

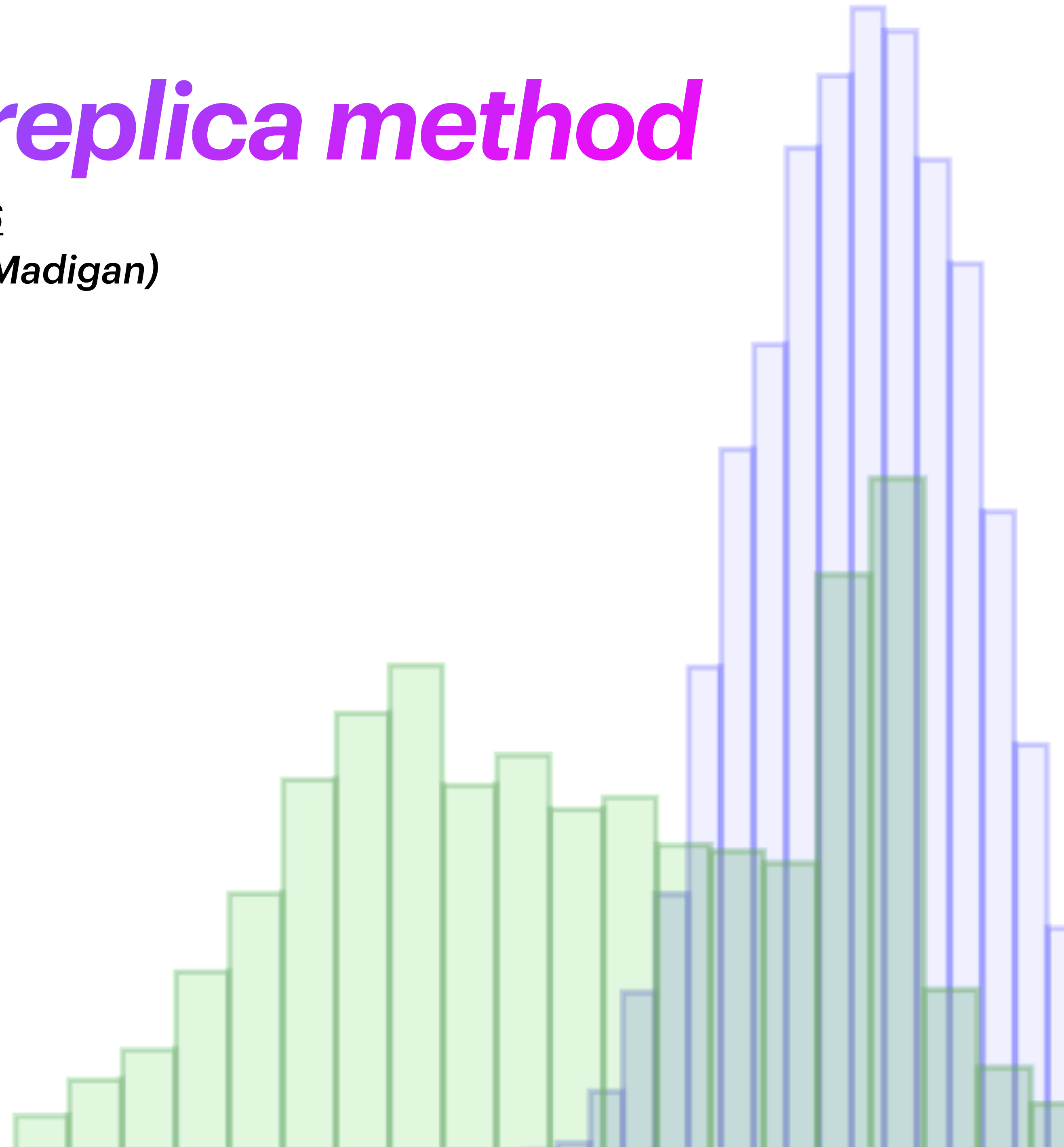
A study of the MC replica method

for PDF4LHC 2024, CERN, based on 2404.10056

(with Mark Costantini, Luca Mantani and Maeve Madigan)

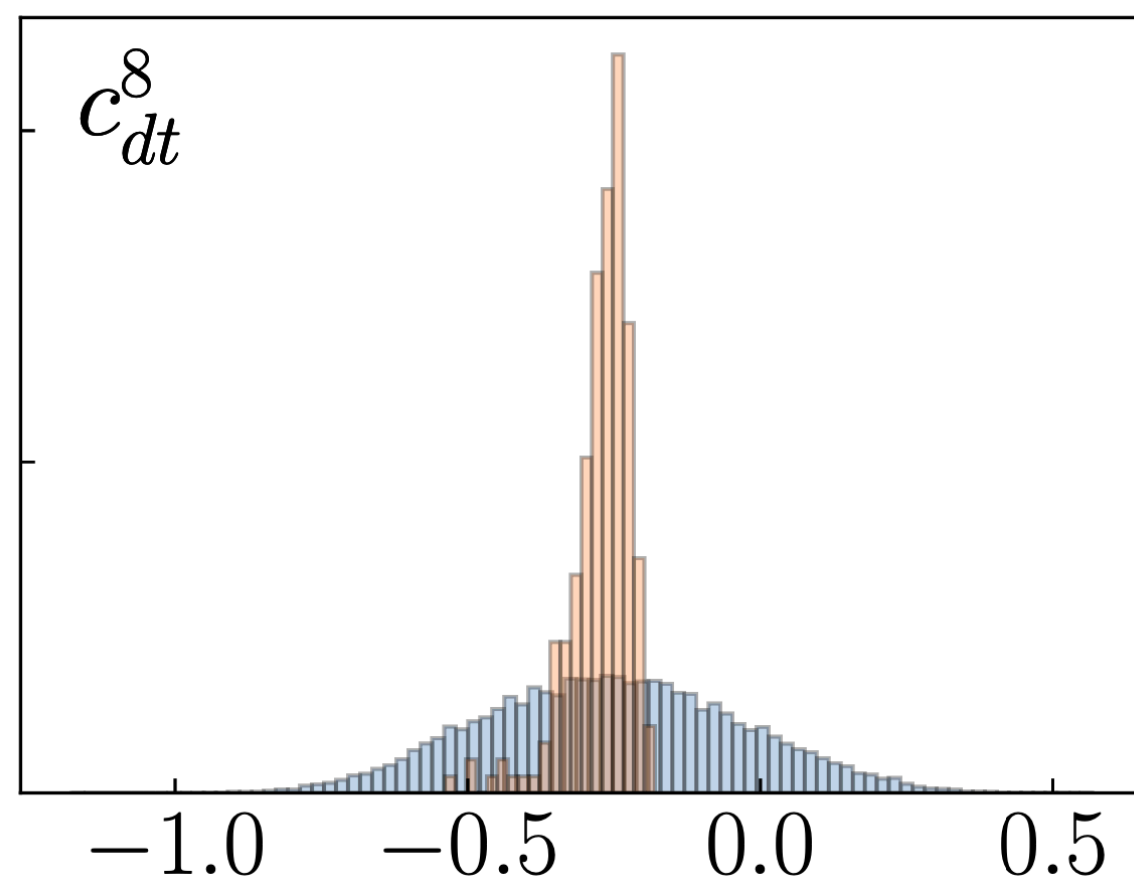


**James Moore,
Lucy Cavendish College,
University of Cambridge**



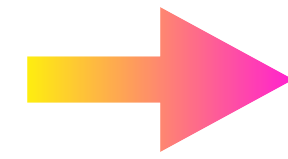
The talk in a nutshell...

1. Why study the Monte Carlo replica method?

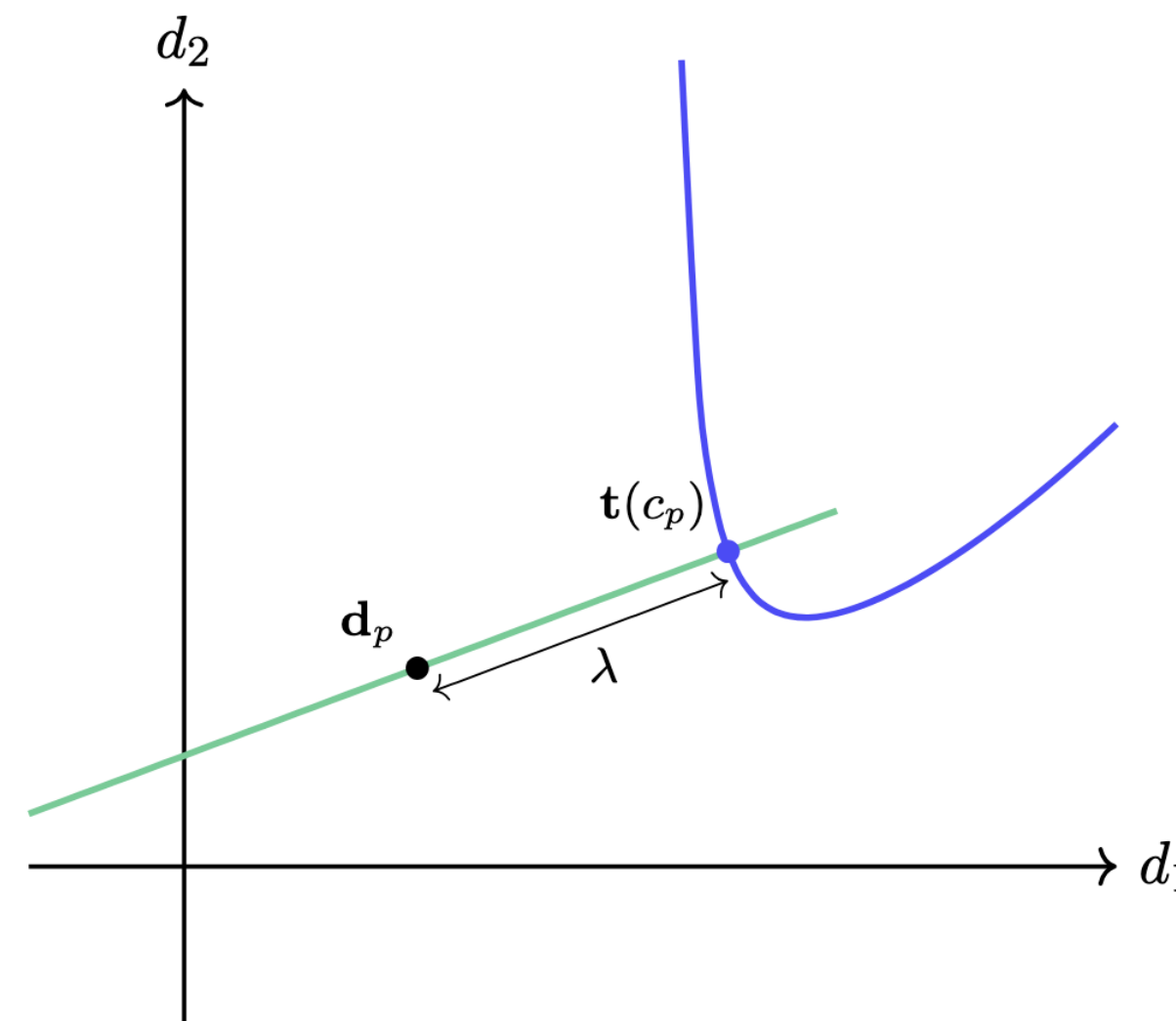
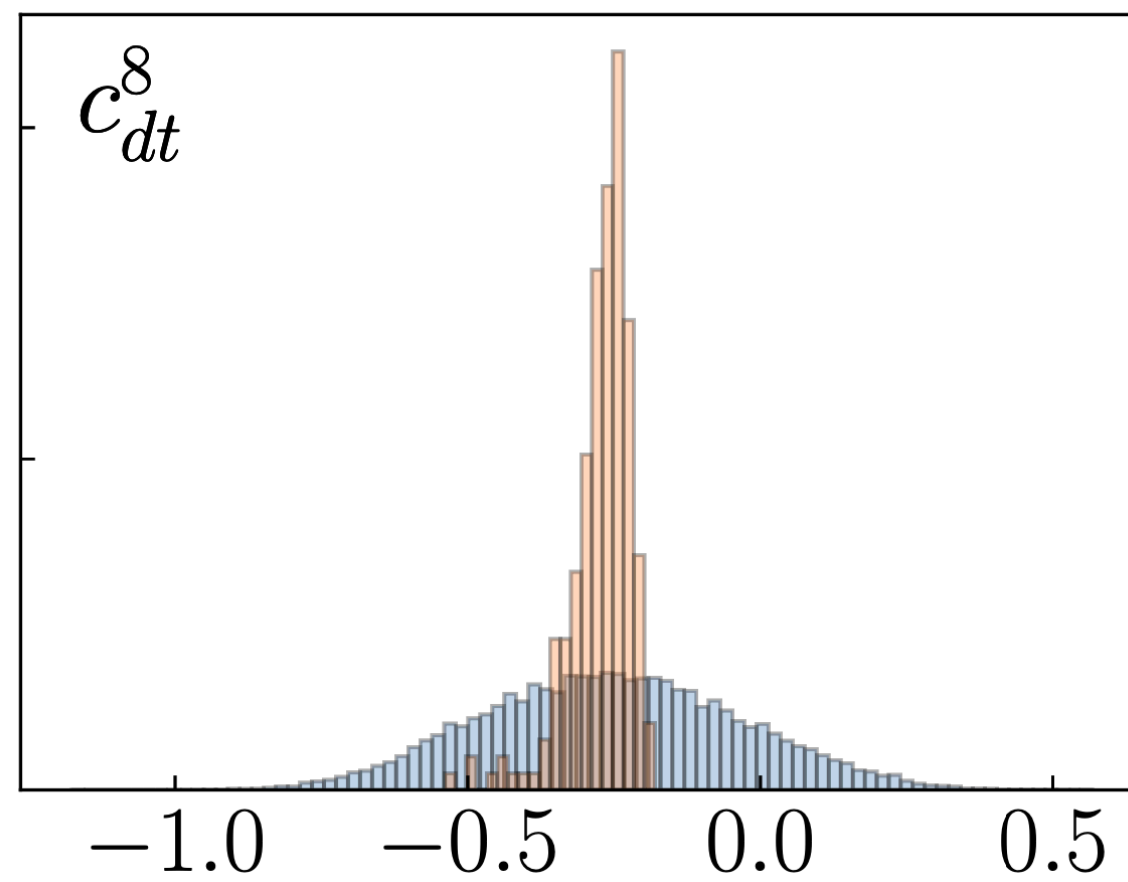


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1. Why study the Monte Carlo replica method?



2. A geometric approach to the MC replica method

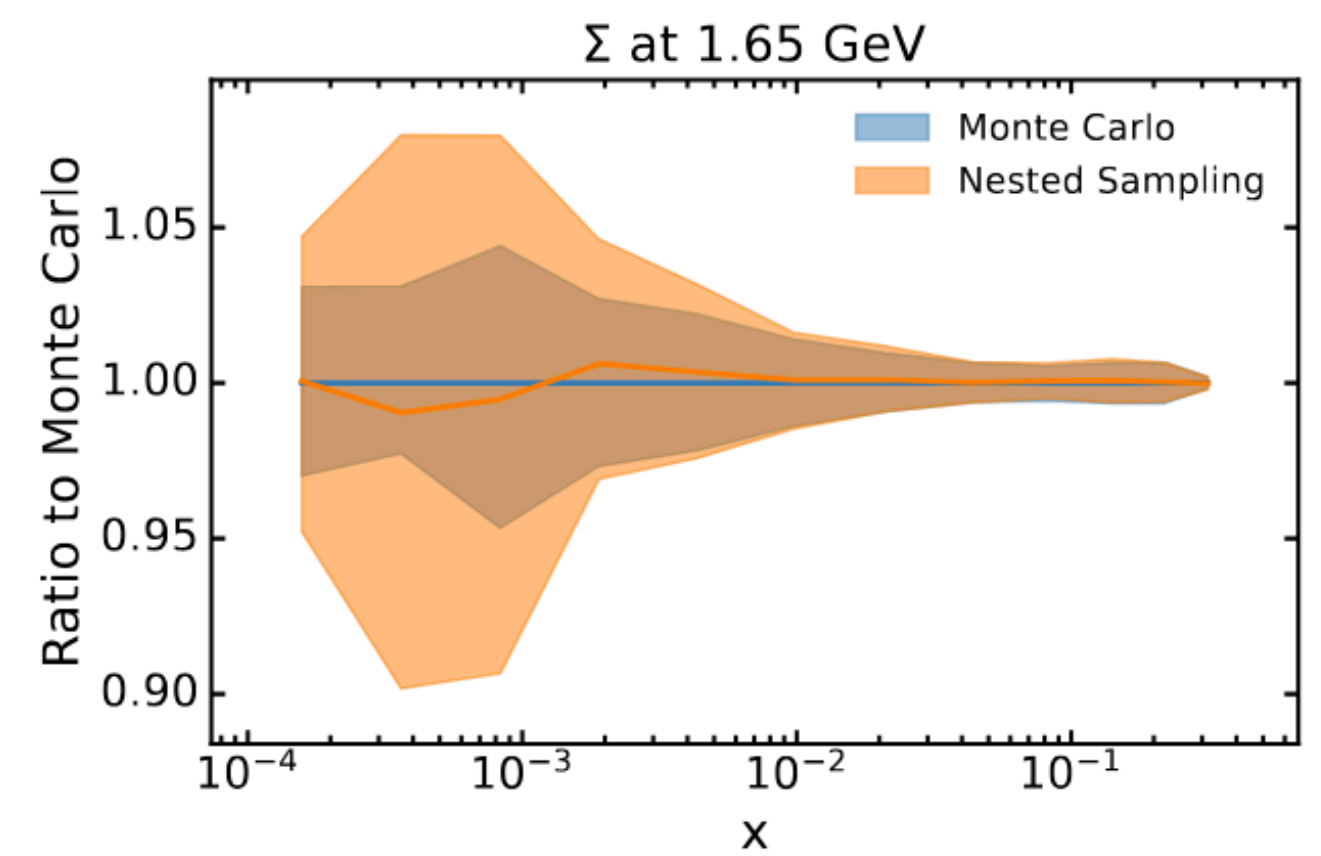
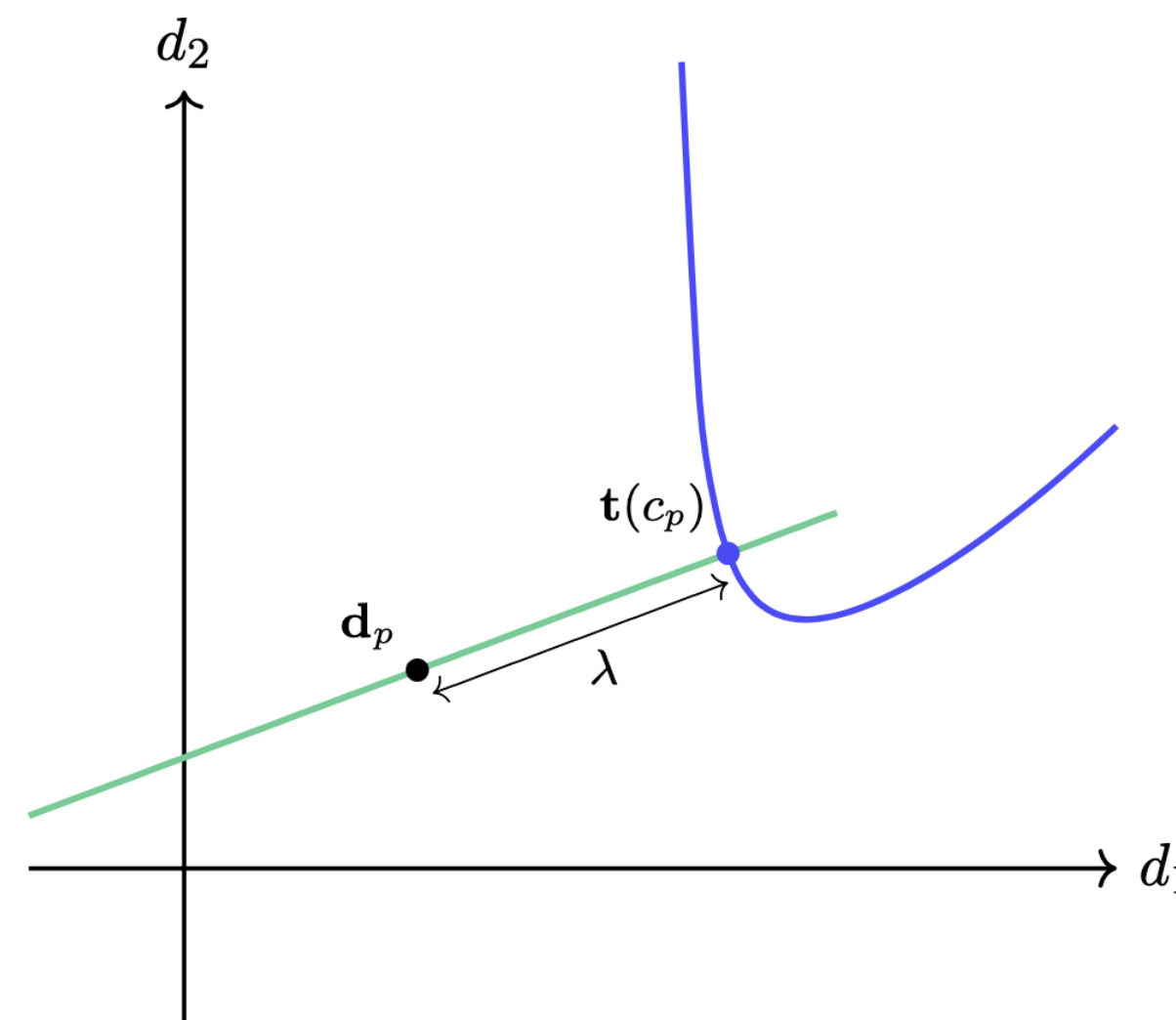
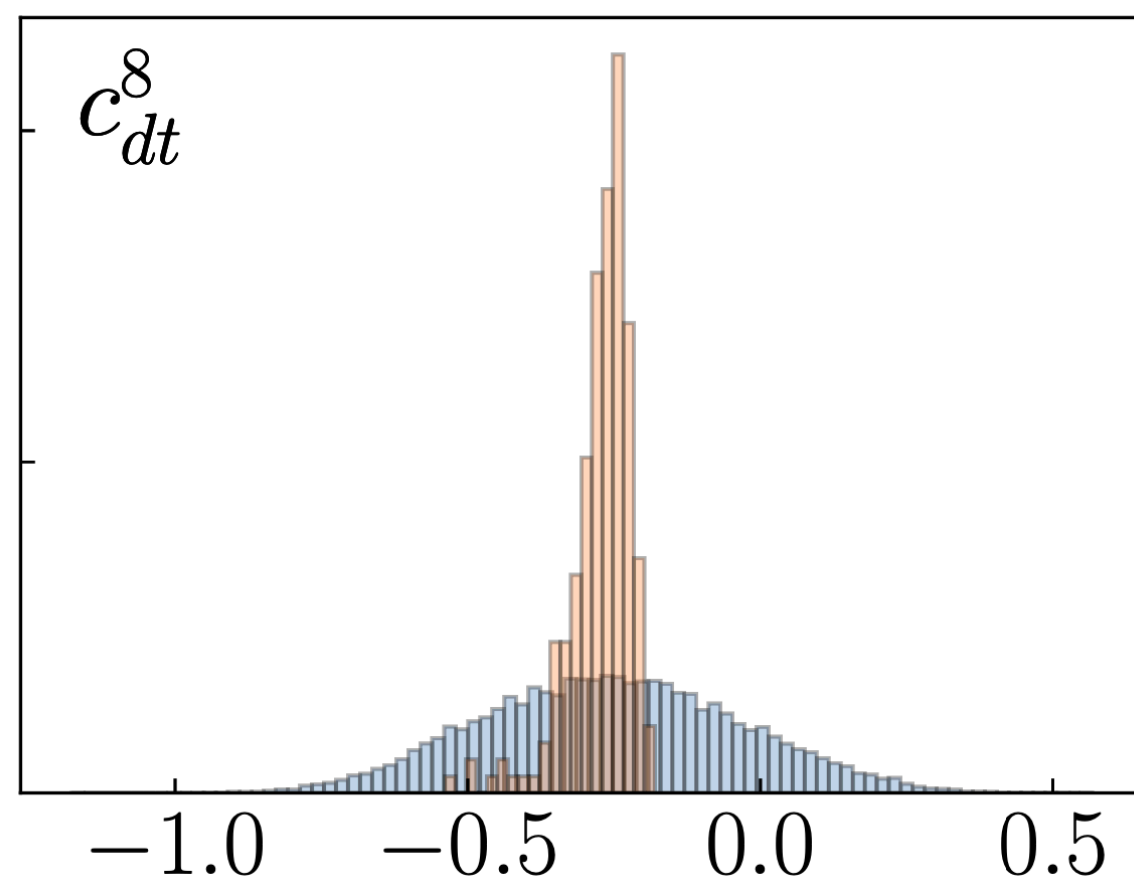


The talk in a nutshell...

1. Why study the Monte Carlo replica method?

2. A geometric approach to the MC replica method

3. Relevance in PDF fits



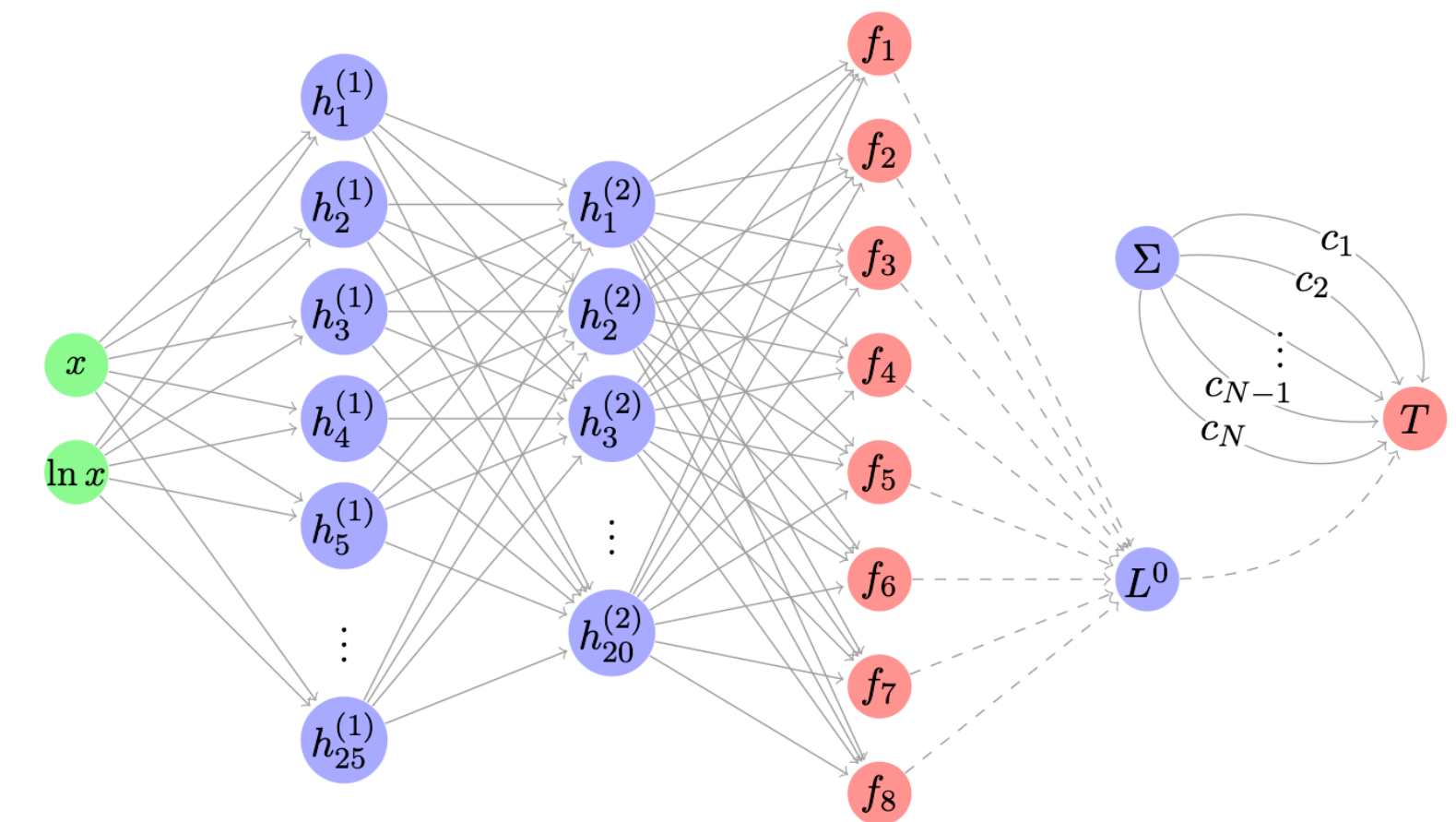
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New Physics in the top sector...?

- Back in the **winter of 2022**, Maria Ubiali's team in Cambridge, together with Juan Rojo, were working on a comprehensive analysis of the impact of **top data** on simultaneous PDF-EFT fits - see [2303.06159](#).

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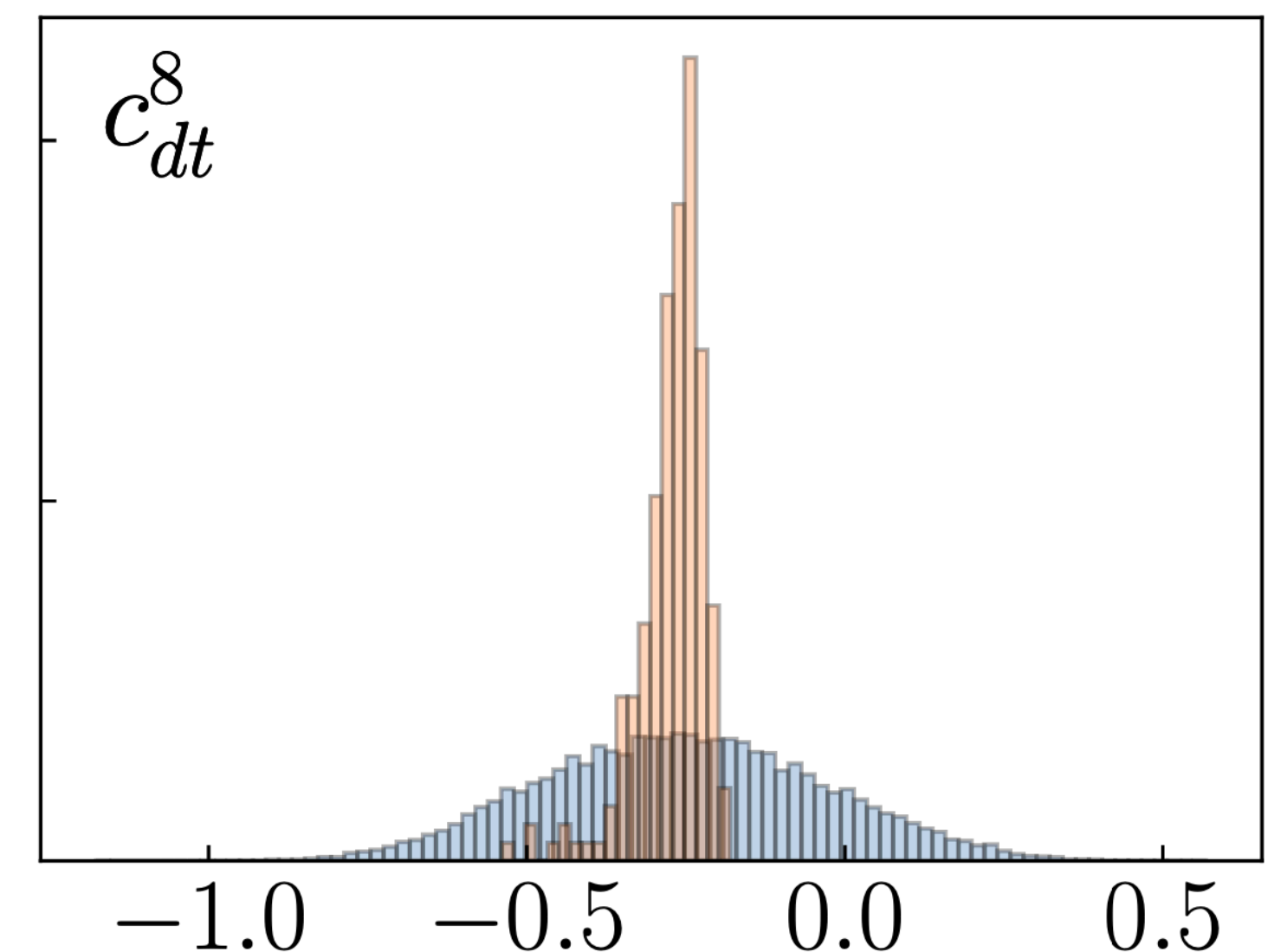
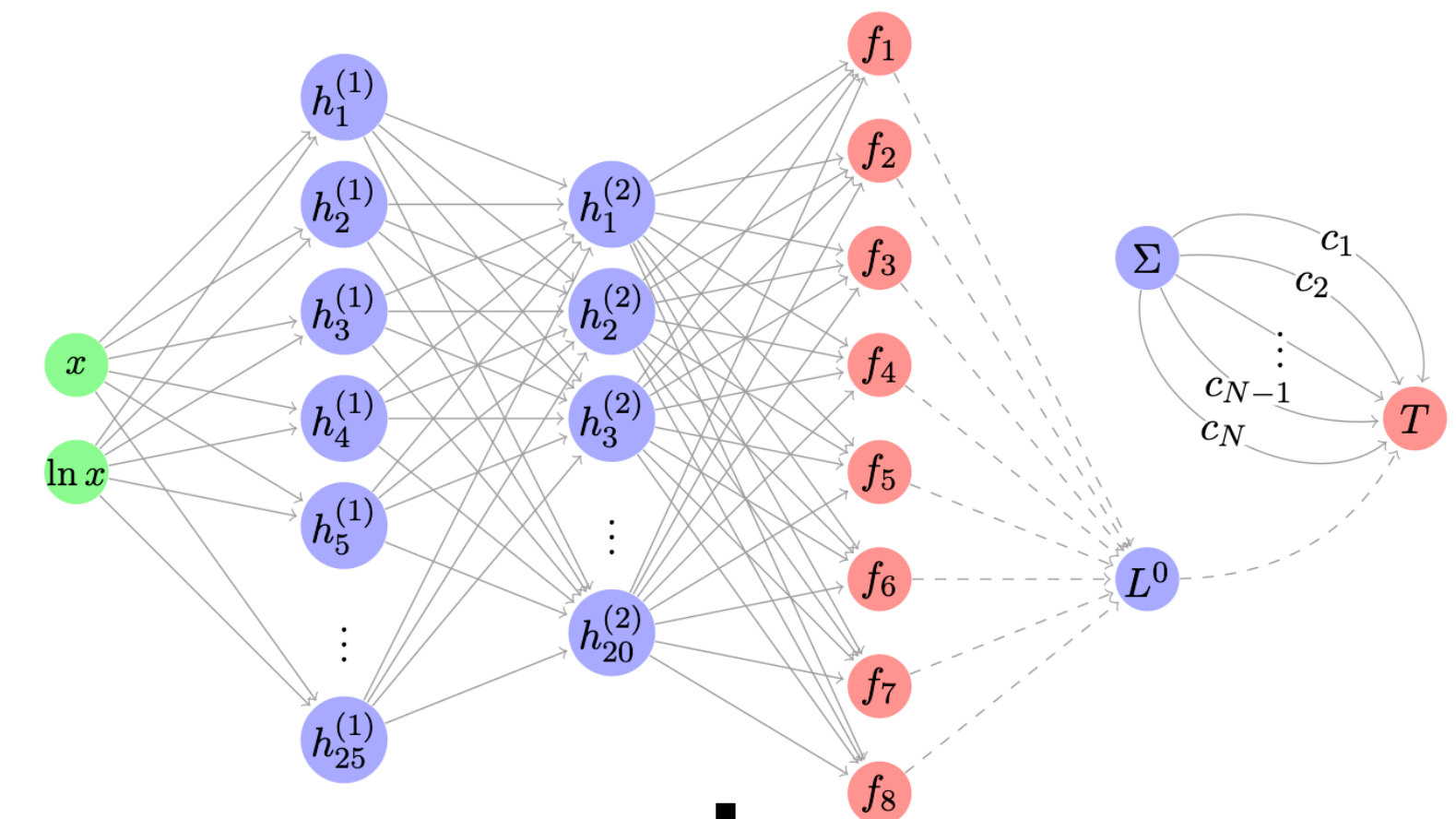
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- This study was facilitated through our fantastic new **SIMUnet code**, which is publicly available for use at: <https://hep-pbsp.github.io/SIMUnet>. **Go check it out!**



SIMUnet

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- We had a bit of a shock, though, when we ran the **quadratic SMEFT results**, and discovered **new physics at the 7σ level!**



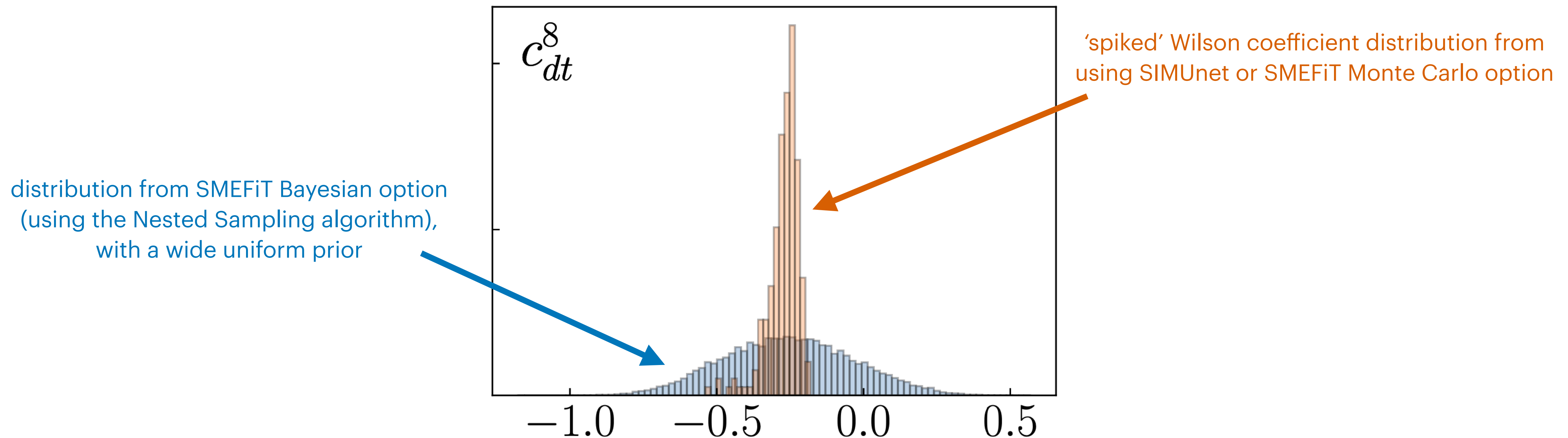
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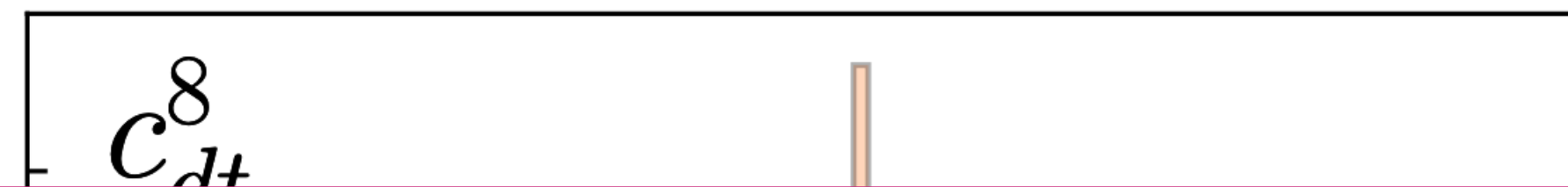
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- We realised this by **cross-checking with the SMEFiT code (2302.06660)** - we saw that **the Monte Carlo option disagreed with the Bayesian option.**



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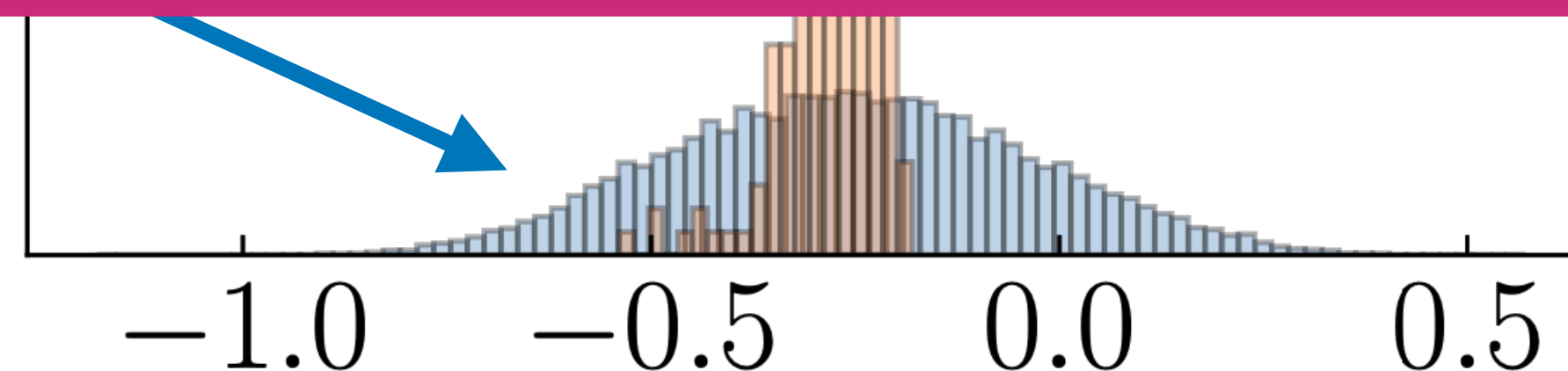
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'spiked' Wilson coefficient distribution from
SMEFiT Monte Carlo option

It appeared that using the Monte Carlo replica method to construct the SMEFT parameter distributions was leading us to make incorrect conclusions.

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- 1. Why was there a disagreement between the distributions - can this be understood mathematically?**
- 2. Could using the Monte Carlo replica method in PDF fits give different distributions, as compared to using a Bayesian methodology?**

2. - A geometric approach to the MC replica method

Bayesian vs MC analyses

- The disagreement shown in our top fit distributions motivated us to compare the Monte Carlo replica method with a fully Bayesian method on a **deeper mathematical level**. To present the results, we first set up the framework.

Bayesian vs MC analyses

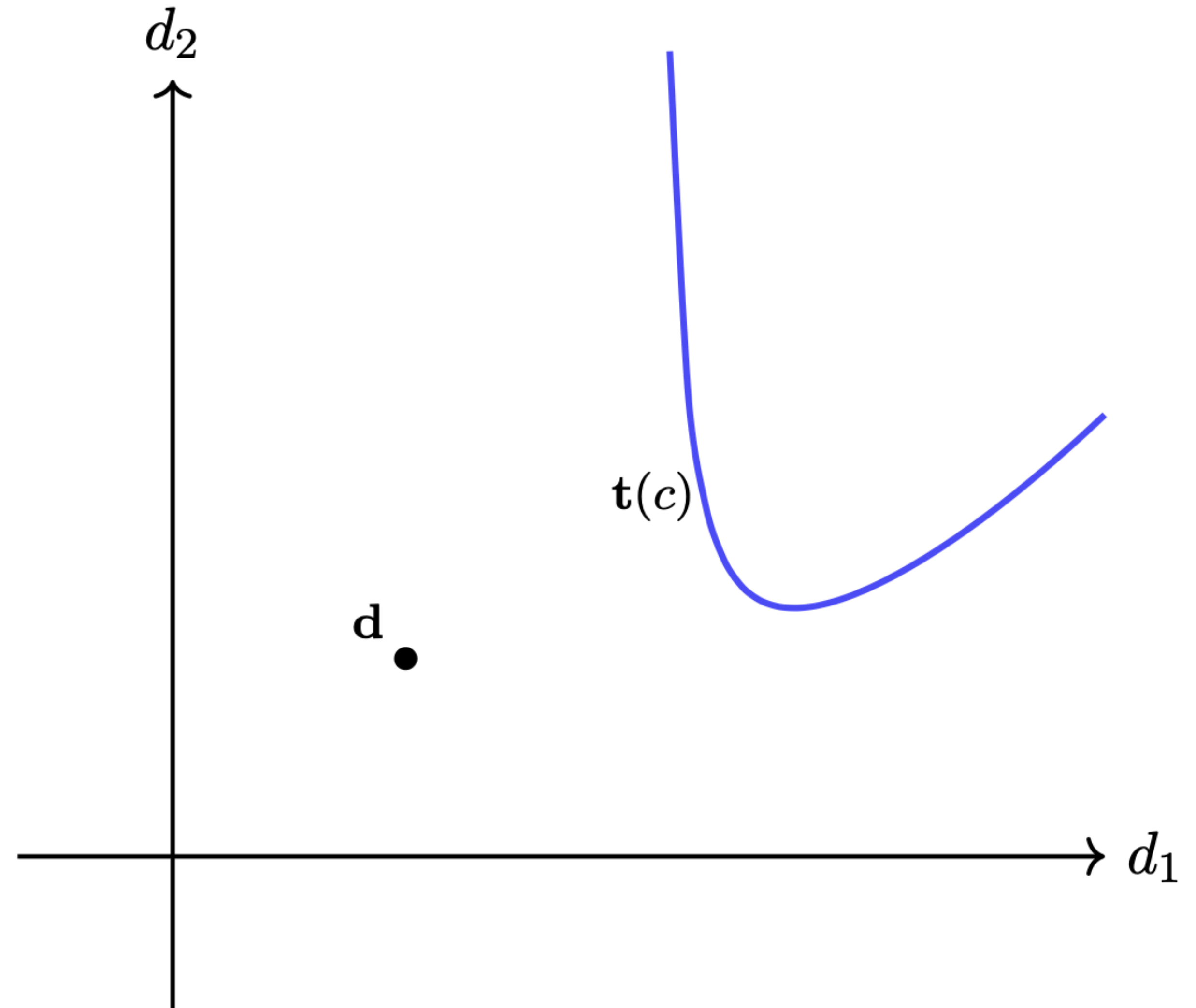
- The disagreement shown in our top fit distributions motivated us to compare the Monte Carlo replica method with a fully Bayesian method on a **deeper mathematical level**. To present the results, we first set up the framework.
- Imagine we are given a vector of experimental central data values $\mathbf{d} \in \mathbb{R}^{N_{\text{dat}}}$ by the experimentalists, together with an experimental covariance matrix Σ . We would like to compare this to our **theory predictions**:

$$\mathbf{t} : \mathbb{R}^{N_{\text{param}}} \rightarrow \mathbb{R}^{N_{\text{dat}}}$$

which can be viewed as a vector function $\mathbf{t}(\mathbf{c})$ of some parameters $\mathbf{c} \in \mathbb{R}^{N_{\text{param}}}$. The parameters in SMEFT fits are **Wilson coefficients**, and in PDF fits they are the **parameters of the PDF model under consideration**.

Bayesian vs MC analyses

- If the number of parameters is smaller than the number of data, we can view the theory function $\mathbf{t}(\mathbf{c})$ as **carving out a surface in data-space** $\mathbb{R}^{N_{\text{dat}}}$.
- *Right:* an example, with **two data points** and a theory prediction (in **blue**) depending on a **single parameter** c . The **observed data** is at the point **d**.



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where $\pi(\mathbf{c})$ is some **prior distribution**, and we use the fact that the experimental data is **normally distributed**.

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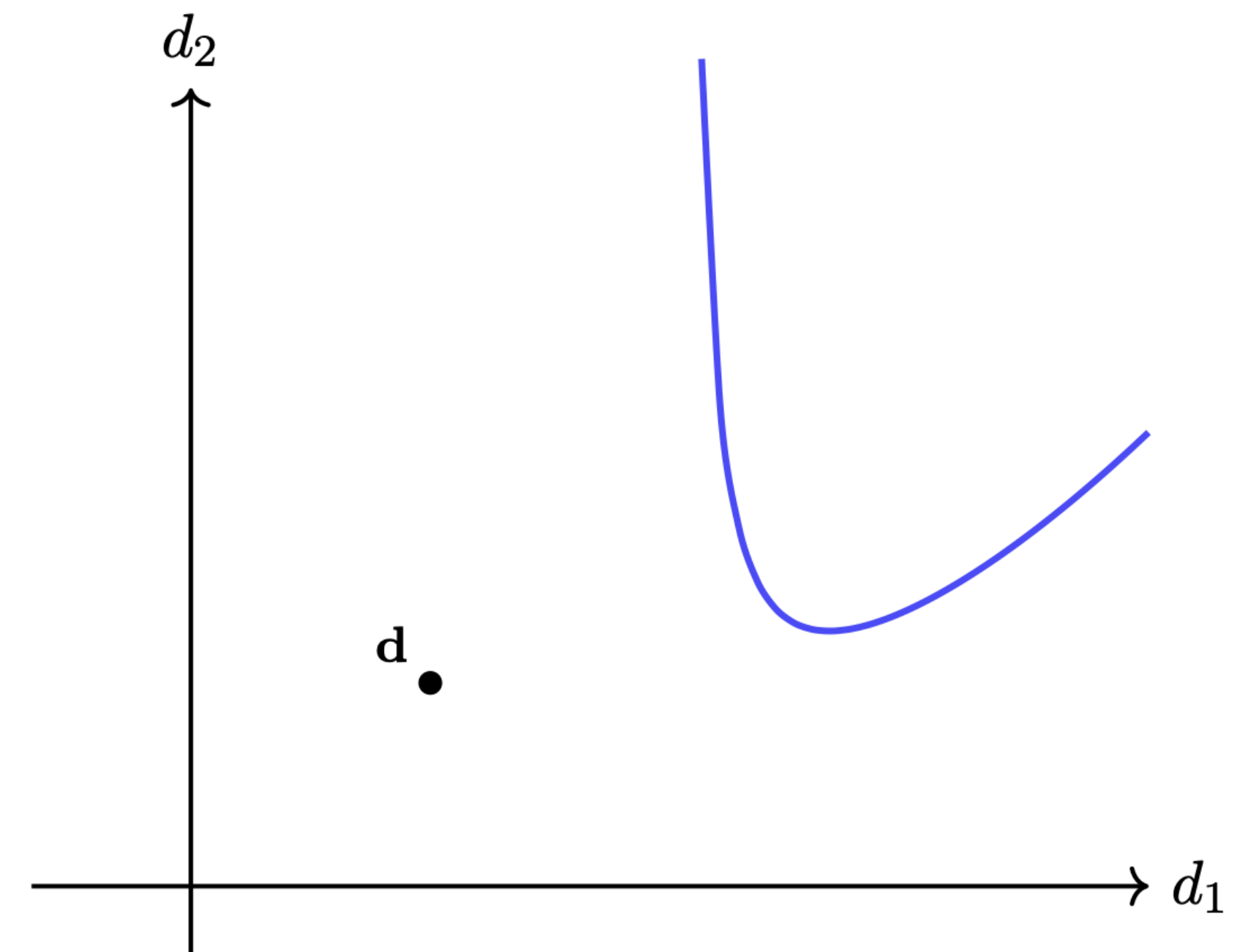
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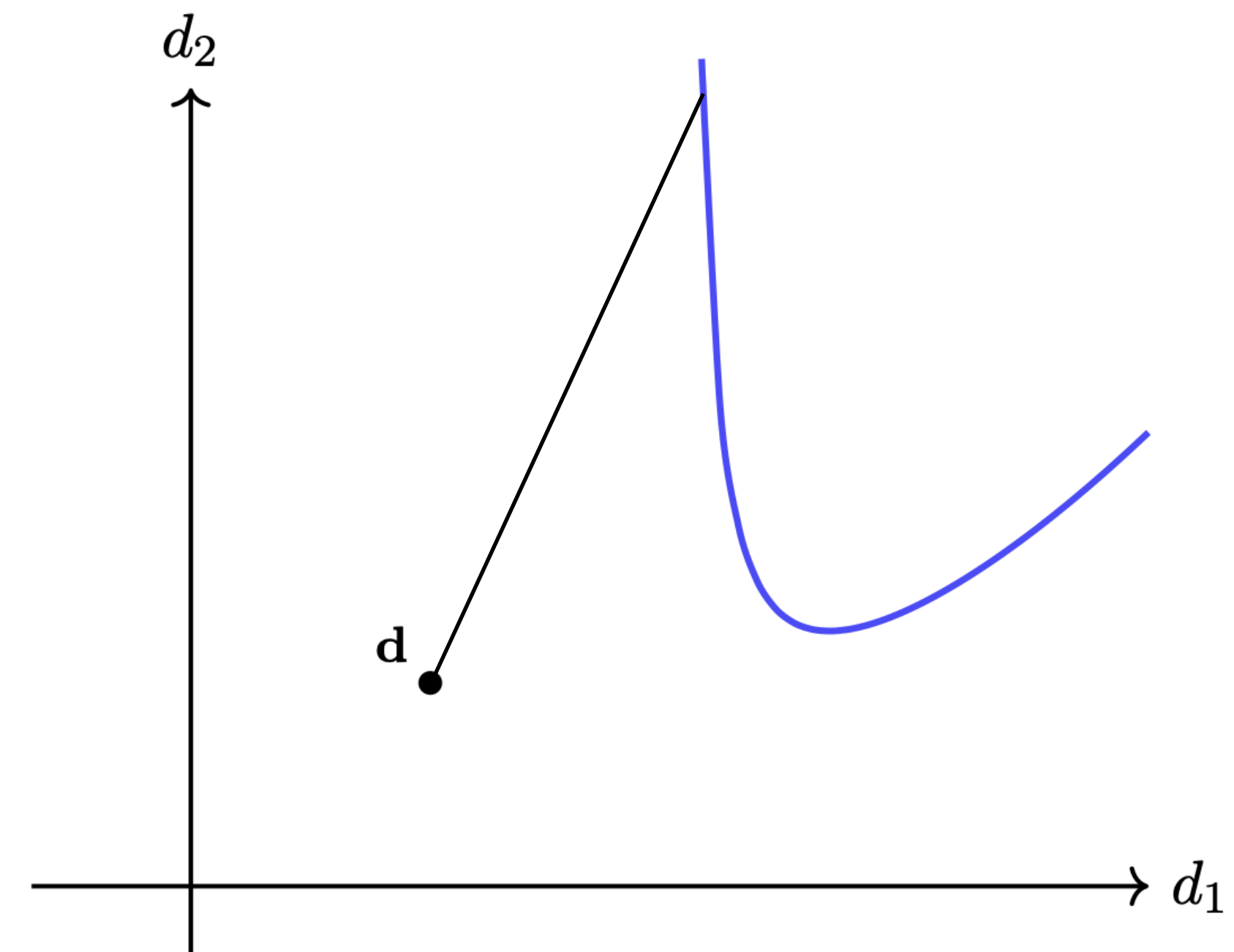
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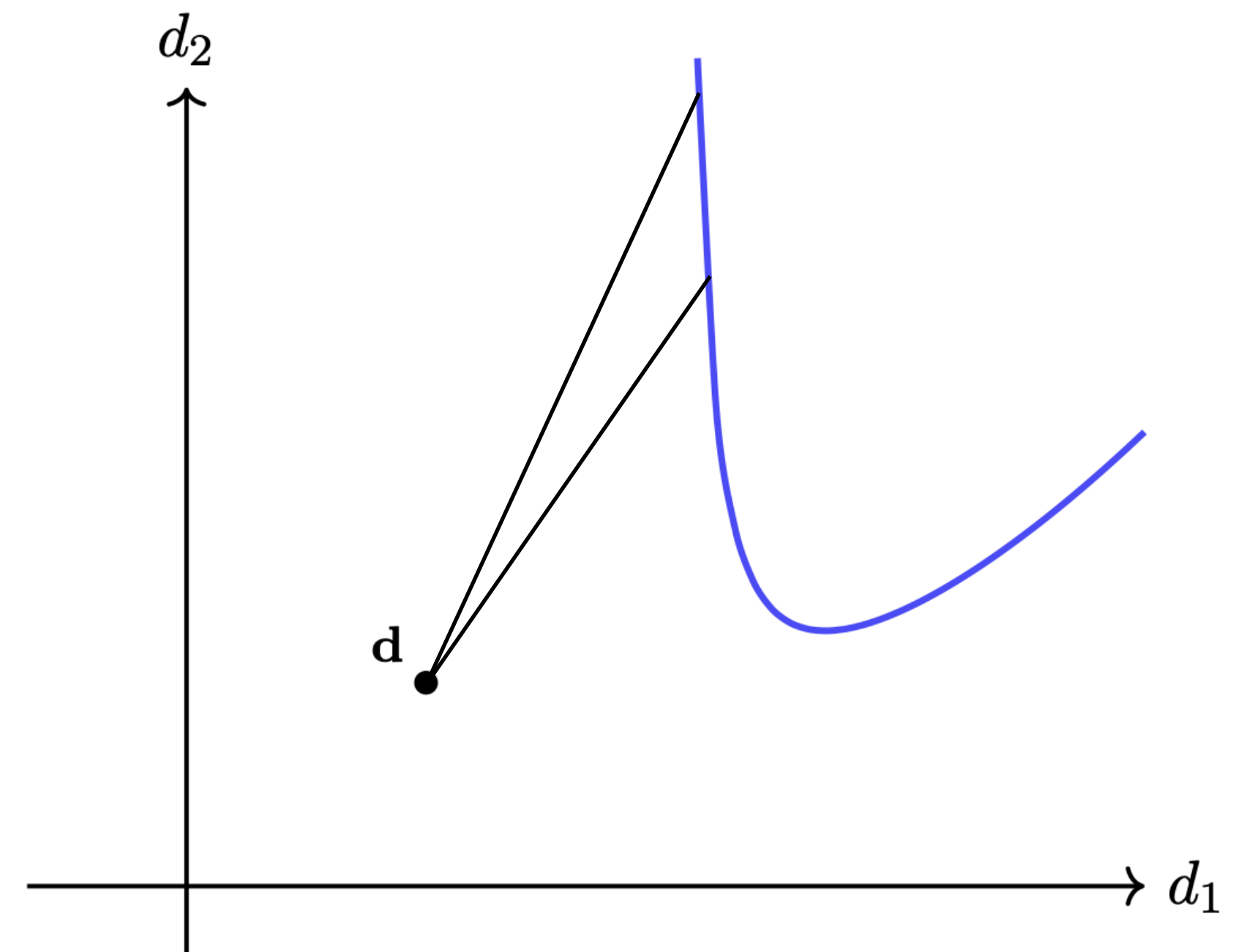
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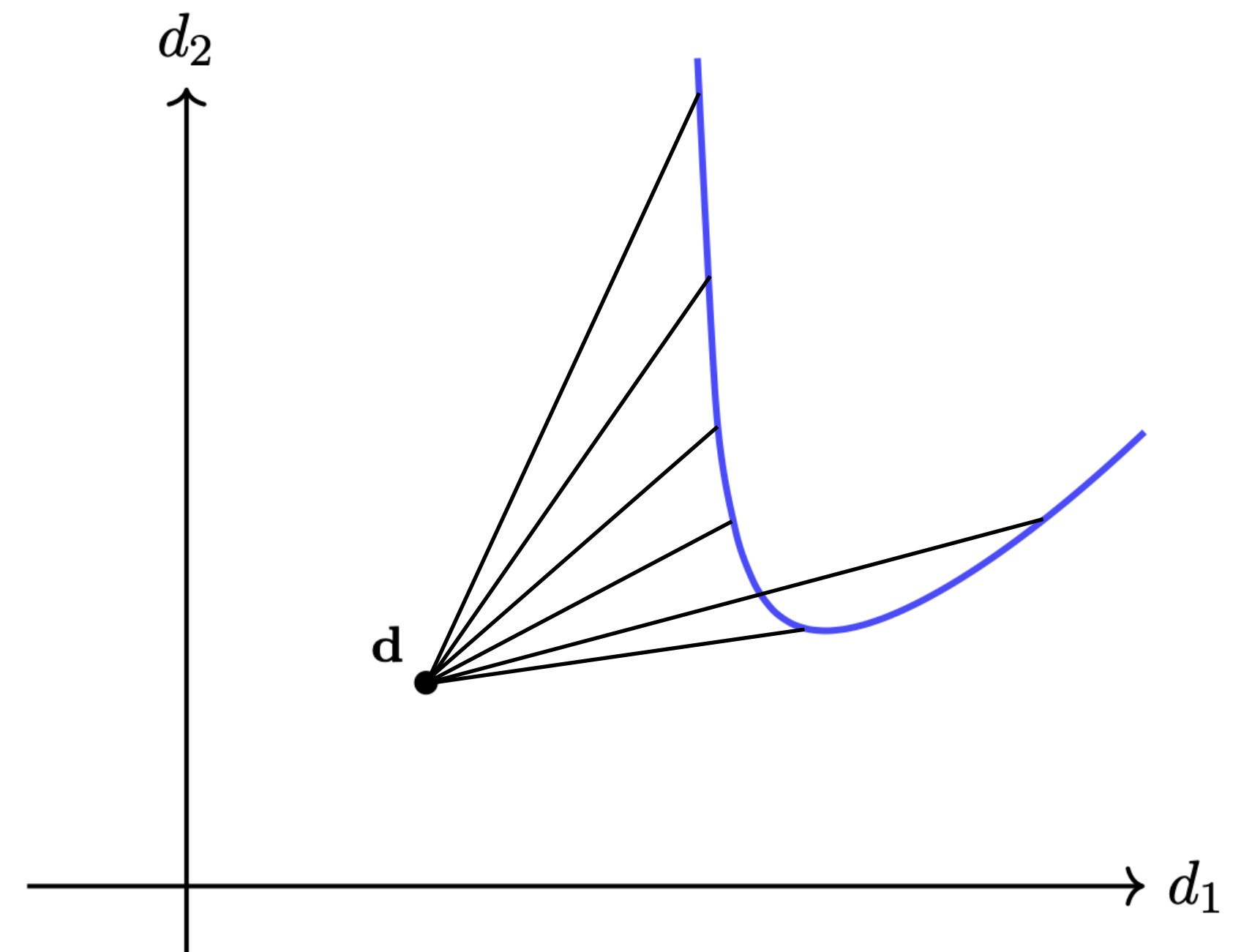
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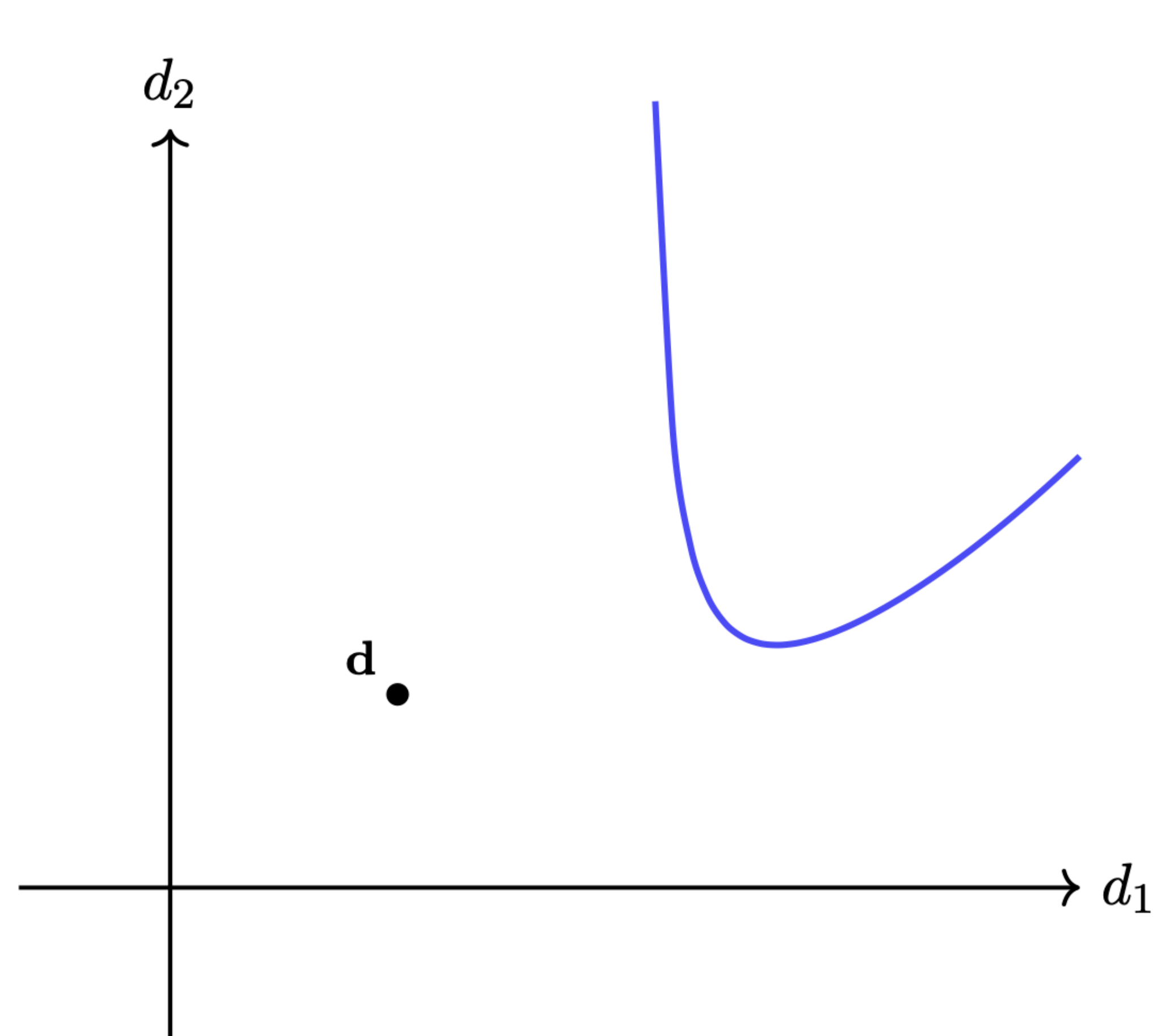
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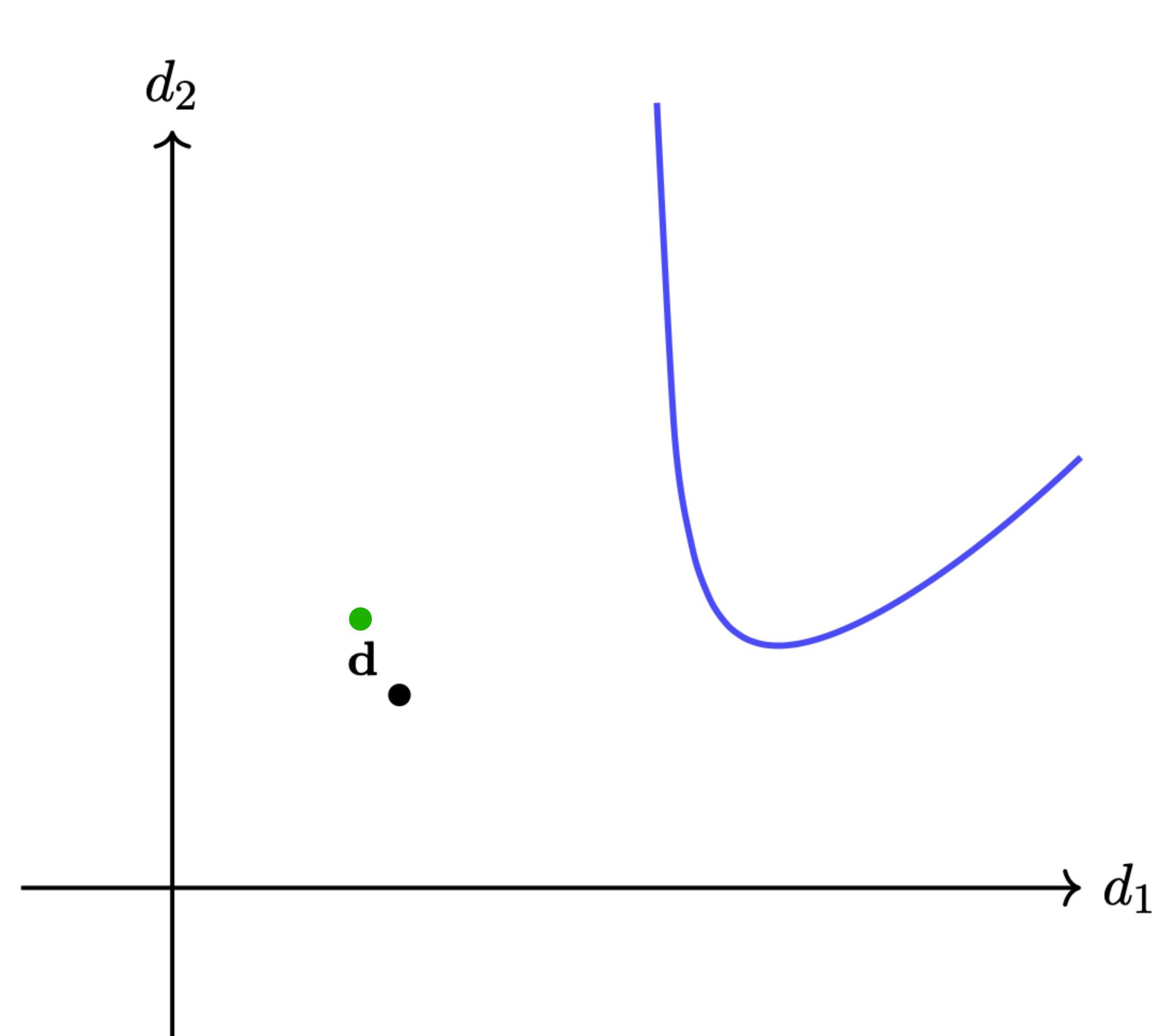


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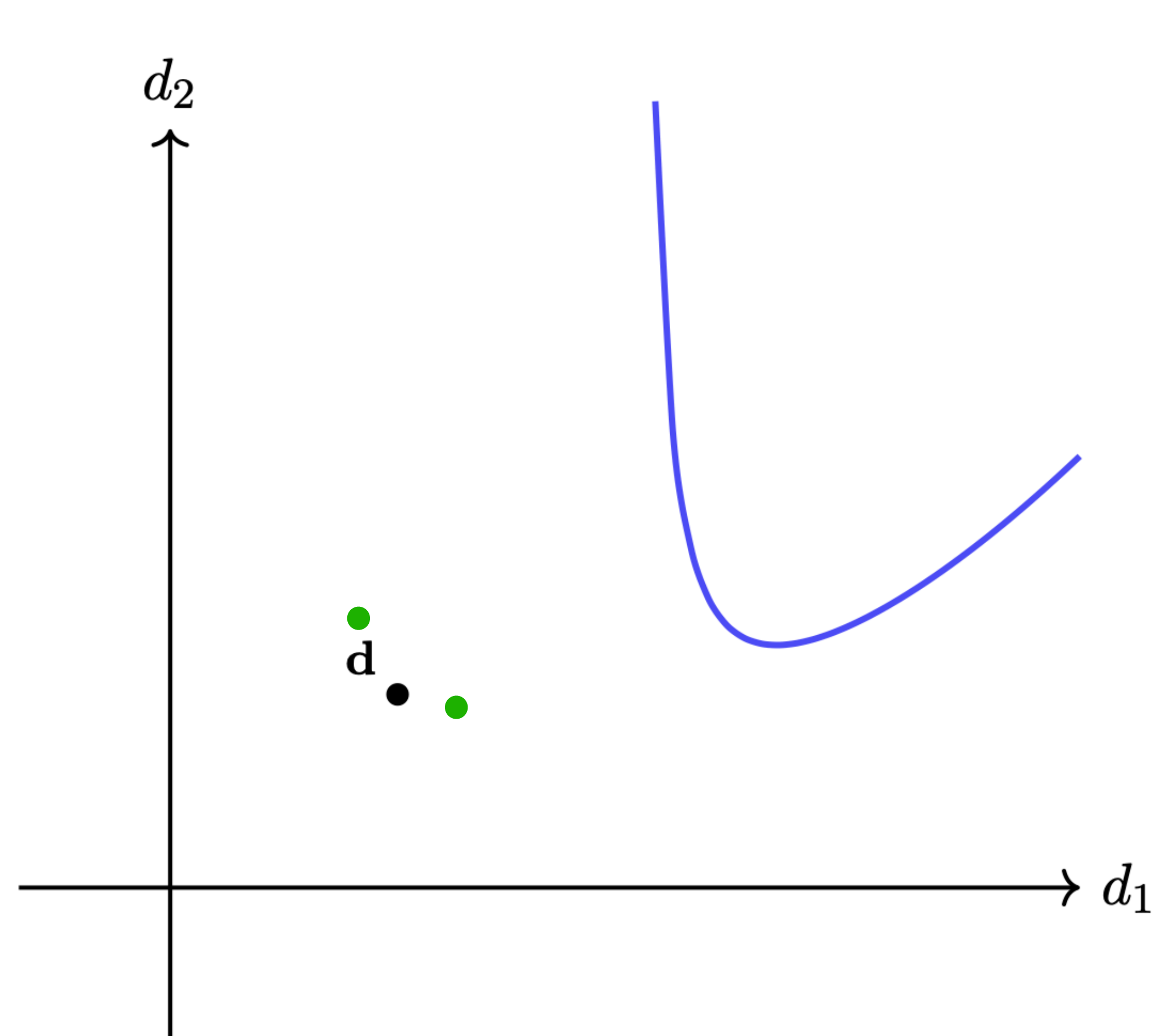


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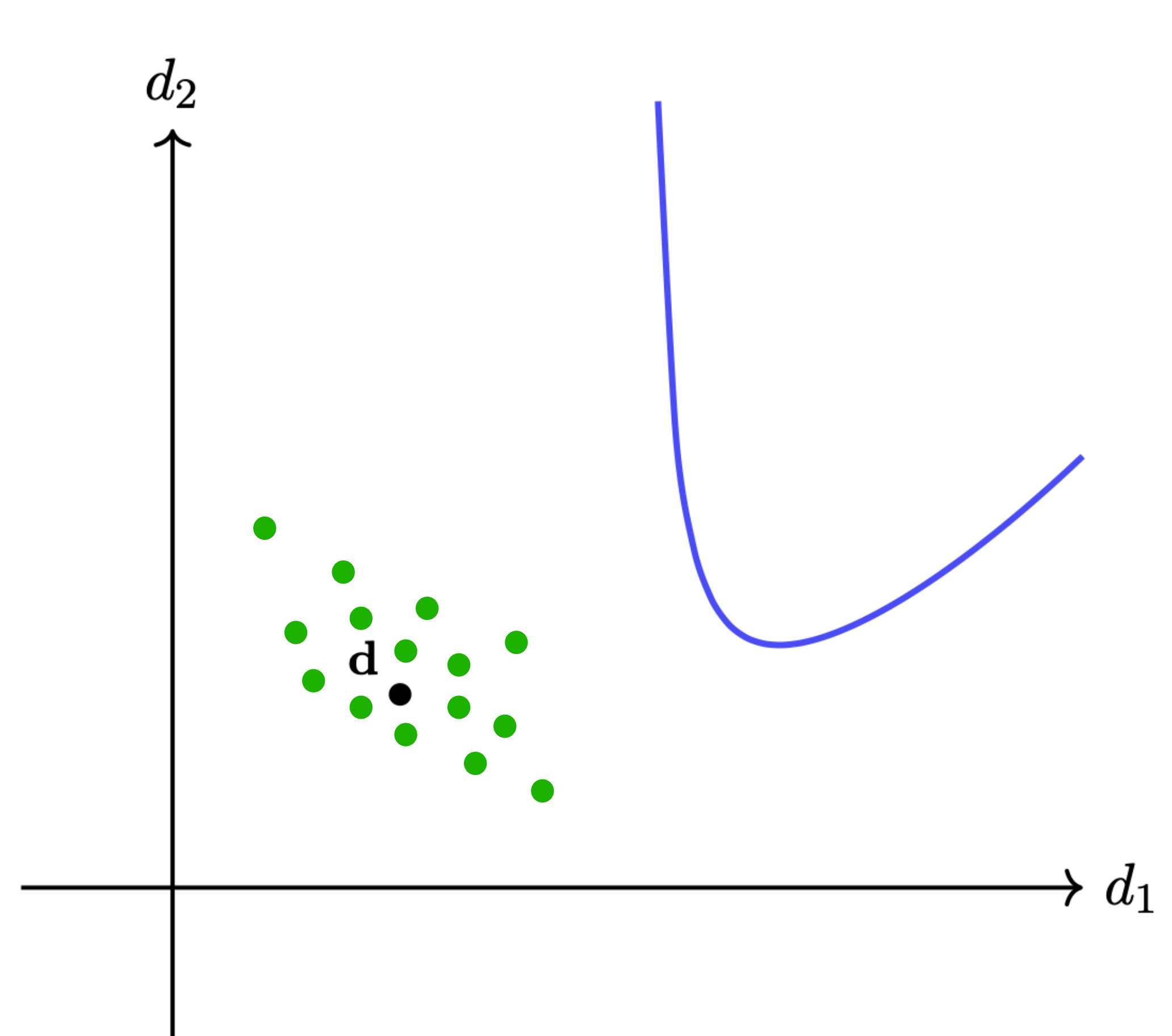


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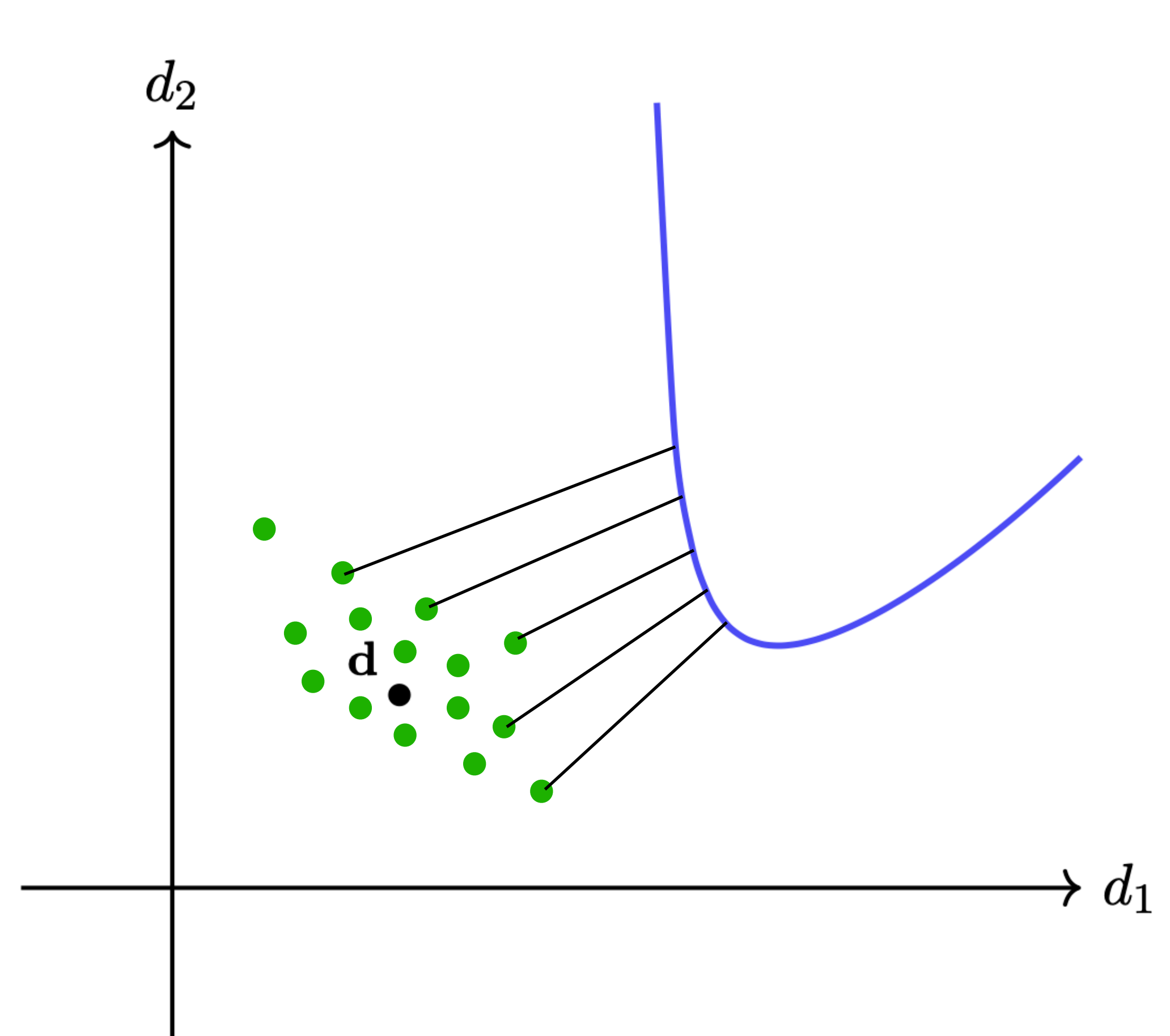


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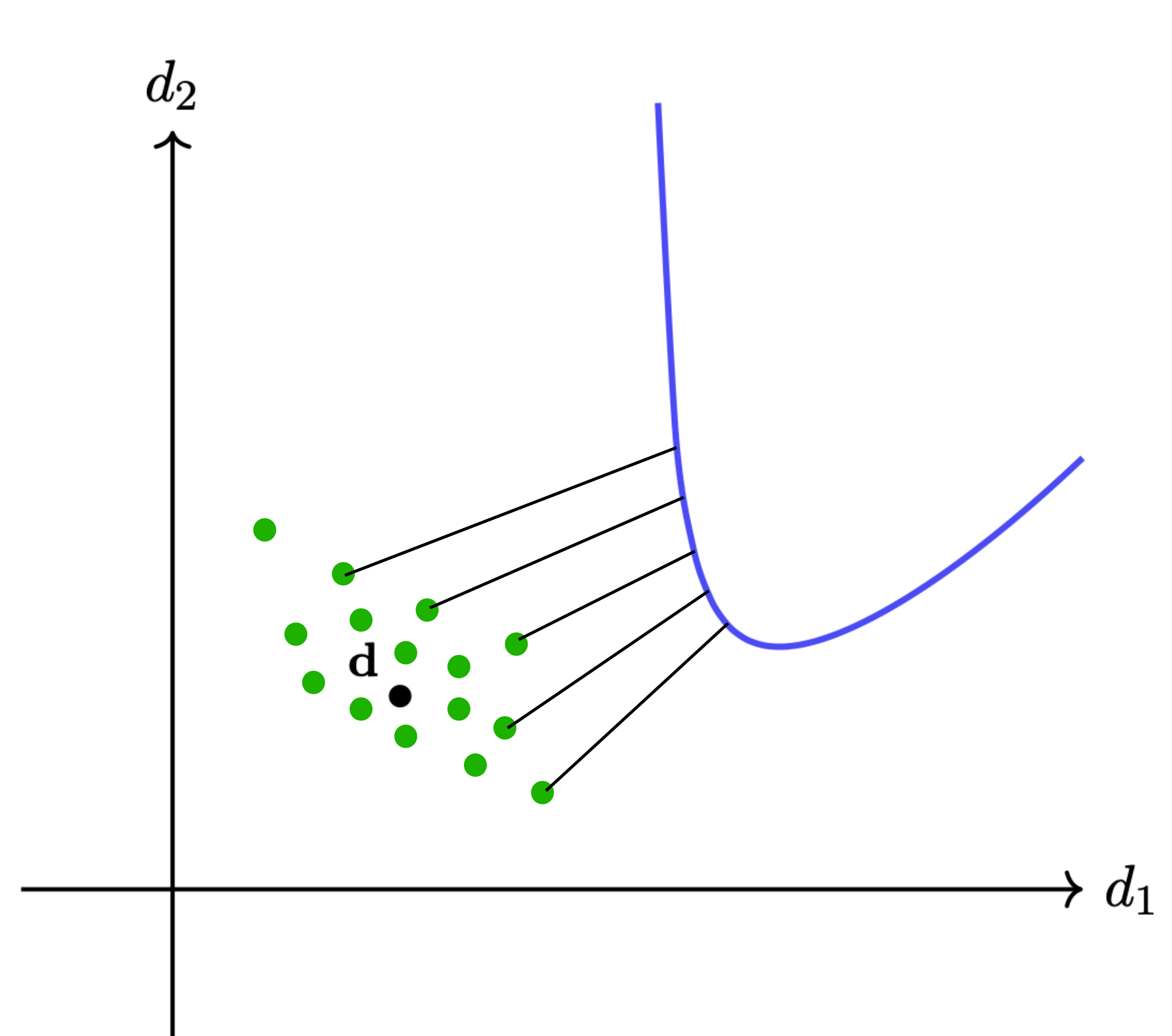


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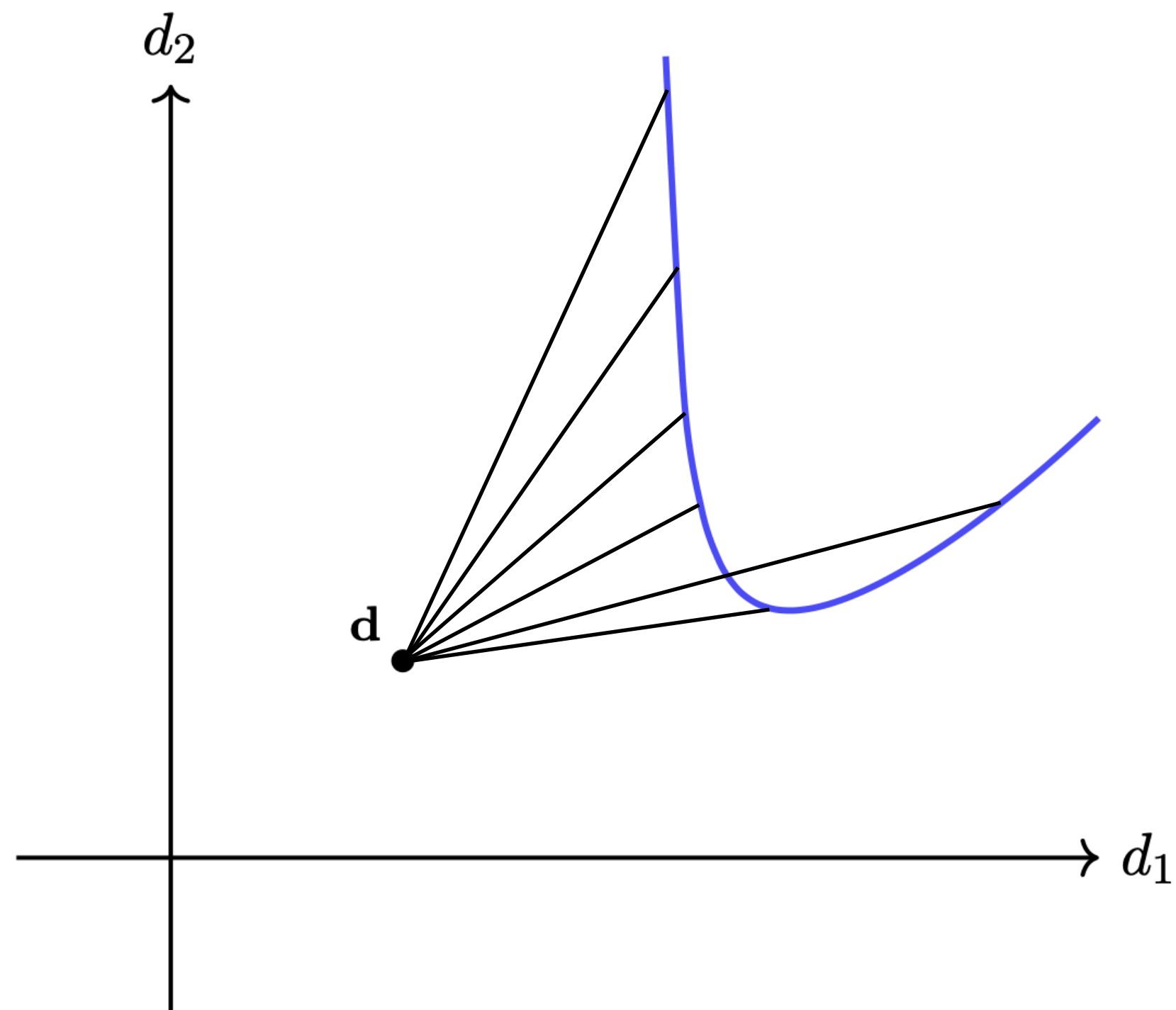
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- For each pseudodata point, we **minimise** - we compute the **closest point** on the theory surface (in the Σ -distance), and thus obtain **associated parameter values**.
- Repeating for **large amounts of pseudodata** gives an **approximation** to the **parameter distributions**.



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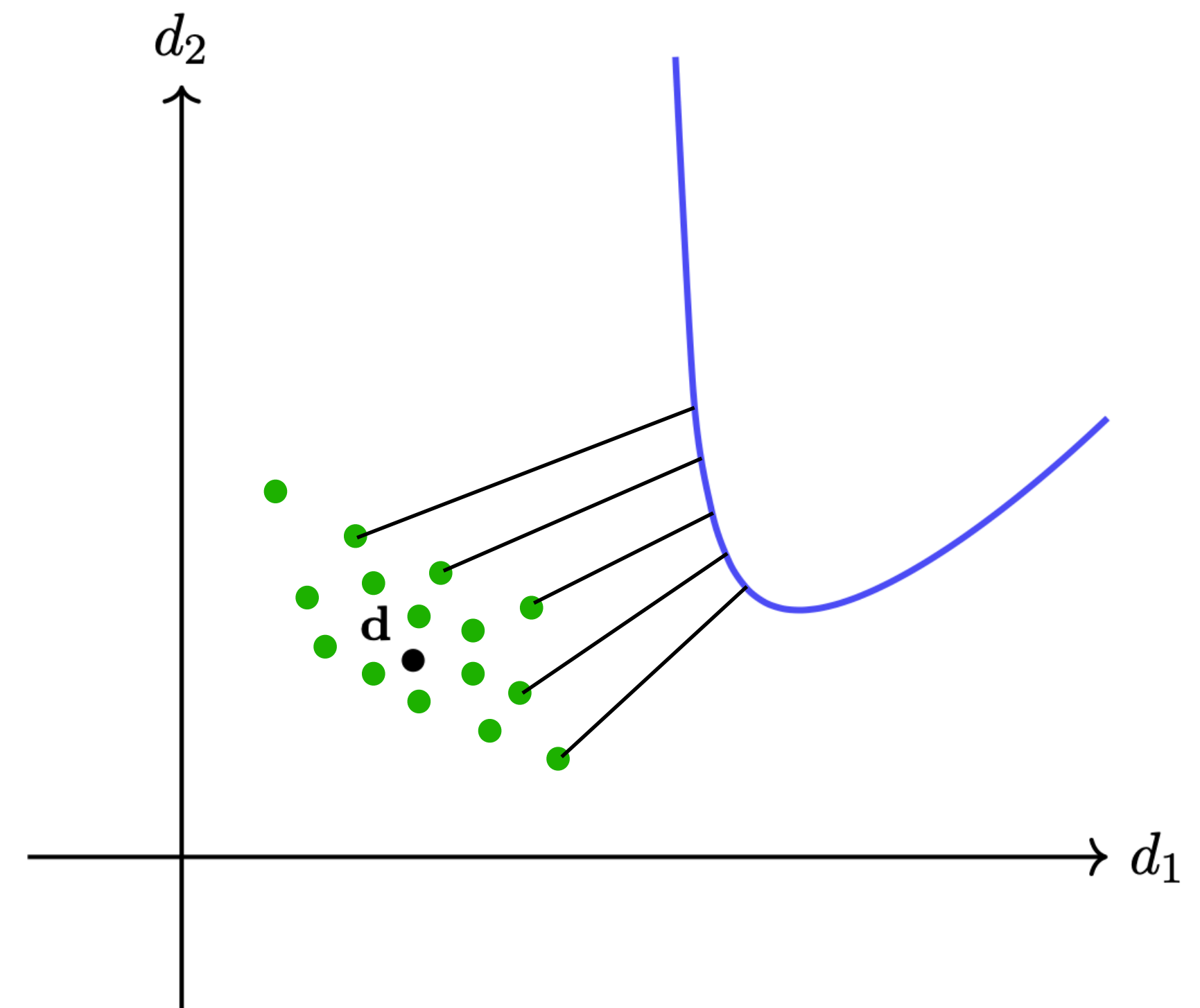
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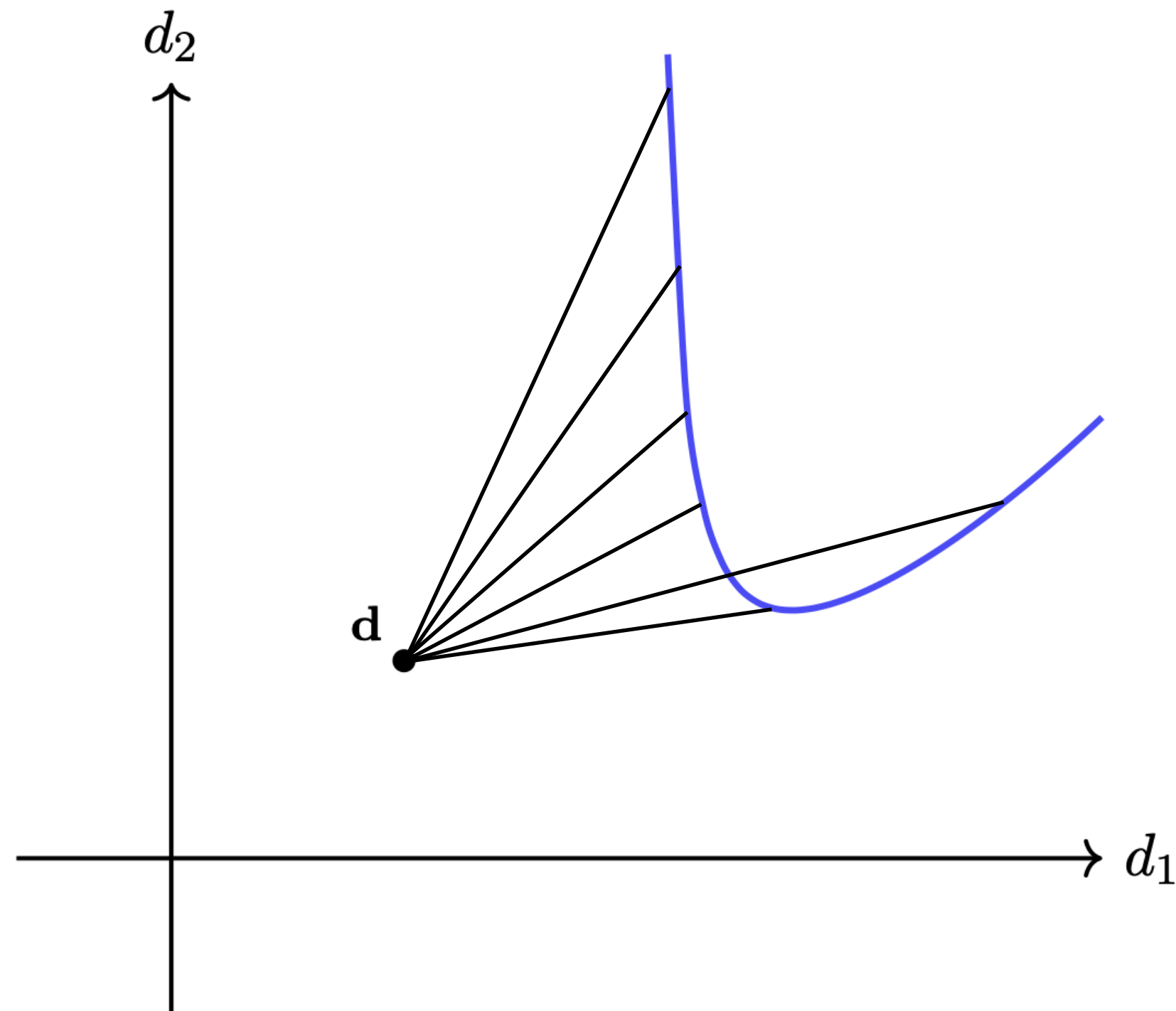
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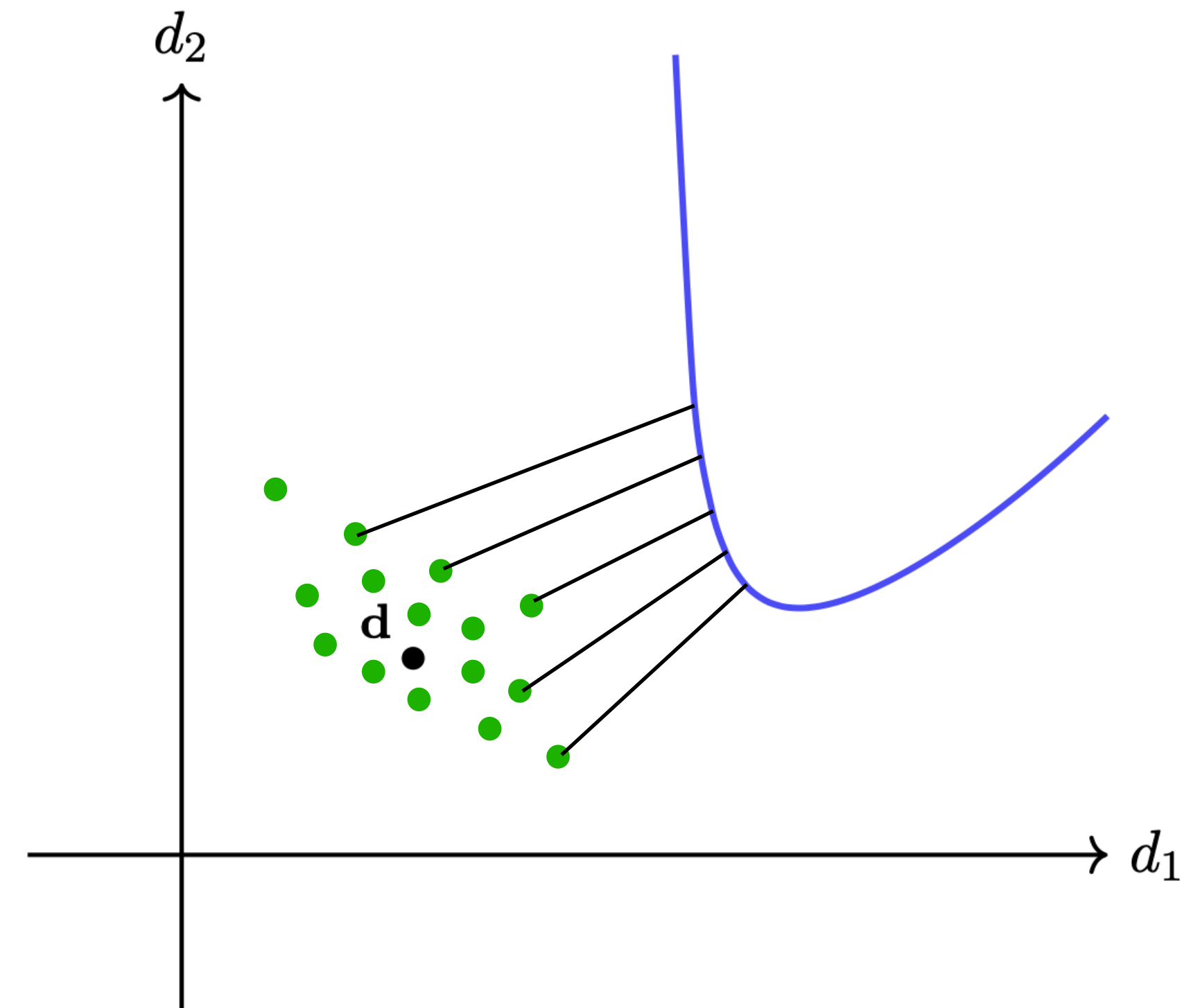
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MC replica method

d_2

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Note:

(1) It is not obvious that these methods agree.

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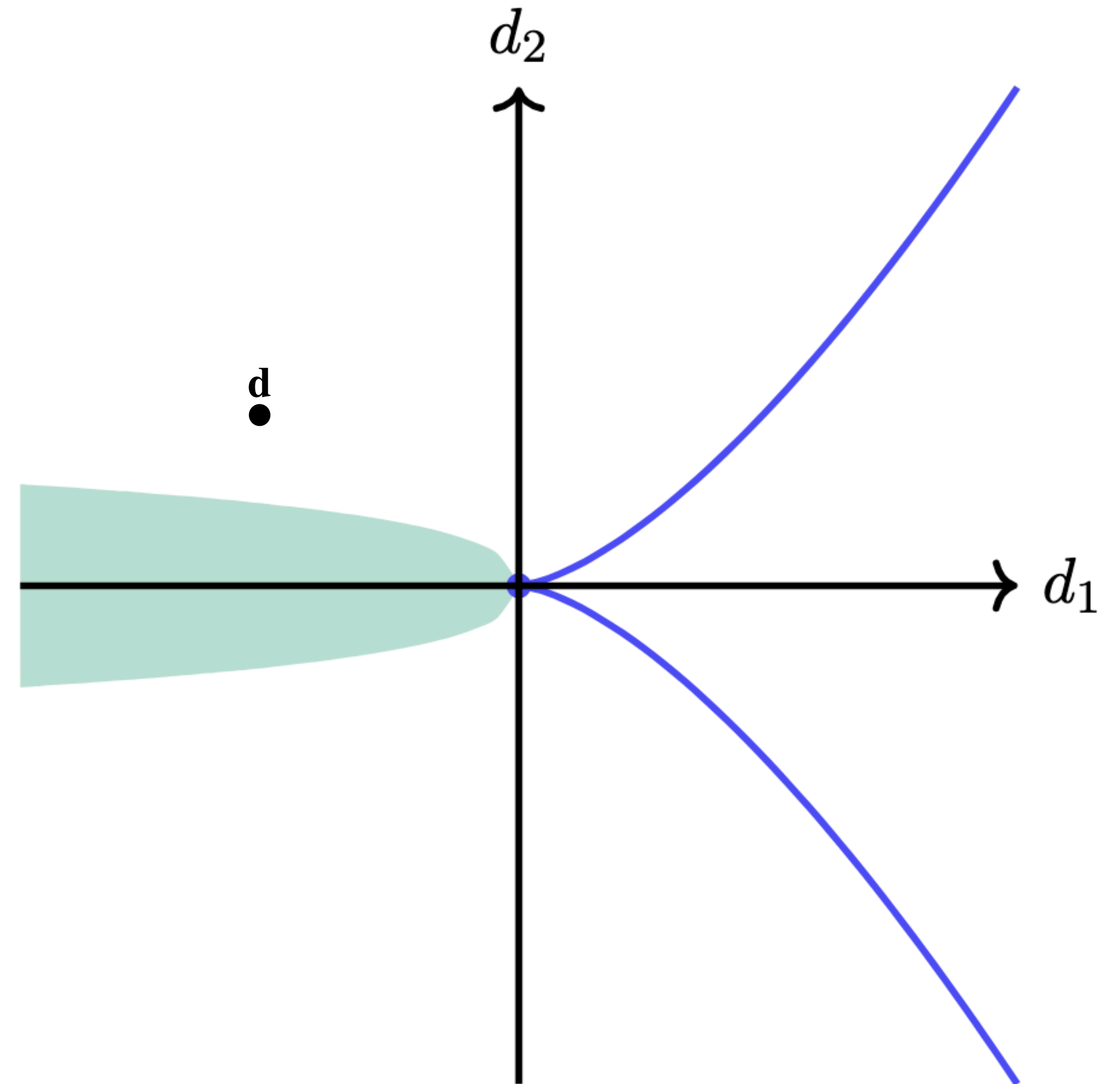
- It is not obvious that these methods agree.*
- We have not derived an explicit expression for the 'Monte Carlo posterior'.*

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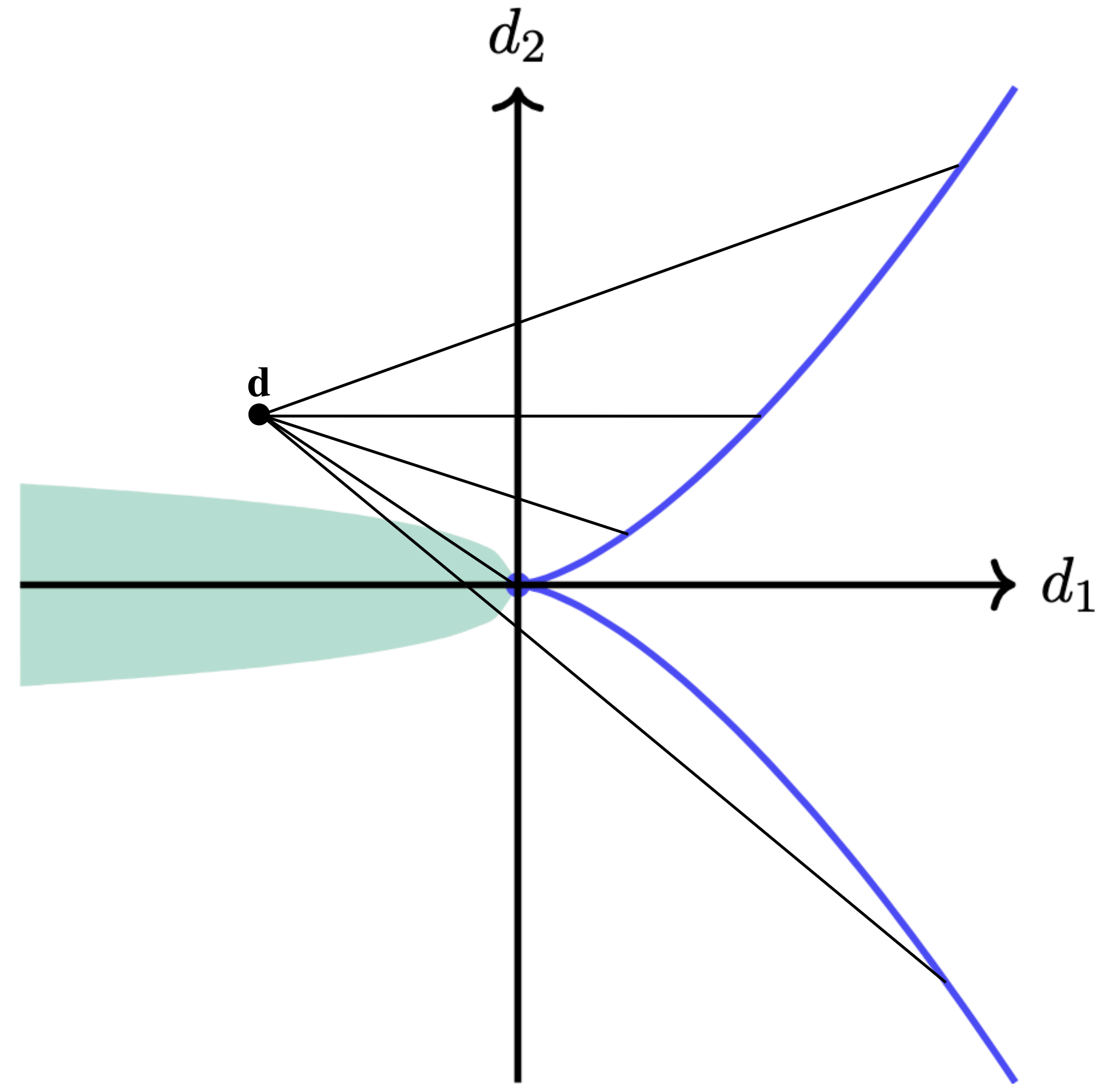
(1) Where might we expect disagreement?

- This geometric understanding helped us to see that we might expect the methods to disagree near a point of **high curvature** on a theory surface.
- On the right, we show the **non-linear theory surface** $\mathbf{t}(c) = (c^2, c^3)^T$ in blue. The green region is the **set of all points** whose **closest point** on the theory surface is the **origin**.



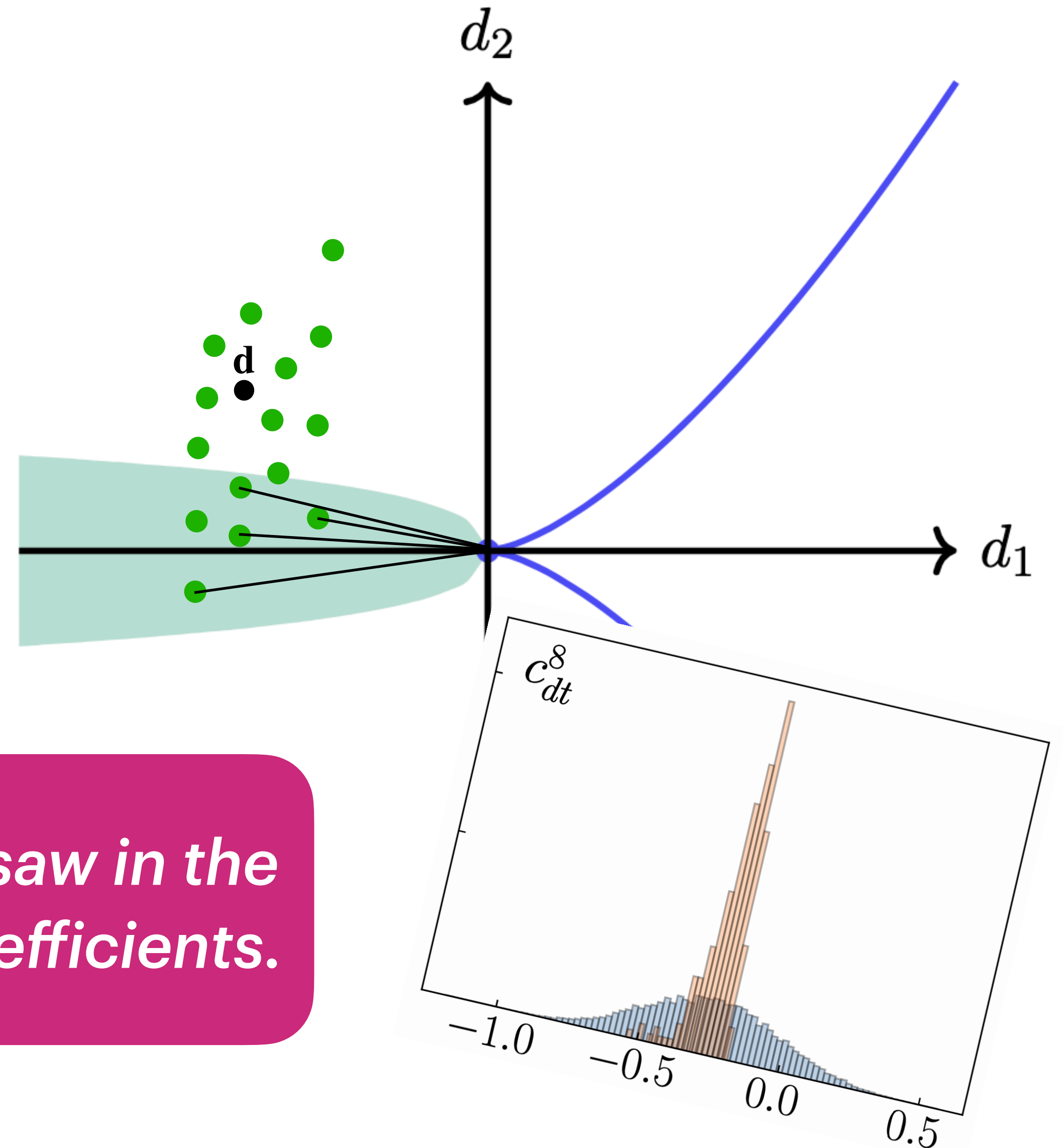
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(1) Where might we expect disagreement?

- If we use the **Bayesian method** to analyse this problem, points near the cusp are treated **like any other points**.
- On the other hand, if we throw pseudodata near **\mathbf{d}** , some proportion enters the **green 'basin of attraction'**, and is unfairly **drawn towards the cusp**.



This is the origin of the 'spiked peaks' we saw in the Monte Carlo distributions of the Wilson coefficients.

(2) The Monte Carlo posterior

- Examples such as the one we have just seen are **characteristic of the general behaviour** of the **Monte Carlo posterior**, which was derived **explicitly** in our paper. The maths is hard, and the result is not easy to understand either:

$$\begin{aligned}
 & \exp\left(-\frac{1}{2}\chi_{\mathbf{d}_0}^2(\mathbf{c})\right) \\
 & \cdot \int d^{N_{\parallel}(\mathbf{c})}\mathbf{u} \delta(\mathbf{c} - \mathbf{f}(\mathbf{u})) \int_{\Lambda(\mathbf{c})} d^{N_{\perp}(\mathbf{c})}\boldsymbol{\lambda} \left| \det\left(\frac{\partial \mathbf{t}}{\partial \mathbf{c}}(\mathbf{f}(\mathbf{u}))\frac{\partial \mathbf{f}}{\partial \mathbf{u}} + \frac{\partial(\Sigma M \boldsymbol{\lambda})}{\partial \mathbf{c}}(\mathbf{f}(\mathbf{u}))\frac{\partial \mathbf{f}}{\partial \mathbf{u}} \Big|_{\Sigma M(\mathbf{f}(\mathbf{u}))}\right) \right| \\
 & \cdot \exp\left(-\frac{1}{2}\boldsymbol{\lambda}^T M(\mathbf{c})^T \Sigma M(\mathbf{c}) \boldsymbol{\lambda} + \boldsymbol{\lambda}^T M(\mathbf{c})^T (\mathbf{d}_0 - \mathbf{t}(\mathbf{c}))\right), \tag{2.17}
 \end{aligned}$$

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Bayesian result, for uniform flat prior

$$\exp\left(-\frac{1}{2}\chi_{\mathbf{d}_0}^2(\mathbf{c})\right)$$

Jacobian 'transformation factor' which is enhanced in regions of high curvature

$$\int d^{N_{\parallel}(\mathbf{c})} \mathbf{u} \delta(\mathbf{c} - \mathbf{f}(\mathbf{u})) \int_{\Lambda(\mathbf{c})} d^{N_{\perp}(\mathbf{c})} \boldsymbol{\lambda} \left| \det \left(\frac{\partial \mathbf{t}}{\partial \mathbf{c}}(\mathbf{f}(\mathbf{u})) \frac{\partial \mathbf{f}}{\partial \mathbf{u}} + \frac{\partial (\Sigma M \boldsymbol{\lambda})}{\partial \mathbf{c}}(\mathbf{f}(\mathbf{u})) \frac{\partial \mathbf{f}}{\partial \mathbf{u}} \middle| \Sigma M(\mathbf{f}(\mathbf{u})) \right) \right|$$

delta functions, which cannot be integrated out in case of singular points, like cusp

$$\cdot \exp\left(-\frac{1}{2}\boldsymbol{\lambda}^T M(\mathbf{c})^T \Sigma M(\mathbf{c}) \boldsymbol{\lambda} + \boldsymbol{\lambda}^T M(\mathbf{c})^T (\mathbf{d}_0 - \mathbf{t}(\mathbf{c}))\right), \quad (2.17)$$

(Read the paper for the full, careful derivation: <https://arxiv.org/pdf/2404.10056>.)

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Bayesian result, for
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Key takeaway from (1) and (2):

As compared to a Bayesian method, the Monte Carlo replica method unfairly favours regions of high curvature. In particular, its validity is not guaranteed in non-linear models.

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case of singular
points, like cusp

3. - Relevance in PDF fits

PDF fitting is *non-linear*

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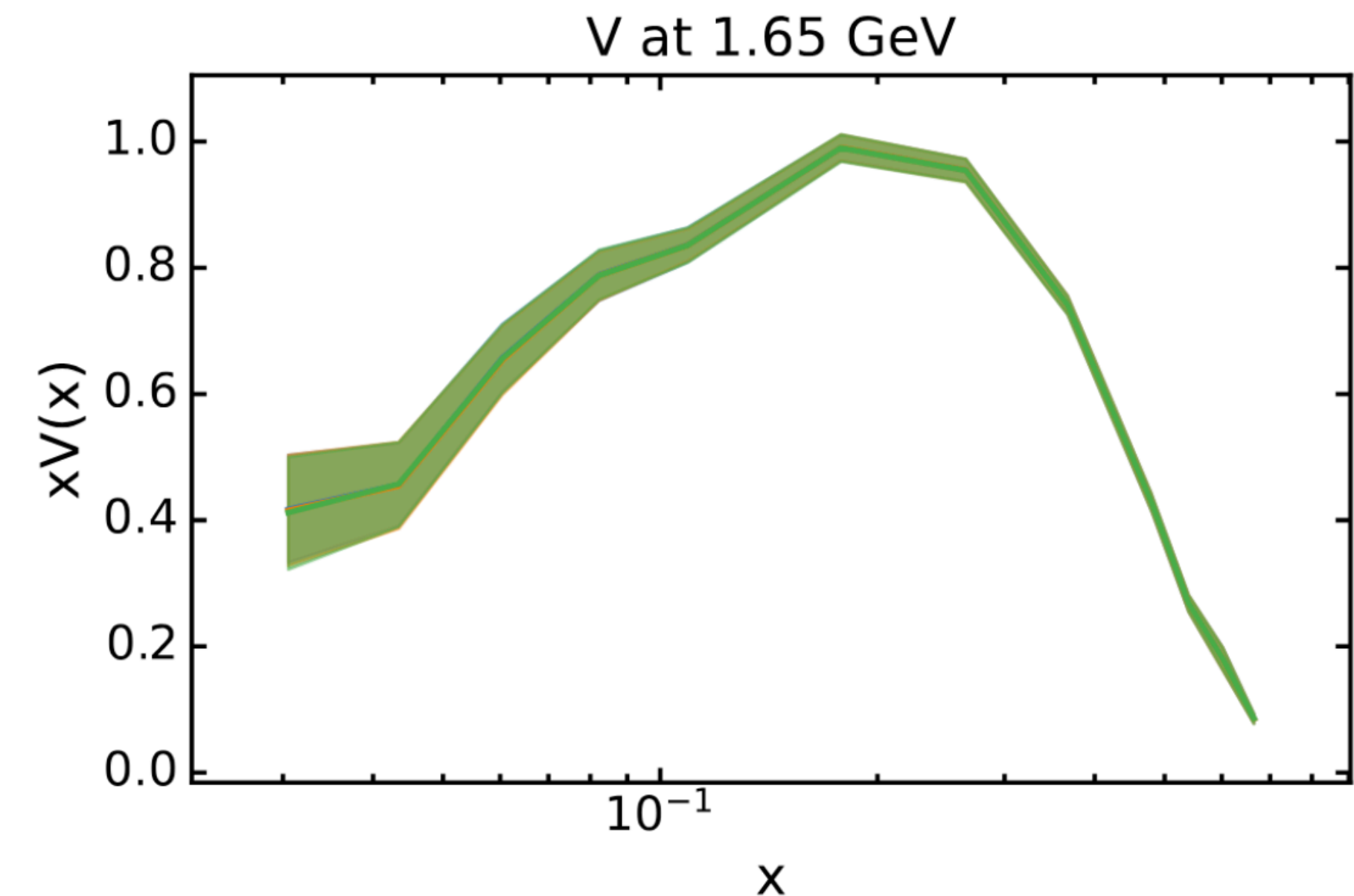
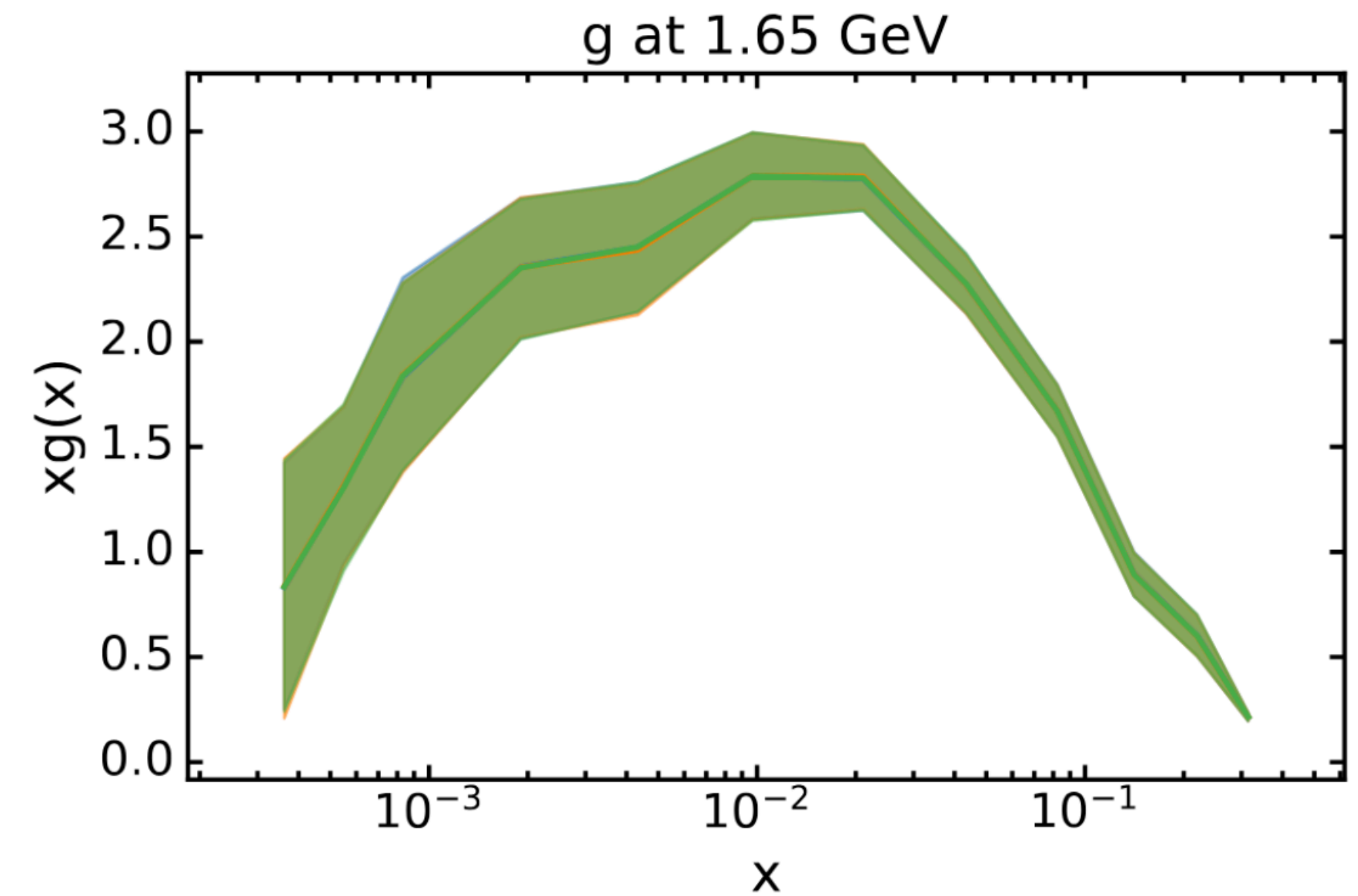
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- It is natural to ask: ***could the use of the Monte Carlo method result in incorrect conclusions about PDF uncertainties, as we add more proton-proton data?***

Fitting 'kinky' PDFs

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- We consider the proton in terms of just the **singlet, gluon** and **valence PDFs**. Further, we parametrise each of the PDFs as **linear interpolants** on an x -grid comprising **12 grid points for each flavour**, hence 36 grid points in total. Examples of our 'kinky' PDFs are shown on the right.



Fitting 'kinky' PDFs

- We generated an **artificial copy of the complete NNPDF4.0 dataset***, with noise, based on a kinky PDF with values at the grid points taken from the NNPDF4.0 central PDF.

(* excluding jets for technical reasons)

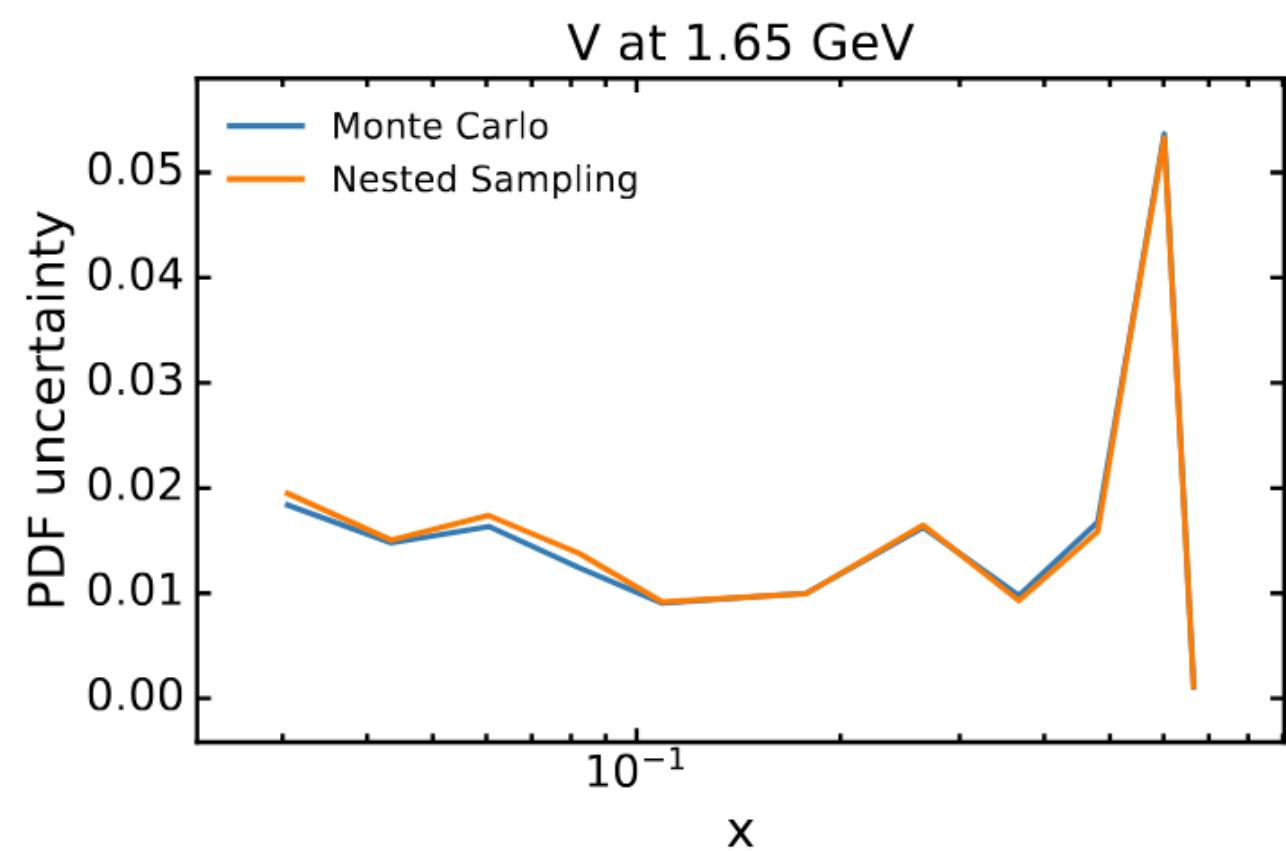
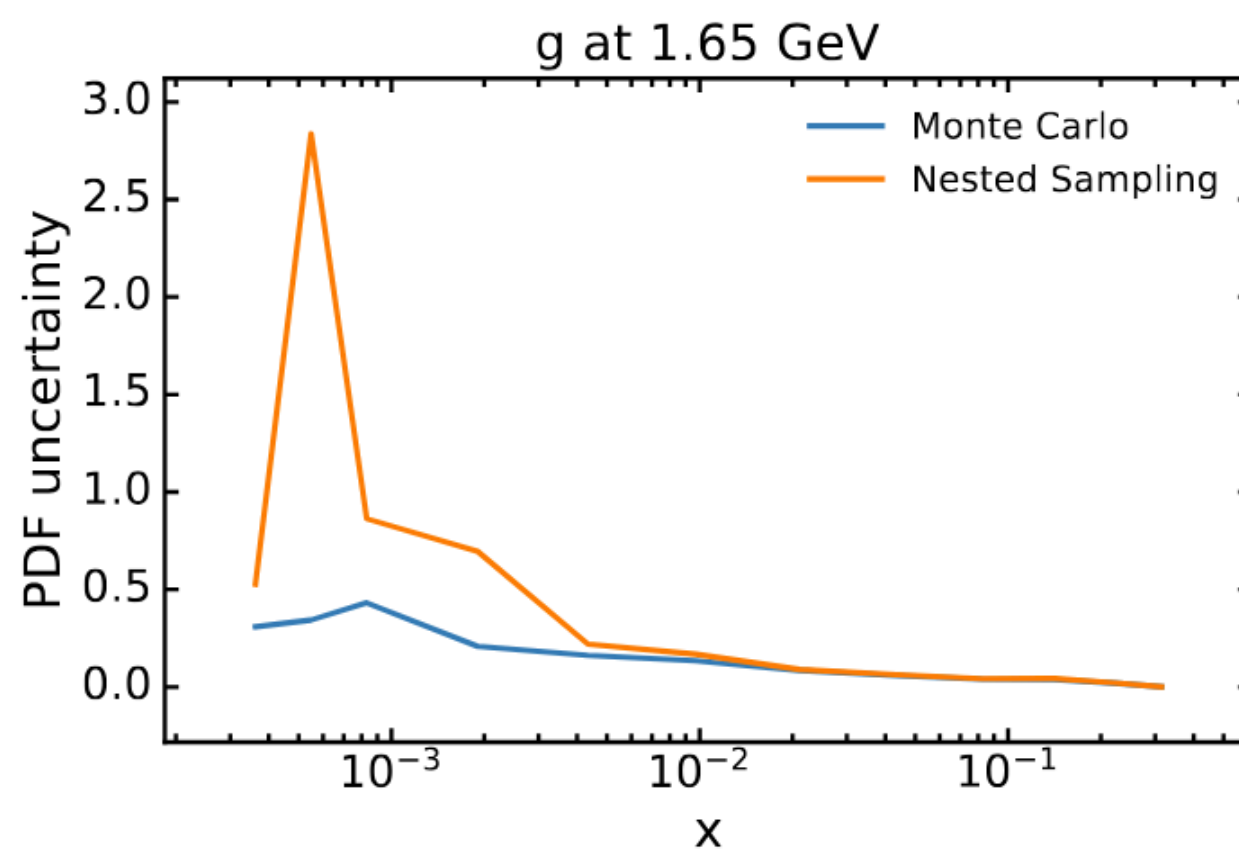
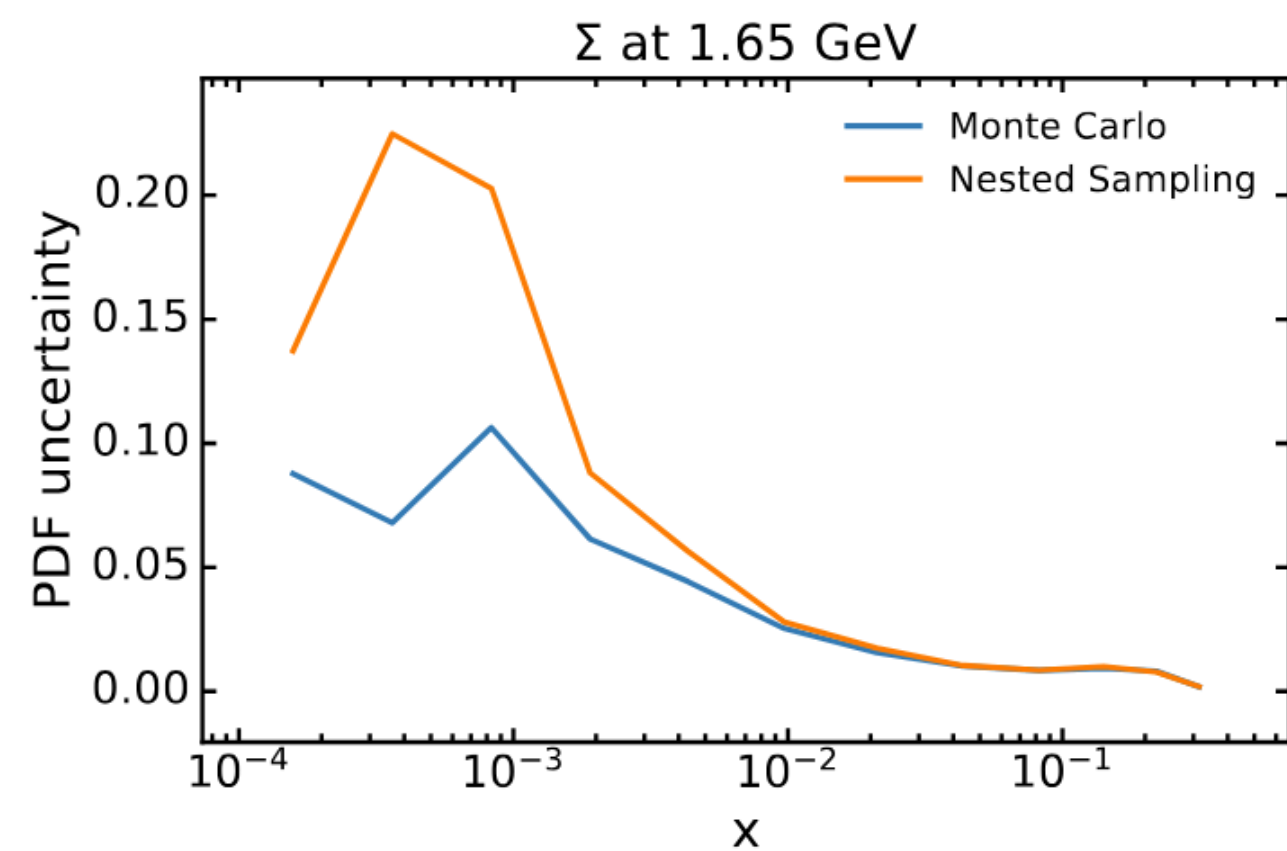
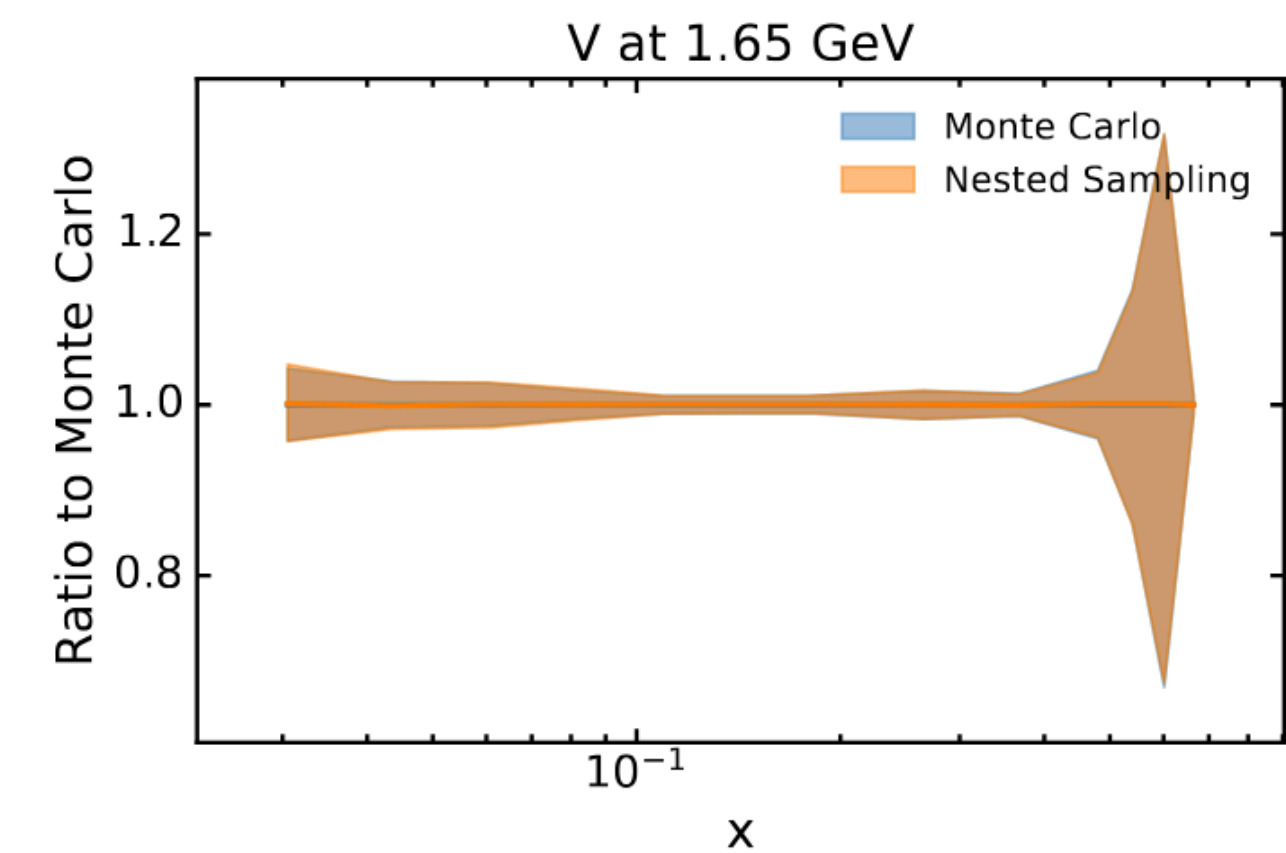
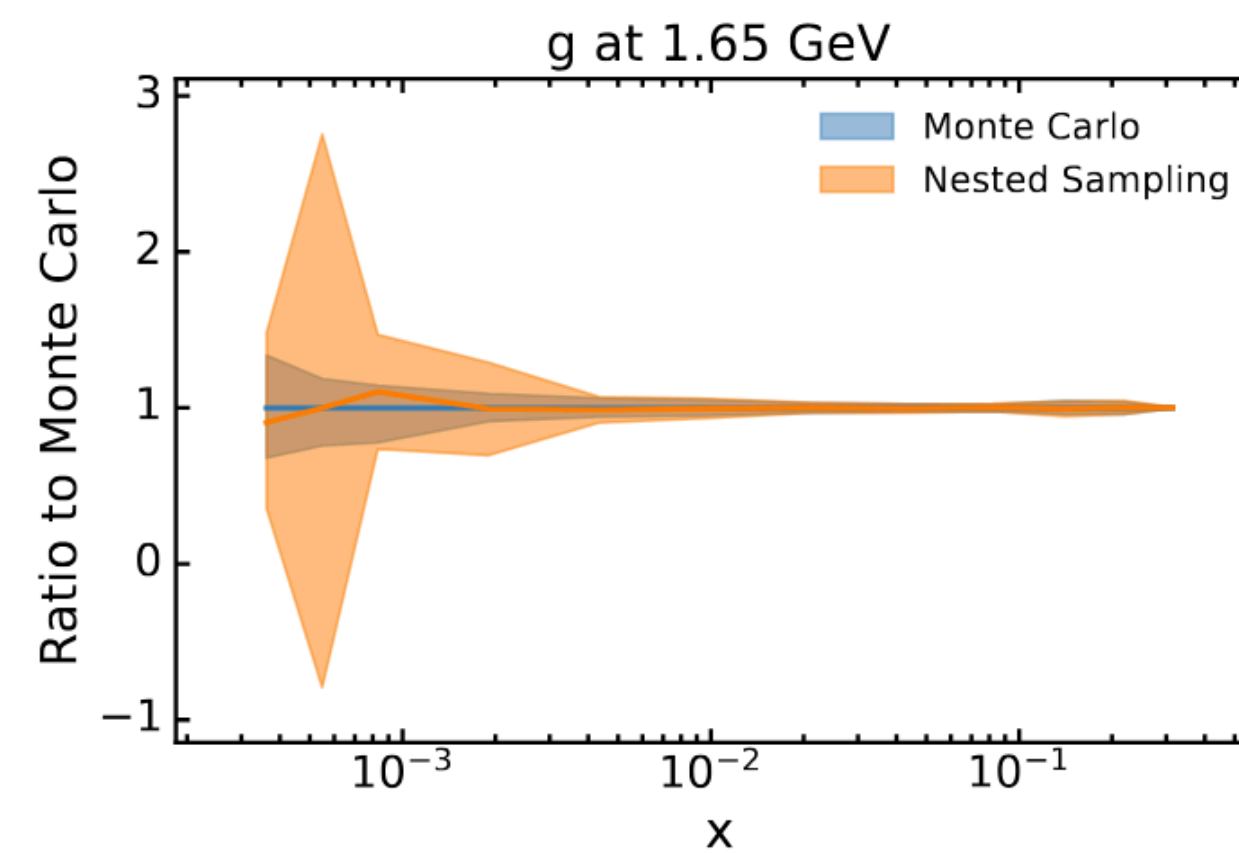
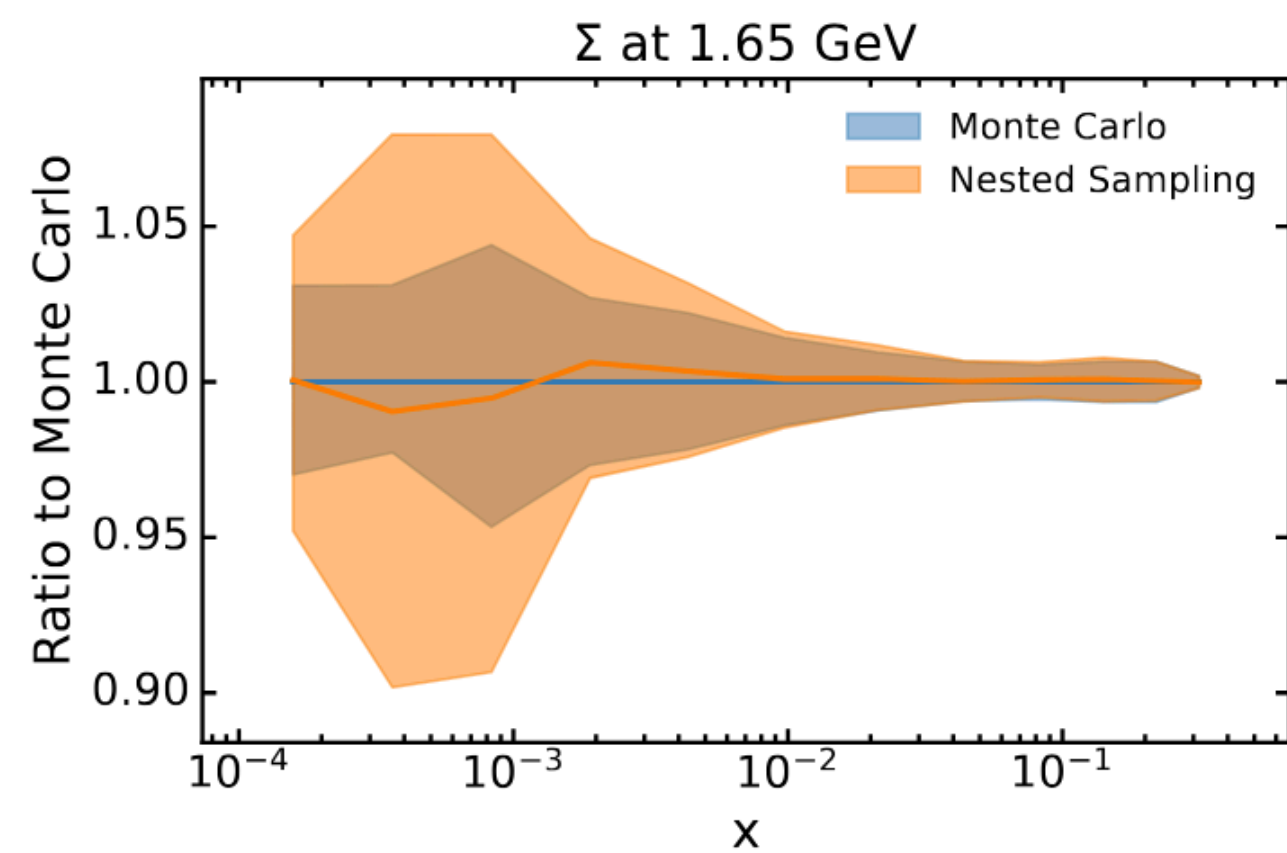
Fitting 'kinky' PDFs

- We generated an **artificial copy of the complete NNPDF4.0 dataset***, with noise, based on a kinky PDF with values at the grid points taken from the NNPDF4.0 central PDF.
- We then fit the **dataset**, using a **fully Bayesian methodology** (with a large uniform prior), and using the **Monte Carlo replica methodology**.

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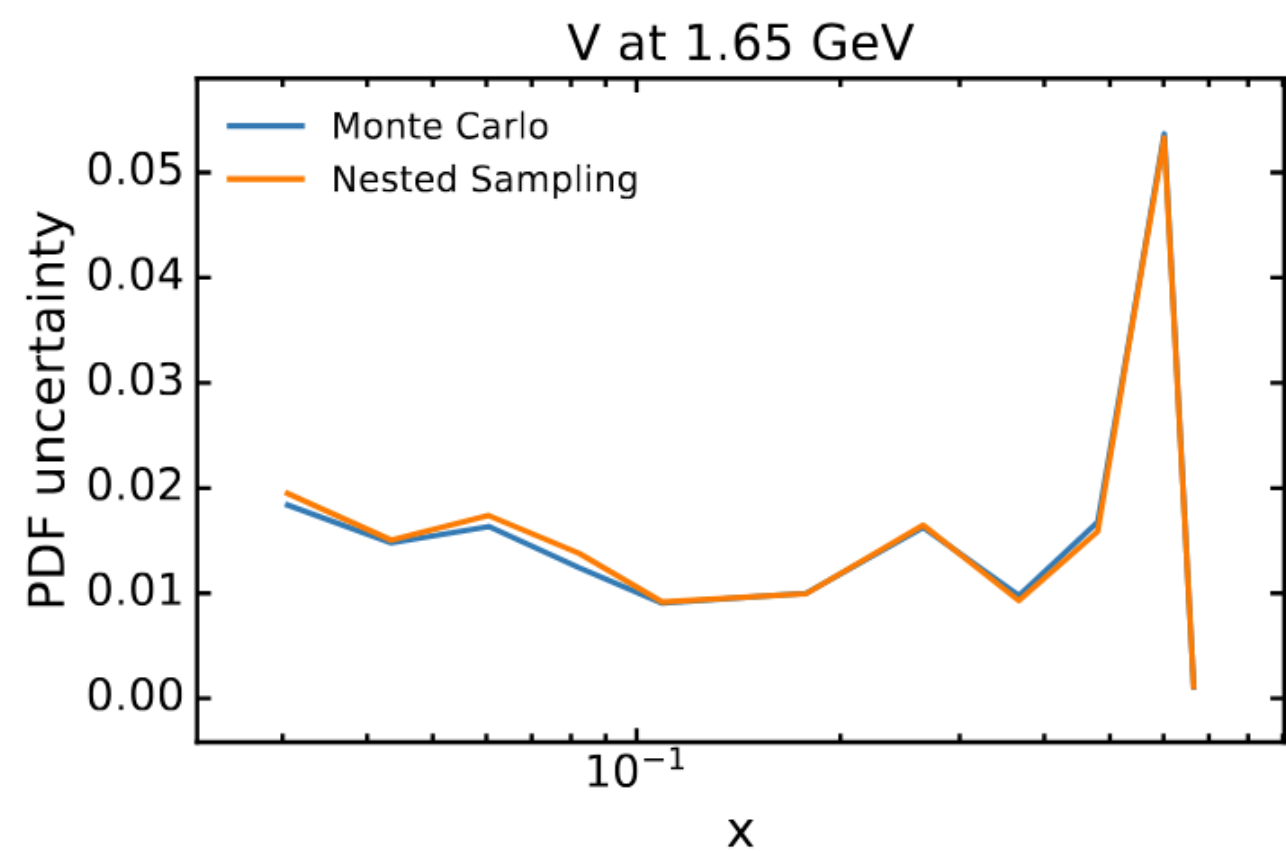
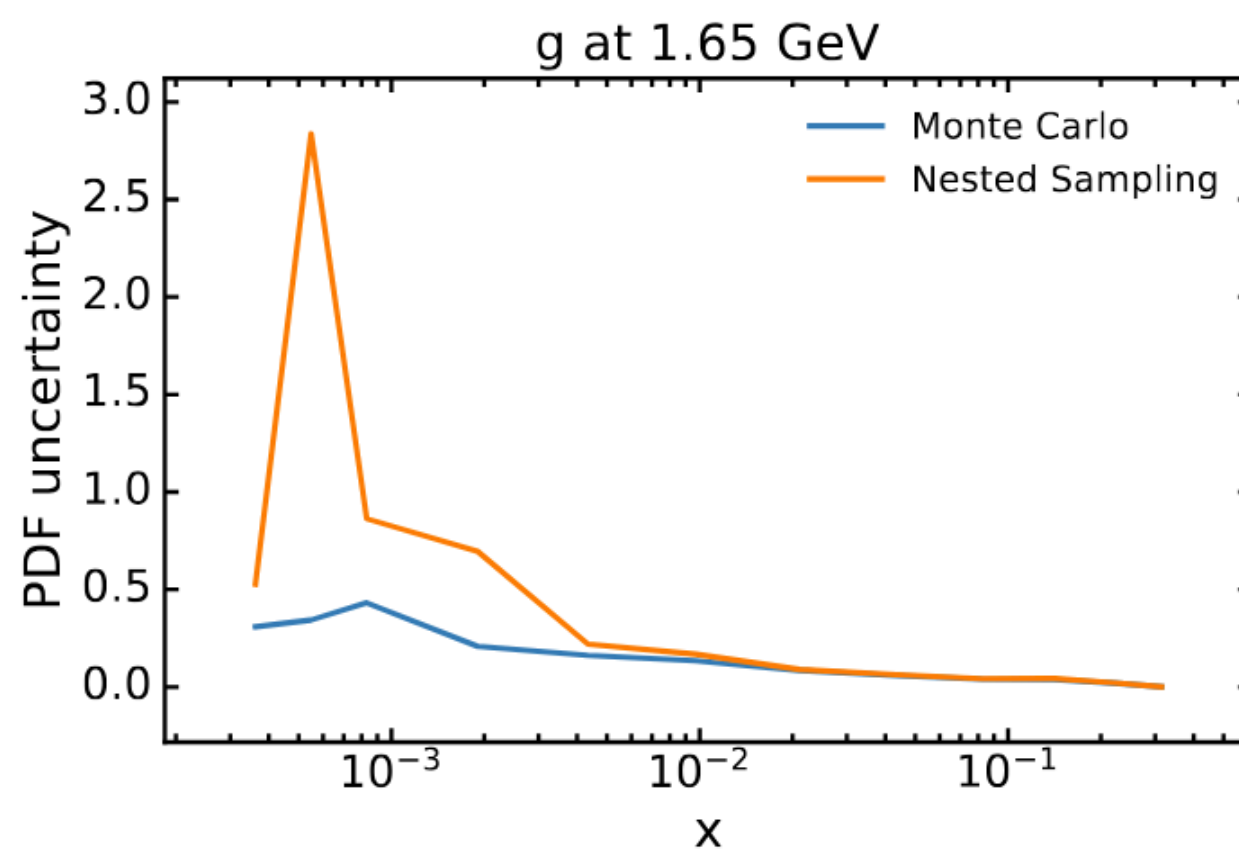
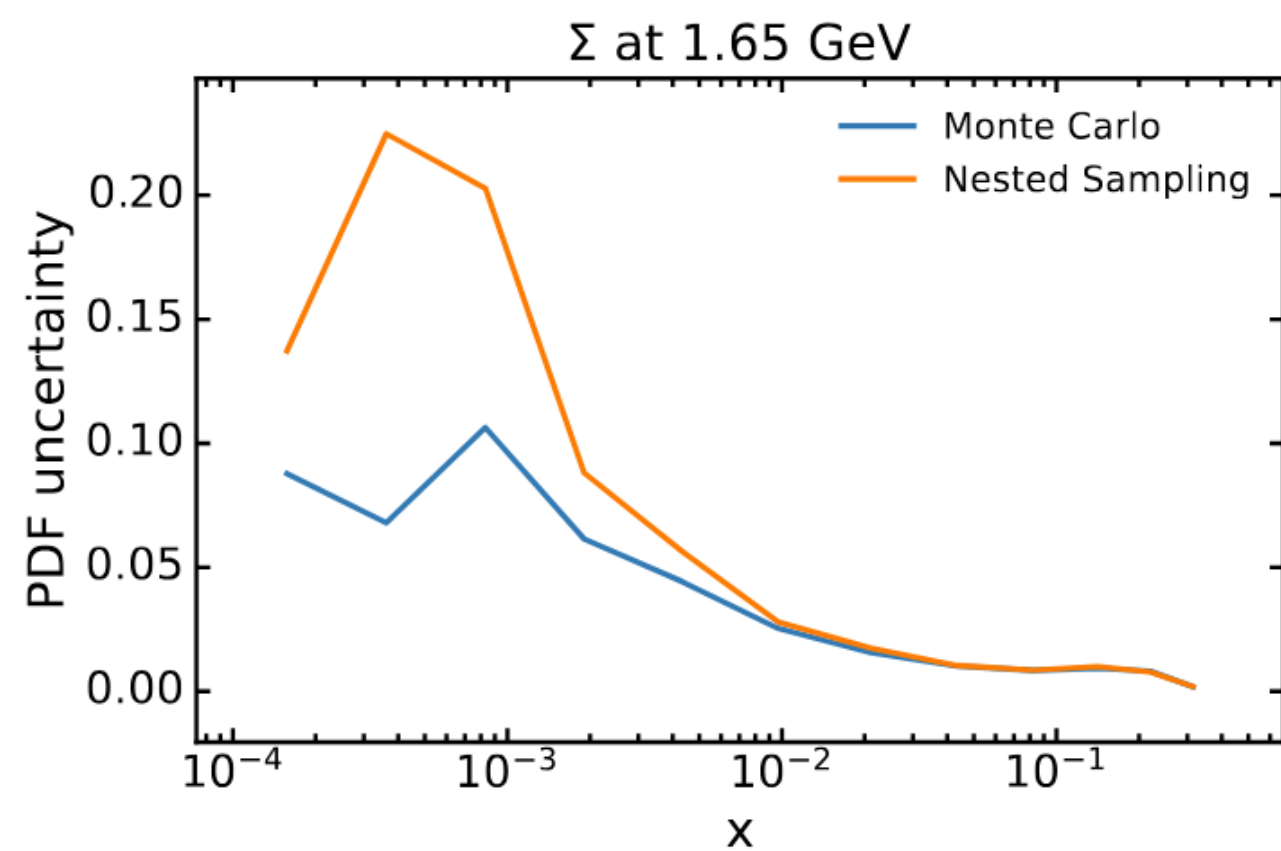
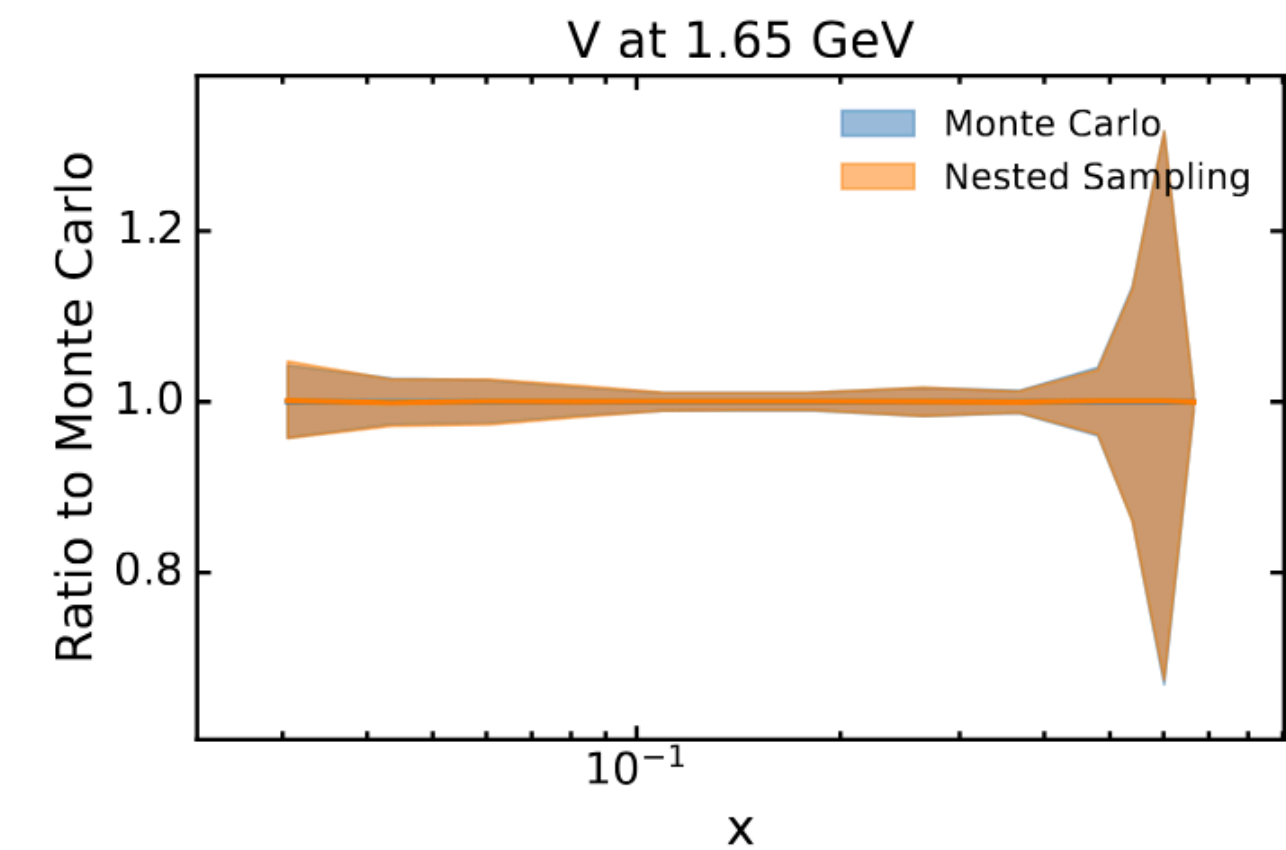
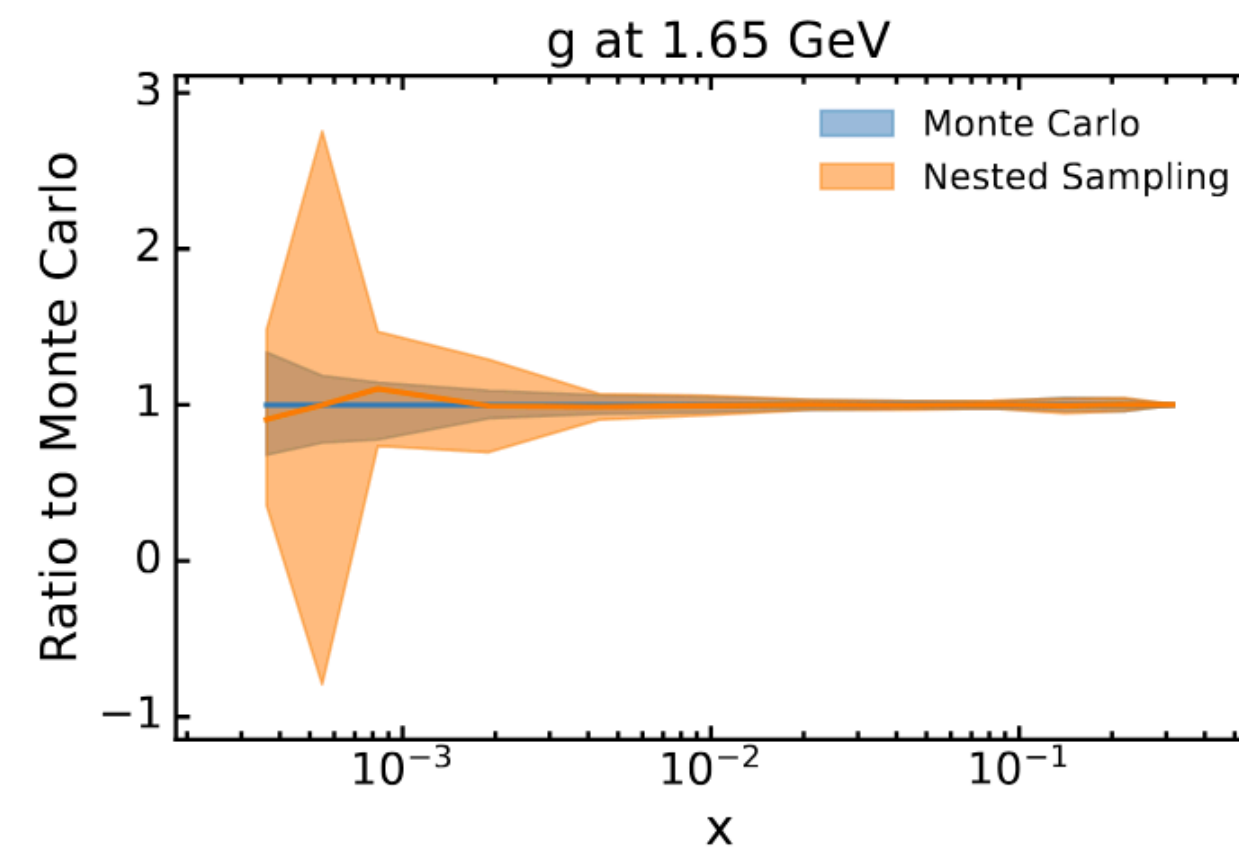
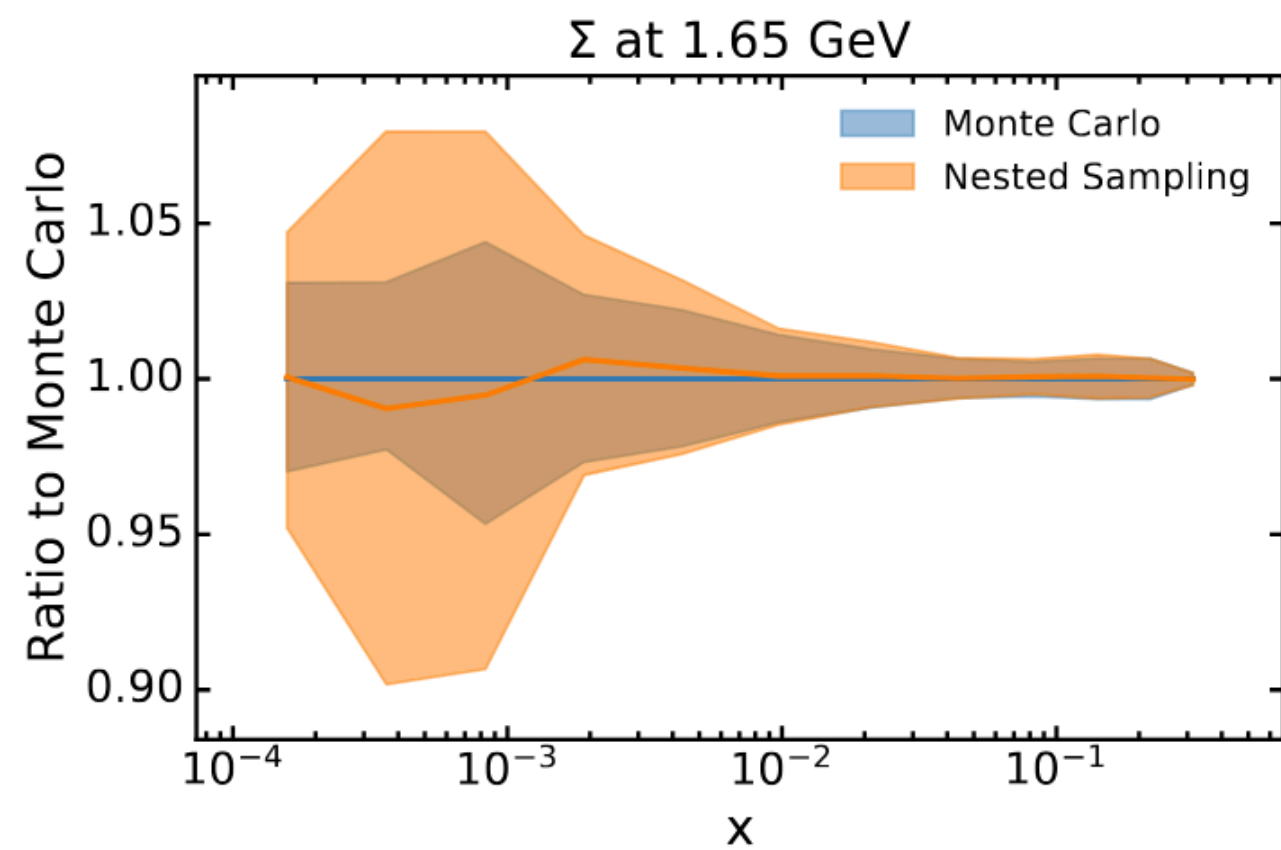
Results

- We found that, in our toy scenario, Monte Carlo does seem to **underestimate** errors relative to a fully Bayesian methodology.



Results

- Particularly at **lower x -values**, we see a reduction in uncertainties of **up to 60%** in the gluon PDF, when using Monte Carlo.



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Hence: existing PDF fits are not invalidated by this study, it merely suggests a clear and present need for a future fully Bayesian PDF analysis.

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- This can lead to **false conclusions** (originally identified in the SMEFT) in **inference problems**.
- In a **toy PDF fit**, we showed that the Monte Carlo and Bayesian approaches **disagree**, with the Monte Carlo method **underestimating** uncertainties. Hence, there is a **clear and present need** to produce a **realistic, fully Bayesian PDF fit** in the near future.

Thanks for listening!
Questions?