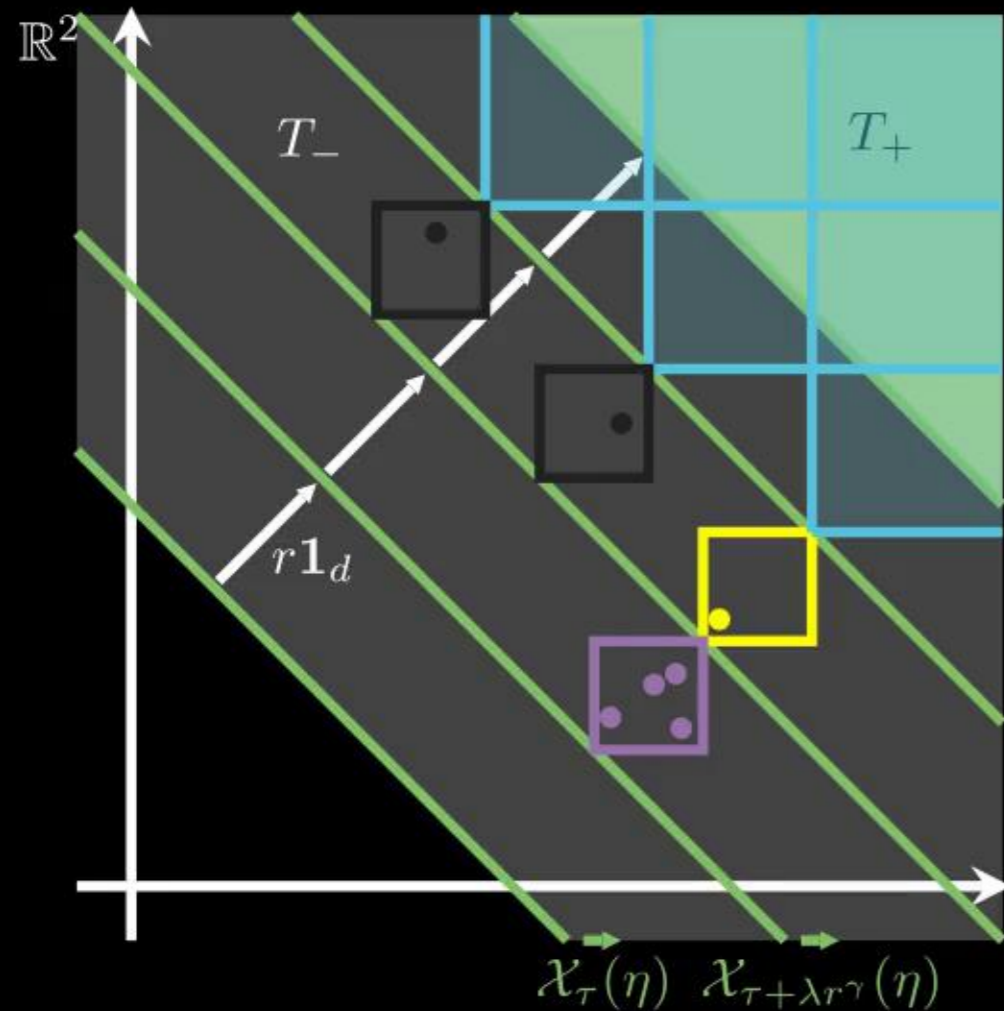
Box contains roughly nr^d points with expected response $\geq \tau + \lambda r^\gamma$.

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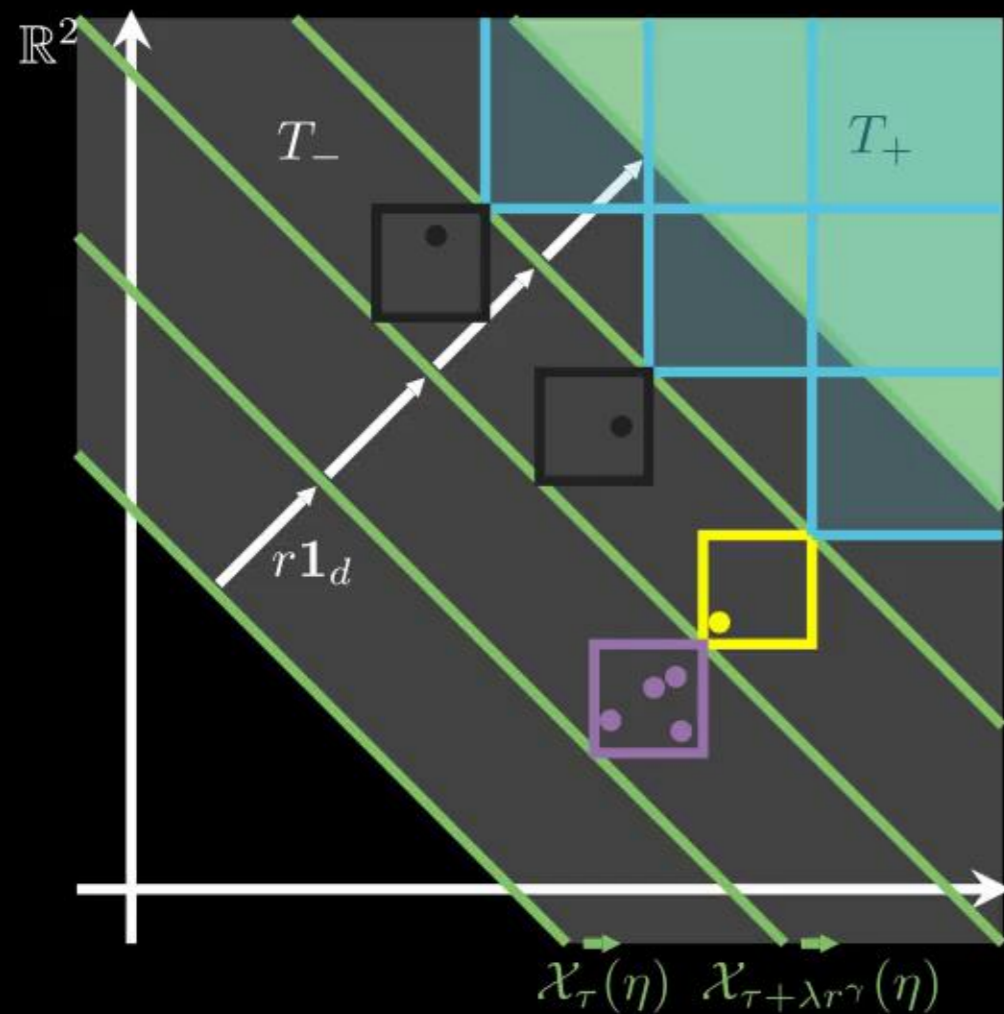
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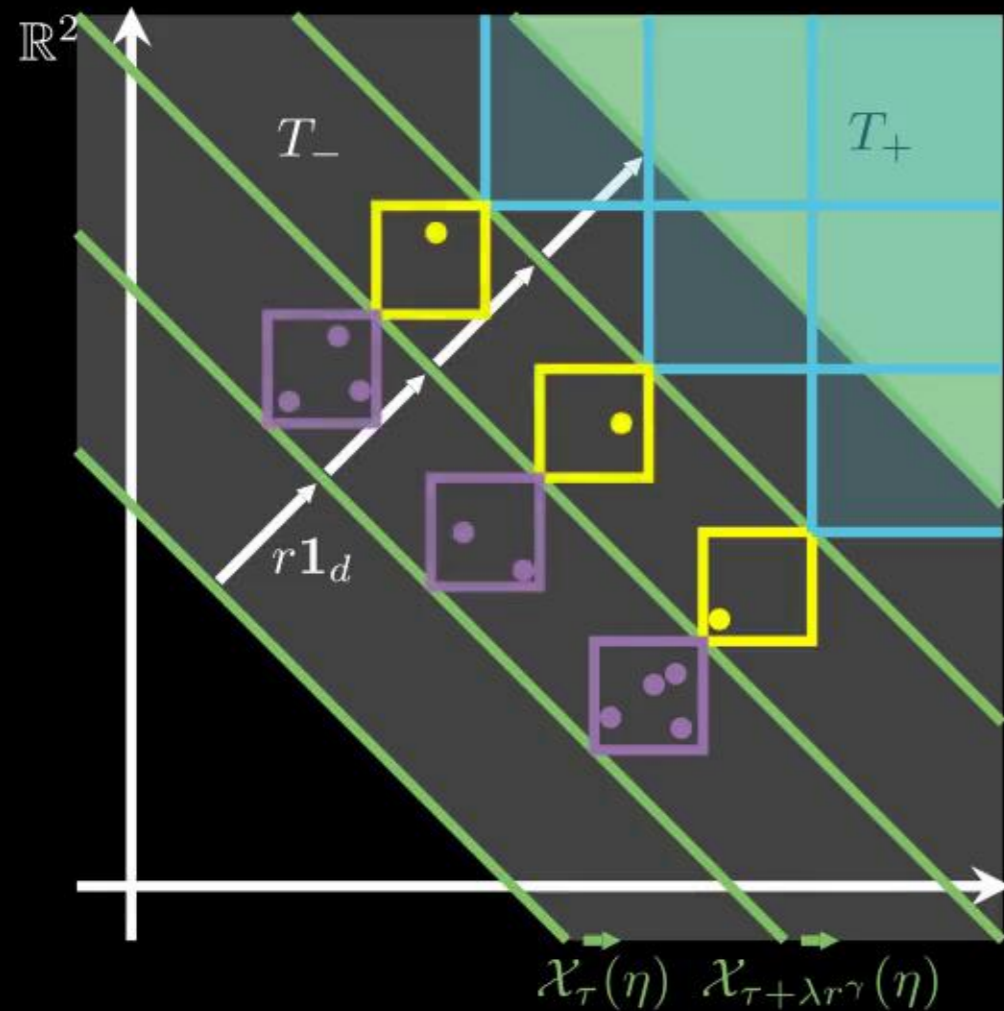
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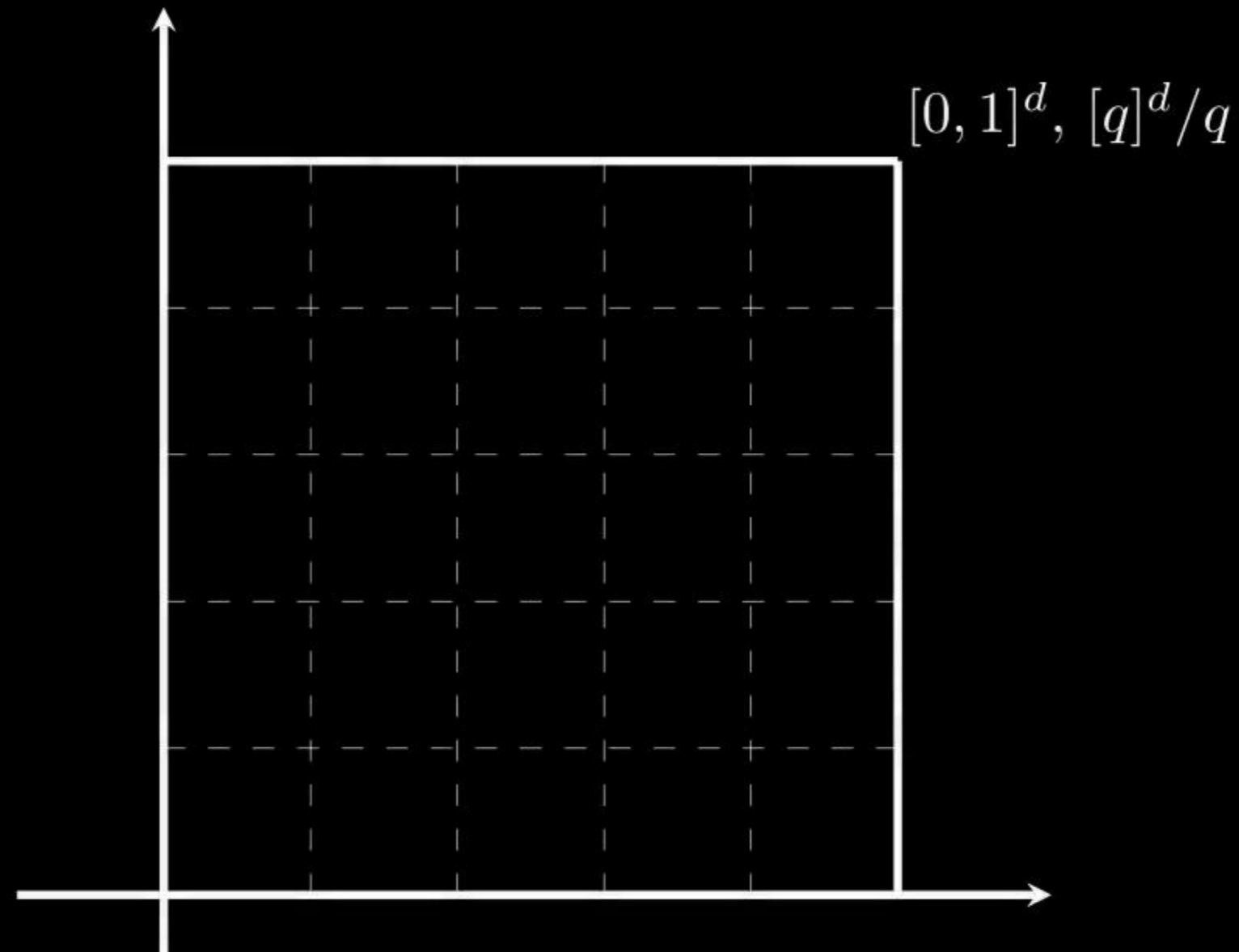
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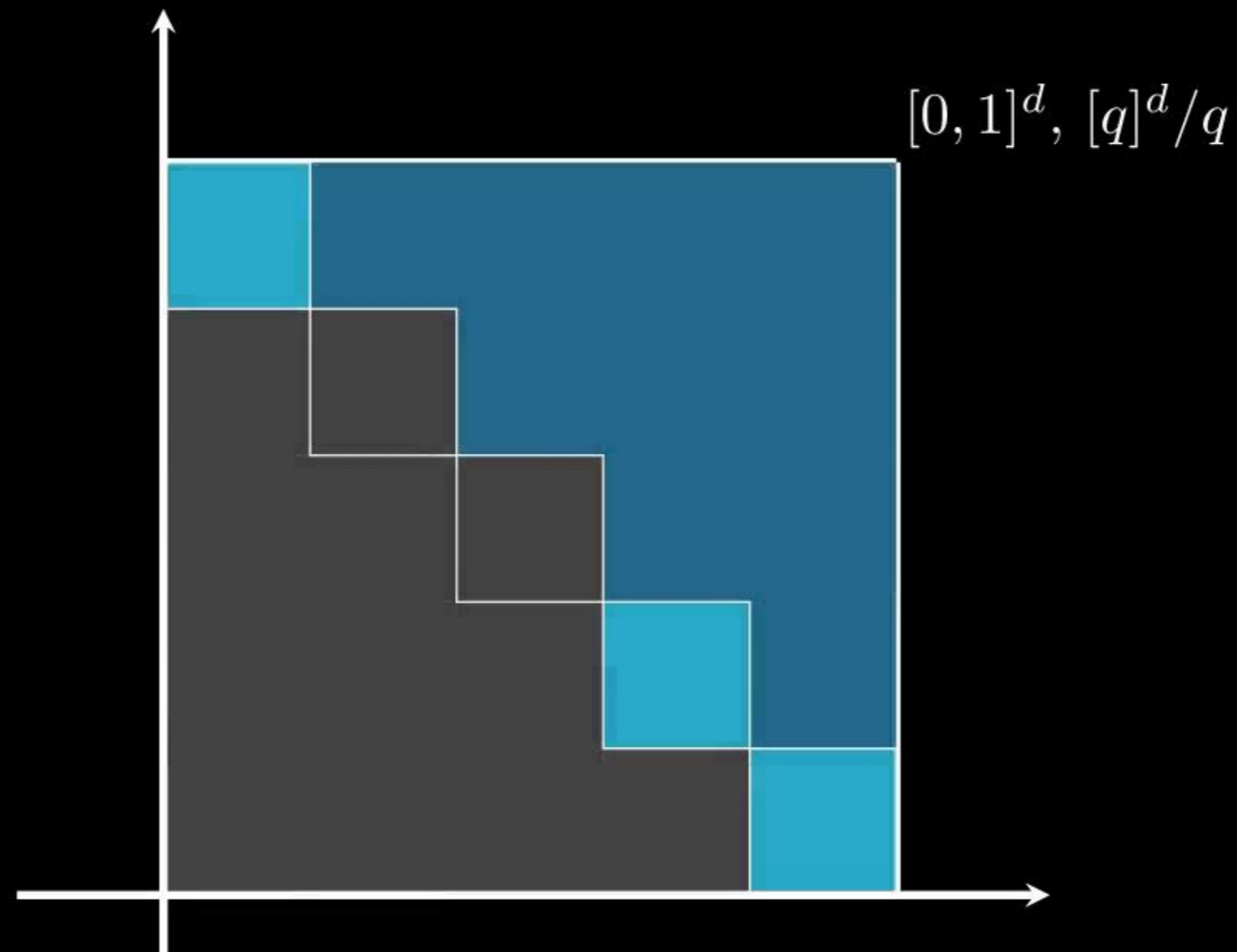
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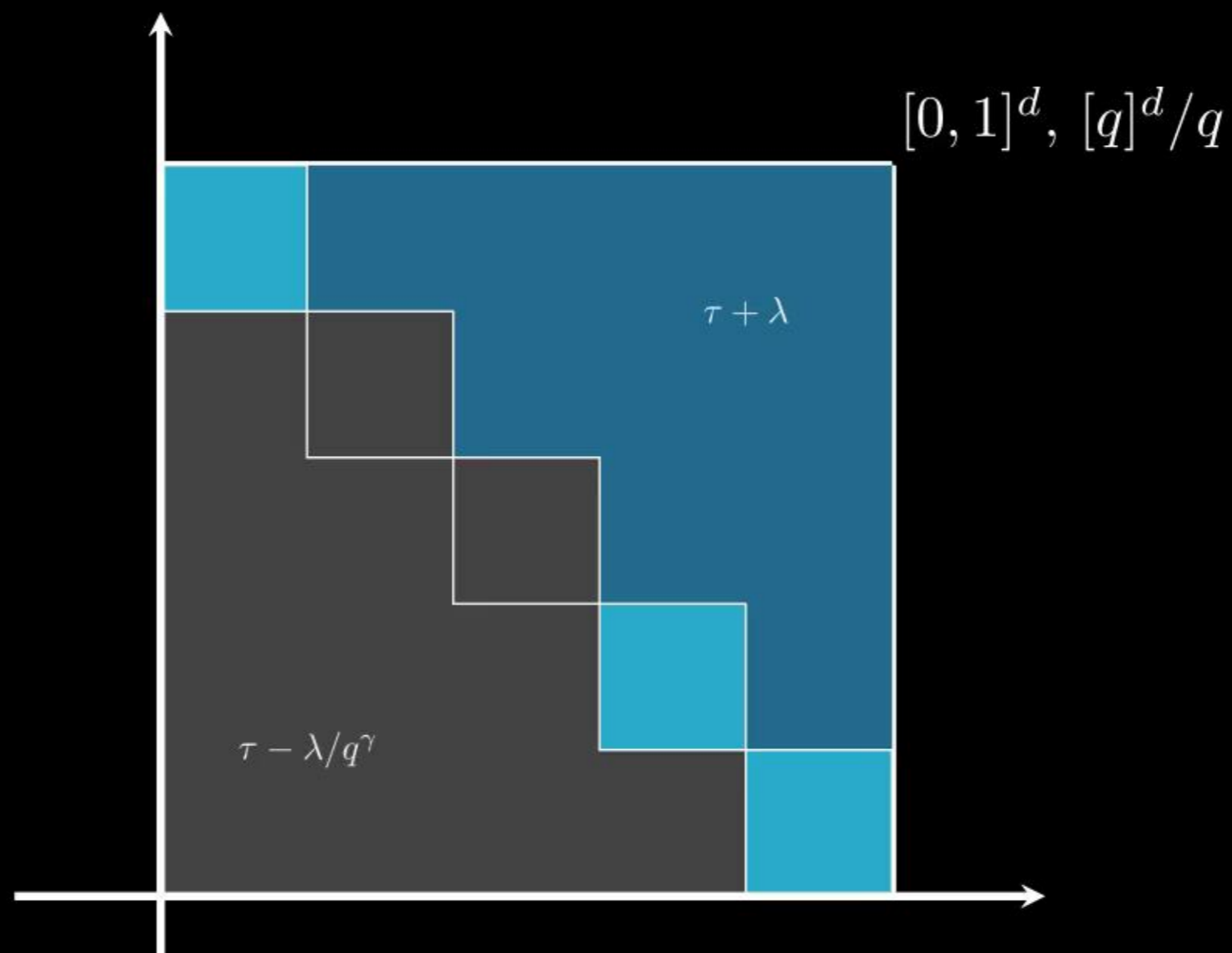
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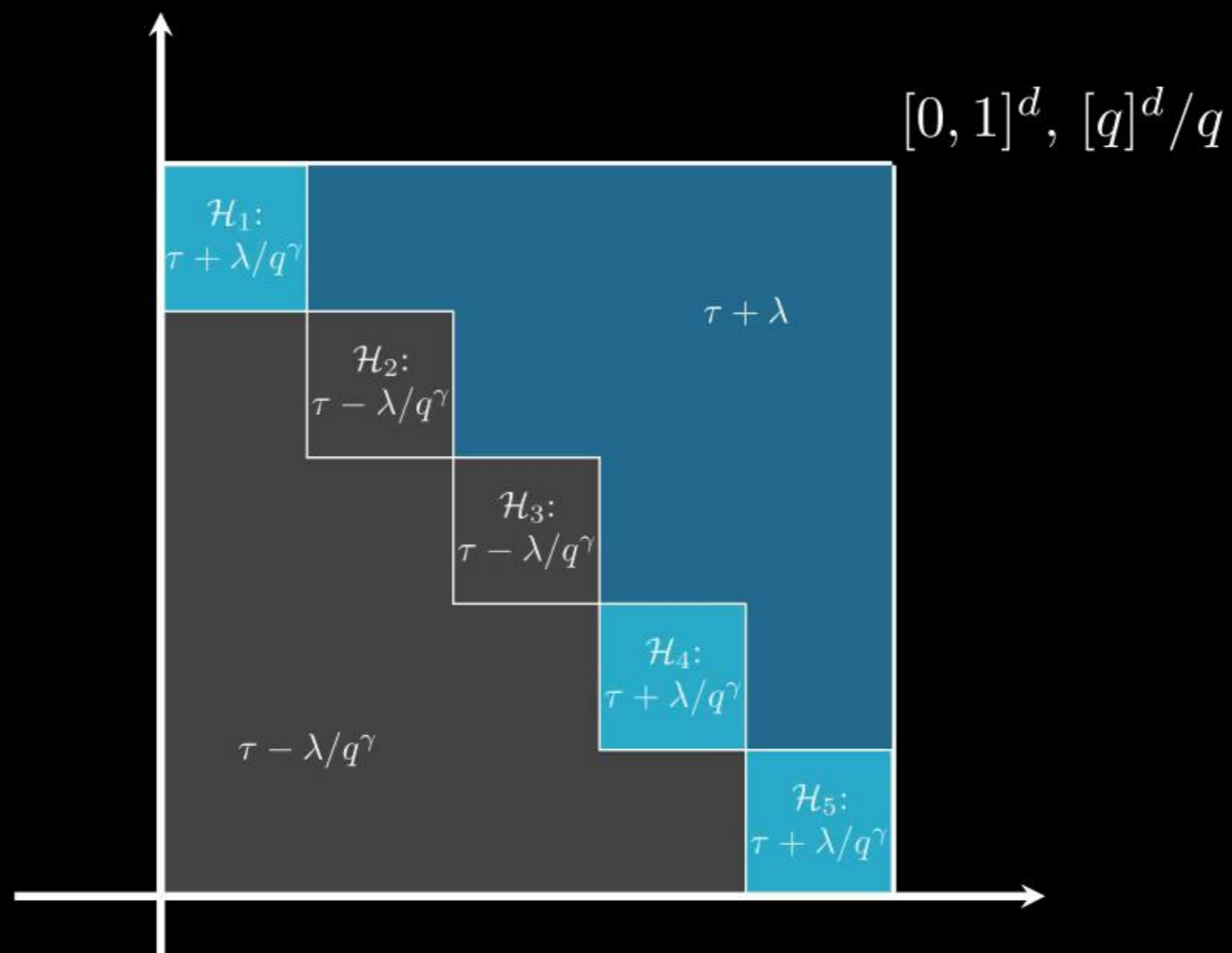
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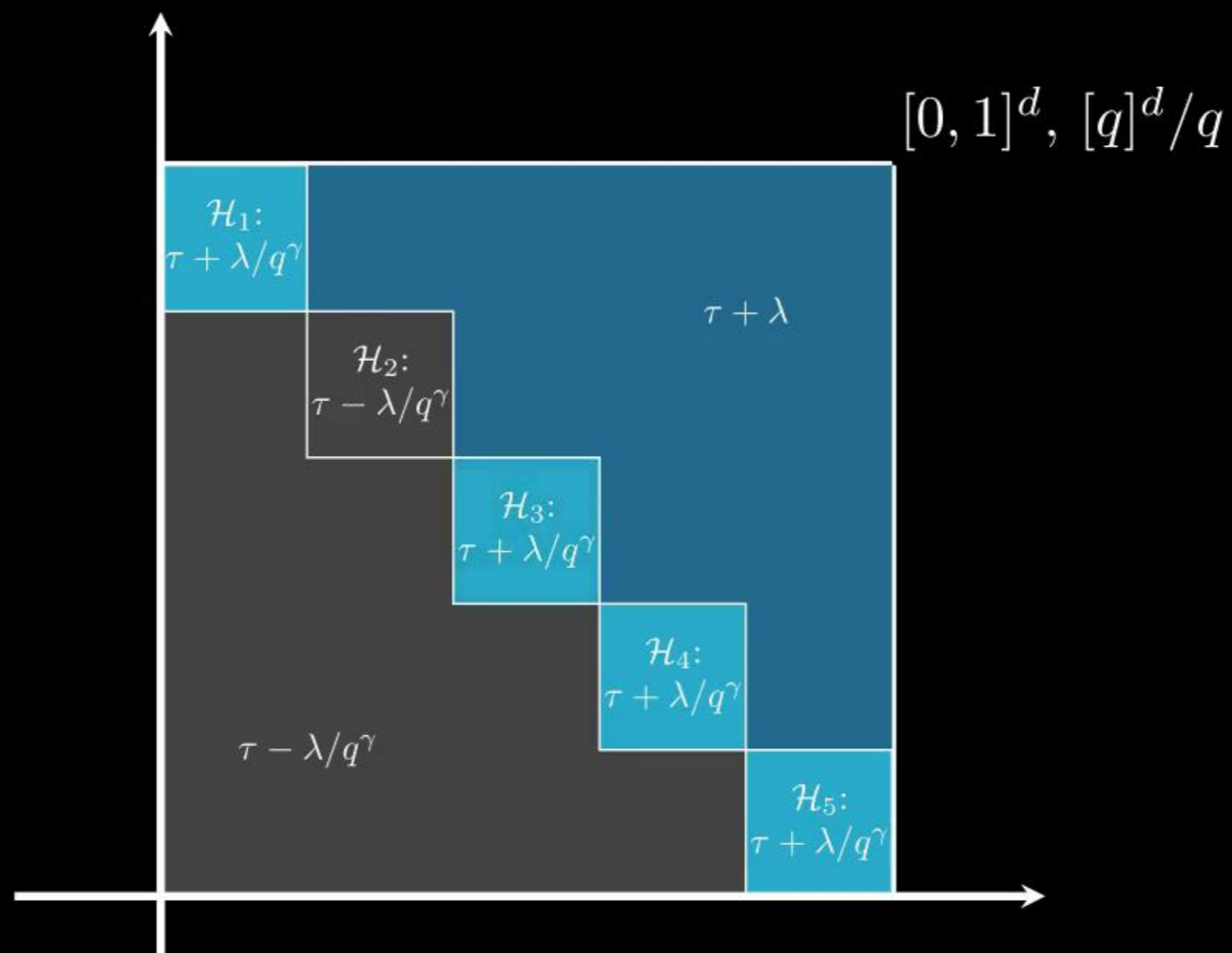
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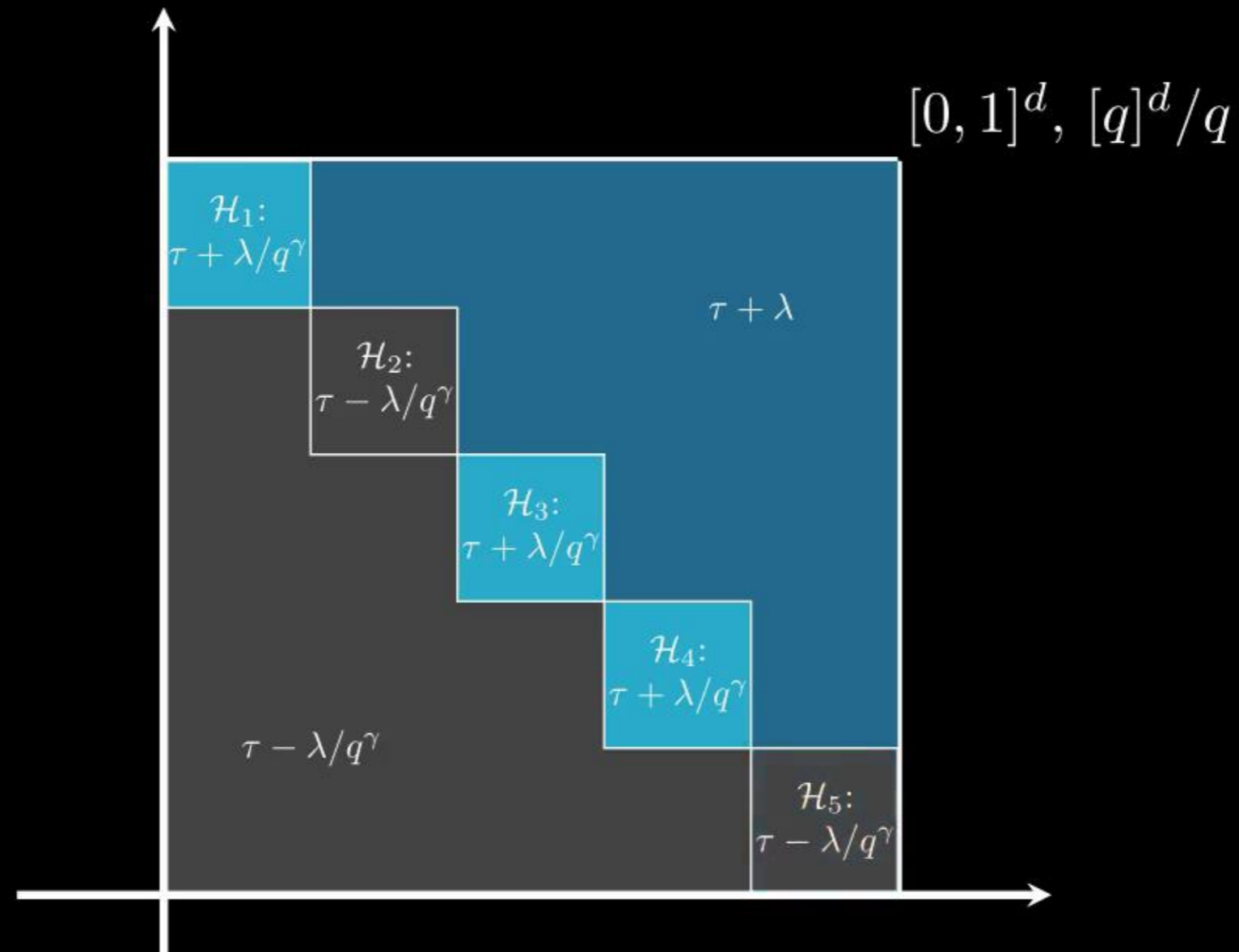
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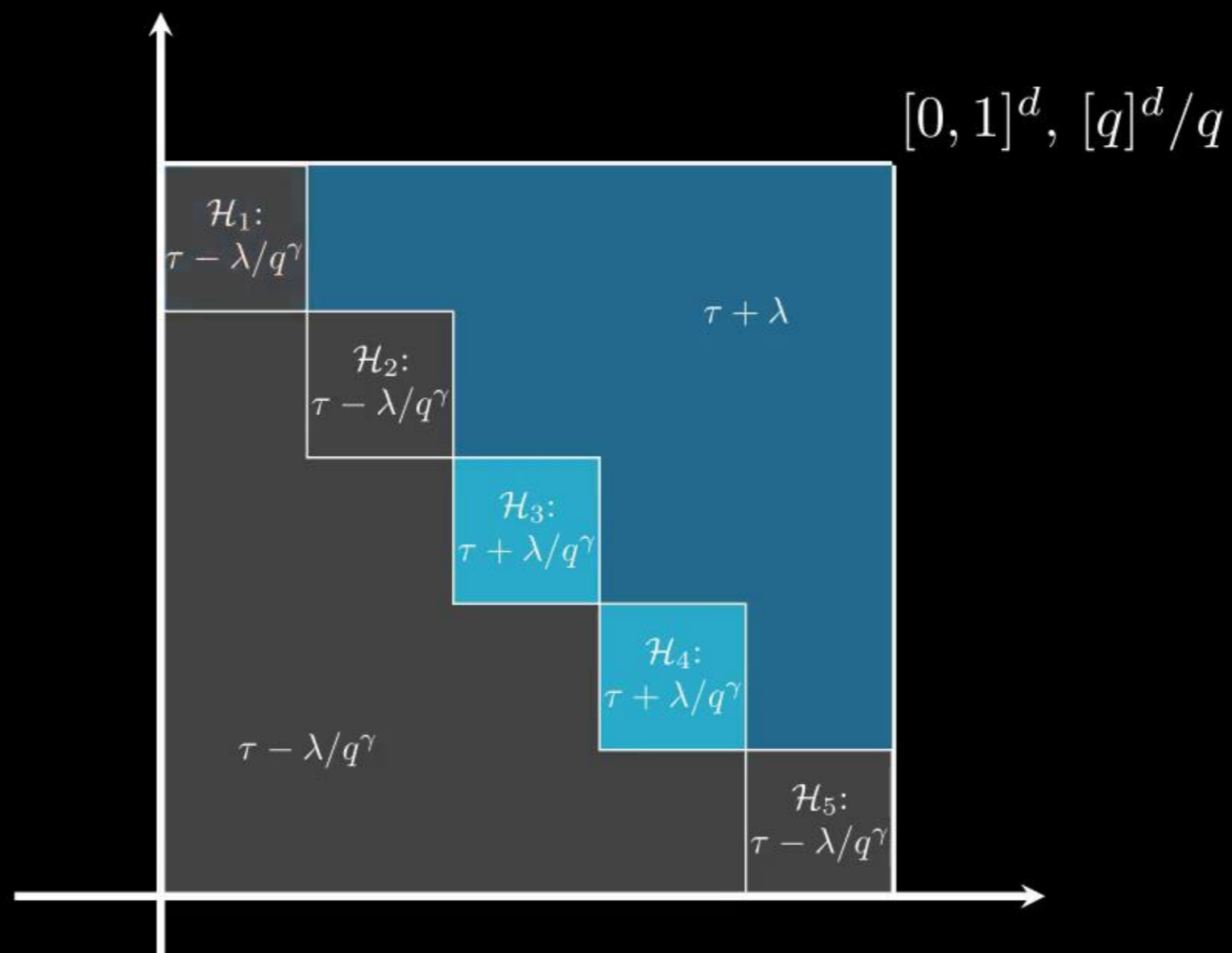
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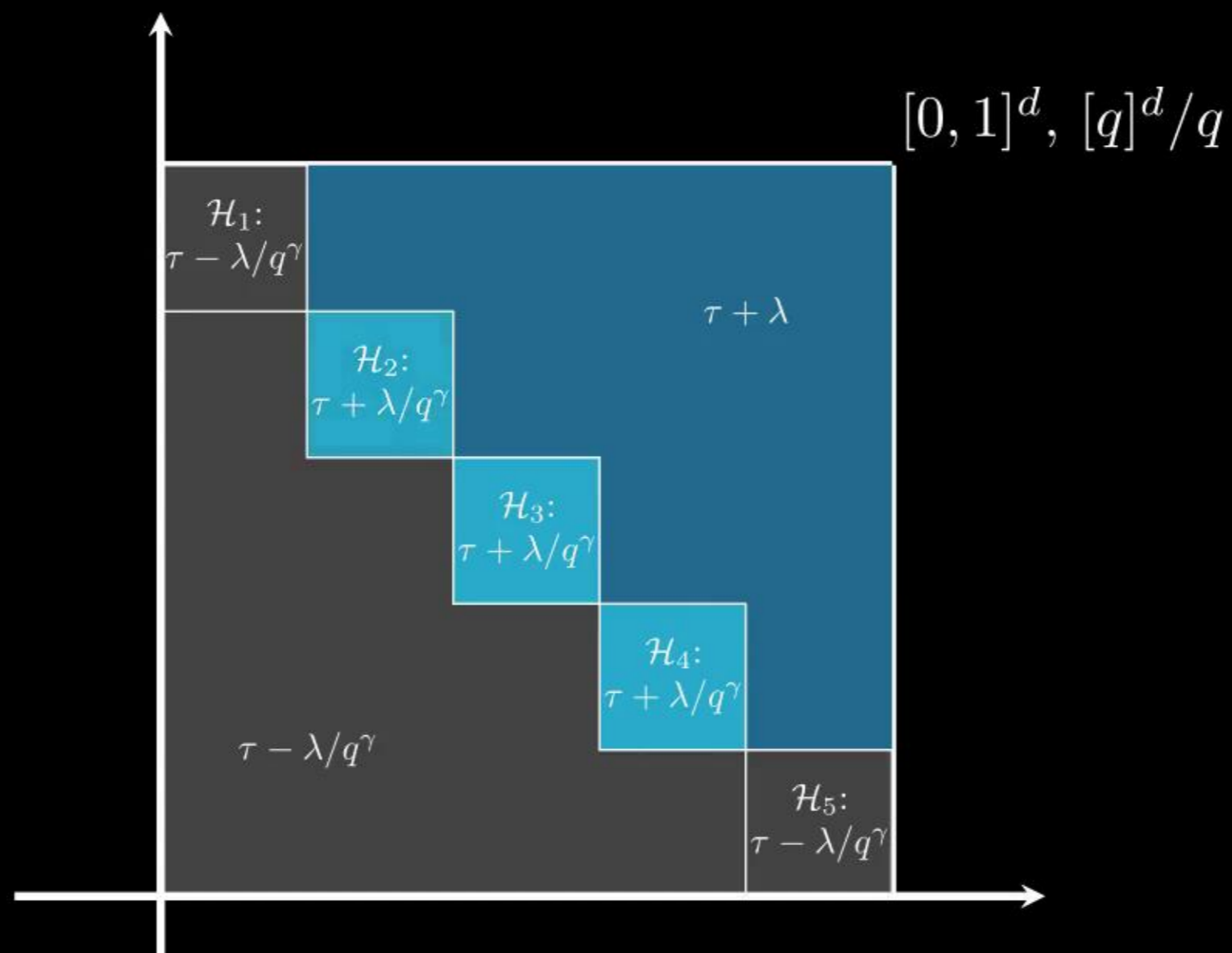
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Write $\pi(x) := \mathbb{P}(T = 1 | X = x)$ for the *propensity score*, and consider the *inverse propensity weighted response*

$$Y := \frac{T - \pi(X)}{\pi(X)(1 - \pi(X))} \cdot \tilde{Y},$$

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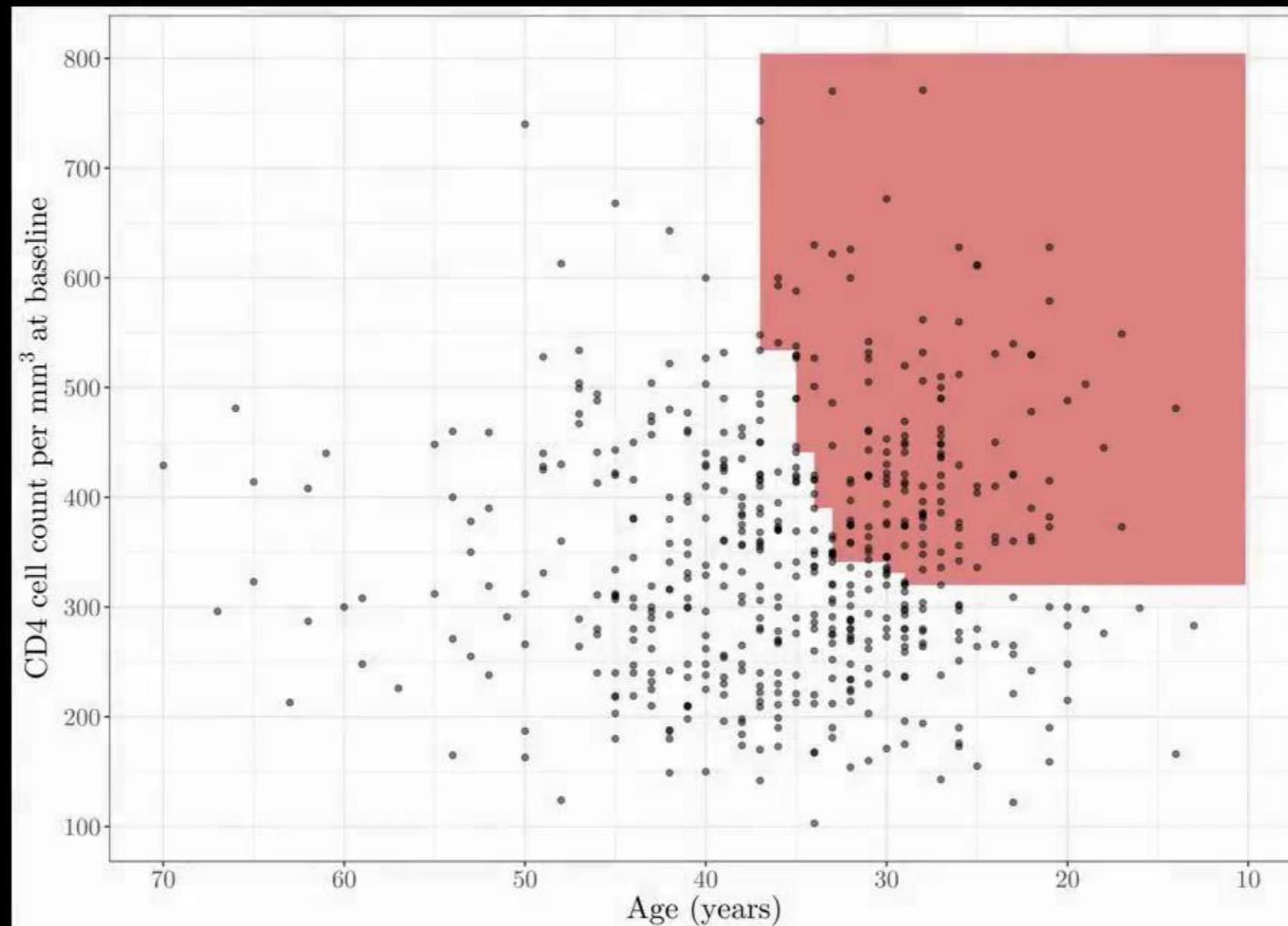
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Hence, if we compute $\mathcal{D} = ((X_1, Y_1), \dots, (X_n, Y_n)) \sim P^n$, then $\mathbb{P}(\hat{A}^{\text{ISS}}(\mathcal{D}) \subseteq \mathcal{X}_\tau(\eta)) \geq 1 - \alpha$ whenever $P \in \mathcal{P}_{\text{Mon},d}(\sigma)$.

Application

Primary endpoint: reduction of the CD4 cell count by 50%, development of AIDS, or death, with median follow-up duration 143 weeks (Hammer et al., 1996). Let $\alpha = 0.05$ and $\tau = 1/2$.



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In common situations, no smoothing-parameters have to be specified.

References and acknowledgement

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Thank you!

Main reference:

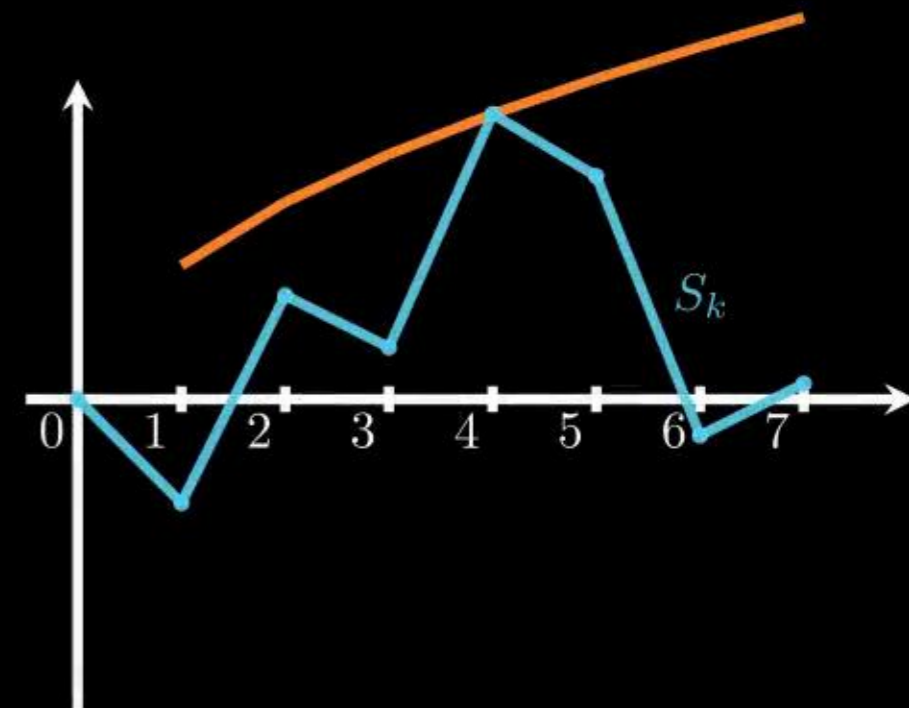
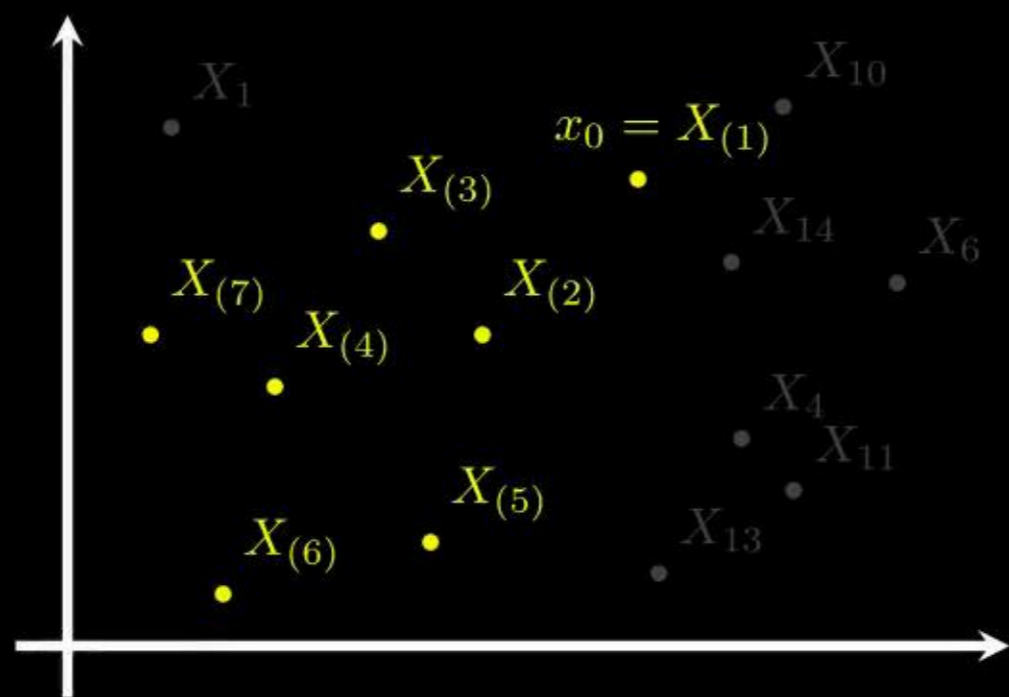
Müller, M. M., Reeve, H. W. J., Cannings, T. I. and Samworth, R. J. (2023) Isotonic subgroup selection. *arXiv preprint arXiv:2305.04852*.

See manuelmueller.github.io for data and R-code.

Appendix

Construct p -values \hat{p}_i for $H_0(X_i)$, $i \in [m]$

Given $x_0 \in \mathbb{R}^d$, we seek a p -value for $H_0(x_0) := \{P \in \mathcal{P}_{\text{Mon},d}(\sigma) : \eta(x_0) < \tau\}$.



Denote $\mathcal{I}(x_0) := \{i \in [n] : X_i \preceq x_0\}$, $n(x_0) := |\mathcal{I}(x_0)|$.

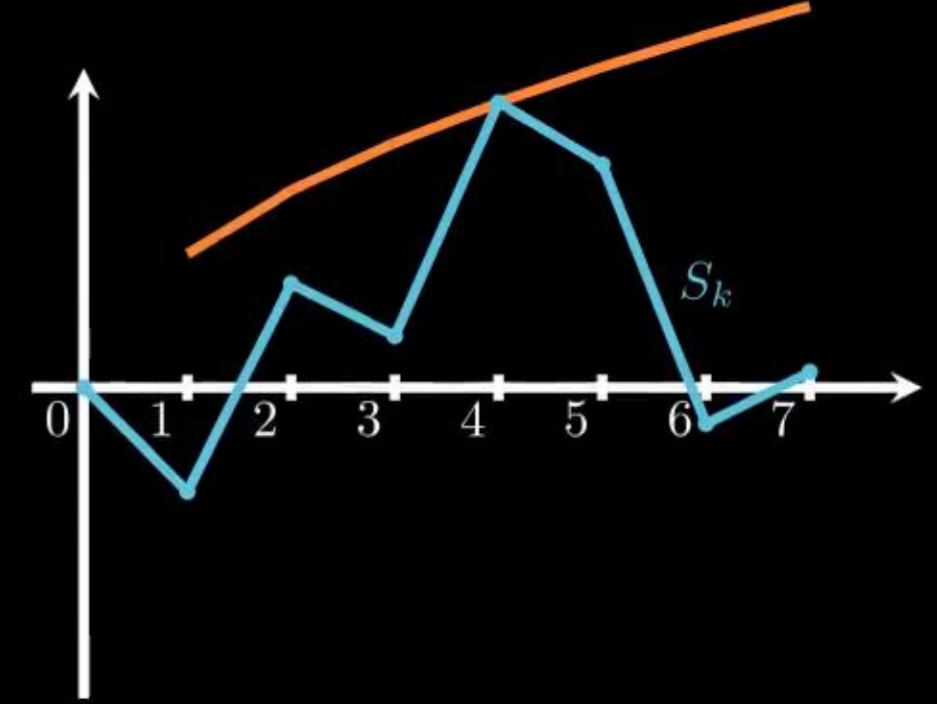
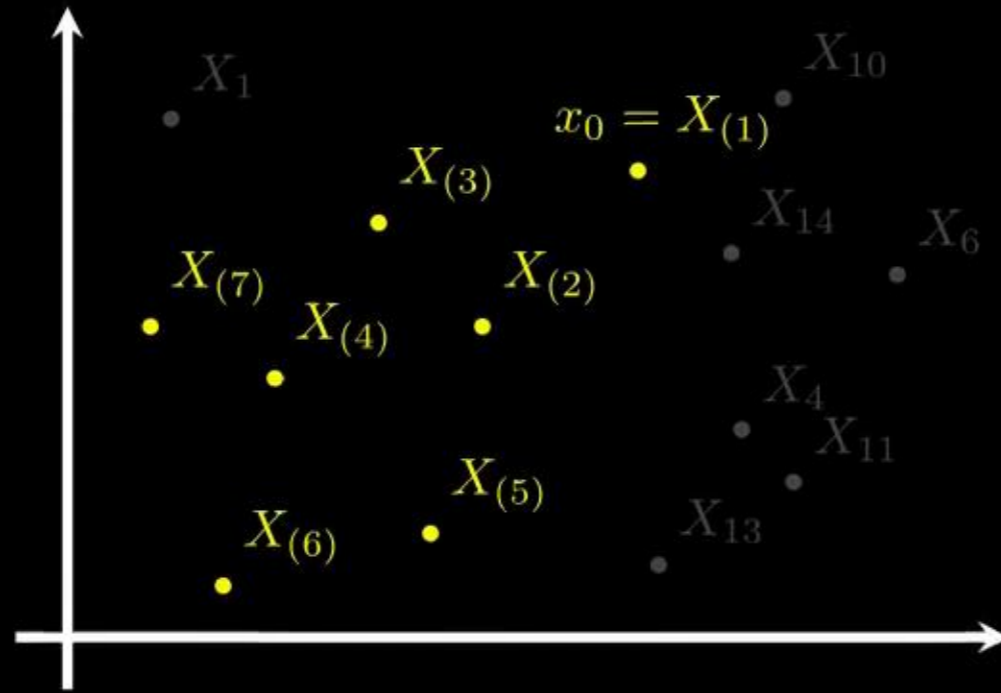
Let $X_{(j)}$ be the j th nearest neighbour of x_0 among X_i , $i \in \mathcal{I}(x_0)$, in sup-norm and let $Y_{(j)}$ be the corresponding response.

Let

$$S_k := \sum_{j=1}^k \frac{Y_{(j)} - \tau}{\sigma}.$$

Then S_k is a supermartingale under $P \in H_0(x_0)$. Combination with time-uniform bounds by Howard et al. (2021) gives p -values from this martingale test (Duan et al., 2020).

Construct p -values \hat{p}_i for $H_0(X_i)$, $i \in [m]$

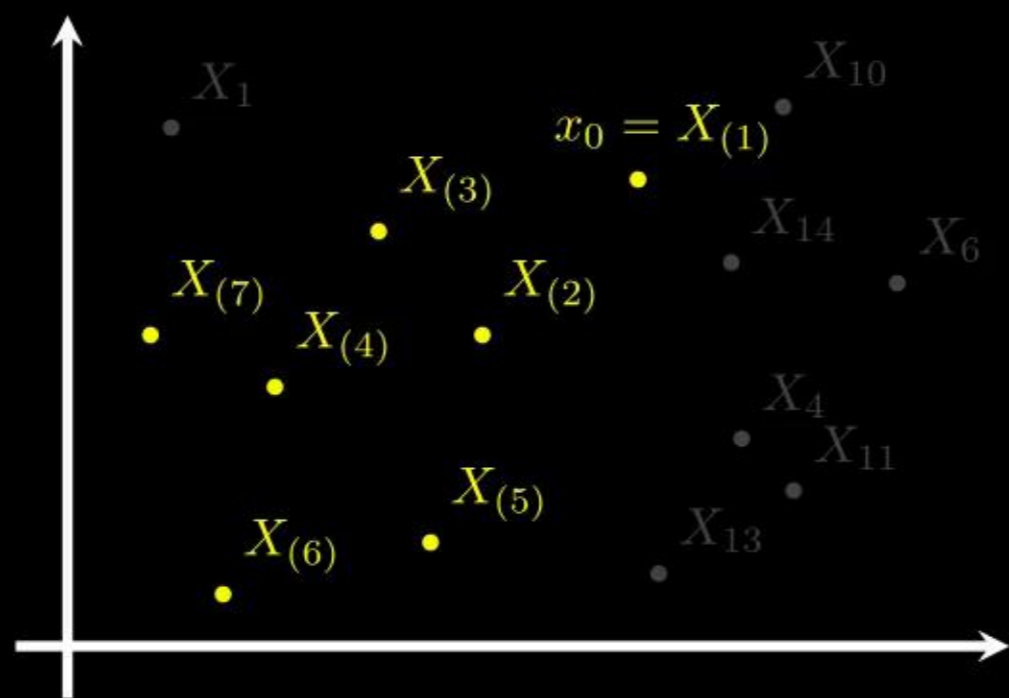


Definition. For $\sigma > 0$, $x \in \mathbb{R}^d$, let

$$\hat{p}_{\sigma, \tau}(x) := 1 \wedge \min_{k \in [n(x)]} 5.2 \exp \left\{ -\frac{(S_k \vee 0)^2}{2.0808k} + \frac{\log \log(2k)}{0.72} \right\},$$

whenever $n(x) > 0$, and $\hat{p}_{\sigma, \tau}(x) := 1$ otherwise.

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Consider adaptation to σ^2 when $Y - \eta(X)|X \sim \mathcal{N}(0, \sigma^2)$.

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Key idea: use an online split likelihood ratio test (Wasserman et al., 2020) for $H_0 : Y_{(k)}|X_{(k)} \sim \mathcal{N}(t_k, \sigma^2), t_k < \tau, \sigma > 0, \forall k \geq 1$.

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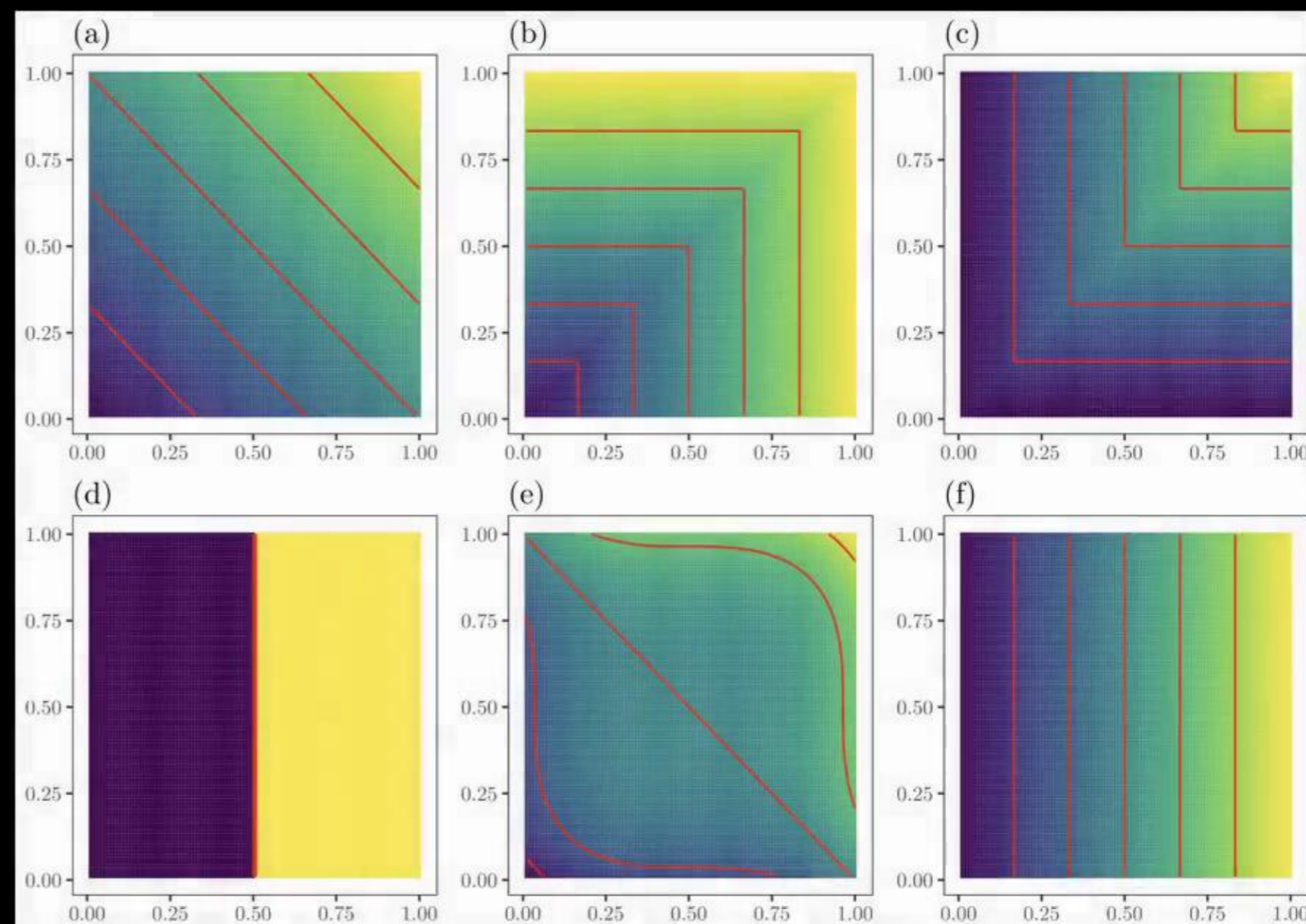
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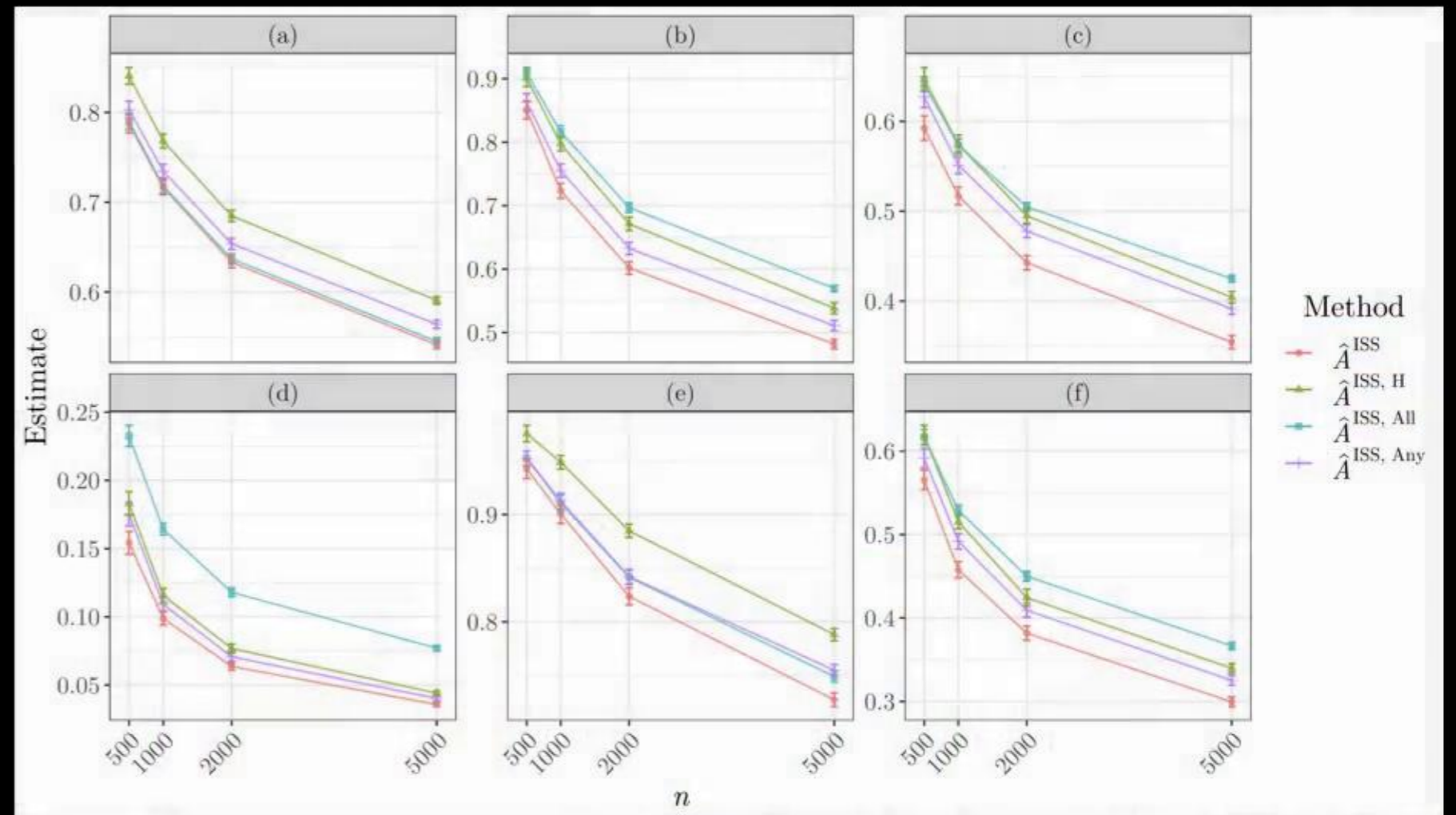
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Here, $d = 2$, $\sigma = 1/4$.

See also Meijer and Goeman (2015).

