Characterizing insect communities within thin-soil environments

Katherine McNamara Manning[1], Kayla I. Perry[2], & Christie A. Bahlai[1]

[1]Department of Biological Sciences, Kent State University, Kent, Ohio, USA
[2]Department of Entomology, Ohio State University, Wooster, Ohio, USA

Abstract

Natural thin-soil environments are those which have little to no soil accumulation atop

hard substrates. Many of these natural thin-soil environments, such as alvars, rocky lakeshores or

glades, cliffs and cliff bluffs, and barrens, are found in the Great Lakes Region of North

America. Due to their ubiquity and ecosystem services they provide, characterizing insects in

sensitive environments such as these is important. This study monitored insects in nine thin-soil

sites, within three regions, on a 630 km latitudinal gradient in the Southeastern Great Lakes

Region of North America from June - August 2019. Over 22,000 insect specimens collected

were identified to order or family, and bee specimens were identified to genus or species. We

found that overall insect community composition and biodiversity characteristics were similar

between the three regions examined. However, the central region had higher taxonomic richness

than the southern region. Although unique bee taxa were observed in each region, diversity

metrics and community composition of bees were similar among sites. This study provides

taxonomic information about the insect, particularly bees, and plant communities in thin-soil

environments in this region, which could support conservation and management efforts.

Introduction

Ecological communities are shaped by the physical attributes of their environments. In

the Great Lakes basin, numerous globally rare habitats occur, including those characterized by

their paucity of surface soil. The ecological class 'Primary' is characterized by having little to no

soil accumulation on top of bedrock, cobble, or exposed mineral soil. There are many natural

thin-soil community types, such as alvars, rocky (cobble or bedrock) lakeshores or glades, cliffs

and cliff bluffs, and barrens (Cohen et al. 2015). These habitats may experience intense wind and

solar radiation, as well as varying precipitation conditions, from heavy rain to drought, due to the

bedrock being at or near the surface (Stephenson and Herendeen 1986, Brunton 1988, Albert

2006). Together, these factors limit the primary producers that can survive there (Lundholm

2006), creating naturally open landscapes typically with low growing vegetation (Reschke et al.

1999). These unique and sensitive habitats are often home to rare plant and insect species

adapted to these unusual environments, making them areas of research interest and conservation

concern (Comer et al. 1997, Reschke et al. 1999, Albert 2006, Neufeld et al. 2018, McMullin

2019).

In the Great Lakes Region of North America, research in these rare thin-soil

environments, mostly alvars, has generally focused on identifying the taxa that occur in them or

are unique to them. Some of the characteristic rare vascular plants on Great Lakes alvars include:

Carex juniperorum W. J. Crins (Juniper Sedge), *Cirsium hillii* (Canby) (Hill's Thistle),

Cypripedium arietinum W. T. Aiton (Ram's Head Lady Slipper), and *Solidago houghtonii* A.

Gray (Houghton's Goldenrod), as well as *Iris lacustris* Nutt. (Dwarf Lake Iris) and *Tetraneuris*

herbacea Greene (Lakeside Daisy), both of which are endemic to the Great Lakes. Insect taxa

that have been known to show particular affinity for alvars include: carabid beetles, cicadellids,

lepidopterans, orthopteroids, and symphytes (Reschke et al. 1999). In alvar surveys in southern

Ontario, three species of ground beetles (Carabidae) were dominant, but rarely collected in other

non-alvar parts of the province: *Agonum nutans* (Say), *Chlaenius purpuricolliis* Randall, and

Pterostichus novus Straneo (Bouchard et al. 1998, 2005).

Insects play a variety of critical roles in ecosystem function and service, one of which is

pollination (Losey and Vaughan 2006, Noriega et al. 2018). Plant-pollinator relationships are

important for biodiversity in natural and managed ecosystems (Ollerton 2017, Wei et al. 2021).

Among insects, certain flies, wasps, beetles, butterflies, and moths perform pollination services,

however, bees are considered to be the most efficient insect pollinator (Potts et al. 2016). Both

wild and managed bee species are vital to at least 35% of crop production (Klein et al. 2007),

which is crucial for human food security (Ollerton 2017). Bees are a diverse group of insects,

with over 22,000 species worldwide. In North America, north of Mexico, there are over 4,000

bee species (Wilson and Messinger Carril 2016). However, in recent years, several high-profile

studies have reported mass declines of insect taxa around the globe (Fox 2013, Hallmann et al.

2017, Lister and Garcia 2018, Loboda et al. 2018, Seibold et al. 2019, Sánchez-Bayo and

Wyckhuys 2019, Wepprich et al. 2019, van Klink et al. 2020, Zattara and Aizen 2020). Bees are

not immune to this global loss of biodiversity, with reports of declines in managed

(vanEngelsdorp et al. 2008, Ellis et al. 2010, Steinhauer et al. 2014) and wild bee species (Colla

and Packer 2008, Grixti et al. 2009, Cameron et al. 2011, Scheper et al. 2014, Koh et al. 2016,

Arbetman et al. 2017, Graham et al. 2021). Because of their important roles as pollinators,

characterizing bee communities in different environments, such as thin-soil environments, is

valuable for management and conservation decisions.

Surveys of insects in Great Lakes thin-soil environments have largely focused on smaller

geographical regions, such as the intense sampling of alvars of Ontario, especially the Saugeen

(Bruce) Peninsula, and in Michigan near the northern shorelines of Lakes Michigan and Huron

(Albert et al. 1994, 1995, Reschke et al. 1999, Bouchard et al. 2005, Albert 2006, Cohen et al.

2015). Few surveys have occurred over the north-south expanses of this ecoregion, and none

have explicitly examined the communities of insects across these thin-soil habitats spanning the

region. Additionally, to our knowledge bee communities in these thin-soil environments have not

been characterized before. To address these knowledge gaps, we conducted a study of the

community composition of insects in thin-soil environments in three regions on a latitudinal

gradient across the Southeastern Great Lakes Region of North America. In this study, we

examined biodiversity metrics of insect communities, including bees as our focal taxa, between

northern, central, and southern regions of this area. We predicted that even if habitats and plant

communities differed, insect communities would be similar across these three regions because

the community will be shaped by the abiotic traits of the landscape.

Methods

Site descriptions

This study was conducted in thin-soil environments along a latitudinal gradient spanning

630 km in eastern North America (Figure 2). Along this gradient, three sites were selected on the

northern Bruce Peninsula in Ontario, Canada, three sites in Cuyahoga County, Northeast Ohio,

USA, and three sites in Hocking County, Southeast Ohio, USA (Figure 1). All sites had thin soils

(about 15 cm or less) and typically lacked direct canopy cover, leaving them relatively open and

exposed to solar radiation, winds, and ranging precipitation conditions. Permits to collect insects

and plants were obtained for all sites in accordance with landholder policy.

Northern sites were located on the Northern Bruce Peninsula, Ontario, Canada along the

Niagara escarpment and owned by The Nature Conservancy of Canada. All sites were

characterized as alvars, with little to no vegetation on limestone bedrock. The Davis alvar is

further defined as a dry lichen-moss open pavement alvar. It had the most open and flat

landscape, with sparse, low-growing vegetation. The Beaton alvar was a creeping juniper and

shrubby cinquefoil dwarf shrub alvar, with a slightly rocky landscape. The Cape Hurd Alvar was

similar to Beaton in vegetation and landscape, but closer to the lakeshore making it subject to

frequent low floods. All sites occurred in a rural landscape, primarily consisting of undeveloped

land, and protected areas, but the area had previously been subjected to heavy logging pressure.

Central sites were located in Cuyahoga County in Northeast Ohio, USA, specifically in

the municipalities of Walton Hills, Parma, and Highland Heights. All sites were embedded in a

greater landscape of mixed use, urban, industrial, and semi-urban land use histories. In Walton

Hills, we sampled a thin-soil mossy barren near a hiking trail, between a meadow and forest,

near a cliff edge over a creek in the Cleveland Metroparks Bedford Reservation. Cleveland

Metroparks also owned the site in Parma, which was a roadside hill with slate shale soil, with

sparse vegetation and trees. The Highland Heights Dusty Goldenrod Preserve is owned by the

West Creek Conservancy, Cuyahoga Soil and Water Conservation District, Friends of Euclid

Creek, and Euclid Creek Watershed Council. The portion of the preserve we sampled was an

open wet meadow in a forest, with tall vegetation, sparse trees, and is the only known home in

Ohio to its namesake, the rare and endangered Dusty Goldenrod (*Solidago puberula* Nutt.). This

site had thin soils due to a history of topsoil harvesting.

Southern sites were all located within the 520 hectare Crane Hollow Preserve in

Rockbridge, Southeast Ohio, USA, part of the unglaciated Allegheny Plateau. These sites

occurred in a highly rural private landscape with sparse human activity and development history.

These sites were located on cliff edges on top of deep ravines with a Black Hand sandstone

bedrock. The "W" ladder and picnic rocks sites were both named for spots along the "W" hollow

and Snyder was along Snyder hollow. All sites were open, xeric, thin-soil areas within forest,

dominated by mosses, lichens, and small shrubs. The two "W" sites were heavily covered in

Reindeer Lichen (*Cladonia* sp.) and American Wintergreen (*Gaultheria procumbens* L.).

Field and laboratory methods

The thin soils in these habitats constrained the methods used to monitor insect

communities, preventing typical pitfall sampling for ground-dwelling insects from being

deployed. From a conservation standpoint, disturbing natural thin soil environments is

undesirable, and from a practical standpoint, some of these sites were bedrock with almost no

soil. Due to these constraints, all sampling had to be done above the surface level. Insects were

surveyed using three types of passive sampling traps: yellow sticky cards (Pherocon, Zoecon,

Palo Alto, CA, USA), bee bowls (also known as pan traps, inspired by Leong & Thorp, 1999),

and ramp traps (ChemTica Internacional S.A., Santo Domingo, Costa Rica) (Figure 3), evenly

spaced at each site for 48 hours once per month in June, July, and August 2019. The spacing and

number of traps was dependent on the area available for sampling at the site (Table S1). Each

bee bowl was an array of three different colored bowls spray painted fluorescent yellow, blue, or

white (Krylon Industrial, Cleveland, OH, USA). Sticky cards and bee bowls were placed at the

height of the plant community to collect flying insects, and ramp traps were placed on the

surface of the ground to collect ground-dwelling insects. Bowls and ramps were filled with soapy

water (Dawn Original Liquid Dish Soap, Procter & Gamble, Cincinnati, OH, USA) and had

quart zipper-top bags filled with sand placed on top to minimize wind disturbance. Upon

collection, bowls and ramps were strained in the field and samples were placed in a gallon

zipper-top bag with 70% ethanol. Sticky cards were placed directly into gallon zipper-top bags.

Vegetation was sampled at the sites in July-August 2019 by placing a meter-squared

quadrat in a well-vegetated location within the site and identifying the vascular plants to species

or genus (det. John Gerrath, Joe Moosburgger, and Glenn Vande Water). The trees that directly

surrounded the sites, as well as prominent plant species that were not inside the quadrat were

also recorded (Table S2).

In the lab, the liquid samples were strained, and specimens were identified and placed in

vials with 70% ethanol for storage. The sticky cards were frozen, and specimens were identified

while remaining in the bag, afterwards they remained stored in the freezer. Specimens were

identified to order, suborder, superfamily, group ("wingless parasitoid wasps"), or family (Table

1). We identified bees (Hymenoptera: Apoidea: Anthophila) to the highest precision possible

(genus or species) using expert identification and various interactive Discover Life keys (det.

MaLisa Spring, http://www.discoverlife.org).

Statistical methods

All statistical analyses were completed using R 4.1.3 (R Core Team, 2022). Model fits

were evaluated for statistical assumptions of normality of residuals and homogeneity of variance.

Taxonomic richness (number of taxa per sample) and Shannon diversity index (Hill 1973) were

calculated using the *vegan 2.5-7* package (Oksanen et al. 2019).

Linear mixed effects models were developed using the *lme4* (Bates et al. 2015) and

lmerTest (Kuznetsova et al. 2017) packages to examine differences in insects among regions.

The response variables examined were taxa richness and Shannon diversity. Each model

included region (north, central, south), sampling date (date of trap collection), and trap type

(yellow sticky card, bee bowl, ramp trap) as categorical fixed effects and trap number nested

within the site as a random effect: *Response variable ~ region+Date+Trap+(1|Site:Replicate).*

The function 'Anova' from the car package (Fox and Weisberg 2019) was used to examine the

effect of region in each model. Tukey pairwise comparisons were performed using the *emmeans*

1.7.4-1 package (Lenth 2021) for all models to compare between regions (north, central, and

south).

To visualize the insect communities collected in each region we used non-metric

multidimensional scaling (NMDS, with Jaccard distance), computed using the *vegan 2.5-7*

package. For this analysis we used presence-absence data pooled by site for each sampling date.

Permutational multivariate analysis of variance (PERMANOVA), analysis of multivariate

homogeneity of group dispersions (BETADISPER), and pairwise multilevel comparison using

the adonis function were performed following the NMDS analysis to assess compositional

dissimilarity between regions. NMDS, PERMANOVA, and BETADISPER were computed

using functions in the *vegan 2.5-7* package. Pairwise adonis (pairwise PERMANOVA) was

performed using the *pairwiseAdonis* package (Martinez Arbizu 2020).

Accumulation curves of bees for each region were created using the *BiodiversityR*

package (Kindt and Coe 2005). To estimate sampling efficiency for each trap type, we used

nonparametric Jackknife order 1 estimator to compare observed and estimated richness. Linear

mixed effects models were used to examine differences in bees among regions, taking the form:

Response variable ~ region+Date+Trap+(1|Site:Replicate). The response variables examined

were bee taxa richness and Shannon diversity. Tukey pairwise comparisons were performed

using the *emmeans 1.7.4-1* package for all models to compare between regions.

Results

Three sampling periods at our nine sites yielded 252 samples: 72 from the north, 107

from the central, and 73 from the south. From these samples, we identified 22,459 specimens:

9,304 from the north sites, 11,107 from central, and 2,048 from the south. In the central and

southern regions, Diptera was the most abundant order. In the northern region, Hymenoptera was

the most abundant order, due to a single emergence event where 6,032 chalcid wasps were

collected, followed by Diptera (Table 1).

Traps at the southern sites produced the fewest unique taxa, followed by northern sites,

although the two regions did not differ statistically from each other in richness. The central sites

captured the most unique taxa, although this did not differ significantly from the richness

captured at northern sites. There were no differences detected for Shannon diversity between any

regions (Figure 4).

Non-metric multidimensional scaling (stress = 0.22) suggested strong overlap in insect

communities between northern, central, and southern regions at this taxonomic resolution

(Figure 5). When examined by sites within each region, the communities at all sites overlapped

one another (Figure S1). The PERMANOVA testing for differences between regions was

statistically significant (p = 0.001). Pairwise PERMANOVA found a statistically significant

difference between southern and central regions (p = 0.01).

Bees

Overall, there were 169 bees collected across all sites during our sampling period. We

identified 151 caught in the bee bowls and ramp traps, excluding 18 individuals caught by the

sticky cards as the glue damaged the specimens and obscured some identifying features. In the

bee bowls and ramp traps there were 26 taxa. Of the identified specimens, 14 individuals from 5

taxa were collected in the northern region, 88 from 19 taxa in central, and 49 from 16 taxa in the

southern. When compared with first order jackknife richness estimates, capture efficiency in the

northern region was 74%, 71% in the central region, and 63% in the southern region (Figure 6).

Region was not found to contribute significantly to bee community variation as there was no

difference in bee richness or Shannon diversity between regions (Figure S2).

The single most abundant species in the north was *Augochlorella aurata* (Smith), but the

genus *Lasioglossum* was more abundant overall. In the central region, *A. aurata* was the most

abundant species. In the southern region, *Lasioglossum versatum* (Robertson) was the most

abundant species, and there were 42 specimens in the genus *Lasioglossum* in total. One genus

was unique to the northern region: Megachilidae: *Heriades*. Eight taxa were unique to the central

region: Andrenidae: *Andrena*; Apidae: *Holcopasites*; Halictidae: *Agapostemon virescens*

Fabricius, *Halictus ligatus* (Say), *Halictus confusus* Smith, *Lasioglossum coeruleum*

(Robertson), *Lasioglossum tegulare* (Robertson), and *Lasioglossum zonulum* (Smith). Seven taxa

were unique to the southern region: Apidae: *Apis mellifera* L., *Ceratina calcarata* Robertson;

Halictidae: *Lasioglossum bruneri* (Crawford), *Lasioglossum cressonii* (Robertson),

Lasioglossum quebecense (Crawford), *Sphecodes*; Megachilidae: *Megachile*.

Plants

Regions varied substantially in plant species present. Overall, 53 vascular plant taxa were

identified across all our sites. There were 21 taxa identified in the northern region, 24 in the

central, and eight in the south. The only species recorded in more than one region was *Acer*

rubrum L. (Red Maple), found in the central and southern regions (Table S2). Red maple was

found at all three sites in the central region and two sites in the southern region.

Discussion

This study aimed to compare insect biodiversity characteristics and trends among thin-

soil environments across a latitudinal gradient in the Southeastern Great Lakes Region of North

America. Our findings suggest that there is strong similarity among the insect communities

within these habitats, at least when using typical sampling and community classification metrics.

We saw no differences in Shannon diversity between any regions when compared over a three

month span. However, the central region had greater insect taxa richness than the southern

region. Visualization with NMDS showed complete overlap of insect community in all three

regions, indicating similar community composition. The significant difference detected between

the central and southern regions with the pairwise PERMANOVA suggests these differences

were likely driven by the greater variability in community composition among sites in the

southern region, while sites in the central region were more similar to each other. Similar trends

were also observed when data were examined at the site level.

Bee analyses followed similar trends to the overall insect community analysis, as we

observed similar taxa richness and Shannon diversity among regions. However, there were

unique taxa collected within each region: one in the north, eight in the central, and seven in the

south. It is important to note that the traps in the northern region captured a very low number of

bees (14), compared to the central (87) and southern (49) regions, and between all the regions

specimen numbers were relatively low, which should be considered when interpreting these

results. However, there is strength in the higher resolution of identification of the bee specimens.

With these data, we can identify some apparent associations of bee taxa with regions and sites.

For example, *Augochlorella aurata*, a species in the family Halictidae, amounted to about 45%

of specimens collected in the central region. All but two of these individuals were captured at

one site, a sparsely vegetated hill with coarse slate and shale soil. *Augochlorella aurata* is a

primitively eusocial, common, generalist, ground-nesting species (Wilson and Messinger Carril

2016). This species was caught in the northern and southern regions as well, but in much lower

numbers. Two genera of cuckoo bee, brood parasites or kleptoparasites (Wilson and Messinger

Carril 2016), were captured during the study. Cuckoo bees lay their eggs in the nests of other bee

species, allowing their offspring to consume the resources meant for the host's offspring, whom

they typically kill (Wilson and Messinger Carril 2016, Litman 2019). Apidae: Nomadinae:

Holcopasites, which typically parasitizes bees in the subfamily Panurginae, was unique to the

central region. Halictidae: *Sphecodes*, which usually parasitizes other bees in their family, was

unique to the southern region. The genus *Lasioglossum*, in the family Halictidae, is the largest

genera of bees and notoriously difficult to identify (Gibbs 2018), however, 80% of *Lasioglossum*

specimens were identified to species in our study. Half of the bees captured in the northern

region were *Lasioglossum*, with three of the seven species collected belonging to the subgenera

Dialictus. Lasioglossum dominated the southern region, amounting to over 85% of bees

captured, most of these (43%) were *Lasioglossum (Dialictus) versatum. Lasioglossum versatum*

is a primitively eusocial, generalist, ground-nesting species (Michener 1966). This species was

captured at all three southern sites, on sandstone cliff bluffs. However, it was not unique to this

region as specimens were captured in the central region as well, mostly at the meadow site.

During the three months of our study, we captured and identified 151 bee specimens

belonging to 26 taxa overall. The northern region had the lowest bee richness and raw abundance

and the central region had the highest bee richness and raw abundance. Though the bee bowls

were the only trap targeting bees, we used the specimens collected in the ramp traps, targeting

ground-dwelling insects, as well. A bee monitoring study that took place on non-thin soil

environments in the county adjacent to our southern region sites (Washington County, OH,

USA) captured 2,753 specimens containing over 130 species using bee bowls and Malaise traps

in April-Oct 2013. When comparing the sampled species richness in only the bee bowls to

jackknife 1 estimates, the capture efficiency was 73% overall, ranging from 70-74% at their

sampling sites (Spring et al. 2017). On the Niagara peninsula, Ontario, Canada, between our

northern and central regions, 15,733 bee specimens containing 124 taxa were sampled in April-

Oct 2003 using pan traps (bee bowls), sweep and targeting netting. All three methods resulted in

an estimated 84% capture efficacy, using ACE estimates. Pan traps alone captured 96 species,

resulting in an estimated 75% capture efficiency (Richards et al. 2011). Although we collected a

relatively low abundance of bees compared to these more intensive studies, using jackknife 1

richness estimates, our capture efficiency was comparable at 69% in total, ranging from 63-74%

for the regions.

Overall, we captured few large-bodied bees, and notably no bumble bees, during the

study, but that is most likely a result of the bias of bee bowls to collect smaller bodied genera

(Wilson et al. 2008, Berglund and Milberg 2019, Portman et al. 2020). In future studies, adding

hand-netting as a complementary sampling method could produce increased richness across sites

and increase the comparability of bee taxa among these regions (Grundel et al. 2011, Prendergast

et al. 2020). For example, Grundel et al. (2011) sampled bee communities in a variety of habitats

in Indiana Nature Preserves using bee bowls and hand-netting techniques. They reported high

abundances of larger species such as *Bombus impatiens* Cresson (Common Eastern Bumble Bee)

and *Apis mellifera* L. (European honey bee) captured with hand nets and smaller-bodied families

such as Halicitidae (sweat bees) captured in the bee bowls. We only captured one *Apis mellifera*

during the study, but this is a common introduced, managed species found over most of the

world (Butz Huryn 1997), and in abundance in other studies in the Great Lakes region (Spring et

al. 2017, Rowe et al. 2020, Turo et al. 2021). Future investigators should consider identifying

other biologically important taxa, such as beetles, to a higher resolution as other groups of

insects may oy may not follow similar patterns as bees if identified to species.

The Davis alvar performed slightly differently from the other northern sites, as we

observed lower Shannon diversity (Figure 3) and higher raw insect abundance. These differences

may be attributable to large numbers of a parasitoid wasp (Trichogrammatidae) which emerged

during our 12-14 August sampling period. Trichogrammatidae (superfamily Chalcidoidea) are

known to parasitize eggs from several insect orders. Some genera, especially Trichogramma, are

used for biological control of pest insects (Jalali et al. 2016). We captured just over 6,000

individuals that all appeared to belong to the same Trichogrammatidae morphological group, at

only this site. When this emergence is removed from the data, the total number of Chalcidoidea

collected in the northern region is much closer to that of the southern region (178 and 179,

respectively). However, our analyses detected nearly identical patterns in comparisons of taxa

richness and Shannon diversity with or without emergence data included. Without the

emergence, the Davis alvar still had the greatest raw abundance of insects of the northern sites.

Presence-absence data was used in the NMDS analysis so this emergence event would not

influence the analysis of insect community composition.

From our observations, plant communities differed between regions with all three regions

producing nearly unique species lists. There were no plant taxa present in all three regions. The

only species we saw in more than one region was red maple, which was present in the central

and southern regions. *Clinopodium arkansanum* (Nutt.) House (Limestone Calamint) was found

at all three sites of the northern region, but absent from the central and southern regions. This

species is associated with lakeshores and has been commonly found in flat rocky alvars near

Lakes Huron and Michigan (Voss and Reznicek 2012, Cohen et al 2015). *Gaultheria*

procumbens (American Wintergreen) and *Pinus virginiana* (Virginia Pine) were present at all

three southern sites but absent from the central and northern regions. American wintergreen

prefers sandy, acidic soils (Homoya and Namestnik 2022), which the sandstone soils of the

southern region provide. Virginia pines are often found in soil derived from sandstone and their

range overlaps with only the southern region (Snow 1965). The southern region, with its xeric,

moss and lichen covered sandstone soils supported the lowest vascular plant richness.

Though the plant communities were unique in each region, we found overall similar

insect community structure, perhaps due to their shared structural characteristic of thin soils. In

general, insect communities were more similar in the north and central regions, with the southern

region having lower insect taxa richness than the central region. Insect richness is often

correlated with plant richness (Haddad et al. 2001), and we found the lowest vascular plant

richness in the southern region as well. In this study, most insects other than bees were identified

to order or family level which could have obscured some community patterns, and this caveat

should be kept in mind when interpreting the findings.

The northern and central sites represent very different recent land-use histories but have

the common history of both being within the extent of glaciation. The former two sites are

situated well within the Great Lakes basin and the latter site is situated at the edge of this region,

as it transitions into the Allegheny plateau. The history and severity of human disturbance at the

sites also have the potential to drive insect community characteristics. One difference between

the regions is their vehicular and pedestrian access and proximity to urban areas. The sites of the

central region are proximate to urban areas while northern or southern region sites are rural and

are on privately held nature preserve property. The central sites are open to the public, however

not in areas typically with high foot traffic. Human disturbance has been well-established to

affect insect abundance and diversity (Winfree et al. 2009, Quintero et al. 2010, Owens et al.

2020, Fenoglio et al. 2021, Raven and Wagner 2021, Wagner et al. 2021), with cities providing

refuge to some insects, including pollinators (Hall et al. 2017). Though the central sites were

located within preserved areas, those sites are still within the urban matrix of the greater

Cleveland, Ohio area and have a history of human disturbance. Hemiptera was the second most

abundant order in the central region due to the large number of aphids, a common crop pest,

captured (Miller and Foottit 2009, Emden and Harrington 2017). This and the higher overall

numbers of insects collected in the central region could be a result of previous disturbance.

Thin-soil environments, while considered important rare habitats, may also serve a more

utilitarian role as model systems. These habitats have been studied as biotic and abiotic templates

for urban green, or vegetated, roof design because of their structural similarities (Lundholm

2006, Lundholm and Walker 2018). Extensive green roofs are a type of living architecture in

which plants are intentionally grown on top of a human-built structure in shallow (typically 15

cm or less) growing medium (Getter and Rowe 2006). Like the natural thin-soil environments,

they may face similar weather patterns as they are in an exposed area: varying precipitation

conditions, intense solar radiation, and wind. Thus, our expectations for biotic function in these

environments should too be modeled after natural systems with similar traits. Green roofs have

the potential to provide urban areas with many services, including stormwater retention, reduced

building energy consumption, habitat for organisms, and more depending on how it is designed

(Dunnett and Kingsbury 2004). Plants and other physical structures from natural thin-soil

environments may be used to design green roofs that feature native plants and landscapes similar

to those found outside of the urban area (Lundholm 2006, Lundholm and Walker 2018). These

habitat patches may aid in conservation of plants and insects found in those natural ground-level

environments and increase certain ecosystem services, such as pollination and pest control, for

the city (Kadas 2006, Colla et al. 2009, MacIvor and Lundholm 2011).

Conclusions

Thin-soil environments, characterized by thin or absent soils atop hard substrates, are

unique and globally rare environments (Reschke et al. 1999, Cohen et al. 2015, Neufeld et al.

2018). Due to the nature of their soils, these open habitats can be quite harsh from an abiotic

standpoint and thus home to unique communities of plants and animals (Albert et al. 1995,

Comer et al. 1997, Reschke et al. 1999, Neufeld et al. 2018, McMullin 2019). While previous

work has examined the composition of these habitats (Albert et al. 1994, 1995, Comer et al.

1997, Bouchard et al. 1998, 2005, Reschke et al. 1999, Albert 2006, Cohen et al. 2015, Neufeld

et al. 2018, McMullin 2019), in this study, we examined how these environments shape

similarities in insect communities. This study leverages a unique design that encompasses a large

area of the Southeastern Great Lakes Region, spanning over 600 km north to south.

By examining the plant and insect communities in three separate regions of this area of

North America, we found that overall insect community composition and biodiversity

characteristics were similar. Though some variation in taxa richness was detected, Shannon

diversity was similar across all three regions. The southern region had the lowest insect taxa

richness and the lowest vascular plant richness. All regions contained unique plant taxa, with

only one species being found in more than one region. Bee identity was similar across all

regions, finding no differences in biodiversity metrics and similar community composition,

though each region contained unique taxa.

This study provides taxonomic information about the insect, particularly bee, and plant

communities in thin-soil environments in this region. This knowledge supports conservation and

management efforts of rare thin-soil habitats such as alvars. Furthermore, this study provides

insights into how communities of insects assemble in habitats with these structural

characteristics. If these habitats are modeled by architects or urban planners for green roofs or

other urban infrastructure, insights can be used to create realistic expectations for biodiversity

outcomes in the built environment.

Data and code: https://github.com/katiemmanning/Thin-soil

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Figures

Figure 1: Sampling sites in each region. North: (A) Davis alvar, (B) Beaton alvar, (C) Cape hurd alvar,

Central: (D) Slate shale hill, (E) Dusty goldenrod meadow, (F) Bedford barren, and South: (G) Snyder, (H)

The "W" – ladder, (I) The "W" – picnic rock.

Figure 2: General location of sampling sites in each region. Figure adapted from Google Maps.

Figure 3: Sticky cards, bee bowls, and ramp traps (left to right) deployed at various field sites in the
Southeastern Great Lakes Region in 2019.

Table 1: Insect abundances, total (T) and standardized (S), by region. Standardized abundances were

calculated by dividing total abundances for each taxon by the total number of traps used in that region.

Order		North		Central		South	
		T	S	T	S	T	S
Blattodea		0	0	1	0.01	0	0
Diptera		2510	34.86	5334	74.08	1142	15.86
	Syrphidae	163	2.26	316	4.39	21	0.29
Ephemeroptera		5	0.07	0	0	0	0
Hymenoptera		6280	87.22	687	9.54	591	8.21
	Anthophila	15	0.21	99	1.38	54	0.75
	Chalcidoidea	6189	85.96	433	6.01	179	2.49
	Chrysididae	0	0	0	0	1	0.01
	Ichneumonoidea	41	0.57	104	1.44	76	1.06
	Dryinidae	0	0	1	0.01	0	0.00
	Formicidae	28	0.39	35	0.49	272	3.78
	Pompilidae	2	0.03	0	0	0	0
	Unknown wasp	2	0.03	15	0.21	9	0.13
	Wingless parasitoid	3	0.04	0	0	0	0
Hemiptera		274	3.81	4681	65.01	127	1.76
	Aleyrodidae	2	0.03	133	1.85	11	0.15
	Aphididae	148	2.06	4317	59.96	13	0.18
	Cercopidae	0	0	7	0.10	1	0.01
	Cicadellidae	110	1.53	163	2.26	70	0.97
	Fulgoroidea	0	0	3	0.04	0	0
	Membracidae	0	0	3	0.04	11	0.15
	Miridae	1	0.01	1	0.01	0	0
	Orius	0	0	1	0.01	0	0
	Psyllidae	7	0.10	24	0.33	2	0.03
	Tingidae	0	0	1	0.01	0	0
	Unknown Hemipteran	6	0.08	28	0.39	19	0.26
Coleoptera		235	3.26	404	5.61	188	2.61
	Buprestidae	0	0	0	0	2	0.03
	Cantharidae	5	0.07	18	0.25	1	0.01
	Carabidae	1	0.01	1	0.01	0	0
	Chrysomelidae	6	0.08	48	0.67	2	0.03
	Coccinellidae	0	0	1	0.01	1	0.01
	Curculionidae	2	0.03	17	0.24	4	0.06
	Elateridae	2	0.03	1	0.01	0	0
	Lampyridae	25	0.35	1	0.01	1	0.01
	Meloidae	1	0.01	0	0	1	0.01
	Mordellidae	8	0.11	25	0.35	14	0.19

	Scarabaeidae	1	0.01	3	0.04	5	0.07
	Staphylinidae	20	0.28	4	0.06	32	0.44
	Unknown Coleoptera	21	0.29	36	0.50	18	0.25
Lepidoptera		38	0.53	17	0.24	1	0.01
Lepismatidae		0	0	0	0	1	0.01
Mecoptera		2	0.03	0	0	0	0
Neuroptera		0	0	5	0.07	0	0
Orthoptera		13	0.18	18	0.25	12	0.17
Psocodea		0	0	2	0.03	12	0.17
Symphyta		2	0.03	0	0	0	0
Thysanoptera		79	1.10	207	2.88	80	1.11
Trichoptera		9	0.13	0	0	1	0.01

Figure 4: Box plots displaying taxa richness and Shannon diversity of insects, comparing sites by region. Letters shared indicate no statistical difference in estimated marginal means by Tukey method, P <0.05.

Figure 5: Non-metric multidimensional scaling figure (stress = 0.22) representing insect communities by region: North, Central, and South.

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Figure 6: Accumulation curves for bee richness in each region. Shading represents standard deviation from the mean.