

1 **TITLE:**

2 The importance of considering common sources of unknown DNA when evaluating findings  
3 given activity level propositions

4

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25

26 **KEY WORDS:**

27 Background DNA; activity level; evidence evaluation; Bayesian networks.

28

29 **HIGHLIGHTS:**

- 30
- 31 • We consider the evaluation of findings from multiple similar items
  - 32 • Bayesian Networks are used to construct evaluative frameworks
  - 33 • We demonstrate the importance of considering multiple sources of unknown DNA
  - 34 • We apply the theory presented to a real case scenario from South Australia
- 35
- 36
- 37
- 38

39 **ABSTRACT:**

40 Evaluating forensic biological evidence considering activity level propositions is becoming  
41 more prominent around the world. In such evaluations it is common to combine results from  
42 multiple items associated with the alleged activities. The results from these items may not be  
43 conditionally independent, depending on the mechanism of cell/DNA transfer being considered  
44 and it is important that the evaluation takes these dependencies into account. Part of this  
45 consideration is to incorporate our understanding of prevalent DNA and of background DNA  
46 on objects and people, and how activities can lead to common sources of unknown DNA being  
47 deposited on items. We demonstrate a framework for evaluation of DNA evidence in such a  
48 scenario using Object-Oriented Bayesian Networks and apply it to a motivating case from  
49 South Australia.

50

51 **INTRODUCTION:**

52 Evaluating forensic DNA results considering activity level propositions is an important task  
53 for the forensic scientist. It puts the DNA results in a case context that is not achieved with the  
54 evaluation of the same results given (sub-) source level propositions. There are guidelines and  
55 publications recommending the evaluation of results given activity level propositions [1, 2].  
56 There are a number of published examples of evaluations using activity level propositions  
57 available [2-6]. Depending on the case circumstances, the propositions of prosecution and  
58 defence may dispute the actor involved in the alleged crime, or the activity of the alleged crime  
59 [7]. If the actor is in dispute, then it is common to consider an alternate offender (often  
60 designated as AO), who is currently unidentified and who may have donated DNA that will  
61 present itself as originating from an unknown individual in the context of the forensic work  
62 being conducted. Another source of unknown DNA is the background DNA present almost  
63 ubiquitously on all items. The level to which the presence or absence of unknown DNA  
64 supports one proposition over the other then becomes a balance between the probabilities of  
65 the results given a DNA transfer from an AO compared to the presence of background DNA.  
66 However, as well as the fact that there is a presence of DNA from an unknown source on  
67 multiple items, there is additional information that can assist evaluations when considering if  
68 the same unknown contributor is present on these items.

69

70 One scenario that has not been explored extensively is how an evaluation should proceed when  
71 there are multiple similar items in a case. In this case there are two aspects to consider that will  
72 affect the evaluation, and the results of the evaluation:

- 73 1) The choice of whether to treat transfers to the similar items as one or multiple events.  
74 2) The probability of a common unknown contributor having donated DNA to the  
75 samples.

76 The first point is related to the issue commonly referred to as the ‘two-trace problem’, which  
77 was originally discussed by Evett [8], and considers when there are two or more stains and two  
78 offenders. This was later extended to generality with respect to the number and type of traces  
79 and the number of offenders by Triggs et al. [9] and tackled using Bayesian networks by  
80 Gittelsohn et al. [10]. A general explanation on the combination of dependant pieces of evidence  
81 was given by Juchli et al. [11]. We draw upon the concepts of Gittelsohn et al. [10] and Juchli

82 et al. [11] to show how propositions can dictate whether DNA transfer to multiple items are  
83 considered one, or multiple events, and how this impacts the evaluation.

84  
85 We also demonstrate the importance of considering the various pathways that unknown DNA  
86 can be deposited on to similar items, particularly if it is being suggested that they have been  
87 handled in a similar way, and by the same person (whether that be the defendant or an alternate  
88 offender). The classic treatment of common unknowns is to consider them as occurring either  
89 because they have come from a single unknown offender, or because they come from different  
90 individuals (as background), whose DNA profiles match. The probability associated with  
91 matching background DNA profile is typically set quite low, due to the discrimination power  
92 of DNA profiling systems. The value can be assigned based on a match probability if the  
93 unknowns are able to be interpreted, or could be based on mixture to mixture comparisons [12,  
94 13] if the unknowns are not resolvable. When common unknown donors are found to be  
95 present, the evaluation will show that the findings provide strong support for the presence of  
96 the alternate offender, as the probability of transfer, persistence and recovery of the DNA of  
97 that donor is much higher than matching profiles from different sources present as background.  
98 However, given our knowledge of the level of background DNA on items from being in  
99 proximity to a person (such as in their home [14], car [15] or workplace [16]), or the presence  
100 of an unknown person's DNA on the hands of the person who touched the item (e.g. such as a  
101 cohabitant [14]) there are other explanations for the presence of common unknowns on multiple  
102 similar items. It becomes quite important to consider these alternate routes for common  
103 unknowns within an evaluation in order to obtain a sensible result.

104  
105 Providing even more power to help to discriminate between propositions is to have the  
106 reference samples of people associated with the persons of interest in a case. Having profiles  
107 from these secondary associated individuals can eliminate, or confirm, certain common donor  
108 DNA transfer mechanisms, and is preferable to dealing with the uncertainty statistically. While  
109 we focus on a case example that has biological evidence, there are natural extensions of this  
110 thinking that can be applied to combine evidence across disciplines. We point the reader to the  
111 recent work of de Koeijer et al. [17] for discussions in this area.

112  
113 We work through the DNA results for the case R v QUIST heard in South Australia in 2016.  
114 The defendant was convicted, but this conviction was appealed and overturned in 2017 [18]  
115 (not on the basis of DNA evidence, but in the way the trial judge instructed the jury in various  
116 non-scientific matters). At the time of this paper being written a retrial has not occurred but is  
117 scheduled. The goal of this work is to demonstrate a Bayesian Network (BN) based method for  
118 evaluating DNA results from multiple similar items, and in particular we show the importance  
119 to the evaluation of considering difference sources of common unknown profiles on the similar  
120 items.

121  
122 We detail the case circumstances and forensic work in sections 2 and 3, and then elaborate on  
123 the different ways that the DNA results could be interpreted or evaluated in section 4.

## 124 125 **2.0 CASE SCENARIO:**

126 2.1 - Background information

127 At 5:15pm on the 23<sup>rd</sup> of December 2013 a fire started in the disabled toilets of a shopping  
128 centre in Parafield Gardens in South Australia. The fire had been started by igniting an open  
129 plastic bottle filled with petrol. The fire was extinguished, and during the processing of the  
130 scene (at approximately 7pm), six additional bottles (all filled with petrol) were found hidden  
131 in the ceiling space above the toilet. These bottles were taken out of the toilet ceiling (without  
132 touching the lid, and wearing appropriate protective clothing to minimise contamination), out  
133 of the toilet block and placed on the ground where the lids were immediately swabbed. There  
134 was no fingerprint detection work carried out on the body or lid of the bottles. The defendant  
135 in this matter was seen around the area at 5:15pm and leaving shortly after with burns to her  
136 body.

137

138 2.2 - DNA results

139 In this case there are seven plastic soft drink bottles involved. One bottle was set alight and  
140 was considered not suitable for DNA sampling. Six bottles were from the ceiling space and not  
141 burnt. South Australian Police took swabs of the lids of these six bottles and submitted them  
142 to Forensic Science SA. Of the six swabs of the lids of the hidden bottles, five of these were  
143 accepted for DNA profiling (it is unclear why one was not accepted). The results of DNA  
144 testing were that three of the five bottles had no DNA detected (and so did not proceed to DNA  
145 profiling) and the other two had approximately 0.8 and 0.7ng of DNA detected. The first of  
146 these (possessing 0.8 ng) yielded a three person mixture, that using STRmix™ V2.6 [19], were  
147 in proportions 82%, 13% and 6%. A DNA profile corresponding to the DNA profile of the  
148 defendant was observed in the major component (with the proportion of 82%) and a likelihood  
149 ratio (*LR*) was calculated using the following sub-source level propositions:

- 150 • Hp: The DNA originated from the defendant and two unknowns
- 151 • Hd: The DNA originated from three unknowns

152 The *LR* being ~ 4.8 billion in support of the defendant being a DNA donor to the sample rather  
153 than not.

154

155 The second sample (possessing 0.7 ng of DNA) yielded a two-person mixture, that using  
156 STRmix™ V2.6, were in proportions 94% and 6%. A DNA profile corresponding to the DNA  
157 profile of the defendant was observed in the major component (with the proportion of 94%)  
158 and an *LR* was calculated using propositions:

- 159 • Hp: The DNA originated from the defendant and an unknown
- 160 • Hd: The DNA originated from two unknowns

161 The *LR* being ~ 1.3 billion in support of the defendant being a DNA donor to the sample rather  
162 than not.

163

164 A mixture-to-mixture comparison was carried out (as per the method in [12, 13]) and an *LR*  
165 calculated considering the propositions:

- 166 • H1: There is a common contributor to the minor components of the mixtures
- 167 • H2: There are no common contributors to the minor components of the mixtures

168 With a  $LR \sim 50$  in support of H2 compared to H1. Therefore, the DNA observations on the  
169 minor components are more likely if different unknown individuals contributed.

170

171 Due to a combination of the heat from the fire, and the very public nature of the crime scene,  
172 no other DNA samples that were taken were deemed suitable for analysis.

173

#### 174 2.3 - The prosecution scenario

175 The prosecution alleges that the defendant filled the six hidden plastic bottles with petrol and  
176 placed them in the ceiling space of the disabled toilet, and then set fire to a seventh open bottle  
177 of petrol on the toilet floor.

178

#### 179 2.4 - The defence scenario

180 The defence alleges that someone other than the defendant (an alternate offender, AO) filled  
181 the six hidden bottles with petrol and placed them in the ceiling space, and then lit the seventh  
182 open bottle on the floor of the toilet. The fire died down due to lack of oxygen, but then  
183 reignited (the ‘flashback’ effect) when the defendant opened the toilet door, burning her. She  
184 then left the scene, and due to her injuries, she shed cellular material in the path she walked  
185 through the shopping centre. When the Police recovered the bottles, and walked them out of  
186 the bathroom to the point where they were sampled, the action of walking through the same  
187 space that had been walked through by the defendant, 90 minutes earlier, ‘reinvigorated’ the  
188 cellular material (presumably meaning that the cellular material was stirred up into the air).  
189 The cellular material in the air then settled on the bottle lids.

190

### 191 **3.0 THE TRIALS AND APPEAL**

192 In the trials held in 2016 and 2019 there were multiple facets of evidence adduced. These  
193 included the DNA evidence, and also evidence from fire experts, chemistry experts (to assist  
194 in identifying the contents of the bottles), CCTV footage, and eyewitnesses. We will only  
195 concentrate on the DNA evidence component of the case.

196

197 During the original trial in 2016, the defence scenario was put to the DNA expert. The expert  
198 replied by stating that:

199 *‘...I couldn’t exclude it as a possibility, but if there was a time delay in between, then I would*  
200 *lean towards it being an unlikely way for DNA to transfer’*

201 She also brought up the point that if shed DNA being stirred up into the air and depositing on  
202 objects was a good explanation for the transfer then it may have been expected to find highly  
203 complex mixtures on the bottle lids (due to the very public nature of the shopping centre where  
204 they were swabbed). Finally, it was brought up by the prosecution that if the ‘air DNA’ scenario  
205 was a good explanation for DNA transfer, then we might expect to see a similar result on all  
206 the bottle lids. Note that this comment goes directly to the heart of whether we consider the  
207 possibility of transfer to the bottle lids as independent events or a single event.

208

209 In the 2017 appeal ruling [18] the aspect of the testimony of original DNA expert witness  
210 obviously weighed heavily in the minds of the Judges. Evidence by the fact that the description  
211 of the defence scenario is given in a section of the ruling entitled:

212 'A highly far-fetched theory of innocent indirect DNA transfer'

213

214 In this paper, we would like to go beyond the intuitively appealing argument (as made by the  
215 Judge) suggesting that the findings support the allegations of the prosecution and that they  
216 would be unlikely under the activities envisaged by the defence. The evaluation is better placed  
217 to be made in a forensic context, rather than as an intuitive opinion. Also, we will show that  
218 the complexity of the evaluation renders difficult for the forensic scientist to offer on the spot,  
219 at trial, a fully articulated response. These cases need to be fully assessed before trial. Object-  
220 oriented Bayesian networks (OOBN) offer the flexibility to handle forensic evaluation problem  
221 with multiple pathways of DNA transfer and multiple items. The literature though is often  
222 limited to one item or a limited set of results. In the case above, we could be tempted to ignore  
223 the three bottles with the low quantity of DNA and concentrate the evaluation on the two bottles  
224 that provided DNA results that have found correspondence with the defendant. In doing so, we  
225 would ignore part of the results, whereas these results, on the three other bottles, were  
226 considered by the reporting officer when questioned in court. We aim at constructing the  
227 inference engine in the form an OOBN that could tackle all the forensic DNA results. It will  
228 allow us to assess if the defence scenario was that fanciful in the sense that the results could be  
229 hardly expected under that view, but more expected under the prosecution view.

230

231 We develop an evaluation framework that shows how findings such as those in this case can  
232 be considered together, and the importance of considering different sources of unknown DNA,  
233 and whether they are from a common source. While we calculate an *LR* under different  
234 assumptions by assigning values to the probabilities of events occurring that are important to  
235 the evaluation, our goal is not to assign an *LR* that would be suitable for this case specifically  
236 (indeed we assign probabilities in some instances with no informative data), but rather to show  
237 the mechanism by which various aspects of the evaluation can be incorporated.

238

#### 239 **4.0 EVALUATION:**

240 Given the case information and the scenario given by prosecution and defence, we consider the  
241 following activity level propositions in the evaluation of the findings in the case:

- 242 • Hp: The defendant filled the soft drink bottles with petrol and placed them in the ceiling  
243 space of the public toilet
- 244 • Hd: An unknown offender filled the soft drink bottles with petrol and placed them in  
245 the ceiling space of the public toilet

246

247 The additional information we use is:

- 248 • The defendant was in the toilet when the fire was lit. She then moved through the  
249 shopping centre to exit the building.
- 250 • The bottles were removed from the ceiling space by Police 90 minutes after the incident  
251 and taken out into the shopping centre area (where the defendant had earlier walked  
252 through) and the bottle lids were sampled for DNA.

253

254 The assumptions we make in the evaluation of the findings in this case are:

- 255 • The same person who filled the bottles with petrol, also placed them into the ceiling  
256 space of the public toilet. This affects the area that we might expect DNA to be  
257 transferred to i.e. to the caps of the bottles from the person who filled them (where the  
258 samples were taken from) or to the body of the bottles from the person who placed them  
259 in the ceiling
- 260 • That the offender (whether this was the defendant or another person) did not wear  
261 gloves when handling the bottles
- 262 • That when the bottles were taken out of the toilet for swabbing that if DNA has been  
263 invigorated into the air, that this is a single event, and not one invigoration event per  
264 bottle given the narrow window of time in which this occurred
- 265 • That due to the fact that there were no issues reported with quality controls within the  
266 laboratory we consider the possibility of contamination to be a separate event for each  
267 bottle (i.e. as opposed to a reagent contaminated with the defendant's DNA, which  
268 would be a single event). We also apply this same reasoning to extend to the work done  
269 by the Police when sampling the bottles
- 270 • That the defendant is a donor of DNA to two bottles
- 271 • That the order of DNA bottle handling by the offender has not had an effect on the  
272 DNA transferred to the bottles
- 273 • That there are no common unknown DNA donors to the two bottles (we will however,  
274 explore the impact of this assumption on the *LR* in section 4.5)
- 275 • Note that we do not model persistence in our evaluation and so knowledge of the  
276 timeframe is not strictly necessary here. However, by not considering persistence the  
277 model assumes the DNA has persisted in the same state as the initial transfer. In making  
278 this assumption, we have used the knowledge of a relatively short timeframe between  
279 offence and sampling (90 mins).

280  
281 We explore the consequences of common unknowns. We will also treat the DNA as being  
282 present or absent on an object, rather than dealing with DNA amounts. This choice is, again,  
283 made simply to keep the OOBN complexity to a minimum, and allowing focus on the main  
284 point of the paper, but note that systems such as that in Taylor et al. [20] could be used. We  
285 construct OOBNs using software HUGIN [21] and follow the OOBN construction method of  
286 Taylor et al. [3].

287  
288 Throughout our paper we consider that background DNA is DNA of unknown origin on an  
289 item that is not being explicitly modelled as coming from a specific person. By "modelled",  
290 we mean that given the sampling location, and the framework of circumstances we account for  
291 its presence or absence in the context of some defined event (such as a transfer or prevalence).  
292 For example, on AO hands we expect to find AO's DNA, but this is not considered background  
293 as we are modelling it as coming from AO (using its expected prevalence). Other DNA of  
294 unknown origin on AO hands however is considered background as it is not modelled as  
295 coming from a designated person. Hence, we consider unknown DNA to be any DNA  
296 (background or modelled) which cannot be accounted for by one of the reference DNA profiles.

297 In the example of DNA on AO hands, both AO DNA and background DNA on AO hands are  
298 considered unknown DNA. As is evident from this example there is some overlap between  
299 background and unknown DNA.

300

301

#### 302 4.1 – Transfer to bottles, one event or many?

303 In order to evaluate the evidence in any case where transfer to multiple items may have  
304 occurred, the level of dependence existing between them must be considered. One possible  
305 course of action is to employ a simplification to the model so that all objects are considered as  
306 one, and transfer is considered to have occurred to the meta-object if it has occurred to any of  
307 the objects. This is a similar suggestion to one of the steps that was suggested in Taylor [3],  
308 and could be used if the items have a very close relationship. For example, it could be that the  
309 handle of a knife was divided into ten parts, which were swabbed separately. These ten swabs  
310 could be considered together (just as though a single swab had been used to sample the entire  
311 handle).

312

313 Such a simplifying assumption may not always be appropriate. In the motivating case the  
314 prosecution alleges that the defendant has placed the bottles in the ceiling cavity and hence had  
315 contact with each separately (in fact it is of no lesser consequence if the defendant has only  
316 placed two of the five bottles in the ceiling). If the defence proposition stipulated that the  
317 defendant had never had contact with the bottles, and never been in the area where the bottles  
318 were found, we could consider that the only possible mechanism for the presence of DNA  
319 matching that of the defendant is if an alternate offender possessed a matching DNA profile  
320 (ignoring the possibility of laboratory contamination, or error). In this case the activity level  
321 *LR* would take the value of the inverse of obtaining chance matching profile (i.e. the same  
322 numerical value as the sub-source *LR*) and the combination of presence or absence of the DNA  
323 matching the defendant on combination of bottles will not affect the *LR* as long as a profile  
324 matching the defendant is on at least one bottle i.e. the *LR* obtained if one bottle had a profile  
325 matching the defendant is the same as if all had matching profiles. In this instance the bottles  
326 could be considered as one meta-object.

327

328 If, however the defence proposition stipulated that the presence of DNA matching the  
329 defendant on any bottle has arisen from contamination in the laboratory, then we may consider  
330 each bottle with the presence of the defendant's DNA as an independent contamination event.  
331 In this case the specific combination of presence or absence of DNA of the accused on the  
332 bottles will have an effect on the *LR* i.e. the *LR* obtained if one bottle had a DNA profile  
333 matching the defendant would be much lower than if all had matching profiles. In the  
334 contamination scenario we would also have to consider that an alternate offender would have  
335 had to place the bottles in the ceiling and so the presence or absence of a common unknown  
336 profile would also become important.

337

338 In the motivating case the defence scenario is that the bottles were placed in an area where the  
339 defendant's DNA was swirled around in the air, which then settled on the bottles. We do not  
340 consider that there have been five separate instances of the defendant's DNA being swirled



341 into the air and landing on one bottle. In this scenario the results of all bottles are linked (or  
342 have dependence with each other). Further, we must consider that even though the transfer of  
343 DNA is considered one event in this instance, due to slight variations in transfer and recovery  
344 of the traces, it is still possible that DNA matching the defendant could be recovered from some  
345 bottles and not from others (indeed this is the situation we have in the motivating case). To  
346 evaluate the DNA findings in the motivating case we consider a single event in which the  
347 defendant's DNA has been swirled back into the air around the bottles. Then the probability  
348 that the defendant's DNA has settled on each bottle (given it is swirling around in the air) can  
349 be assigned considering the event for each bottle as conditionally independent. Note here we  
350 are considering an 'event' as the movement of DNA, i.e. from the ground into the air, or from  
351 the air onto a bottle.

352

353 As with the scenario previously put forward that considered laboratory contamination as the  
354 source of the defendant's DNA on the bottles, the *LR* obtained from the air swirling scenario  
355 will be sensitive to the number of bottles that have the defendant's DNA and/or unknown DNA  
356 as there is both a parent factor (swirling), and independent settling factors that will be assigned  
357 probabilities that are different to the probability of DNA transfer from handling.

358

#### 359 4.3 – Creating class networks

360 In Figure 1 we show the Object-Oriented BN (OOBN) that has been developed to evaluate the  
361 evidence in this case. There are five parts, in three layers of class networks. At the highest layer  
362 is the main BN (Part A of Fig 1), which is where the user would interact with the BN to  
363 instantiate information about the case and get the *LR*. In the main BN, the structure has been  
364 added to automatically calculate the *LR* and its inverse, purely for convenience, as described  
365 in [3, 22]. In the main network there are DNA profile results nodes for each of the bottles  
366 (nodes 20 – 24), but then there is also a series of nodes dedicated to whether common unknown  
367 contributors have been observed (nodes 10 – 19). There are a couple of ways that this could  
368 have been carried out, one would be to have a single node that considered all possibilities of  
369 common unknown configurations, i.e. considering the unknowns on bottles B1 to B5, the node  
370 would have states:

- 371 •  $B1 \neq B2 \neq B3 \neq B4 \neq B5$
- 372 •  $B1 = B2, B1 \neq B3 \neq B4 \neq B5$
- 373 • ...
- 374 •  $B1 = B2, B3 = B4, B1 \neq B3 \neq B5$
- 375 • ...
- 376 •  $B1 = B2 = B3 = B4 = B5$

377 The alternate possibility, and the one used in our OOBN construction, is to have one node per  
378 possible pairing, each with states of 'Y' or 'N'. We chose the latter option due to the number  
379 of options and resulting BN table size required for the first configuration. Because the  
380 invigoration of the cells into the air is assumed to be one event, it sits up at the main, outermost  
381 BN layer, as do the activity nodes (in blue) and the root nodes (in grey) relating to the presence  
382 of background DNA on the hands of the defendant (D) or alternate offender (AO). We explain  
383 later the reason for including the presence of unknown background DNA on the hands of an

384 alternate offender, who themselves also has an unknown profile. Finally, root nodes for the  
385 probability of profiles of the AO (or an unknown on the hands of the AO) matching the  
386 defendant are at the highest level of the BN as there are global values that apply to all possible  
387 matching scenarios.

388

389 Within the main BN are class objects that relate to each bottle, and also a class network that  
390 relates to matching unknowns. The Bottle class BN (Part B of Fig 1) possesses nodes that  
391 consider the various DNA transfers that could occur from people that may have handled the  
392 bottles, the background DNA on the bottles, the potential for laboratory contamination<sup>1</sup>, and  
393 the potential for various sources of DNA to match key actors in the evaluation. The Bottle BN  
394 also considers the probability of DNA being recovered, as the various bottles samples are taken  
395 as separate extracts and may have different extraction efficiencies. The outputs of the Bottle  
396 class BN are the DNA results from that bottle (which are then carried back out to the main BN  
397 layer so that the user need only interact with the BN on that level) and the presence of specific  
398 sources of unknown DNA that will then be used in the unknown DNA matcher. Note that the  
399 Bottle class BN also have DNA profile matcher class BNs (Part D of Fig 1) that, at the third  
400 layer of class network, deals with the possibility of by-chance matching profiles. As the DNA  
401 profile matcher has the matching probability as an input, these must be passed in from the  
402 Bottle class network. In some instances, these come from inputs, taking values from the  
403 outermost layer of the BN, but in the case of background DNA we pass in separate probabilities.

404

405 The main BN class network also possesses an unknown matcher class BN (Part C of Fig 1) that  
406 takes the unknown DNA output nodes from the bottle class BN and compares pairs of these  
407 for each possible bottle pairing, within a Single-bottle Unknown Matcher class BN (Part E of  
408 Fig 1). At this third layer of class BN in the Single-bottle Unknown Matcher the possibility of  
409 by-chance matching background DNA is considered as a root node (node 45). This root node  
410 could be set up as an input and passed in a value from the unknown matcher class BN for each  
411 bottle, if different values were desired (perhaps if relatively neutral *LRs* had been obtained from  
412 mixture to mixture matching, e.g. see [12]), however we have made the simplifying  
413 architectural decision to use a single value for all unknowns across all bottles.

414

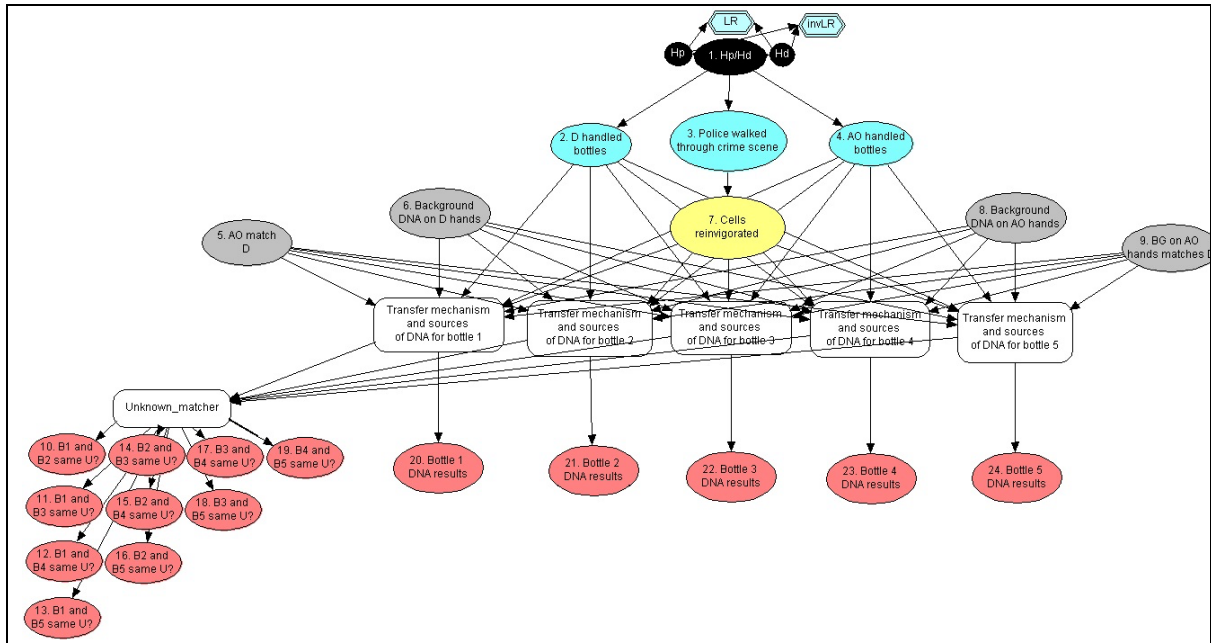
415 The assignment of probabilities to the nodes within the BN shown in Figure 1 is not the main  
416 focus of the paper and so, although important, we do not spend time on explaining the  
417 assignments, or carrying out sensitivity analyses. We provide the conditional probability tables  
418 as supplementary material (and the OOBN itself) as supplementary material for the interested  
419 reader.

420

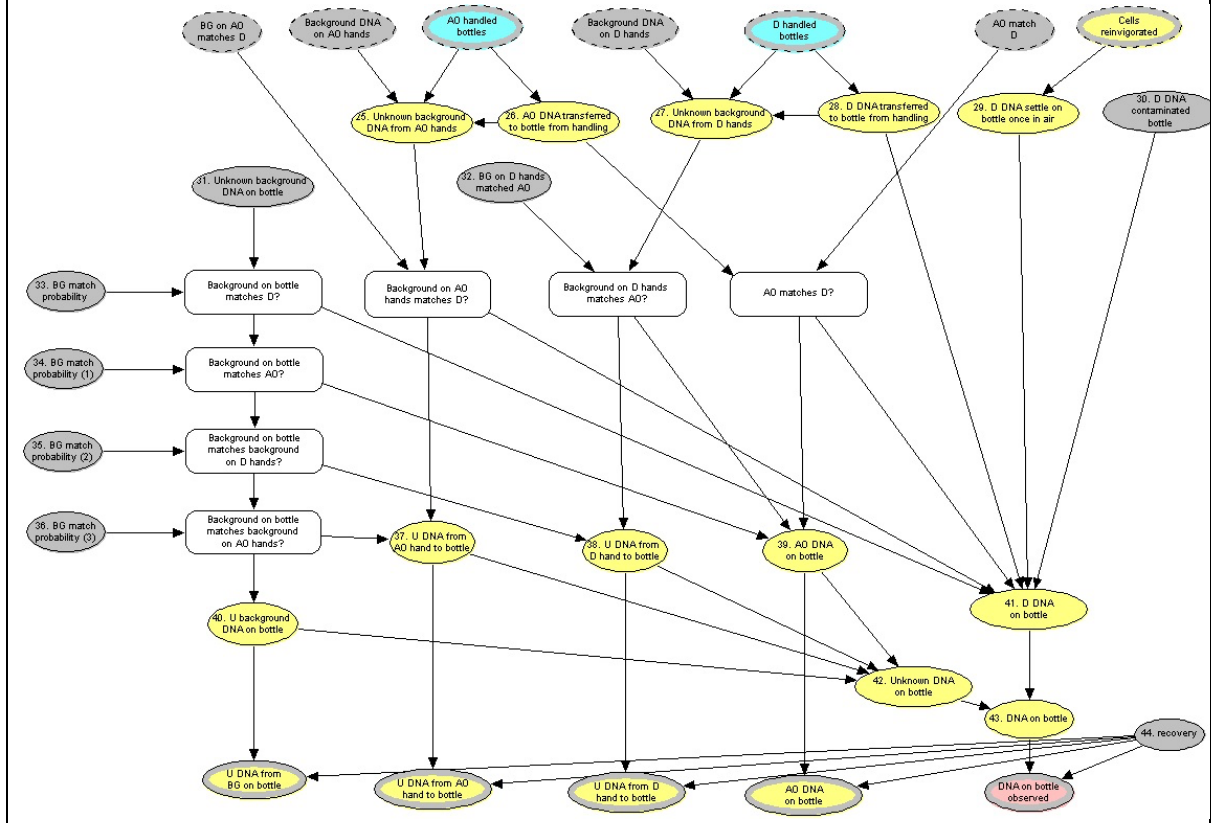
A) Main BN

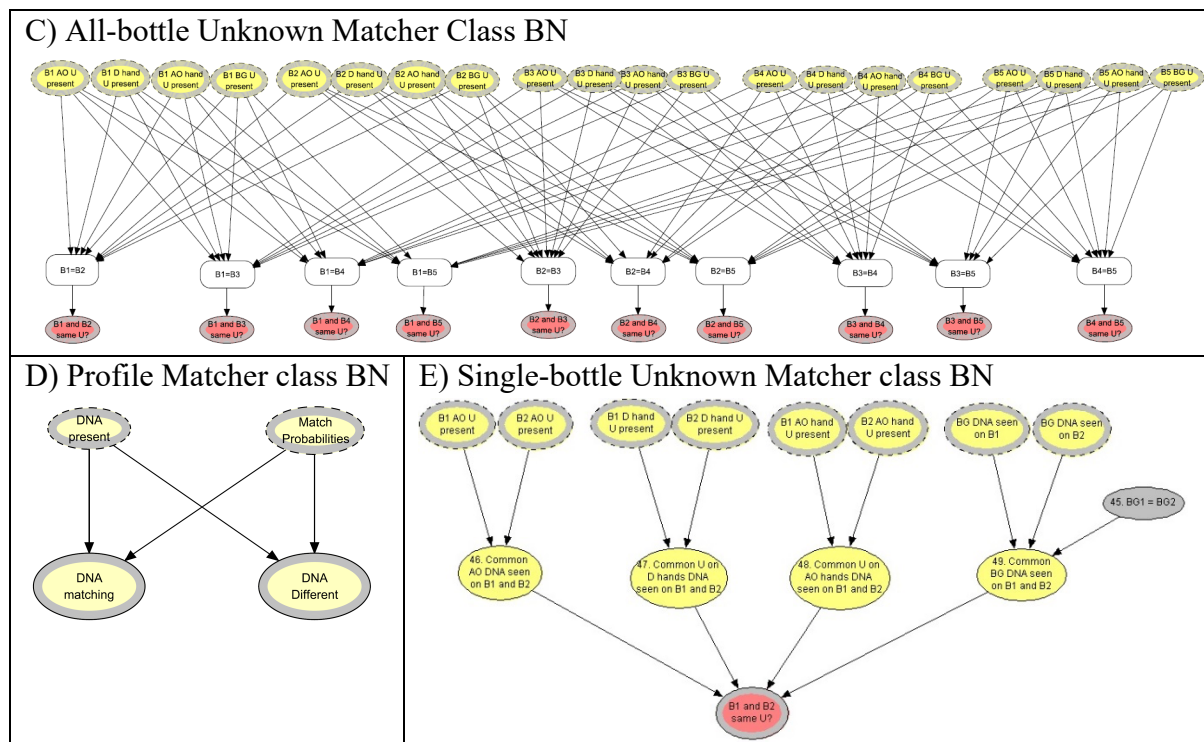
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<sup>1</sup> Note that the ‘air DNA’ mechanism itself should be considered an occurrence of contamination that has occurred as part of the processing of the crime scene. The additional contamination node relates to the probability of other, more conventional, contamination routes e.g. transfer via gloves.



**B) Bottle class BN**





422 *Figure 1: BN considering results for all bottles in the case, and whether common unknowns*  
 423 *were present. Panel A shows the over-arching main BN, parts B and C the second layer of*  
 424 *class BNs dealing with individual bottles and unknown profile pairwise comparisons and parts*  
 425 *D and E show the third layer of class networks dealing with profile and unknown matching.*  
 426 *Input nodes are signified by grey border with a dashed outline, and output nodes are signified*  
 427 *by a grey border with a solid outline. The function nodes ‘LR’ and ‘Inv LR’ allow for*  
 428 *calculation of the ratio of the posterior probabilities for each proposition and are not an*  
 429 *integral part of the network.*

430

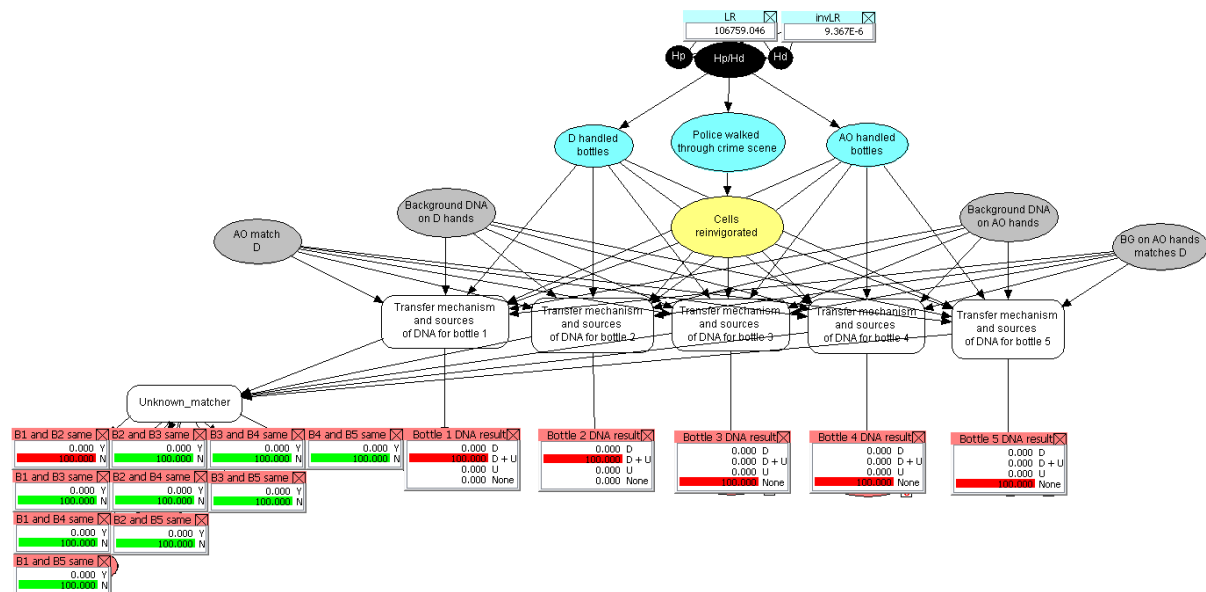
#### 431 4.4 – Providing the results from the case

432 Given the mixture-to-mixture result (i.e. an  $LR = 50$  in support of the propositions that the  
 433 bottles did not possess a common donor compared to them possessing a common donor, other  
 434 than the defendant) we assume that a common unknown donor does not exist. Given this, and  
 435 the other results previously described, the following case results can be instantiated:

- 436
- 437 • B1 and B2 same U? – N
  - 438 • Bottle 1 DNA results – D+U
  - 439 • Bottle 2 DNA results – D+U
  - 440 • Bottle 3 DNA results – None
  - 441 • Bottle 4 DNA results – None
  - 441 • Bottle 5 DNA results – None

442 and the  $LR$  obtained is approximately 107 000 in support of  $H_p$  over  $H_d$ . We treat ‘U’ as being  
 443 the presence of any unknown DNA (from one or multiple sources), and that matching  
 444 unknowns is matching between any of these ‘U’s on the bottles being compared. This is a  
 445 simplification of reality, in that it is possible to have different matching unknowns on different  
 446 sets of bottles, with one or more bottles having multiple unknowns belonging to multiple sets.

447 Our BN does not account for such refinement of unknown matching. The instantiated form of  
 448 the main BN is shown in Figure 2.  
 449



450  
 451 *Figure 2: Main BN seen in Part A of Figure 1 with case findings instantiated*  
 452

453 In Figure 2 it can be seen when the result of the DNA profiling on bottles are instantiated to  
 454 ‘None’ then the unknown matching nodes only have the possibility of ‘N’ (no unknowns  
 455 matching) due to the fact that there is no unknown DNA to match.  
 456

457 The *LR* shown in Fig 2 is quite high compared to many evaluations that consider latent DNA  
 458 samples [23]. The strength comes from the fact that the probability of the results under the  
 459 defence routes are driven by:

- 460 • the air invigoration mechanism, which is an accumulation of an air invigoration (with  
 461 a probability of 1 in 1000) and two instances of cells landing on the bottle tops (each  
 462 with a probability of 1 in 100).
- 463 • Contamination of the two bottles with the defendant’s DNA (each with a probability of  
 464 1 in 1000)

465 Which is quite small compared to the probabilities for the results given the prosecution DNA  
 466 transfer route, which is two transfer events from direct touching (each with a probability of  
 467 0.5). Indeed, given the values we have assigned to the probabilities of DNA air invigoration or  
 468 DNA (once invigorated) settling back onto the bottles, it is not surprising that the level of  
 469 support given to Hp over Hd is large. With such assignments we could have intuitively been  
 470 able to predict such an outcome. It is likely this intuition that led to the appeal judge’s  
 471 description of the scenario as far-fetched or guided the answers of the forensic scientist in court.  
 472 In our assignments of these probabilities we have not relied on any data, as the assignment  
 473 itself is not the focus of our work (rather the architecture of the evaluation), however if the case  
 474 were evaluated for court purposes it would require more rigorous testing, and likely an  
 475 investigation into the sensitivity of the *LR* to the assignment, which would inform the analyst  
 476 as to whether the evaluation was robust.

477

478 4.5 – Dealing with the presence of unknowns in the evaluation

479 Having seen the results of the evaluation using the full network, we will now demonstrate why  
480 the seemingly extensive consideration of the presence of unknown DNA is important in this  
481 evaluation.

482

483 In the motivating case, the defence proposition stipulates that if the defendant has not placed  
484 the bottles in the ceiling then an AO has. As a specific person has not been suggested, we  
485 cannot obtain a reference and their DNA would be interpreted as unknown. If an AO has placed  
486 the bottles in the ceiling then we may expect to see a common unknown profile on the bottle  
487 top, however there are other mechanisms (both under the prosecution or defence propositions)  
488 by which we may also find a common unknown profile on multiple bottles:

- 489 • An unknown background DNA source is present on the defendant's hands (UH) and  
490 this is transferred to the bottles when they are touched by the defendant.
- 491 • There is background DNA (BG) on the bottles from different sources and these have  
492 the same alleles (or similar enough to result in support for a common donorship)
- 493 • UH is transferred to some bottles and matches the BG on others
- 494 • BG is on the bottles and matches the AO on others
- 495 • There is unknown background DNA on the AO hands and this is transferred to the  
496 bottles when they are touched by the AO
- 497 • BG is on the bottles and matches the unknown background on the AO hands

498

499 We do not consider aspects of matchings such as the defendant having background DNA on  
500 their hands that matches themselves. We also do not consider the possibility of background  
501 DNA being present on the first bottle, and then transferred to the hands of the offender and  
502 then on to subsequent bottles. This mechanism could be subsumed into the common unknown  
503 background DNA on the hands of D or AO being transferred to the bottles, as long as the first  
504 bottle is the source of the transferred background. If it is not then we are in a position that we  
505 need to consider the order of the bottles being handled, and as this is not known then we need  
506 to model uncertainty in the bottle handing order, which adds additional complexity into the  
507 evaluation.

508

509 The consideration that a common unknown comes from the hands of the defendant as  
510 background is important in order to obtain a result that is intuitive with our understanding of  
511 DNA in the real world. It is not uncommon for individuals to carry non-self DNA on their  
512 hands and transfer this through touching or handling an object [24-26]. Without this  
513 consideration, then any presence of matching unknown DNA on the bottles must be described  
514 under the prosecution proposition as occurring by the chance matching of alleles (an extremely  
515 unlikely event) and will drive the *LR* very strongly towards support for the defence proposition.  
516 In the example seen in Figure 2, if we instantiate the node 'B1 and B2 are same U?' to 'Y' then  
517 the *LR* increases to approximately 388 000 in favour of Hp over Hd. While this may initially  
518 seem counter-intuitive, it is a result of the fact that given D's DNA has been found on the  
519 bottles, the most probable source of a common unknown is now unknown BG on D's hands

520 (hence adding more support to Hp). If the node ‘Background DNA on D hands’ is instantiated  
 521 to ‘N’ (mimicking a BN that did not consider the BG on D’s hands as a potential source of  
 522 common unknown DNA) then the *LR* becomes approximately 4300 in favour of Hd over Hp  
 523 i.e. if we hadn’t considered the possibility of a common unknown source of DNA coming from  
 524 D’s hands in this BN then the effect would have been a shift in the *LR* by a factor of  
 525 approximately 1 billion. This is because without the common BG on D’s hands the only  
 526 possibility for obtaining a common unknown under Hp is via chance profile matching  
 527 background occurring (with probabilities of 1 in 1 billion).

528  
 529 If the DNA on the bottles is only unknown (i.e. there is no DNA from the defendant), then the  
 530 additional information of those unknowns having a common source provides more support to  
 531 Hd. We show several alternative instantiations of (fictitious) results in Table 1.

532

Bottles	Common U DNA	Background hands		<i>LR</i> (support for)
		D hands	AO hands	
B1 – D+U B2 – D+U B3 – None B4 – None B5 – None	B1≠B2	Allowed	Allowed	106 759 (Hp over Hd)
		No	Allowed	180 712 (Hp over Hd)
		Allowed	No	105 878 (Hp over Hd)
		No	No	179 220 (Hp over Hd)
B1 – D+U B2 – D+U B3 – None B4 – None B5 – None	B1=B2	Allowed	Allowed	388 193 (Hp over Hd)
		No	Allowed	4301 (Hd over Hp)
		Allowed	No	374 157 (Hp over Hd)
		No	No	4463 (Hd over Hp)
B1 – U B2 – U B3 – U B4 – U B5 – U	B1=B2=B3=B4=B5	Allowed	Allowed	6 400 000 (Hd over Hp)
		No	Allowed	1.6 x 10 <sup>46</sup> (Hd over Hp)
		Allowed	No	6 400 000 (Hd over Hp)
		No	No	1.6 x 10 <sup>46</sup> (Hd over Hp)
B1 – U B2 – U B3 – U B4 – U B5 – U	B1≠B2≠B3≠B4≠B5	Allowed	Allowed	11 (Hd over Hp)
		No	Allowed	12 (Hd over Hp)
		Allowed	No	10 (Hd over Hp)
		No	No	11 (Hd over Hp)
B1 – D B2 – D B3 – D B4 – D B5 – D	not relevant as there is no unknown DNA present	Allowed	Allowed	1 x 10 <sup>9</sup> (Hp over Hd)
		No	Allowed	2 x 10 <sup>9</sup> (Hp over Hd)
		Allowed	No	5 x 10 <sup>8</sup> (Hp over Hd)
		No	No	1 x 10 <sup>9</sup> (Hp over Hd)

533 *Table 1: LR obtained for varying instantiations of results*

534

535 Note that in our BN construction we have chosen to include the consideration of common  
536 unknowns arising from BG DNA on the AO hands. This may seem odd, given that the AO  
537 themselves will already appear as unknown DNA. The reason for this is that in order to  
538 maintain sensible behaviour in the BN we seek to treat the defendant and the AO equally i.e.  
539 whatever the probability of DNA transfer is for one, so too should it be for the other. In  
540 somewhat special circumstances, we could have different transfer probabilities for D compared  
541 to AO (for example D could be a very poor shedder and we use an average shedder for AO).  
542 In the BN shown in Figure 1 the presence of unknown DNA on the AO hands has little effect  
543 on the *LR* (as seen in Table 1 in the differences between *LRs* when both unknowns on hands is  
544 allowed as background vs when the unknown on AO hands is not allowed) and so could  
545 arguably have been left out. However, our own anecdotal experience with workshopping  
546 counterintuitive BN behaviour has sometimes arisen from unbalanced treatment of defendant  
547 and AO in the architecture of the BN.

548

#### 549 **DISCUSSION:**

550 The BN we have constructed in Figure 1 takes into account the possibility of contamination of  
551 exhibits with the defendant's DNA, and also the possibility for chance matching DNA profiles  
552 between different sources of DNA. While the defence do not specifically mention these as part  
553 of their proposition or case argument, we do not believe there is a need for them to do so.  
554 Indeed, whether or not they mention these occurrences will not have affected the probability  
555 of their occurrence. In many evaluations the consideration of these relatively improbable events  
556 will not have a significant effect on the *LR*, as the presence of DNA matching the defendant is  
557 much more probable given alternate explanations, and so will limit the size of the *LR*. In  
558 evaluations such as the one demonstrated, where the scenario being put forward by defence  
559 requires one or more quite improbable events to occur, the inclusion of events such as  
560 contamination or chance matching DNA profiles become more important to include.

561

562 There are various other ways in which the BN in Figure 1 could be extended. We could consider  
563 DNA amounts throughout the network rather than simply the presence or absence. This would  
564 allow us to consider extraction and sampling efficiency distributions as in [23]. This would  
565 also allow a more detailed evaluation with regards to the air DNA mechanism and how the  
566 amount of DNA expected to be deposited from such a route compares to the amount of DNA  
567 expected from a direct transfer. Incorporating DNA amounts into the BN would also allow us  
568 to make use of the fact that the defendant's DNA was the major component of the mixtures  
569 observed in the two bottles that had DNA profiles generated. The difficulty with extending the  
570 BN to consider DNA amounts comes from the lack of knowledge regarding DNA amounts that  
571 relate to different transfer mechanisms in this study, particularly the air-DNA mechanism for  
572 which there is little to no relevant literature available. Hence, even if adding DNA amounts  
573 may have offered increased discrimination between propositions, adopting a presence/absence  
574 strategy is the adequate level of granularity allowing to maintain operational use. As with any  
575 evaluation there is always additional factors that could be considered, and there is sometimes  
576 a choice as to the level of complexity required to provide meaningful guidance to the court.

577

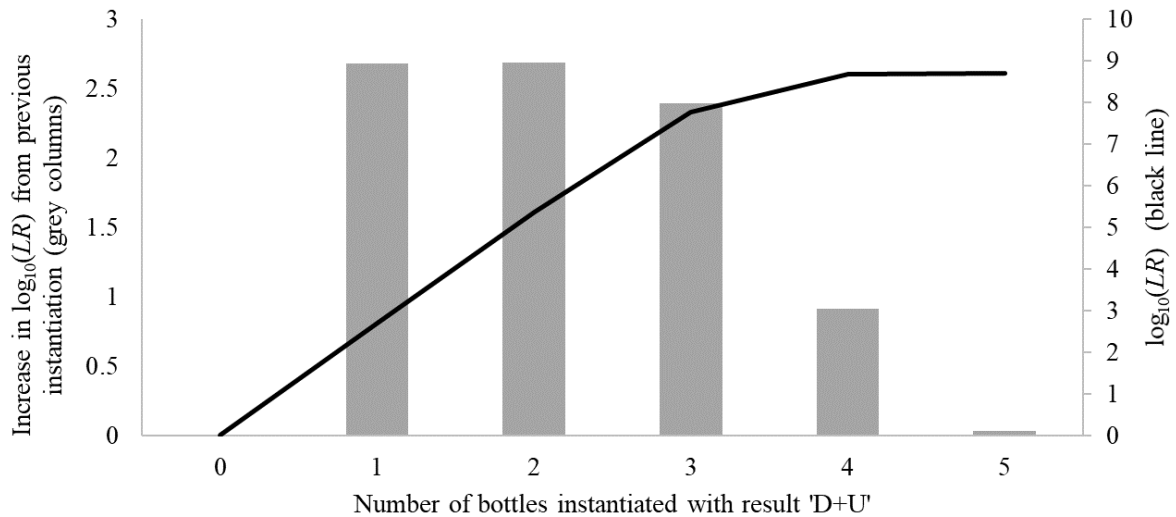


578 Another consideration could be that if the air-DNA mechanism is to resuspend DNA from the  
579 surroundings into the air then we could consider the background DNA in the environment, so  
580 that in the BN if the defendant's DNA was resuspended, so too might we expect unknown  
581 DNA to be also. In this instance the evaluation could have been assisted by taking background  
582 DNA swabs (i.e. of the floor or area around where the bottles were swabbed by police) to see  
583 if the defendant's DNA was indeed present. We note though that this is not usual practice for  
584 most forensic crime scene processing, and the relevance of doing so (i.e. the knowledge of the  
585 defence proposition) was not known for three years after the offence. Given that indirect  
586 transfer mechanisms are increasingly proposed as 'explanations' for finding DNA on items of  
587 interest, anticipating such scenarios by incorporating collection of background samples in  
588 routine crime scene procedure may be advisable [27].

589  
590 Note the importance of including all the bottles in the evaluation, even though no DNA was  
591 obtained from some of the extractions (note though that there is no benefit to including the  
592 bottle swab that was not analysed in the BN as a complete lack of knowledge of the DNA result  
593 provides no power to discrimination between propositions; a lack of knowledge is quite  
594 different from a lack of DNA). This consideration would be particularly important if DNA  
595 amounts were used in the BN, but even just considering the presence or absence of DNA, there  
596 is an effect of knowing that little to no DNA was obtained from three of the bottles. In the case  
597 scenario the  $LR$  when providing all case information was 107 000 in support of  $H_p$  over  $H_d$   
598 (from section 4.4). If the bottles that did not yield any DNA were not included in the BN  
599 (mimicked by not instantiating those nodes) then the  $LR$  is 104 000. This slight decrease comes  
600 about from the fact that as knowledge of no DNA being found is taken away then the  
601 probability of the cells being invigorated into the air slightly increases compared to the  
602 probability of DNA presence given direct transfer and so the  $LR$  slightly decreases. While the  
603 same probability is assigned under both propositions, the decrease in  $LR$  comes from the fact  
604 that  $\Pr(E|H_p)$  is the same, but  $\Pr(E|H_d)$  has slightly increased.

605  
606 As iteratively more bottles are instantiated to include the defendant's DNA the increase in the  
607  $LR$  in support of  $H_p$  over  $H_d$  lessens. This again comes down partly to the effect described  
608 above (i.e. an increase in the invigoration mechanism), but as more bottle results are added the  
609 dominant defence proposition becomes an AO having a matching profile with the defendant.  
610 The effect can be seen in Figure 3, where from zero to five bottles are instantiated with result  
611 'D+U' and common unknown nodes are not instantiated. When the first bottle is instantiated  
612 the support for  $H_p$  over  $H_d$  is approximately 2.6 bans ( $LR$  expressed in  $\log_{10}$ ), whereas by the  
613 time the fifth bottle is instantiated virtually no additional support is gained, and the  $LR$  plateaus  
614 at the probability of the defendant and the AO having matching profiles.

615



616

617 *Figure 3: increase in LR in support for  $H_p$  over  $H_d$  when zero to five bottles are instantiated*  
 618 *with result 'D+U' and common unknown nodes are not instantiated. The total LR is shown in*  
 619 *black on the graph (and relates to the right-hand axis). The increase in the total  $\log_{10}(LR)$  with*  
 620 *the addition of each bottle is shown with grey bars (and relates to the left axis).*

621

622 This same effect of the chance matching profiles of the defendant and AO can be seen in the  
 623 LRs in Table 1 all being around the inverse of the match probabilities when all result are  
 624 instantiated to 'D'. The slight deviations from exactly 1 billion with the different instantiations  
 625 of background DNA on the hands of the defendant or AO come from the multiple possible  
 626 routes of matching background DNA within the BN.

627

628 In our BN construction we have used the general value of 1 in 1 billion for all profile match  
 629 probabilities (i.e. between D and AO, but also between unknowns and AO or D). In carrying  
 630 out this approximation it allows the use of a single class network for the profile matching.  
 631 However, we could consider different probabilities for each match, which was related to the  
 632 level of DNA profile information obtained. To do so the profile matching class network would  
 633 require an additional input node, which would have passed into it a probability of matching,  
 634 and which then could be set individually per matching type (or per matching type and per bottle  
 635 if the profile matching probabilities were passed down from the main BN). It depends on the  
 636 position and type of matching that is being considered but given higher probabilities of  
 637 matching profiles could lead to the LR plateauing at a lower point than shown in Figure 3.

638

639 Also note that when there is seemingly unrelated unknown DNA found on all bottles (i.e. only  
 640 unknown DNA with no indication of any matching unknowns) then the LR becomes quite small  
 641 in comparison to other scenarios in Table 1, only providing slight support for  $H_d$  over  $H_p$ ,  
 642 which largely comes from probabilities of non-transfer of the defendant's DNA.

643

644 **CONCLUSION:**

645

646 We have shown here an evaluation given activity level propositions of the forensic DNA  
 647 profiling results from five bottles, all of which have been treated in a very similar manner

648 within the framework of circumstances of the case. Our focus was not on the probability  
649 assignment (although this an important aspect of any evaluation), but rather the appropriate  
650 treatment of the DNA on these items, and particularly the importance of the correct treatment  
651 of unknown DNA. Given the sensitivity of modern DNA profiling techniques, and our ever-  
652 increasing knowledge of the prevalence of DNA on items, it is possible to deduce multiple  
653 pathways for a common unknown DNA profile to be present on multiple similar items. We  
654 have shown the importance of including the key pathways for common unknowns to exist  
655 under both prosecution and defence propositions. In our BN, if the presence of unknown  
656 background DNA on the hands of the defendant is not considered then the transition from  
657 unknowns that are not common, to unknowns that are common between bottles, gives rise to  
658 an unrealistically dramatic change in support for the propositions.

659  
660 In an extension to our BN we could consider DNA amounts rather than the presence or absence  
661 of DNA, and we could consider environmental unknown DNA that could have also been  
662 resuspended back into the air and these would produce an even more discriminating  $LR$  given  
663 the case circumstances and findings (as long as appropriate data existed to assist with  
664 probability assignment).

665  
666 In the motivating case the defence proposition of an air-DNA transfer is not very well supported  
667 by the observed results compared to the prosecution proposition of a direct contact (as  
668 illustrated by its description by an appeal judge as '*A highly far-fetched theory of innocent*  
669 *indirect DNA transfer*') and the evaluation of the case results provides very strong support for  
670 the evidence given  $H_p$  rather than given  $H_d$ , if the probabilities assigned are valid.

671 Even with much higher probabilities for the DNA swirling into the air, and then landing on  
672 bottles (set at some values that an analyst feels may represent the upper bound of a reasonable  
673 range, which intuition tells us must be lower than the probability of transfer from a direct  
674 contact), the evaluation will still favour  $H_p$  over  $H_d$ . The detailed probabilistic analysis  
675 developed in this paper confirms the broad intuitive assessment of both the reporting scientist  
676 and the court. It may lead some to conclude that such a complex analysis is not needed. We  
677 disagree with that view. The benefit of the above approach is to be able to actually number the  
678 probabilities of the observations given each of the allegations. It enables to qualify what is  
679 meant by '*highly far-fetched*' or '*being an unlikely way for DNA to transfer*'.

680

681

682 **Supplementary material:**

683 1. The OOBN from Figure 1

684 2. A description on the setup of each node and the population of the conditional  
685 probability tables with data

686

687

688

689

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769

770

771

772 **Supplementary: Probability assignments for BNs in Figure 1**

773 Main BN (Part A of Fig 1)

774

775 Node 1 (Hp/Hd): The proposition node with possible states of ‘Hp’ for prosecution or ‘Hd’ for  
 776 defence. The ‘Hp’ option is the prosecution scenario and ‘Hd’ considers the defence scenario  
 777 as given in section 4.0. These two options are assigned with equal prior probabilities. Note that  
 778 this does not mean the prior odds in this case are equal; equal prior probabilities are applied for  
 779 the propositions so that the values obtained by the BN inform the likelihood ratio for the  
 780 scientific evidence only.

**Hp/Hd**

Hp	0.5
Hd	0.5

781

782 Node 2 (D handled bottles): considers the activity of D placing the bottles in the ceiling space,  
 783 which has occurred under Hp and not under Hd.

784

**Hp/Hd**  
  
**D handled bottles**

	Hp	Hd
Yes	1	0
No	0	1

785

786 Node 3 (Police walked through crime scene): There is no dispute that this activity has occurred  
 787 and therefore 1 is assigned for state ‘yes’ under both Hp and Hd.

788

**Hp/Hd**  
  
**Police walked through crime scene**

	Hp	Hd
Yes	1	1
No	0	0

789

790 Node 4 (AO handled bottles): considers the activity of the AO placing the bottles in the ceiling  
 791 space, which has occurred under Hd and not under Hp.

792

**Hp/Hd**  
  
**D handled bottles**

	Hp	Hd
Yes	0	1
No	1	0

793

794 Node 5 (AO match D): This root node shows the probability that an unknown person will share  
 795 the same DNA profile and here we use the profile probability for the defendant of 1 in 1 billion.

796

**AO match D**

Match	1.0E-9
No match	1 – 1.0E-9

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Node 6 (Background DNA on D hands): Considers how often non-self DNA is found on the hands. For illustration of the BN performance we use values of 0.5 for ‘yes’ and ‘no’ but concede the presence of background DNA on hands has been shown to be more prevalent as demonstrated in (Szkuta et al. 2017).

**Background DNA on D hands**

Yes	0.5
No	0.5

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Node 7 (Cells reinvigorated): The probability that DNA relocation can occur via shed cells in the pathway of investigators processing a crime scene. We consider the possibility of cell relocation during evidence collection as one event, and allocate a probability of 1 in 1000. If investigators did not walk through the pathway, then cell relocation cannot occur, therefore 0 is assigned for state ‘yes’ and 1 is assigned for state ‘no’.

Police walked through crime scene

**Cells reinvigorated**

	Yes	No
Yes	0.001	0
No	0.999	1

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Node 8 (Background DNA on AO hands): The same treatment as non-self DNA on the hands of the defendant so states will have the same values as node 6.

**Background DNA on AO hands**

Yes	0.5
No	0.5

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Node 9 (BG on AO hands matches D): The same treatment as node 5. Although already accounting for the presence of unknown DNA from the AO, this node considers unknown DNA present as background on this unknown person’s hands.

**BG on AO hands matches D**

Match	1.0E-9
No match	1 – 1.0E-9

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Node 10-19 (B1 and B2 same U?, B1 and B3 same U? ..., B4 and B5 same U?): Considers the different sources of unknown DNA for bottle 1 and bottle 2. This accounts for whether the same unknown DNA that may be present from background DNA on the defendant’s hands, unknown DNA from the AO, background DNA on the AO’s hands, and background DNA on the bottles. The same reasoning then extends to the remaining nodes for all pairwise comparisons for each bottle.

Unknown_matcher		Yes	No
		1	0
<b>B1 and B2 same U?</b>	Yes	1	0
	No	0	1

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Node 20-24 (Bottle 1 DNA results ..., Bottle 5 DNA results): These nodes combine the parental node values to consider the different transfer mechanisms and sources of DNA for each bottle (example of the conditional probability table is shown here for bottle 1). Here we use presence or absence of DNA for the various sources and consider possibilities of the defendant's DNA (D), defendant and unknown DNA (D+U), unknown DNA only (U) or no DNA at all (None).

Transfer mechanisms and sources of DNA for bottle 1		D	D+U	U	None
		1	0	0	0
<b>Bottle 1 DNA results</b>	D	1	0	0	0
	D+U	0	1	0	0
	U	0	0	1	0
	None	0	0	0	1

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Bottle class BN (Part B of Fig 1)

Node 25 (Unknown background DNA from AO hands): If the AO handled the bottles (Y), with background DNA present on their hands (Y) and DNA had transferred to the bottle from handling (Y), then we expect the probability to find unknown background DNA to be quite high. We have assigned this value as 0.95 for state 'yes' and 0.05 for state 'no'.

Alternatively, if the AO handled the bottles with background DNA on the hands, however their DNA did not transfer to the bottle from handling (N), then we expect the probability of unknown DNA to also transfer to be quite low. The values assigned in this scenario are 0.05 for state 'yes' and 0.95 for state 'no'.

Finally, if the AO did not handle the bottle (N), or background DNA was not present on the hands (N), then unknown background DNA cannot be present via the hands. All values here are assigned 0 for state 'yes' and 1 for state 'no'.

AO handled bottles		Y				N			
		Y		N		Y		N	
Background DNA on AO hands		Y	N	Y	N	Y	N	Y	N
Y		0.95	0.05	0	0	0	0	0	0



<b>Unknown background DNA from AO hands</b>	N	0.05	0.95	1	1	1	1	1	1
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Node 26 (AO DNA transferred to bottle from handling): Considers the probability for DNA to transfer via a direct contact to the bottle surface. For illustration we use a rate of transfer of 0.5 so from handling the five bottles directly, we would expect to obtain a DNA profile from the handler two or three times. Again, if the bottles were not handled, then DNA cannot transfer and we assign 0 for state ‘yes’ and 1 for state ‘no’.

		AO handled bottles	
		Yes	No
AO DNA transferred to bottle from handling	Yes	0.5	0
	No	0.5	1

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Node 27 (Unknown background DNA from D hands): If the defendant handled the bottles (Y), with background DNA present on the defendant’s hands (Y) and DNA had transferred to the bottle from handling (Y), then we expect the probability to find unknown background DNA to be quite high. We have assigned this value as 0.95 for state ‘yes’ and 0.05 for state ‘no’.

Alternatively, if the defendant handled the bottles with background DNA on the hands, however the defendant’s DNA did not transfer to the bottle from handling (N), then we expect the probability of unknown DNA to also transfer to be quite low. The values assigned in this scenario are 0.05 for state ‘yes’ and 0.95 for state ‘no’.

Finally, if the defendant did not handle the bottle (N), or background DNA was not present on the hands (N), then unknown background DNA cannot be present via the hands. All values here are assigned 0 for state ‘yes’ and 1 for state ‘no’.

D handled bottles		Y				N			
		Y		N		Y		N	
Background DNA on D hands		Y		N		Y		N	
D DNA transferred to bottle from handling		Y	N	Y	N	Y	N	Y	N
Unknown background DNA from D hands		Y	0.95	0.05	0	0	0	0	0
		N	0.05	0.95	1	1	1	1	1

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Node 28 (D DNA transferred to bottle from handling): The same treatment as node 26, to inform the node on this DNA transfer mechanism we expect similar findings for the AO and the defendant. Indeed shedder status, time since hand washing and opportunity for DNA loading will all impact this value, and we do not have case specific information to inform us of the shedder status of the defendant. We are left guided by scientific literature alone.

		Yes	No
		0.5	0
<b>D DNA transferred to bottle from handling</b>	Yes	0.5	0
	No	0.5	1

880

881 Node 29 (D DNA settle on bottle once in air): We consider DNA to settle on each of the five  
 882 bottles as individual events, therefore we have the ‘cells reinvigorated’ node on the outer layer  
 883 of the BN as an input node, and the ‘D DNA settle on bottle once in air’ node present in the  
 884 class OOBN inner layer. We have assigned a probability of 1 in 100 for each bottle so state  
 885 ‘yes’ is 0.01 and state ‘no’ is 0.99. Again, if cells were not present in the air environment then  
 886 the defendant’s DNA cannot settle on a bottle (state ‘no’ = 1).

887

		Yes	No
		0.01	0
<b>D DNA settle on bottle once in air</b>	Yes	0.01	0
	No	0.99	1

888

889 Node 30 (D DNA contaminated bottle): Considers the probability of laboratory contamination  
 890 and we assign a probability of 1 in 1000 (state ‘yes’ = 0.001 and ‘no’ = 0.999)

891

<b>D DNA contaminated bottle</b>	Yes	0.01
	No	0.99

892

893 Node 31 (Unknown background DNA on bottle): Considers the prevalence of background  
 894 DNA on bottles and we assign a value of 0.5 for state ‘yes’ and 0.5 for state ‘no’.

895

<b>Unknown background DNA on bottle</b>	Yes	0.5
	No	0.5

896

897 Node 32 (BG on D hands matched AO): Considers the probability of background DNA on the  
 898 defendant’s hands to have same alleles as an unknown donor as we use the profile probability  
 899 of 1 in 1 billion.

900

<b>BG on D hands matched AO</b>	Match	1.0E-9
	No match	1 – 1.0E-9

901

902 Node 33-36 (BG match probability): This series of nodes considers whether background DNA  
 903 on the bottle is the same as the defendant, alternate offender, background DNA on the

904 defendant's hands, or background DNA on the AO's hands. Again we use the profile  
905 probability as with node 32.  
906

**BG match probability**

Match	1.0E-9
No match	1 – 1.0E-9

907  
908 Node 37-43 are accumulation nodes of the various sources of unknown DNA, or DNA from  
909 the Defendant, which may be present on the bottle.

910  
911 Node 44 (Recovery): Considers the ability to recover DNA from the outer surface of the bottle  
912 lids using a swabbing technique followed by DNA extraction. We consider the ability to obtain  
913 a DNA profile with efficiency of 0.8 for state 'yes' and 0.2 for state 'no'.  
914

**Recovery**

Yes	0.8
No	0.2

915  
916 Single-bottle unknown matcher class BN (Part E of Fig 1)

917  
918 Node 38 (BG1 = BG2): Considers whether background DNA observed on two bottles will  
919 share a common unknown donor. Again, we use the profile probability for whether the two  
920 sources of unknown DNA will share the same alleles.  
921

**BG1 = BG2**

Match	1.0E-9
No match	1 – 1.0E-9

922  
923 Node 46-49 are the accumulation of common sources of DNA that may be observed from  
924 background, and again we do not describe the probability assignments as data is not required  
925 to inform these nodes.  
926