Bias-Variance Trade-off and Model Selection for Proton Radius Extractions

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Intuitively, a scientist might assume that a more complex regression model will necessarily yield a better predictive model of experimental data. Herein, we disprove this notion in the context of extracting the proton charge radius from charge form factor data. Using a Monte Carlo study, we show that a simpler regression model can in certain cases be the better predictive model. This is especially true with noisy data where the complex model will fit the noise instead of the physical signal. Thus, in order to select the appropriate regression model to employ, a clear technique should be used such as the Akaike information criterion or Bayesian information criterion, and ideally selected previous to seeing the results. Also, to ensure a reasonable fit, the scientist should also make regression quality plots, such as residual plots, and not just rely on a single criterion such as reduced χ². When we apply these techniques to low four-momentum transfer cross section data, we find a proton radius that is consistent with the muonic Lamb shift results. While presented for the case of proton radius extraction, these concepts are applicable in general and can be used to illustrate the necessity of balancing bias and variance when building a regression model and validating results, ideas that are at the heart of modern machine learning algorithms.

I. INTRODUCTION

High-precision Lamb shift experiments on muonic hydrogen atoms have determined the proton radius to be 0.84087(39) fm [1,2]. This result is in stark contrast to the CODATA-2014 recommended value of 0.8751(61) fm [3] which comes from electron scattering results and atomic transition frequencies [4–8]. The discrepancy between these radius values has become known as the proton radius puzzle [9–14]. While initial efforts to understand this puzzle focused on the details of the muon experiment, attention has now turned to re-examining the atomic and electron scattering results [15–18].

For the electron scattering data, the proton charge radius, r\(_p\), is extracted from cross section data by determining the slope of the electric form factor, G\(_E\), in the limit of four-momentum transfer, Q\(^2\), approaching zero:

\[
r_p \equiv \left( -6 \frac{dG_E(Q^2)}{dQ^2} \right|_{Q^2=0} \right)^{1/2}.
\]

This definition of the radius has been shown to be the same as the one used by the Lamb shift measurements [19]. Unfortunately, electron scattering experiments cannot reach the Q\(^2\) = 0 limit; thus, an extrapolation is required to determine the charge radius from the experimental data.

The various proton radius values that have been extracted from electron scattering data are shown in Fig. 1. In general, r\(_p\) extractions using simple statistical modeling of the low Q\(^2\) data [20–23] tend towards a smaller radius while the more complex statistical model, with many free parameters, tend to extract a large proton radius [24,25]. Nuclear theory constrained extractions of the radius tend to favor a smaller radius [29,31]. One technical detail that is affecting all these results is how the normalization of the experimental data is handled.

![FIG. 1. Shown are the radii extracted from electron scattering data [24–50]. The vertical bands indicate the value and uncertainty of the proton radius from muonic hydrogen [11,2] and CODATA-2014 [3]].](image-url)
examining the very low $Q^2$ region where slope and normalization are systematically linked together. A general criticism of the small radius extractions of the proton radius is the presence of statistical bias [55, 57], with an implication that bias needs to be avoided in order to successfully extract the true radius from the data. The use of Monte Carlo methods to find bias and then reject simple proton radius extraction methods originated with the classic Monte Carlo study of Borkowski et al. [49] where linear extrapolations were flatly rejected in favor of quadratic extrapolations. Interestingly, that work ignored the variance of the more complex function.

We will show in this work that when using a Monte Carlo study to test a model’s ability to extract the proton radius one needs to consider not only bias but also variance, and find an appropriate balance between the two. After all, it is better to have a slightly biased watch to tell the hour, than a broken one that is unbiased because it overestimates and underestimates the time symmetrically.

We will also illustrate that one must consider the range, quantity and precision of the data when determining the best predictive statistical model and show that simple biased statistical models can have a higher predictive validity than unbiased more complex models [58]. We will then apply these ideas to model selection with real data, where instead of millions of Monte Carlo results, we get but a single realization of the possible outcomes.

II. BIAS

In the English language, bias is often used as a pejorative term. In the context of regression, it is simply an offset from the mean from the true central value. Since it is part of a distribution, it is not a property of a single realization but can be determined by repeated sampling. In the context of the proton radius extractions, bias was nicely illustrated by Borkowski et al. [49] and we will describe their procedure in the following paragraphs.

Form factor pseudo-data is first systematically generated from 0.1 fm$^{-2}$ to 0.4, 0.8, 1.2, and 1.6 fm$^{-2}$ in steps of 0.05 fm$^{-2}$ using the standard dipole function:

$$G_D(Q^2) = (1 + Q^2/(18.23 \text{ fm}^{-2}))^{-2}, \tag{2}$$

where the cutoff parameter of 18.23 fm$^{-2}$ corresponds to a radius of 0.8113 fm. In this study, the authors of the original work erroneously concluded that the linear models should always be rejected in favor of the lower-bias quadratic function.

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The procedure was repeated with $10^6$ sets of pseudo data to precisely determine the mean of the extracted radii for these two functions. Since the standard dipole was used as the input function, one would expect an unbiased function to return a radius of 0.8113 fm. Table I reproduces the original result [49]. As the table shows, the mean values of $a_0$ and $r_p$ show a clear bias. Based on this study, the authors of the original work erroneously concluded that the linear models should always be rejected in favor of the lower-bias quadratic function.

III. VARIANCE

While the linear fit does exhibit a bias, bias is not the only quantity that must be considered when selecting an appropriate statistical model to use. In particular, along with the offset from the true value, bias, one must also consider the distribution of the outcomes, the variance. Where variance is the square of the standard deviation, $\sigma$, of the statistical distribution. Table II shows a more complete picture of the simulation results where the $\sigma$ of the results is shown along with the bias and is graphically represented in Fig. 2.

For all $Q^2$ intervals, the linear fits provide significantly smaller $\sigma$ than the more complex quadratic fits. Though picking the function to use based solely on variance would also be incorrect. Thus, this seemingly simple example has turned into a nearly textbook illustration of the trade-off between variance and bias with the linear fits having a relatively high bias with a low variance, while the quadratic fits have a low bias and high variance. Of course, to calculate the bias requires we know the true value which often isn’t the case in a real experiment.

| Interval $Q^2$ fm$^{-2}$ | Linear fit $a_0$ [fm] | Radius [fm] | Quadratic fit $a_0$ [fm] | Radius [fm] |
|--------------------------|----------------------|-------------|--------------------------|-------------|
| 0.1 – 0.4                | 1.000                | 0.79        | 1.000                    | 0.81        |
| 0.1 – 0.8                | 0.999                | 0.78        | 1.000                    | 0.81        |
| 0.1 – 1.2                | 0.997                | 0.77        | 1.000                    | 0.81        |
| 0.1 – 1.6                | 0.996                | 0.76        | 1.000                    | 0.81        |
TABLE II. An expanded version of Table I, where instead of just showing the mean offset of the fit results for $a_1/a_0$, the bias, we also indicate the width of the fit results, $\sigma$. Recall that the point uncertainty is fixed at 0.5%. For the simulated radius of 0.8113 fm, one would expect an unbiased fit to give $a_1/a_0$ of 0.1097 fm$^{-2}$; thus, the difference of that value from the mean fitted value of $a_1/a_0$ is the bias and the width of the distribution, $\sigma$. Also shown is the root mean square error, RMSE, which can be used to quantify the best function for a given interval taking into account both bias and variance.

| Data Interval Points | $a_0$ Radius [fm] | $a_1/a_0$ Bias [fm] | $\sigma$ RMSE [fm$^2$] | $a_0$ Radius [fm] | $a_1/a_0$ Bias [fm] | $\sigma$ RMSE [fm$^2$] |
|----------------------|-------------------|---------------------|---------------------|-------------------|---------------------|---------------------|
| 0.1 – 0.4 7          | 0.9995 0.7949      | -0.1053             | -0.0044 0.0184      | 0.0189            | 1.0000 0.8065       | -0.1084 -0.0013      |
| 0.1 – 0.8 15         | 0.9987 0.7827      | -0.1021             | -0.0076 0.0057      | 0.0095            | 1.0000 0.8094       | -0.1092 -0.0005      |
| 0.1 – 1.2 22         | 0.9975 0.7711      | -0.0991             | -0.0106 0.0030      | 0.0110            | 0.9999 0.8087       | -0.1090 -0.0017      |
| 0.1 – 1.6 31         | 0.9959 0.7601      | -0.0963             | -0.0134 0.0019      | 0.0136            | 0.9998 0.8075       | -0.1087 -0.0010      |

TABLE III. Same as Table II, but now with equal number of data points for each range.

| Data Interval Points | $a_0$ Radius [fm] | $a_1/a_0$ Bias [fm] | $\sigma$ RMSE [fm$^2$] | $a_0$ Radius [fm] | $a_1/a_0$ Bias [fm] | $\sigma$ RMSE [fm$^2$] |
|----------------------|-------------------|---------------------|---------------------|-------------------|---------------------|---------------------|
| 0.1 – 0.4 31         | 0.9995 0.7952      | -0.1054             | -0.0043 0.0098      | 0.0107            | 1.0000 0.8091       | -0.1091 -0.0006      |
| 0.1 – 0.8 31         | 0.9987 0.7827      | -0.1021             | -0.0076 0.0041      | 0.0086            | 1.0000 0.8098       | -0.1093 -0.0004      |
| 0.1 – 1.2 31         | 0.9974 0.7711      | -0.0991             | -0.0106 0.0026      | 0.0109            | 0.9999 0.8091       | -0.1091 -0.0006      |
| 0.1 – 1.6 31         | 0.9959 0.7601      | -0.0963             | -0.0134 0.0019      | 0.0136            | 0.9998 0.8075       | -0.1087 -0.0010      |

IV. GOLDFILOCKS DILEMMA

As was noted by George Box, all models are wrong, thus, the goal is to find the most useful model [59]. For any given set of statistical models, the goal is to find the optimal balance between bias and variance. In general, this can be written as

$$\frac{d\text{Bias}^2}{d\text{Complexity}} \approx -\frac{d\text{Variance}}{d\text{Complexity}},$$

which as illustrated in Fig. 3. Thus, to quantify the goodness of the fits, we choose Root Mean Square Error (RMSE) [60, 61],

$$\text{RMSE} = \sqrt{\text{bias}^2 + \text{Sigma}^2} = \sqrt{\text{Bias}^2 + \text{Variance}}.$$  

Using the RMSE values in Table II, one can now quantify that for this example the 0.1–0.8 fm$^{-2}$ interval is the preferred range for the linear model while the 0.1–1.6 fm$^{-2}$ interval is the preferred range for the quadratic model. Going to even higher $Q^2$ will require even more complex models as illustrated in Fig. 3. This is in contrast to the conclusion one draws when one only considers bias as presented in Table I, though consistent with the observation that the optimal specific form of the parameterization may depend on the $Q^2$ region being fit [62].

It is interesting to repeat the Monte Carlo simulation for equal number of data points within each range especially since, for elastic scattering, cross sections are significantly higher at lower values of $Q^2$ and thus, it is easy to obtain more low $Q^2$ data. This is shown in Table III and now the picture is even grayer as the RMSE of the linear fit is nearly equal to the quadratic, thus, assuming that the standard dipole was the true generating function, an experiment with 31 data points and an uncertainty of 0.005 per point over a range of 0.1 to 0.8 fm$^{-2}$ and a different experiment over a range of 0.1 to 1.6 fm$^{-2}$ would have an equal probability of reproducing the correct radii if all other things were equal (given that the modeler in the first group adjusts a line and the one in the second a parabola). This is visualized in Fig. 5 where the linear fit is clearly biased but has a small variance compared to the unbiased, large variance quadratic fit.

The choice of the parsimonious modeler to use the low $Q^2$ data would likely be driven by the recognition of the fact that as $Q^2$ increases the extraction of the charge form factor is complicated by the growing influence of the magnetic form factor. The choice to use a larger $Q^2$ range would likely be driven by a desire to form a more complete picture of the proton’s structure. For example, the parsimonious modeler may only be interested in the proton radius while another modeler may be interested in higher order moments [63]. Thus, the tension in the extractions of the proton radius from electron scattering data is really about the fact that modelers using the low $Q^2$ are generally getting a systematically different result than the modelers doing fits which include high $Q^2$ data, and perhaps points to a systematic problem with our knowledge of the magnetic form factor and/or the functional form of the form factors.

V. THE BEST PREDICTIVE MODEL

Selecting between a linear or quadratic regression of the more complex standard dipole function may seem a bit contrived, as one might naively think that just using
FIG. 2. A graphic representation of the Monte Carlo results showing how the linear fits tend to have a relatively high bias though a low variance, while the quadratic fits tend to have a relatively low bias but a large variance. The black line/point represents the true value.

the generating function itself would always yield the best results. In section D we prove that for non linear fits there is an induced bias even when using the generating function.

When taking into account the variance, we can also show that this is not always the case with a simple set up: we use the lowest $Q^2$ range, 0.1 − 0.4 fm$^{-2}$, and replace the quadratic function with the generating function and a floating normalization term:

$$f_{\text{DipoleFit}}(Q^2) = n_0 (1 - b_1 Q^2 / 2)^{-2},$$  \hspace{1cm} (7)

where $n_0$ is the normalization factor and the radius is given by $\sqrt{-6b_1}$. Pseudo data is generated for absolute random errors of 0.01, 0.005 and 0.003 with three different spacings: 0.05 fm$^{-2}$ spacing with 7 points, 0.01 fm$^{-2}$ spacing with 31 points, and 0.005 fm$^{-2}$ spacing with 61 points. The results of fitting these pseudo data sets are shown in Table IV. For a given row in the Table, the linear fit has the greater bias and the dipole fit always has the greater variance; bringing the root mean square error very close for all the test cases. This example also makes it clear that it is not just the number of points that matter, but the size of the uncertainties and the range of the data that will also be critical parameters in model
selection. In the cases where the MSE of the line was smaller than the ones of the Dipole, the line was being a better “predictive” model for the proton radius, while the Dipole was a better “descriptive” model of the form factor as whole.

Of course for real data, nature hides the true generating function from us, so perhaps it is reassuring to know that a reasonable approximation is able to reveal the underlying physics just as well as, if not better than, the true function. To be clear, the lesson is not that one function is better than another; it is that for a given set of data, the scientist is challenged to use the most appropriate model (either descriptive or predictive) for the task at hand. Further details on the general mathematics behind these example problems can be found in [58].

Using an inappropriate model can lead to erroneous conclusions (e.g. due to different normalizations of the data). In Fig. 6, we show an example of one set of the pseudo data fit with a linear and quadratic where the prior that the $G_E(0) = 1$ has been applied (i.e. the data has been divided by the normalization term from the regression so the function goes to the known limiting value at the origin). Since here we know the function that generated the pseudo data, it is clear that the quadratic fit can cause the data to be inappropriately shifted and can generate an large variations in the radius. Of course in the real world, the true function is not known; thus, with real data a model selection technique is required in order to determine the appropriate function.

VI. MODEL SELECTION

While this classic Monte Carlo example problem is over 40 years old, it actually points to exactly the split in the current electron scattering proton radius extraction procedures. The parsimonious modelers, who are focused solely on extracting a radius, have focused on the low $Q^2$ region accepting a slightly higher level of bias in exchange for low variance; on the other hand, those modelers who are interested in extracting more information about the proton (e.g. higher order moments) fit longer $Q^2$ ranges and have focused on complex models which, while lower in bias, come at a cost of higher variance.
Also, since we do not know the true model, one cannot in general calculate the RMSE. So, while Monte Carlo exercises like the one described herein are extremely useful for finding reasonable models to consider and understanding expected uncertainties, the data must be used to select the appropriate model. For this one can rely on statistical modeling selection techniques such as an F-test for nested models [65, 67] or the more general Akaike information criterion (AIC) [68] or Bayesian information criterion (BIC) [69] to guide our selection of the most appropriate model to describe a given set of data. These statistical criteria are calculated as follows:

\[
\chi^2 = \sum_{i=1}^{N} (\frac{\text{data}_i - \text{model}}{\sigma_i})^2, \quad (8)
\]

\[
\text{reduced } \chi^2 = \frac{\chi^2}{N - N_{\text{var}}}, \quad (9)
\]

\[
\text{AIC} = N \log(\chi^2/N) + 2N_{\text{var}}, \quad (10)
\]

\[
\text{BIC} = N \log(\chi^2/N) + \log(N)N_{\text{var}}, \quad (11)
\]

where \( N \) is the number of data points, \( \text{data}_i \) and \( \sigma_i \) are measured values and estimated uncertainties, and \( N_{\text{var}} \) is the number of model parameters. Further details about current model selection techniques can be found in [70].

One should also keep in mind that the input models in Monte Carlo simulations are always just an approximation and one needs to be careful about drawing too strong inferences from the simulated results. For example, just because the linear model has a negative bias when compared to the standard dipole, does not imply that it has a negative bias with respect to all possible models.

### VII. REAL DATA

This brings us to the real data and the current proton radius extractions. Many different functions have been tried over the years from simple linear fits [53, 55] and continued fractions [43] to high order polynomials with [29] and without constraints [26]. Since obtaining sub-percent level absolute cross sections is nearly impossible, a normalization parameter is included to allow an entire set of data to shift as was done in the Monte Carlo simulations. Again, this just allows that the prior, \( G_E(0) = 1 \), can be applied. To be clear, in this work, when referring to a set of experimental data we are referring to a group of data with a single normalization parameter.

The measured cross sections, \( \sigma_{\text{Meas}} \), are related to the charge and electric form factors via

\[
\frac{\sigma_{\text{Meas}}}{\sigma_{\text{Mott}}} = \frac{n_0}{\varepsilon(1 + Q^2/4M^2)} \left[ \varepsilon G_E^2(Q^2) + \frac{Q^2}{4M^2} G_M^2(Q^2) \right], \quad (12)
\]

where \( n_0 \) is a normalization factor and the kinematic quantities \( Q^2 \) and \( \varepsilon \) are given by

\[
Q^2 = \frac{2ME^2(1 - \cos \theta)}{M + E(1 - \cos \theta)}, \quad (13)
\]

\[
\varepsilon = \left[ 1 + 2 \left( 1 + \frac{Q^2}{4M^2} \right) \tan^2 \theta \right]^{-1} \quad (14)
\]

where \( E \) is the incident electron beam energy, \( M \) is the proton mass, \( \theta \) is the measured electron scattering angle. The Mott cross section, \( \sigma_{\text{Mott}} \), with the recoil factor included, is given by

\[
\sigma_{\text{Mott}} = \frac{\alpha^2}{4E^2 \sin^4(\theta/2)(1 + \frac{Q^2}{M(1 - \cos \theta)})}. \quad (15)
\]

From Eq. 12 it is clear that as \( Q^2 \) goes to zero and \( \varepsilon \) goes to one (forward angle electron scattering), the terms which include the magnetic form factor, \( G_M \), become relatively unimportant.

To illustrate the current tension between fits done with different models, Fig. 8 shows 104 data points from one full subset from a modern electron scattering experiment [71] that covers a range similar to the range studied herein. By using a single set, only one floating normalization parameter is required. This one set of data covers...
FIG. 8. Shown is one set of modern cross section data to illustrate the current tension between different models and the effect of the normalization parameter. In electron scattering data, the point-to-point uncertainties can be rather small, compared to the typical normalization uncertainty of a few percents. To make the curves, the standard dipole magnetic form factor has been used along with charge form factors from several recent publications: a bounded 13th-order polynomial with a radius of 0.88 fm [25], an unbounded 10th-order polynomial with a radius of 0.88 fm [25], a continued fraction with a radius of 0.84 fm [22], and a dipole function with a radius of 0.84 fm [20].

a range similar to the 85 data point fits with 6 floating normalizations [21]. Along with the data, four representative functions are shown using Eq. 12 with a standard magnetic form factor. Two functions give radii that agree with the CODATA value for the proton radius [22] and two that agree with the muonic Lamb shift measurements [22]. We note that this figure looks oddly similar to the pseudo data in Fig. 6, but here the true function is unknown so it is not clear which curves are shifted with respect to the true reduced cross section values.

Following the logic of this work, we try fitting the cross section data shown in Fig. 8 with Eq. 12 where $G_E$ has been approximated by Eq. 3 and Eq. 4 though with the normalization now subsumed in Eq. 12. Additionally, we performed fits with the following two commonly used functions:

$$f_{\text{cubic}}(Q^2) = 1 + a_1 Q^2 + a_2 Q^4 + a_3 Q^6,$$

$$f_{\text{rational}}(Q^2) = \frac{1 + n_1 Q^2}{1 + m_1 Q^2},$$

where the radius for the rational function is given by $\sqrt{-6(n_1 - m_1)}$ and for the polynomials by $\sqrt{-6a_1}$. The low order rational function ($n = m = 1$) can easily be extended to give the expected asymptotic behavior as $Q^2 \to \infty$ by using a more complex rational function where $m = n + 2$ such as in [12, 73, 74].

In order to check on the influence of the magnetic form factor on the results, several different magnetic form factor functions were used. First, the fits were done with a standard dipole magnetic form factor and then repeated with the magnetic form factor from [25] and [72]. The $F$-test, AIC, or BIC model selection techniques all slightly prefer fits with Eq. 3. The use of statistical criteria for model selection helps avoids confirmation bias, though one could still be using an inappropriate function for the problem at hand. Uncertainties were determined by applying a statistical bootstrap to the data [75]. This is done by repeatedly randomly sampling the true data with replacement to generate thousands of new sets of N points and refitting those new sets. This allows one to get uncertainty distributions using the data itself and avoids a number of assumptions that are required for $\chi^2$ uncertainty techniques to be valid [75] and, unlike $\chi^2$ techniques, is also sensitive to over-fitting [76].

It is of particular note that the quadratic fit is the same function over a similar range as found in the original 1976 work opting for the quadratic fit [25] though herein we use a single floating normalization instead of three and the point-to-point errors are smaller. It is perhaps distressing that published values of the radius extracted from electron scattering remained basically unchanged since the 1976 work [19] while the functions used to make the extrapolation and obtain that same radius became increasingly complex and convoluted. Oddly enough the standard dipole magnetic form factor quadratic fit has the lowest AIC and BIC values and gives a result consistent with the muonic Lamb shift; though the rational function is nearly as good and is nearly exactly between the CODATA and muonic Lamb shift values.

VIII. GRAPHS IN STATISTICAL ANALYSIS

Beyond simply checking statistic criteria, it is important to check the statistical analysis graphically [77]. The graphs help ensure that our underlying assumptions about the data are reasonably correct and help establish that the results aren’t being overly influenced by a single point. This is particularly important when doing $\chi^2$ minimizations where a clear outlier can easily have an undue weight, given the quadratic structure of $\chi^2$ in the errors. This is not to imply that one should simply remove an outlier, but it does mean that one might wish to study the effect of fitting with and without the outlining point to clearly show its influence on the result. The researcher can also review their notes to ensure that nothing odd happened during the taking of that data point. To illustrate these points clearly, one can find in Appendix A an updated version of the classic Anscombe example [77] where we have added uncertainties so all the examples give a reduced $\chi^2$ of unity yet the distribution of the data are in fact all clearly different.

In the top half of Fig. 9 the reported data is shown along with the quadratic fit: best fit from Table V based on AIC and BIC. In the bottom half, the residual, a difference between the data and the model, is shown. While these two plots are perhaps the most common statistical
TABLE V. Using four different magnetic form factor parameterizations, we extract the normalization and electric radius using a linear, quadratic, cubic, and rational approximation for the $G_E$ function in Eq. [12]. To avoid multiple floating normalizations, the single set of 104 data points shown in Fig. 5 is used. The parameter uncertainties were obtained by performing statistical bootstraps of the data. As seen during the Monte Carlo studies, the linear fit over this interval produces a small variance; but is clearly biased from the true 0.84–0.88 fm proton radius, whereas the quadratic fit has a larger variance but gives less biased result. The cubic fit function over-fits and thus, produces a huge variance. The rational function is nearly as good as the quadratic through produces a systematically larger radius though nicely in the range we expect. In the Table df stands for degrees of freedom.

| $G_E$ Function | df | $\chi^2$ without Coulomb correction | AIC | BIC | Norm Extrapolated Radius [fm] | $\chi^2$ with Coulomb correction | AIC | BIC | Norm Extrapolated Radius [fm] |
|----------------|----|----------------------------------|-----|-----|--------------------------------|---------------------------------|-----|-----|--------------------------------|
| linear         | 102 | 162.6 1.594 50.47 55.74 0.988(1) 0.785(2) | 167.2 1.639 53.35 58.64 0.985(1) 0.790(2) |
| quadratic      | 101 | 119.4 1.182 20.35 28.29 1.000(2) 0.852(10) | 119.0 1.178 20.00 27.93 0.998(2) 0.860(10) |
| cubic          | 100 | 117.3 1.173 20.51 31.13 0.990(6) 0.785(57) | 117.1 1.171 20.33 30.90 0.990(6) 0.797(57) |
| rational       | 101 | 120.0 1.188 20.83 28.76 1.001(2) 0.860(10) | 119.6 1.184 20.50 28.64 0.999(2) 0.869(10) |

| $G_E$ Function | df | $\chi^2$ without Coulomb correction | AIC | BIC | Norm Extrapolated Radius [fm] | $\chi^2$ with Coulomb correction | AIC | BIC | Norm Extrapolated Radius [fm] |
|----------------|----|----------------------------------|-----|-----|--------------------------------|---------------------------------|-----|-----|--------------------------------|
| linear         | 168.1 1.648 53.95 59.24 0.986(1) 0.780(2) | 173.0 1.669 53.35 58.64 0.985(1) 0.785(2) |
| quadratic      | 101 | 119.4 1.182 20.33 28.27 1.000(2) 0.852(10) | 119.0 1.178 20.00 27.91 0.998(2) 0.860(10) |
| cubic          | 100 | 117.1 1.171 20.36 30.94 0.985(6) 0.798(57) | 116.9 1.169 20.14 30.72 0.983(6) 0.613(57) |
| rational       | 101 | 120.0 1.188 20.89 28.82 1.001(2) 0.861(10) | 119.6 1.184 20.87 28.80 1.000(2) 0.870(10) |

| $G_E$ Function | df | $\chi^2$ without Coulomb correction | AIC | BIC | Norm Extrapolated Radius [fm] | $\chi^2$ with Coulomb correction | AIC | BIC | Norm Extrapolated Radius [fm] |
|----------------|----|----------------------------------|-----|-----|--------------------------------|---------------------------------|-----|-----|--------------------------------|
| linear         | 168.1 1.647 53.85 59.14 0.986(1) 0.791(2) | 168.0 1.647 53.85 59.14 0.986(1) 0.791(2) |
| quadratic      | 101 | 119.5 1.183 20.44 28.37 1.000(2) 0.854(10) | 119.0 1.179 20.08 28.00 1.000(4) 0.862(10) |
| cubic          | 100 | 117.1 1.173 20.52 31.13 0.992(6) 0.786(56) | 117.1 1.171 20.33 30.91 0.990(6) 0.797(56) |
| rational       | 101 | 120.0 1.189 20.93 28.86 1.001(2) 0.871(13) | 119.7 1.185 20.60 28.54 0.999(2) 0.871(13) |

| $G_E$ Function | df | $\chi^2$ without Coulomb correction | AIC | BIC | Norm Extrapolated Radius [fm] | $\chi^2$ with Coulomb correction | AIC | BIC | Norm Extrapolated Radius [fm] |
|----------------|----|----------------------------------|-----|-----|--------------------------------|---------------------------------|-----|-----|--------------------------------|
| linear         | 167.2 1.639 53.35 58.64 0.987(1) 0.781(2) | 172.0 1.686 56.33 61.62 0.984(1) 0.796(2) |
| quadratic      | 101 | 119.4 1.182 20.33 28.26 1.000(2) 0.852(10) | 119.0 1.190 19.97 27.91 0.998(2) 0.860(10) |
| cubic          | 100 | 117.3 1.173 20.55 31.12 0.992(6) 0.785(55) | 117.1 1.171 20.33 30.91 0.992(6) 0.797(55) |
| rational       | 102 | 119.8 1.188 20.87 28.80 1.001(2) 0.860(14) | 119.6 1.184 20.54 28.48 0.999(2) 0.870(14) |

Although these fits look beautiful, the extraction of the radius depends on our belief that we can reasonably ex-

IX. LOWEST $Q^2$ DATA

While the set of Mainz data studied above had a range similar to the large range of our Monte Carlo studies, it is not the lowest $Q^2$ set of data. The lowest set comes from a run with 180 MeV [79], and covers a range from 0.090 to 0.330 fm$^{-2}$. This range is particularly intriguing as it has been shown to be low enough that even a linear regression should be able to extract the radius; though instead of just relying on the Monte Carlo study to make that choice, we once again systematically look at the data with various function and variance choices for the magnetic form factor in Table [V]. The residuals and normal Q-Q plots for the AIC and BIC selected models are shown in Fig. [11] and [12] respectively.

analysis graphs, they are not found in many cited proton radius papers. In fact, some papers have no data quality plots at all (e.g. [21]). As the human eye will find patterns in statistical noise, the normal Q-Q plot is an important tool to check whether or not the data is normally distributed [78]. This is done in Fig. 10 where the sorted residuals are plotted against, and are shown to follow, a normal distribution. There are of course more formal statistical tests one can apply, but in general the visualizations of the data can quickly reveal if there are any major problems. In particular, with least squares fitting, one should be mindful that a single outlier can skew the results.
TABLE VI. Using four different magnetic form factor parameterizations, we extract the normalization and electric radius using a linear, quadratic, cubic, and rational approximation for the $G_E$ function in Eq. [12]. To avoid multiple floating normalizations, the single set of 106 data points shown in Fig. 8 is used. The parameter uncertainties were obtained by performing statistical bootstraps of the data. As seen during the Monte Carlo studies, the linear fit over this interval produces a small variance; but is clearly biased from the true 0.84–0.88 fm proton radius, whereas the quadratic has a larger variance but gives less biased result. The cubic fit function is over-fitting which is why it produces a huge variance. The rational function is nearly as good as the quadratic through produces a systematically larger radius though nicely in the range we expect. In the Table df stands for degrees of freedom.

| Standard Magnetic Form Factor | without Coulomb correction | with Coulomb correction |
|------------------------------|---------------------------|------------------------|
| $G_E$ Fit df df Function     | $\chi^2$ reduced AIC BIC Norm Extrapolated Radius [fm] | $\chi^2$ reduced AIC BIC Norm Extrapolated Radius [fm] |
| linear 104                   | 69.2 0.666 $-41.14 -35.81$ 1.000(1) 0.826(8) | 69.6 0.670 $-40.53 -35.21$ 0.997(1) 0.842(8) |
| quadratic 103                | 67.0 0.651 $-42.58 -34.59$ 1.006(3) 0.923(50) | 66.9 0.649 $-42.80 -34.81$ 1.004(3) 0.948(50) |
| cubic 102                    | 66.3 0.650 $-41.79 -31.13$ 0.996(6) 0.598(nan) | 66.2 0.649 $-41.95 -31.30$ 0.994(6) 0.645(nan) |
| rational 103                 | 63.8 0.632 $-43.77 -30.46$ 0.934(3) 0.936(55) | 67.0 0.650 $-42.63 -34.64$ 1.004(3) 0.964(55) |

| Kelly Magnetic Form Factor   | without Coulomb correction | with Coulomb correction |
|------------------------------|---------------------------|------------------------|
| $G_E$ Fit df df Function     | $\chi^2$ reduced AIC BIC Norm Extrapolated Radius [fm] | $\chi^2$ reduced AIC BIC Norm Extrapolated Radius [fm] |
| linear 104                   | 69.3 0.666 $-41.03 -35.70$ 1.000(1) 0.824(8) | 69.7 0.670 $-40.41 -35.08$ 0.998(1) 0.840(8) |
| quadratic 103                | 67.0 0.651 $-42.58 -34.59$ 1.002(2) 0.923(50) | 66.9 0.649 $-42.80 -34.81$ 1.004(2) 0.948(50) |
| cubic 102                    | 66.2 0.649 $-41.86 -31.21$ 0.995(6) 0.525(nan) | 66.1 0.648 $-42.03 -31.38$ 0.993(6) 0.578(nan) |
| rational 103                 | 67.1 0.652 $-42.43 -34.44$ 1.006(3) 0.937(55) | 67.0 0.651 $-42.62 -34.63$ 1.005(3) 0.965(55) |

| Bernauer Magnetic Form Factor | without Coulomb correction | with Coulomb correction |
|------------------------------|---------------------------|------------------------|
| $G_E$ Fit df df Function     | $\chi^2$ reduced AIC BIC Norm Extrapolated Radius [fm] | $\chi^2$ reduced AIC BIC Norm Extrapolated Radius [fm] |
| linear 104                   | 69.2 0.665 $-41.19 -35.87$ 1.001(1) 0.826(8) | 69.9 0.669 $-40.59 -35.26$ 0.998(1) 0.842(8) |
| quadratic 103                | 67.0 0.651 $-42.58 -34.59$ 1.006(2) 0.923(50) | 66.9 0.649 $-42.80 -34.81$ 1.004(4) 0.948(50) |
| cubic 102                    | 66.2 0.650 $-41.79 -31.13$ 0.996(6) 0.598(nan) | 66.2 0.649 $-41.95 -31.30$ 0.994(6) 0.645(nan) |
| rational 103                 | 67.1 0.652 $-42.44 -34.44$ 1.007(3) 0.936(55) | 67.0 0.650 $-42.63 -34.64$ 0.999(3) 0.964(55) |

| Ye Magnetic Form Factor      | without Coulomb correction | with Coulomb correction |
|------------------------------|---------------------------|------------------------|
| $G_E$ Fit df df Function     | $\chi^2$ reduced AIC BIC Norm Extrapolated Radius [fm] | $\chi^2$ reduced AIC BIC Norm Extrapolated Radius [fm] |
| linear 104                   | 69.3 0.666 $-41.04 -35.71$ 1.000(1) 0.824(8) | 69.7 0.670 $-40.42 -35.09$ 0.998(1) 0.841(8) |
| quadratic 103                | 67.0 0.651 $-42.58 -34.59$ 1.006(2) 0.923(50) | 66.9 0.649 $-42.80 -34.81$ 1.004(2) 0.949(50) |
| cubic 102                    | 66.3 0.650 $-41.79 -31.13$ 0.996(6) 0.598(nan) | 66.2 0.649 $-41.95 -31.30$ 0.994(6) 0.645(nan) |
| rational 103                 | 67.1 0.652 $-42.43 -34.44$ 1.007(3) 0.936(55) | 67.0 0.651 $-42.62 -34.63$ 1.005(3) 0.964(55) |

As has been shown by theory calculations such as Alarcon and Weiss [63, 64] the moments of the generating function are likely far too complex to be constrained by descriptive fitting of experimental data. Also, as pointed out on page 378 of [67], moments do not uniquely define functions, so the best we can do for the higher order terms is determine the shape of the data and then compare with theory.

One can of course try to combine multiple sets of data, though this quickly turns into a Bayesian exercise with no unique solution and has an inherent human-in-the-loop factor [65]. One could also try using a physical model and just simply compare the model with a descriptive fit of the data [63, 85, 66, 69].

In addition, in Appendix B we show an example of using a machine learning significance testing technique, forward stepwise regression, to find the best polynomial fit function. And in Appendix C we also show the effect of conformal mapping in order to clearly show that even after doing a mapping one will still need to apply a rigorous model selection criteria.
FIG. 9. The cross section data and the quadratic fit curve are shown in the upper plot and the residual, the difference between the data and fit, are shown in the lower plot. If an appropriate fit function was used, the residuals should be randomly dispersed around the horizontal axis.

FIG. 10. In order to check if the distribution of the residuals are plausibly normally distributed, one can make a normal quantile-quantile, Q-Q, plot. This is done by sorting the residuals from lowest to highest and plotting the ordered residuals against a theoretical normal distribution. The resulting plot is a beautiful example of normally distributed residuals.

FIG. 11. The cross section data and the quadratic fit curve are shown in the upper plot and the residual, the difference between the data and fit, are shown in the lower plot.

FIG. 12. The low $Q^2$ data quite as normally distributed residuals though still reasonable close to what one would expect.

X. SUMMARY

This work was not meant to be an exhaustive study of the world electron scattering data; but instead an exploration of bias-variance trade-off as it relates to the extraction of the proton radius from electron scattering data. To do this, we have revisited a classic Monte Carlo study [49] and shown that simply rejecting models with a bias is incorrect. Although adding more regression parameters usually reduces the bias, it comes at the cost of increased variance. We also illustrated how parsimonious models can have better predictive power than even the true underlying model in certain situations (see also section D for semi analytical calculations on this regard).

Next, we carefully studied high precision sets of Mainz low $Q^2$ data [24] which covered a range very similar to the Monte Carlo study. Here we have defined a set to be a group of data with a single normalization parameter. The model selection techniques presented herein provide a rigorous method of selecting an appropriate model to describe a given set of data. We have illus-
treated that these model selection techniques along with some key statistical analysis plots can help to ensure a reasonable fit.

These ideas and techniques of model selection are not limited to the physical sciences, but also extend to quantitative analysis [60], and sit at the heart of statistical learning [60]. It has been argued that over-fitting is perhaps more problematic now than in the past due modern computing power [91]. It is amusing to note it was just 1985 when Feynman noted that computers could not beat humans at the game of Go [92], while today computers dominate even this complicated game [93, 94]. With all this computer power available, it is perhaps more necessary than ever to keep in mind the power and importance of parsimonious modeling as nicely summarized by the renowned statistician George Box: “Since all models are wrong the scientist cannot obtain a ‘correct’ one by excessive elaboration. On the contrary, following William of Occam, [the scientist] should seek an economical description of natural phenomena. Just as the ability to devise simple but evocative models is the signature of the great scientist so over-elongation and over-parameterization is often the mark of mediocrity.” [59]

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We would like to thank Miha Mihovilović for many extremely useful discussions and help developing a number of the figures. The regressions done in the body of this work were done in Python making use of the outstanding LMFIT package [95] to interface with SciPy libraries [90]. Non-linear regressions were done using the LevenbergMarquardt method [97, 98]. Special thanks to Edward Tufte and his efforts to help improve the visualization of evidence [96]. Non-linear regressions were done using the LMFIT package [95] to interface with SciPy libraries [90]. These ideas and techniques of model selection are not limited to the physical sciences, but also extend to quantitative analysis [60], and sit at the heart of statistical learning [60]. It has been argued that over-fitting is perhaps more problematic now than in the past due modern computing power [91].

Appendix A: Anscombe’s Quartet

With the power of modern computing, one can be tempted to blindly assume the results of a calculation are correct; but this can be extremely misleading [103]. To illustrate this point we use the Anscombe quartet [77], though as nuclear physicists tend to use reduced $\chi^2$ instead of $R^2$, we have taken the 1973 example problem and added uncertainties to the points as shown in Table VII.

These four sets of (x,y,dy) values give to three significant figures the same statistical quantities: mean, variance, $\chi^2$, reduced $\chi^2$, etc. So if one fails to make graphical checks, one can be completely fooled into thinking the fits are all equally good; but by simply graphing (see Fig. 13) one can see that only the first set of data is distributed in an ideal way around the fit function.

| Obs. no. | x   | y   | y   | x   | y   | dy  |
|---------|-----|-----|-----|-----|-----|-----|
| 1       | 10.0| 8.04| 8.14| 7.46| 8.0 | 6.58|
| 2       | 8.0 | 6.95| 8.14| 6.77| 8.0 | 5.76|
| 3       | 13.0| 7.58| 8.74|12.74|8.0 | 7.71|
| 4       | 9.0 | 8.81| 8.77| 7.11| 8.0 | 8.84|
| 5       | 11.0| 8.33| 9.26| 7.81| 8.0 | 8.47|
| 6       | 14.0| 9.96| 8.10| 8.84| 8.0 | 7.04|
| 7       | 6.0 | 7.24| 6.13| 6.08| 8.0 | 5.25|
| 8       | 4.0 | 4.26| 3.10| 5.39| 19.0|12.50|
| 9       | 12.0|10.84|9.13| 8.15| 8.0 | 5.56|
| 10      | 7.0 | 4.82| 7.26| 6.42| 8.0 | 7.91|
| 11      | 5.0 | 5.68| 4.74| 5.73| 8.0 | 6.89|

Data set two clearly has a curved residual yet has exactly the same mean, errors, $\chi^2$ as the first fit. This suggests that the fitter should likely add a quadratic term to the regression as well as check that the uncertainties have been correctly reported.

Data set three illustrates the effect of an outlier on the regression. Of course, the scientist doesn’t simply throw out an outlier. Instead one should report on the outlier’s influence on the result. For example, in data set three it would be worth noting that if the outlier is removed, the data exactly follow a line of $y = 4 + 0.346x$ and that that measurement should be repeated.

Set four looks unsatisfactory, since all the information about the slope comes from one observation. This is very different from data set one where any one point can be removed and one will obtain nearly the same result. Thus, for this data set four it should be pointed out that a single observation plays a critical role in the result.

FIG. 13. Graphs of the four sets of data.

Table VII. Four data sets of (x,y,dy) values.

| Data set | 1-3 | 1 | 2 | 3 | 4 | 4 | 1-4 |
|----------|-----|---|---|---|---|---|-----|
| Variable | x   | y | y | x | y | dy |     |
| Obs. no. |     |   |   |   |   |    |     |
| 1        | 10.0| 8.04|9.14|7.46|8.0 |6.58|1.235|
| 2        | 8.0 | 6.95|8.14|6.77|8.0 |5.76|1.235|
| 3        | 13.0|7.58|8.74|12.74|8.0 |7.71|1.235|
| 4        | 9.0 | 8.81|8.77|7.11|8.0 |8.84|1.235|
| 5        | 11.0|8.33|9.26|7.81|8.0 |8.47|1.235|
| 6        | 14.0|9.96|8.10|8.84|8.0 |7.04|1.235|
| 7        | 6.0 | 7.24|6.13|6.08|8.0 |5.25|1.235|
| 8        | 4.0 | 4.26|3.10|5.39|19.0|12.50|     |
| 9        | 12.0|10.84|9.13|8.15|8.0 |5.56|1.235|
| 10       | 7.0 | 4.82|7.26|6.42|8.0 |7.91|1.235|
| 11       | 5.0 | 5.68|4.74|5.73|8.0 |6.89|1.235|
Appendix B: Stepwise Regression

In this paper, we have proceeded as is natural for a human statistical modeler, starting from a simple linear approximation and then adding additional terms in $Q^2$ as needed. Of course nothing says that this type of fitting determines the terms of the moments the true generating function but simply that we have found an appropriate function for fitting the data. In fact, using experimentally determined “moments” as a regression constraint can lead to circular logic as the normalization parameters and moments are linked in a very complex way.

In Ref. [23] the authors used forward stepwise regression on the Mainz Rosenbluth data [24] and the code returned a beautiful alternating sign power series which is what one would expect if the power series fit was in fact extracting moments of a generating function. Nevertheless, without a physical theory in mind, no claim was made that these were the moments of the generating function.

In fact, if one applies the same stepwise regression code to very low $Q^2$ data it can give some seemingly nonsensical results. While the linear term will remain essentially unchanged one no longer get the series of alternating . As an example, using R [104] with the CAR package [105] to perform stepwise regression with AIC as the model selection criteria on the $G_E$ data from Griffioen, Carlson and Maddox [22] with $Q^2 < 0.8 \text{ fm}^{-2}$, to match a range studied in the Monte Carlo simulations, one obtains the following result:

```
Start: AIC=36.47
data$y ~ data$x

Df Sum of Sq RSS AIC
+ I(data$x^4) 1 9.1204 357.98 30.090
+ I(data$x^3) 1 9.1063 358.00 30.104
+ I(data$x^5) 1 8.9767 358.13 30.224
+ I(data$x^2) 1 8.8697 358.23 30.324
+ I(data$x^6) 1 8.7322 358.37 30.451
+ I(data$x^7) 1 8.4320 358.67 30.730
+ I(data$x^8) 1 8.1084 359.00 31.030
+ I(data$x^9) 1 7.7822 359.32 31.333
+ I(data$x^{10}) 1 7.4655 359.64 31.626
+ I(data$x^{11}) 1 7.1643 359.94 31.905
<none> 367.10 36.468

Step: AIC=30.09
data$y ~ data$x + I(data$x^4)

Df Sum of Sq RSS AIC
<none> 357.98 30.090
+ I(data$x^2) 1 0.028238 357.96 32.064
+ I(data$x^3) 1 0.021000 357.96 32.071
+ I(data$x^7) 1 0.020418 357.96 32.071
+ I(data$x^8) 1 0.020399 357.96 32.071
+ I(data$x^9) 1 0.019865 357.96 32.073
+ I(data$x^6) 1 0.018854 357.96 32.073
+ I(data$x^{10}) 1 0.018804 357.96 32.073
+ I(data$x^{11}) 1 0.016573 357.97 32.075

Weighted Residuals:
```

FIG. 14. The $G_E$ data of [22] shown along with a residual of the best fit using forward stepwise regression.

Unlike the previous studies in this work where tests were simply made by increasing the order of the power series by one, the forward stepwise regression scans over all possible next terms in the power series up to a user defined 11th order. The results shown above are in the algebraic notation of R and the x term is $Q^2$. Starting from a linear function, different combinations of higher order functions are tried by adding and removing terms, in order to find the best performing predictive model. The final result of the fit is illustrated in Fig. 14 and code to perform the fits can be found on github [106].

Appendix C: Conformal Mapping

It has been argued that the use of conformal mapping can improve the situation; but while some authors of this technique confine themselves to low $Q^2$ and model constrained parameters, others seem to use the mapping to obfuscate clear over-fitting. For example, in Table 1 of [40] the authors do various fits up to fifth order with no clear model selection technique in mind. By simply using
the selection criteria provided herein, AIC or BIC, one finds that in general the two parameter fits are sufficient and the data used therein give 0.86(2)\,fm for the radius. By model selection criteria, all the fits beyond second order are simply over-fitting the data.

On the other hand, the idea of conformal mapping is very powerful. From a pure mathematical point of view, it can take a function, like \( G_E(Q^2) \) that goes from zero in infinity and map it onto a finite range where one could use orthogonal polynomial to describe data. It can also be used to avoid the effects of poles when doing polynomial regressions, though it is worth noting that rational functions also have that property \[107\].

The transformation from \( Q^2 \) to the mapped parameter \( z \) is done by:

\[
    z(t, t_{\text{cut}}, t_0) = \frac{\sqrt{t_{\text{cut}} - t} - \sqrt{t_{\text{cut}} - t_0}}{\sqrt{t_{\text{cut}} - t} + \sqrt{t_{\text{cut}} - t_0}},
\]

where \( t = -Q^2, t_{\text{cut}} = 4m^2, \) and \( t_0 \) is a free parameter representing the point being map onto \( z = 0 \). Herein we have used \( t_0 = 0 \). It is argued without proof that "the curvature is smaller in the \( z \) variable than in the \( Q^2 \) variable" \[10\], yet by taking exactly the same data as before and fitting with stepwise regression one gets the following result:

Start: AIC=62.02
data$y - data$x

| Df | Sum of Sq | RSS | AIC |
|----|-----------|-----|-----|
| 2  | 36.992    | 358.23 | 31.779 |
| 3  | 35.577    | 359.65 | 33.092 |
| 4  | 33.518    | 361.71 | 34.993 |
| 5  | 31.295    | 363.93 | 37.034 |
| 6  | 29.136    | 366.09 | 39.003 |
| 7  | 27.125    | 369.95 | 40.827 |
| 8  | 25.277    | 371.82 | 42.495 |
| 9  | 23.582    | 373.20 | 44.017 |
| 10 | 22.026    | 374.63 | 45.408 |
| 11 | 20.595    | 374.63 | 46.683 |
|<none>| 395.23 | 62.019 |

Step: AIC=31.78
data$y - data$x + I(data$x^2)

| Df | Sum of Sq | RSS | AIC |
|----|-----------|-----|-----|
| 2  | 0.46895   | 357.77 | 33.828 |
| 3  | 0.46636   | 357.77 | 33.830 |
| 4  | 0.46557   | 357.77 | 33.831 |
| 5  | 0.46025   | 357.77 | 33.836 |
| 6  | 0.45531   | 357.78 | 33.840 |
| 7  | 0.45194   | 357.78 | 33.844 |
| 8  | 0.45177   | 357.78 | 33.844 |
| 9  | 0.41901   | 357.82 | 33.874 |
| 10 | 0.36310   | 357.87 | 33.926 |
|<none>| 358.23 | 31.779 |

Step: AIC=31.78
data$y - data$x + I(data$x^2)

| Df | Sum of Sq | RSS | AIC |
|----|-----------|-----|-----|
| 2  | 0.46636   | 357.77 | 33.830 |
| 3  | 0.46557   | 357.77 | 33.831 |
| 4  | 0.45531   | 357.78 | 33.840 |
| 5  | 0.45194   | 357.78 | 33.844 |
| 6  | 0.45177   | 357.78 | 33.844 |
| 7  | 0.41901   | 357.82 | 33.874 |
| 8  | 0.36310   | 357.87 | 33.926 |
|<none>| 358.23 | 31.779 |

Weighted Residuals:

\[
\begin{array}{cccc}
\text{Min} & \text{1Q} & \text{Median} & \text{3Q} & \text{Max} \\
-3.07706 & -0.75955 & -0.09171 & 0.67092 & 3.04833 \\
\end{array}
\]

Coefficients:

\[
\begin{array}{ccc}
\text{Estimate} & \text{Std. Error} & \text{t value} & \text{Pr}(>|t|) \\
(\text{Intercept}) & 0.9992836 & 0.0005757 & 1735.683 & < 2e-16 \\
data$x & -0.9741340 & 0.0256596 & -37.964 & < 2e-16 \\
I(data$x^2) & 0.2627947 & 0.9992836 & 1.5340681 & 0.0265696 & -37.964 & < 2e-16 \\
\end{array}
\]

Residual standard error: 1.042 on 330 degrees of freedom
Multiple R-squared: 0.9932, Adjusted R-squared: 0.9932
F-statistic: 2.41e+04 on 2 and 330 DF, p-value: < 2.2e-16

FIG. 15. Illustration of the result of performing the \( z \) transformation on the standard dipole function shown the solid line. For the standard dipole at low \( Q^2 \), this turns a slightly concave function into a convex function. The dashed line is simply a straight line to make the concavity of the functions clear. As described in the text, this same effect is seen with real \( G_E \) data.

Thus, the stepwise regression shows that the transformation has increased the curvature of the data and turned the slightly convex data in \( Q^2 \) to clearly concave data in \( z \). Assuming that a quadratic can be used to make a reasonable extrapolation, one can use this R re-

FIG. 16. The \( z \)-transformed \( G_E \) data shown along with a residual of the best fit.

Thus, the stepwise regression shows that the transformation has increased the curvature of the data and turned the slightly convex data in \( Q^2 \) to clearly concave data in \( z \). Assuming that a quadratic can be used to make a reasonable extrapolation, one can use this R re-
gression result to extract a radius using the formula:

\[ r_p = \frac{\sqrt{-6(b - d)} - (data \cdot x)}{4m_n}. \]  

(C2)

Thus, one finds 0.85(1) fm, a result again consistent with the muonic Lamb shift result, as well as the other results shown in this work. These ideas can be taken further and one could use ideas such as Gaussian process regression to calculate many possible extrapolating paths and then assign relative probabilities \[108\] as recently done by Zhou et al. \[109\].

Appendix D: Semi-analytical Calculations

In this section we demonstrate a semi-analytical procedure to calculate the bias and variance induced by the noise in our estimation of the slope at zero. This framework can be used to reproduce the graph shown in Fig. 2 and Fig. 3 without the repeated Monte Carlo procedure of creating many noisy data sets and sampling from the fitted functions. The virtue of this method is that it can give us explicitly the influence each data point is having. Before proceeding we must realize that the bias in our estimation of the slope has two sources. The first one is due to the noise in the data and will increase as the noise increases. The second is due to the possible fact that we are not using the “true” function to fit the data, as was shown in Table IV when the line had a clear bias even when the noise was very small. We therefore split our bias as:

\[ \text{bias} = \text{bias}_\sigma + \text{bias}_0, \]  

(D1)

where \( \text{bias}_\sigma \) denotes the bias that scales with the noise, while \( \text{bias}_0 \) will always be present even when \( \sigma \rightarrow 0 \), where \( \sigma \) represents the size of the noise.

In this section we will show how to calculate the first source, while the second can only be calculated if we know the “true” value of the slope. Let us define our quantity of interest, the slope at zero, as \( m(X, Y, \epsilon) \), where \( X \) and \( Y \) are the lists of \( n \) data points and \( \epsilon \) is a particular realization of the noise, which we assume to be Gaussian with mean zero and deviation \( \sigma \). Here \( m(X, Y, \epsilon) \) can be a closed expression for the slope given the data, as is the case for linear models, or it can be treated as a numerical routine that returns the slope when fitting a non-linear model, like the Dipole.

1. Noise Bias

We want to find the average value of \( m(X, Y, \epsilon) \), \( \langle m(X, Y) \rangle \), once all the possible realizations of the noise have been taken into account, weighted correctly by their Gaussian distribution. The result is given by the integral:

\[ \langle m(X, Y) \rangle \equiv \int P(\epsilon) m(X, Y, \epsilon) \, d\epsilon, \]  

(D2)

where \( d\epsilon = d\epsilon_1 d\epsilon_2 \ldots d\epsilon_n \), and \( P(\epsilon) \) denotes the Gaussian probability distribution for the noise given by:

\[ P(\epsilon) = \left( \frac{1}{(\sqrt{2\pi}\sigma^2)} \right)^n e^{(\epsilon_1^2 + \epsilon_2^2 + \ldots + \epsilon_n^2)/2\sigma^2}. \]  

(D3)

The integral in Eq. \( \text{D2} \) is effectively taking into account all possible noise realizations, weighted by their corresponding probabilities. Since in most cases we do not have an available expression for \( m(X, Y, \epsilon) \), we can expand it using a multivariate Taylor expansion and evaluate each term directly under the integral sign in Eq. \( \text{D2} \). The first three terms in this expansion are:

\[ m(X, Y, \epsilon) \approx m_0 + [\nabla \epsilon m]_0 \cdot \epsilon + \frac{1}{2} \epsilon^T \cdot [H_m]_0 \cdot \epsilon, \]  

(D4)

where \( m_0 = m(X, Y, \epsilon = 0) \) is the value of the slope when the noise is zero. We denote \([\nabla \epsilon m]_0 \) the gradient of \( m \) taking the errors \( \epsilon \) as variables, \( [\nabla \epsilon m]_0 \equiv \left( \frac{\partial m}{\partial \epsilon_1}, \frac{\partial m}{\partial \epsilon_2}, \ldots, \frac{\partial m}{\partial \epsilon_n} \right) \),  evaluated at \( \epsilon = 0 \). Finally, \([H_m]_0 \) denotes the Hessian matrix of \( m \), \( [H_m]_{i,j} \equiv \frac{\partial^2 m}{\partial \epsilon_i \partial \epsilon_j} \), again taking the \( \epsilon \) as variables and evaluating at \( \epsilon = 0 \).

Equation \( \text{D4} \) says that for some small realization of the errors \( \epsilon_i \), the value of \( m \) is approximately its value when \( \epsilon = 0 \), plus a linear correction in \( \epsilon \) by the gradient and finally a second order correction proportional to \( \epsilon^2 \) that involves the Hessian. Both of these quantities, the gradient and the Hessian are evaluated at \( \epsilon = 0 \) and are therefore just a list and a matrix of fixed numbers, respectively. These two groups of numbers can be obtained numerically by taking finite differences on \( m \) as an approximation to derivatives.

Once we have expanded our slope function \( m \) we proceed to calculate the integral \( \text{D2} \):

\[ \langle m(X, Y) \rangle \approx \int \left( m_0 + [\nabla \epsilon m]_0 \cdot \epsilon + \frac{1}{2} \epsilon^T \cdot [H_m]_0 \cdot \epsilon \right) P(\epsilon) \, d\epsilon. \]  

(D5)

The linear correction, being proportional to \( \epsilon \), would integrate to zero. The term \( m_0 \) does not depend on the noise and since \( P(\epsilon) \) integrates to 1 it would just appear as it is in the final result. The last term with the Hessian is a quadratic form that would look like \( \epsilon_1^2 H_{11} + 2\epsilon_1 \epsilon_2 H_{12} + \ldots + \epsilon_n^2 H_{nn} \), where the constants
\( H_{ij} \) are the \((i,j)\) elements of the Hessian. Since our noise additions are mutually independent (\( \epsilon_i \) and \( \epsilon_j \) do not correlate) the integration of terms involving mixing of different \( \epsilon_i \) will also yield zero. The integration over terms that involve the same \( \epsilon_i \) are by definition the variance of the noise, \( \sigma^2 \). Therefore we have:

\[
\langle m(X,Y) \rangle \approx m_0 + \frac{1}{2} \int (\nabla^T \cdot [H_m]_0 \cdot \nabla) P(\epsilon) \, d\epsilon = m_0 + \frac{1}{2} \sigma^2 \left[ \frac{\partial^2 m}{\partial \epsilon_1^2} + \frac{\partial^2 m}{\partial \epsilon_2^2} + \ldots + \frac{\partial^2 m}{\partial \epsilon_n^2} \right]_0.
\]

This equation shows that there will be a noise-induced bias in the average estimation of our quantity \( m \), which grows proportionally to \( \sigma^2 \). This bias will not be present if the function that estimates \( m \) in terms of the data is linear in the observations \( Y \) (and therefore linear in the noise \( \epsilon \)), since second derivatives of \( m \) will be zero from the start. This means that, regardless of the size of the noise, linear fits like a straight line, or any polynomial, will not gain a noise-induced bias, and the distributions for \( m \) will always center at the zero noise point.

For non-linear fits on the other hand, there will be a bias that will grow quadratically in the noise. This can apply to the actual function that generated the data, like the Dipole in our study. Once the noise gets too high, the true generating function might not be the most reliable, and therefore linear in the observations \( Y \). This bias will grow quadratically in the noise. This can apply to the actual function that generated the data, like the Dipole in our study. Once the noise gets too high, the true generating function might not be the most reliable, and therefore linear in the observations \( Y \).

The term accompanying \( \sigma^2 \), the trace of the Hessian matrix, can be interpreted as the Laplacian of \( m \) at zero noise. Consistent with our findings, the Laplacian operator of \( f(x) \) at a point \( a \) can be related to the rate of change of the average of \( f \), \( \langle f \rangle \), at a small sphere centered at \( a \) compared to \( f(a) \).

2. Standard Deviation

In order to calculate the standard deviation \( s = \sqrt{\langle m^2 \rangle - \langle m \rangle^2} \) of our distribution for \( m \) we must compute the expected value of \( m^2 \), since we already have \( \langle m \rangle \). We can calculate this quantity up to second order in \( \sigma \), using expression [D1]

\[
\langle m(X,Y)^2 \rangle \approx \int \left[ m_0 + \left[ \nabla m \right]_0 \cdot \nabla + \frac{1}{2} \sigma^2 + \frac{1}{2} \left[ \nabla^T \cdot [H_m]_0 \cdot \nabla \right] \right] P(\epsilon) \, d\epsilon = \int \left[ m_0^2 + 2 m_0 \left[ \nabla m \right]_0 \cdot \nabla + \left[ \nabla^T \cdot [H_m]_0 \cdot \nabla \right] \right] P(\epsilon) \, d\epsilon + 2 m_0 \sigma^2 + \mathcal{O}(\epsilon^4) + \mathcal{O}(\epsilon^2) \, d\epsilon.
\]

The first term does not depend on \( \epsilon \) and can go out of the integral. The second term is proportional to \( \epsilon \) and integrates to zero. The Hessian term we already know how to integrate and we have neglected its square, which will be order \( \epsilon^4 \). The third term, the gradient square, reads:

\[
\left( \left[ \nabla m \right]_0 \cdot \nabla \right)^2 = \left( \frac{\partial m}{\partial \epsilon_1} \right)_0^2 \epsilon_1^2 + \left( \frac{\partial m}{\partial \epsilon_2} \right)_0^2 \epsilon_2^2 + \ldots + \left( \frac{\partial m}{\partial \epsilon_n} \right)_0^2 \epsilon_n^2.
\]

Once we expand the square we will have two types of terms, those of the form \( \left( \frac{\partial m}{\partial \epsilon_i} \right)_0^2 \epsilon_i^2 \) and those with mixed \( \epsilon_i \), \( \left( \frac{\partial m}{\partial \epsilon_i} \frac{\partial m}{\partial \epsilon_j} \right)_0 \epsilon_i \epsilon_j \). Since our noise components are independent, only the first type of terms will give non-zero results under integration: \( \sigma^2 \). Therefore we have:

\[
\langle m(X,Y)^2 \rangle \approx m_0^2 + m_0 \sigma^2 \text{Tr}[H_m]_0 + \sigma^2 \left( \frac{\partial m}{\partial \epsilon_1} \right)_0^2 + \left( \frac{\partial m}{\partial \epsilon_2} \right)_0^2 + \ldots + \left( \frac{\partial m}{\partial \epsilon_n} \right)_0^2 \right].
\]

The first coefficient accompanying \( \sigma^2 \) is the trace of the Hessian matrix, which will cancel the same exact term that appears in the square of the mean value for \( m \):

\[
\langle m \rangle^2 \approx (m_0 + \frac{1}{2} \sigma^2 \text{Tr}[H_m]_0)^2 \approx m_0^2 + m_0 \sigma^2 \text{Tr}[H_m]_0.
\]

The second coefficient accompanying \( \sigma^2 \) is the magnitude of the gradient of \( m \), evaluated at zero noise. Our result for the standard deviation \( s \), if we only keep terms up to first order in \( \sigma \) for \( \langle m \rangle^2 \) then reads:

\[
s = \sqrt{\langle m^2 \rangle - \langle m \rangle^2} = \sigma \left[ \nabla m \right]_0 \cdot \nabla.
\]

Therefore, our calculations show that no matter how small the noise in the data, there would be an appreciable, proportional, standard deviation in our estimate of \( m \). This is true unless the gradient is zero, meaning that our quantity \( m \) is not sensitive to change in the data to first order.

All the calculations presented here can easily be generalized to include correlations between the \( \epsilon_i \) and to include different sizes of noise for each point: \( \sigma \) instead of a single \( \sigma \). The only requirement is that the noise is considered Gaussian and the integrals would still be easily solvable.

It is also possible to extend the Taylor expansion of \( m \) in equation [D4] to include more orders in the noise \( \epsilon \). Since the Taylor expansion will contain some combination of different \( \epsilon_i \), the Gaussian integrals can still be performed in closed form, term by term.

3. Maximizing Information Gained

Some fundamental questions that arises in experimental designs and theory of information are: for a given relation \( f(x) = y \) between two quantities, what are the
most relevant values of $x$ such that, when $y$ is measured, the relation $f$ is constrained the most? How much new information one gains by adding more points or reducing the statistical uncertainties on the existing ones? In other words, what are the $(x, y)$ values that, considered as a set, contain the most amount of information regarding $f$, for a given set of experimental uncertainties.

With respect to the proton radius extraction that we have studied here, a question of interest would be: for a given range in $Q^2$, what specific values $Q_i^2$ give us the most amount of information regarding the slope at zero, or the proton radius.

Using the semi-analytical framework described, we can attempt to give a quantitative answer to this question. Both Eqs. [D6] and [D9] have explicit the contribution each point makes to the noise bias and the standard deviation, respectively. Therefore, we could directly identify how a reduction in uncertainty on each particular point would propagate to our estimate of the radius.

Alternatively, if we are stuck on some level of uncertainty, we could estimate where the $n$ measurements should be made such that, again, our information gain is maximal. In order to test these ideas, we minimized the RMSE defined in Eq. [D1] as a function of the location of $n$ $Q^2$ points, and studied how much our uncertainty in the slope at zero is reduced as $n$ increased.

Our $y$ points were again generated by the Dipole function defined in Eq. [2] and we studied the fit of a line in the range $0.1 - 0.8$ fm$^{-2}$ and the fit of a parabola in the range $0.1 - 1.6$ fm$^{-2}$, both of which have proven to be optimal in our simulations on section IV.

Since the slope estimate from both models is linear in the observations $y$, we know that there will be no noise-induced bias, and the only source of bias will be a constant term, see Eq. [D1] By “constant” we mean that it will not grow with the noise, but it would change depending on the points $Q_i^2$ used.

We seek therefore to minimize:

$$\text{RMSE}^2(X) = \text{bias}^2(X) + s^2$$

$$= (m_{true} - m_0(X))^2 + \sigma^2||\nabla_y m(X)||_0^2,$$  \hspace{1cm} \text{(D11)}

where $X = (Q_1^2, ... Q_n^2)$ are the $n$ locations of the “observed” points, and we have omitted the $Y$ dependence on our quantities since we are calculating $Y$ directly from our points $X$, and $m_{true} = -0.1097$ is the fixed value of the slope at zero, in fm$^2$.

Now, the bias quantity $(m_{true} - m_0(X))$ will indeed depend on $X$, and is a number that is usually hidden to us by nature, since we do not know the true value. Nevertheless, this exercise is helpful in showing that some points have more information than others, and we will see how both terms evolve as $n$ grows.

Figures [17] and [18] show the obtained results when the points $Q_i^2$ are not distributed uniformly but are rather moved to positions that, respecting the assigned range, minimize the RMSE given by Eq. [D11]. It is very interesting to note that, as we expected from our analysis in section IV, the RMSE of the parabola is primarily driven by the standard deviation contribution, while the line is more balanced, although towards higher bias.

The main result that we can extract from these two graphs is that it takes only 10 points in the case of the line and 13 in the case of the parabola to obtain the same value of RMSE compared to 31 uniformly spaced points, see Table III. If we use 31 points but locate them in optimal positions we obtain an RMSE of 0.0068 fm$^2$ for the line and an RMSE of 0.0056 fm$^2$ for the Parabola. The first one represents an improvement of 21% while the second represents and improvement of 31%.

As we have already mentioned, calculating (and optimizing with respect to) the bias requires the true value to be known, which is usually not the case in interesting problems. Nevertheless, the standard deviation we have...
defined is computable only as a function of the observed values and errors, and therefore it is a quantity we can use when facing the real data. As can be seen in Figure 15, the bias of the parabola does not seem to change much for any configuration of points, but the \( \sigma \), and therefore the RMSE, can be improved substantially.

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