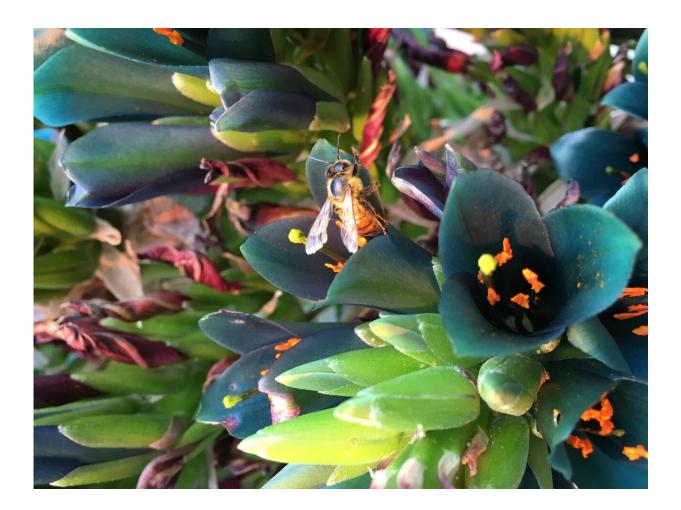


Genome sequencing shows climate barrier to spread of Africanized bees

October 21 2020, by Andy Fell



A wild honeybee in Davis, California. European honeybees (Apis mellifera) were introduced to the Americas in the 1600s. Since 1957, "Africanized" honeybees (Apis mellifera scutellata) have interbred with other honeybees. New genomics research shows that climate forms a barrier to the spread of these scutellata genes to the north and south. Credit: Erin Calfee



Since the 1950s, "Africanized" honeybees have spread north and south across the Americas until apparently coming to a halt in California and northern Argentina. Now genome sequencing of hundreds of bees from the northern and southern limits shows a gradual decline in African ancestry across hundreds of miles, rather than an abrupt shift.

"There's a gradual transition at the same latitude in North and South America," said Erin Calfee, graduate student in the Department of Evolution and Ecology at the University of California, Davis, and first author on the paper, published Oct. 19 in *PLOS Genetics*. "There's a <u>natural barrier</u> that is likely maintained by many different genetic loci."

That barrier is mostly likely climate. Bees with majority African <u>ancestry</u> are unable to survive colder winters.

European colonists brought European species of honeybees (Apis mellifera) with them to the Americas as early as the 1600s. In addition to apiaries, these bees established in the wild alongside native bees.

In 1957, imported African honeybees of the subspecies Apis mellifera scutellata swarmed out of experimental hives in Brazil and started to rapidly spread, interbreeding with the resident European bees. Native to southern and eastern Africa, scutellata bees are known for their defensive behavior. They also carry some useful traits for beekeepers, such as resistance to Varroa mites.

Calfee, working with Professor Graham Coop and Associate Professor Santiago Ramirez from the Department of Evolution and Ecology and collaborators in Argentina, sequenced the genomes of bees collected at the northern and southern edges of the scutellata expansion.

They found that the bees at the northern and southern edges of the range have a highly variable mix of scutellata and European bee ancestry. The



higher the latitude, the less scutellata ancestry is in the mix.

"The whole genome is tracking latitude and climate," Calfee said. There are likely many genes involved in climate sensitivity and winter survival, she said. But Calfee also finds that in some parts of the genome scutellata ancestry has spread far beyond these climate limits in both North and South America, evidence that some scutellata genes are advantageous and not tied to climate sensitivity. In contrast, the researchers did not find any evidence for selection for European ancestry in the spread of scutellata bees.

The findings challenge the idea of a binary difference between "Africanized" and "European" honeybees, Ramirez said. In fact, all of these introduced honeybees are hybrids to some degree.

Diversity a resource for breeding bees

The results of the study could be of interest for breeding bees for desirable traits, such as resistance to pathogens. Although the researchers looked only at gene sequences and not the resulting phenotypes (except for one, wing length), the results do show which genetic loci are important because they are under selection in the hybrid zones.

Scutellata bees and hybrids have a lot more genetic diversity than European bees, and the researchers found that they lost very little of this diversity in their expansion from Brazil to California.

"Beekeepers can potentially draw from this genetic variation to breed for desirable traits," Ramirez said. What started out as an invasion may become part of the solution to declining bee health.

"It makes sense but it's kind of surprising because we have 1.5 million (honeybee) colonies being brought into California every spring to



pollinate crops," Ramirez said. Those domesticated bees are then trucked around the country, but they do not appear to have a large effect on the spread of African ancestry into feral <u>honeybee</u> populations.

More information: Erin Calfee et al, Selection and hybridization shaped the rapid spread of African honey bee ancestry in the Americas, *PLOS Genetics* (2020). DOI: 10.1371/journal.pgen.1009038

Provided by UC Davis

Citation: Genome sequencing shows climate barrier to spread of Africanized bees (2020, October 21) retrieved 13 May 2024 from <u>https://phys.org/news/2020-10-genome-sequencing-climate-barrier-africanized.html</u>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.