

# Hidden Multiple Comparisons Increase Forensic Error Rates

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**When wires are cut, the tool produces striations on the cut surface; as in other forms of forensic analysis, these striation marks are used to connect the evidence to the source that created them. Here, we argue that the practice of comparing two wire cut surfaces introduces complexities not present in better-investigated forensic examination of toolmarks such as those observed on bullets, as wire comparisons inherently require multiple distinct comparisons, increasing the expected false discovery rate. We call attention to the multiple comparison problem in wire examination and relate it to other situations in forensics that involve multiple comparisons, such as database searches.**

Forensic Evidence | Statistics | Wire cuts | Toolmark analysis

In forensic evaluations, a single conclusion often relies on many comparisons, either implicitly or explicitly. Multiple comparisons arise persistently when developing statistical methods to address scientific problems (1), and greatly increase the probability of false discoveries. Now that vast databases and efficient algorithms are routinely used in forensic evaluations to propose matches to crime scene items, the problem of close non-matches (2) due to multiple comparisons becomes critically important. This often ignored issue increases the false discovery rate, and can contribute to the erosion of public trust in the justice system through conviction of innocent individuals. The multiple comparison problem is not new: it has been raised in the past with regard to DNA (3) and latent print evaluations (4). One of the root causes (5) leading to the wrongful accusation of Brandon Mayfield in the 2004 Madrid train bombing case was that the large size of the IAFIS database used to search for similar prints made it possible to locate ‘unusually’ close non-matches. As database size increases, so does the probability of finding a close non-match.

Compounding this issue, the use of algorithms also results in a large number of comparisons that are not obvious to the user. For example, the cross-correlation function (6), which computes the correlation for each alignment of two sequences, was one of the first measures proposed to quantify the similarity between two patterns in response to the 2009 NRC report (7), and continues to be used in many pattern searching algorithms to find the best alignment between two images and to quantify their overall similarity. Finding the best alignment often consists in sliding one surface across the whole length (for one-dimensional patterns, such as striations) or area (for two dimensional sources, such as impression marks) of the other item while keeping track of the value of a similarity measure. This mirrors the forensic examination process: the examiner visually rotates and shifts items under a comparison microscope to align two surfaces. In order to avoid false accusations and the corresponding impact on public perception of forensics, we must address the problem of multiple comparisons in database and alignment searches and control their effect on false discovery rates.

Here, we consider the multiple comparisons problem that arises from a relatively simple toolmark examination: matching a cut wire to a wire-cutting tool. We describe the comparison approach, estimate the (minimal) number of comparisons that are needed to carry out the examination, and discuss how the false discovery rate changes with the number of comparisons involved, using error rates derived from published black-box studies.

## Examination Process

A forensics examiner tasked with determining whether a wire in evidence was cut by a recovered tool will create one or more blade cuts, which are then compared to the cut surface of the wire recovered from the scene. These cuts are made in

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125 a sheet of material matching the wire composition, and may  
 126 be performed at multiple angles, as the angle of the tool to  
 127 the substrate can affect which striations are recorded on the  
 128 substrate surface. The blade cuts will then be compared to  
 129 the wire under a comparison microscope, though eventually,  
 130 automatic comparison algorithms may also be validated for  
 131 lab use. Each side of each blade cut will be compared to  
 132 each side of the wire; different tool designs have between 2  
 133 and 4 cutting surfaces in contact with the substrate.

## 135 Calculating the Number of Comparisons

136 In order to calculate the number of comparisons carried out  
 137 in the course of one examination, we define  $b$  to be the length  
 138 of the blade cut, and  $d$  to be the diameter of the wire. We  
 139 assume that the wire is covered with striations suitable for  
 140 comparison across its full diameter  $d$ . If this is not the case,  
 141 we reduce the value  $d$ . Both the blade and the wire are either  
 142 digitally scanned at resolution  $r$  mm per pixel, or visually  
 143 examined using a microscope with a digital resolution that  
 144 can be expressed as  $r$  equivalent to the digital scan. An  
 145 illustration of the sliding comparison process is shown in  
 146 Figure 1. Imagine that we move the cut wire along the  
 147 blade cut in order to assess whether striations on the blade  
 148 cut match the striations on the wire. We can move the wire  
 149 unit-by-unit, or we can move the wire by its full length, with  
 150 no overlap to the previous comparison.

151 The first option gives us the maximum number of  
 152 comparisons ( $b/r - d/r + 1$ ), while the second option gives us  
 153 the minimum number of comparisons  $b/d$ . In the first case,  
 154 sequential comparisons share much of the same physical data  
 155 and are highly related; in the second case, no data are shared  
 156 between physical comparisons and we can expect that they  
 157 are statistically independent, though empirically there will  
 158 be nonzero correlations due to physical similarities between  
 159 striations. For simplicity, let us consider the number of  
 160 comparisons to lie somewhere between these two estimates.  
 161 Note that when  $b/d \approx 1$ , as in some toolmark comparisons,  
 162 the upper number of comparisons goes to 1. Finally, we must  
 163 consider the number of surfaces which must be compared:  
 164 the wire may have one or two sets of striae and there may  
 165 be two to four blade cut surfaces to examine, depending on  
 166 the tool. This results in a multiplier of as much as 8.

167 **A concrete example.** Let us consider a wire-cutting tool with  
 168 a 1.5 cm razor blade that meets a cast surface (one such  
 169 tool is shown in Figure 1); the wire is held against this  
 170 rectangular cast surface as the blade is pushed into the wire,  
 171 splitting it in two. This is a minimal scenario - the wire  
 172 will acquire striations from one side of the blade, while the  
 173 blade itself has two cutting edges, which we will call side A  
 174 and side B. A blade cut of a sheet of aluminum will thus  
 175 produce two striated edges corresponding to side A and side  
 176 B which are compared to cut wires to assess similarity. We  
 177 also have a 12 gauge aluminum wire (2 mm diameter) which  
 178 may have been cut by the wire-cutting tool described above.  
 179 Class characteristics, which are shared by all tools of similar  
 180 manufacture, appear to match: there is a flat impression on  
 181 one side of the wire corresponding to the cast metal backstop  
 182 of the tool, and the wire is cut such that the blade and the  
 183 backstop appear to be perpendicular (that is, the wire  
 184 appears to have been cut with a tool of similar configuration).



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**Fig. 1.** (Top) A comparison between a wire and a blade cut requires sliding the wire along the entire blade cut length to determine the best match (or whether there is a match). Surfaces shown are rendered 2D topographical scans of a wire and blade cut taken with a confocal light microscope. (Bottom) RJ45 Crimp tool with a 1.5 cm razor blade used for cutting. 1 mm and 2 mm diameter aluminum wires cut with the pliers are shown in a box in the top right corner.

In this example,  $b = 15$  mm,  $d = 2$  mm, and there are at least  $b/d = 7.5$  comparisons between a wire cut and a blade cut. As there are two blade cuts (side A and side B), the minimal number of comparisons is 15, as these comparisons are non-overlapping and independent (on average).

Assuming a resolution of  $0.645\mu\text{m}$  per pixel, the maximum number of comparisons per blade cut is around 20,000; thus, we need 40,000 comparisons in order to find the optimal alignment between the wire and the blade cut. These comparisons are implicit in the calculation of cross-correlation, which is the first and often the only step used to quantitatively assess the similarity between striated evidence such as bullets, aperture shear, and firing pin impressions. Implicit comparisons are not unique to algorithms; an examiner would need to physically align the wire and the blade cut by searching along the length of the cut to visually match striations, performing the same process physically that the algorithm performs computationally. While these sequential comparisons are highly auto correlated, and we cannot assume sequential independence when calculating the probability of an error, they serve as an upper bound on the number of comparisons which could be performed. As the number of comparisons increases, the probability of encountering a coincidental match increases. Statisticians call this the *family-wise error rate E*; it is an important quantity to control when conducting a series ("family") of tests(8).

## Probability of False Discoveries

There are at least two components of the false discovery rate (FDR): identifying two pieces of evidence that have similar characteristics but are from different sources (a coincidental match) and procedural failures (e.g. lab process errors) (2, p 50). In objective disciplines with standardized evaluation rules (e.g. DNA), these sources can be distinguished. However, in toolmark examination, no objective evaluation rules are used; examiners testify based on subjective rules for how much similarity is sufficient for an identification. Assuming that lab procedure errors are not a factor in studies, we use reported error rates from three open-set studies of striated evidence(9–11) to obtain a ballpark estimate of the coincidental match rate of a single wire-cut

		False Discoveries (%) in N comparisons				
	Study	FDR $e$	$E_{10}$	$E_{100}$	$E_{1,000}$	$E_N < 0.1$
252	Mattijssen (2021)	7.24%	52.8	99.9	100.0	1
253	Pooled Error	2.00%	18.3	86.7	100.0	5
254	Bajic (2020)	0.70%	6.8	50.7	99.9	14
255	Best (2022)	0.45%	4.5	36.6	98.9	23
256		1 in 1,000	1.0	9.5	63.2	105
257		1 in 10,000	0.1	1.0	9.5	1,053
258		1 in 100,000	$10^{-4}$	0.1	1.0	10,535

259 **Table 1.** Table showing the relationship between false discovery  
260 rates and the chance of a false discovery in  $N$  comparisons for a  
261 set of different FDRs and different number of comparisons. The last  
262 column gives the number of comparisons allowed while ensuring a  
263 familywise false discovery percentage of at most 10%.

264 comparison. These studies have FDRs between 0.0045 (11)  
265 and 0.072 (10); pooling data from these studies weighted by  
266 sample size yields an FDR of 0.02. For a single-comparison  
267 FDR of  $e$ , the family-wise FDR for  $n$  comparisons,  $E_n$  is  
268  $1 - [1 - e]^n$ . Table 1 shows the impact the number of  
269 comparisons has on these published error rates. With an  
270 error rate of 0.07, as suggested by Bajic (2020), examiners  
271 can make up to 14 comparisons, i.e. even the simple example  
272 in this paper exceeds an upper bound of 10% for the family  
273 wise false discovery error. To conduct a search of a modestly  
274 sized database with 1000 entries, the initial FDR cannot  
275 exceed 1 in 10,000 to guarantee a family-wise total false  
276 discovery error of at most 10%.

277 Under these constraints, the accuracy of an examination  
278 involving multiple comparisons between a wire and a tool  
279 will be low, as the number of candidate alignments that  
280 must be examined is high. Even the most innocuous example  
281 (small blade, only 2 cutting surfaces, and a relatively large  
282 wire) involves a minimum of 15 comparisons. Examiners  
283 would make cuts under multiple angles (12), increasing the  
284 number of comparisons and making a false discovery even  
285 more probable. As a result, it is questionable whether  
286 wire comparisons made under current protocols are reliable  
287 enough to be presented at trial.

288 Clearly, studies for wire evidence, and larger studies for  
289 striated evidence in general, are necessary. Moving away  
290 from binary assessments toward quantification of striation  
291 similarity and observed pattern frequency will also reduce  
292 the severity of this issue and allow examiners to assign  
293 unusual striation patterns more weight in the process.

## 294 Discussion & Conclusions

295 Forensic practitioners often report the findings from their  
296 examinations in the form of a categorical conclusion  
297 reflecting a single decision. This is misleading when  
298 the decision relies on multiple comparisons which are not  
299 individually presented in reports or testimony. In this short  
300 contribution, we have shown that the implicit comparisons  
301 performed during forensic analysis of wire cuts increase the  
302 family-wise error rate.

303 We describe a simple scenario where a wire is cut using  
304 a two-sided blade, but findings apply to any situation  
305 where a forensic evaluation involves multiple comparisons,  
306 including, e.g., database searches. Forensic practitioners

307 should understand how the number of comparisons can affect  
308 the accuracy of their final conclusion. We propose three  
309 strategies to enhance transparency and enable more reliable  
310 estimates of examination-specific error rates.

311 First, examiners should report (or defense attorneys  
312 should request) the overall length or area of surfaces  
313 generated during the examination process, along with the  
314 total consecutive length or area of the recovered evidence.  
315 These pieces of information will take the place of  $b$  and  $d$   
316 and facilitate calculation of examination-wide error rates.

317 Second, researchers should conduct studies relating the  
318 length/area of comparison surface to the error rate. For  
319 instance, we have pooled studies looking at bullet striations  
320 and firing pin shear marks because we could not find black-  
321 box error rate studies of wire cuts. The striated surfaces  
322 are of orders of magnitude different lengths, but represent  
323 the best estimate of the error rate for striated materials.  
324 New studies should be designed to assess error rates (false  
325 discovery and false elimination) when examiners are making  
326 difficult comparisons.

327 Finally, when databases are used at any stage of  
328 the forensic evidence evaluation process (from suitability  
329 assessment and triage to reports which will be used at trial),  
330 the number of database items searched (or comparisons  
331 made) and the number of results returned must be reported.  
332 Additionally, the number of results used for further manual  
333 comparison should also be reported. For example, if a  
334 firearms examiner searches a local NIBIN database with 1000  
335 entries, requests the 20 closest matches to her evidence, and  
336 then carries out a physical examination of five exemplars  
337 from the list of 20, all of those values should be clearly  
338 reported to enable estimation of the familywise error rate.  
339 This will help make the multiple comparison issue accessible  
340 to everyone involved in evaluating the value of forensic  
341 evidence: examiners, lawyers, jurors, and judges.

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1. Y Benjamini, Y Hochberg, Controlling the False Discovery Rate: A Practical and Powerful Approach to Multiple Testing. *J. Royal Stat. Soc. Ser. B (Methodological)* **57**, 289–300 (1995).
2. President's council of advisors on science and technology, *Forensic science in criminal courts: Ensuring scientific validity of feature-comparison methods*. (Executive Office of the President of the United States, President's Council), (2016).
3. WC Thompson, F Taroni, CGG Aitken, How the probability of a false positive affects the value of dna evidence. *J. Forensic Sci.* **48**, 2001171 (2003).
4. JJ Koehler, S Liu, Fingerprint error rate on close non-matches. *J. Forensic Sci.* **66**, 129–134 (2021).
5. GA Fine, A review of the fbi's handling of the brandon mayfield case unclassified executive summary. (Washington DC), Technical report (2006).
6. T Vorburger, et al., Applications of cross-correlation functions. *Wear* **271**, 529–533 (2011).
7. NRC, *National Research Council: Strengthening Forensic Science in the United States: A Path Forward*. (National Academies Press), (2009).
8. J Tukey, Multiple comparisons. *J. Am. Stat. Assoc.* **48**, 624–625 (1953).
9. S Bajic, LS Chumbley, M Morris, D Zamzow, Validation study of the accuracy, repeatability, and reproducibility of firearm comparisons, (Ames, IA), Technical report (2020).
10. EJAT Mattijssen, et al., Firearm examination: Examiner judgments and computer-based comparisons. *J. Forensic Sci.* **66**, 96–111 (2021) \_eprint: <https://onlinelibrary.wiley.com/doi/10.1111/1556-4029.14557>.
11. BA Best, EA Gardner, An assessment of the foundational validity of firearms identification using ten consecutively button-rifled barrels. *AFTF J.* **54**, 28–37 (2022).
12. M Baker, Toolmark variability and quality depending on the fundamental parameters: Angle of attack, toolmark depth and substrate material. *Forensic Sci. Int.* p. 10 (2015).