# Approximating decomposed likelihood ratios using machine learning

# K Cranmer<sup>1</sup>, J Pavez<sup>2</sup> and G Louppe<sup>1</sup>

- <sup>1</sup> Physics Department, New York University, New York, NY 10003, U.S.A.
  - <sup>2</sup> Informatics Department, Universidad Técnica Federico Santa María, 1240 Av. España, Valparaíso, Chile

E-mail: juan.pavezs@alumnos.usm.cl

Abstract. Likelihood ratio test are a key tool in many fields of science. In order to evaluate the likelihood ratio the likelihood function is needed. However, it is common in fields like high energy physics to have complex simulations that describe the distribution while not having a description of the likelihood that can be directly evaluated. In this setting it is impossible or computationally expensive to evaluate the likelihood. It is possible to construct an equivalent version of the likelihood ratio that can be easily evaluated by using discriminative classifiers. We show how this can be used to approximate the likelihood ratio when the underlying distribution is a weighted sum of probability distributions (e.g. signal plus background model). We demonstrate how the results can be considerably improved by decomposing the test and use a set of classifiers in a pairwise manner on the components of the mixture model and in which way this can be used to estimate the unknown coefficients of the model, such as the signal contribution.

### 1. Introduction

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In High Energy Physics (HEP) and many other fields hypothesis testing is a key tool when reporting results from an experiment. Likelihood ratio test are the main technique for hypothesis testing and they are the most powerful statistic for simple hypothesis testing. For composite hypothesis testing the profile or generalized likelihood ratio test is commonly used. When computing the likelihood ratio the data distribution  $p(x|\theta)$  must be evaluated, where  $\theta$  are parameters of the probability distribution. However, it is common in HEP to have physics simulations that allows to sample high dimensional vectors from the distribution  $p(x|\theta)$  while not having a description that can be directly evaluated. Commonly it is impossible or computationally expensive to evaluate the likelihood ratio in this setting.

It has been recently proved that discriminative classifiers can be used to solve this problem by constructing an equivalent version of the likelihood ratio in this likelihood-free setting [1].

In this work we show how these results can be used to approximate the likelihood ratio when the underlying distribution is a weighted sum of probability distributions (mixture model). This case is common in physics when identifying a signal process over one or more background process. We also show that by training a set of classifiers in a pairwise manner on the components of the mixture model it is possible to considerably improve the results of the approximation.

## 2. Approximating likelihood ratios using discriminative classifiers

A common use of likelihood ratios in HEP is signal process identification. In this task the hypothesis testing procedure is used to evaluate the signal process significance by contrasting the only background (null) hypothesis versus the signal plus background (alternative) hypothesis. In this setting, the underlying distribution can be seen as a signal and background mixture model defined as

$$p(x \mid \mu, \nu) = \mu p_s(x \mid \nu) + (1 - \mu) p_b(x \mid \nu), \tag{1}$$

where  $p_x(x \mid \nu)$  correspond to the signal distribution and  $p_b(x \mid \nu)$  is the background distribution, both parametrized by nuisance parameters  $\nu$  which describe uncertainties in underlying physics predictions or response of measurement devices. The parameter  $\mu$  is the mixture coefficient corresponding to the signal component of the distribution. In this case the generalized likelihood ratio test takes the form of

$$T(D) = \prod_{e=1}^{n} \frac{p(x_e | \mu = 0, \hat{\nu})}{p(x_e | \hat{\mu}, \hat{\nu})},$$
(2)

where D is a data set of i.i.d observations  $x_e$ ,  $\hat{\nu}$  is the conditional maximum likelihood estimator for  $\nu$  under the null hypothesis  $\theta_0$  ( $\mu = 0$ ) and  $\hat{\nu}$ ,  $\hat{\mu}$  are the maximum likelihood estimators for  $\nu$  and  $\mu$ . This approach has been used extensively to assert the discovery of new particles in HEP [2], such as in the discovery of the Higgs boson [3, 4].

As previously mentioned, the original distributions for signal and background can only be approximated by forward simulations. Most of the likelihood ratio tests at the LHC are made on the distribution of a single feature that discriminate between signal and background observations. For this, the simulated data is used together with interpolations algorithms in order to approximate the parametrized model and then use it in the hypothesis testing procedure [5].

Noticeably, it has been shown that a discriminative classifier trained to classify between signal and background can be used to obtain an equivalent likelihood ratio test [1]. Let  $s(x; \theta_0, \theta_1)$  to represent the classification score learned by the classifier, parametrized by  $\theta_0$  and  $\theta_1$  the parameters of the statistical model, let  $p(s(x; \theta_0, \theta_1)|\theta)$  the probability distribution of the score on the data from the original distribution  $p(x|\theta)$ . Then, the likelihood ratio test is equivalent to the test on the conditional distributions of the score,

$$T(D) = \prod_{e=1}^{n} \frac{p(x_e|\theta_0)}{p(x_e|\theta_1)} = \prod_{e=1}^{n} \frac{p(s(x_e;\theta_0,\theta_1)|\theta_0)}{p(s(x_e;\theta_0,\theta_1)|\theta_1)},$$
(3)

where  $\theta_0$  correspond to the null hypothesis and  $\theta_1$  to the alternative hypothesis. The only requirement is that the discriminative classifier learns a monotonic function of the per event ratio  $p(x_e|\theta_0)/p(x_e|\theta_1)$  [1]. This is the usual case since many of the commonly used classifiers learn to approximate some monotonic function of the regression function  $s(x) \sim p(y|x) = p(x|\theta_1)/(p(x|\theta_0) + p(x|\theta_1))$  which is monotonic to the desired ratio.

# 3. Decomposed likelihood ratio test for mixture models

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A generalized version of the signal and background mixture model of eq. (1) for several components is

$$p(x|\theta) = \sum_{i=1}^{k} w_i(\theta) p_i(x|\theta), \tag{4}$$

where  $w_i(\theta)$  are the mixture coefficients for each one of the components parametrized by  $\theta$ . In [1] it is shown that the likelihood ratio between two mixture models

$$\frac{p(x|\theta_0)}{p(x|\theta_1)} = \frac{\sum_{i=1}^k w_i(\theta_0) p_i(x|\theta_0)}{\sum_{i=1}^{k'} w_i(\theta_1) p_j(x|\theta_1)},\tag{5}$$

is equal to the composition of pairwise ratios for each one of the components which by eq. (3) is equivalent to the composition of ratios on the score distribution of pairwise trained classifiers

$$\frac{p(x|\theta_0)}{p(x|\theta_1)} = \sum_{i=1}^k \left[ \sum_{j=1}^{k'} \frac{w_j(\theta_1)}{w_i(\theta_0)} \frac{p_j(x|\theta_1)}{p_i(x|\theta_0)} \right]^{-1} = \sum_{i=1}^k \left[ \sum_{j=1}^{k'} \frac{w_j(\theta_1)}{w_i(\theta_0)} \frac{p_j(s_{i,j}(x;\theta_0,\theta_1)|\theta_1)}{p_i(s_{i,j}(x;\theta_0,\theta_1)|\theta_0)} \right]^{-1}.$$
 (6)

In the case that the only free parameters of the mixture model are the coefficients  $w_i(\theta)$ , then each distribution  $p_i(s_{i,j}(x;\theta_0,\theta_1)|\theta)$  is independent of  $\theta$  and can be pre-computed and used after in the evaluation of the likelihood ratio. Moreover, in this case each ratio  $p_j(s_{i,j}(x;\theta_0,\theta_1)|\theta_1)/p_i(s_{i,j}(x;\theta_0,\theta_1)|\theta_0)$  with i=j can be replaced by 1. Also, for a two-class classifier the values of  $s_{j,i}(x;\theta_0,\theta_1)$  for one of the classes can be replaced by the values of  $s_{i,j}(x;\theta_0,\theta_1)$  for the opposing class, then it is only necessary to train the classifiers for i< j. This saves a lot of computation time and avoid possible variance that can be introduced by differences between  $s_{i,j}(x;\theta_0,\theta_1)$  and  $s_{j,i}(x;\theta_0,\theta_1)$  due to imperfect training. It is common that in the case of only background hypothesis versus signal plus background hypothesis the signal coefficient  $w_j(\theta_1)$  is a very small number compared to the background coefficients under the alternate hypothesis. In this conditions a classifier trained in data from the full mixture model will have a lot of problems to identify the signal since most of the useful discriminative data will lay in a small region of the feature space while the decomposed model will not face this issues.

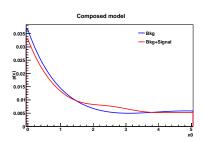
It is possible to estimate the signal and background coefficients by using maximum likelihood on the ratios. This can be done by keeping the denominator fixed and estimate the parameters on the numerator using the maximum likelihood method.

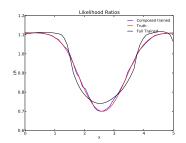
The complete algorithm to approximate the likelihood ratio using pairwise trained classifiers can be separated into three independent stages: classifier training, score distribution estimation and composition formula computation. Since this steps are independent, each one can be solved as a different problem. In the first step any classifier satisfying the monotonic requirement can be used. In the second stage the probability distribution of the score on data from  $\theta_0$  or  $\theta_1$  can be estimated using any univariate density estimation technique such as histograms or kernel density estimation [6, 7].

#### 4. Experiments

In this section two examples of how the method works on data generated from known distributions will be presented. In both cases we will study how the decomposition formula works using pairwise trained classifiers and we will compare the results to the true (in this case known) likelihood ratio and to the likelihood ratios obtained by training a classifier in data from the full mixture model and using directly eq. (3). All studies were conducted using a simple multilayer perceptron model. This classifier shows a good tradeoff between quality of the ratios and simplicity of the model (good results were also obtained using boosted decision trees, logistic regression and support vector machines). The probability models were implemented with RooFit probabilistic programming language and the classifiers were implemented using Theano(a framework to build neural network models) and scikit-learn(a general framework for machine learning in python) [6, 8, 9](the code is available for replication of the results at https://github.com/jgpavez/systematics).

First, we present a simple case in which each component is an univariate distribution. We consider a mixture model consisting of three distributions, where  $p_0(x)$  and  $p_1(x)$  are univariate gaussian distributions while  $p_2(x)$  is a decaying exponential. The mixture models are composed by the weighted sum of those distributions where  $p_0(x)$  correspond to the signal component. The mixture models with coefficients  $W(\theta_0) = \{0., 0.3, 0.7\}$  for the only background hypothesis and  $W(\theta_1) = \{0.1, 0.27, 0.63\}$  for the background plus signal hypothesis is shown in Figure 1.





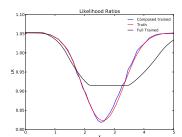


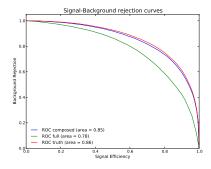
Figure 1. The mixture models.

**Figure 2.** Density ratio comparison for a signal weight of 0.1.

Figure 3. Density ratio comparison for a signal weight of 0.05.

Three neural networks were trained on data sampled from the pairs of single distributions and one on data from the full model. The distribution of the score is estimated using histograms (the number of bins were carefully chosen in order to allow a good approximation while minimizing poisson fluctuations). The composed ratio using eq. (6), the ratio estimated using a classifier trained on data from the full model and the true density ratios are shown in Figure 2 and Figure 3 for different values of the signal contribution (0.1 and 0.05) while keeping the ratio between the background contributions fixed.

It can be seen that the ratios obtained by the composition method are better (closer to the true ratios) than the obtained using models trained in data from the full mixture model and this become clearer when the signal contribution is smaller.



**Figure 4.** ROC curve comparison for a signal weight of 0.1.

**Figure 5.** ROC curve comparison for a signal weight of 0.05.

The same experiments are repeated but now considering a much harder mixture model consisting of three distributions, each one composed by the sum of three 10-dimensional multivariate gaussian distributions. This is more similar to what can be found in real physics experiments. To evaluate the estimated ratios the ROC curves on data from  $\theta_0$  and  $\theta_1$  are used with the density ratio used as discriminative variable. Again, the ROC curves for each one of the three cases (decomposed, full and true) is shown for a signal contribution of 0.1 and 0.05 in Figure 4 and Figure 5.

The values of each one of the coefficients of the mixture model can be estimated by using the method of maximum likelihood as explained in Section 3. In Figure 6 the contour plot for the likelihood ratio values given the signal and one background weight, obtained using the estimated density ratios and the true density ratios is shown. Histogram for the fitted values (estimated

and true) of each one of the coefficients on 80 different pseudo-datasets of size 1000 are shown in Figure 7 and Figure 8. It can be seen on the histograms that the estimations are unbiased.

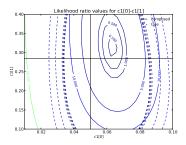


Figure 6. Likelihood ratio values given signal (c1[0]) and bkg. (c1[1]) weights.

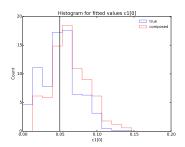
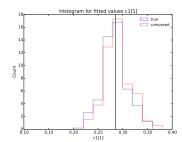


Figure 7. Histogram of fitted values for the signal weight (c1[0]).



**Figure 8.** Histogram of fitted values for the bkg. weight (c1[1]).

#### 5. Conclusions

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We have shown the power of using discriminative classifiers in order to approximate the likelihood ratio. In the case of mixture models we proved that the decomposed version of the ratio can greatly improve the quality of the results. Using the same method it was shown how to estimate the unknown parameters of the model. Initial experiments have been conducted on simulated data from different Higgs production mechanisms obtaining encouraging results. An open source Python package was implemented allowing to easily use the presented method (the package can be found at https://github.com/diana-hep/carl).

#### 6. Acknowledgments

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