Principle Component Analysis & Mosquitos



Eve198

Week 8

Maddie Armstrong & Rachael Bay

Article

Current Biology

Climate and Urbanization Drive Mosquito Preference for Humans

Highlights

 African populations of Ae. aegypti vary in preference for human versus animal odor

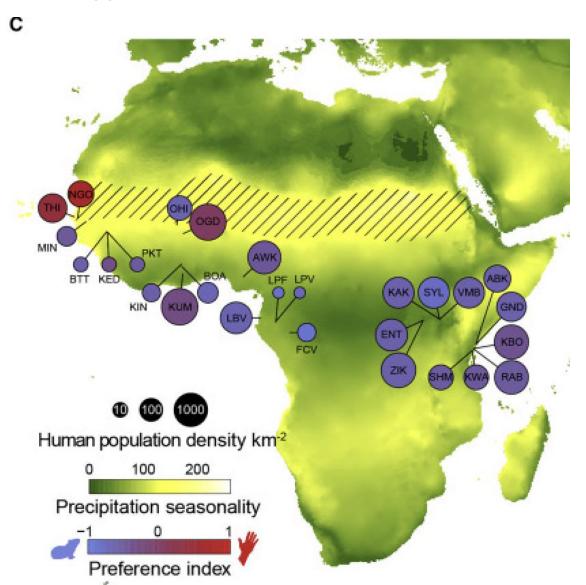
Authors

Noah H. Rose, Massamba Sylla, Athanase Badolo, ..., Bradley J. White, Jacob E. Crawford, Carolyn S. McBride



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Collected mosquito eggs with ovitraps
Hatched eggs in lab for human preference experiments



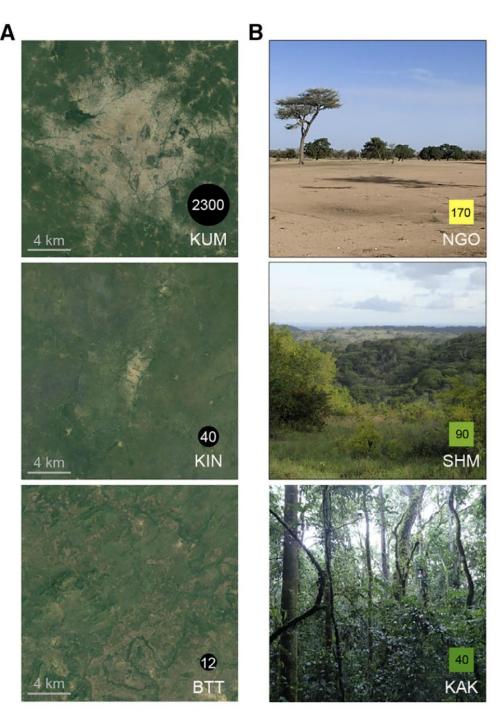


Fig 1: Preference for human odor varies widely in Aedes aegypti mosquitos across Africa

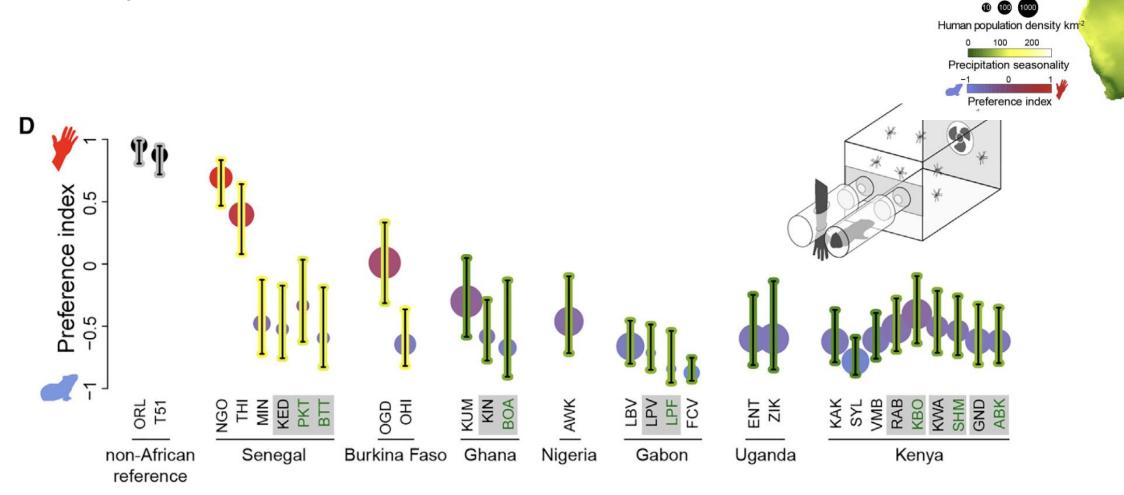
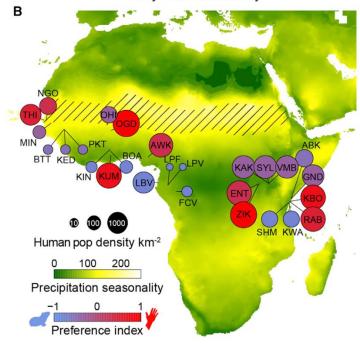
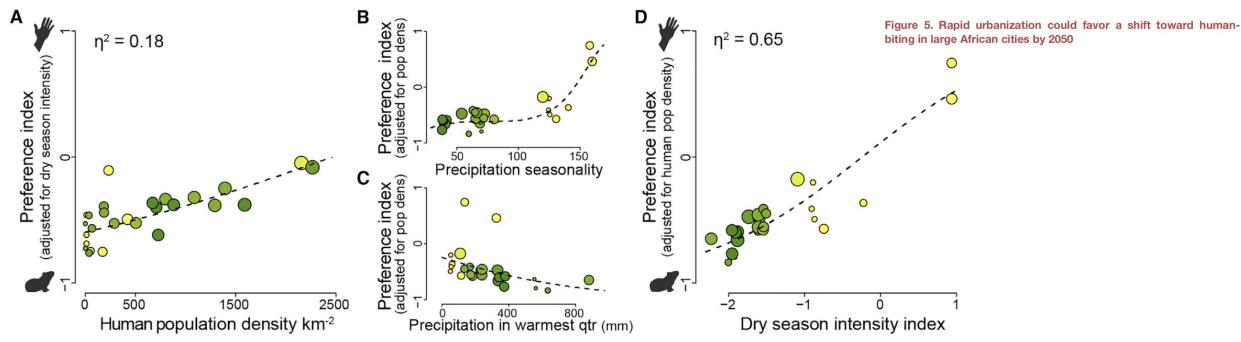


Fig 2: Human Population Density + Dry Season Explain Variation in Preference





Bioinformatic Pipeline:

Extracted DNA from 480 field collected mosquitos



Whole Genome Sequencing



Fastqc



Called variants using ANGSD and bcftools



Aligned to reference genome with GATK



Adapters trimmed with cutadapt



Variant Call Format (VCF) file to interpret genetic data

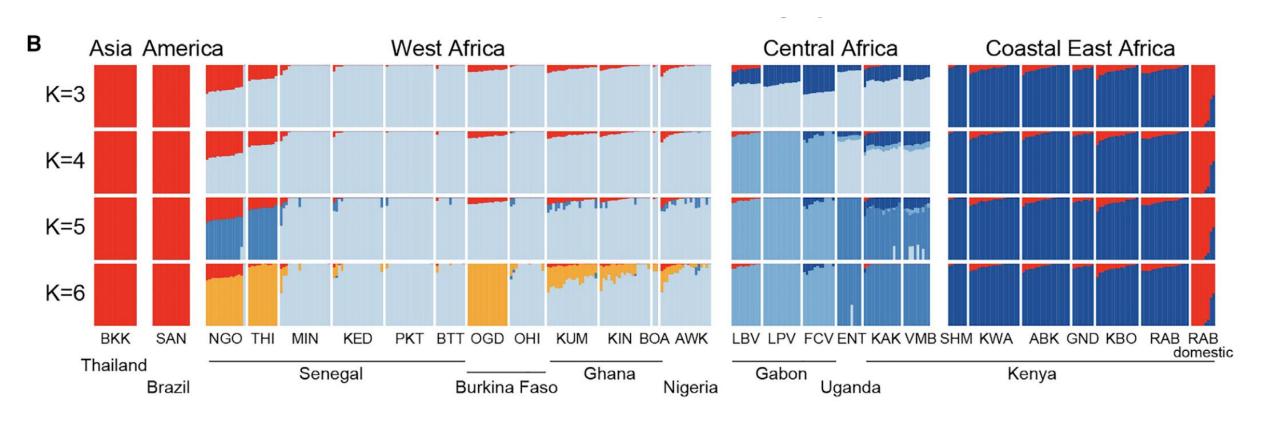


Assessed admixture and conducted PCA based analyses



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Fig 3: Revisiting Population Structure:

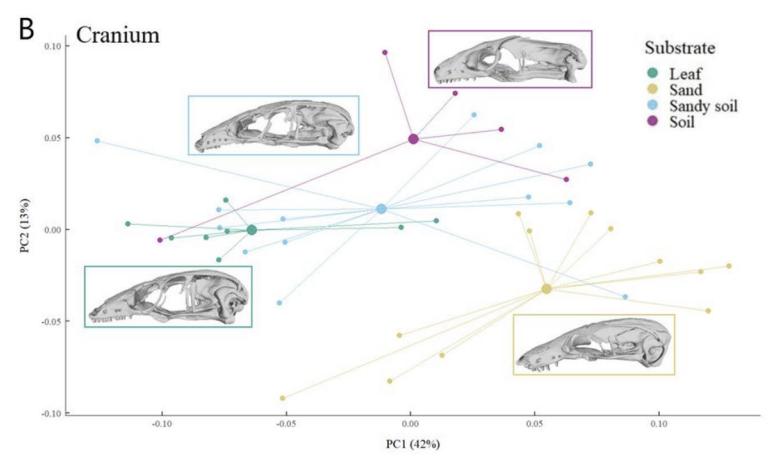


What is a Principle Component Analysis?

A method to reduce the dimensionality of large datasets

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A method to reduce the dimensionality of large datasets



Phylogenetic history influences convergence for a specialized ecology: comparative skull morphology of African burrowing skinks (Squamata; Scincidae)

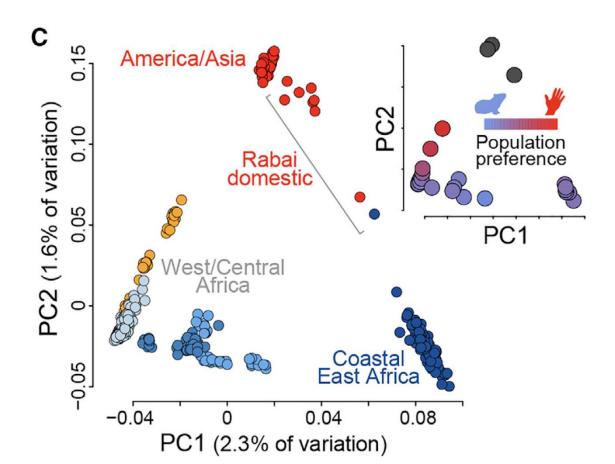
Natasha Stepanova^{1,2*} and Aaron M. Bauer¹



(Stepanova & Bauer, 2021)

What is a Principle Component Analysis?

- A method to reduce the dimensionality of large datasets
 - Genomic Data= many SNPs for many individuals





Steps involved in running a PCA

- Data standardization
 - PCA is sensitive to variances in the data
 - For example, if one site is much more variable than any other



$$\begin{bmatrix} Cov(x,x) & Cov(x,y) & Cov(x,z) \\ Cov(y,x) & Cov(y,y) & Cov(y,z) \\ Cov(z,x) & Cov(z,y) & Cov(z,z) \end{bmatrix}$$

- Compute the eigen vectors and eigen values of the covariance matrix
 - This is where we identify the principal components of the data!

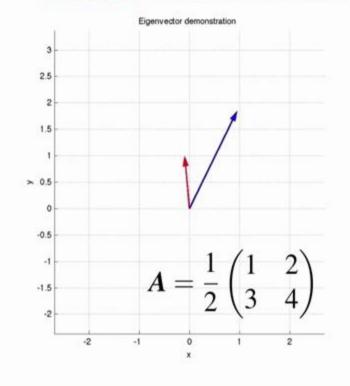


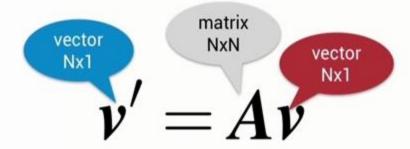
Eigenvalues and eigenvectors

Basically:

An eigenvector points in the direction it is stretched by the transformation and the eigenvalue is how much it was stretched.

Vector transformation



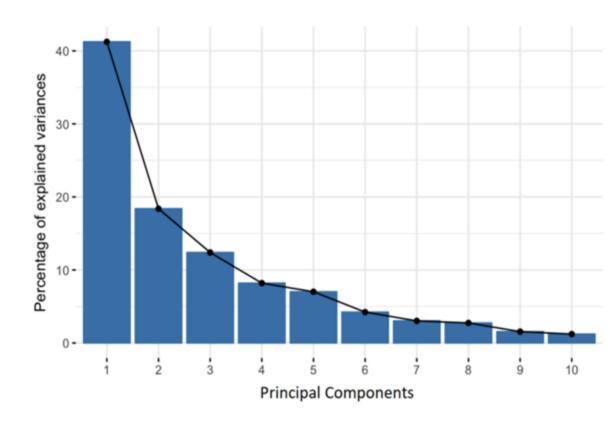


- For every matrix A there are N vectors whose direction is unchanged by A
 - these are the eigenvectors
 - their length is scaled by the eigenvalue

What is a principal component?

 New variables constructed from a mixture of our initial variable.

- A PCA will return to you the same dimension of data you gave it.
 - But most of the information should be in the first few PCs
 - For 10 individuals genotyped at 100 loci you will get 10 principal components

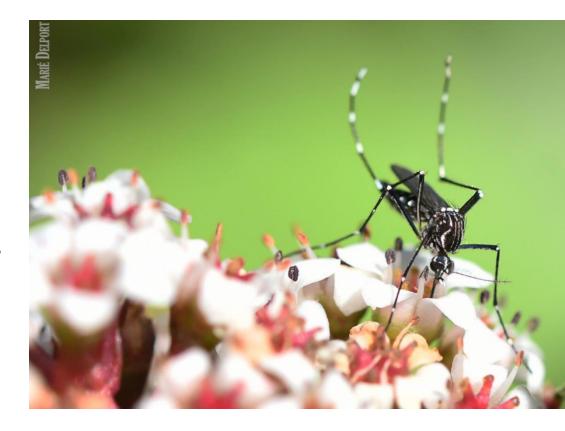


Why do we care about genetic diversity?

Tells us about population demography, gene flow

Can inform conservation strategies

 Evolution requires variation in phenotypes and genotypes

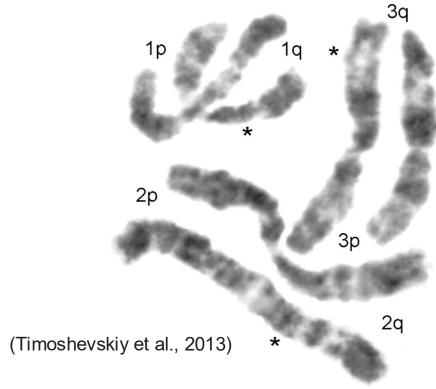


Working with a subset of data today:

Our methods differ from those used in this paper, primarily for convenience and time (ours).

Just a section of chromosome 1





An Integrated Linkage, Chromosome, and Genome Map for the Yellow Fever Mosquito *Aedes aegypti*

Vladimir A. Timoshevskiy, David W. Severson, Becky S. deBruyn, William C. Black, Igor V. Sharakhov, Maria V. Sharakhova 🖪