STATISTICAL CHALLENGES FOR SEARCHES FOR NEW PHYSICS AT THE LHC

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Because the emphasis of the LHC is on 5σ discoveries and the LHC environment induces high systematic errors, many of the common statistical procedures used in High Energy Physics are not adequate. I review the basic ingredients of LHC searches, the sources of systematics, and the performance of several methods. Finally, I indicate the methods that seem most promising for the LHC and areas that are in need of further study.

1 Introduction

The Large Hadron Collider (LHC) at CERN and the two multipurpose detectors, ATLAS and CMS, have been built in order to discover the Higgs boson, if it exists, and explore the theoretical landscape beyond the Standard Model.\textsuperscript{1,2} The LHC will collide protons with unprecedented center-of-mass energy ($\sqrt{s} = 14$ TeV) and luminosity ($10^{34}$ cm$^{-2}$s$^{-1}$); the ATLAS and CMS detectors will record these interactions with $\sim 10^8$ individual electronic readouts per event. Because the emphasis of the physics program is on discovery and the experimental environment is so complex, the LHC poses new challenges to our statistical methods – challenges we must meet with the same vigor that led to the theoretical and experimental advancements of the last decade.

In the remainder of this Section, I introduce the physics goals of the LHC and most pertinent factors that complicate data analysis. I also review the formal link and the practical differences between confidence intervals and hypothesis testing.

In Sec. 2, the primary ingredients to new particle searches are discussed. Practical and toy examples are presented in Sec. 3, which will be used to assess the most common methods in Sec. 4. The remainder of this paper is devoted to discussion on the most promising methods for the LHC.

1.1 Physics Goals of the LHC

Currently, our best experimentally justified model for fundamental particles and their interactions is the standard model. In short, the physics goals of the LHC come in two types: those that improve our understanding of the standard model, and those that go beyond it.

The only particle of the standard model that has not been observed is the Higgs boson, which is key for the standard model’s description of the electroweak interactions. The mass of the Higgs boson, $m_H$, is a free parameter in the standard model, but there exist direct experimental lower bounds and more indirect upper bounds. Once $m_H$ is fixed, the standard model is a completely predictive theory. There are numerous particle-level Monte Carlo generators that can be interfaced with simulations of the detectors to predict the rate and distribution of all experimental observables. Because of this predictive power, searches for the Higgs boson are highly tuned and often employ multivariate discrimination methods like neural networks, boosted decision trees, support vector machines, and genetic programming.\textsuperscript{3,4,5}

While the Higgs boson is key for understanding the electroweak interactions, it introduces a new problem: \textit{i.e.} the hierarchy problem. There are several proposed solutions to the problem, one of which is to introduce a new fundamental symmetry, called supersymmetry (SUSY), between bosons and fermions. In practice, the minimal supersymmetric extension to the standard model (MSSM), with its 105 parameters, is not so much a theory as a theoretical framework.

They key difference between SUSY and Higgs searches is that, in most cases, discovering SUSY will not be the difficult part. Searches for SUSY often rely on robust signatures that will show a deviation from the standard model for most regions of the SUSY parameter space. It will be much more challenging to demonstrate that the deviation from the standard model is SUSY and to measure the fundamental parameters of the theory.\textsuperscript{6} In order to restrict the scope of these proceedings, I shall focus LHC Higgs searches, where the issues of hypothesis testing are more relevant.
1.2 The Challenges of LHC Environment

The challenges of the LHC environment are manifold. The first and most obvious challenge is due to the enormous rate of uninteresting background events from QCD processes. The total interaction rate for the LHC is of order $10^9$ interactions per second; the rate of Higgs production is about ten orders of magnitude smaller. Thus, to understand the background of a Higgs search, one must understand the extreme tails of the QCD processes.

Compounding the difficulties due to the extreme rate is the complexity of the detectors. The full-fledged simulation of the detectors is extremely computationally intensive, with samples of $10^7$ events taking about a month to produce with computing resources distributed around the globe. This computational limitation constrains the problems that can be addressed with Monte Carlo techniques.

Theoretical uncertainties also contribute to the challenge. The background to many searches requires calculations at, or just beyond, the state-of-the-art in particle physics. The most common situation requires a final state with several well-separated high transverse momentum objects (e.g. $t\bar{t}jj \rightarrow b\ell vbjjjj$), in which the regions of physical interest are not reliably described by leading-order perturbative calculations (due to infra-red and collinear divergences), are too complex for the requisite next-to-next-to-leading order calculations, and are not properly described by the parton-shower models alone. Enormous effort has gone into improving the situation with next-to-leading order calculations and matrix-element–parton-shower matching. While these new tools are a vast improvement, the residual uncertainties are still often dominant.

Uncertainties from non-perturbative effects are also important. For some processes, the relevant regions of the parton distribution functions are not well-measured (and probably will not be in the first few years of LHC running), which lead to uncertainties in rate as well as the shape of distributions. Furthermore, the various underlying-event and multiple-interaction models used to describe data from previous colliders show large deviations when extrapolated to the LHC. This soft physics has a large impact on the performance of observables such as missing transverse energy.

In order to augment the simulated data chain, most searches introduce auxiliary measurements to estimate their backgrounds from the data itself. In some cases, the background estimation is a simple sideband, but in others the link between the auxiliary measurement to the quantity of interest is based on simulation. This hybrid approach is of particular importance at the LHC.

While many of the issues discussed above are not unique to the LHC, they are often more severe. At LEP, it was possible to generate Monte Carlo samples of larger size than the collected data, QCD backgrounds were more tame, and most searches were not systematics-limited. The Tevatron has much more in common with the LHC; however, at this point discovery is less likely, and most of the emphasis is on measurements and limit setting.

1.3 Confidence Intervals & Hypothesis Testing

The last several conferences in the vein of PhyStat 2005 have concentrated heavily on confidence intervals. In particular, 95% confidence intervals for some physics parameter in an experiment that typically has few events. More recently, there has been a large effort in understanding how to include systematic errors and nuisance parameters into these calculations.

LHC searches, in contrast, are primarily interested in $5\sigma$ discovery. The $5\sigma$ discovery criterion is somewhat vague, but usually interpreted in a frequentist sense as a hypothesis test with a rate of Type I error $\alpha = 2.85 \cdot 10^{-7}$.

There is a formal link between confidence intervals and hypothesis testing: frequentist confidence intervals from the Neyman construction are formally inverted hypothesis tests. It is this equivalence that links the Neyman-Pearson lemma to the ordering rule used in the unified method of Feldman and Cousins. Furthermore, this equivalence will be very useful in translating our understanding of confidence intervals to the searches at the LHC.

In some cases, this formal link can be misleading. In particular, there is not always a continuous parameter that links the fully specified null hypothesis $H_0$ to the fully specified alternate $H_1$ in any

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9The lemma states that, for a simple hypothesis test of size $\alpha$ between a null $H_0$ and an alternate $H_1$, the most powerful critical region in the observable $x$ is given by a contour of the likelihood ratio $L(x|H_0)/L(x|H_1)$.
physically interesting or justified way. Furthermore, the performance of a method for a 95% confidence interval and a 5σ discovery can be quite different.

2 The Ingredients of an LHC Search

In order to assess the statistical methods that are available and develop new ones suited for the LHC, it is necessary to be familiar with the basic ingredients of the search. In this section, the basic ingredients, terminology, and nomenclature are established.

2.1 Multiple Channels & Processes

Almost all new particle searches do not observe the particle directly, but through the signatures left by the decay products of the particle. For instance, the Higgs boson will decay long before it interacts with the detector, but its decay products will be detected. In many cases, the particle can be produced and decay in many different configurations, each of which is called a search channel (see Tab. 1). There are may be multiple signal and background processes which contribute to each channel. For example, in $H \to \gamma\gamma$, the signal could come from any Higgs production mechanism and the background from either continuum $\gamma\gamma$ production or QCD backgrounds where jets fake photons. Each of these processes have their own rates, distributions for observables, and uncertainties. Furthermore, the uncertainties between processes may be correlated.

In general the theoretical model for a new particle has some free parameters. In the case of the standard model Higgs, only the mass $m_H$ is unknown. For SUSY scenarios, the Higgs model is parametrized by two parameters: $m_A$ and $\tan\beta$. Typically, the unknown variables are scanned and a hypothesis test is performed for each value of these parameters. The results from each of the search channels can be combined to enhance the power of the search, but one must take care of correlations among channels and ensure consistency.

The fact that one scans over the parameters and performs many hypothesis tests increases the chance that one finds at least one large fluctuation from the null-hypothesis. Some approaches incorporate the number of trials explicitly, some approaches only focus on the most interesting fluctuation, and some see this heightened rate of Type I error as the motivation for the stringent 5σ requirement.

2.2 Discriminating Variables & Test Statistics

Typically, new particles are known to decay with certain characteristics that distinguish the signal events from those produced by background processes. Much of the work of a search is to identify those observables and to construct new discriminating variables (generically denoted as $m$). Examples include angles between particles, invariant masses, and particle identification criterion. Discriminating variables are used in two different ways: to define a signal-like region and to weight events.

The usage of discriminating variables is related to the test statistic: the real-valued quantity used to summarize the experiment. The test statistic is thought of as being ordered such that either large or small values indicate growing disagreement with the null hypothesis.

A simple “cut analysis” consists of defining a signal-like region bounded by upper- and lower-values of these discriminating variables and counting events in that region. In that case, the test statistic is simply the number of events observed in the signal like region. One expects $b$ background events and $s$ signal events, so the experimental sensitivity is optimized by adjusting the cut values. More sophisticated techniques use multivariate algorithms, such as neural networks, to define more complicated signal-like regions, but the test statistic remains unchanged.
In these number counting analyses, the likelihood of observing \( n \) events is simply given by the Poisson model.

There are extensions to this number-counting technique. In particular, if one knows the distribution of the discriminating variable \( m \) for the background-only (null) hypothesis, \( f_b(m) \), and the signal-plus-background (alternate) hypothesis, \( f_{s+b}(m) = [sf_s(m) + bf_b(m)]/(s + b) \), then there is a more powerful test statistic than simply counting events. This is intuitive, a well measured ‘golden event’ is often more convincing than a few messy ones. Following the Neyman-Pearson lemma, the most powerful test statistic is

\[
Q = \frac{L(m|H_1)}{L(m|H_0)}
\]

\[
= \prod_{i}^{N_{chan}} \text{Pois}(n_i|s_i + b_i) \prod_{j}^{N_{chan}} \text{Pois}(n_i|b_i) \prod_{j}^{N_{chan}} f_0(m_{ij})
\]

\[
= \prod_{i}^{N_{chan}} \text{Pois}(n_i|b_i) \prod_{j}^{N_{chan}} f_0(m_{ij})
\]

\[
q = \ln Q = -s_{tot} + \sum_{i}^{N_{chan}} \sum_{j}^{N_{chan}} \ln \left( 1 + \frac{s_i f_s(m_{ij})}{b_i f_0(m_{ij})} \right).
\]

The test statistic in Eq. 2 was used by the LEP Higgs Working Group (LHWG) in their final results on the search for the Standard Model Higgs.\(^{14}\)

At this point, there are two loose ends: how does one determine the distribution of the discriminating variables \( f(m) \), and how does one go from Eq. 2 to the distribution of \( q \) for \( H_0 \) and \( H_1 \). These are the topics of the next subsections.

2.3 Parametric and Non-Parametric Methods

In some cases, the distribution of a discriminating variable \( f(m) \) can be parametrized and this parametrization can be justified either by physics arguments or by goodness-of-fit. However, there are many cases in which \( f(m) \) has a complicated shape not easily parametrized. For instance, Fig. 2 shows the distribution of a neural network output for signal events. In that case kernel estimation techniques can be used to estimate \( f(m) \) in a non-parametric way from a sample of events \( \{m_i\} \).\(^{15}\) The technique that was used by the LHWG\(^{14}\) was based on an adaptive kernel estimation given by:

\[
f_{\hat{i}}(m) = \sum_{i}^{n} \frac{1}{nh(m_{i})} K \left( \frac{m - m_{i}}{h(m_{i})} \right),
\]

where

\[
h(m_{i}) = \left( \frac{4}{3} \right)^{1/5} \sqrt{\frac{\sigma}{f_0(m_{i})}} n^{-1/5},
\]

\[
\sigma = \text{standard deviation of } \{x_i\}, K(x) \text{ is some kernel function (usually the normal distribution), and } f_0(x) \text{ is the fixed kernel estimate given by the same equation but with a fixed } h(m_{i})
\]

\[
h^* = \left( \frac{4}{3} \right)^{1/5} \sigma n^{-1/5}.
\]

The solid line in Fig. 2 shows that the method (with modified-boundary kernels) works very well for shapes with complicated structure at many scales.

2.4 Numerical Evaluation of Significance

Given, \( f_s(m) \) and \( f_b(m) \) the distribution of \( q(x) \) can be constructed. For the background-only hypothesis, \( f_b(m) \) provides the probability of corresponding values of \( q \) needed to define the single-event pdf \( \rho_1 \).

\[
\rho_{1,b}(q_0) = \int f_b(m) \delta(q(m) - q_0) dm
\]

For multiple events, the distribution of the log-likelihood ratio must be obtained from repeated convolutions of the single event distribution. This convolution can either be performed implicitly with approximate Monte Carlo techniques,\(^{16}\) or analytically with a Fourier transform technique.\(^{17}\) In the Fourier domain, denoted with a bar, the distribution of the log-likelihood for \( n \) events is

\[
\bar{p}_n = \bar{p}_1^n
\]

Thus the expected log-likelihood distribution for background with Poisson fluctuations in the number

\(\text{The integral is necessary because the map } q(m) : m \to q \text{ may be many-to-one.}\)
of events takes the form

$$\rho_b(q) = \sum_{n=0}^{\infty} \frac{e^{-b/n}}{n!} \rho_{n,b}(q)$$

(8)

which in the Fourier domain is simply

$$\overline{\rho_b(q)} = e^{b[\rho_{1,s}(q)-1]}.$$  \hspace{1cm} (9)

For the signal-plus-background hypothesis we expect $s$ events from the $\rho_{1,s}$ distribution and $b$ events from the $\rho_{1,b}$ distribution, which leads to the expression for $\rho_{s+b}$ in the Fourier domain

$$\overline{\rho_{s+b}(q)} = e^{b[\rho_{1,s}(q)-1]+s[\rho_{1,b}(q)-1]].$$  \hspace{1cm} (10)

This equation generalizes, in a somewhat obvious way, to include many processes and channels.

Numerically the computations are carried out with the Fast Fourier Transform (FFT). The FFT is performed on a finite and discrete array, beyond which the function is considered to be periodic. Thus the range of the $\rho_1$ distributions must be sufficiently large to hold the resulting $\rho_b$ and $\rho_{s+b}$ distributions. If they are not, the “spill over” beyond the maximum log-likelihood ratio $q_{\text{max}}$ will “wrap around” leading to unphysical $\rho$ distributions. Because the range of $\rho_b$ is much larger than $\rho_{1,b}$ it requires a very large number of samples to describe both distributions simultaneously. The implementation of this method requires some approximate asymptotic techniques that describe the scaling from $\rho_{1,b}$ to $\rho_b$.\footnote{Perhaps it is worth noting that $\overline{\rho(q)}$ is a complex valued function of the Fourier conjugate variable of $q$. Thus numerically the exponentiation in Eq. 9 requires Euler’s formula $e^{i\theta} = \cos \theta + i \sin \theta$.}

The nature of the FFT results in a number of round-off errors and limit the numerical precision to about $10^{-16}$ which limit the method to significance levels below about 8\,$\sigma$. Extrapolation techniques and arbitrary precision calculations can overcome these difficulties,\footnote{Some systematic errors provided by Sinervo.} but such small $p$-values are of little practical interest.

From the log-likelihood distribution of the two hypotheses we can calculate a number of useful quantities. Given some experiment with an observed log-likelihood ratio, $q^*$, we can calculate the background-only confidence level, $CL_b$:

$$CL_b(q^*) = \int_{q^*}^{\infty} \rho_b(q')\,dq'$$

(11)

In the absence of an observation we can calculate the expected $CL_b$ given the signal-plus-background hypothesis is true. To do this we first must find the median of the signal-plus-background distribution $\overline{q_{s+b}}$. From these we can calculate the expected $CL_b$ by using Eq. 11 evaluated at $q' = \overline{q_{s+b}}$.

Finally, we can convert the expected background confidence level into an expected Gaussian significance, $Z\sigma$, by finding the value of $Z$ which satisfies

$$CL_b(\overline{q_{s+b}}) = \frac{1 - \text{erf}(Z/\sqrt{2})}{2}. \hspace{1cm} (12)$$

where $\text{erf}(Z) = (2/\pi)^{1/2} \int_{0}^{Z} \exp(-y^2)\,dy$ is a function readily available in most numerical libraries. For $Z > 1.5$, the relationship can be approximated\footnote{Also of interest is the classification of systematic errors provided by Sinervo.} as

$$Z \approx \sqrt{u - \ln u} \text{ with } u = -2\ln(CL_b/\sqrt{2\pi}) \hspace{1cm} (13)$$

2.5 Systematic Errors, Nuisance Parameters & Auxiliary Measurements

Sections 2.3 and 2.4 represent the state of the art for HEP in frequentist hypothesis testing in the absence of uncertainties on rates and shapes of distributions. In practice, the true rate of background is not known exactly, and the shapes of distributions are sensitive to experimental quantities, such as calibration coefficients and particle identification efficiencies (which are also not known exactly). What one would call a systematic error in HEP, usually corresponds to what a statistician would refer to as a nuisance parameter.

Dealing with nuisance parameters in searches is not a new problem, but perhaps it has never been as essential as it is for the LHC. In these proceedings, Cousins reviews the different approaches to nuisance parameters in HEP and the professional statistical literature.\footnote{Also of interest is the classification of systematic errors provided by Sinervo.} Also of interest is the classification of systematic errors provided by Sinervo.\footnote{Also of interest is the classification of systematic errors provided by Sinervo.} In Sec. 4, the a few techniques for incorporating nuisance parameters are reviewed.

From an experimental point of view, the missing ingredient is some set of auxiliary measurements that will constrain the value of the nuisance parameters. The most common example would be a sideband measurement to fix the background rate, or some control sample used to assess particle identification efficiency. Previously, I used the variable $M$ to denote this auxiliary measurement\footnote{Also of interest is the classification of systematic errors provided by Sinervo.}, while Linnemann,\footnote{Also of interest is the classification of systematic errors provided by Sinervo.} Cousins,\footnote{Also of interest is the classification of systematic errors provided by Sinervo.} and Rolke, Lopez, and Conrad used $y$. Additionally, one needs to know
the likelihood function that provides the connection between the nuisance parameter(s) and the auxiliary measurements.

The most common choices for the likelihood of the auxiliary measurement are \( L(y|b) = \text{Pois}(y|\tau b) \) and \( L(y|b) = G(y|\tau b, \sigma_y) \), where \( \tau \) is a constant that specifies the ratio of the number of events one expects in the sideband region to the number expected in the signal-like region.\(^d\)

A constant \( \tau \) is appropriate when one simply counts the number of events \( y \) in an “off-source” measurement. In a more typical case, one uses the distribution of some other variable, call it \( m \), to estimate the number of background events inside a range of \( m \) (see Fig. 3). In special cases the ratio \( \tau \) is independent of the model parameters. However, in many cases (e.g. \( f(m) \propto e^{-am} \)), the ratio \( \tau \) depends on the model parameters. Moreover, sometimes the sideband is contaminated with signal events, thus the background and signal estimates can be correlated. These complications are not a problem as long as they are incorporated into the likelihood.

The number of nuisance parameters and auxiliary measurements can grow quite large. For instance, the standard practice at BaBar is to form very large likelihood functions that incorporate everything from the parameters of the unitarity triangle to branching fractions and detector response. These likelihoods are typically factorized into multiple pieces, which are studied independently at first and later combined to assess correlations. The factorization of the likelihood and the number of nuisance parameters included impact the difficulty of implementing the various scenarios considered below.

3 Practical and Toy Examples

In this Section, a few practical and toy examples are introduced. The toy examples are meant to provide simple scenarios where results for different methods can be easily obtained in order to expedite their comparison. The practical examples are meant to exclude methods that provide nice solutions to the toy examples, but do not generalize to the realistic situation.

3.1 The Canonical Example

Consider a number-counting experiment that measures \( x \) events in the signal-like region and \( y \) events in some sideband. For a given background rate \( b \) in the signal-like region, say one can expect \( \tau b \) events in the sideband. Additionally, let the rate of signal events in the signal-like regions – the parameter of interest – be denoted \( \mu \). The corresponding likelihood function is

\[
L_P(x, y|\mu, b) = \text{Pois}(x|\mu + b) \cdot \text{Pois}(y|\tau b). \tag{14}
\]

This is the same case that was considered in Refs. \(^{20,22,23,24}\) for \( x, y = O(10) \) and \( \alpha = 5\% \). For LHC searches, we will be more interested in \( x, y = O(100) \) and \( \alpha = 2.85 \cdot 10^{-7} \). Furthermore, the auxiliary measurement will rarely be a pure number counting sideband measurement, but instead the result of some fit. So let us also consider the likelihood function

\[
L_G(x, y|\mu, b) = \text{Pois}(x|\mu + b) \cdot G(y|\tau b, \sqrt{\tau b}). \tag{15}
\]

As a concrete example in the remaining sections, let us consider the case \( b = 100 \) and \( \tau = 1 \). Operationally, one would measure \( y \) and then find the value \( x_{\text{crit}}(y) \) necessary for discovery. In the language of confidence intervals, \( x_{\text{crit}}(y) \) is the value of \( x \) necessary for the \( 100(1 - \alpha)\% \) confidence interval in \( \mu \) to exclude \( \mu_0 = 0 \). In Sec. 4 we check the coverage (Type I error or false-discovery rate) for both \( L_P \) and \( L_G \).

Linnemann reviewed thirteen methods and eleven published examples of this scenario.\(^{19}\) Of the

\(^d\)Note that Linnemann used \( \alpha = 1/\tau \) instead, but in this paper \( \alpha \) is reserved for the rate of Type I error.
published examples, only three (the one from his reference 18 and the two from 19) are near the range of \(x, y\), and \(\alpha\) relevant for LHC searches. Linnemann’s review asks an equivalent question posed in this paper, but in a different way: what is the significance \((Z\) in Eq. 12) of a given observation \(x, y\).

### 3.2 The LHC Standard Model Higgs Search

The search for the standard model Higgs boson is by no means the only interesting search to be performed at the LHC, but it is one of the most studied and offers a particularly challenging set of channels to combine with a single method. Figure 1 shows the expected significance versus the Higgs mass, \(m_H\), for several channels individually and in combination for the ATLAS experiment.\(^{25}\) Two mass points are considered in more detail in Tab. 1, including results from Refs.\(^1\),\(^{25}\),\(^{26}\). Some of these channels will most likely use a discriminating variable distribution, \(f(m)\), to improve the sensitivity as described in Sec. 2.3. I have indicated the channels that I suspect will use this technique. Rough estimates on the uncertainty in the background rate have also been tabulated, without regard to the classification proposed by Sinervo.

The background uncertainties for the \(t\bar{t}H\) channel have been studied in some detail and separated into various sources.\(^{26}\) Figure 4 shows the \(m_{bb}\) mass spectrum for this channel.\(^5\) Clearly, the shape of the background-only distribution is quite similar to the shape of the signal-plus-background distribution. Furthermore, theoretical uncertainties and \(b\)-tagging uncertainties affect the shape of the background-only spectrum. In this case the incorporation of systematic error on the background rate most likely precludes the expected significance of this channel from ever reaching 5\(\sigma\).

Similarly, the \(H \rightarrow \gamma\gamma\) channel has uncertainty in the shape of the \(m_{\gamma\gamma}\) spectrum from background processes. One contribution to this uncertainty comes from the electromagnetic energy scale of the calorimeter (an experimental nuisance parameter), while another contribution comes from the theoretical uncertainty in the continuum \(\gamma\gamma\) production. Figure 5 shows two plausible shapes for the \(m_{\gamma\gamma}\) spectrum from “Born” and “Box” predictions.

### 4 Review of Methods

Based on the practical example of the standard model Higgs search at the LHC and the discussion in Sec. 2, the list of admissible methods is quite short. Of the thirteen methods reviewed by Linnemann, only five are considered as reasonable or recommended. These can be divided into three classes: hybrid Bayesian-frequentist methods, methods based on the Likelihood Principle, and frequentist methods based on the Neyman construction.

#### 4.1 Hybrid Bayesian-Frequentist Methods

The class of methods frequently used in HEP and commonly referred to as the Cousins-Highland technique (or secondarily Bayes in statistical literature)
are based on a Bayesian average of frequentist $p$-values as found in the first equation of Ref.\textsuperscript{28}. The Bayesian average is over the nuisance parameters and weighted by the posterior $P(b|y)$. Thus the $p$-value of the observation $(x_0,y_0)$ evaluated at $\mu$ is given by

$$
p(x_0,y_0|\mu) = \int_0^\infty db p(x_0|\mu,b) P(b|y_0)
$$

(16)

where

$$
P(x|\mu,y_0) = \int_0^\infty db P(x|\mu,b) \frac{P(y_0|b) P(b)}{P(y_0)}
$$

(17)

The form in Eq. 16, an average over $p$-values, is similar to the form written in Cousins \& Highland’s article; and it is re-written in Eq. 17 to the form that is more familiar to those from LEP Higgs searches.\textsuperscript{16,17} Actually, the dependence on $y_0$ and the Bayesian prior $P(b)$ shown explicitly in Eq. 18 is often not appreciated by those that use this method.

The specific methods that Linnemann considers correspond to different choices of Bayesian priors. The most common in HEP is to ignore the prior and use a truncated Gaussian for the posterior $P(b|y_0)$, which Linnemann calls $Z_N$. For the case in which the likelihood $L(y|b)$ is known to be Poisson, Linnemann prefers to use a flat prior, which gives rise to a Gamma-distributed posterior and Linnemann’s second preferred method $Z_{\Gamma}$, which is identical to the ratio of Poisson means $Z_{Bi}$ and can be written in terms of (in)complete beta functions as\textsuperscript{19}

$$
Z_{\Gamma} = Z_{Bi} = B(1/(1+\tau),x,y+1)/B(x,y+1).
$$

(19)

The method Linnemann calls $Z_{\tilde{\gamma}}$ can be seen as an approximation of $Z_N$ for large signals and is what ATLAS used to assess its physics potential.\textsuperscript{1} The method not recommended by Linnemann and was critically reviewed in Ref.\textsuperscript{29}.

$$
x_{\text{crit}}(y) = y/\tau + Z\sqrt{y/\tau(1+1/\tau)}
$$

(20)

4.2 Likelihood Intervals

As Cousins points out, the professional statistics literature seems less concerned with providing correct coverage by construction, in favor of likelihood-based and Bayesian methods. The likelihood principle states that given a measurement $x$ all inference about $\mu$ should be based on the likelihood function $L(x|\mu)$. When nuisance parameters are included, things get considerably more complicated.

The profile likelihood function is an attempt to eliminate the nuisance parameters from the likelihood function by replacing them with their conditional maximum likelihood estimates (denoted, for example, $\hat{b}$ ). The profile likelihood for $L_P$ in Eq. 14 is given by $L(x,y|\mu_0,\hat{b}(\mu_0))$, with

$$
\hat{b}(\mu_0) = \frac{x + y - (1 + \tau)\mu_0}{2(1 + \tau)} + \sqrt{(x + y - (1 + \tau)\mu_0)^2 + 4(1 + \tau)y\mu_0}/2(1 + \tau)
$$

(21)

The relevant likelihood ratio is then

$$
\lambda_P(\mu|x,y) = \frac{L(x,y|\mu_0,\hat{b}(\mu_0))}{L(x,y|\hat{\mu},\hat{b})},
$$

(22)

where $\hat{\mu}$ and $\hat{b}$ are the unconditional maximum likelihood estimates.

One of the standard results from statistics is that the distribution of $-2\ln \lambda$ converges to the $\chi^2$ distribution with $k$ degrees of freedom, where $k$ is the
number of parameters of interest. In our example $k = 1$, so a $5\sigma$ confidence interval is defined by the set of $\mu$ with $-2 \ln \lambda(\mu|x,y) < 25$. Figure 6 shows the graph of $-2 \ln \lambda(\mu|x,y)$ for $y = 100$ at the critical value of $x$ for a $5\sigma$ discovery.

At PhyStat2003, Nancy Reid presented various adjustments and improvements to the profile likelihood which speed asymptotic convergence properties. Cousins considers these methods in more detail from a physicist perspective.

Only recently was it generally appreciated that the method of Minuit commonly used in HEP corresponds to the profile likelihood intervals. The coverage of these methods is not guaranteed, but has been studied in simple cases. These likelihood-based techniques are quite promising for searches at the LHC, but their coverage properties must be assessed in the more complicated context of the LHC with weighted events and several channels. In particular, the distribution of $q$ in Eq. 10 is often highly non-Gaussian.

4.3 The Neyman Construction with Systematics

Linnemann’s preferred method, $Z_{Bi}$, is related to the familiar result on the ratio of Poisson means. Unfortunately, the form of $Z_{Bi}$ is tightly coupled to the form of Eq. 14, and can not be directly applied to the more complicated cases described above. However, the standard result on the ratio of Poisson means and Cousins’ improvement are actually special cases of the Neyman construction with nuisance parameters (with and without conditioning, respectively).

Of course, the Neyman construction does generalize to the more complicated cases discussed above. Two particular types of constructions have been presented, both of which are related to the profile likelihood ratio discussed in Kendall’s chapter on likelihood ratio tests & test efficiency. This relationship often leads to confusion with the profile likelihood intervals discussed in Sec. 4.2.

The first method is a full Neyman construction over both the parameters of interest and the nuisance parameters, using the profile likelihood ratio as an ordering rule. Using this method, the nuisance parameter is “projected out”, leaving only an interval in the parameters of interest. I presented this method at PhyStat2003 in the context of hypothesis testing, and similar work was presented by Punzi at this conference. This method provides coverage by construction, independent of the ordering rule used.

The motivation for using the profile likelihood ratio as a test statistic is twofold. First, it is inspired by the Neyman-Pearson lemma in the same way as the Feldman-Cousins ordering rule. Secondly, it is independent of the nuisance parameters; providing some hope of obtaining similar tests. Both Punzi and myself found a need to perform some “clipping” to the acceptance regions to protect from irrelevant values of the nuisance parameters spoiling the projection. For this technique to be broadly applicable, some generalization of this clipping procedure is needed and the scalability with the number of parameters must be addressed.

The second method, presented by Feldman at the Fermilab conference in 2000, involves a Neyman construction over the parameters of interest, but the nuisance parameters are fixed to the conditional maximum likelihood estimate: a method I will call the profile construction. The profile construction is an approximation of the full construction, that does...
not necessarily cover. To the extent that the use of the profile likelihood ratio as a test statistic provides similar tests, the profile construction has good coverage properties. The main motivation for the profile construction is that it scales well with the number of nuisance parameters and that the “clipping” is built in (only one value of the nuisance parameters is considered).

It appears that the CHOOZ experiment actually performed both the full construction (called “FC correct syst.”) and the profile construction (called “FC profile”) in order to compare with the strong confidence technique.36

Another perceived problem with the full construction is that bad over-coverage can result from the projection onto the parameters of interest. It should be made very clear that the coverage probability is a function of both the parameters of interest and the nuisance parameters. If the data are consistent with the null hypothesis for any value of the nuisance parameters, then one should probably not reject it. This argument is stronger for nuisance parameters directly related to the background hypothesis, and less strong for those that account for instrumentation effects. In fact, there is a family of methods that lie between the full construction and the profile construction. Perhaps we should pursue a hybrid approach in which the construction is formed for those parameters directly linked to the background hypothesis, the additional nuisance parameters take on their profile values, and the final interval is projected onto the parameters of interest.

5 Results with the Canonical Example

Consider the case \( b_{\text{true}} = 100, \tau = 1 \) (i.e. 10% systematic uncertainty). For each of the methods we find the critical boundary, \( x_{\text{crit}}(y) \), which is necessary to reject the null hypothesis \( \mu_0 = 0 \) at 5\( \sigma \) when \( y \) is measured in the auxiliary measurement. Figure 7 shows the contours of \( L_G \), from Eq. 15, and the critical boundary for several methods. The far left curve shows the simple \( s/\sqrt{b} \) curve neglecting systematics. The far right curve shows a critical region with the correct coverage. With the exception of the profile likelihood, \( \lambda_P \), all of the other methods lie between these two curves (i.e. all of them under-cover). The rate of Type I error for these methods was evaluated for \( L_G \) and \( L_P \) and presented in Table 2.

![](image.png)

Figure 7. A comparison of the various methods critical boundary \( x_{\text{crit}}(y) \) (see text). The concentric ovals represent contours of \( L_G \) from Eq. 15.

The result of the full Neyman construction and the profile construction are not presented. The full Neyman construction covers by construction, and it was previously demonstrated for a similar case (\( b = 100, \tau = 4 \)) that the profile construction gives similar results.22 Furthermore, if the \( \lambda_P \) were used as an ordering rule in the full construction, the critical region for \( b = 100 \) would be identical to the curve labeled “\( \lambda_P \) profile” (since \( \lambda_P \) actually covers).

It should be noted that if one knows the likelihood is given by \( L_G(x, y|\mu, b) \), then one should use the corresponding profile likelihood ratio, \( \lambda_G(x, y|\mu) \), for the hypothesis test. However, knowledge of the correct likelihood is not always available (Sinervo’s Class II systematic), so it is informative to check the coverage of tests based on both \( \lambda_G(x, y|\mu) \) and \( \lambda_P(x, y|\mu) \) by generating Monte Carlo according to \( L_G(x, y|\mu, b) \) and \( L_P(x, y|\mu, b) \). In a similar way, this decoupling of true likelihood and the assumed likelihood (used to find the critical region) can break the “guaranteed” coverage of the Neyman construction.

It is quite significant that the \( Z_N \) method under-covers, since it is so commonly used in HEP. The degree to which the method under-covers depends on the truncation of the Gaussian posterior \( P(b|y) \). Linenmann’s table also shows significant under-coverage (over estimate of the significance \( Z \)). In order to obtain a critical region with the correct coverage, the
author modified the region \( x_{\text{crit}}(y) = x_{\text{crit}}^{Z_N}(y) + C \) and found \( C = 16 \) provided the correct coverage. A discrepancy of 16 events is not trivial.

Table 2. Rate of Type I error interpreted as equivalent \( Z \sigma \) for various methods designed for a \( 5 \sigma \) test. Monte Carlo events are generated via either \( L_G \) or \( L_P \). The critical \( x \) for \( y = 100 \) is also listed for easy comparison.

| Method   | \( L_G \) (\( Z \sigma \)) | \( L_P \) (\( Z \sigma \)) | \( x_{\text{crit}}(y = 100) \) |
|----------|-----------------|-----------------|-----------------|
| No Syst  | 3.0             | 3.1             | 150             |
| \( Z_{\text{bi}} \) | 4.1             | 4.1             | 171             |
| \( Z_N \) (Sec. 4.1) | 4.2             | 4.2             | 178             |
| ad hoc   | 4.6             | 4.7             | 188             |
| \( Z_{\text{B}} \) \( \equiv \) \( Z_{\text{Bi}} \) | 4.9             | 5.0             | 185             |
| profile \( \lambda_P \) | 5.0             | 5.0             | 185             |
| profile \( \lambda_G \) | 4.7             | 4.7             | \( \sim 182 \) |

Notice that for large \( x, y \) the Bayesian-frequentist hybrid \( Z_N \) approaches \( Z_{\text{bi}} \), where the critical region is of the form \( x_{\text{crit}}(y) = y/\tau + n\sqrt{y}/\tau \). Because the boundary is very nearly linear around \( y_0 \), one can find the value of \( n \) that gives the proper coverage with a little geometry. In particular, the number \( n \) needed to get a \( Z \sigma \) test gives

\[
x_{\text{crit}}(y) = y/\tau + Z \sqrt{1 + 1/\tau m^2} \sqrt{y}/\tau	ag{23}
\]

where

\[
m = \left(1 + \frac{Z}{2\sqrt{y/\tau}}\right)^{-1}	ag{24}
\]

The \( m^2 \) factor can be seen as a correction to the \( Z_{\text{bi}} \) and \( Z_N \) results. Notice that the correction is larger for higher significance tests. As an ad hoc method, I experimented with the \( Z_N \) method replacing \( \tau \) with \( \tau m^2 \) in the posterior \( P(b|y) \). The coverage of this ad hoc method is better than \( Z_N \), but not exact because \( x, y \) are not sufficiently large.

6 Conclusions

I have presented the statistical challenges of searches at the LHC and the current state of the statistical methods commonly used in HEP. I have attempted to accurately portray the complexity of the searches, explain their key ingredients, and provide a practical example for future studies. Three classes of methods, which are able to incorporate all the ingredients, have been identified: hybrid Bayesian-frequentist methods, methods based on the Likelihood Principle, and frequentist methods based on the Neyman construction.

The Bayesian-frequentist hybrid method, \( Z_N \), shows significant under-coverage in the toy example considered when pushed to the \( 5 \sigma \) regime. While Bayesian might not care about coverage, significant under-coverage is undesirable in HEP. Further study is needed to determine if a more careful choice of prior distributions can remedy this situation—especially in more complex situations. The improved coverage of \( Z_T \) may give some guidance.

The methods based on the likelihood principle have gained a great deal of attention from HEP in recent years. While the methods appear to do well in the toy example, it requires further study to determine their properties in the more realistic situation with weighted events.

Slowly, the HEP community is coming to grips with how to incorporate nuisance parameters into the Neyman construction. Several ideas for reducing the over-coverage induced by projecting out the nuisance parameters and reducing the computational burden have been presented. A hybrid approach between the full construction and the profile construction should be investigated in more detail.

Finally, it seems that the HEP community is approaching a point where we appreciate the fundamental statistical issues, the limitations of some methods, and the benefits of others. Clearly, the philosophical debate has not ended, but there seems to be more emphasis on practical solutions to our very challenging problems.

Acknowledgments

I would like to thank the many people that helped in preparing this review. In particular, Bob Cousins, Jim Linnemann, Gary Feldman, Jan Conrad, Fredrik Tegenfeldt, Wolfgang Rolke, Nancy Reid, Gary Hill, and Stathes Paganis. I would also like to thank Louis Lyons for his continuing advice and the invitation to speak at such an enjoyable and productive conference.

This manuscript has been authored by Brookhaven Science Associates under Contract No. DE-AC02-98CH1-886 with the U.S. DOE. The U.S. Government retains, and the publisher, by accepting the article for publication, acknowledges, a world-wide license to publish or reproduce the published form of this manuscript, or allow others to do so, for the U.S. Government purposes.
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