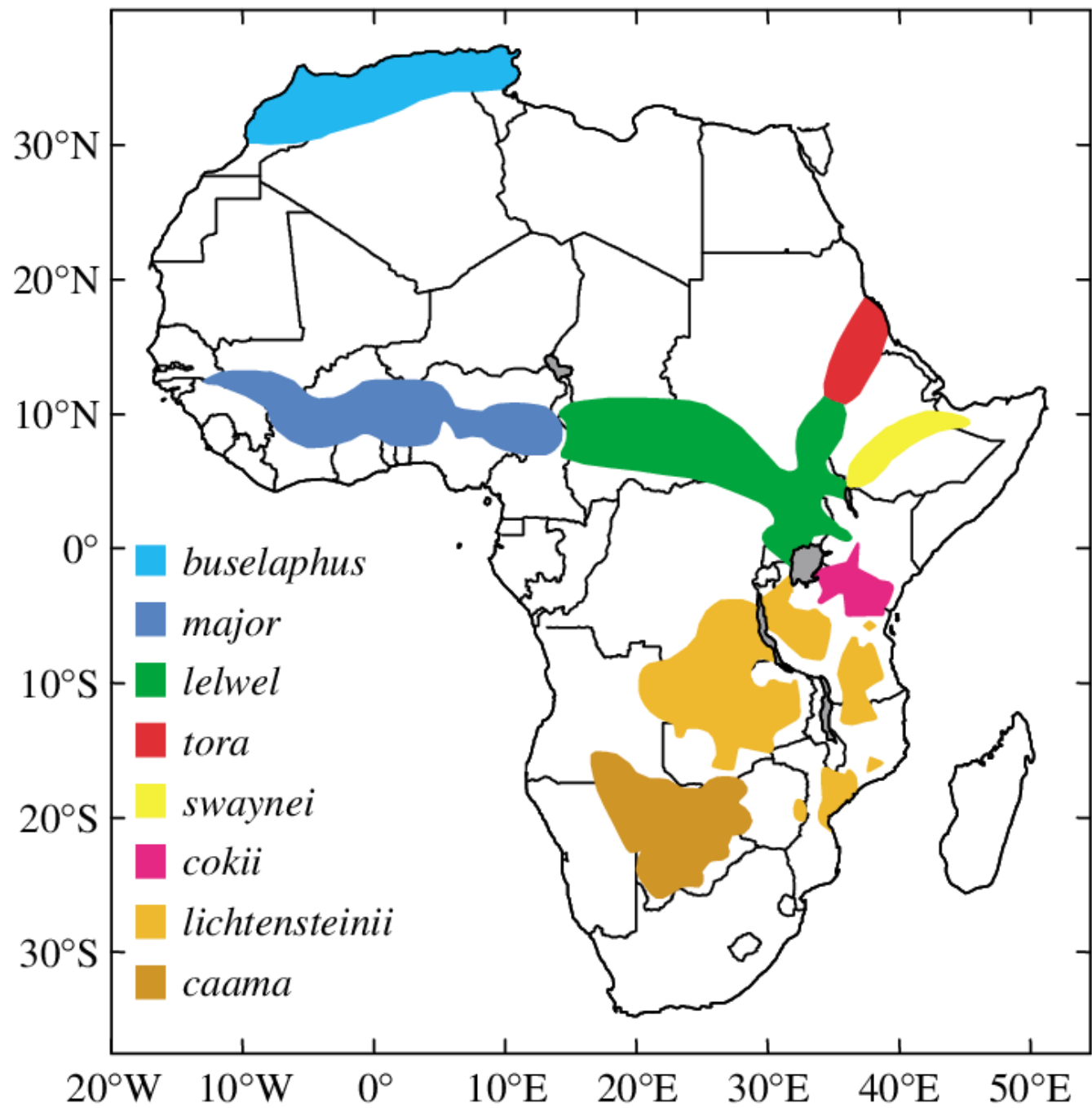
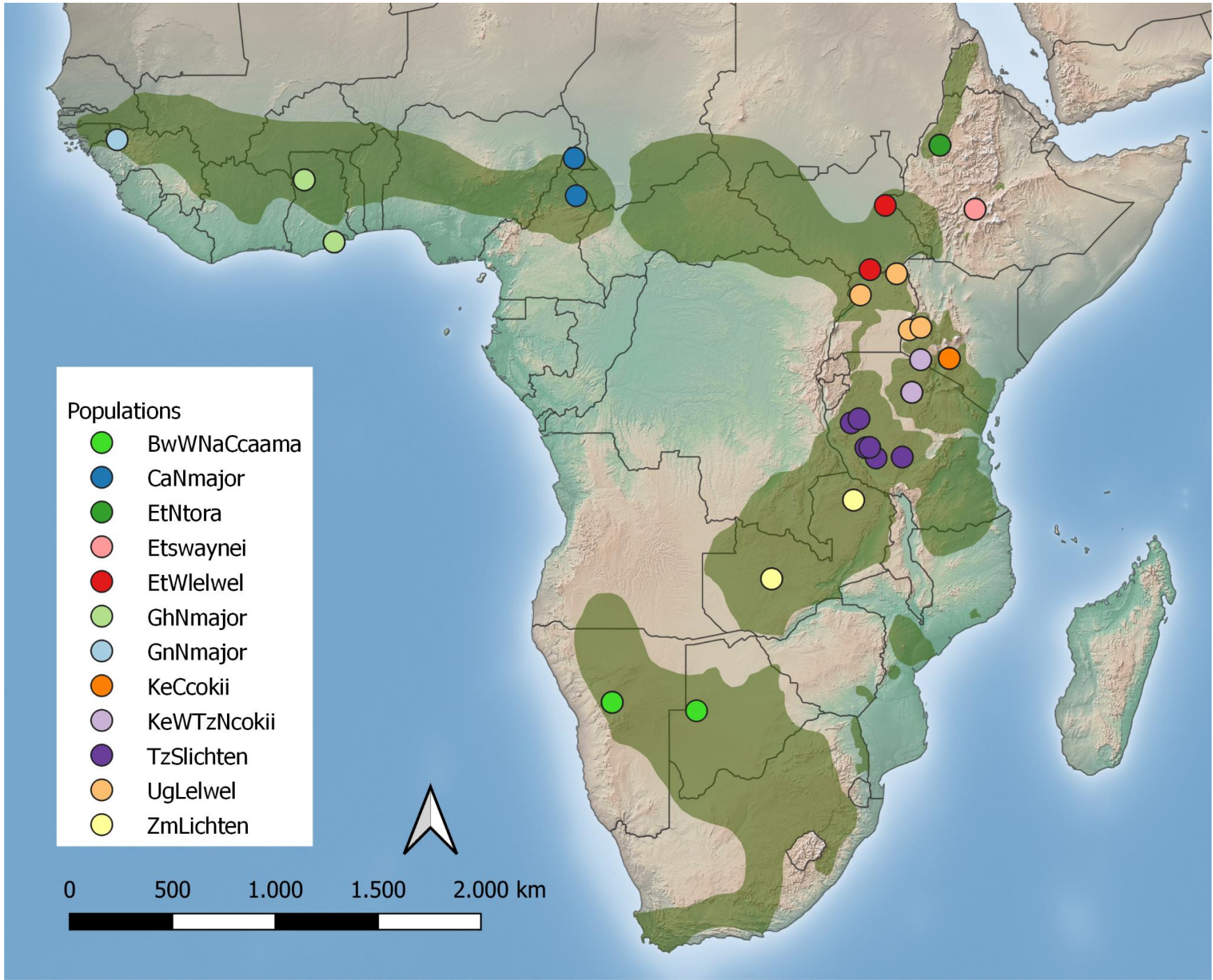


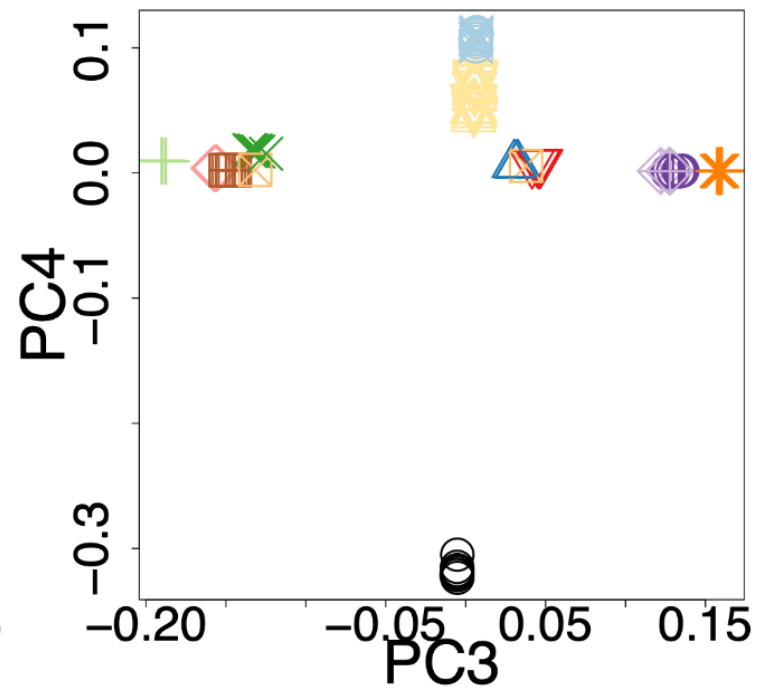
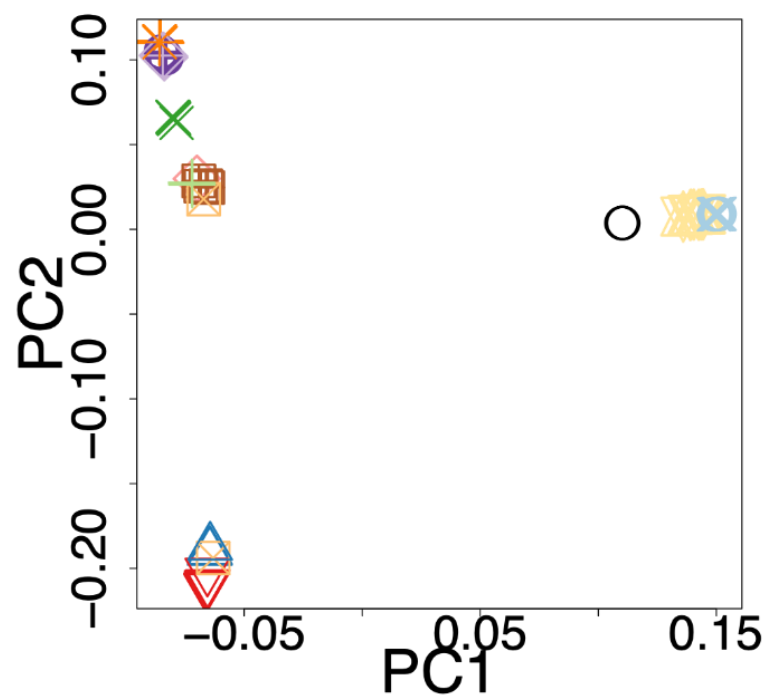
# Population structure in African wildlife

Day 3, Open Institute workshop, Kilifi, August 9<sup>th</sup> 2024

Assoc. Prof. Rasmus Heller, UCPH

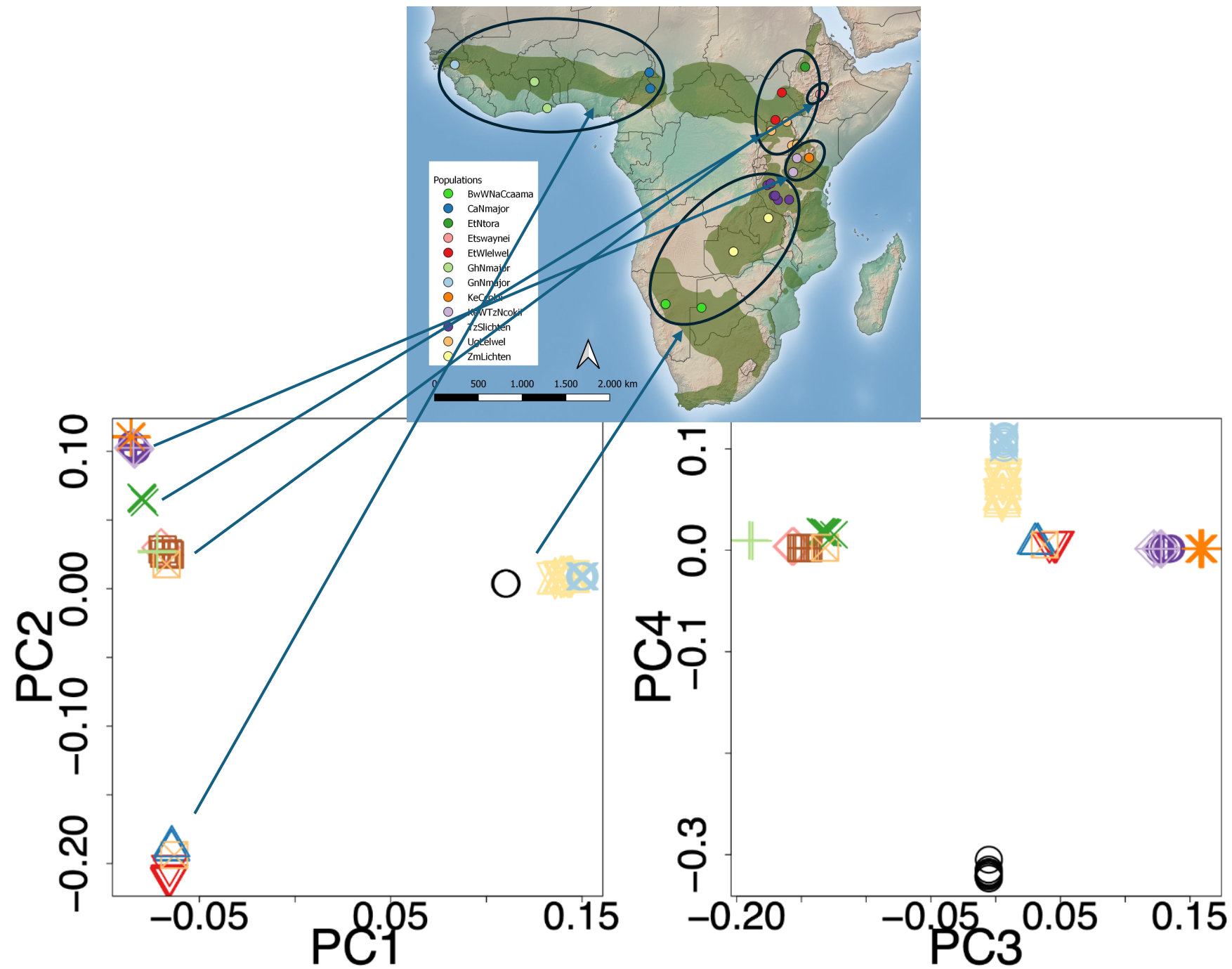


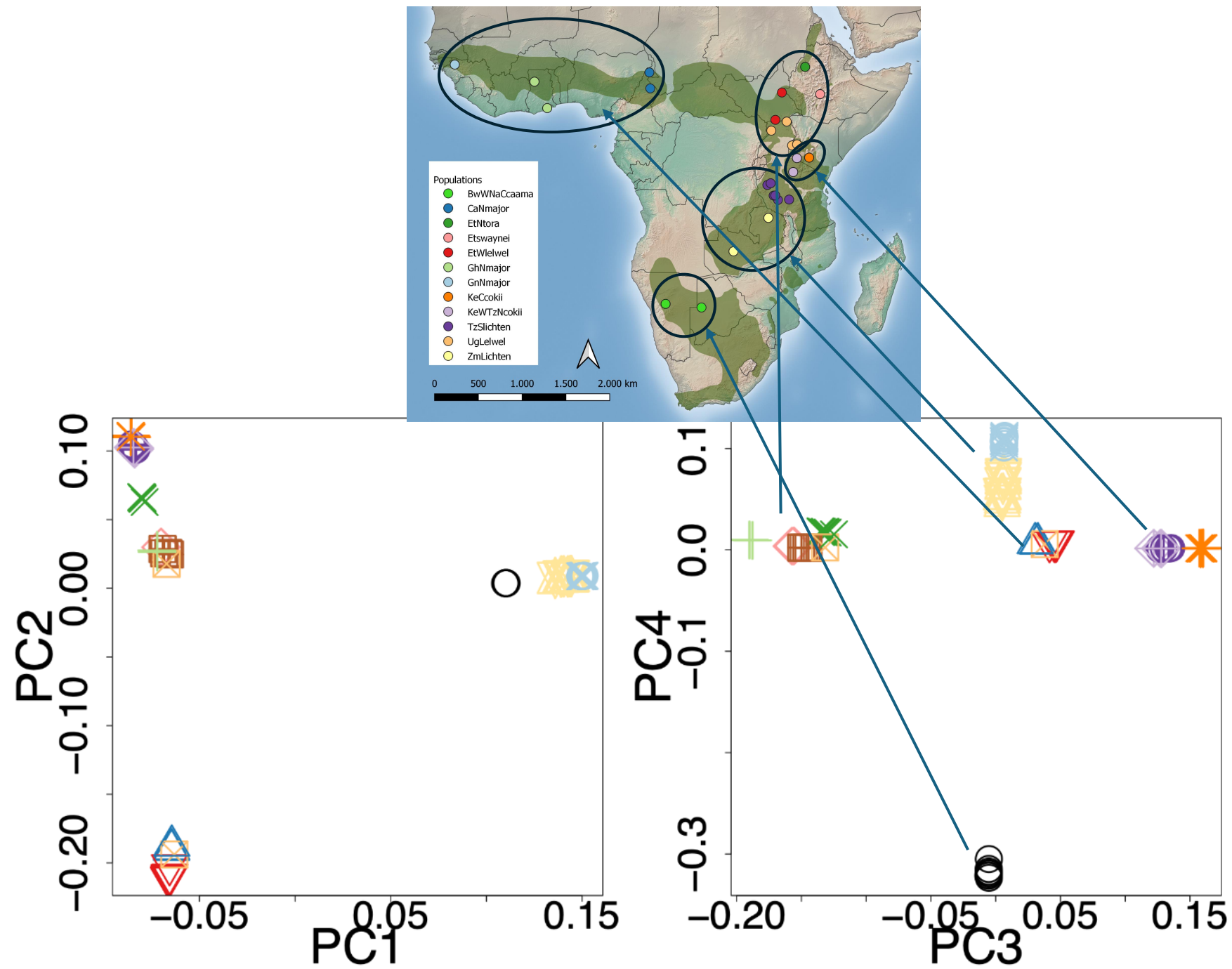




- BNcaama
- △ CaNorth
- + EtNorth
- × EtSwaynei
- ◇ EtWest
- ▽ GhNorth
- ⊠ GnNorth
- ✱ KeCentral
- ⬠ KeWcokii
- ⊕ TzNcokii
- ☆ TzSouth
- ▣ UgLelwel
- ⊞ ZmLichten

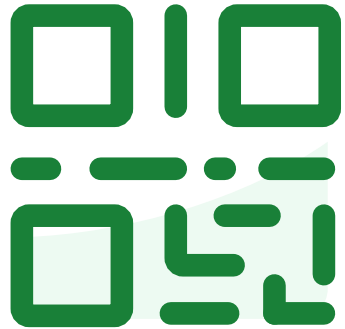






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**How would you characterize population structure in hartebeests (is it strong/weak, geographically sorted, does it correspond to the described subspecies, etc.).**

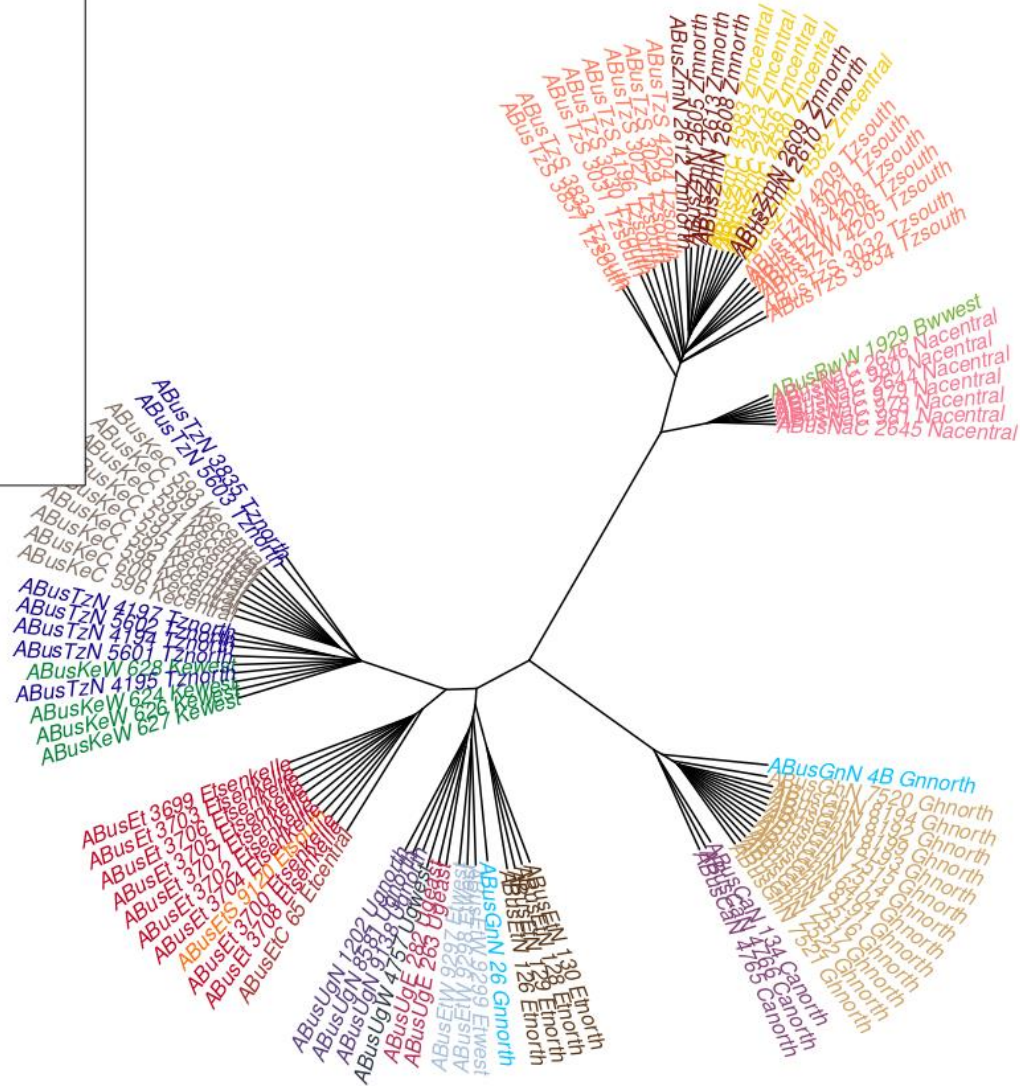
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Sub-Localities Plink-distance square ibs

- BwWest
- CaNorth
- EtNorth
- EtSouth
- EtWest
- EtSenkelle
- GhNorth
- KeCentral
- KeWest
- NaCentral
- TzNorth
- TzSouth
- UgNorth
- UgWest
- UgEast
- ZmCentral
- ZmNorth
- GnNorth
- EtCentral



<b>BwWNaCcaama</b>	0.41	0.53	0.52	0.52	0.51	0.49	0.46	0.56	0.52	0.25	0.3
<b>ZmLichten</b>	0.44	0.56	0.55	0.55	0.54	0.53	0.49	0.59	0.56	0.05	
<b>TzSlichten</b>	0.41	0.53	0.52	0.51	0.5	0.5	0.46	0.56	0.52		
<b>KeWTzNcokii</b>	0.22	0.42	0.41	0.3	0.22	0.26	0.23	0.06			
<b>KeCcokii</b>	0.27	0.46	0.45	0.35	0.25	0.31	0.29				
<b>UgLelwel</b>	0.1	0.33	0.32	0.16	0.19	0.07					
<b>EtWlelwel</b>	0.13	0.36	0.36	0.16	0.2						
<b>Etswaynei</b>	0.18	0.38	0.38	0.2							
<b>EtNtora</b>	0.16	0.37	0.37								
<b>GhNmajor</b>	0.09	0.07									
<b>CaNmajor</b>	0.09										
	<b>GnNmajor</b>	<b>CaNmajor</b>	<b>GhNmajor</b>	<b>EtNtora</b>	<b>Etswaynei</b>	<b>EtWlelwel</b>	<b>UgLelwel</b>	<b>KeCcokii</b>	<b>KeWTzNcokii</b>	<b>TzSlichten</b>	<b>ZmLichten</b>

$F_{ST}$  measures  
differentiation

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**Can you deduce the deepest  
evolutionary divergence in  
hartebeests? What about the second  
deepest?**

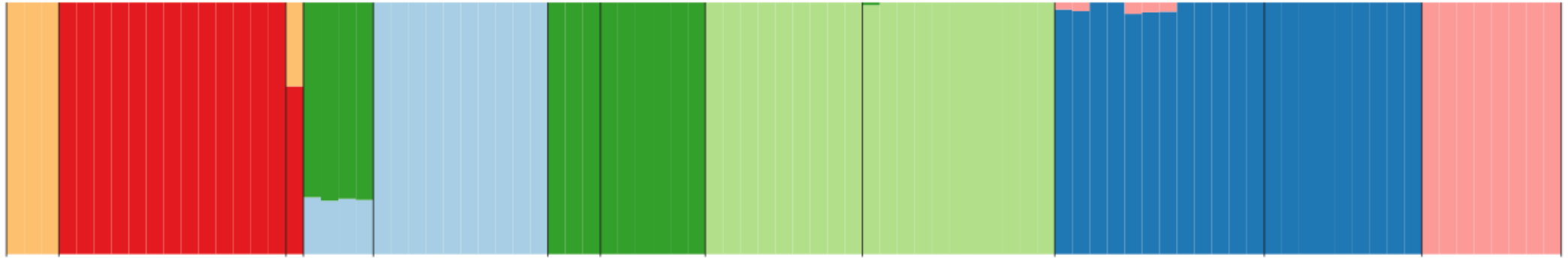
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# Hartebeest

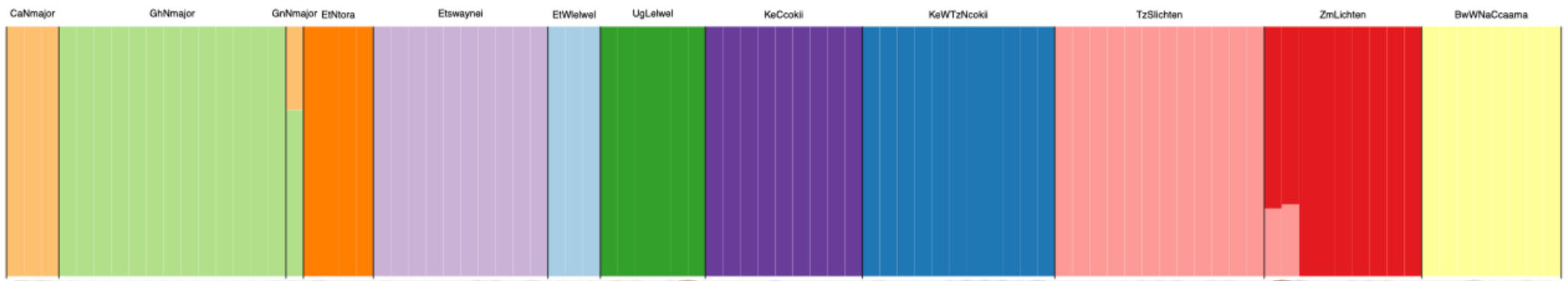


evalAdmix  
Correlations

Admixture proportions



K=7



K=11

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**How many hartebeest  
populations are there in this  
data? Are they admixing?**

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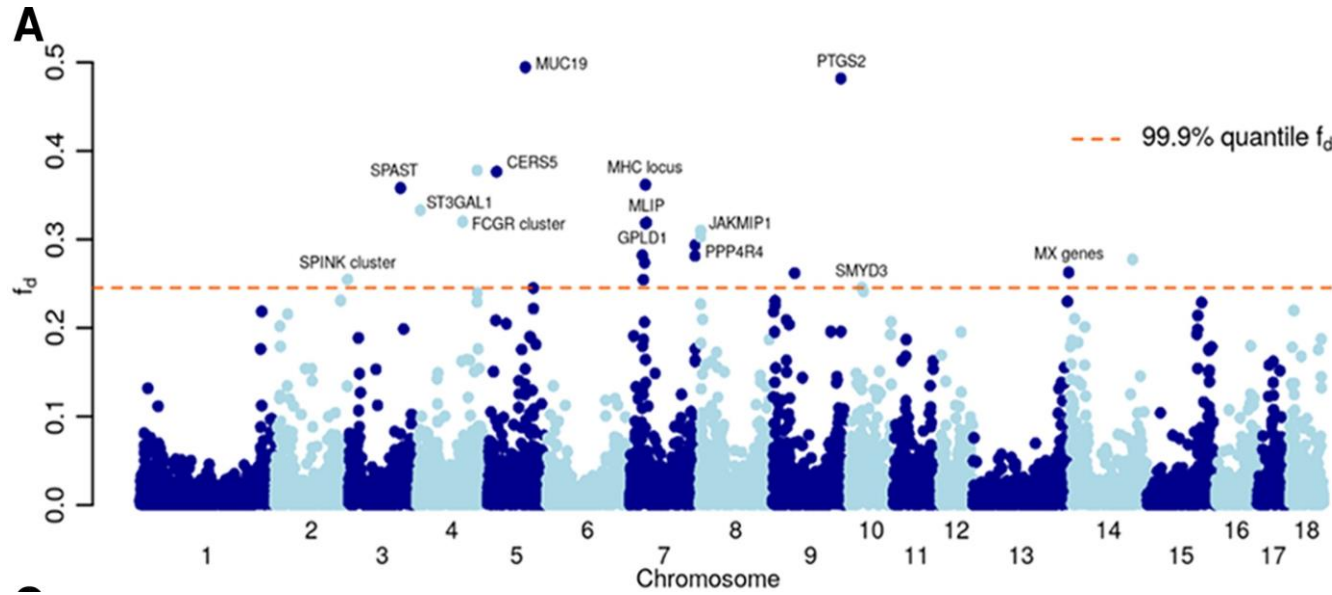


# Examples of admixture in African wildlife

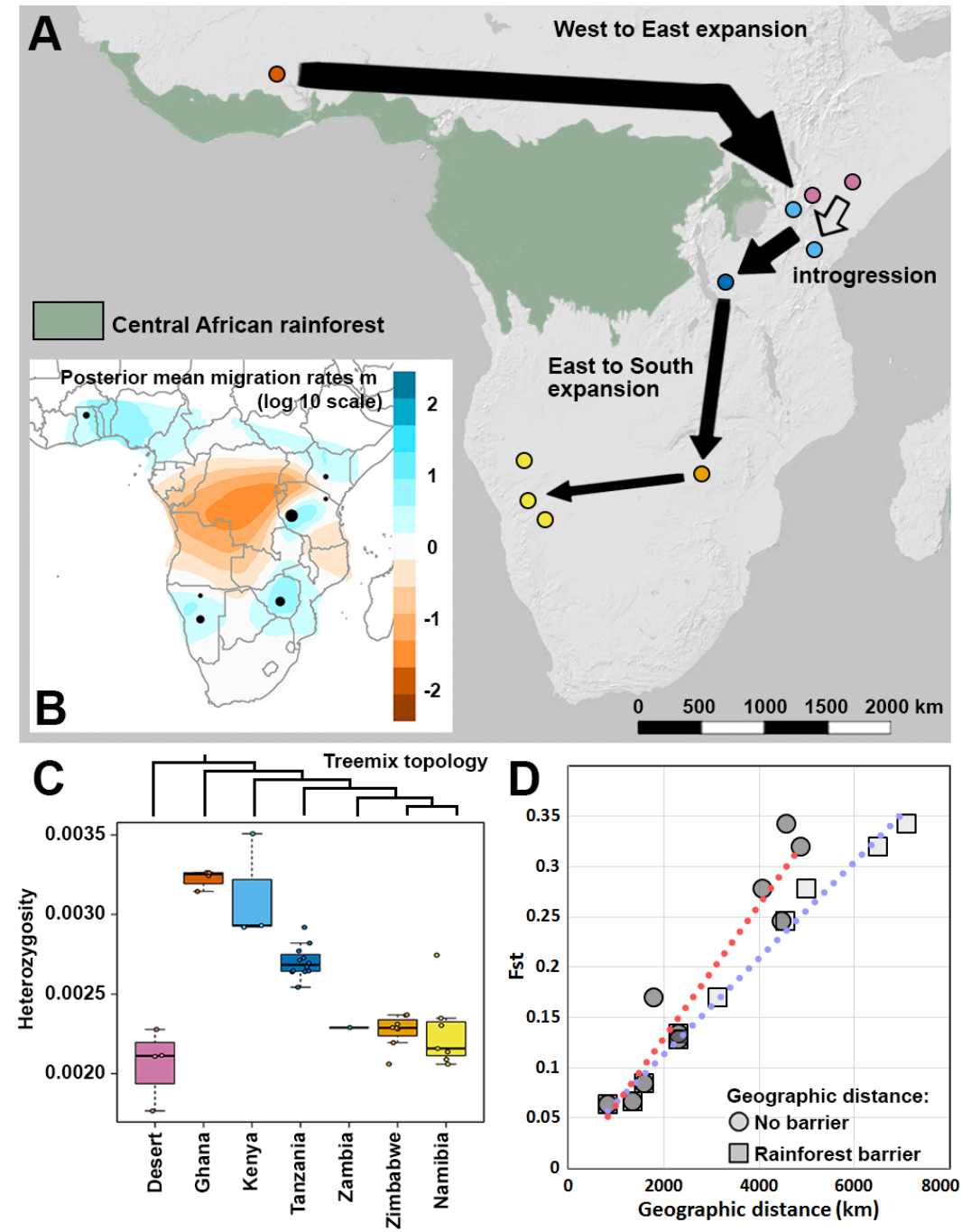
# Case: warthogs



# Case: warthogs



Immune system genes introgressed from desert to common warthog

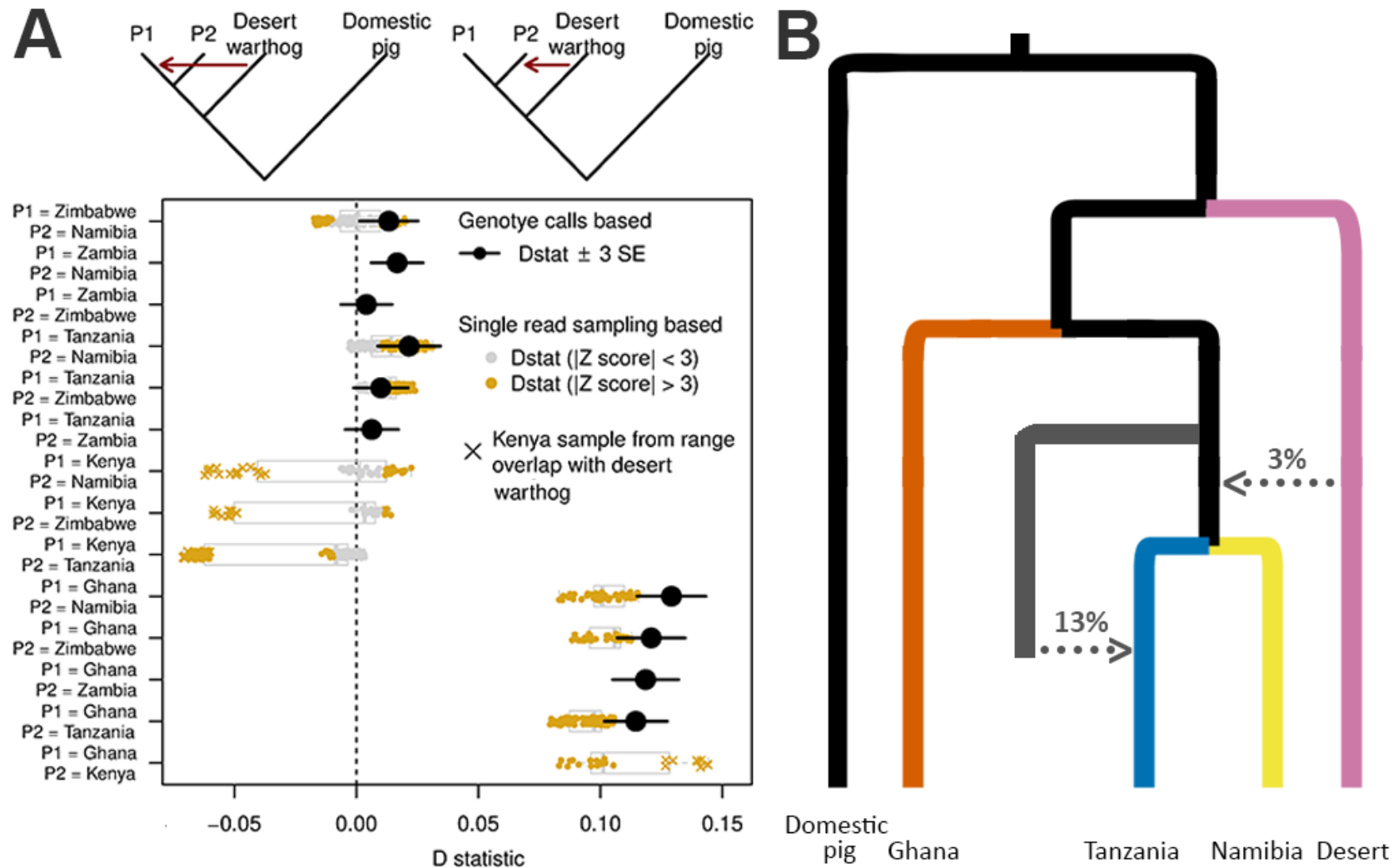




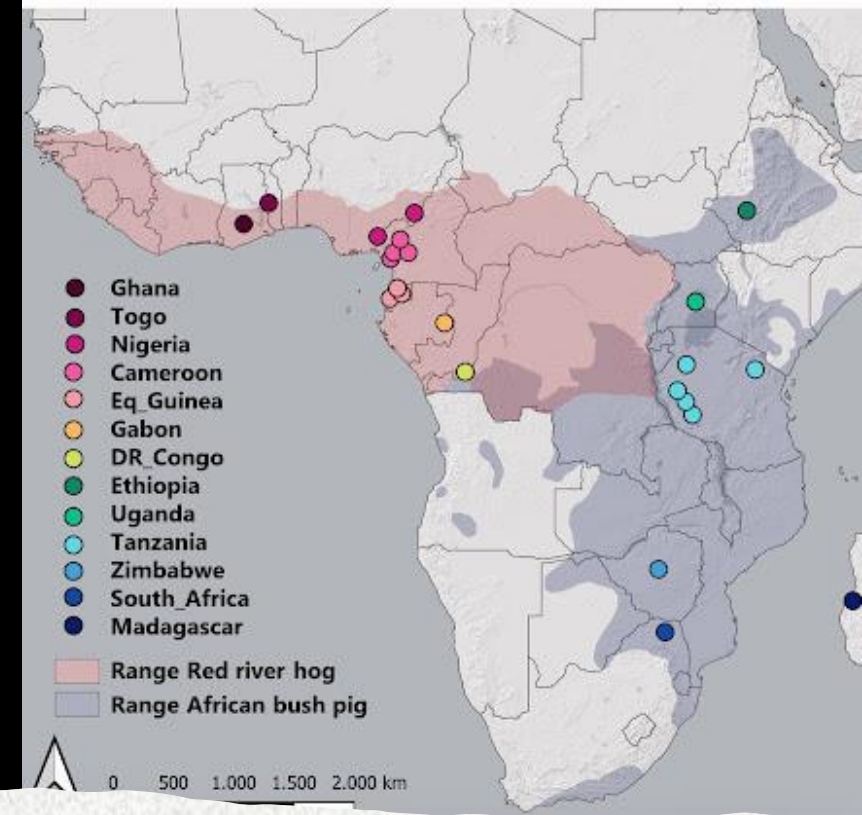
# Why would immune genes introgress as common warthogs expanded their range?

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# How do we know there was introgression?







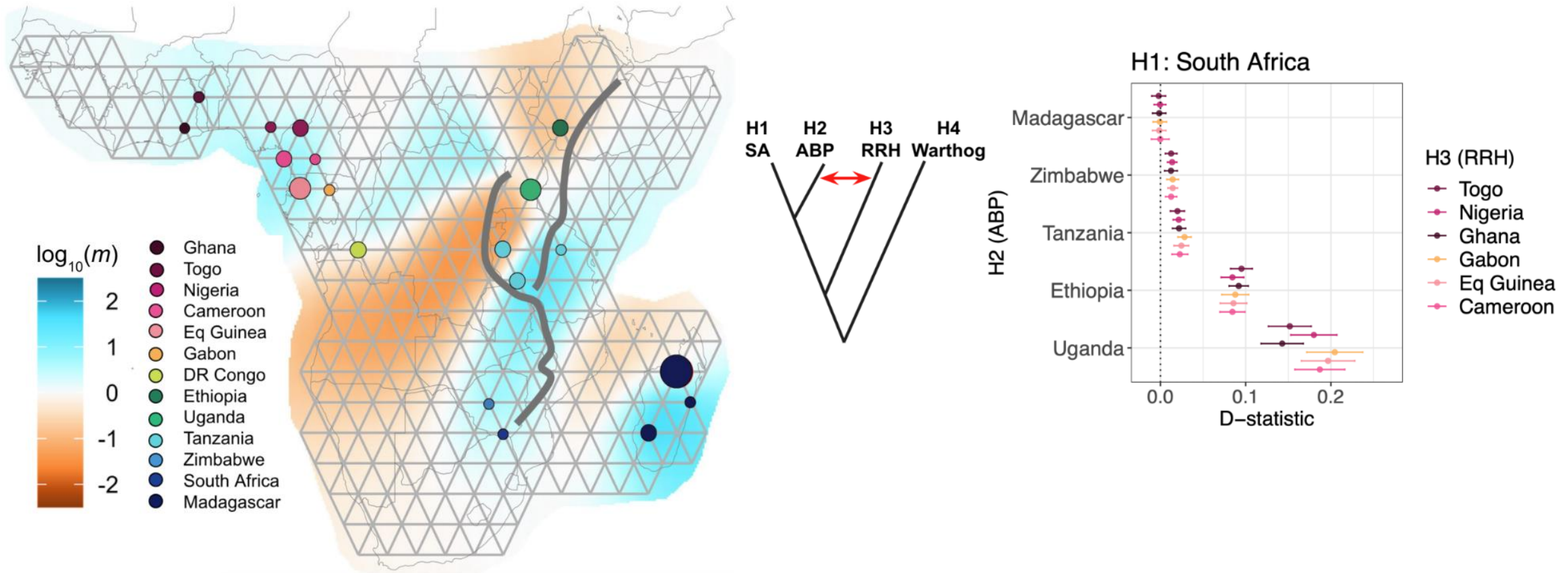
## Case: river hogs

*“...the two populations are morphologically distinct with no indication of gene flow”*

Peter Grubb, 1993

# Case: river hogs

b



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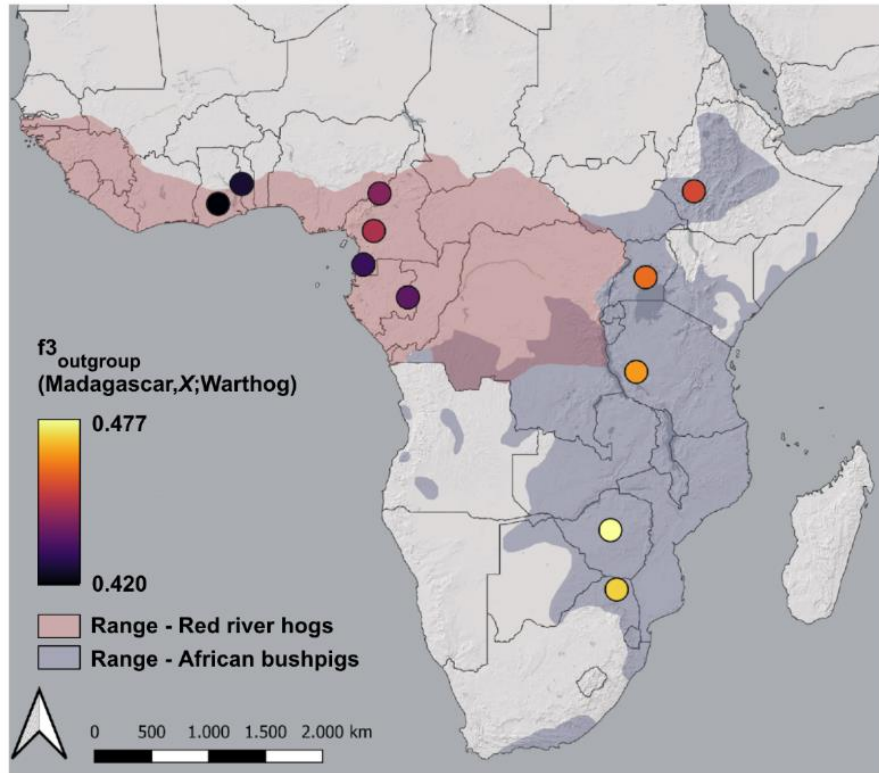
**Are there two species of  
Potamochoerus, and how does  
it influence their conservation?**

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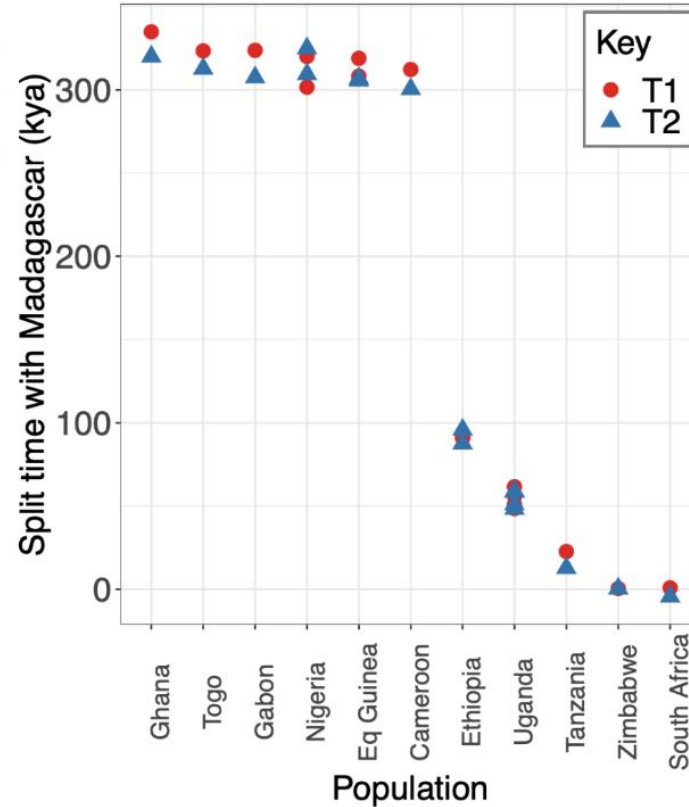


# Case: river hogs

**a**



**b**



**c**

