1	Title: Defining end user requirements for a field-based molecular detection system for
2	wildlife forensic investigations.
3	Alice Masters ^a , Rob Ogden ^{b,c} , Jon H Wetton ^d , Nick Dawnay ^{a,*}
4	^a School of Pharmacy and Biomolecular Sciences, Liverpool John Moores University, Byrom
5	Street, Liverpool, L3 3AF, UK,
6	^b Royal School of Veterinary Studies and the Roslin Institute, University of Edinburgh,
7	Midlothian, EH25 9RG, UK.
8	°TRACE Wildlife Forensics Network, EH12 6LE, Edinburgh, UK
9	^d Department of Genetics & Genome Biology, University of Leicester, University Road,
10	Leicester, LE1 7RH, UK.
11	
12	* Corresponding author: Tel +44 151 231 2485.
13	E-mail address: n.m.dawnay@ljmu.ac.uk

15 Abstract

The increasing use of non-laboratory-based DNA and protein detection methods promise to 16 provide rapid investigative intelligence and support sample prioritisation. Primarily developed 17 18 for human forensic or medical applications, current systems may also show utility in the field 19 of wildlife forensic science. However, it is currently unknown whether the requirements of the wildlife forensic community can be met by current non-laboratory based tools. Given the 20 21 diverse array of stakeholders and sample types commonly encountered, it is necessary to first identify the needs of the community and then try and map their needs to current instrumentation. 22 By using a market research style questionnaire, this study identified key requirements for a non-23

laboratory-based system following feedback from the wildlife forensic community. Data 24 25 showed that there is strong support for field-based detection methods while highlighting concerns including contamination risks and reduced quality assurance associated with non-26 laboratory testing. Key species and applications were identified alongside hurdles to 27 implementation and adoption. Broadly, the requirements align with many of the developmental 28 drivers that have led to the rise of in-field portable detection instrumentation, specifically rapid 29 detection within one hour, ease-of-use, and $\geq 95\%$ accuracy. Several existing platforms exist that met some of the identified requirements but not all. With further collaboration between 31 industry partners and the wildlife forensic community it is possible that new field-based systems 32 33 can be developed and applied routinely.

34 Key words: Field-based testing; molecular; wildlife; forensic; industry; development

35

36 **<u>1. Introduction</u>**

37 The illegal wildlife trade (IWT) poses a huge threat to the survival of many species. The blackmarket trade in endangered species is estimated between US\$5 billion and US\$20 billion a year 38 and disrupting the trade requires a multi-faceted approach [1, 2]. Challenges in understanding 39 40 IWT include the covert and transnational nature of the trade [3], coupled with difficulties associated with discovering and then identifying illicit items by non-specialist regulatory 41 officers [4]. This is typically achieved using traditional investigative approaches, such as 42 intelligence-led international operations [5, 6], or through random searches of items at borders 43 [7]. Confirming the species identity of seized items, or determining whether or not they contain 44 45 derivatives of an endangered species, is then necessary to support a criminal prosecution [8]. However, given the heavily processed nature of many of the samples recovered, or the lack of 46 species distinguishing characters between immature specimens of many species, diagnostic 47

identification needs to be performed. Currently this is conducted by specialist laboratories with
expertise in morphological or molecular identification techniques [9-11], but the development
and future implementation of field-based analytical equipment may allow on-site-analysis
saving both time and investigative resources.

52 Portable rapid detection methods can detect either DNA or proteins unique to the sample of interest and be developed to match the end-user requirements depending on the field of 53 54 research. The potential for application in forensic science has long been recognised by the 55 human forensic community, where consultation with stakeholders has revealed a number of clearly defined end user requirements [12, 13]. These requirements have allowed industry 57 groups to develop and commercialise several DNA and immunoassay approaches [e.g. 14-17]. Such advancements now allow analysis of forensically relevant samples by police officers and 58 Crime Scene Investigators out of the laboratory. While a large proportion of this work has 59 focussed on human forensic applications, there is evidence that similar approaches may be useful in the wildlife forensic arena [18-20]. However, the application of such portable 61 62 approaches to wildlife forensics is likely to be complicated by the diverse array of sample types encountered in casework and the ability of any of the existing instrumentation to fulfil the 63 requirements of the end user. Furthermore, the timeframe for development, validation and 64 65 implementation of any approach in a wildlife forensic context is very difficult to predict given the diverse array of jurisdictions and the individual needs of specialist forensic groups. It is therefore possible that for the foreseeable future field-based approaches are restricted to 67 presumptive test applications, complimenting subsequent confirmatory analysis at a laboratory; 68 that said, it seems likely that wildlife forensic applications will reach the field at some point. 69

This study seeks to identify the key requirements of a field-based detection system as required by potential end users and wider stakeholder groups in the wildlife forensic and law enforcement arena. In doing so, the community's needs can either be mapped to identify a compatible instrument or the need for more bespoke instrumentation and support from industrydevelopers.

75

76 **2. Methodology**

An online questionnaire (supplemental material 1) was distributed using SurveyMonkey Inc (San Mateo, California, USA) to participants at the 2017 Society for Wildlife Forensic Science (SWFS) conference in Edinburgh and to postgraduate students studying at the Liverpool Centre for Advance Policing (LCAP). The survey was voluntary, anonymised and no personal information was collected. The research was granted ethical approval prior to being conducted (Approval Number 17/PBS/004).

83 In total, 100 individuals participated in the survey; 78 SWFS participants and 22 LCAP participants. Average completion rate of the questionnaire was 74%. Response data was 84 exported to Excel and weighted averages applied to all rank questions. Preliminary analysis 85 86 allowed the grouping of individuals into four broad categories based on their profession; laboratory-based researcher (n=27; consisting of university or government researchers), 87 **laboratory-based practitioner** (n=25; consisting of scientists employed to provide analytical 88 services, e.g. forensic caseworkers, food standards, conservation), field-based practitioner 89 (n=35; consisting of customs/border control, field-based wildlife crime investigators, 90 police/enforcement officers and postgraduate students in policing and criminal investigation) 91 and desk-based individuals (n=13; consisting of charity/NGO/policy representatives and R&D 92 project managers).

94

95 4. Results and Discussion

96 4.1. Stakeholder Awareness

The data shows a knowledge gap may exist between user groups regarding awareness of field-97 based DNA systems (Table 1A). The data shows that ~68% of field-based practitioners have 98 99 'some' or 'very little' knowledge of current field-based detection systems compared to ~50% of desk-based individuals who described themselves as being 'very familiar' or 'familiar'. A 100 similar proportion was also seen in the lab-based practitioner group, ~48% of whom also 101 102 identified as being 'very familiar' or 'familiar' while the most aware were the lab-based researchers, ~67% of whom were 'very familiar' or 'familiar' with current field based systems. 103 One possible explanation for the lack of familiarity observed in the field-based practitioner 104 105 group is that many of the field-based systems are only recently out of the R&D phase. As such, much of the information available has been disseminated through scientific publications with 106 little targeted knowledge transfer to field-based end-users. Similar knowledge gaps have been 107 reported between the enforcement and research communities with other technology [21, 22], 108 and has been cited as a reason for the slow adoption of pioneering research by enforcement 109 110 groups.

111	Table 1. Ranking of the issues preventing the use of field-based instrumentation in wildlife forensic casework and participant's level of familiarity
112	with current, field-based DNA instruments.

Topic under evaluation and response options	Field-ba Practitic		Lab-ba Practit		Desk- Individ		Lab-ba Resea		All Partici	pants
A) Participants level of familiarity with field instrumentation	Percent	(%)			-		-		-	
1) Very familiar with current, field-based DNA instruments	0.0		20.0		16.7		25.0		12.6	
2) Familiar with some platforms	14.7		28.0		33.3		41.7		26.3	
3) Some literature-based knowledge	35.3		40.0		25.0		8.3		30.5	
4) Very little known	32.4		8.0		20.8		16.7		21.1	
5) Not previously aware of field-based DNA instrumentation	17.6		4.0		4.2		8.3		9.5	
B) Issues preventing the use of field-based instrumentation	Weighted average of the scores (rank)									
1) Cost	2.15	(1)	1.46	(1)	0.80	(1)	1.66	(1)	6.07	(1)
2) Lack of funding for purchasing	1.89	(2)	1.26	(4)	0.73	(2)	1.35	(2)	5.23	(3)
3) Accuracy of the test and instrument	1.80	(3)	1.44	(2)	0.68	(3)	1.35	(2)	5.28	(2)
4) Sensitivity of the test and instrument	1.54	(6)	1.46	(1)	0.65	(4)	1.24	(3)	4.89	(4)
5) Lack of an instrument that suits my needs	1.77	(4)	1.30	(3)	0.41	(7)	1.20	(4)	4.68	(5)
6) Lack of an assay that I can use	1.28	(7)	1.18	(5)	0.44	(6)	1.12	(5)	4.02	(7)
7) Ease of use	1.59	(5)	1.12	(6)	0.51	(5)	0.89	(6)	4.11	(6)
8) The colour of the instrumentation	0.72	(8)	0.43	(7)	0.16	(8)	0.40	(7)	1.71	(8)

A - Results are the calculated percentage of participants (%) based on the number of responders. Number of responders to question was 34 for field-based practitioner, 25 for lab-based practitioner, 12 for desk-based individual, 24 for lab-based researcher.

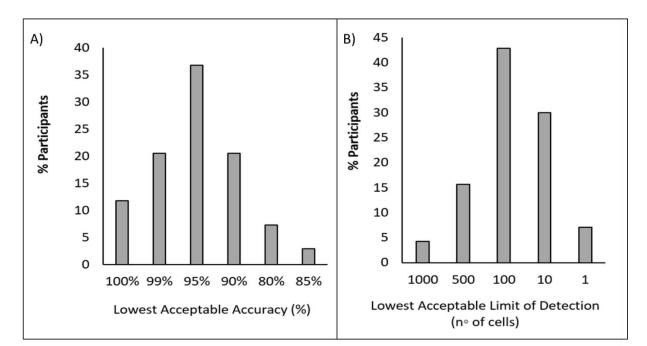
B - Results were ranked using a weighted average of the scores (1-8) entered by participants giving more importance to the issues selected first. Number of responders to question was 30 for field-based practitioner, 22 for lab-based practitioner, 10 for desk-based individual, 21 for lab-based researcher.

4.2. Perceived issues regarding the adoption of field-based instrumentation

Participants from all groups selected 'cost' as the primary issue preventing the adoption of field-115 based instrumentation (Table 1B). Regarding the maximum per sample cost of analysis the data 116 117 reveals that 2% of participants would consent to paying £100 per sample; 14% would pay £50 per sample; 37% would pay £20; 32% would pay £10; while 14% identify £1 as the maximum 118 per sample analysis cost. Together the data suggests that a consumable cost of £20 per sample 119 120 will satisfy 53% of users. With regards to maximum instrumentation cost, the data shows that none of the participants would pay £100,000 for a field-based detection system; 3% would pay 121 £50,000; 16% would pay £10,000; 41% would pay £5,000; 26% would pay £1,000; while only 122 123 14% of are looking for instrumentation that costs £100. Together the data suggests that an instrumentation cost of £5,000 per unit will satisfy 60% of surveyed users. Results indicate that 124 assay and instrument cost are key issues for commercial development groups to consider if they 125 want to expand into the wildlife forensic marketplace. Data also shows that the funding needed 126 to purchase such instrumentation would be secured from a variety of different sources; 42% 127 128 from government grants; 27% from academic funding bodies; 15% from internal institutional 129 based funding calls; and 15% from NGO or charity funding. The emphasis on central government financing suggests there may be a need for specific funding to facilitate the 130 adoption of field-based instrumentation. 131

The second overall hurdle to implementing field-based testing as a strategy was the instrument and test 'accuracy' (Table 1B). Analysis shows that 67% of respondents would be satisfied with a test accuracy of 95%, while only 33% of participants require a test with 99-100% accuracy (Figure 1a). Test accuracy is a measure of the agreement between the 'information' obtained from the sample under evaluation and a controlled standard or voucher specimen. The type of 'information' provided will depend on the purpose of the test (see section 4.4 below), although diagnostic accuracy can be expressed in many ways, including '*Sensitivity*' and '*Specificity*'

[23, 24]. Under this definition, the number of True Positives, False Positives, True Negatives 139 and False Negatives are recorded. These numbers are used to report on the test Sensitivity (the 140 proportion of true positives that are correctly identified by the test) and Specificity (the 141 proportion of true negatives that are correctly identified by the test) with very accurate tests 142 show a high percentage scores for both. The number of false negatives recorded can vary as a 143 function of the system's Limit of Detection (LOD) and reduce the overall measure of accuracy. 144 The data in Figure 1b shows that 4% of participants suggest an LOD of \leq 1000 cells; 16% 145 suggest LOD of \leq 500 cells; 43% suggest an LOD of 100 cells; while 37% suggest an LOD of 146 <10 cells. Together, the data shows that 63% of respondents consider detection of \leq 100 cells 147 148 an acceptable LOD. This is largely in line with the limit of detection displayed by human 149 forensic tests. One explanation for the different requirements is that each stakeholder group likely process different types of biological sample, ranging from DNA rich items such as tissue 150 and blood to extremely low concertation samples such as powdered derivatives or trace 151 material. 152



153

Figure 1. Test Accuracy (a) was identified as a hurdle to implementing field-based systems together with the test Limit of Detection (b). Number of responders to question was 26 for fieldbased practitioner, 14 for lab-based practitioner, 12 for desk-based individual, and 18 for labbased researcher.

Portable rapid detection tests are typically described as either being 'presumptive' or 158 159 '*diagnostic*'. Presumptive tests will produce a higher number false positive and false negative results and are therefore less accurate than diagnostic tests used in the laboratory [12, 25]. There 160 is no strict classification of what is required to classify a test as being either presumptive or 161 diagnostic based on its accuracy although the data suggests that there is room for the 162 development of presumptive tests with 95% accuracy at 100 cells input which may include 163 164 affordable and easy-to-use immunoassay-based approaches [e.g. 18] as well as more sensitive and specific DNA based approaches [e.g. 19, 20]. 165

Other highly ranked issues included increasing contamination events as PCR moves out of the 166 167 laboratory and a reduction in QA/QC as processes become field-based (Figure 2). These represent serious concerns to the adoption of field-based testing even when using tests with a 168 high reported accuracy as the QA/QC practices of a testing laboratory may differ markedly 169 from the processes employed at a crime scene, in the field, or at boarders. However, it should 170 be recognised that the necessity to adopt ISO17025 standards during sample collection is not 171 172 unique to wildlife forensic investigations as both Crime Scene Investigators and Sexual Assault Referral Centre Staff handling human casework samples have only in the last few years begun 173 adopting and defining sector specific standards [26]. As such, it is considered likely that the 174 175 wildlife community follow suit and that training and knowledge transfer events be organised in preparation for the adoption of field-based testing supported by community working groups, 176 government regulators and special interest groups. Such training will need to also look at the 177 validation of the novel technology prior to use in forensic investigations. The validation process 178 is universally recognised by laboratory analysists and validation guidelines and 179 180 recommendations are available [26, 27]. However, with field-based technology the responsibility for validation will fall on the shoulders of enforcement teams who may have little 181 experience in this area. 182

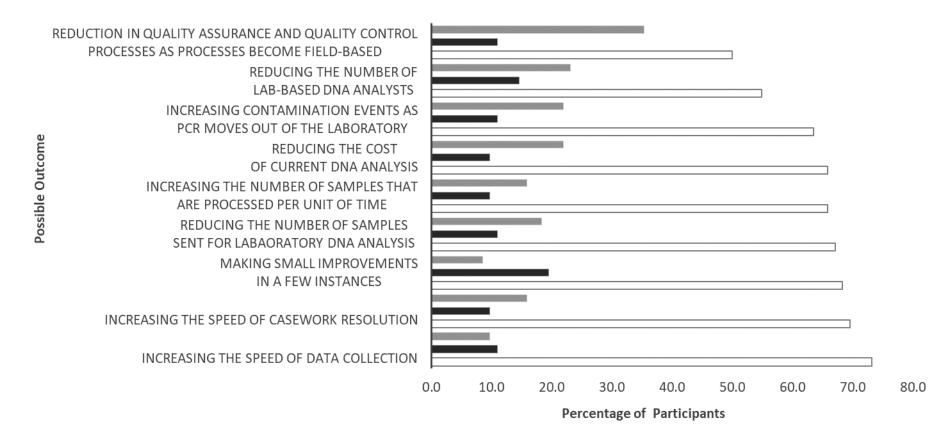


Figure 2. Possible outcomes when adopting field-based DNA instrumentation in the field. Respondents answered either 'likely' (white bars); 'impartial' (black bars); or 'unlikely (grey bars) when asked about each of the possible outcomes. Number of responders to question was 30 for field-based practitioner, 22 for lab-based practitioner, 9 for desk-based individual, and 20 for lab-based researcher.

190 *4.3. Perceived benefits regarding the adoption of field-based instrumentation*

Analysis shows that participants believe that once introduced, the impact of the field-based 191 intervention would be positive (Table 3). Ranking of possible outcomes by participants shows 192 193 that increasing the speed of data collection, increasing the speed of casework resolution, and increasing the number of samples processed per unit of time were identified as 'likely' 194 outcomes (Figure 2). When asked 'how long should it take to prepare a sample for analysis?' 195 196 36% of participants selected \leq 30 minutes; 21% selected \leq 10 minutes; 40% selected \leq 5 minutes; and 3% selected ≤ 1 minute. As such 97% of potential users would be satisfied if 197 sample preparation time was within 5 minutes. When asked 'how long should it take to generate 198 199 usable and understandable data?' 8% of participants selected ≤ three hours; 29% selected 60-90 minutes; 36% identified 30-60 minutes; and 27% selected less than 30 minutes. As such the 200 data suggest that 73% of participants would be happy with a test that runs within 1 hour. 201

Typically, developers have increased the speed of current processes by integrating sample 202 203 purification (DNA extraction) and sample amplification (PCR) steps [19, 28, 29]. This has often been in due to the high demands in law enforcement to analyse more DNA samples faster at 204 less expense to increase the speed of casework resolution [13, 30, 31]. One mechanism explored 205 in human forensic analysis is the idea of using field-based testing for sample triage at the crime 206 scene which can bring objectivity to evidential assessment and can reduce the number of 207 samples sent for analysis prior to obtaining a result [32]. Such benefits may also be translated 208 to practitioners of wildlife forensic casework which remains expensive due to the cost related 209 to the development of in-house protocols and the low sample throughput which raises the cost 210 of analysis per sample. A development target for commercial groups has been to perform DNA 211 analysis in under an hour from the point of sample collection. The data presented here supports 212 this as a developmental goal. 213

Another key developmental driver has been on ease-of-use. When asked 'what level of user 214 215 expertise should field-based instrumentation be aimed at?' 28% of participants selected 'DNA aware CSI'; 22% selected 'Forensic Aware Enforcement Officer'; 22% selected 'anyone with 216 5 minutes training'; 16% selected 'Good DNA knowledge'; and 12% selected 'DNA expert' 217 (Table 3). Interestingly, it was the desk-based and the field-based practitioner groups who 218 selected 'DNA Expert' as an acceptable descriptor of an end user in contrast to the lab-based 219 220 practitioners and lab-based researchers who did not select this descriptor at all. Overall, the data suggests that there is a clear expectation that the instrumentation should be aimed at non-221 laboratory-based individuals. Ease of use also relates to data interpretation. When asked 'what 222 223 features of the analysis and software are required' 21% of participants selected 'graduated percentage confidence in the result' and 19% selected 'software-based interpretation'. Such 224 functionality would make it extremely easy for field practitioners, especially as the percentage 225 226 match result is already provided through existing sequence similarity searches. Interestingly, 14% of the participant's selected 'expert based interpretation' suggesting that there is a desire 227 for some further verification of the result by another individual. Also, 12% selected 'binary 228 yes/no answer'; 12% selected 'probabilistic result' and 11% selected 'raw accessible data'. 229 Interestingly, only 10% selected 'weighted and phrased for use in forensic casework' which 230 231 suggests that participants currently see little need for the analysis software to format the data ready for submission as evidence. This may be due to the existing reliance on forensic 232 laboratory staff to present data in court and unwillingness by the community to automate the 233 interpretation process. However, it should be noted that such automation has already been 234 partially achieved with DNA data in the form of the STRmixTM expert software [33] and 235 validation guidelines exist to support software developers [34]. 236

Study Question	Response	Desk-based Individual	Lab-based Researcher	Lab-based Practitioner	Field-based Practitioner	Total Average (%)
		Percent (%)				
() Impact of	Positive Effect	80.0	85.0	86.4	87.0	85.6
	No Effect	0.0	15.0	9.1	6.5	8.4
A) Impact of intervention B) Expertise descriptors for possible end-users C) Location for field- based deployment	Negative Effect	20.0	0.0	4.5	6.5	6.0
	DNA Expert	25.0	0.0	0.0	22.2	11.8
B) Expertise	Good DNA Knowledge	8.3	27.8	21.4	7.4	16.2
descriptors for	DNA Aware CSI	16.7	33.3	28.6	33.3	28.0
possible end-users	Forensic Aware Enforcement	16.7	16.7	28.6	25.9	22.0
	Anyone	33.3	22.2	21.4	11.2	22.0
		Weighted Pe	ercent (%)			
	Offices	0.0	10.5	20.0	13.3	13.0
C) Location for field	Customs and Border Stations	38.5	39.5	37.1	28.9	35.1
	Vehicles	7.7	18.4	17.1	17.8	16.8
based deployment	Field Sheltered	53.8	18.4	22.9	24.4	25.2
	Field Unsheltered	0.0	13.2	2.9	15.6	9.9
	Software based interpretation	18.5	16.0	20.0	22.2	19.3
	Graduated % confidence in result	22.2	16.0	26.7	20.4	20.5
D) Features of analysis	Expert based interpretation	18.5	12.0	13.3	14.8	14.3
and interpretation	Binary Yes/No Answer	7.4	12.0	10.0	16.7	12.4
desired	Probabilistic	7.4	14.0	13.3	13.0	12.4
	Raw data accessible	11.1	16.0	10.0	7.4	11.2
	Weighted and phrased for use in casework	14.8	14.0	6.7	5.6	9.9

Table 3. Participant groups response to impact of intervention, end-user expertise, and optimal location for deployment.

Results are the percentage based on the number of responders. Number of responders to question A) was 31 for field-based practitioner, 22 for lab-based practitioner, 10 for desk-based individual, 20 for lab-based researcher. Number of responders to question B) was 27 for field-based practitioner, 14 for lab-based practitioner, 12 for desk-based individual, 18 for lab-based researcher. Number of responders to question C) was 27 for field-based practitioner, 14 for lab-based practitioner, 12 for desk-based individual, 18 for lab-based researcher. Number of responders to question D) was 26 for field-based practitioner, 14 for lab-based practitioner, 14 for lab-based practitioner, 14 for lab-based practitioner, 12 for desk-based individual, 18 for lab-based researcher. Number of responders to question D) was 26 for field-based practitioner, 14 for lab-based practitioner, 12 for desk-based individual, 18 for lab-based researcher.

With regards to the most suitable location for field-based testing, 35% of participant's selected 239 240 'customs and boarder stations'; 25% selected 'field sheltered'; 17% selected 'vehicles'; 13% selected 'offices'; and 10% selected 'field unsheltered'. This represents a possible division in 241 relation to what an instrument is expected to do. It is likely that customs, border posts and 242 offices have electric power supplies which would allow the use of any instrumentation that 243 requires power, including larger desk-based instrumentation. Field stations may require a 244 generator or require battery powered instrumentation or utilise methods that require no power 245 source for analysis such as lateral flow and immunoassay-based devices. 246

When polled on 'how many samples would be run each week using field-based instrumentation' 247 248 70% of total participants stated they would analyse at least five samples a week (Figure 3) with the greatest usage identified in the field-based practitioner group. Usage was identified in other 249 groups also, although it is difficult to assess whether this represents a true need or whether 250 participants were responding from the point of view of a field practitioner. It is likely that usage 251 will vary between different applications and jurisdictions so further insight may be required as 252 253 specific species of interest and enforcement groups are identified who may become early adopters of field-based analysis. 254

The data reveals that respondents broadly favour the adoption of field-based, office-based or 255 non-laboratory-based instrumentation. Furthermore, there is support for deployment at borders 256 and ports suggesting that detection of trafficked items is the preferred application. With regard 257 to data interpretation (Table 3) the two most common requests, representing almost 40% of 258 respondents, was for 'software based interpretation' with a 'graduated % confidence in the 259 result', directly relating to accuracy or percentage similarity akin to DNA sequence similarity 260 searching [35]. This would suggest that the greatest proportion of individuals would like 261 minimal hands on data analysis with fewer individuals wanting access to the raw data. 262

Frequency of sample analysis 40 Field-based Practitioner 35 Lab-based Practitioner 30 % Participants Desk-based 25 Lab-based Researcher 20 15 10 5 0 1 2 5 10 25 50 100 Number of samples run per week

Figure 3. Cumulative total showing the number of participants (%) that would run at least 1,
2, 5, 10, 25, 50 or 100 samples per week using field-based DNA analysis. Number of responders
to question was 27 for field-based practitioner, 14 for lab-based practitioner, 12 for desk-based
individual, and 18 for lab-based researcher.

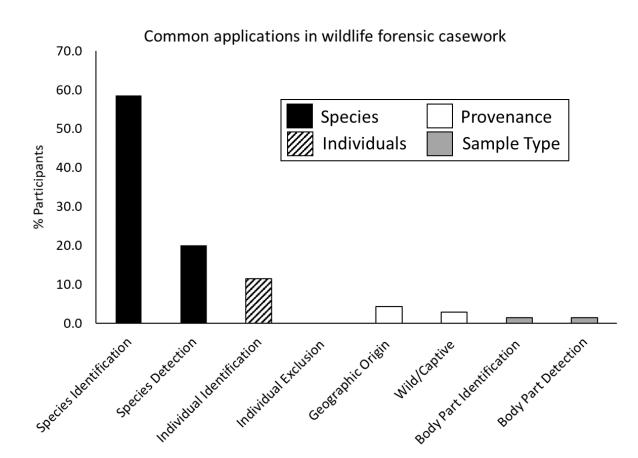
268

269 4.4. Investigative Questions, Species and Sample Types

Beyond legal casework to directly support prosecutions, wildlife forensic science includes a 270 range of stakeholders working in areas of academic research, trade monitoring, supply chain 271 verification and intelligence gathering. The development of a single solution to field-based 272 testing is therefore complicated by the different species, objectives and priorities in play. Our 273 results show that determining species identity is currently the most common form of analysis 274 performed (Figure 4). For this type of analysis forensic providers match the DNA sequence of 275 the unknown sample to 'known' DNA sequences stored on open-access databases [36, 37]. 276 However, even this common approach suffers from limitations as the databases are unregulated, 277 278 leading to uncertainty in the result, and are sometimes not populated with data from the species of interest. To combat this problem, the wildlife forensic community are developing the ForCyt 279 DNA database [38], a fully-regulated database of species that are commonly encountered in 280 forensic investigations. Such a database would make the development of a field-test more 281

achievable, but may still require different design strategies depending on whether the question
is one of identification (what species is it?) or detection (is it Tiger?). Typically, molecular tests
are developed to detect a specific analyte, addressing the closed form of the question [i.e. 3942]. When an open identification question is asked, the emphasis shifts toward building a test
capable of identifying every single species of interest and consequently becomes more difficult.
The preference to ask open questions often severely limits what a laboratory can do and
investigators are often asked to be more specific with their request.

289



290

Figure 4. Common applications in wildlife forensic casework. Number of responders to question was 25 for field-based practitioner, 20 for lab-based practitioner, 8 for desk-based individual, and 19 for lab-based researcher.

Analysis pertaining to individual identification and determination of geographic origin or wild/captive assessment are less commonly required because the tests are expensive to develop,

niche in application and often require de-novo collection of appropriate population reference databases [43-45]. The least common question is to identify whether a sample belongs to a specific part of an animal. Given that determining species identity through open or closed questions is required in the majority of instances, it is seems sensible that industry groups develop approaches that seek to address this type of question.

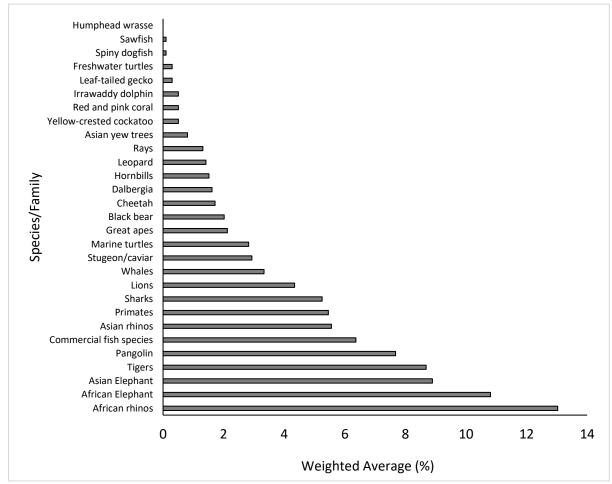
When asked to identify 'which group of fauna/flora is most often encountered', 63% of 302 participant's selected 'mammal'; 13% selected 'birds'; 9% selected 'fish'; 7% selected 'invertebrates'; 4% selected 'timber' 1% selected 'reptiles'; and 4% selected 'various'. With 304 regards to the 'type' of sample commonly encountered, 35% of participants selected 'meat/body parts/organs'; 21% selected 'bones/teeth/scales'; 16% selected 'live animal'; 15% selected 306 'skins/pelts/furs/wools'; 14% selected 'liquid mixtures'; 14% selected 'whole dead animals'; 307 selected 'powdered derivatives'; 10% selected 'horns/ivory'; and 7% selected 11% 'pods/seeds'. The range of sample types highlights a problem for developers of field-based molecular approaches for wildlife forensic applications. Developing a detection platform that 310 311 works across an entire range of samples types is difficult, and in some instances has limited the 312 use of non-lab based systems to a single form of analysis, such as individual identification, on a single sample type, such as buccal swabs [e.g. 15]. Other systems have also recommended 313 additional expertise and time spent on pre-processing steps [e.g. 46] to allow complete analysis. 314 Indeed, the challenging samples encountered by forensic scientists continue to be the focus of 315 development for laboratory processing, let alone field-based applications [47]. 316

To further understand taxon importance, participants were asked to rank a list of flora and fauna (pre-identified by the authors as forensically relevant) and thereby identify which is most likely to benefit from a field-based detection system. The weighted data shows the top five groups identified are African rhinos, African elephant, Asian elephant, tiger and pangolin (Figure 5). The identification of four flagship taxa and pangolin requires some discussion. The main

forensically relevant samples collected from elephants is ivory, which is often unprocessed and 322 323 exported to Asia where it is in high demand, especially in China [48]. While it can be readily identified morphologically, ivory from African and Asian elephants can be mixed making it 324 difficult to distinguish between the two groups without laboratory testing. Forgers are also 325 becoming increasingly adept at creating fake ivory pieces from bone, teeth, horn, plastic and 326 resin which are sold as real ivory [49, 50]. Furthermore, the use of population assignment 327 328 approaches has been used to identify poaching hotspots [51] suggesting that population assignment may be the primary application when considering field-based molecular approaches 329 for ivory. Pangolin scales can also be identified morphologically, although not to sub-species 330 331 level. As with elephants, the broad distributions of certain pangolin species from both Asia and Africa, may suggest that the primary application of any test is to differentiate between 332 geographic regions to support investigations and identify poaching hotspots. It should be noted 333 that any test capable of population assignment in these species will also simultaneously perform 334 species identification which remains an important consideration. Forensically relevant samples 335 of the two remaining top-ranked species can include tiger bone and rhino horn, both of which 336 can be ground up into powders for inclusion in Traditional Asian Medicines (TAM) [52]. The 337 lack of any identifying characteristics when handling these processed samples suggests that a 338 339 simple field-test for species identification would support investigations that involve the analysis of TAM products.

The inclusion of commercial fish species as the sixth most likely to benefit from field-based testing suggests that the illegal fishing, landing and species substitution of high value species with low value species also represents a clear development goal as species detection and verification is something that is required throughout the food chain [53, 54]. To understand and develop an assay for commercial fish species further research is needed with regard to species prioritisation. Research has listed demersal fish, salmon, trout and smelt as having the highest levels of illegal fishing [55] but it remains difficult to identify a single species to target with 54% of the stock/species categorised as being at high risk of illegal, unregulated and unreported fishing [56]. Indeed, it is likely that the development of field-based testing in fisheries and food supply chains will be prioritised over methods developed for critically endangered species, as fish identification represents a larger end-user market and has an immediate relevance to human health and food safety. It is also considered likely that the development of a system that works on fish species will be easier to apply, given the samples commonly encountered include single source, DNA rich, tissue and muscle.

355



356

Figure 5. Ranked species in order of most likely to benefit from a field-based DNA profiling
system. Each participant was asked to rank what they thought were the top 5 species. Results
were ranked using a weighted average giving more importance to the species selected first.
Number of responders to question was 21 for field-based practitioner, 21 for lab-based
practitioner, 8 for desk-based individual, and 20 for lab-based researcher.

363 **<u>5. Summary</u>**

This questionnaire has identified a need for non-laboratory detection applications in wildlife forensic science. The results highlight a series of end user expectations and concerns that industry groups and developers can address either through mapping requirements to existing systems or developing entirely bespoke assays or instruments. The key elements identified are broadly in alignment with the expectations placed on human-based detection platforms:

369

• Results within one hour from the start of sample processing

- Easy to use tool with simplified data interpretation
- 371

• 95% accuracy of identification

At this moment in time there are a number of systems that are close to fulfilling some of the 372 requirements outlined by this research but no assay or instrument currently fulfils all the 373 requirements. Instruments of note include, the Oxford Nanopore Technologies MinION [57-374 59], a highly portable system that meets cost requirements and can be used in the generation of 375 data suitable for species identification. It meets limit of detection requirements, but the current 376 377 end-users require a high level of experience at sample preparation and result interpretation 378 although simple disposable consumables and software are under development to address these limitations. The ParaDNA system [60, 61] has shown potential as a forensic screening system 379 and has been developed specifically for end users with no laboratory experience. Data interpretation is by automated software which requires no expertise to interpret. Accuracy is 381 high but the system is only within the budget of a small portion of the participants of this 382 questionnaire. Furthermore, it only runs pre-developed assays which may reduce the likelihood that a wildlife assay can be used in conjunction with the system without collaboration from the 384 industry developers. Immunoassays [62-64] are low cost, easy to use and suitable for field and indoor conditions. However, issues exist regarding sensitivity and specificity and they do not always work with degraded samples. Typically, molecular detection tests with low cost show
questionable accuracy. However, it is important to recognise that such tests have an important
function in forensic casework as presumptive tests.

Both presumptive and diagnostic tests have utility in an investigative framework but there is yet to be a test that combines low cost with high accuracy. Further research looking at 391 mechanisms to achieve this are ongoing [65, 66] but are likely 5-10 years away from being 392 commonly used. As such, if non-laboratory-based detection systems are to be utilised in the interim period it is likely to be done on an ad-hoc basis with each end user group identifying 394 the system that specifically suits their needs and collaborating with industry developers to 395 396 understand ways in which it can improved to better suit their purpose. A likely stepping stone towards true field-based tools is the early adoption of some of these technologies within forensic 397 laboratories in low and middle income countries which currently lack relatively expensive 398 genetic analysis instrumentation and are the sources of many of the species involved in the illegal wildlife trade. Adoption of cheaper and faster tests will significantly enhance regional 400 401 enforcement action by initially building capacity within such dedicated facilities whilst the 402 developments required for deployment outside of a laboratory are validated. Finally, it is essential that community groups help develop a series of guidelines for the field-based 403 404 validation of detection systems that can be readily used by enforcement groups and nonlaboratory trained individuals. In doing so, many of the concerns identified during this study 405 will addressed in preparation for the widespread adoption of future field-based analysis 406 systems. 407

408

409 Acknowledgements

The authors would like to thank all those who participated in the questionnaire. Funding for this
research was provided by the Peoples Trust in Endangered Species (PTES) Internship funding
scheme.

413

414 **References**

- [1] Rosen GE, Smith KF. Summarizing the evidence on the international trade in illegal
 wildlife. *EcoHealth.* 7 (2010) 24-32.
- [2] Wyler LS, Sheikh PA. International illegal trade in wildlife: Threats and US policy. Library

418 of Congress Washington DC Congressional Research Service.

- [3] Warchol GL. The transnational illegal wildlife trade. *Criminal Justice Studies*. 17 (2004)
 57-73.
- [4] Sollund R, Maher J. The Illegal wildlife trade: A case study report on the illegal wildlife
 trade in the United Kingdom, Norway, Colombia and Brazil. A study compiled as part of the
 EFFACE project. (2015). *http://efface. eu/sites/default/files/EFFACE_Illegal% 20Wildlife%*20Trade_revised. pdf. Accessed on 11.2.2018.
- 425 [5] Europol. Illicit trafficking in endangered animal species.
 426 https://www.europol.europa.eu/crime-areas-and-trends/crime-areas/environmental-
- 427 *crime/illicit-trafficking-in-endangered-animal-species*. Accessed on 6.6.2018.
- 428 [6] Interpol. Environmental Compliance and Enforcement Committee (ECEC) and Working
- 429 Groups. https://www.interpol.int/Crime-areas/Environmental-crime/Committee-and-Working-
- 430 *Groups/Wildlife-Crime-Working-Group*. Accessed on 6.6.2018.
- [7] Eastern and Southern Africa Anti-Money Laundering Group. A Special Typologies Project
 Report on Poaching, Illegal Trade in Wildlife and Wildlife Products and Associated Money

433 Laundering in the ESAAMLG Region. http://esaamlg.org/reports/TYPOLOGIES-REPORT-

434 ON-THE-WILDLIFE-CRIMES-AND-RELATED-ML..pdf Accessed on 6.6.2018.

- [8] Johnson R, Wilson-Wilde L, Linacre A. Current and future directions of DNA in wildlife
- 436 forensic science. *Forensic Science International: Genetics.* **10** (2014) 1-11.
- 437 [9] Ogden R. Forensic science, genetics and wildlife biology: getting the right mix for a wildlife
- 438 DNA forensics lab. *Forensic Science, Medicine, and Pathology.* **6** (2010) 172-179.
- [10] Ogden R, Dawnay N, McEwing R. Wildlife DNA forensics—bridging the gap between
- 440 conservation genetics and law enforcement. *Endangered Species Research.* **2** (2009) 179-95.
- [11] Dawnay N, Ogden R, McEwing R, Carvalho GR, Thorpe RS. Validation of the barcoding
- gene COI for use in forensic genetic species identification. *Forensic Science International*. 173
 (2007) 1-6.
- [12] Morrison J, Watts G, Hobbs G, Dawnay N. Field-Based Detection of Biological Samples
- for Forensic Analysis: Established Techniques, Novel Tools, and Future Innovations. *Forensic*
- 446 *Science International.* **285** (2018) 147-160.
- [13] Mennell J, Shaw I. The future of forensic and crime scene science. Part I. A UK forensic
 science user and provider perspective. *Forensic Science International*. **157** (2006) S7-S12.
- [14] Dawnay N, Stafford-Allen B, Moore D, Blackman S, Rendell P, Hanson EK, Ballantyne
- 450 J, Kallifatidis B, Mendel J, Mills DK, Nagy R. Developmental Validation of the ParaDNA®
- 451 Screening System-A presumptive test for the detection of DNA on forensic evidence items.
- 452 Forensic Science International: Genetics. **11** (2014) 73-9.
- 453 [15] Salceda S, Barican A, Buscaino J, Goldman B, Klevenberg J, Kuhn M, Lehto D, Lin F,
- 454 Nguyen P, Park C. Pearson F. Validation of a rapid DNA process with the RapidHIT® ID

455 system using GlobalFiler® Express chemistry, a platform optimized for decentralized testing
456 environments. *Forensic Science International: Genetics*. 28 (2017) 21-34.

[16] Old J.B, Schweers B.A, Boonlayangoor P.W, Reich K.A. Developmental Validation of
RSIDTM-Saliva: A Lateral Flow Immunochromatographic Strip Test for the Forensic Detection
of Saliva. *Journal of forensic sciences*. 54 (2009) 866-873.

- [17] Old J, Schweers B.A, Boonlayangoor P.W, Fischer B, Miller K.W, Reich K.
 Developmental Validation of RSIDTM-Semen: A Lateral Flow Immunochromatographic Strip
 Test for the Forensic Detection of Human Semen. *Journal of forensic sciences*. 57 (2012) 489463 499.
- [18] Peppin L, McEwing R, Webster S, Rogers A, Nicholls D, Ogden R. Development of a
 field test for the detection of illegal bear products. *Endangered Species Research.* 9 (2008) 263270.
- [19] Dawnay N, Hughes R, Syndercombe-Court D, Duxbury N. Species detection using
 HyBeacon ® probe technology: Working towards rapid onsite testing in non-human forensic
 and food authentication applications. *Forensic Science International: Genetics.* 20 (2016) 103111.
- [20] Pomerantz A, Penafiel N, Arteaga A, Bustamante L, Pichardo F, Coloma LA, BarrioAmoros CL, Salazar-Valenzuela D, Prost S. Real-time DNA barcoding in a remote rainforest
 using nanopore sequencing. *bioRxiv*. (2017) 189159.
- [21] Weisburd D, Neyroud P. Police science: Toward a new paradigm. *Australasian Policing*.
 5 (2013) 13-21.
- [22] Rojek J, Alpert G, Smith H. The utilization of research by the police. *Police Practice and Research.* 13 (2012) 329-341.

- 478 [23] Bossuyt P.M, Reitsma J.B, Bruns D.E, Gatsonis C.A, Glasziou P.P, Irwig L.M, Lijmer J.G,
- 479 Moher D, Rennie D, de Vet H.C. Standards for Reporting of Diagnostic Accuracy. Towards
- complete and accurate reporting of studies of diagnostic accuracy: the STARD initiative. *British Medical Journal.* 326 (2003) 41-44.
- [24] Altman DG, Bland JM. Diagnostic tests. 1: Sensitivity and specificity. *British Medical Journal.* 308 (1994) 1552.
- [25] Virkler K, Lednev I. Analysis of body fluids for forensic purposes: From laboratory testing
 to non-destructive rapid confirmatory identification at a crime scene. *Forensic Science International.* 188 (2009) 1-17.
- [26] Forensic Science Regulator (2017). Codes of Practice and Conduct for forensic science 487 providers and practitioners in the Criminal Justice System. 488 Issue 4. https://assets.publishing.service.gov.uk/government/uploads/system/uploads/attachment_data/ 489 file/651966/100 - 2017 10 09 - The Codes of Practice and Conduct -490
- 491 <u>Issue_4_final_web_web_pdf_2_.pdf</u>. Accessed on 18.4.2019.
- 492 [27] SWGDAM Validation Guidelines for DNA Analysis Methods.
 493 <u>https://docs.wixstatic.com/ugd/4344b0_813b241e8944497e99b9c45b163b76bd.pdf</u>. Accessed
 494 on 19.2.2019.
- [28] Liu P, Scherer JR, Greenspoon SA, Chiesl TN, Mathies RA. Integrated sample cleanup
 and capillary array electrophoresis microchip for forensic short tandem repeat analysis. *Forensic Science International: Genetics.* 5 (2011) 484–492.
- 498 [29] Hird HJ, Brown MK. Design, optimisation and preliminary validation of a human specific
- 499 loop-mediated amplification assay for the rapid detection of human DNA at forensic crime
- scenes. *Science and Justice*. **57** (2017) 409-414.

- [30] Butler JM. The future of forensic DNA analysis. *Phil. Trans. R. Soc. B.* **370** (2015)
 20140252.
- 503 [31] Gold S. RapidDNA: a game changer in the law enforcement identification
 504 stakes. *Biometric Technology Today*. *Jan* (2013) 7-10.
- 505 [32] Mapes AA, Kloosterman AD, de Poot CJ, van Marion V. Objective data on DNA success
- rates can aid the selection process of crime samples for analysis by rapid mobile DNA
 technologies. *Forensic Science International.* 264 (2016) 28-33.
- 508 [33] Bright JA, Taylor D, McGovern C, Cooper S, Russell L, Abarno D, Buckleton J.
- 509 Developmental validation of STRmix[™], expert software for the interpretation of forensic DNA
- 510 profiles. *Forensic Science International: Genetics*. **23** (2016) 226-39.
- 511 [34] OSAC Best Practice Recommendation for Validation of Forensic DNA Software.
- 512 https://www.nist.gov/sites/default/files/documents/2018/07/17/best_practice_recommendation
- 513 <u>for validation of forensic dna_software.pdf</u>. Accessed on 19.2.2019.
- [35] Altschul SF, Gish W, Miller W, Myers EW, Lipman DJ. Basic local alignment search tool.
- *Journal of molecular biology.* **215** (1990) 403-410.
- [36] Sujeevan R, Hebert PA. BOLD: the Barcode of Life Data System. *Molecular Ecology Notes.* 7 (2007) 355-364.
- 518 [37] Pruitt KD, Tatusova T, Maglott DR. NCBI reference sequences (RefSeq): a curated non-
- redundant sequence database of genomes, transcripts and proteins. *Nucleic acids research*. 35
 (2006) D61-65.
- 521 [38] Ahlers N, Creecy J, Frankham G, Johnson RN, Kotze A, Linacre A, McEwing R, Mwale
- 522 M, Rovie-Ryan JJ, Sitam F, Webster LM. 'ForCyt'DNA database of wildlife species. Forensic
- 523 Science International: Genetics Supplement Series. 6 (2017) e466-8.

524	[39] Aabo S, Rasmussen OF, Roseen L, Sørensen PD, Olsen JE. Salmonella identification by
525	the polymerase chain reaction. <i>Molecular and cellular probes</i> . 7 (1993) 171-178.
526	[40] Cavrois M, de Noronha C, Greene WC. A sensitive and specific enzyme-based assay
527	detecting HIV-1 virion fusion in primary T lymphocytes. Nature biotechnology. 20 (2002)
528	1151.
529	[41] Old J, Schweers BA, Boonlayangoor PW, Fischer B, Miller KW, Reich K. Developmental
530	Validation of RSID TM -Semen: A Lateral Flow Immunochromatographic Strip Test for the
531	Forensic Detection of Human Semen. Journal of Forensic Sciences. 57 (2012) 489-499.
532	[42] Karabasanavar NS, Singh SP, Kumar D, Shebannavar SN. Detection of pork adulteration
533	by highly-specific PCR assay of mitochondrial D-loop. Food chemistry. 145 (2014) 530-534.
534	[43] Manel S, Berthier P, Luikart G. Detecting wildlife poaching: identifying the origin of
535	individuals with Bayesian assignment tests and multilocus genotypes. Conservation Biology.
536	16 (2002) 650-659.
537	[44] Wasser SK, Mailand C, Booth R, Mutayoba B, Kisamo E, Clark B. Stephens M. Using
538	DNA to track the origin of the largest ivory seizure since the 1989 trade ban. Proc. Natl. Acad.
539	<i>Sci. U. S. A.</i> 104 (2007) 4228–4233
540	[45] Ogden R, Linacre A. Wildlife forensic science: a review of genetic geographic origin
541	assignment. Forensic Science International: Genetics. 18 (2015) 152-159.
542	[46] Lu H, Giordano F, Ning Z. Oxford Nanopore MinION sequencing and genome assembly.
543	Genomics, Proteomics & Bioinformatics. 14 (2016) 265-279.
544	[47] Kraemer M, Prochnow A, Bussmann M, Schere, M, Peist R, Steffen, C. Developmental
545	validation of QIAGEN Investigator® 24plex QS Kit and Investigator® 24plex GO! Kit: Two

- 6-dye multiplex assays for the extended CODIS core loci. *Forensic Science International: Genetics.* 29 (2017) 9-20.
- [48] Stiles, D. China faces a conservation challenge: the expanding elephant and mammoth
- 549 ivory trade in Beijing and Shanghai. *Pachyderm.* **56** (2015) 122-126.
- 550 [49] Santiapillai C, Silva A, Karyawasam C, Esufali S, Jayaniththi S, Basnayake M, Unantenne
- 551 V, Wijeyamohan S. Trade in Asian elephant ivory in Sri Lanka. Oryx. 33 (1999) 176.
- [50] Buddhachat K, Brown J, Thitaram C, Klinhom S, Nganvongpanit K. Distinguishing real
 from fake ivory products by elemental analyses: A Bayesian hybrid classification
 method. *Forensic Science International.* 272 (2017) 142-149.
- [51] Wasser SK, Clark WJ, Drori O, Kisamo ES, Mailand C, Mutayoba B, Stephens M.
 Combating the illegal trade in African elephant ivory with DNA forensics. Conservation
 Biology. 22 (2008) 1065-1071.
- [52] Ellis R. Tiger bone & rhino horn: the destruction of wildlife for traditional Chinese
 medicine. Island Press (2013) 11-27.
- 560 [53] Rasmussen R, Morrissey M. Application of DNA-Based Methods to Identify Fish and
- Seafood Substitution on the Commercial Market. *Comprehensive Reviews in Food Science and Food Safety.* 8 (2009) 118-154.
- [54] Helyar S, Lloyd H, de Bruyn M, Leake J, Bennett N, Carvalho G. Fish Product
 Mislabelling: Failings of Traceability in the Production Chain and Implications for Illegal,
 Unreported and Unregulated (IUU) Fishing. *PLoS ONE*. 9 (2014) p.e98691.
- 566 [55] Agnew D, Pearce J, Pramod G, Peatman T, Watson R, Beddington J, Pitcher T. Estimating
- the Worldwide Extent of Illegal Fishing. *PLoS ONE*. **4** (2009) e4570.

- 568 [56] Freitas B. Illegal Fishing: Which fish species are at highest risk from illegal and 569 unreported fishing? (2015). Washington DC, United States, 1-95. Available at:
- 570 https://c402277.ssl.cfl.rackcdn.com/publications/834/files/original/Fish_Species_at_Highest
- 571 _*Risk_from_IUU_Fishing_WWF_FINAL.pdf?1446130921.* Accessed on 25.2.2018.
- 572 [57] Juul S, Izquierdo F, Hurst A, Dai X, Wright A, Kulesha E, Pettett R, Turner DJ. What's in
- 573 my pot? Real-time species identification on the MinION. *bioRxiv*. **Jan** (2015) 1:030742.
- [58] Lindberg MR, Schmedes SE, Hewitt FC, Haas JL, Ternus KL, Kadavy DR, Budowle B.
 A comparison and integration of MiSeq and MinION platforms for sequencing single source
 and mixed mitochondrial genomes. *PloS ONE*. 9 (2016) e0167600.
- 577 [59] Benítez-Páez A, Portune KJ, Sanz Y. Species-level resolution of 16S rRNA gene
 578 amplicons sequenced through the MinION[™] portable nanopore sequencer. *GigaScience*. 5
- 579 (2016).
- [60] Blackman S, Dawnay N, Ball G, Stafford-Allen B, Tribble N, Rendell P, Neary K, Hanson
- EK, Ballantyne J, Kallifatidis B, Mendel J. Developmental validation of the ParaDNA®
 Intelligence System—A novel approach to DNA profiling. *Forensic Science International: Genetics.* 17 (2015) 137-48.
- [61] Dawnay N, Flamson R, Hall MJ, Steadman DW. Impact of sample degradation and
 inhibition on field-based DNA identification of human remains. *Forensic Science International: Genetics.* 37 (2018) 46-53.
- [62] Hsieh YH, Woodward BB, Ho SH. Detection of species substitution in raw and cooked
 meats using immunoassays. *Journal of Food Protection*. 58 (1995) 555-559.
- [63] Asensio L, González I, García T, Martín R. Determination of food authenticity by enzyme-
- ⁵⁹⁰ linked immunosorbent assay (ELISA). *Food control.* **19** (2008) 1-8.

- [64] Ayaz Y, Ayaz ND, Erol I. Detection of species in meat and meat products using Enzyme-
- Linked Immunosorbent Assay. *Journal of Muscle Foods*. **17** (2006) 214-20.
- [65] Pardee K, Green AA, Ferrante T, Cameron DE, DaleyKeyser A, Yin P, Collins JJ. Paper
- 594 based synthetic gene networks. *Cell.* **159** (2014) 940-954.
- 595 [66] Gootenberg JS, Abudayyeh OO, Kellner MJ, Joung J, Collins JJ, Zhang F. Multiplexed
- and portable nucleic acid detection platform with Cas13, Cas12a, and Csm6. *Science*. 360
 (2018) 439-444.

612 Supplementary Material 1

.			
Q1		h of the following roles most closely matches your current position	Tick
	A	Lab-based, forensic case worker	
	В	Field-based, wildlife crime investigator	
	C	Lab scientist - other (food standards, conservation etc)	
	D	R&D Project Manager (Academic/Industry)	
	E	Customs/Boarder Control	
	F	Police/Enforcement Officer	
	G	Charity/NGO/Policy Representative	
	Н	University researcher	
	1	Other (please state)	
	_		
Q2	Of the wi	ldlife samples you work with, what percentage requires some form of laboratory based DNA analysis?	Tick
QZ	A	None	TICK
	В	<20%	
	C	20-40%	
	D	40-60%	
	E	60-80%	
	F	100%	4
	P	100%	<u> </u>
Q3	How fam	liar are you with current, field-based, DNA instruments?	Tick
45	A	Very familiar with current technology and approaches	TICK
	B	Familiar with some platforms	
	C	Some literature based knowledge	
	D	Very little known	
	E	Not previously aware of field-based DNA instrumentation	
	L	Not previously aware of netu-based DNA instrumentation	
	Rank eac	h of the following issues (1-8) regarding how they prevent the use of current field-based instrumentation in wildlife	Tick
Q4	-	u can't rank them equally and you have to use all values 1-8 in your selection.	TICK
	A A	Cost	
	В	The colour of the instrumentation	
	C	Ease of use	
	D	Lack of funding for purchasing	
	F	Accuracy of the test and instrument	
	F	Sensitivity of the test and instrument	
	G	Lack of an instrument that suites my needs	
	н	Lack of an assay that I can use	
Q5	How helr	J ful would field-based DNA instrumentation be in your current work?	Tick
4.5	A	Extremely Useful	TICK
	B	Useful	
	ь С	Slightly useful	
	D	No effect	
	E	Slightly unhelpful	
	F	Unhelpful	
	F G	Extremely unhelpful	<u> </u>
<u> </u>	Score ear		
Q6		nely likely, 2= Very likely, 3= Likely, 4= Neither likely or unlikely, 5= Unlikely, 6= Very unlikely, 7= Extremely unlikely	Tick
<u> </u>	A	Reducing the cost of current DNA analysis	1
	B	Reducing the number of samples sent for laboratory DNA analysis	1
	C	Increasing the speed of data collection	
	C D		
	E	Increasing the speed of casework resolution	
	F	Increasing the number of samples that are processed per unit of time	
		Reducing the number of lab-based DNA analysts	<u> </u>
	G	Making small improvements in a few instances	
1	н	Reduction in quality assurance and quality control processes as processes become field-based	
		Increasing contamination events as PCR moves out of the laboratory	

Q7	Tick th	e wildlife group that you most commonly encounter in your role	Tick
	А	Reptiles	
	В	Mammals	
	С	Birds	
	D	Fish	
	E	Invertebrates	
	F	Amphibians	
	G	Timber	
Q8	Assign	a percentage score (0-100%) to each of the following sample descriptions based on how often you come across these	Tick
	A	powdered derivatives	İ
	В	Live animals	
	С	meat/body parts/organs	
	D	whole dead animals	
	E	pods/seeds	
	F	skins/pelts/furs/wools	
	G	horns/ivory	
	Н	liquid mixtures	
	1	bones/teeth/scales	
Q9	Rank t	he following forensic casework questions (1 most common - 8 least common) in terms of which is the most often asked in	Tick
	A	What species is it?	
	В	Is it species XXXX?	
	С	Can you identify the individual animal who left the sample using a DNA database or match probability calculation?	
	D	Can you exclude individual XXXX as the animal who left the sample?	
	E	Where did the animal come from?	1
	F	Did the animal come from the wild?	
	G	What part of the species does the sample come from?	1
	Н	Does the sample come from the XXXX part of the animal?	

Q10	Rank the following species (1-5) in order of most likely to benefit from a field based DNA profiling system							
	Asian Elephant	Primates	Hump head Wrasse					
	African Elephant	Pangolin	Sawfish					
	Asian rhinos	Leaf-tailed Gecko	Red and pink coral					
	African Rhinos	Hornbills	Spiny dogfish					
	Lions	Yellow-Crested Cockatoo	Sturgeon/caviar					
	Tigers	Whales	Commercial Fish Species					
	Leopard	Irrawaddy Dolphin	Asian Yew Trees					
	Cheetah	Freshwater turtles	Dalbergia					
	Black bear	Marine Turtles						
	Great Apes	Sharks						

Q11	If you had to select a single species to prioritise developing a field based DNA assay for, what species would it be and why?
	Answer

Q12	What love	el of user expertise should field-based DNA instrumentation be aimed at?	Tick
QIZ			TICK
	A B	DNA Expert Good knowledge of DNA approaches	
	C	DNA aware Forensic Investigators Forensic Aware Enforcement Officers	
	D F		
	E	Anyone with 5 minutes training	
Q13	Whore do	veu see field besed instrumentation being deployed?	Tick
QIS		you see field-based instrumentation being deployed? Offices	ПСК
	A		
	B	Customs and border stations Vehicles	
	C		
	D E	Field sheltered Field unsheltered	
	E		
014	Herriane	ah an Iditatele at menero o comple for explusions field beaudinate metation?	Tiele
Q14		should it take to prepare a sample for analysis on field based instrumentation?	Tick
	A	1 minute	
	В	5 minutes	
	C	10 minutes	
L	D	Within 30 minutes	Ļ
Q15		of samples should the instrument and test work on?	Tick
	A	Blood	
	В	Powdered derivatives	
	C	meat/body parts/organs	
	D	horns/ivory	
	E	liquid mixtures	
	F	bones/teeth/scales	
	G	Degraded samples	
	Н	Samples mixed with environmental contaminants (e.g. soil/fauna)	
Q16	How long	should it take to generate useable and understandable data from the time you collect the sample?	Tick
	A	<30 minutes	
	В	30-60 minutes	
	С	60-90 minutes	
	D	3 hrs	
Q17	How accu	rate does the test need to be?	Tick
	A	80% Accurate	
	В	85% Accurate	
	С	90% Accurate	
	D	95% Accurate	
	E	99% Accurate	
	F	100% Accurate	
	How sens	itive does the test need to be (how much biological material does it need to detect)?	
Q18	NOTE: Mo	st Current laboratory DNA tests can routinely detect between 10 and 100 cellular copies of nuclear DNA, less if mtDNA	Tick
	is being u	sed	
	A	Single cell or 6.6pg DNA	1
	В	10 cells or 66pg DNA	1
	С	100 cells or 660pg DNA	
	D	500 cells or 3.3ng DNA	İ
	E	1000 cells or 6.6ng DNA	İ
Q19	What feat	ures of the analysis and interpretation are required?	Tick
	A	Software based interpretation	
	В	Expert based interpretation	<u> </u>
	C	Binary Yes/No Answer	
	D	Graduated % confidence in result	
	E	Probabilistic	
	F	Raw data accessible	
	F G		
	9	Appropriately weighted and phrased for use in forensic casework	

Q20	What	s the maximum you would pay for a single field-based DNA instrument?	Tick
420	Δ	£100	
	В	£1,000	
	C	£5,000	
	D	£10,000	
	E	£50,000	
	с с	£100,000	
	'	1100,000	'
Q21	Of the	wildlife samples you work with, what percentage would you consider using field-based DNA analysis methods on?	Tick
	А	None	
	В	<20%	
	С	20-40%	
	D	40-60%	
	E	60-80%	
	F	100%	
Q22		s the maximum you would pay for a set of reagents to perform your wildlife test	Tick
	A	£1 Per Sample	
	В	£10 Per Sample	
	С	£20 Per Sample	
	D	£50 Per sample	
	E	£100 Per sample	
	F	£200 Per Sample	
	How li	kely are you to buy a field based DNA instrument if it performed according to your requirements and was within your	
Q23	budge		Tick
	А	Very Likely	
	В	Likely	
	С	Unlikely	
	D	Very unlikely	
Q24	How n	nany samples would you run per week?	Tick
	А	1	
	В	2	
	С	5	
	D	10	
	E	25	1
	F	50	I
	G	100	
Q25		rould you secure funds to purchase field-based DNA instrumentation	Tick
	A	Government Grants	ļ
	В	Research Funding Bodies	
	С	Internal Institutional Based Funding Calls	
	D	NGO/Charity Funding	