



## CSI Wildlife: Using Genetics to Hunt Elephant Poachers

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Student Worksheet

### INTRODUCTION

Forensic scientists collect and analyze scientific evidence to solve crimes. One type of evidence they use is genetic data. In this activity, you will use DNA analysis to solve several crimes related to elephant conservation, a field of science known as wildlife forensics. This worksheet complements the Click & Learn [CSI Wildlife](#).

### MATERIALS

- access to the Click & Learn [CSI Wildlife](#)
- four-function calculator

### PROCEDURE

Congratulations! You have just been hired by an international police force to work as a forensic scientist dedicated to protecting wildlife and investigating illegal activity where protected animals are involved. In particular, you will focus on the protection of African elephants. You are excited to put your scientific skills to work. To help you learn more about what you will do in your new position, your boss asks you to complete HHMI BioInteractive's Click & Learn [CSI Wildlife](#). But before you begin, she asks you to answer the following questions.

1. Elephants are a keystone species and play a pivotal role in shaping the forests and savannas in which they live. Knowing the importance of elephants, an international group of scientists conducted a census to estimate the number of African elephants. The Great Elephant Census data were released in August 2016, and unfortunately, the results reveal that African elephant populations have decreased by 30% in just the past seven years. Why do you think the number of elephants is declining?
  
2. You have likely heard of investigators using DNA fingerprinting to identify individuals in crime scene investigations. Briefly describe your current understanding of the process of DNA fingerprinting.

**Part 1**

Now that you have shared with your boss what you already know, she is ready to send you for training. Complete the activities in **Case One** of the *CSI Wildlife* Click & Learn.

- Your colleagues in the legal department are preparing to take a case to trial. To help them, rewrite your description of the process of DNA fingerprinting so a jury could understand it.

Now you are ready for your first major assignment. Your colleagues are investigating a newly seized tusk and wonder if the tusk is from a recently discovered crime where 10 elephants from the same population were found dead with their tusks removed. The genotype of the seized tusk for five short tandem repeat (STR) loci is shown in Table 1. Allele names for each of the STR loci correspond to their size in base pairs (bp). Table 2 lists the genotypes of the same five STR loci for 10 recently slaughtered elephants.

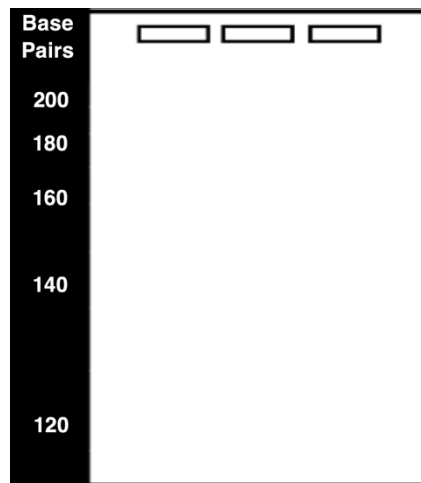
Seized tusk	FH19	FH60	FH67	FH71	FH129
	193	147	97	62	152
	193	147	103	62	160

**Table 1.** Genotype for five STR loci recovered from a seized elephant tusk.

	FH19	FH60	FH67	FH71	FH129
Elephant 1	189	147	97	62	152
	193	147	97	64	154
Elephant 2	185	147	95	62	152
	193	147	103	62	152
Elephant 3	183	147	95	62	152
	185	147	97	64	162
Elephant 4	193	147	91	62	152
	193	147	105	64	152
Elephant 5	187	145	95	62	152
	187	147	95	64	160
Elephant 6	187	145	91	64	152
	193	147	97	64	160
Elephant 7	193	149	97	60	156
	193	151	97	62	160
Elephant 8	193	147	97	62	152
	193	147	103	62	160
Elephant 9	193	147	97	62	160
	193	147	97	62	160
Elephant 10	189	147	95	62	152
	193	147	105	62	156

**Table 2.** Genotypes using five STR loci for 10 slaughtered elephants.

4. Did the seized tusk come from one of the recently slaughtered elephants? \_\_\_\_\_ What evidence supports your finding?
5. Draw the expected pattern of bands on a gel that shows the alleles for the seized tusk and the elephant you identified as a potential match for STR loci FH60 and FH129. Include a DNA ladder that has DNA fragments that are 125, 150, and 175 bp in size.



6. Your next challenge is to calculate the probability that another elephant from the same population would have the same genotype as the one from the seized ivory sample.
- a. Calculate the frequency of each of the alleles genotyped in the ivory sample using data from the 10 slaughtered elephants. Start by examining the data in Table 2 for STR locus FH19. The ivory sample was homozygous for the 193-bp allele. In the sample of 10 elephants, 12 alleles out of a total of 20 were also 193 bp in size. Look at how these data are used to determine the frequency of the allele in Table 3. Use the same approach to calculate the frequency of the alleles for the other four STR loci and record your answers in the table. Some of the data is provided for you.

STR locus	FH19	FH60	FH67	FH71	FH129
# of copies of Allele 1 in the group of 10 elephants studied	12	16	9		
Total number of alleles in the group of 10 elephants studied	20			20	
Frequency of Allele 1	$\frac{12}{20}$ = <b>0.60</b>				
Number of copies of Allele 2 in the group of 10 elephants studied	12		2		
Total number of alleles in the group of 10 elephants studied	20		20		
Frequency of Allele 2	$\frac{12}{20}$ = <b>0.60</b>				

**Table 3.** Frequency of alleles for five different STR loci in the group of slaughtered elephants.

- b. Use the formulas below to calculate the probability of another individual having the same genotype for each individual locus. Enter the formulas and probability in Table 4.
- If the ivory sample is **homozygous** for an STR locus, the probability ( $P_{(\text{genotype for locus } X)}$ ) that another individual has the same genotype for the STR locus is:

$$P_{(\text{genotype for locus } X)} = (\text{frequency of Allele 1})^2$$

- If the ivory sample is **heterozygous** for an STR locus, the probability ( $P_{(\text{genotype for locus } X)}$ ) that another individual has the same genotype for the STR locus is:

$$P_{(\text{genotype for locus } X)} = 2 \times (\text{frequency of Allele 1}) \times (\text{frequency of Allele 2})$$

STR locus	FH19	FH60	FH67	FH71	FH129
Formula with filled-in values for the probability	(0.60) <sup>2</sup>				
Calculated probability for the locus	0.36				

**Table 4.** The probability that another elephant in the same population has the same genotype at each individual STR locus, based on the allele frequencies in Table 3.

- c. To determine the probability that another individual has the same genotype for each of the five STR loci (in other words, the same genetic fingerprint), multiply all the individual probabilities together. What is the probability that all five match? \_\_\_\_\_
7. Impressed with your work, your boss now asks you to apply what you have learned to the genotype for the ivory sample for all 16 STR loci, listed in Table 5 below. The first five STR loci are the same as the ones listed in Table 4. In addition, your colleague has been able to collect additional samples of elephants from the same population as the ones shown in Table 2. He now has allele frequency data based on 80 alleles, shown in the table below.

STR locus	FH19	FH60	FH67	FH71	FH129	FH39	FH40	FH48
Allele 1 size (bp)	193	147	97	62	152	239	242	170
Allele 1 frequency	0.28	0.66	0.36	0.51	0.39	0.31	0.68	0.05
Allele 2 size (bp)	193	147	103	62	160	243	242	172
Allele 2 frequency	0.28	0.66	0.10	0.51	0.25	0.41	0.68	0.38
STR locus	FH94	FH102	FH103	FH126	FH127	FH153	S03	S04
Allele 1 size (bp)	225	177	149	94	247	169	141	154
Allele 1 frequency	0.65	0.64	0.13	0.18	0.04	0.011	0.55	0.8
Allele 2 size (bp)	225	181	151	110	247	171	143	154
Allele 2 frequency	0.65	0.14	0.61	0.11	0.04	0.08	0.39	0.8

**Table 5.** Allele size and frequency for 16 STR loci from a sample of 40 elephants (80 alleles).

- a. Compare the frequency of the alleles for STR loci FH19, FH60, FH67, FH71, and FH129 that you completed in Table 3 with the values in Table 5. Why do you think the values differ?
- b. Calculate the probability that another individual has the same genotype at all 16 STR loci by using the same procedure as Questions 6b and 6c. What is the probability of an identical match with another elephant at all 16 STRs? \_\_\_\_\_

**Part 2**

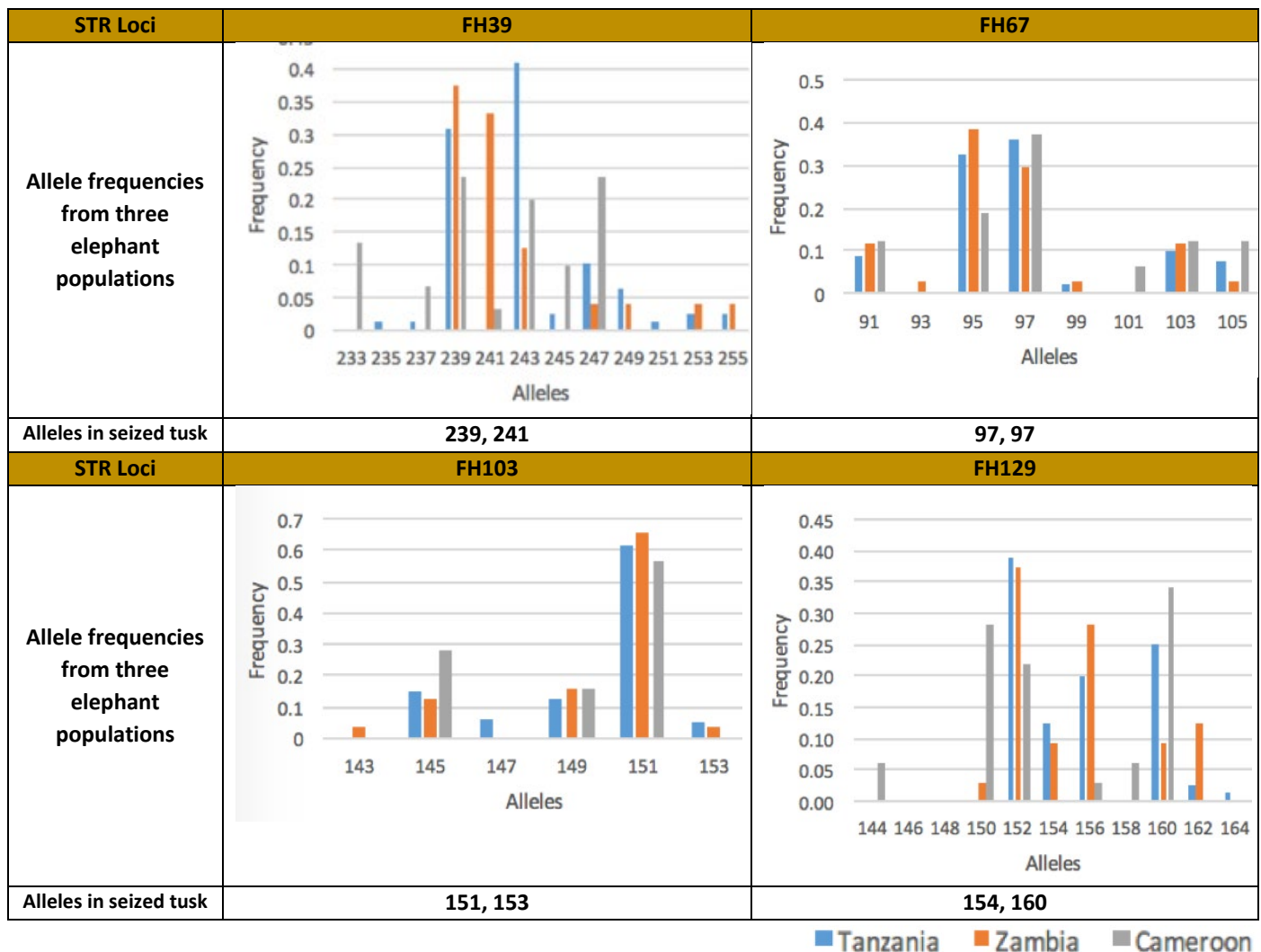
Your colleagues can tell that you have the necessary skills to help stop the killing of elephants for ivory and to bring poachers to justice. Your boss promotes you to lead a new case. To solve this case, you first need to complete **Case Two** of the *CSI Wildlife* Click & Learn.

**The details of the case:** Another shipment of ivory has been seized. Law-enforcement officials captured one of the people shipping the ivory. Through interviews with this individual and his contacts, the police determine the timing of the poaching event, but they still do not know the location. At the time the ivory was collected, three suspects were each caught poaching elephants in parks in three different parts of Africa. Suspect 1 was poaching in Tanzania, Suspect 2 in Cameroon, and Suspect 3 in Zambia.

FH39	FH67	FH103	FH129
239	97	151	154
241	97	153	160

**Table 6.** The genotype for four STR loci of a seized tusk.

Your boss would like you to link the genotype of one of the seized tusks to a reference population taken from one of the three countries. You determine the genotype of the seized tusk for four STR loci, shown in Table 6. Your lab assistant summarizes the allele frequency data for the three possible source populations in Figure 1 below.

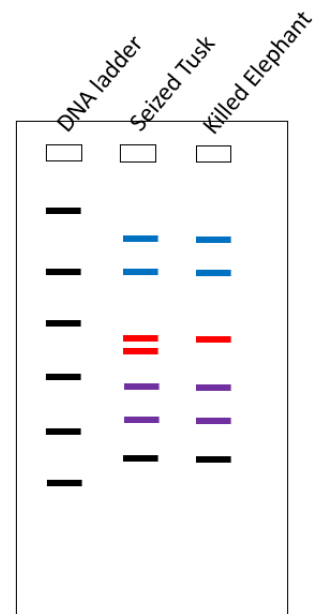


**Figure 1.** Comparison of the alleles from the ivory tusk to those of the reference populations in Tanzania, Zambia, and Cameroon.

- Make a claim about which country in Africa is most likely to be near the source population of the seized tusks. Your claim will be used in court, so make sure to highlight the evidence you used to make your claim.

**Part 3**

Late one night, rangers in a park hear a gunshot and quickly move to investigate. One elephant has been slain, but the quick actions of the rangers make the poacher flee, which saves the lives of many elephants. Eventually, a suspect is captured with one pair of elephant tusks. Your boss gives you the tusk and a blood sample from the killed elephant. She asks you to determine whether the tusks came from the elephant that was killed. You analyze the samples and the results are shown in Figure 1. Each STR locus is shown in a different color.



**Figure 2.** The gel with the alleles for four different STR loci for the elephant tusk and the killed elephant.

- Use the evidence in Figure 2 to determine whether the tusk came from the slain elephant. Write a brief summary of your findings for the case file.
- It is very important for scientists to consider sources of error in an experiment and alternative explanations. What sources of error or alternative explanations might change the answer you gave in Question 9?
- After completing this activity, do you believe that DNA fingerprinting is useful as evidence in criminal cases and can help in reducing elephant poaching? Provide three pieces of evidence from this experience to support your answer.