# Estimating error rates for bullet comparisons in forensic science

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## Abstract

This is the abstract.

It consists of two paragraphs.

Text based on elsarticle sample manuscript, see http://www.elsevier.com/author-schemas/latex-instructions#elsarticle

## 1. Introduction

Firearm examination is an important topic in forensic science to help reveal the underlying pattern of firearm evidence. XXX go into a bit more detail on what you mean by the underlying pattern Currently, those examinations are executed in forensic labs by Firearms and Toolmark Examiners (FTEs) under the regulation of AFTE (AFTE Criteria For Identification Committee, 1992). However, this process is in its in nature based on human decisions and therefore subjective. XXX Change from passive voice to active, i.e. The PCAST report and the NRC called out this subjectivity and ... The subjectivity in the conventional firearm evidence identification processes is called to be reduced or to be complemented by more objective procedures by the National Research Council (National Research Council, 2009). The President's Council of Advisers on Science and Technology (PCAST) also emphasized the importance to establish the scientific validity and reliability of subjective forensic feature-comparison methods by blind empirical test (President's Council of Advisors on Science and Technology (PCAST), 2016). The PCAST report also recognized the importance of developing of objective computer-based algorithms in its following addendum (President's Council of Advisors on Science and Technology (PCAST), 2017).

In response to this criticism, subsequent research on cartridge case comparisons introduced the method of congruent matching cells (CMC) (Song, 2015), which allows a quantitative assessment of the similarity of two breech face impressions. Error rates were reported by [@Song2018; @Zhang2019] on XXX which studies were used in these papers? based on distributional models XXX

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could you expand on those distributional models?. XXX xiao-hue Tai's paper should be mentioned. XXX mention the data that all of these methods are based on.

## XXX i'm not sure you need to go into bullet comparisons

In bullet comparisons, an automatic matching algorithm was proposed by Hare (Hare et al., 2016). The discussion about the degraded land comparison was also made in the following paper (Hare et al., 2017). The empirical error rates associated with this method based on three test sets are discussed by Vanderplas (Vanderplas et al., 2020). The most prominent single feature in the random forest proposed by Hare (Hare et al., 2016) is the cross-correlation function (Vorburger et al., 2011) according to the empirical studies (Hare et al., 2016; Vanderplas et al., 2020). Other automated comparison methods or improvement such as Robust LOESS in groove engraved area identification (Rice et al., 2020), Chumbley score method (Krishnan and Hofmann, 2019) are also proposed.

## XXX What does this however refer to?

However, due to the limitation of sample sizes and designs of study, a sound distributional assumption about the resulting similarity scores was not well established in the bullet comparison field for automated matching algorithms. The error rates are estimated empirically from dozens of bullets which were originally designed for examiner proficiency tests. And many of those tests are also criticized for the closed-set design and potential information provided to the examiners as pointed out by PCAST (President's Council of Advisors on Science and Technology (PCAST), 2017, 2016). It's important to understand the underlying distributions of the similarity scores to estimate the theoretical error rates of any automatic algorithm. And that also helps later to choose proper thresholds for the methods instead of simply choosing 0.5 (in a 0-1 range of similarity scores) as a belief.

A sketch of a bullet and a barrel would be good here - showing rifling (grooves and lands) and striation marks. Introduce rifling, Describe LEA and GEA, and structure of LEAs

A fired bullet has several LEAs according to the barrels' design.

Generally, in forensic science, we consider the problem of distinguishing two pieces of evidence coming from a same source or different sources. Specifically, in bullet comparisons, we want to know if two bullets were fired from the same gun. When shooting a bullet, micro-imperfections inside the barrel leave markings on the surface of a bullet in form of striations. Striations on land engraved areas (LEAs) are used by firearms and toolmarks examiners in comparisons to determine same source of two pieces of evidence (AFTE Criteria For Identification Committee, 1992).

include paragraph on PCAST and NRC reports: lack of objectivity and error rates

Automatic matching algorithms were introduced to address the criticism by PCAST and NRC.

The next paragraph needs to be worked out in more detail. You are getting to densities too fast. First define scores as outcomes of an algorithm, then describe

same source density and different source density and (score based) likelihood ratios

And the LEA is also used in the developed computer-based methods. So, we are not doing one step to identify bullets, instead we consider each LEA and then combine the land level conclusions to a bullet level one. Accordingly, we start our discussion about underlying distributions and error rates from the land level.

There are two types of distributions we would consider in the bullet comparison cases. One is called known-match (KM) distributions for those actually matched lands. The other is called known-non-match distributions (KNM) for those actually not matched lands. Those two distributions are only available in experimental settings where we know the ground truth. They are the starting point for any discussion about theoretical error rates. And particularly, we consider the cross-correlation function (CCF) for bullet signatures as a typical similarity score with range 0-1 used to distinguish between same source and different source comparisons. The cross-correlation function (CCF) is calculated after extracting the signatures of bullet lands by maximizing the cross correlation of two sequence of signatures by horizontally adjusting the signatures (Vorburger et al., 2011).

In section 2... In section 3...

#### 2. The distributional forms of similarity scores

The quantitative methods used to objectively measure the similarity between LEAs report various quantities, such as counts, correlations, distances, probabilities and more general similarity scores (Hare et al., 2016; Ma et al., 2004; Nichols, 2003, 1997). To understand how those quantities reflect the strength of evidence and to study the underlying error rates in making decisions based on those quantities, distributional forms are usually set up (Song et al., 2018; Zhang, 2019). Particularly, we are focusing on the similarity scores which range from 0 to 1 and the probabilities reported as the likelihood of an actual match. The similarity scores reported in the forensic researches are classified into two categories as known matches (KM) and known non-matches (KNM), and the corresponding distributions are named as KM distributions and KNM distributions. When we make any decisions based on any quantitative measurement, we are actually making a distinguish between those two potential distributions. The strength of any identification process is also measured by the disparities of those distributions. However, in practice, we can hardly discriminate those two distributions entirely, thus, we are never 100% sure which distribution the observed score comes from. This is where the identification error raises.

The cross-correlation function (CCF) is the most prominent single feature in the automatic random forest algorithm (Hare et al., 2016) which has theoretical range from -1 to 1. But in real applications, this similarity score always has values in 0 to 1, e.g. in our case, none of 121992 comparisons has negative values. So, it is selected as a representative of similarity scores with range from 0 to 1 in this paper for further analysis. The similarity scores in 0 to 1 can be explained

as probabilities that quantify the likelihood that a pair of LEAs are actually a match. Or we can think of them as general similarity measurement. As the name indicated, the higher the similarity score is, the stronger evidence is to support the same source assumption. For different combination of ammunition and firearms, the scores are distributed differently. It is expected that systematic differences exist there for different cases (Vanderplas et al., 2020). So, it is necessary to study the scores under controlled conditions.

We can see from the Figure ... (a figure of nonparametric fit of the CCF?), which is a typical one we usually have for the similarity scores, that the distributions of KM and KNM are apart for the majorities. In the bullet LEA comparison problems, we usually have a well separated bullet scores but for the land scores, there are usually some overlaps (Vanderplas et al., 2020). We propose beta distributions for those scores. Because the beta distribution is a well-used distribution in statistics to describe a quantity from 0 to 1 which is usually a probability or proportion quantifying our knowledge for another distribution in Bayesian analysis. And it is very flexible to capture unimodal asymmetric shapes in 0 to 1. However, it may not be adequate to explain a heavy tail or even a second mode. Thus, we further consider the beta mixture distribution which is a more complex distribution than the beta distribution as a special case. In the beta mixture distribution, we introduce a hierarchical structure with a prior probability to combine a few beta distributions as one. The two-component beta mixture distributions are defined below.

The reported similarity scores are denoted as  $Y_{ij}$  for  $j^{th}$  LEA comparison within class i, where i = 1 is KM and i = 2 is KNM.  $Y_{ij}$ 's are considered independent and identically distributed within each class, i.e.

$$Y_{1j} \stackrel{iid}{\sim} Betamix(p_1, \mu_{11}, \phi_{11}, \mu_{12}, \phi_{12})$$
  
 $Y_{2j} \stackrel{iid}{\sim} Betamix(p_2, \mu_{21}, \phi_{21}, \mu_{22}, \phi_{22})$ 

where  $\mu_{ik}$  and  $\phi_{ik}$  are distribution parameters for  $i^{th}$  class and  $k^{th}$  component, k=1 or 2. And  $p_i$  is the prior probability for the first component in  $i^{th}$  class, thus,  $1-p_i$  is the prior probability for the second component in  $i^{th}$  class. Note that we are using the mean and precision parameterization of beta distributions which simplifies the math in calculation and is more intuitive ( $\mu$  is the mean, and the variance is roughly proportional to the reciprocal of  $\phi$ ). It's equivalent to the usual  $\alpha$  and  $\beta$  parameterization through the following transformation:

$$\mu = \frac{\alpha}{\alpha + \beta}$$
$$\phi = \alpha + \beta$$

## 3. LAPD data set and the estimated distributions

For the following sections of the paper, we will base our analysis on the LAPD data sets. This is a large data set of ... It is the first time such a large

data set available to the researchers, which makes it possible for a statistical analysis for the distribution of similarity scores. (More on the data. Is there any formal documentation for LAPD? And number of land comparisons for different cases should be discussed here)

We consider cross correlation functions (CCF) for land comparisons, which is produced by calculating the maximized CCF between two signatures extracted from a pair of bullet LEAs.

The estimation was done using Nelder-Mead algorithm in R [reference]. This is a general purpose numeric method which works reasonably well for multidimensional optimization problems. In our case, the objective function is the log likelihood function of the beta mixture distribution. Therefore, we finally found the maximum likelihood estimates (MLE). Instead of a simpler beta model, we start with the more complex two-component beta mixture model, see how well it fits the data and test if some components are necessary. We estimated beta mixture distributions for both KM and KNM as in Table 1 and Table 2 respectively. For the beta distributions, the KM distribution has a mean at 0.635 and the KNM distribution has a mean at 0.404. We can also see that for the two component beta mixture distributions for both KM and KNM, there are components with mean around 0.5 and the other components more separated from each other. This indicates the model successfully accounted for different situations of comparisons involving tank rash and random identification etc. This is an ideal property we would like to see and make use in explaining the similarity scores. Another point about the estimates that worth mentioning is that the  $\phi$  of the KM beta distribution is much smaller than any of the two components of the two-component KM beta mixture distribution. This indicates that the two component distribution could be better because of smaller variances for components.

Full Data KM Distribution Estimation										
Model	Component	$\mu$	$\phi$	logLik	p-value	BIC				
	Prior									
	Probability									
	0.247	0.406	24.097							
3-comp	0.521	0.664	13.342	6278		-12479				
	0.232	0.819	33.823							
2-comp	0.419	0.466	15.812	6243	0	-12438				
	0.581	0.759	18.476			-12430				
Beta		0.635	6.529	5712	0	-11404				

Table 1: Parameter estimations for the Beta distribution (1-component beta mixture distribution), 2-component and 3-component distributions for the KM CCF. "2-comp" refers to the 2-component beta mixture distribution, and the same for "3-comp". Column "logLik" is the maximized log likelihood for each distribution. Column "p-value" is the p-value for asymptotic likelihood ratio tests between the current model and one-step more complex model, where 0 indicates that we would reject the hypothesis that the current one is sufficient to decribe the data

Full Data KNM Distribution Estimation									
Model	Component	$\mu$	$\phi$	logLik	p-value	BIC			
	Prior								
	Probability								
	0.650	0.345	35.017						
3-comp	0.314	0.490	24.979	78961		-157830			
	0.032	0.667	18.454						
2-comp	0.674	0.358	39.908	78808	0	-157558			
	0.336	0.494	13.324			-101000			
Beta		0.404	15.801	75349	0	-150675			

Table 2: Parameter estimations for the Beta distribution (1-component beta mixture distribution), 2-component and 3-component distributions for the KNM CCF. "2-comp" refers to the 2-component beta mixture distribution, and the same for "3-comp". Column "logLik" is the maximized log likelihood for each distribution. Column "p-value" is the p-value for asymptotic likelihood ratio tests between the current model and one-step more complex model, where 0 indicates that we would reject the hypothesis that the current one is sufficient to decribe the data

The estimated beta distributions are shown in the Figure 1 and the estimated two-component beta mixture distributions are also shown in the Figure 2. As expected, the estimated distributions show the properties we desire for the similarity scores. The majorities of the estimated distributions are apart from each other, while the minority part between the two distributions has some overlap. It's worth to note that both curves have heavy tails to the farther boundaries and the KM curve has the heavier tail compared to KNM. We can see the two-component distributions fit the data very well, while the single beta distributions are not as good as the two-component distributions for both KM and KNM. The single beta distribution for KM clearly failed to capture the potential second mode of the histogram indicating it is not sufficient. The single beta distribution for KNM is not too bad but still failed to capture the distributional information for some parts. This indicates that even though beta distributions are flexible for variables in 0 to 1, they are still restricted too much to be able to describe the cases here. The two-component beta mixture distributions look more promising.

Three candidate distributions for each of KM and KNM are considered. For the increasing of complexity, we have single beta, two-component beta mixture and three-component beta mixture distributions. And they are also nested in that order. Naturally, we first look at the maximized log-likelihood of each model, and we can do asymptotic log-likelihood ratio chi-square tests for single beta against two-component beta mixture distributions, and two-component against three-component beta mixture distributions. The p-values of those test are shown in the Table 1 and Table 2 as the column "p-value". Surprisingly (or not), we found all those p-values are 0 which strongly suggests a more complex model when there is one. However, considering the size of the data we are using to fit these models, we can expect the statistical significance will be easily achieved

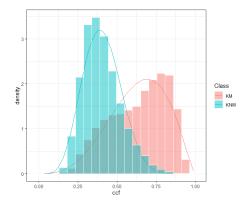


Figure 1: Estimated beta distributions for full data

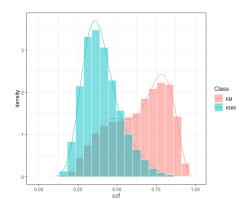


Figure 2: Estimated two-component beta mixture distributions for full data

since any small difference of the sufficiency will be detected. So we have to take the sample size effect into consideration. As Bayesian information criterion (BIC) is a well used criterion which takes the model complexity, sufficiency and the sample size (by a log function) into account. As shown in the column "BIC" in the tables, the BICs for the single beta distributions are larger than that of the two-component beta mixtures by relatively large proportions. And the BICs for the two-component beta mixture distributions are a little larger than that of the three-component beta mixture distributions for both KM and KNM. Obviously, we would prefer the two-component beta mixture distributions, but we would cast a doubt when it comes to the three-component beta mixture distributions. We still prefer the two-component one instead of the three-component one. The reasons are: 1) these differences of BICs between two and three component distributions are really small in proportion (by 0.3% for KM and 0.017% for KNM), 2) the estimation cost is not accounted, which will be higher for a more complex model, 3) the BICs still don't take the sample size effect fully into

account since the log function for sample size goes to flat when the sample sizes are large. By simple calculation, we can see that the log-likelihood increased by a multiplicative factor more than 10 when the sample sizes increased from KM case to KNM case, however, at the same time, the log of sample size only roughly increased from 9 to 11. So we choose the two-component beta mixture distributions for both KM and KNM. And also as we have seen, these distributions have potential good forensic interpretations.

It's also helpful to see how the individual components look like in the beta mixture distributions as in Figure 3. The KNM and KM distributions seem to share a common component while keeping the other components far apart from each other. The separated components represent the ideal cases where the bullet land engraved areas preserved the information of source well and result in clearly distinct separation. The shared components represent the cases where the KM results in lower scores because of some degree of tank rash, pitting, breakoff or other damages on the bullets and the KNM results in higher score because of the random identification effect. According to the estimated prior probabilities, both distributions put less weight on the common component while putting larger weight on the the components characterizing the differences of KM and KNM respectively, which agrees on our expectation that majorities of the distributions are separated while the minorities overlapped. These properties together well explained the observed empirical distribution of similarity scores in our cases. Particularly, the heavier tail of the KM comparisons is explicitly included in the form of the model by one of the components.

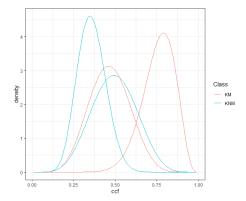


Figure 3: Estimated components

## 4. Determining source

In current practice, FTEs fire two or three test bullets from a suspected firearm using ammunition that matches the evidence found on the crime scene as closely as possible. Test fires are checked for consistency of striation marks

between fires. Generally, the bullet with the more well exopressed striations is used to compare to the evidence.

Here, we distinguish between two scenarios: (1) use scores from comparisons of test fires to estimate densities for same source and different source, and (2) use scores with known provenance from a database of similar ammunition and firearms.

similar to NIST approach of using a database to derive densities for same source (KM) and different source pairs (KNM) (in a setting of cartridge cases).

In either of the two scenarios, we derive scores by comparing test fires and the questioned bullet.

We evaluate the accuracy under each scenario by using a comparable set of tests with questioned bullets of known origin.

#### 5. Evaluate the error rates

(report theoretical error rates)

- given the cutoff point where the two distributions have equal density (also a likelihood ratio with value 1, which does an optimal classification under 0-1 loss? or at the point where the probabilities equal to each other instead of the densities)
- Consider four types of error rates (will need a table to illustrate this): false positive error rate, false negative error rate, false identification error rate, false exclusion error rate.
- Given the iid assumption, could we develop a further model for the bullet level distribution and estimate the corresponding error rates?

## 6. Estimations with changing sample size

(quantify the variation, reproduce some results in the previous sections with changing sample size)

## 7. Conclusion

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