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TITLE:

A Method for Quantifying Foliage-Dwelling Arthropods

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SUMMARY:

We describe how to quantify leaf dwelling arthropods by sealing the leaves and end of branches in a bag, clipping and freezing the bagged material, and rinsing the previously frozen material in water to separate arthropods from the substrate for quantification.

ABSTRACT:

Terrestrial arthropods play an important role in our environment. Quantifying arthropods in a way that allows for a precise index or estimates of density requires a method with high detection probability and a known sampling area. While most described methods provide a qualitative or semi-quantitative estimate adequate for describing species presence, richness, and diversity, few provide an adequately consistent detection probability and known or consistent sampling areas to provide an index or estimate with adequate precision to detect differences in abundance across environmental, spatial, or temporal variables. We describe how to quantify leaf-dwelling arthropods by sealing the leaves and end of branches in a bag, clipping and freezing the bagged material, and rinsing the previously frozen material in water to separate arthropods from the substrate and quantify them. As we demonstrate, this method can be used at a landscape scale to quantify leaf-dwelling arthropods with adequate precision to test for and describe how spatial, temporal, environmental, and ecological variables influence arthropod richness and abundance. This method allowed us to detect differences in density, richness, and diversity of leaf-dwelling arthropods among 5 genera of trees commonly found in southeastern deciduous forests.

INTRODUCTION:

Terrestrial arthropods play an important role in our ecosystem. In addition to being of scientific interest arthropods can be both detrimental and beneficial to crops, horticultural plants, and natural vegetation as well as provide an important trophic function in food webs. Thus,

understanding the factors that influence arthropod community development and abundance is critical to farmers, pest control managers, plant biologists, entomologists, wildlife ecologists, and conservation biologists that study community dynamics and manage insectivorous organisms. Understanding factors that influence arthropod communities and abundances often requires the capture of individuals. Capture techniques can generally be categorized into qualitative techniques that only detect presence of a species for estimates of species range, richness, and diversity, or semi-quantitative and quantitative techniques that allow for an index or estimate of abundance and density of individuals within a taxonomic group.

Qualitative techniques that only allow inference regarding presence of a species or community structure have an unknown or intrinsically low detection probability or are lacking in providing inference regarding detection probability and size of area sampled. Because detection probability with these techniques is low, variability associated with detection precludes adequate precision for inferring how explanatory variables influence arthropod population metrics. Qualitative techniques used to estimate presence include suction sampling¹, light traps², emergence traps³, feeding patterns on roots⁴, brine pipes⁵, baits⁶, pheromone³, pitfall traps⁷, Malaise traps⁸, window traps⁹, suction traps¹⁰, beating trays¹¹, spider webs¹², leaf mines, frass¹³, arthropod galls¹⁴, vegetation and root damage¹⁵.

Alternatively, semi-quantitative and quantitative techniques allow researchers to estimate or at least consistently sample a specified sample area and estimate probability of detection or assume detection probability is non-directional and adequate as to not obscure the researcher's ability to detect spatial or temporal variation in abundance. Semi-quantitative and quantitative techniques include sweep nets¹⁶, suction or vacuum sampling¹⁷, systematic counting of visible arthropods¹⁸, sticky traps¹⁹, various pot-type traps²⁰, entrance or emergent holes²¹, chemical knockdown²², sticky and water filled color traps²³, and branch bagging and clipping²⁴.

Recent anthropogenic-induced changes to climate and disturbance regimes have led to dramatic changes in plant communities, making interactions between plant-community species composition and arthropod communities an active area of study. Understanding how arthropod communities vary with plant species composition is a critical component for understanding the potential economic and environmental impacts of changes to plant communities. Semi-quantitative or quantitative methods of quantifying arthropod abundance with adequate precision to detect differences among species of plants are needed. In this article, we describe a method for indexing foliage-dwelling arthropods that, with reasonable effort, provided adequate precision to identify differences in individual abundance and biomass, diversity, and richness among 5 taxa of trees commonly found in the southeastern deciduous forests of North America²⁵. This approach provided precision adequate for estimating abundance to allow inference as to how changes in species composition of forest plant communities due to anthropic modified disturbance regimes influence composition of arthropods, potentially influencing abundance and distribution of higher trophic insectivorous birds and mammals. More specifically, by using a modified bagging technique first described by Crossley et al.²⁴, we estimated density of surface, foliage-dwelling arthropods and tested the prediction that we would detect differences in diversity, richness, and abundance of arthropods in the foliage of faster growing more xeric

species of trees relative to slower growing more mesic species. The goal of this article is to provide detailed instructions of the technique.

We conducted the study on the Shawnee National Forest (SNF) in southern Illinois. The SNF is a 115,738-ha forest located in the Central Hardwoods region of the Ozarks and Shawnee Hills natural divisions²⁶. The forest comprises a mosaic of 37% oak/hickory, 25% mixed-upland hardwoods, 16% beech/maple, and 10% bottomland hardwoods. The SNF is dominated by second growth oak/hickory in upland xeric areas and sugar maple, American beech, and tulip tree (*Liriodendron tulipifera*) in sheltered mesic valleys^{27,28}.

Site selection for this method will be dependent on the overarching goals of the study. For example, the primarily goal of our original study was to provide insight into how changes in tree community might influence higher trophic organisms by comparing foliage-dwelling arthropod community metrics between mesic and xeric adapted tree communities. Thus, our primary objective was to quantify the arthropod community on individual trees located within the xeric or mesic tree community. We selected 22 study sites along an oak/hickory (xeric) to beech/maple (mesic) dominated gradient using USFS stand cover maps (allveg2008.shp) in ArcGIS 10.1.1. To prevent potential confounding effects, we selected sites using the following criteria: not located in riparian areas, ≥ 12 ha, and located within contiguous upland-deciduous forest habitat (i.e., elevation above 120 m). All sites contained mature trees >50 years old in hilly terrain, thus comprised similar slopes and aspects. While beech/maple site boundaries were distinguished based on the transition of tree communities, oak/hickory site boundaries were identified artificially using SNF cover maps and ArcGIS 10.1.1. All sites were large forest blocks within unglaciated terrain; their differences in tree species composition were not due to differences in location on the landscape but were representative of past land usage (e.g., clear cuts or selective harvest). We ground-truthed the maps by uploading discrete polygon shapefiles of each study site to a handheld Global Positioning System (GPS) and verifying tree species composition. We randomly selected sampling points ($n = 5$) at each site. At each point, we sampled three trees from 0600–1400 hours during 23 May to 25 June 2014. To locate sample trees, we searched outward to a 30 m radius from vegetation points until mature trees (>20 cm d.b.h.) with branches low enough to sample were found. Typically, the three mature trees that represented three of the five genera (*Acer*, *Carya*, *Fagus*, *Liriodendron*, and *Quercus*) of interest and were closest to the center point were sampled.

PROTOCOL:

1. Building the sampling device prior to going to the field

1.1. Using bolt cutters, large wire cutters, or an electric grinding disk, remove the bottom 1/3 of the 30 cm wire tomato cage so that it is approximately 55 cm in length.

1.2. Cut two, 30 cm braces made from aluminum tube or similarly semi-rigid material to use as attachment rods and braces on each side of the largest end of the tomato cage. Attach one end of each of the two attachment rods to opposite sides of the tomato cage with zip ties and duct

or electrical tape ensuring the tape is wrapped around at least 6 cm of the cage and rod. Be certain to wrap the tape around the cage and rod numerous times to ensure the cage is permanently attached to the rod.

1.3. Attach the other end of each of the two attachment rods on the opposite sides of the end of an extendable pole with zip ties and duct or electrical tape. As before, wrap the tape multiple times to affix it permanently ensuring the tape overlaps the pole and rods by at least 6 cm. Be certain the opening of the cage is in contact with the end of the telescoping pole when the cage is attached.

1.4. Attach the cage directly onto the end of the pole using zip ties and electrical or duct tape. Attach hook-and-loop fastener strips at 3 points to the opening of the cage.

NOTE: These strips will be use later to keep the bag open.

2. Enclosing the branch

2.1. Attach 3 pieces of hook-and-loop fastener to the opening of the bag so they align with the hook-and-loop fastener attached to the opening of the cage. These will be used to hold the opening of the bag in place while it is brought over a sample branch. Be certain the hook-and-loop fastener is aligned so when the bag is inserted and attached, the opening to the pull strings of the bag run parallel to the telescoping pole.

2.2. Insert a ~49 L kitchen garbage bag in the wire tomato cage. Place one gator clip on each respective side of the bottom of the bag and attach the clips to both the bag and wire cage to hold the bag against the cage. Repeat the same procedure for the top of the bag with gator clips and attach the opening of the bag to the opening of the cage.

2.3. Orient the gator clips so they are perpendicular to both the telescoping pole and the openings to the draw strings as to run parallel with the telescoping pole. Be certain to leave adequate space near the opening of the bag to allow the bag to close when the draw string is pulled.

2.4. Attach para cord to each of the bag's two draw strings. Cut plastic tubing in one-inch sections and attach with duct or electrical tape at three equidistance locations (i.e., one each along the top, middle and bottom) along the non-extendable outside of the telescoping pole. Run both para cord strings together through the plastic tubing inserts to hold them in place.

2.5. For each sample tree, use a random number generator to select a sample height that is within the height of the extension pole when extended at maximum length. Use a random number generator to select a sample distance from the tree trunk. Identify a branch that will fit in the bag with minimal disturbance to the foliage and is the height and distance from the trunk based on the numbers generated from the random number generator.

2.6. Raise the sampling pole to a height parallel with the desired branch. Quickly slide the bag over the branch then rapidly pull the para cord strings attached to the draw strings on the bag to seal the bag. Practice this a few times prior to the first attempt to become efficient at incorporating the foliage with minimal disturbance to the leaves.

2.7. Have a second person clip the branch at the location adjacent to the bag's opening with the extension pole pruner. Carefully bring the sample bag to the ground and rapidly tie the bag's draw strings closed. Attempt to complete the bagging, cutting, and bag-tying steps as quickly as possible to prevent insects from escaping.

2.8. Store the bagged branch in a freezer until ready to conduct the laboratory arthropod analysis.

3. Arthropod analysis

3.1. Hold the frozen bag and branch upright and shake the sample branch while in the bag to dislodge arthropods into the bag. Carefully remove the branch and rinse in large collection pan to remove remaining arthropods. Empty remaining material from the bag into the collection pan. Remove any non-arthropod debris.

3.2. Separate arthropods into desired taxonomic groups. Note differences between larvae and adults.

3.3. Quantify arthropods as desired. If biomass is of interest, either measure length of arthropods and use published length mass table to estimate biomass, or place arthropods in small drying pans, dry in drying oven for 24 h at 45 °C, and weigh on an electronic balance.

4. Estimating density

4.1. To estimate density and control for variation in leaf structure and leaf density between samples within tree species and among tree species either:

4.1.1. Count and measure the surface area of the leaves from each sample.

4.1.2. Dry the leaves in a drying oven for 48 h at 45 °C and weigh the leaves on an electronic balance.

4.1.3. Measure the length of all woody branch within the sample.

NOTE: Diel differences occur in arthropod communities, so sampling should be conducted throughout the entire period of inference.

REPRESENTATIVE RESULTS:

We collected 626 samples from 323 individual trees composing 5 tree groups. For estimates of total arthropod biomass per meter of branch sampled, the standard error ranged from 12% to 18% of the mean for the 5 tree groups (**Table 1**). This level of precision was adequate to detect variation among tree groups and a quadratic change in biomass with date²⁵. This technique provided more precision when estimating guild diversity as demonstrated by the standard error of arthropod guild diversity (H') ranging from 3% to 7% of the mean diversity across the 5 tree groups (**Table 1**). Precision at this level was adequate to detect variation across the 5 tree groups²⁵. Precision of estimates of richness was also very good as demonstrated by standard errors that ranged from 3% to 6% of the mean richness among the 5 tree groups (**Table 1**). This level of precision was adequate to identify variation among tree groups, a quadratic association with date, decrease in richness with height on the tree, and a positive relationship between arthropod richness and distance from the tree trunk²⁵.

TABLE LEGENDS:

Table 1: Parameter estimates from most parsimonious model²⁵. The mean (X), standard error of the mean (SE), and percentage of the mean of the standard error for each community metric of foliage dwelling-arthropods captured on 5 groups of trees using the described branch clipping method in the Shawnee National Forest in southern Illinois.

DISCUSSION:

Two necessities of accurately quantifying arthropod communities are relatively high detection probabilities and known or consistent sampling areas. When sampling for arthropods, less than 100% detection probability can be attributed to either individual arthropods avoiding traps or some individuals that were trapped being undetected during processing. Interceptor traps that intercept flying arthropods (Malaise/window traps, sticky traps, etc.) appear to be the most frequently used approach to enumerate arthropod communities in the forest canopy^{29,30,31}. These types of traps can be placed throughout the canopy, are effective at intercepting flying arthropods, and typically preserve arthropods for long periods (weeks or months) for later identification and quantification^{29,30,31}, though they are typically limited in their ability to trap crawling arthropods³¹. Interceptor traps that attract arthropods using light or pheromones have additional limitations in that they trap only night flyers and their attractiveness varies with taxon, moonlight, background illumination, and cloud cover impact^{32,33}. Additionally, because arthropods captured in interceptor traps are from unknown distances, the area trapped is unknown. As such, although interceptor traps are effective for indexing flying arthropods across an environmental gradient, data produced from interceptor traps cannot be used to estimate arthropod density²⁵.

An additional method frequently used to monitor foliage arthropods is chemical knockdown^{34,35}. Chemical knockdown can be very effective for collecting diverse groups of arthropods providing accurate estimates of taxonomic richness and diversity. However, this method is expensive and time consuming, is non-specific as it samples all arthropods on the tree including those on the bark and branches, may have unintended environmental impacts due to wind drift, and is illegal in some areas^{36,37,38,39}.

Branch bagging has been demonstrated as an effective method to estimate arthropod density from surface tree foliage with an adequately high capture probability to detect variation across various environmental gradients^{24,40}. The wire tomato cages and 49 L garbage bags used in this study allowed researchers to encompass the branch fully with little to no disturbance prior to the closure of the opening of the bag. As such, it is important that researchers are careful to not disturb foliage of the desired branch sample prior to enclosing it with the sampling bag. Thus, a critical step is to bring the sampling bag parallel with the desired sampling bag and swiftly enclose, seal, and tie the bag after each sample is collected. Sample collection is limited to the maximum height the researcher can hold an extended telescoping pole at (8 m in our study), although the same branch-bagging equipment and methodology can be used in other situations such as suspension in the canopy. Some authors have suggested that when using this procedure active, flighted-arthropods are underrepresented^{40,41,42}. However, we believe that as long as the foliage remains undisturbed until it is enclosed by the sampling bag, it is unlikely that a substantial number arthropods present in or on the foliage at the time escaped capture. The results of our study support this assertion in that when sampling a reasonable number of trees (323), the standard error was at most 17% of the relative arthropod biomass mean (*Cayra* = 11%, *Acer* = 12%, *Fagus* = 17%, *Liriodendrum* = 15%, and *Quercus* = 11%). Similarly, when considering guild richness and diversity, the most variable estimate was diversity on *Fagus*, with a standard error that was 7% of the mean. Clearly these estimates provided adequate precision to model differences among tree genera groups as well as other ecological or environmental variables. A limitation to our results, however, is that although we are confident that detection probability with this method is high, i.e., likely nearing 100%, we do not have a method of independently verifying this assertion. Thus, while we demonstrated detection probability is adequate to detect variation across an environmental variable, which in this case was tree genera, biomass estimates produced from this methodology have the potential to be biased low by some unknown amount⁴⁰.

Most authors have examined the content of the bag in the field^{36,42,43,44,45}. We believe a critical step to maximizing detection is to freeze the bag as we did, then examine and quantify the content in the laboratory under controlled conditions. We believe this approach will decrease measurement error by minimizing the number of trapped arthropods that are overlooked or are misidentified.

Estimating the area sampled for comparison of density among tree species may be problematic if leaf structure varies considerably among tree species, as was the case in our study. In past studies, when authors were interested in quantifying foliage-dwelling arthropods, they often estimated sampling area by weighing leaves to estimate the amount of substrate available for arthropods^{46,47,48}. The various species of oak trees, however, tend to have thicker waxy leaf cuticles than other tree species. Thus, the mass to surface area ratio for oaks is greater than other species⁴⁹. Because the mass to surface area ratio is greater in oaks, using the mass of leaves as an estimate of substrate for foliage dwelling arthropods would overestimate sampling area and underestimate arthropod density for oak trees relative to tree species with less thick leaf cuticles. Additionally, if the ability to support arthropods varies among tree species, the surface area of

the landscape covered by a given tree species will dictate the level of substrate supported within a specified landscape. Because the amount of surface area a given tree occupies is determined by crown spread (i.e., branch spread outwards from the trunk) and leaf density varies among trees, we believe when quantifying arthropods for consumption by insectivores, total branch length sampled is more appropriate than leaf biomass when estimating total area sampled. Our results again appear to support this assertion in that we detected differences among tree groups consistent with the predicted pattern based on previous studies²⁵. We believe arthropod abundance or biomass per measure of branch length is most appropriate when the primary objective is to compare resources provided for insectivores among tree species. If, however, individuals are comparing tree species that produce leaves with similar leaf cuticle thickness, using leaf biomass as an estimate of sampling area may be more appropriate. Regardless of whether researchers use actual leaf area, leaf area as estimated by leaf biomass, or total branch length as a quantifiable metric, by using the bagging technique, a measurable quantity of arthropods at a specific point in time on a measurable surface area is captured per sample. This allows researchers to use leaf surface area, leaf area as estimated by leaf biomass, or total branch length as a quantifiable metric. This method provides a consistent estimate for comparing quantified arthropods among spatial or temporal variables and an estimate of arthropod density²⁵.

In general, the sampling method described in this article appears to be effective at allowing for spatial or temporal comparisons of foliage-dwelling arthropod metrics. This approach is affordable and feasible at the landscape scale. Furthermore, although freezing the entire branch requires substantial freezer space, freezing the branch then rinsing the branch in water is an effective way to separate arthropods from foliage with minimal effort, therefore providing a cost-efficient approach to obtaining arthropod metrics. Finally, because the primary objective of our original study was to better understand how mesophication of southeastern deciduous forests is likely to impact forest-dwelling insectivorous birds and mammals we grouped arthropods into guilds based on diagnostic morphological features. However, we do not see a reason why these capture techniques cannot be used to quantify arthropods at the species or any other taxonomic level.

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DISCLOSURES:

The authors have nothing to disclose.

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Table 1.

Tree species	Richness			Biomass			Shannon Diversity		
	X	SE	% of mean	X	SE	% of mean	X	SE	% of mean
Maple spp. N = 140	3.54	0.17	5%	0.003	0.0004	13%	0.86	0.05	6%
Hickory spp. N = 141	4.62	0.20	4%	0.013	0.002	15%	1.10	0.04	4%
Tulip Poplar N = 70	4.32	0.20	5%	0.011	0.002	18%	1.12	0.05	4%
American Beach N = 67	3.23	0.22	7%	0.002	0.0003	15%	0.81	0.06	7%
Oak spp. N = 208	4.77	0.15	3%	0.006	0.0007	12%	1.10	0.03	3%

	Richness			Biomass			Shannon Diversity		
	X	SE	% of mean	X	SE	% of mean	X	SE	% of mean
Tree species									
Maple spp. (N = 140)	3.54	0.17	5%	0.003	0.0004	13%	0.86	0.05	6%
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Oak spp. (N = 208)	4.77	0.15	3%	0.006	0.0007	12%	1.10	0.03	3%

Name of Material/Equipment	Company	Catalog Number	Comments/Description
13 gallon garbage bags	Glad	78374	
Aluminum rod	Grainger	48ku20	
Pruner	Bartlet arborist supply	pp-125b-2stick	
Telescoping pole	BES	TPF620	
Tomato Cage	Gilbert and Bennet	42 inch galvanized	



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Author(s):

Eichholz & Sierzega

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Thank you for allowing us the opportunity to revise our manuscript. Our responses can be found following the comments below.

1. Please take this opportunity to thoroughly proofread the manuscript to ensure that there are no spelling or grammar issues. The JoVE editor will not copy-edit your manuscript and any errors in the submitted revision may be present in the published version.

Done

2. Please revise lines 64-67 and 73-86 to avoid textual overlap with previously published work.

Text on lines 64-67 revised from:

“We conducted our study in the forested, un-glaciated landscape of southern Illinois throughout the Shawnee National Forest (SNF). The 286,000ac SNF is located in the Central Hardwoods region within the Ozarks and Shawnee Hills natural divisions²⁶. Current forest composition is a mosaic of 37% oak/hickory, 25% mixed-upland hardwoods, 16% beech/maple, and 10% bottomland hardwoods and is dominated by second growth oak/hickory in the uplands and sugar maple, American beech, and tulip tree (*Liriodendron tulipifera*), in sheltered mesic valleys^{27,28}.”

To:

We conducted the study on the Shawnee National Forest (SNF) in southern Illinois. The SNF is a 115,738 ha forest located in the Central Hardwoods region of the Ozarks and Shawnee Hills natural divisions²⁶. The forest comprises a mosaic of 37% oak/hickory, 25% mixed-upland hardwoods, 16% beech/maple, and 10% bottomland hardwoods and is dominated by second growth oak/hickory in upland xeric areas and sugar maple, American beech, and tulip tree (*Liriodendron tulipifera*), in sheltered mesic valleys^{27,28}.”

Lines 71-86 have been revised from:

“To ensure the factors influencing the arthropod community described on an individual tree was limited to those within our scope of interest, we selected 22 study sites along an oak/hickory dominance gradient by identifying either oak/hickory or beech/maple dominated sites with the USFS stand cover map for the SNF (allveg2008.shp) in ArcGIS 10.1.1 (ESRI, Redlands, CA, USA). We used the following criteria for site selection to prevent potential confounding effects: located within contiguous upland-deciduous forest habitat (i.e., elevation above 120 m), ≥ 12 ha, and not located in riparian areas. Both oak/hickory and beech/maple sites contained mature trees > 50 years old and were situated in hilly terrain, thus comprised similar slopes and aspects. Boundaries of beech/maple sites were distinguished based on the transition of tree communities while boundaries of oak/hickory sites were identified artificially using SNF cover maps and ArcGIS 10.1.1 (ESRI, Redlands, CA, USA). As the SNF is primarily second growth timber, differences in tree species composition among our sites were representative of past land usage (e.g., clear cuts or selective harvest), not due to differences in location on the landscape; our sites

were large forest blocks within un-glaciated terrain. Our study sites have not experienced landscape level disturbance (i.e., logging) for > 50 years. We uploaded discrete polygon shapefiles of each study site to a handheld Global Positioning System (GPS) for ground-truthing purposes. We sampled three trees from 0600-1400 hours during 23 May to 25 June 2014. We located trees by searching outward to a 30 m radius around each vegetation point until a mature tree > 20 cm d.b.h. with branches low enough to sample was found. In general, the three mature trees closest to the center point that represented three of the five genera (*Acer*, *Carya*, *Fagus*, *Liriodendron*, and *Quercus*) of interest were sampled. “

To:

“We selected 22 study sites along an oak/hickory (xeric) to beech/maple (mesic) dominated gradient using USFS stand cover maps (allveg2008.shp) in ArcGIS 10.1.1 (ESRI, Redlands, CA, USA). We selected sites using the following criteria to prevent potential confounding effects: located within contiguous upland-deciduous forest habitat (i.e., elevation above 120 m), ≥ 12 ha, and not located in riparian areas. All sites contained mature trees > 50 years old in hilly terrain, thus comprised similar slopes and aspects. Beech/maple site boundaries were distinguished based on the transition of tree communities while oak/hickory site boundaries were identified artificially using SNF cover maps and ArcGIS 10.1.1 (ESRI, Redlands, CA, USA). Differences composition of tree species among our sites were a function of past land usage (e.g., clear cuts or selective harvest), not differences in location on the landscape; all sites were large forest blocks within un-glaciated terrain. The study sites have not experienced disturbance at a landscape level (i.e., logging) for > 50 years. We ground-truthed the maps by uploading discrete polygon shapefiles of each study site to a handheld Global Positioning System (GPS) and verifying tree species composition. We sampled three trees from 0600-1400 hours during 23 May to 25 June 2014. To locate sample trees, we searched outward to a 30 m radius from vegetation points until mature trees (> 20 cm d.b.h.) with branches low enough to sample were found. Typically, the three mature trees that represented three of the five genera (*Acer*, *Carya*, *Fagus*, *Liriodendron*, and *Quercus*) of interest and were closest to the center point were sampled.”

3. 1.1: What tool is used here?

Changed text from:

” 1.1 remove the bottom 1/3 of the 30 cm wire tomato cage so that it is approximately 55 cm in length

To:

“1.1 Using bolt cutters, large wire cutters, or an electric grinding disk, remove the bottom 1/3 of the 30 cm wire tomato cage so that it is approximately 55 cm in length”

4. Please combine some of the shorter Protocol steps so that individual steps contain 2-3 actions and maximum of 4 sentences per step.

Modified text from:

- 1.1 Using bolt cutters, large wire cutters, or an electric grinding disk, remove the bottom 1/3 of the 30 cm wire tomato cage so that it is approximately 55 cm in length
- 1.2 Cut two, 30 cm braces made from aluminum tube or similarly semi-rigid material to use as attachment rods and braces on each side of the largest end of the tomato cage
- 1.3 Attach one end of each of the two attachment rods to opposite sides of the tomato cage with duct or electrical tape ensuring the tape is wrapped around at least 6 cm of the cage and rod. Be certain to rap the tape around the cage and rod numerous times to ensure the cage is permanently attached to the rod.
- 1.4 Attach the other end of each of the two attachment rods on the opposite sides of the end of an extendable pole with duct or electrical tape. As before wrap the tape multiple times to affix it permanently ensuring the tape overlaps the pole and rods by at least 6 centimeters. Be certain the opening of the cage is in contact with the end of the telescoping pole when the cage is attached
- 1.5 Attach the cage directly onto the end of the pole using the electrical or duct tape.
- 1.6 Attach Velcro strips at 3 points to the opening of the cage. These Velcro strips will be use later to hold the opening of the bag open.

2 Enclosing the branch

- 2.1 Attach 3 pieces of Velcro to the opening of the bag so they align with the Velcro attached to the opening of the cage. These will be used to hold the opening of the bag in place while it is being placed of the branch. Be certain the Velcro is aligned so when the bag is inserted and attached, the opening to the pull strings of the bag run parallel to the telescoping pole.
- 2.2 Insert a 13-gallon kitchen garbage bag in the wire tomato cage and, using 2 gator clips, one on each side of the bottom of the bag, attach the clips to both the bag and wire cage to hold the bag against the cage.
- 2.3 Using 2 gator clips, attach the opening of the bag to the opening of the cage. Orient the gator clips so they are perpendicular to the telescoping pole and the openings to the draw strings so the run parallel to the telescoping pole. Be certain to leave adequate space near the opening of the bag to allow the bag to close when the draw string is pulled.
- 2.4 Attach para cord to each side of the bags draw string.
- 2.5 Use a random number generator to randomly select a height of the tree within the height of the extension pole when it is extended to its maximum length.
- 2.6 Use a random number generator to randomly select a distance from the tree trunk of the tree.
- 2.7 Identify a branch that will fit in the bag with minimal disturbance to the foliage and is the height and distance from the trunk based on the numbers generated from the random number generator.

2.8 Quickly slide the bag over the branch and pull the cords attached to the draw string on the bag to seal the bag. Practice this a few time prior to the first attempt to become efficient at incorporating the foliage with minima disturbance to the leaves.

2.9 Clip the branch adjacent to the bag opening with the extension pole pruner

2.10 Store the bagged branch in a freezer until you are ready to conduct the laboratory arthropod analysis.

3. Arthropod analysis

3.1 Hold frozen bag and branch upright and shake branch while in the bag to dislodge arthropods into the bag.

3.2 Carefully remove the branch and rinse in large collection pan to remove remaining arthropods.

3.3 Empty remaining material from the bag into the collection pan.

3.4 Remove any non-arthropod debris

3.5 Separate arthropods into desired taxonomic groups recording larvae and adults

3.6 Quantify arthropods as desired. If biomass is of interest either:

3.6a Measure length of arthropods and use published length mass table to estimate biomass.

3.6b Place arthropods in small drying pans, dry in drying oven for 24 hours at 45° C, and weigh on an electronic balance.

4 Estimating density

4.1 To estimate density and control for variation in leaf structure and leaf density between samples within tree species and among tree species either:

4.1a Count and measure the surface area of the leaves from each sample.

4.1b Dry the leaves in a drying oven for 48 hours at 45° C and weigh the leaves on an electronic balance.

4.1c Measure the length of all woody branch within the sample.

Diel differences occur in arthropod communities so sampling should be conducted throughout the entire period of inference.

To:

1 Building the sampling device - prior to going to the field

1.1 Using bolt cutters, large wire cutters, or an electric grinding disk, remove the bottom 1/3 of the 30 cm wire tomato cage so that it is approximately 55 cm in length

1.2 Cut two, 30 cm braces made from aluminum tube or similarly semi-rigid material to use as attachment rods and braces on each side of the largest end of the tomato cage. Attach one end of each of the two attachment rods to opposite sides of the tomato cage with duct or electrical tape ensuring the tape is wrapped around at least 6 cm of the cage and rod. Be certain to rap the tape around the cage and rod numerous times to ensure the cage is permanently attached to the rod.

1.3 Attach the other end of each of the two attachment rods on the opposite sides of the end of an extendable pole with duct or electrical tape. As before wrap the tape multiple times to affix it permanently ensuring the tape overlaps the pole and rods by at least 6 centimeters. Be certain the opening of the cage is in contact with the end of the telescoping pole when the cage is attached

1.4 Attach the cage directly onto the end of the pole using the electrical or duct tape. Attach Velcro strips at 3 points to the opening of the cage. These Velcro strips will be use later to hold the opening of the bag open.

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2.1 Attach 3 pieces of Velcro to the opening of the bag so they align with the Velcro attached to the opening of the cage. These will be used to hold the opening of the bag in place while it is being placed of the branch. Be certain the Velcro is aligned so when the bag is inserted and attached, the opening to the pull strings of the bag run parallel to the telescoping pole.

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2.3 Use a random number generator to randomly select a height of the tree within the height of the extension pole when it is extended to its maximum length. Use a random number generator to randomly select a distance from the tree trunk of the tree. Identify a branch that will fit in the bag with minimal disturbance to the foliage and is the height and distance from the trunk based on the numbers generated from the random number generator.

2.4 Quickly slide the bag over the branch and pull the cords attached to the draw string on the bag to seal the bag. Practice this a few time prior to the first attempt to become efficient at incorporating the foliage with minima disturbance to the leaves. Clip the branch adjacent to the bag opening with the extension pole pruner. Store the bagged branch in a freezer until you are ready to conduct the laboratory arthropod analysis.

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4.1c Measure the length of all woody branch within the sample.

Diel differences occur in arthropod communities so sampling should be conducted throughout the entire period of inference.

5. Please include single line spacing between each numbered step or note in the protocol.

Done

6. After you have made all the recommended changes to your protocol section (listed above), please highlight in yellow up to 2.75 pages (no less than 1 page) of protocol text (including headers and spacing) to be featured in the video. Bear in mind the goal of the protocol and highlight the critical steps to be filmed. Our scriptwriters will derive the video script directly from the highlighted text.

Done

7. Discussion: As we are a methods-based journal, please discuss critical steps in the protocol, modifications and troubleshooting of the method, and limitations of the method.

Modified the text on line 162-168 to make it more clear a limitation to this method is active or flighted-arthropods may be under-represented. We then caution researchers a critical step of this procedure is to ensure foliage remains undisturbed until it is enclosed.

Lines 177-180: We identify assessing samples in the field as a critical issue when trying to maximize detection probability and recommend a procedure we devised for doing so in the lab. We modified the text to better emphasize this.

Lines 181-199: Discuss the issue difficulty and limitation of using this procedure to estimate arthropod density and provide recommended approaches.

8. Table 1: Please upload it to your Editorial Manager account as an .xlsx file. Avoid any coloring or formatting in the tables.

Done

9. Table of Materials: Please ensure that it has information on all relevant supplies, reagents, equipment and software used, especially those mentioned in the Protocol. Please sort the materials alphabetically by material name.

Done

Reviewers' comments:

Reviewer #2:

Manuscript Summary:

The revision of the manuscript entitled "A method for quantifying foliage-dwelling arthropods" has been improved greatly. Thus, I suggest that it can be accepted with the present version.

Major Concerns:

No.

Minor Concerns:

No.

Reviewer #5:

The ms describe the benefits of the branch bagging methodology to quantify density, richness and diversity of foliage dwelling arthropods in 5 groups of tree species in Shawnee National Forest, Illinois. The freezing of the sample branch followed by it rinse with water can reduce the time spent looking for individuals, as well decrease the chances of lost some of them.

The estimating means and standard errors were used to know the level of precision from the samplings. The results showed that this method can be precise to test the influence of environmental variables on richness, diversity and density of species, since the values of percentage of mean of the standard errors for community metrics were low.

The method procedure was clear and described in detail. The method is sound, I believe that the ms will benefit field researchers providing a substantive way to estimate differences in arthropods abundance and biomass, diversity and richness of foliage-dwelling arthropods communities.