

# Analysis of darknoise measurements in Valencia

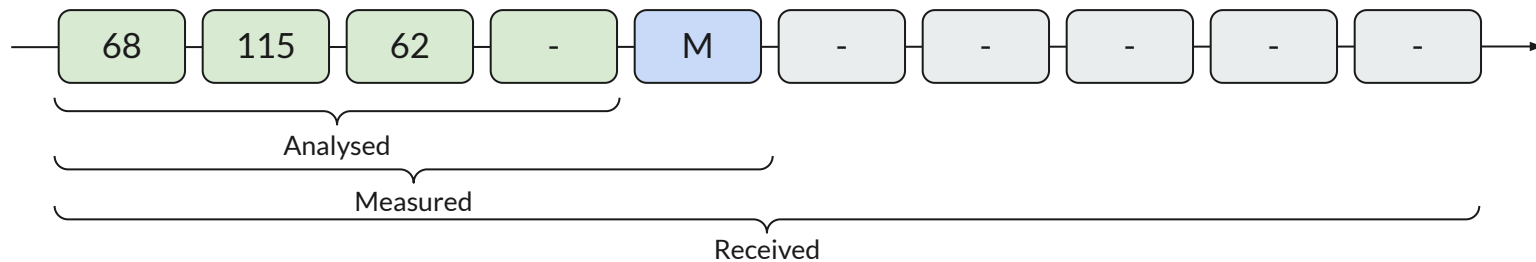
Statistical characterization of HPK SiPMs for FD1

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Photosensor WG meeting - 17 Dec 2024

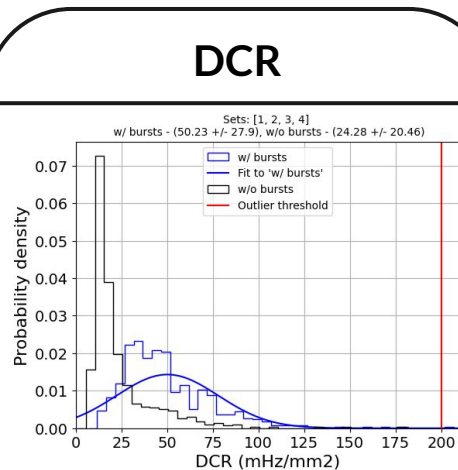
# Introduction

- 5 trays have been measured so far
- The first three trays (68, 115 and 62) have already been analysed and presented
  - Results from tray 68 (set 1), Photosensor WG meeting on 2024/02/13 - <https://indico.fnal.gov/event/63323/>*
  - Results from tray 68 (set 1), Photosensor WG meeting on 2024/02/27 - <https://indico.fnal.gov/event/63509/>*
  - Results from tray 115 (set 2), Photosensor WG meeting on 2024/05/07 - <https://indico.fnal.gov/event/64609/>*
  - Results from tray 62 (set 3), Photosensor WG meeting on 2024/05/07 - <https://indico.fnal.gov/event/66372/>*
- The fourth tray contains a mixture of boards from different trays. From now on, we will refer to the former-trays as *sets*.
- Although we have used the results from the analysis of the fourth set in this presentation, we have not devoted a particular presentation to show them yet. We will do so in coming photosensor meetings.
- In this presentation we show the results of an statistical study performed to assess the number of SiPMs which we need to measure in order to achieve a certain relative error in the statistical characterization of the DCR, XTP and APP for the full HPK production of FD1

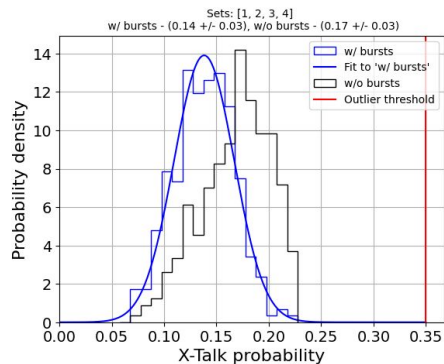


# DCR, XTP and APP distributions so far (sets 1, 2, 3 & 4)

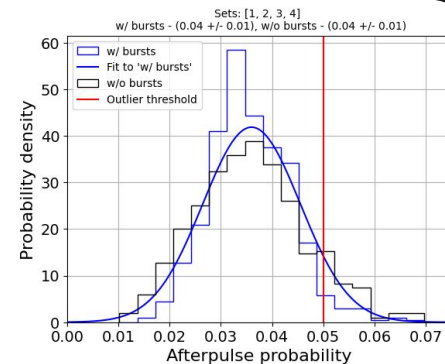
Normal fit



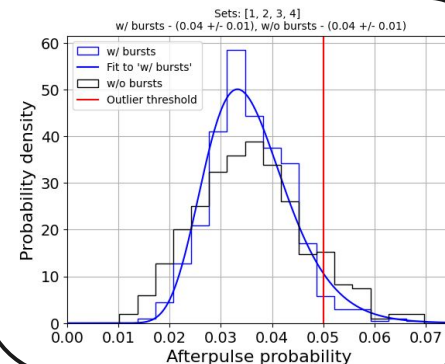
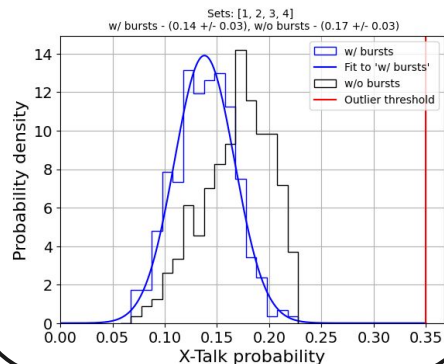
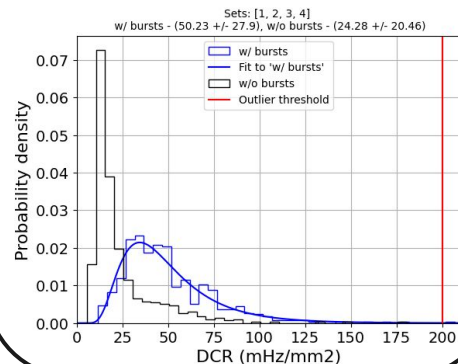
### XTP



### APP



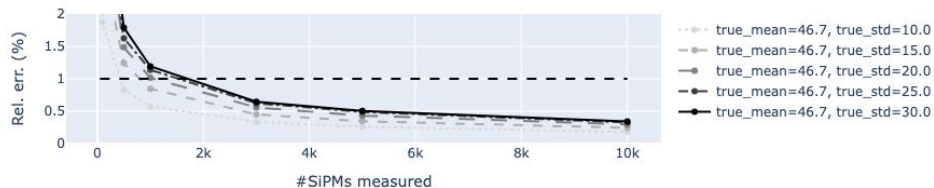
Lognormal fit



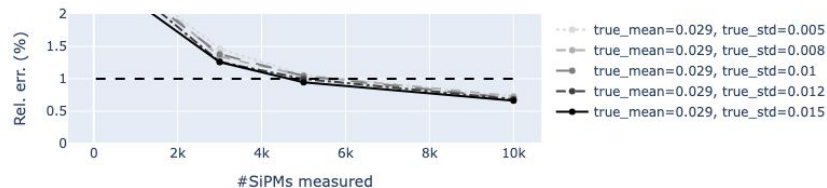
# Procedure

- For every distribution in ('normal', 'lognormal'):
  - For every magnitude in ('DCR', 'XTP', 'APP'):
    - Fit the chosen distribution to the histogram of the chosen magnitude using our results from the first four trays
    - Compute a set of mean-values and sigma-values covering a vicinity of the fitted mean and sigma
    - For every (mean, sigma) pair:
      - For every number-of-SiPMs, N, in (50, 100, ..., 5000, 10000):
        - Repeat 100 times:
          - Sample the fixed distribution with the fixed (mean, sigma) pair N times
          - Histogram the resulting samples and fit them to a distribution of the chosen kind
          - Compute the fit error for the fit mean and sigma
          - Make it relative to the (true) fixed mean and sigma of the underlying distribution
        - Compute the average relative fit errors for mean and sigma over the 100 experiments

Vertical axes: relative error (%) in the mean of DCR; Number of experiments: 100

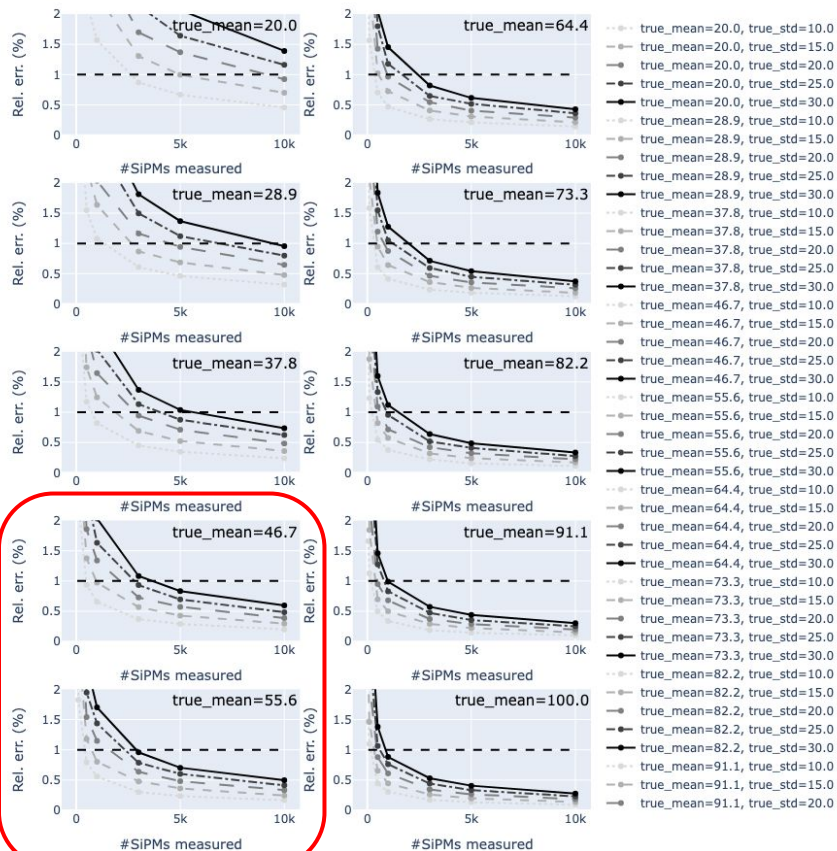


Vertical axes: relative error (%) in the std of APP; Number of experiments: 100

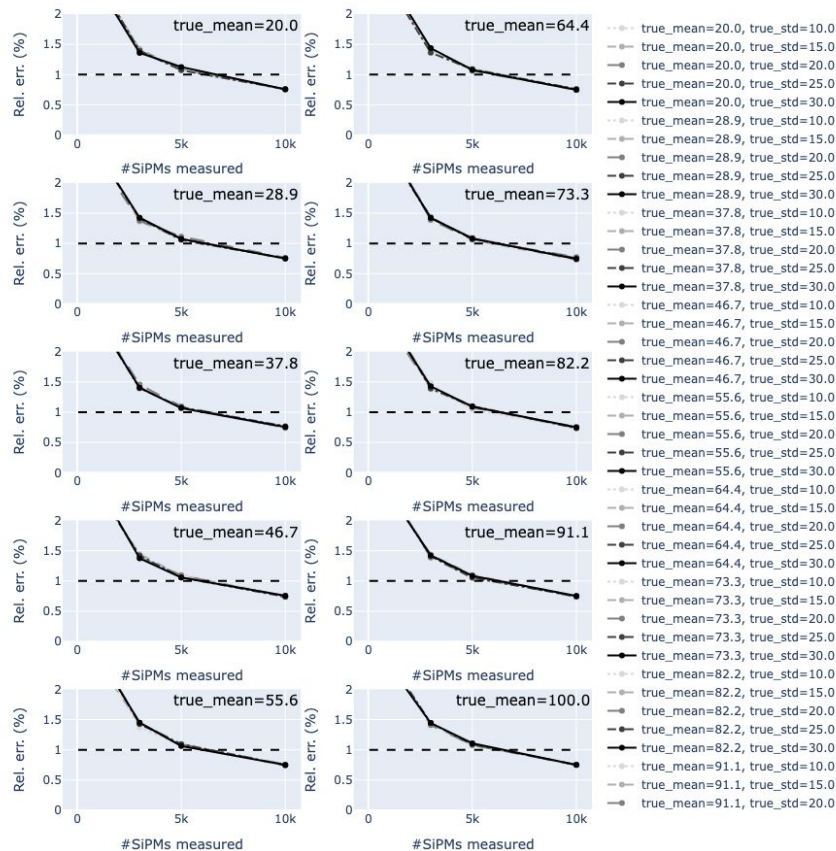


# DCR (normal fit)

Vertical axes: relative error (%) in the mean of DCR; Number of experiments: 100

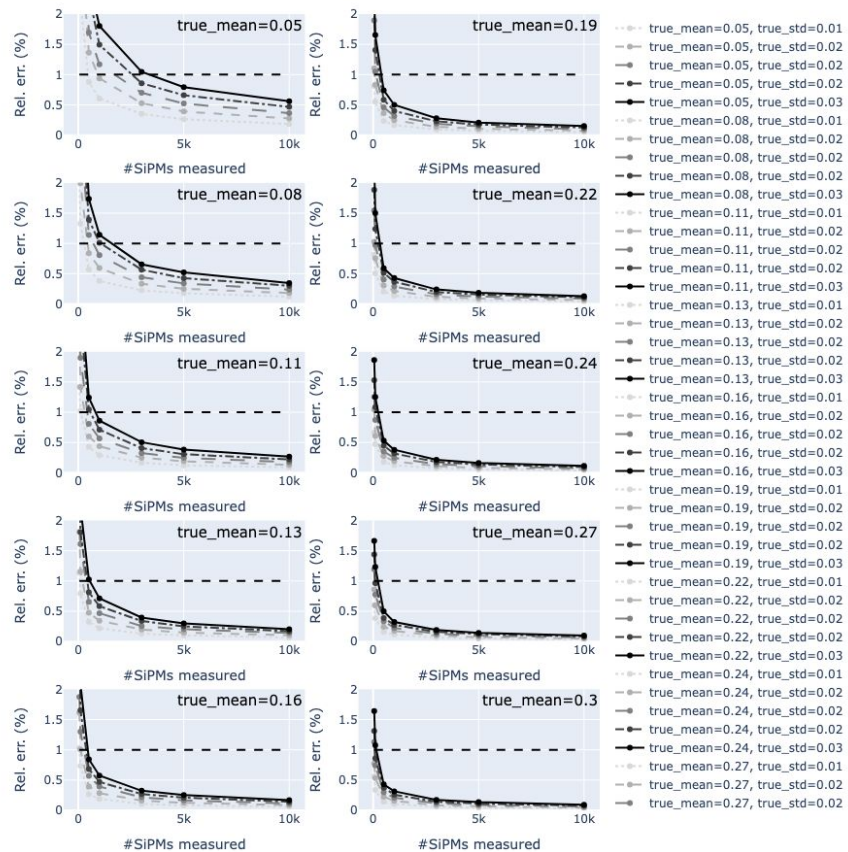


Vertical axes: relative error (%) in the std of DCR; Number of experiments: 100

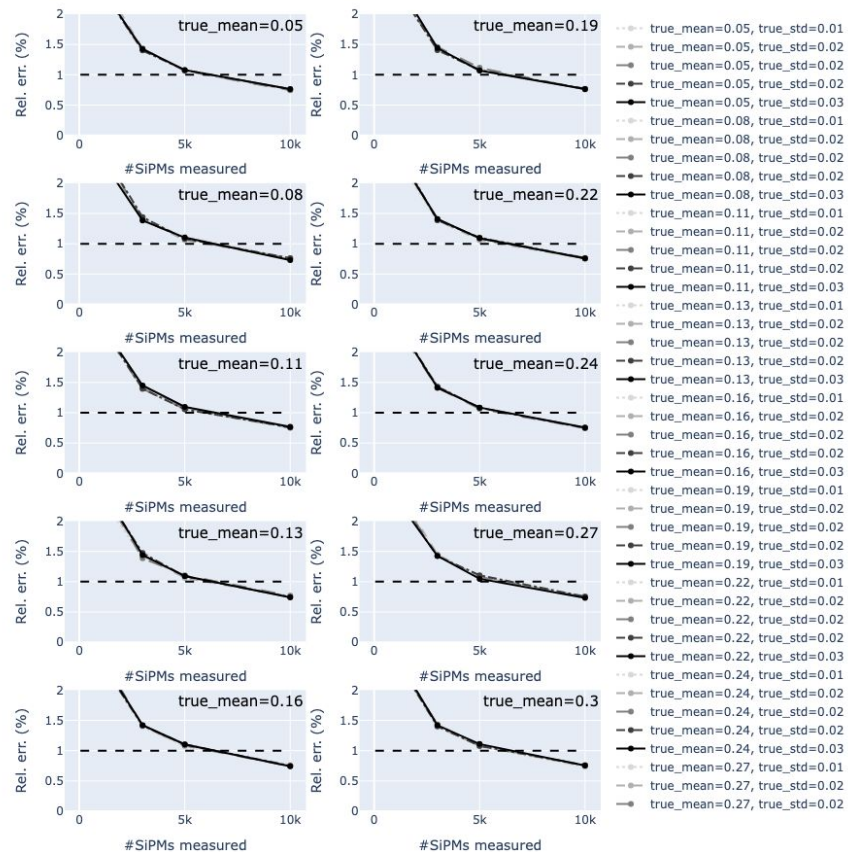


# XTP (normal fit)

Vertical axes: relative error (%) in the mean of XTP; Number of experiments: 100

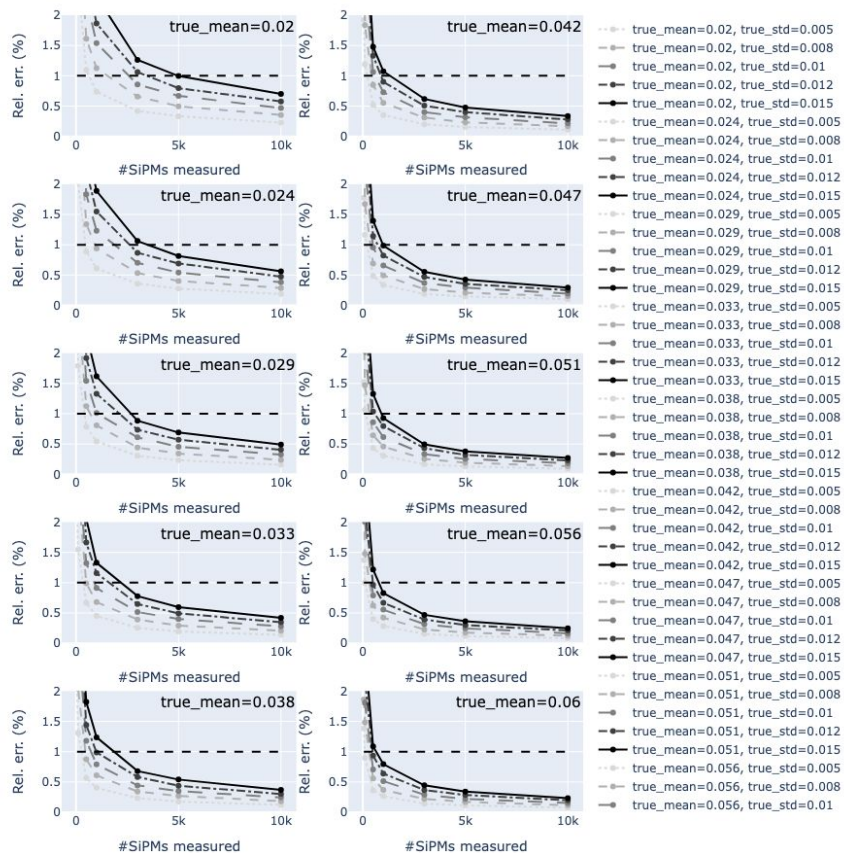


Vertical axes: relative error (%) in the std of XTP; Number of experiments: 100

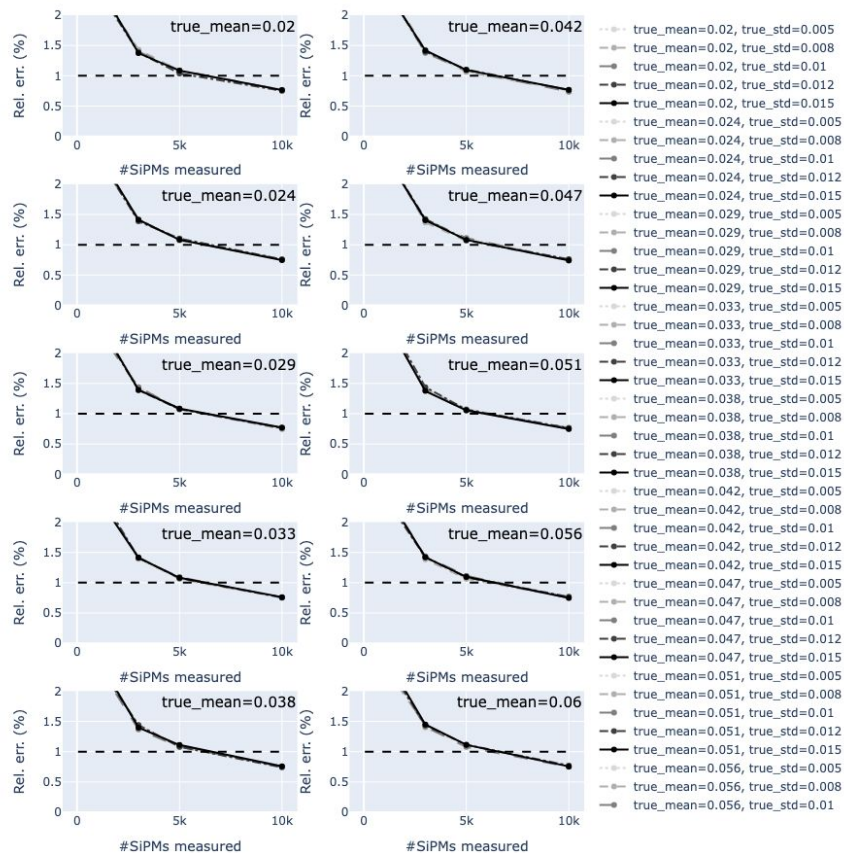


# APP (normal fit)

Vertical axes: relative error (%) in the mean of APP; Number of experiments: 100

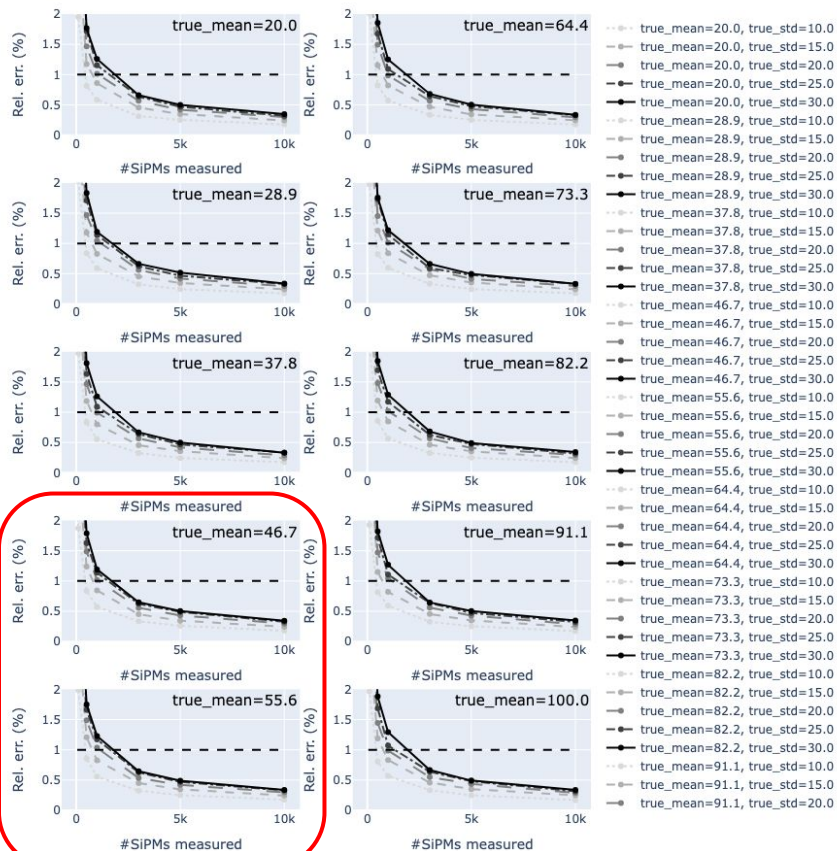


Vertical axes: relative error (%) in the std of APP; Number of experiments: 100

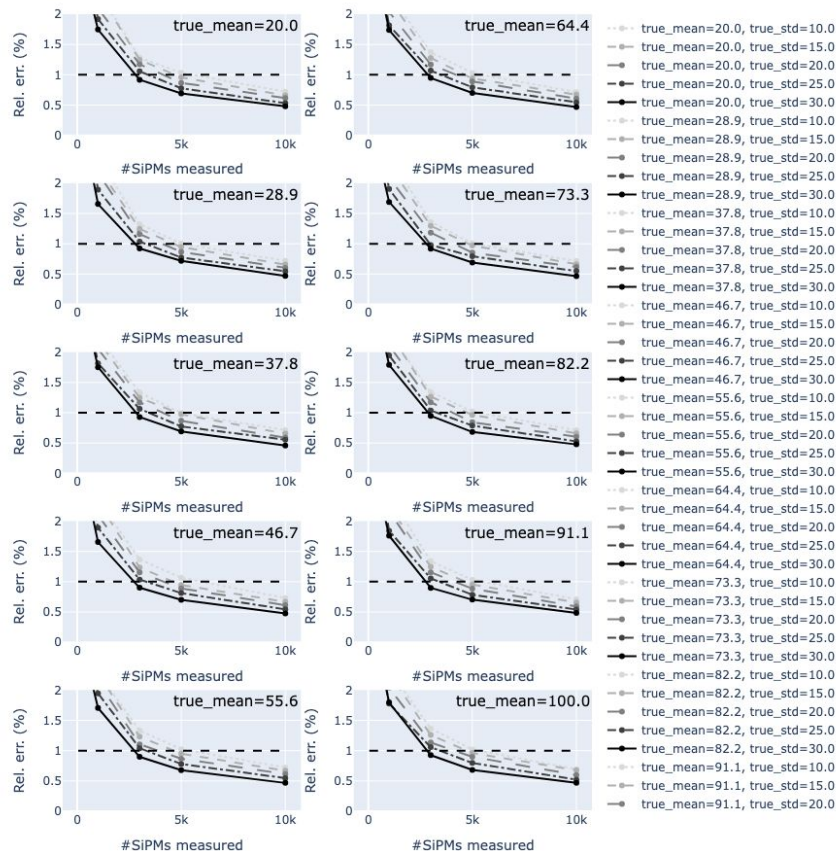


# DCR (log-normal fit)

Vertical axes: relative error (%) in the mean of DCR; Number of experiments: 100



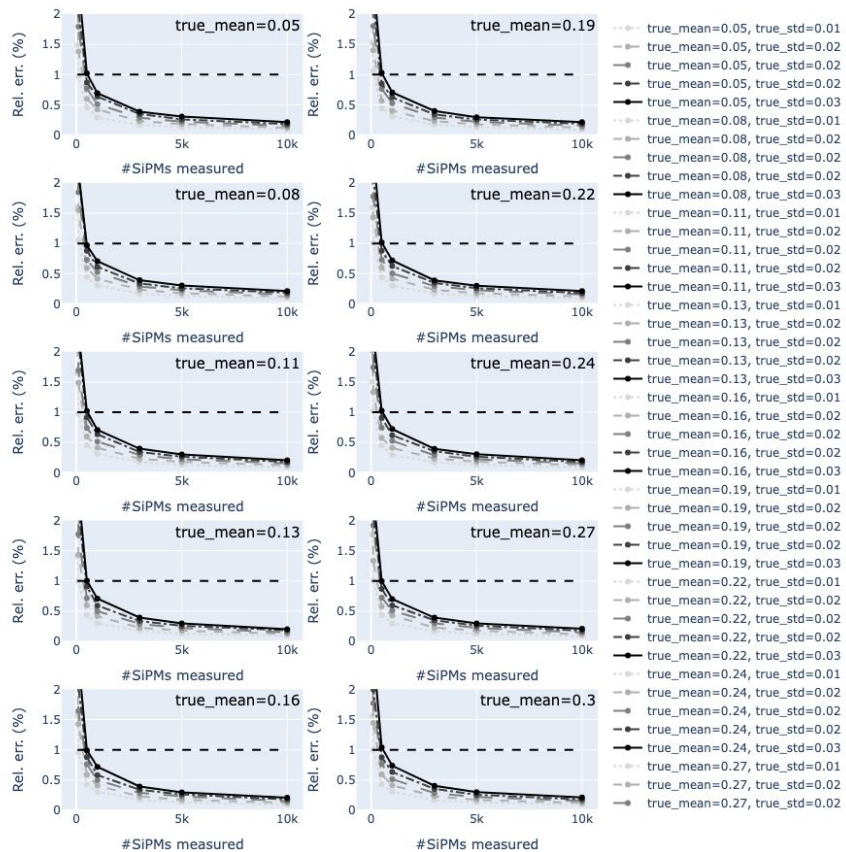
Vertical axes: relative error (%) in the std of DCR; Number of experiments: 100



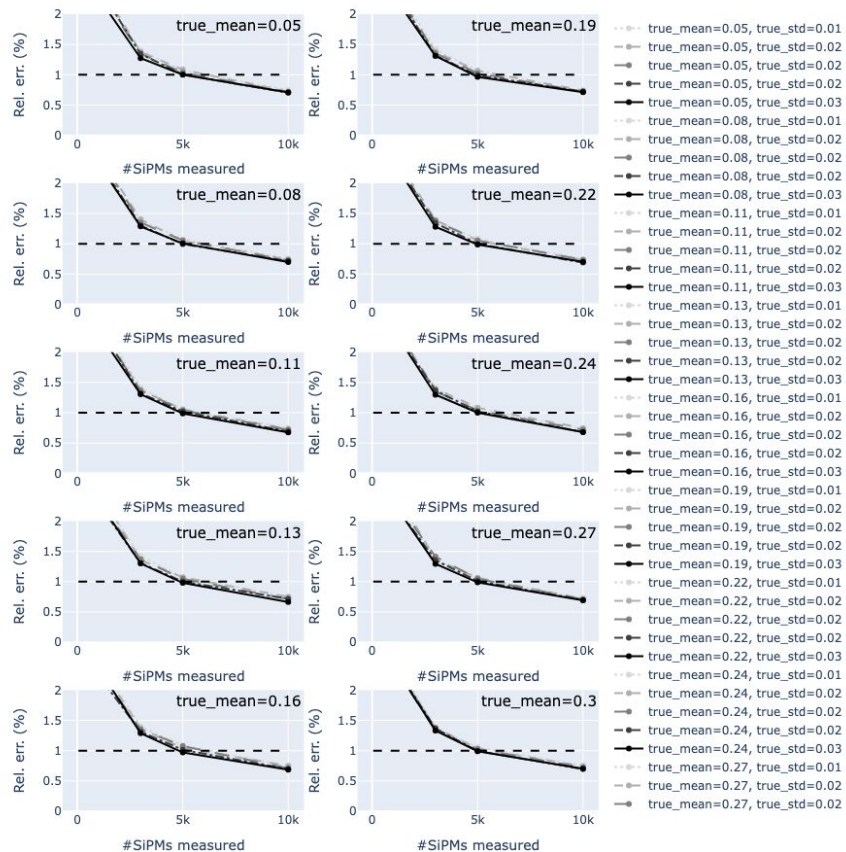


# XTP (log-normal fit)

Vertical axes: relative error (%) in the mean of XTP; Number of experiments: 100

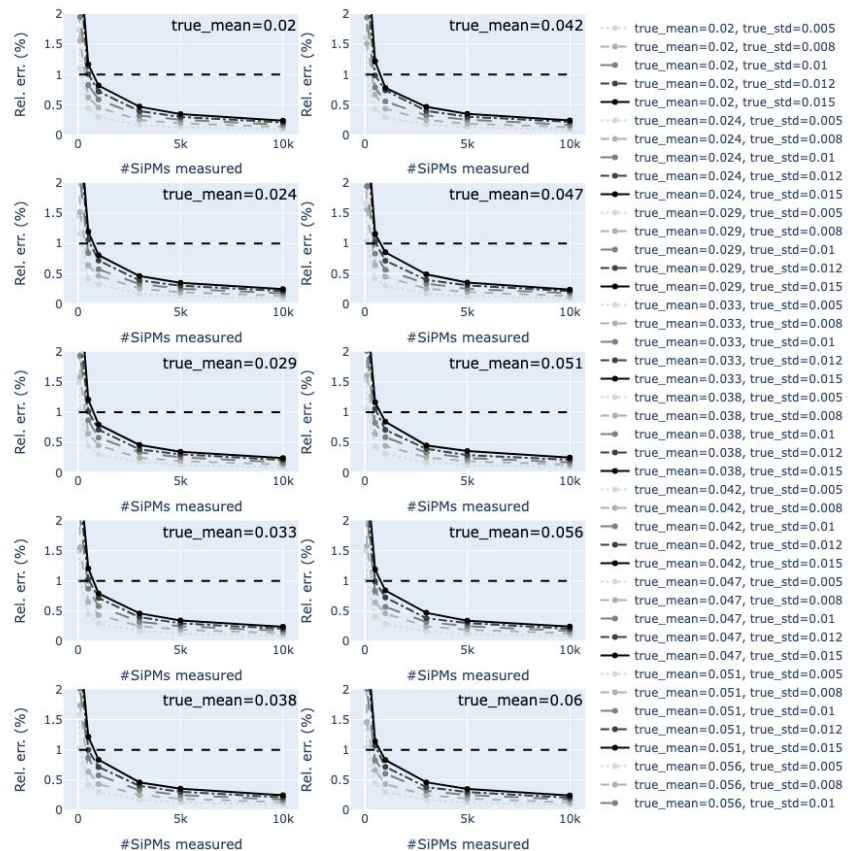


Vertical axes: relative error (%) in the std of XTP; Number of experiments: 100

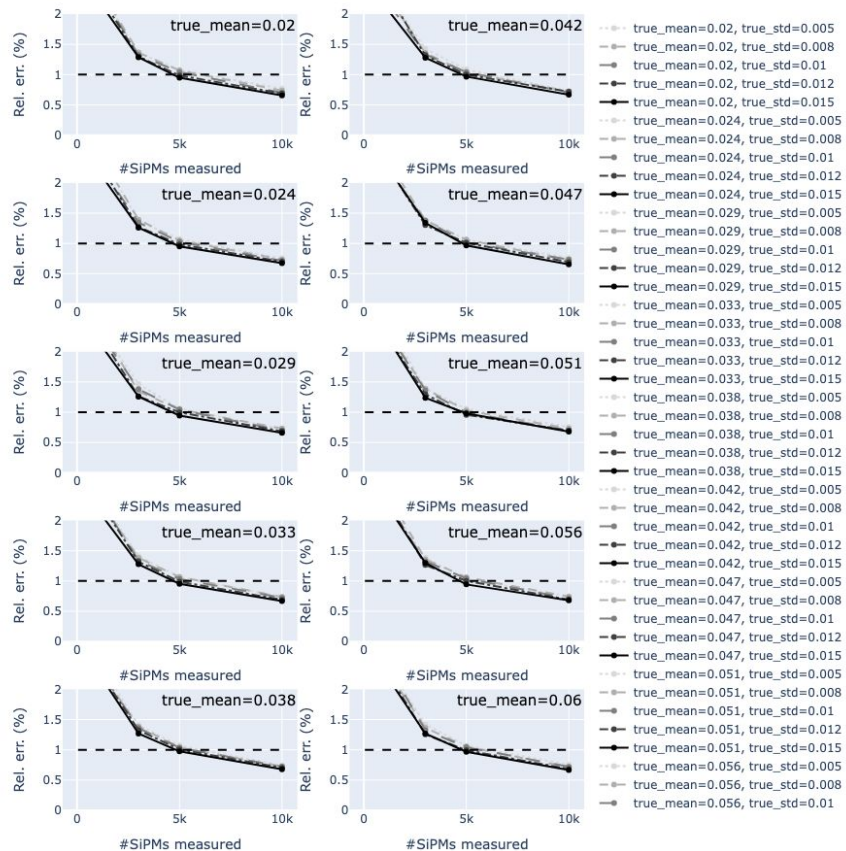


# APP (log-normal fit)

Vertical axes: relative error (%) in the mean of APP; Number of experiments: 100



Vertical axes: relative error (%) in the std of APP; Number of experiments: 100



# How many SiPMs do we need to measure? (normal/log-normal distribution assumption)

- As far as the **mean** is concerned, the limiting factor (the one which requires a bigger number of measured SiPMs to achieve a fixed relative error) is the DCR. Assuming a DCR of  **$\sim 50 \text{ mHz/mm}^2$  mean with a  $28 \text{ mHz/mm}^2$  sigma** (typical value from sets 1, 2, 3 and 4) we would need to measure 3000 SiPMs to achieve a  $\sim 1\%$  relative error in the mean.
- The relative error for the **sigma** seems *almost* unaffected by the true mean and sigma. In whichever case, we would need to measure 3000 SiPMs to achieve a  $\sim < 1.5\%$  relative error, and 5000 SiPMs to achieve a  $\sim 1\%$  relative error.

3000 SiPMs add up to a  $\sim 1\%$  of the FD1 SiPMs.  
Equivalently, they are 25 trays (so far we have received 10 trays, from which Carlos has measured 5 and I have analysed 4)

From *DUNE TDR Vol. I: Introduction to DUNE*

An **SP module** is instrumented with three module-length (58.2 m) anode planes constructed from 6 m high by 2.3 m wide **anode plane assemblies (APAs)**, stacked two **APAs** high and 25 wide, for **50 APAs per plane, and 150 total**. Each **APA** consists of an aluminum frame with three layers of active wires, strung at angles chosen to reduce ambiguities in event reconstruction, that form a grid on each side of the **APA**. The relative voltage between the layers is chosen to ensure

Introduction to DUNE

The DUNE Technical Design Report

Novel **silicon photomultiplier (SiPM)** based **photon detectors (PDs)** called **ARAPUCAs<sup>®</sup>** are placed in the inactive space between the innermost wire planes of the **APAs**, installed through slots in the **APA** frame. Each **APA** holds ten **PD** modules, for a total of 1500 per **SP module**. Of these, 500 are mounted in the **APAs** of the central anode plane and collect light from both directions, and 500 each are mounted in the outer **APA** frames and collect light from only the inner-facing direction.

48 SiPMs/supercell x 4 supercells/PD module x 10 PD modules/APA x 50 APAs/plane x 3 planes/FD1 = **288000 SiPMs/FD1**

# Summary and further questions

- The measured DCR distribution is better fit by a lognormal PDF rather than a normal one. Although less significant, XTP and APP also show heavier tails towards bigger values. For the moment we lack a fundamental reason to explain this.
- Based on the 480 SiPMs measured and analyzed in Valencia so far, we estimate that **in order to characterize the DCR, XTP and APP distributions within a ~1% error for the full HPK FD1 production** (without batch distinction) **we need to measure 3000 (HPK) SiPMs**. This is our current baseline goal.
- **Additionally, do we want to discriminate between batches?**
  - If so, how do we define a production batch?
    - Can we track production batches? I.e. SiPMs that were manufactured simultaneously
    - If that's not the case, then do we want to define a batch as the 1600 boards that are sent by HPK in the same delivery?
      - Is this definition useful from an sipm-characterization point of view?
- Do we want to discard entire batches based on the measurements?
  - If so, what should be the threshold for the fraction of SiPMs not fulfilling requirements in order to discard a batch?
- Do we want to discard individual boards based on measurements?
  - In this case, notice that our results would not be representative of the accepted sample (we'll be only measuring a 1% of the HPK production)