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Original Article

TIMING OF INVASION BY AFRICANIZED BEES COINCIDES WITH LOCAL EXTINCTION OF A SPECIALIZED POLLINATOR OF A RARE POPPY IN UTAH, USA

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Abstract

The introduction of exotic species can have profound impacts on mutualisms between native species in invaded areas. However, determining whether a new invader has impacted native species depends on accurately reconstructing the invasion timing. The arrival of Africanized honey bees (AHB) in southern Utah at some point between 1994 and 2011 has recently been implicated in the local extinction of *Perdita meconis*, a native specialist pollinator of an endangered poppy, *Arctomecon humilis*. Although AHBs were purportedly first detected in southern Utah in 2008, their presence in nearby Nevada, Arizona, and New Mexico by 1998-2001 suggests that they may have been present in Utah much earlier. We refined the arrival date of AHBs in southern Utah by using a molecular marker to determine maternal ancestry of museum specimens collected between 2000 and 2008. We found that AHBs were present in southern Utah from 2000 onwards, advancing the arrival date of this invader by at least 8 years. This lends credence to the hypothesis that AHBs played a critical role in the local extinction of *P. meconis* in Utah. This work also highlights the importance of vouchering even common species such as honey bees in museum collections to serve future research needs.

Keywords: *Apis mellifera* L, *Arctomecon*, disrupted mutualism, museum collections, molecular diagnostics, Perdita meconis

NTRODUCTION

The introduction of exotic species can have profound impacts on mutualisms in native ecosystems (Traveset & Richardson, 2014). For example, the replacement of a native ant seed disperser by the invasive Argentine ant has greatly changed seed dispersal dynamics by plants of coastal scrublands in Australia (Rowles & O'Dowd, 2009). There are also many examples of disruption by invasives of pollination systems with significant alterations to both pollinator and plant components of the mutualism (Traveset & Richardson, 2014).

Recently, Portman, et al. (2018) presented evidence that implicated Africanized honey

bees (AHB) as the primary force that drove a population of the highly specialized bee, Perdita meconis, to extinction in Southern Utah, USA. This small, native bee is a specialist pollen collector and pollinator of poppies in the genera Arctomecon and Argemone (Griswold, 1993), including the endangered Dwarf bear poppy, Arctomecon humilis, an endemic of Washington County, Utah (USFWS, 1979). The isolated population of *P. meconis* in southern Utah disappeared between 1994 and 2012 (Tepedino, et al., 2014; Portman et al., 2018), a broad span of time that overlaps with the advance of AHBs into the region. However, assigning an important role to the AHB in the local extinction of *P.meconis* requires that we establish a convincing timeline

of invasion; current reports claim AHBs did not arrive in Utah until 2008 (Hodgson et al., 2010; UDAF, 2017). If this invasion date is correct, then AHBs drove a specialized native bee population to extinction in a maximum of four years. Alternatively, AHBs may have been present in southern Utah much earlier than 2008 but escaped detection.

AHBs are one of the prime exemplars of rapid advance by an invasive species. Apis mellifera scutellata queens were first imported into Rio Claro, Sao Paulo, Brazil in 1956 (Moritz, Härtel, & Neumann, 2005). Swarms from an experimental apiary escaped containment, and 26 queens, along with the paternal genes of their approximately 200 mates, escaped and spread through the landscape at an astonishing rate (Michener, 1975). The initial proliferation of AHBs throughout the New World was estimated at over 480 km per year in the first few decades, but slowed after they reached the United States (Visscher, Vetter, & Baptista, 1997). This spread has been accomplished through gene flow, with current feral honey bee populations in the invaded areas consisting largely of hybrids between European and Africanized lineages (Quezada-Euán, Pérez-Castro, & May-Itzá, 2003; Kraus, Franck, & Vandame, 2007).

AHBs were first reported in the United States in 1990, when a feral swarm was trapped near Hidalgo, Texas; by 1993, they had reached southern New Mexico and Arizona; they were found in southern California in 1994, in southern Nevada in 1998, and by 2001 had been recorded throughout Arizona (Winston, 1992; Visscher, Vetter, & Baptista, 1997; Hicks, 1999; Hodgson et al., 2010). However, despite monitoring efforts in the neighboring state of Utah beginning in 1994, AHBs were not reported until 2008 when the first AHB colony was detected in Cedar City in Iron County, with additional colonies detected in Washington and Kane Counties the following vear (Burfitt, 2009; UDAF, 2017). Their delayed arrival in Utah is curious considering the absence of geographical barriers along its borders with Arizona and Nevada.

Our objective here is to examine arrival dates of AHBs in Utah more closely. An earlier arrival

date would strengthen our contention that AHBs played a critical role in the local extinction of *P. meconis* in Utah (Portman et al., 2018). To further probe arrival dates, we used honey bee specimens obtained from museum collections. Natural history collections and molecular diagnostics are increasingly being used to refine the known distributions of insects, track changes due to climate change, document species declines, and capture information on insect movement through time (Shaffer, Fisher, & Davidson, 1998; Lees et al., 2011; Lister & Climate Change Research Group, 2011).

MATERIAL AND METHODS

Honey bee specimens collected in 1994-2011 from southern Utah were sought from the following national and regional collections: The National Pollinating Insect Collection at USDA-ARS Bee Biology and Systematics Laboratory (BBSL), Monte L. Bean Life Science Museum Collection at Brigham Young University (BYU), Garth and Jerri Frehner Museum of Natural History Collection at Southern Utah University (SUU), University of Utah Natural History Museum Collection (UMNH), and the Dixie State University Natural Science Museum Collection (AHBC). Specimens were shipped to Utah State University for genetic classification, and label data including plant hosts and elevation were recorded for each specimen.

Each specimen was assigned to either European or African maternal ancestry following the methods of Portman et al. (2018). In brief, a single mid-leg was removed from each specimen and DNA was extracted with a salting-out method (Sambrook & Russell, 2001). These extracts were then subjected to a cytochromeb-based, multiplex PCR test (Szalanski & McKern, 2007). Each sample exhibiting amplicons at both 385 and 485 bp was classified as Africanized (AHB), while those with a single amplicon at 385 bp were classified as European (EHB). This detection method is based on a mitochondrial marker, therefore it cannot detect the presence of AHB genes inherited along paternal lines. However, mitochondria are inherited intact

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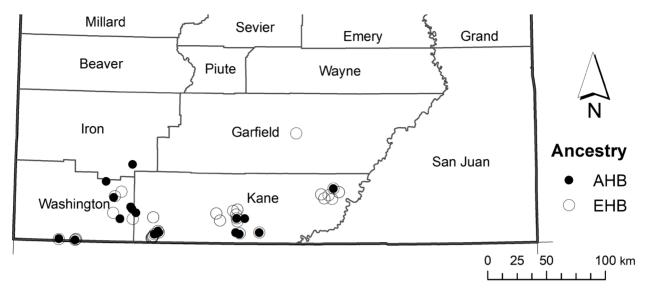


Fig. 1. Map of *Apis mellifera* sample locations and maternal ancestry test results throughout the counties of southern Utah. Solid circles indicate Africanized bees (AHB); open circles indicate European bees (EHB).

Table 1.

Honeybees mitotyped by Utah County, elevation (m) and year of collection, EHB=European honeybee; AHB=Africanized honeybees

County	МТуре	Number	Elevation	Years
Garfield	EHB	1	1718	2002
Iron	AHB	1	2450	2003
Kane	EHB	69	1153-1915	2000-2003
	AHB	17	1326-1882	2000-2001, 2003
Washington	EHB	4	1264-2408	2006-2007
	AHB	6	1254-2107	2006-2007

along the matriline, so a single marker accurately represents an individual's matriline.

RESULTS

A total of 98 worker and drone specimens collected between 2000 and 2007 were obtained from Kane, Garfield, Iron, and Washington counties in southwestern Utah (Fig. 1). Specimens were not available from San Juan, the southeastern-most county, nor for the 1994-1999 period. No specimens meeting our time and location criteria were available from the UMNH or AHBC collections.

Ninety-seven of 98 specimens were successfully mitotyped, with 24 (25%) classified as AHB and 73 (75%) as EHB (Tab. 1). AHBs were found in every year for which specimens were tested. In 2000, the earliest year for which honeybees were available, four of 14 Kane County specimens tested positive for Africanization; all were collected in Coral Pink Sand Dunes State Park from 1153-1886 m elevation (Tripodi et al., 2018). Over all sites and years, AHBs were found at elevations ranging from 1153-2450 m (Tripodi et al., 2018). Floral records were available for 54 specimens, representing 15 plant species (Tripodi et al., 2018). AHBs were found on both native and exotic plants in the region, including the protected native endemic Asclepias welshii and the ubiquitous invasive sweetclovers (Meliltus alba and M. officinalis) and saltcedars (Tamarix sp.).

TRIPODI ET AL.

DISCUSSION

Our results clearly establish that AHBs had invaded southern Utah by 2000 and not in 2008 as previously reported (UDAF, 2017). Although we employed a mitochondrial marker that did not survey paternal contributions, the earliest samples available showed that AHB was present by 2000. The addition of nuclear data would not alter this result. Evidently invasion took place sometime after 1993: in a previous study, no specimens tested from Washington County dating 1981-1993 were Africanized (Portman et al., 2018). AHBs were present throughout adjacent Arizona by 2001 and were first reported in Clark County, Nevada in 1998 (Hicks, 1999). Clark County is 17.4 km from the Utah border in Washington County (UTM-12N, ArcMap v.10.3.1, ESRI, Redding, CA), and there are no obvious geographic barriers to impede the expansion of AHBs from Arizona and Nevada into Utah. Combined, these data suggest that AHBs may have invaded southern Utah as early as 1998.

Advancing the date of arrival in Utah by 8-10 years greatly strengthens our hypothesis that AHB played a critical role in the local extinction of *P. meconis* (Portman et al., 2018). To briefly recap that reconstruction, P. meconis was present in association with its host plant, the endangered A. humilis, which produces ample pollen but no nectar, at one site in Washington Co. through 1993 (Tepedino, et al., 2014). It has been absent in every census since 2012 and has been replaced by the pollen-avid AHB (Winston, 1992) which is an active and more numerous forager earlier in the morning. The AHB has also displaced the generalist native bee *Eucera quadricincta* which is now a rare poppy flower visitor. The period in which P. meconis has become extinct in Utah and E. quadricincta has become uncommon in Washington Co. (sometime between 1994 and 2011) coincides with the arrival of the AHB.

While the local extinction of an isolated population of *P. meconis* appears to be the only potential example of a local extinction caused by the AHB, there are numerous reports of honey

bees, both Africanized and European, displacing native flower-visitors or changing their foraging distributions across available floral resources, and thereby reconstituting native ecosystems. For example, Santos et al. (2012), described significant changes in the structure of pollination networks in a xeric biome in Brazil following AHB invasion as did Kato et al. (1999) for the EHB in the Bonin Islands, and Kato and Kawakita (2004) for New Caledonia. In these and many other examples (reviewed in Geslin et al., 2017; Mallinger, Gaines-Dav, & Gratton, 2017), generalist honey bee foragers overrun the preferred floral resources of many native bees and force natives to either visit secondary or tertiary choices, if available, migrate, or expire. Although it is difficult to be certain, the sum of evidence (see Portman et al., 2018) suggests that once the A. humilis Beehive Dome site was invaded by pollen-hungry AHBs, P. meconis individuals were doomed: they could not switch hosts nor could these small, weak-flying bees migrate through the fragmented, rapidly developing, Washington Co. landscape to other poppy populations.

This example invites caution before accepting the general hypothesis that a specialist mutualism, such as the *P. meconis-A. humilis* association, "might be more protected against any alien entering the pollination web" (Traveset & Richardson, 2006). While our finding hardly debases their general prediction, it does suggest that the vulnerability of specific mutualisms to invasion lies in the details. In our novel example, AHB has displaced both a highly specialized bee and an intent generalist from a globally rare, though occasionally locally abundant, plant with an uncommon reward structure in the presence of livestock grazing (Portman, Tepedino, & Tripodi, 2018).

Early models of AHB in the USA predicted there would be climate-based limits of its northern expansion; winter conditions were expected to prevent the spread of AHB to high elevations (Taylor, 1977; Taylor & Spivak, 1984; Winston, 1992). One model based on the natural range of *A. m. scutellata* in Africa predicted that AHBs would not venture north of the 16°C January

isotherm (Taylor & Spivak, 1984). Another model, based on physiological cold tolerance, predicted that the AHB would expand much farther north, being stymied only by areas having a maximum of 120 days with a maximum temperature of 10°C or less, an area that encompasses all but the northernmost portion of Utah (Southwick, Roubik, & Williams, 1990). More recent models have incorporated the known occurrence of AHBs in the USA thus far, as well as other environmental factors like precipitation and even patterns of floral resource availability (Jarnevich et al., 2014; Gill & Sangermano, 2016). Today, AHB are found as far north as Napa and Sacramento Counties in California (Lin et al., 2018), and at a similar latitude (~39° N) as far north as Emery and Grand Counties in Utah (UDAF, 2017; Cleary et al., 2018). In this study, AHB mitotypes were more heavily concentrated in the southwestern corner of Utah, which agrees well with modern models of AHB distribution (Jarnevich et al., 2014; Gill & Sangermano, 2016). Nevertheless, the presence of AHB at high elevations such as 2450 m in Iron Co. in 2003 was unexpected.

Recognition of the ability of AHB to hybridize with EHB, which are better adapted to overwintering survival in cooler climates, has largely dispelled the notion that high elevations would be free of AHB. These hybrids exhibit the AHB mitotype inherited along their matrilines, and vet have inherited a number of EHB-derived genes that provide the requisite adaptations needed to survive harsh climates and long winters. Studies of AHB-EHB hybridization in the invaded neotropics have shown that honey bees in high elevation regions exhibit characteristics of both lineages. For example, the mitotypes of high elevation (2500-3500 m) honey bees in Peru are primarily AHB (56%), yet most (42%) exhibit intermediate phenotypes indicative of hybridization, rather than pure AHB (23%) or EHB (35%) morphology (Quezada-Euán, Pérez-Castro, & May-Itzá, 2003). However, high-elevation conditions in tropical regions are not equivalent to those in the temperate region, where winters are colder and longer, and the dynamics of hybridization may differ under these selection pressures (Spivak,

1992). In central Mexico, the mitotypes of highelevation (1800-2800 m) populations of honey bees are almost exclusively AHB (96%), yet only 59% of nuclear markers (microsatellites) are of African origin, indicating extensive hybridization not captured by mitotyping (Kraus, Franck, & Vandame, 2007). In the temperate regions of northern Mexico, EHB mitotypes are more common (75-88%) than AHB mitotypes, although AHB phenotypes are present in more variable proportions (14-79%) (Domínguez-Ayala et al., 2016; Medina Flores et al., 2017). Today, the prevailing hypothesis is that AHB can move up mountain slopes as they hybridize with their EHB counterparts but extensive tests have not been conducted in temperate regions of the USA. Africanized honey bees are now common in southern Utah, with half of samples collected between 2008-2017 testing positive for AHB (Cleary et al., 2018). However, our findings of multiple AHBs at locations above 1800 m in Utah including one at 2450 m in Iron Co. in 2003, tends to support range expansion into higher elevations of the temperate zone.

Finally, our study highlights the importance of museum collections and shows that even common species represent information that may become relevant to future research in unforeseen ways (reviewed in Suarez & Tsutsui, 2004). Unfortunately, increasingly underfunded and understaffed museum collections often, of necessity, ignore or even discard such specimens as relatively unimportant distractions in a kind of triage operation. Hopefully, this study will help both to dispel such notions and to demonstrate the importance of funding natural history collections: although active monitoring was taking place in Utah since the 1990s (Hodgson et al., 2010), AHBs went unrecorded for at least eight years.

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TRIPODI ET AL.

and highest elevation specimen; Christy Bills (UMNH), and Marshall Topham (AHBC) searched their respective collections but to no avail. They have our sincere thanks. We also extend our appreciation to Olivia Messinger-Carril and Harold lkerd for collection location clarifications. Dr. Tripodi contributed to this article in her personal capacity. The views expressed are her own and do not necessarily represent the views of the Agricultural Research Service or the United States Government.

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