

Article

Evaluation of Cowpea (*Vigna unguiculata*) in an Intercropping System as Pollinator Enhancer for Increased Crop Yield

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Abstract: Production of pollinator-dependent crops (PDCs) is increasing. However, pollinators are declining partly due to loss of floral resources. There is urgency to mitigate this decline and the potential risks to the production of PDCs and food security. One way is by promoting farming systems that enhance flower-rich habitats. In a two-year study, Pinkeye Purple Hull and Whippoorwill cowpea varieties attractive to pollinators were intercropped with three PDCs (squash, okra, and watermelon). We evaluated whether cowpea intercrop increases the abundance and diversity of pollinators, other beneficial insects, and crop yield, and decreases the abundance of the brown marmorated stink bug (BMSB). Diverse pollinators were recorded in both years for a total of 80,379, representing seven pollinator families (Andrenidae, Apidae, Crabronidae, Formicidae, Halictidae, Tachinidae, and Vespidae) captured in pan traps, six families (Apidae, Crabronidae, Halictidae, Pyralidae, Tachinidae, and Vespidae) from sticky traps, and five pollinator types (bumble bees, carpenter bees, honeybees, butterflies and moths, and wasps) through direct visual count. Pollinator abundance and diversity was highest on the cowpea-intercropped treatments than controls. PDCs intercropped with cowpea recorded more beneficial insects than BMSB. Okra, squash, and watermelon intercrops produced 7%, 27%, and 54% more fruits than the control, respectively. Our findings indicate that intercropping cowpeas with PDCs attracted more and diverse pollinators and resulted in increased crop yield. However, to optimize pollination, factors such as planting dates to synchronize the flowering of both cowpeas and PDCs should be taken into consideration.

Keywords: cowpea; pollinator; *H. halys*; brown marmorated stink bug; BMSB; intercropping; floral resources; pollinator-dependent crops

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1. Introduction

Insect pollinators are an important component of biodiversity that provides ecosystem services, including the pollination of many crops. Pollination mediated by insects provides essential support for food security and ecosystem sustainability. Globally, about 90% of flowering plants that produce seeds require insect pollination [1], and about 35% of the human diet is dependent on insect pollination [2]. However, the abundance and diversity of bees and other pollinating insects have been on a decline [3]. The loss of floral resources and other factors including agricultural intensification, spread of pathogens, climate change, landscape modification, non-native species invasion, and the widespread use of pesticides [3] have contributed to this decline to some extent. This has led to an increased reliance on managed species, most often honeybees, to mediate pollination services despite recorded low yields in some cases [4].

Specific plant species provide unique seasonal pattern in the amount, diversity, and composition of floral resources, which together create phenological complementary forage resources for pollinators [5–7]. A review of previous studies indicated that plant diversity in and around crop fields improved habitat for bees as well as other beneficial

insects [8]. In this mutualistic relationship, pollinators receive nectar and pollen, which provides them with carbohydrates, proteins, fats, vitamins, and minerals [9], and in return plants receive the service of pollination. In recent years, there has been a rise in the production of pollinator-dependent crops (PDCs) which include many vegetables such as squash, pumpkin, cucumber, watermelon, and okra [1,10]. Unfortunately, honeybees, which are the most important insect pollinators worldwide, as well as some native bee species, are on the decline. This raises concerns about the production of pollinator-dependent crops. Research has shown that the diverse assemblage of insect pollinators as a result of the provision of floral resources promotes pollination and crop yield [7,9,11]. For example, flower strips [12–14], intercropping with flowering plants [15], and hedgerows [16] increased pollinator abundance in crop fields.

Cowpea (*Vigna unguiculata* L. Walp) has been recognized for its nectariferous nature and its role in attracting pollinators [17–20]. The plant has extrafloral nectaries on its petioles and leaflets that are nectar-releasing sites, which attract beneficial insects, including pollinators. Cowpea is widely grown in the tropics and subtropical regions as an intercrop [21] to enhance soil fertility through its ability to fix atmospheric nitrogen. It is consumed as a major source of protein; the leaves contain about 34.9% protein and the seeds about 25% protein [22,23]. In the United States, cowpea is grown widely in California, Texas, and in the southern states, mainly for frozen, canned, and fresh peas [24,25]. The brown marmorated stink bug (BMSB), *Halyomorpha halys* (Heteroptera: Pentatomidae), has emerged as a serious pest of many crops [26], including cowpea [27,28]. According to [26], *H. halys* causes millions of dollars' worth of economic loss to fruits and vegetables annually. Currently, pesticides are the main strategy for the management of *H. halys*. Considering the dangers of frequent pesticide use to humans and the environment, there is a need to investigate other environmentally friendly strategies.

Intercropping is an important cultural practice based on the principle of reducing insect pests, increasing diversity in an ecosystem, and increasing crop yield. However, there are inconsistent reports on crop yield associated with intercropping. For example, cowpea intercropped with corn [29], maize-cassava [30], and okra [31,32] resulted in increased crop yield compared to sole cropping. On the other hand, intercropping cowpea with tomato and okra resulted in a 45–55% loss in marketable fruit yields in both vegetables [33,34]. Most often, the ability of cowpea to fix atmospheric nitrogen and its contribution to the replenishment of soil fertility has been associated with the increase in crop yield [35]. Several factors, including pollinators in the cropping system, may also play a crucial role in the yield outcome. It is therefore important to assess the contribution of pollinators in an intercropping system on crop yield. In this study, two cowpea varieties (Pinkeye Purple Hull and Whippoorwill) reported to be highly attractive to diverse pollinators [19] were intercropped with three pollinator-dependent crops (squash, okra, and watermelon), including controls (PDCs with no cowpeas and sole cowpea). We determined the effect of cowpea intercrop on the abundance and diversity of pollinators and other beneficial insects, *H. halys*, and crop yield.

2. Materials and Methods

2.1. Study Site and Experimental Plants

The study was conducted at the North Carolina Agricultural and Technical State University Research Farm in Greensboro, North Carolina, USA (longitude 36.0586243° N and latitude 79.78932° W) during the summers of 2018 and 2019. Three pollinator-dependent crops (PDCs)—squash (*Cucurbita pepo* L.) [S] (Tempest Yellow Summer variety), okra (*Abelmoschus esculentus*) [O] (Clemson Spineless variety), and watermelon (*Citrullus lanatus*) [W] (Little Baby variety in 2018 and Mini Love variety in 2019)—were obtained from Johnny's Seeds (Winslow, ME, USA). Together, these PDCs are referred to as SOW in this paper. They were selected because they are commonly grown by limited-resource

farmers in North Carolina. The PDCs were intercropped with two cowpea varieties: Pink-eye Purple Hull [PPH] procured from Hummert Seed Company (Earth City, MO, USA) and Whippoorwill [WHIP] from Southern Exposure Seed Exchange (Mineral, VA, USA).

2.2. Experimental Design and Setup

In both years, there were three cowpea intercrop treatments (PPH + SOW + PPH, WHIP + SOW + WHIP, and PPHWHIP + SOW + PPHWHIP), two sole cowpea treatments (PPH-only and WHIP-only), and one sole PDC treatment (SOW), for a total of six treatments. The cowpea intercrop treatments comprised six rows of PDCs (two rows of each PDC) bordered on each side by either two rows of PPH or WHIP, or one row each of PPH and WHIP. The sole cowpea treatments comprised five rows each of PPH and WHIP. The sole PDC treatment comprised three rows each of squash, okra, and watermelon without cowpea (control) and was set up 50 m away from the cowpea and cowpea intercrop treatments. The experiments were set up as a randomized complete block design (RCBD) and replicated four times. For each treatment, each row measured 5 m long with 1 m inter-row spacing.

In 2018, all seeds were planted from 1 June to 12 June and according to the 2018 *Southeastern Vegetable Handbook* [36]. During the season, we observed that cowpea flowered before okra but after squash and watermelon. In 2019, the planting date was staggered for the different species based on the date of flowering recorded in 2018 to synchronize the flowering of PDCs with that of cowpeas. Therefore, okra was planted on 4 June, WHIP on 18 June, PPH on 25 June, watermelon on 12 July, and squash on 19 July. In both years, the seeds were planted by the block manually and the entire experimental setup was mulched with wheat straw (Rankin Farms Inc. Ellerbe, NC, USA). Conventional recommendations of growing vegetables in the Southeast were followed [36]. No insecticides were applied, and plots were irrigated (drip irrigation system) as needed.

2.3. Insect Sampling

Pollinators, including other beneficial insects, and *H. halys* were sampled on both the PDCs and the cowpeas. Sampling was conducted using direct visual count, sticky traps, and pan traps (blue, white, and yellow).

2.3.1. Insect Sampling Using Direct Visual Counts

Direct visual counts of pollinators and *H. halys* were carried out by counting the number of pollinator types (bumble bees, carpenter bees, honeybees, butterflies and moths, and wasps) and *H. halys* was observed on each 5 m row for two minutes. For each treatment, a visual count of pollinators began 37 days after planting (DAP), corresponding to the first incidence of pollinators on the experimental plots weekly between 08:00 to 13:00 for five weeks from 11 July to 8 August 2018, and 17 August to 11 September 2019. Visual counts of *H. halys* began 46 DAP, the time when *H. halys* was first observed on the experimental plots, and conducted weekly from 10:00 to 12:00 for five weeks from 20 July to 13 August 2018 and 16 August to 9 September 2019.

2.3.2. Insect Sampling Using Sticky Traps

Double-sided yellow sticky traps (8 × 13 cm²) (Sticky Strips, Olson Products, Medina, OH, USA) were used to monitor smaller pollinators and other beneficial insects that were likely to be missed with visual counts and pan trap sampling. They also provided an additional alternative sampling method. Beginning 34 DAP, weekly sticky trap sampling was conducted from 9 July to 30 July, 2018, and from 12 August to 2 September, 2019. One sticky trap affixed to a metal stake was placed at the center of each row of the cowpea intercrop treatments, the sole PDC treatment, and the inner rows of the sole cowpea treatment for a total of two sticky cards on two rows. Traps were placed just below the canopy height. The traps were removed after 24 h and placed in Ziploc® bags and taken to the

Integrated Pest Management (IPM) Laboratory, where the insects were identified using a microscope (AmScope Stereozoom trinocular microscope, SZMT2 Series, WF10X/20; United Scope LLC, Irvine, CA, USA) to the family level.

2.3.3. Insect Sampling Using Pan Traps

Pollinators were sampled within each treatment using three colored (blue, white, and yellow) pan traps. Sampling began 45 DAP with the presence of flower visitors. As described by [33], the traps consisted of 16 oz squat polypropylene deli bowls (BioServ, Frenchtown, NJ, USA) painted with UV-bright fluorescent blue paint (blue trap) or yellow paint (yellow trap) and unpainted 12 oz white Styrofoam bowls (white trap) (Uline, Pleasant Prairie, WI, USA). The trap setup was carried out by gluing individual unpainted 16 oz polypropylene deli bowls onto a 36" plant prop and three of these were placed 0.2 m apart between the two rows of the PDC and cowpeas for each treatment for the entire sampling period. One of each of the colored traps was then placed inside each of the unpainted bowls on the prop and filled with approximately 250 mL of soapy water solution [33]. The traps were placed so that they were at the same level as the crop canopy. Traps were set out weekly from 19 July to 16 August 2018, and from 22 August to 19 September 2019, between 08:00 to 10:00 and collected after 24 h. During collection, each pan trap was drained, and the contents transferred to vials containing 70% ethanol and taken to the IPM laboratory, where they were stored in a refrigerator and later identified using a microscope (AmScope Stereozoom trinocular microscope, SZMT2 Series, WF10X/20; United Scope LLC, Irvine, CA, USA) to the family level. The traps were collected in the order they were placed to ensure that all traps were available to insects for approximately equal time.

2.4. Crop Yield Data Collection

At maturity, all crops in each treatment including the control were harvested with the aid of a handheld pruner of 10 cm × 2.5 cm × 27 cm dimensions (Fiskars Bypass Pruner, The Home Depot®, NC, USA). Squash was harvested weekly from 9 July (39 DAP) to 6 August in 2018 and from 26 August (39 DAP) to 23 September in 2019. Okra was harvested weekly from 6 August (67 DAP) to 7 September in 2018 and from 23 August (67 DAP) to 26 September in 2019. Watermelon was harvested once on 30 July (60 DAP) in 2018 due to excessive rainfall that led to the bursting of the fruits and subsequent wilting of the plants, and every two weeks for a total of three times from 11 September (59 DAP) to 9 October in 2019. Twenty-five fresh pods from each of the five-cowpea treatments were randomly harvested from each 5 m row within each treatment on 10 August in 2018 and 5 September in 2019. Harvested fruits were weighed using an Ohaus T51P weighing scale (Pine Brook, NJ, USA) and data were recorded.

2.5. Data Analysis

The diversity of pollinators reported from direct visual counts, sticky traps, and pan traps on the various treatments was assessed using the Shannon–Weaver Diversity Index (H') [37,38]. For each sampling method, the total number of pollinators in each family or pollinator type captured on each treatment for the entire sampling period was used to calculate the Diversity and Evenness indices.

The index is expressed as:

$$H' = - \sum pi \ln(pi)$$

where H' = Shannon–Weaver diversity index

pi = the proportion of individuals found in species i

pi is estimated as $pi = ni/N$

where ni = number of individuals in species i

N = total number of individuals in the community

The Shannon Equitability (Evenness) Index (E) normalizes the Shannon Diversity Index (H') to a value between 0 and 1. It provides an idea about the evenness of the distribution of groups of organisms in a community. An index value of 1 means that all groups have the same frequency.

The Evenness Index (E) is expressed as:

$$E = H'/\log(k)$$

where E = Evenness,

H' = Shannon–Weaver Diversity Index

k = number of species/groups in the community

For each sampling method, the total number of insects belonging to each insect family or pollinator type that was captured throughout the sampling period was obtained.

The mean number of insect pollinators and other beneficial insects, *H. halys*, and the yield of fresh cowpeas and PDCs within the 5 m row sampling area were analyzed using generalized linear mixed models using PROC GLIMMIX in SAS 9.4 (SAS Institute Inc, Cary, NC) with block as a random effect. Crops, years, transformation, distribution, and use of the repeated measures for each data type are presented in Table S1. The data of the PDCs were sorted by each crop (i.e., squash, okra, and watermelon) using the PROC SORT function. Some of the data were analyzed as repeated measures with an autoregressive order 1 (AR(1)) repeated measures correlation structure. Polled data for all sampling methods and data from both years combined were not analyzed using repeated measures because of the unequal number of sampling weeks (each year sampling using direct visual count and pan trap was done weekly for five weeks compared to four weeks for sampling using sticky traps). For the weekly distribution of pollinators on the cowpeas and PDCs, pollinator counts obtained during the fifth sampling week were eliminated because data were collected only by direct counts and pan traps and not with sticky traps, unlike in the first four weeks where counts were done using all three sampling methods. Pollinator counts obtained during sampling were transformed either by square root or log-transformation or without any transformation to meet the assumptions of normality (Table S1). Some of the data fit well with negative binomial distribution with logit link function (Table S1). The degrees of freedom (df) were assessed using the Kenward–Rogers degrees of freedom approximation. Multiple comparisons of treatments were made comparing the least-squares means (LSM) with a Tukey–Kramer test at a 5% level of significance.

The relationship between the number of pollinators and the yield of each PDC for each treatment was determined using Pearson's correlation and regression analysis. Data were fitted to the equation:

$$y = a + bx$$

where y is the crop yield and x is the number of pollinators.

All the above analyses except the diversity and evenness assessments were performed using SAS 9.4 (SAS Institute Inc., Cary, NC, USA).

3. Results

3.1. Abundance and Diversity of Pollinators Associated with Cowpea Treatments and Controls

Five pollinator types (bumble bees, carpenter bees, honeybees, butterflies and moths, and wasps) were recorded through direct visual counts in both years on all treatments (Table 1). A total of 4476 and 7417 pollinators were recorded on all treatments in 2018 and 2019, respectively. In 2018, the highest number of pollinators (1159) was recorded on PPH + SOW + PPH and the lowest number (364) on SOW (control) (Table 1). The number captured ranged from 364 on the control to 1159 on PPH + SOW + PPH with numbers on PPH + SOW + PPH > PPHWHIP + SOW + PPHWHIP > PPH-only > WHIP + SOW + WHIP > WHIP-only > SOW. Similarly, in 2019, the highest number of pollinators (2242) was recorded on the PPH + SOW + PPH, and the lowest (104) on the control with numbers on

PPH + SOW + PPH > PPHWHIP + SOW + PPHWHIP > WHIP + SOW + WHIP > PPH-only > WHIP-only > SOW (Table 1).

From sticky cards, a total of 30,114 and 25,163 pollinators representing six families (Apidae, Crabronidae, Halictidae, Pyralidae, Tachinidae, and Vespidae) were captured on all treatments in 2018 and 2019, respectively (Table 2). In 2018, the number captured ranged from 2980 on the control to 6650 on PPH + SOW + PPH with numbers on PPH + SOW + PPH > PPHWHIP + SOW + PPHWHIP > WHIP + SOW + WHIP > PPH-only > WHIP-only > SOW. Similarly, in 2019, the number of pollinators captured ranged from 1581 on the control to 5706 on PPHWHIP + SOW + PPHWHIP with numbers on PPHWHIP + SOW + PPHWHIP > PPH + SOW + PPH > WHIP + SOW + WHIP > PPH-only > WHIP-only > SOW (Table 2).

A total of 7884 and 5295 insects representing seven pollinator families (Andrenidae, Apidae, Crabronidae, Formicidae, Halictidae, Tachinidae, and Vespidae) were captured in pan traps in all treatments in 2018 and in 2019, respectively (Table 3). In 2018, the number of pollinators captured ranged from 802 on PPH-only to 1789 on WHIP + SOW + WHIP with numbers on WHIP + SOW + WHIP > PPHWHIP + SOW + PPHWHIP > PPH + SOW + PPH > SOW > WHIP-only > PPH-only. In 2019, the number of pollinators ranged from 391 on the control to 1251 on WHIP + SOW + WHIP with numbers on WHIP + SOW + WHIP > PPH + SOW + PPH > PPHWHIP + SOW + PPHWHIP > WHIP-only > PPH-only > SOW.

Diversity and evenness varied among the sampling methods and year. In 2018, insects recorded from visual counts were more diverse and evenly distributed on the five-cowpea treatments than control with the highest diversity ($H' = 1.40$) and evenness ($E = 0.87$) recorded on cowpea PPH-only. However, in 2019, insects were more diverse and evenly distributed in the control (Table 1). Insects from sticky traps were diverse and evenly distributed on the control treatment in 2018 and 2019 (Table 2). Species diversity and evenness from pan trap capture were similar among all treatments in 2018. However, in 2019, all five-cowpea treatments were more diverse and evenly distributed than the control (Table 3).

Table 1. The total number of insects recorded through direct visual counts on cowpea treatments and controls in 2018 and 2019.

Order	Pollinator Type (Common Name)	Treatments and Number of Pollinators (2018)						Treatments and Number of Pollinators (2019)					
		PPH + SOW + PPH	WHIP + SOW + WHIP	PPHWHI P + SOW + PPHWHI P	PPH-Only	WHI P-Only	SOW	PPH + SOW + PPH	WHIP + SOW + WHIP	PPHWHIP + SOW + PPHWHIP	PPH-Only	WHIP-Only	SOW
	Bumble bee	526	322	405	298	40	227	145	108	199	62	63	10
Apidae	Carpenter bee	36	24	16	8	11	8	4	1	4	0	1	0
	Honeybee	203	182	220	197	103	100	589	342	495	259	232	42
Hymenoptera	Wasps	260	74	135	233	80	9	1289	814	957	538	417	18
Lepidoptera	Butterfly and moth	134	143	121	159	182	20	215	175	209	94	101	34
	Total	1159	745	897	895	416	364	2242	1440	1864	953	814	104
	(H)	1.36	1.36	1.33	1.40	1.35	0.98	1.08	1.12	1.19	1.08	1.17	1.26
	(E)	0.84	0.85	0.83	0.87	0.84	0.61	0.67	0.70	0.74	0.78	0.72	0.91

Table 2. The total number of insects captured on sticky traps on cowpea treatments and controls in 2018 and 2019.

Pollinator Family	Treatments and Number of Pollinators (2018)						Treatments and Number of Pollinators (2019)					
	PPH + SOW + PPH	WHIP + SOW + WHIP	PPHWHIP + SOW + PPHWHIP	PPH-Only	WHIP-Only	SOW	PPH + SOW + PPH	WHIP + SOW + WHIP	PPHWHIP + SOW + PPHWHIP	PPH-Only	WHIP-Only	SOW
Apidae	14	9	23	7	2	21	12	15	9	8	7	2
Crabronidae	550	673	579	476	383	392	134	169	127	72	68	34
Halictidae	1019	1018	1114	642	654	679	606	644	662	369	377	190
Pyralidae	25	22	34	15	16	45	66	77	77	29	34	153
Tachinidae	4995	4444	4636	3234	2276	1757	4577	3951	4595	3397	2566	1182
Vespidae	47	74	78	41	34	86	167	160	236	192	179	20
Total	6650	6240	6464	4415	3365	2980	5562	5016	5706	4067	3231	1581
(H')	0.78	0.86	0.86	0.82	0.91	1.12	0.66	0.76	0.71	0.63	0.74	0.84
(E)	0.43	0.48	0.48	0.46	0.51	0.62	0.37	0.42	0.40	0.35	0.41	0.47

Table 3. The total number of insects captured in pan traps on cowpea treatments and controls in 2018 and 2019.

Pollinator Family	Treatments and Number of Pollinators (2018)						Treatments and Number of Pollinators (2019)					
	PPH + SOW + PPH	WHIP + SOW + WHIP	PPHWHIP + SOW + PPHWHIP	PPH-Only	WHIP-Only	SOW	PPH + SOW + PPH	WHIP + SOW + WHIP	PPHWHIP + SOW + PPHWHIP	PPH-Only	WHIP-Only	SOW
Andrenidae	16	20	19	8	15	23	1	6	1	1	0	3
Apidae	77	72	76	25	30	82	22	34	24	14	15	7
Crabronidae	138	165	112	79	73	20	68	93	73	68	68	19
Formicidae	37	33	38	21	20	119	1	5	3	3	2	4
Halictidae	669	787	740	407	465	556	410	489	420	228	254	118
Tachinidae	400	649	535	210	328	501	377	397	378	216	215	197
Vespidae	79	63	44	52	50	31	249	227	178	190	174	43
Total	1416	1789	1564	802	981	1332	1128	1251	1077	720	728	391
(H')	1.40	1.32	1.30	1.35	1.32	1.34	1.33	1.38	1.32	1.41	1.39	1.25
(E)	0.72	0.68	0.67	0.69	0.68	0.69	0.68	0.71	0.70	0.72	0.77	0.64

3.2. Pollinators on Cowpeas and Pollinator-Dependent Crops

3.2.1. Pollinators on Cowpeas

Among the five-cowpea treatments, the mean number of pollinators recorded on the cowpeas through direct visual counts over the entire sampling period was significantly different ($F_{4,18.7} = 5.94, p = 0.002$) in 2018 and ($F_{4,16.8} = 4.92, p = 0.008$) in 2019. In 2018, PPH + PPH and PPH-only recorded the highest number of pollinators and the number of pollinators ranged from 17.5 ± 1.4 on WHIP + WHIP to 44.8 ± 10.8 in PPH-only treatment (Figure 1a). In 2019, PPH + PPH and PPHWHIP + PPHWHIP cowpeas recorded the highest number of pollinators of 87.7 ± 7.6 and 77.7 ± 5.8 , respectively (Figure 1a). Sticky trap catches indicated there were significant differences in the numbers of pollinators captured among the cowpeas in 2018 ($F_{4,24.42} = 4.59, p = 0.007$) and in 2019 ($F_{4,23.13} = 6.94, p = 0.0008$). In 2018, the highest number of pollinators were captured on PPH-only cowpea (220.8 ± 10.2) and the lowest on WHIP + WHIP cowpea (167.1 ± 8.4) (Figure 1b). Similarly, in 2019 the highest number of pollinators were captured on PPH-only (203.4 ± 16.0) and the lowest on WHIP + WHIP (156.5 ± 6.3) (Figure 1b). Pollinators recorded from pan traps were not significantly different among the cowpeas in 2018 ($F_{4,21.01} = 2.2, p = 0.1$) or in 2019 ($F_{4,21.14} = 0.23, p = 0.92$) (Figure 1c). However, in 2018, pollinator counts were highest (71.2 ± 13.6) on WHIP + WHIP > WHIP-only (61.4 ± 10.9) > PPHWHIP + PPHWHIP (60.9 ± 10.8) > PPH + PPH (50.4 ± 8.1) cowpeas and lowest (50.2 ± 6.8) in PPH-only cowpea. Pollinator counts in 2019 ranged from 44.2 ± 4.1 on PPHWHIP + PPHWHIP to 47.5 ± 4.2 in WHIP + WHIP cowpeas (Figure 1c).

Combining pollinators from all sampling methods for each year, our results show that the total number of pollinators captured in 2018 was not significantly different ($F_{4,12} = 2.46, p = 0.10$) among the cowpeas. However, PPH-only (315.7 ± 25.9) and PPH + PPH (304.3 ± 39.3) recorded the highest numbers of pollinators (Figure 1d). In 2019, significant differences ($F_{4,15} = 10.4, p = 0.0003$) were recorded among the cowpeas, with counts ranging from 247.8 ± 7.7 on WHIP-only to 323.1 ± 15.8 on PPH + PPH (Figure 1d).

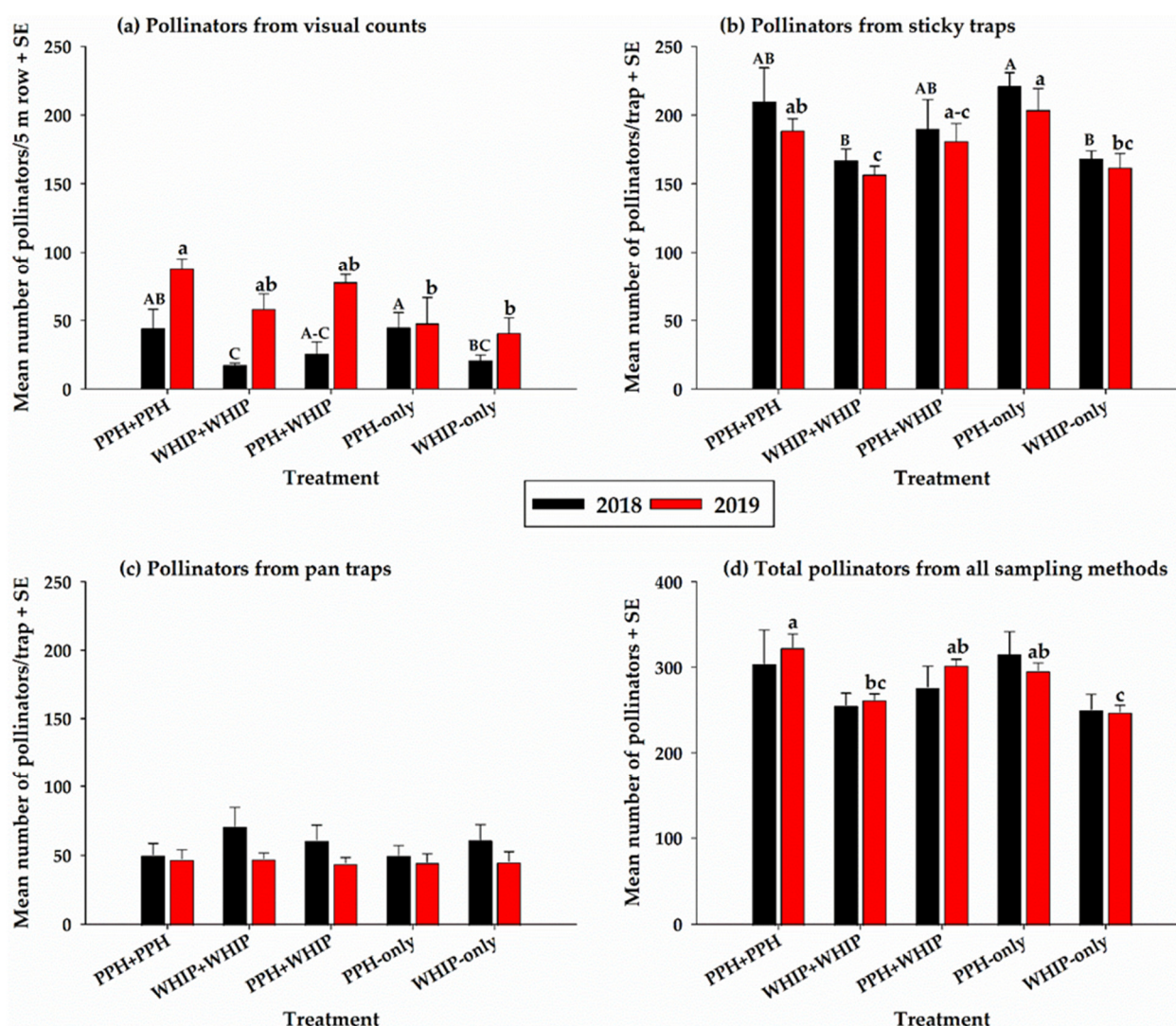


Figure 1. Pollinator abundance on cowpeas from direct visual counts (a), sticky traps (b), pan traps (c), and mean total from all sampling methods combined (d) in 2018 and 2019. For each year and within each graph, treatment means with the same letters are not significantly different (Tukey–Kramer Test, $p > 0.05$).

3.2.2. Pollinators on Pollinator-Dependent Crops (PDCs)

Pollinators recorded from the three sampling methods on each pollinator-dependent crop bordered with cowpea and without cowpea in 2018 and 2019 are presented in Figures 2 and 3, respectively. In 2018, the number of pollinators recorded through direct visual counts on squash intercropped with cowpea was significantly different ($F_{3,23.9} = 8.91$, $p = 0.0004$) than the control (squash without cowpea). Figure 2a shows that more pollinators were recorded on squash that was intercropped with cowpea compared to the control without cowpea. Similarly, in 2019, there was significant difference ($F_{3,15.8} = 4.08$, $p = 0.025$) in the number of pollinators recorded on squash intercropped with cowpea and the control. Figure 3a shows that more pollinators were recorded on squash intercropped with PPH + PPH and PPHWHIP + PPHWHIP cowpeas than on the control. For okra, there was no significant difference ($F_{3,16.5} = 1.78$, $p = 0.19$) in the number of pollinators recorded among all treatments in 2018 (Figure 2a). However, in 2019, there was significant difference among the treatments ($F_{3,12.1} = 4.15$, $p = 0.031$), with okra intercropped with PPH + PPH cowpea recording the highest number of pollinators (Figure 3a). For watermelon, there was no significant difference in the number of pollinators recorded on all treatments in both 2018 ($F_{3,15.14} = 3.08$, $p = 0.0059$) and 2019 ($F_{3,11.77} = 1.72$, $p = 0.22$). However, in both years the count was higher on watermelon intercropped with cowpea (Figures 2a and 3a).

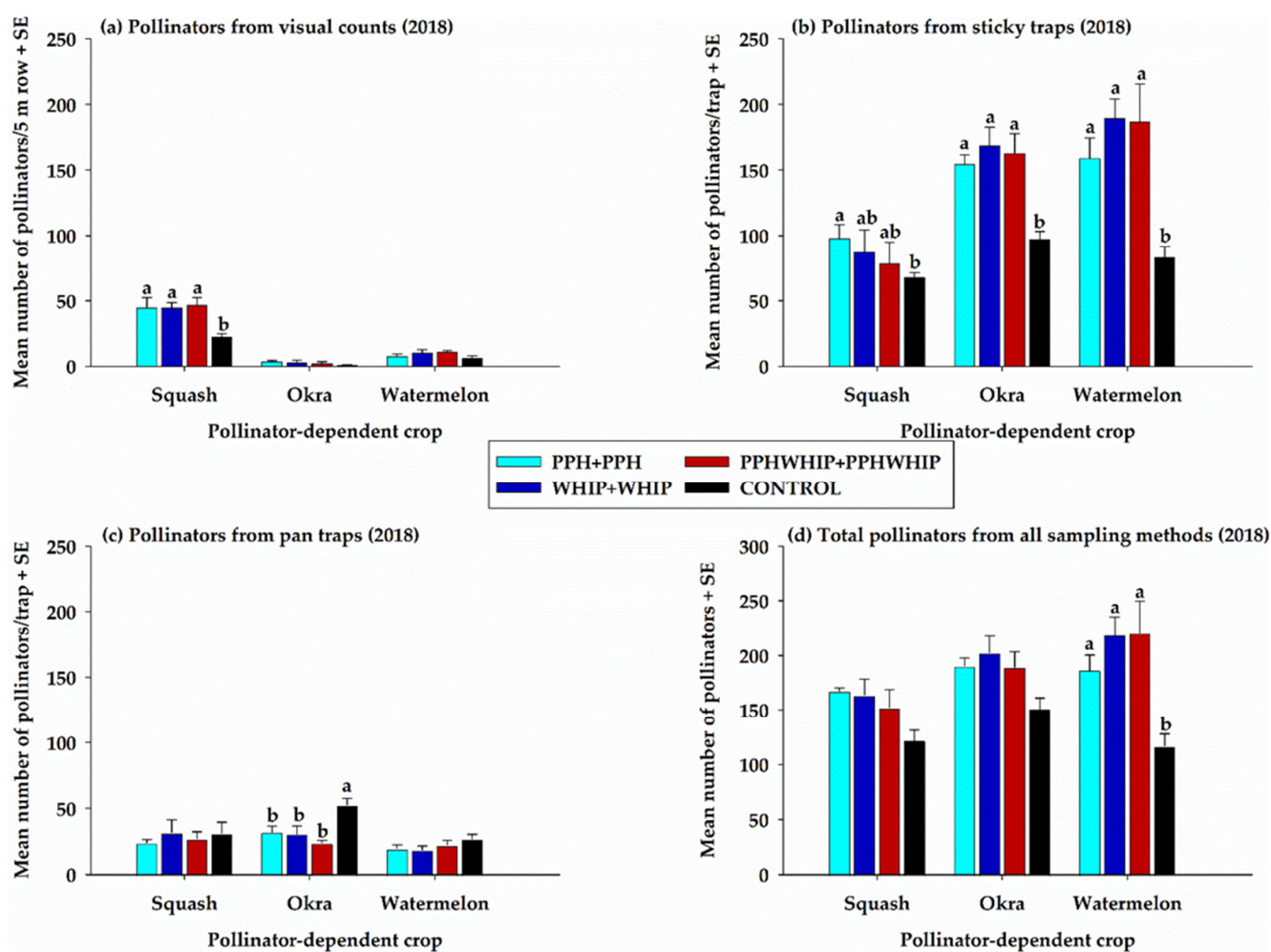


Figure 2. Pollinator abundance on pollinator-dependent crops (PDCs) from direct visual counts (a), sticky traps (b), pan traps (c), and all sampling methods combined (d) in 2018. For each crop and within each graph, treatment means with the same letters are not significantly different (Tukey–Kramer test, $p < 0.05$).

Results from sticky traps show that in 2018 there was a significant difference ($(F_{3,16.6} = 3.45, p = 0.041)$, $(F_{3,20.4} = 8.60, p = 0.0007)$, and $(F_{3,14.3} = 11.90, p < 0.0004)$) among intercropped squash, okra, and watermelon, respectively, and the control. Similarly, in 2019, there was a significant difference ($(F_{3,11.7} = 9.48, p = 0.0002)$, $(F_{3,13.3} = 6.33, p = 0.0007)$, and $(F_{3,10.9} = 9.77, p = 0.002)$) among intercropped squash, okra, and watermelon, respectively, and the control. In both years, more pollinators were recorded on the pollinator-dependent crops (squash, okra, and watermelon) that were intercropped with cowpea than on PDCs without cowpea (Figures 2b and 3b).

From pan traps, there was no significant difference ($(F_{3,15.0} = 0.64, p = 0.60)$ and $(F_{3, 16.7} = 2.60, p = 0.09)$) in the number of pollinators recorded on squash intercropped with cowpea and the control in 2018 and 2019, respectively. Our results show that in 2018, the number of pollinators on the okra control was significantly higher ($F_{3,16.3} = 9.65, p = 0.0007$) than on intercropped okra. However, in 2019, there was no significant difference ($F_{3, 14.6} = 1.09, p = 0.38$) among the treatments, and pollinator counts were higher on the okra intercropped with WHIP + WHIP than on the control (Figure 3c). In 2018 and 2019, pollinator counts on watermelon were not significantly different ($(F_{3,12.9} = 1.02, p = 0.41)$ and $(F_{3,12.9} = 2.92, p = 0.07)$ respectively). However, in 2019, counts were higher on intercropped watermelon than on the control (Figure 3c).

Combining pollinators from all sampling methods on each PDC intercropped with cowpea and controls without cowpea intercropped shows that in 2018, there was no significant difference in squash ($F_{3,9} = 3.28, p = 0.07$) or okra ($F_{3,9} = 3.54, p = 0.06$) in the number

of pollinators recorded on the PDC intercropped with cowpea and the control. However, more pollinators were recorded on squash and okra intercropped with cowpea than the control (Figure 2d). In 2019, there was a significant difference in squash ($F_{3,9} = 8.15$, $p = 0.006$), okra ($F_{3,13} = 12.52$, $p = 0.002$), and watermelon ($F_{3,9} = 8.47$, $p = 0.006$) in the number of pollinators recorded on these PDCs intercropped with cowpea than their controls (Figure 3d).

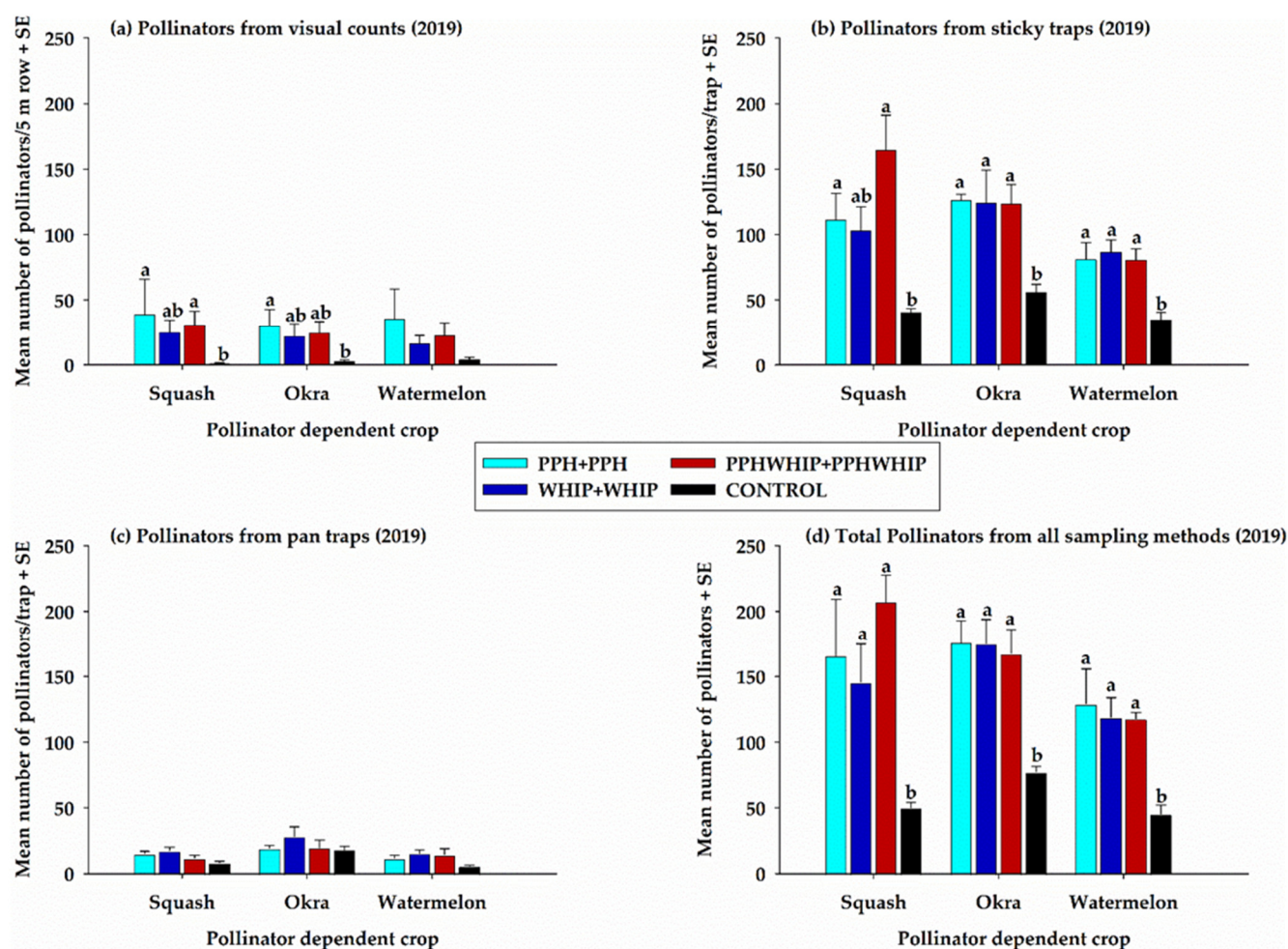


Figure 3. Pollinator abundance on pollinator-dependent crops (PDCs) from direct visual counts (a), sticky traps (b), pan traps (c), and all sampling methods combined (d) in 2019. For each crop and within each graph, treatment means with the same letters are not significantly different (Tukey–Kramer test, $p < 0.05$).

3.2.3. Total Pollinators on Cowpeas and on Pollinator-Dependent Crops and Controls

Total pollinators recorded from all sampling methods in both years on cowpeas from each of the five-cowpea treatments and on the pollinator-dependent crops are presented in Figure 4a,b, respectively. Among the five-cowpea treatments, the number of pollinators recorded on the cowpeas was significantly different ($F_{4,12.3} = 9.6$, $p = 0.001$), with pollinator counts higher on PPH + PPH = PPH-only \geq PPH + WHIP \geq WHIP + WHIP $>$ WHIP-only and ranging from 498.1 ± 18.2 on WHIP-only to 627.4 ± 48.9 on PPH + PPH (Figure 4a). Pollinator counts were not significantly different among the three intercropped pollinator-dependent crops but were significantly different than their controls for squash ($F_{3,9} = 6.84$, $p = 0.01$), okra ($F_{3,12} = 11.8$, $p = 0.0007$), and watermelon ($F_{3,9} = 13.0$, $p = 0.001$). The controls recorded the lowest counts, with 162.3 ± 9.3 in watermelon, 172.3 ± 11.0 in squash, and 228.2 ± 8.4 in okra (Figure 4b).

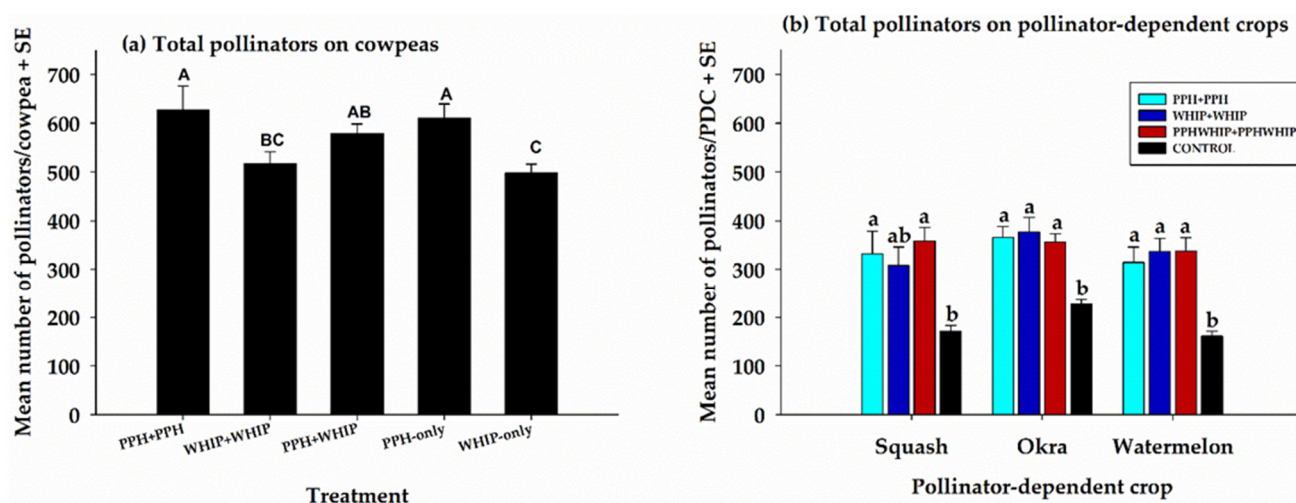


Figure 4. Total pollinators captured on cowpeas (a) and pollinator-dependent crops and control (b) in both years (2018 and 2019) during the entire study period. In subfigure (a), treatment means with the same letters are not significantly different ($p > 0.05$). In subfigure (b), within each crop, treatment means with the same letters are not significantly different ($p > 0.05$).

Comparing the number of pollinators on cowpeas from each of the five-cowpea treatments with the number on squash intercropped with cowpea and on the squash control, our results show there was a significant difference ($F_{8,24} = 41.0, p < 0.001$) among treatments. The highest pollinator counts were recorded on the cowpeas of the five-cowpea treatments, and counts on intercropped squash were higher than counts on the squash control (Figure 5a). There was a significant difference ($F_{8,24} = 38.3, p < 0.001$) between the number of pollinators on cowpeas from each of the five-cowpea treatments and the number on okra intercropped with cowpea and on the okra control. Similar to squash, the highest pollinator counts were recorded on the cowpeas of the five-cowpea treatments, also counts on okra intercropped were higher than counts on the okra control (Figure 5b). On watermelon, there was a significant difference ($F_{8,24} = 62.9, p < 0.001$) among treatments. Similar to squash and okra, pollinator counts on intercropped watermelon were similar and higher than counts on the control but lower than the counts on cowpeas (Figure 5c).

Similarly, during each sampling week, more pollinators were recorded on cowpeas compared to the PDCs intercropped with cowpea (Figure 6a–c). Among the four sampling weeks for each treatment, pollinator counts dropped on week three on both the cowpeas and PDCs and peaked at week four for the PPH + SOW + PPH and PPHWHIP + SOW + PPHWHIP treatments (Figure 6a,b).

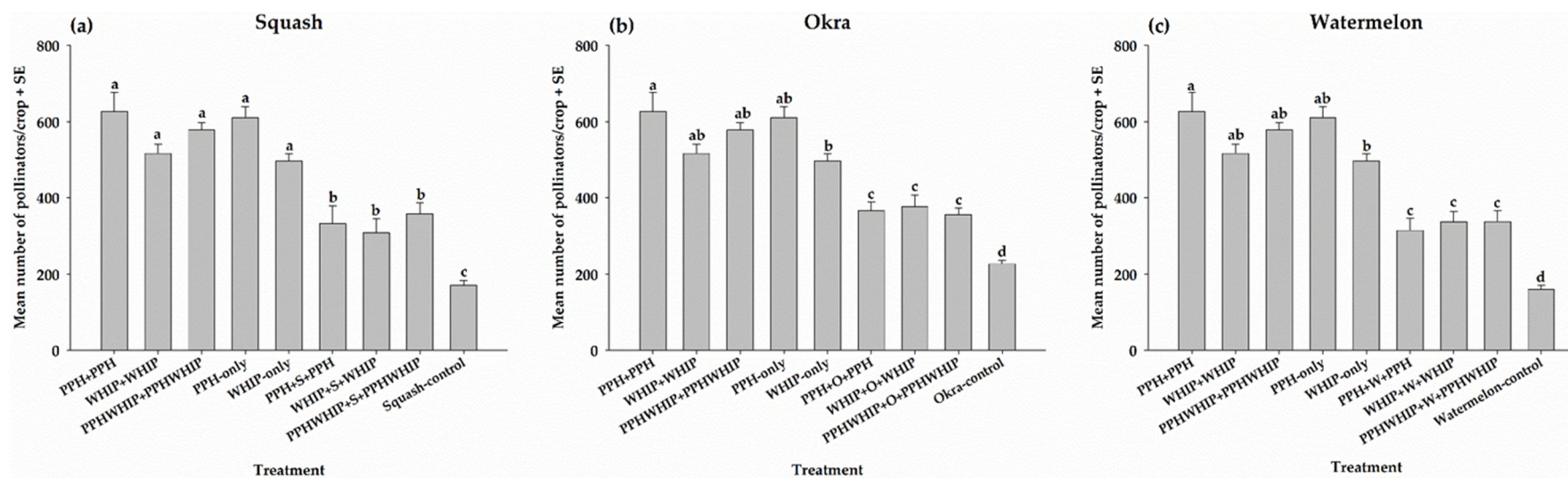


Figure 5. Total pollinators captured on cowpeas and each pollinator-dependent crop ((a) squash, (b) okra, and (c) watermelon) in an intercropping system and control for the entire study period. For each graph, treatment means with the same letters are not significantly different ($p > 0.05$).

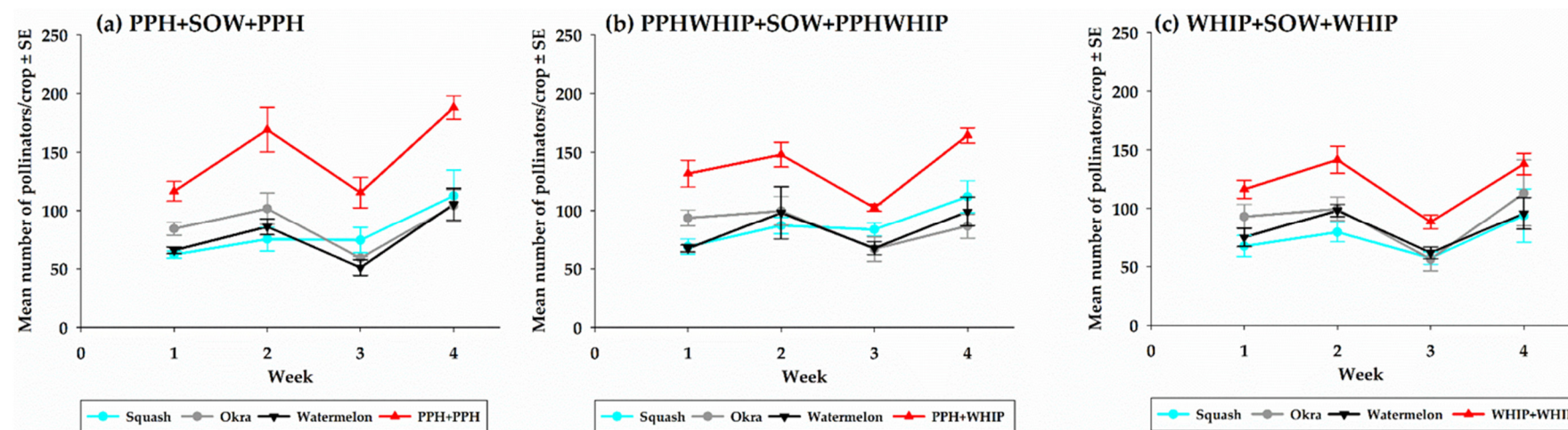


Figure 6. Weekly distribution of pollinators on cowpeas and pollinator-dependent crops for each intercrop treatment ((a) PPH + SOW + PPH treatment, (b) PPHWHIP + SOW + PPHWHIP treatment, and (c) WHIP + SOW + WHIP treatment)).

3.3. Other Beneficial Insects and *H. halys* on Cowpeas and PDCs

The number of *H. halys* on the cowpeas was not significantly different ($F_{4,12} = 1.05$, $p = 0.42$) among the five-cowpea treatments (Figure 7). The number on each of the PDCs intercropped with cowpea and control was not significantly different among treatments; squash ($F_{3,12} = 0.71$, $p = 0.56$), okra ($F_{3,12} = 2.11$, $p = 0.15$), and watermelon ($F_{3,12} = 0.88$, $p = 0.48$), which recorded zero *H. halys*. However, combining all PDCs (SOW), there was a significant difference ($F_{3,9} = 8.87$, $p = 0.0047$) in the number of *H. halys* among treatments, with more recorded on the PDC intercrop (probably due to no *H. halys* recorded on watermelon) than on the control (Figure 8).

Beneficial insects, including Anthocoridae, Berytidae, Dolichopodidae, and Geocoridae, were recorded on all treatments. On the cowpeas, the number of beneficial insects was significantly different ($F_{4,12} = 118.7$, $p < 0.001$), with more on cowpeas that were intercropped than on the sole cowpeas (Figure 7). The number recorded on each of the PDCs intercropped with cowpea was significantly higher on squash ($F_{3,9} = 10.10$, $p = 0.003$) and watermelon ($F_{3,9} = 7.42$, $p = 0.008$) than on the control without cowpea, and similar on okra ($F_{3,9} = 1.99$, $p = 0.19$) and the control. Combining beneficial insects on all PDCs (SOW), there was a significant difference ($F_{3,9} = 17.76$, $p = 0.004$) in the number of beneficial insects on the PDCs intercropped with cowpea than the control without cowpea (Figure 8).

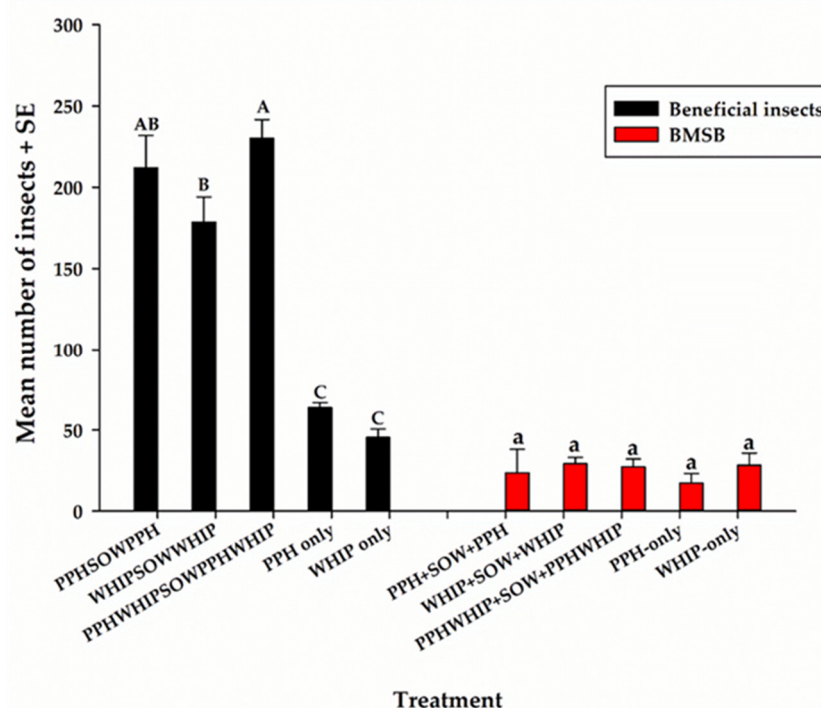


Figure 7. Beneficial insects and *H. halys* recorded on cowpeas planted solely or intercropped with pollinator-dependent crops for the entire study period. Treatment means with the same upper-case letters (for beneficial insects) or the same lower-case letters (*H. halys*) are not significantly different ($p > 0.05$).

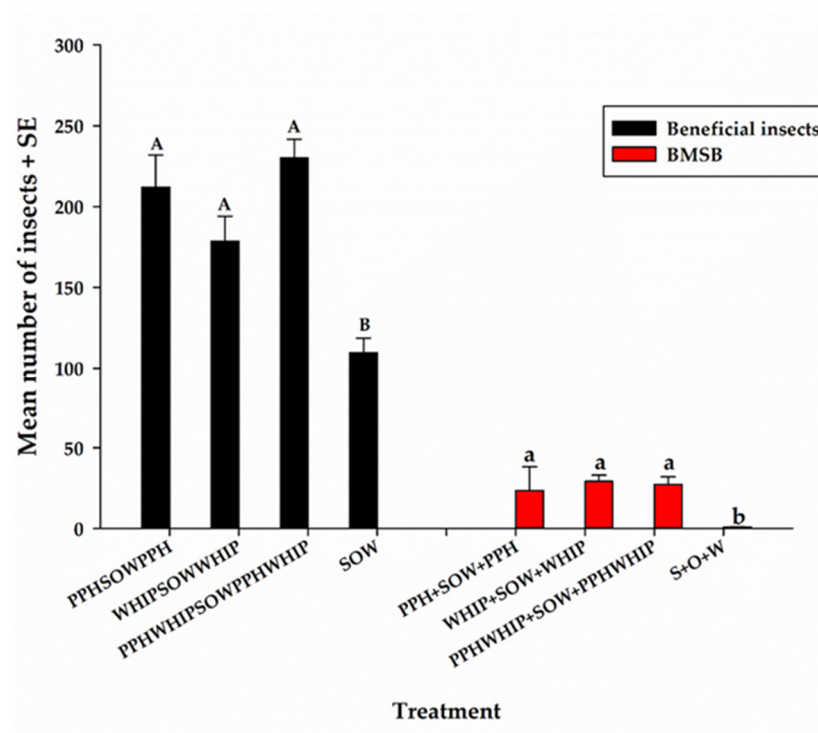


Figure 8. Beneficial insects and *H. halys* recorded on pollinator-dependent crops planted solely or intercropped with cowpea for the entire study period. Treatment means with the same upper-case letters (for beneficial insects) or the same lower-case letters (*H. halys*) are not significantly different ($p > 0.05$).

3.4. Crop Yield

Our results show that in 2018, there was no significant difference ($F_{3,9} = 2.55, p = 0.12$) in squash yield among the three cowpea treatments and the control. However, in 2019 the yield of squash control was significantly lower ($F_{3,12} = 23.9, p < 0.0001$) than the yield in any of the three cowpea intercropped treatments (Table 4). Overall, the squash intercrop produced 27% more fruits than the control. The yield among okra intercropped with cowpea and the control was not significantly different among treatments in 2018 ($F_{3,9} = 1.93, p = 0.20$) or in 2019 ($F_{3,12} = 0.39, p = 0.76$). However, overall, okra intercropped with cowpea produced 7% more fruits than the control (Table 4). The yield of watermelon intercropped with cowpea was significantly higher than the control in 2018 ($F_{3,9} = 14.57, p < 0.0008$) and in 2019 ($F_{3,12} = 11.6, p = 0.0008$). Overall, watermelon intercropped with cowpea produced 54% more fruits than the control. Among the cowpeas, the mean total from both years combined indicates that there was a significant difference ($F_{4,12} = 24.62, p < 0.0001$) in yield (kg), with PPH-only cowpea recording the highest yield (2.0 ± 0.1) > WHIP-only (1.7 ± 0.04) > PPH + PPH (1.6 ± 0.03) > PPH + WHIP (1.5 ± 0.04) > WHIP + WHIP (1.4 ± 0.1).

Table 4. Mean (+ SE) yield (per 5 m row) of pollinator-dependent crops (kg) intercropped with cowpea and control (no cowpea) in 2018, 2019, and combined for both years. For each crop and within each time period, means followed by the same letters are not significantly different ($p > 0.05$).

Treatment	Yield of Pollinator-Dependent Crop (kg)			
	Squash	Okra	Watermelon	
2018	PPH + PPH	29.8 ± 5.8	4.1 ± 0.4	5.5 ± 1.0 ^a
	WHIP + WHIP	34.6 ± 1.8	3.6 ± 0.5	4.9 ± 0.7 ^a
	PPH + WHIP	38.3 ± 4.8	3.0 ± 0.4	5.7 ± 1.6 ^a
	CONTROL	21.2 ± 4.4	3.1 ± 0.1	1.2 ± 0.4 ^b
2019	PPH + PPH	4.6 ± 0.6 ^a	2.4 ± 0.5	11.5 ± 1.9 ^a
	WHIP + WHIP	4.4 ± 0.5 ^a	2.5 ± 0.5	10.1 ± 1.6 ^a
	PPH + WHIP	5.8 ± 0.6 ^a	2.6 ± 0.4	12.2 ± 1.9 ^a
	CONTROL	1.6 ± 0.2 ^b	2.0 ± 0.3	3.8 ± 0.3 ^b
Mean of both years	PPH + PPH	17.2 ± 3.1	3.2 ± 0.3	8.5 ± 1.2 ^a
	WHIP + WHIP	19.5 ± 1.0	3.0 ± 0.5	7.5 ± 1.1 ^a
	PPH + WHIP	22.0 ± 2.5	2.8 ± 0.4	9.0 ± 0.2 ^a
	CONTROL	11.4 ± 2.2	2.6 ± 0.2	2.5 ± 0.1 ^b
Percent change in inter-crop vs. control	+ 26.5%	+ 7.1%	+ 53.7%	

3.5. Relationship between Pollinator Abundance and Crop Yield

There was a positive relationship between pollinator abundance and squash yield for all treatments (Figure 9). A strong and significant relationship was recorded in the squash control ($r = 0.95$; $p = 0.0002$) and squash intercropped with PPH + WHIP cowpea ($r = 0.79$; $p = 0.02$). The relationship in squash intercropped with PPH + PPH cowpea ($r = 0.03$; $p = 0.95$) and WHIP + WHIP cowpea ($r = 0.21$; $p = 0.61$) was weak and not significant. About 95%, 79%, 3%, and 21% of the variation in squash yield can be explained by the relationship between yield and the number of pollinators in the squash control, PPHWHIP + Squash + PPHWHIP, PPH + Squash + PPH, and WHIP + Squash + WHIP cowpea treatments, respectively.

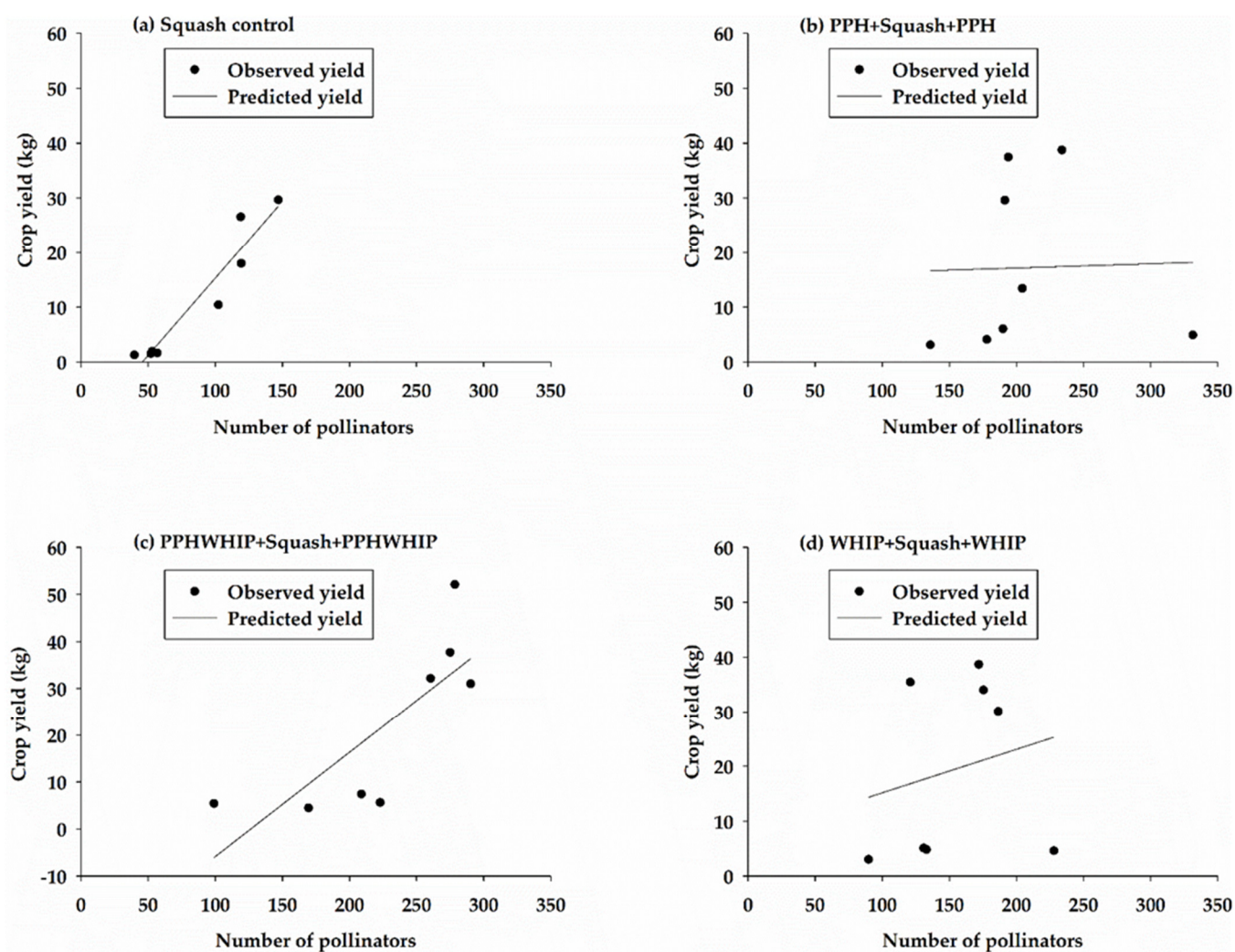


Figure 9. Linear regression showing the relationships between crop yield and number of pollinators recorded on (a) sole squash ($y = -12.8 + 0.3x$; $n = 8$; $r^2 = 0.91$; $p = 0.0002$), (b) squash intercropped with PPH cowpea ($y = 15.6 + 0.01x$; $n = 8$; $r^2 = 0.001$; $p = 0.94$), (c) squash intercropped with PPH + WHIP cowpea ($y = -27.9 + 0.22x$; $n = 8$; $r^2 = 0.62$; $p = 0.02$), and (d) squash intercropped with WHIP cowpea ($y = 7.23 + 0.08x$; $n = 8$; $r^2 = 0.05$; $p = 0.61$).

There was a positive but non-significant relationship between pollinator abundance and yield in the okra control ($r = 0.67$; $p = 0.07$) and in the PPH + Okra + PPH intercrop ($r = 0.47$; $p = 0.24$). A weak negative and non-significant relationship was recorded in PPHWHIP + Okra + PPHWHIP ($r = 0.12$; $p = 0.78$) and in WHIP + Okra + WHIP ($r = 0.05$; $p = 0.91$) (Figure 10). About 67%, 47%, 12%, and 5% of the variation in okra yield can be explained by the relationship between yield and number of pollinators in the okra control, PPH + Okra + PPH, PPHWHIP + Okra + PPHWHIP, and WHIP + Okra + WHIP cowpea treatments, respectively.

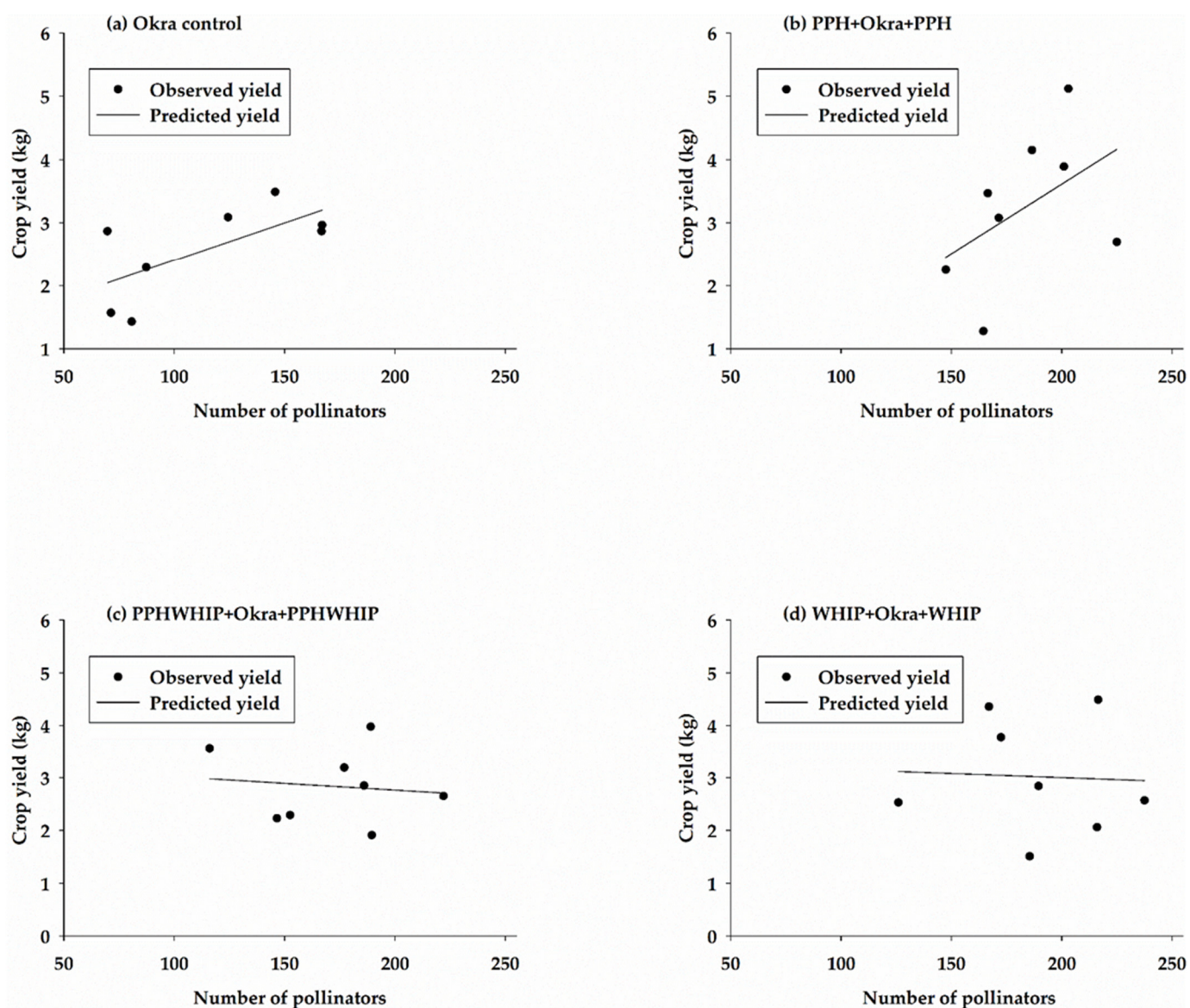


Figure 10. Linear regression showing the relationships between crop yield and number of pollinators recorded on (a) sole okra ($y = 1.2 + 0.01x$; $n = 8$; $r^2 = 0.45$; $p = 0.07$), (b) okra intercropped with PPH cowpea ($y = -0.8 + 0.02x$; $n = 8$; $r^2 = 0.22$; $p = 0.24$), (c) okra intercropped with PPH + WHIP cowpea ($y = 3.3 - 0.003x$; $n = 8$; $r^2 = 0.01$; $p = 0.78$), and (d) okra intercropped with WHIP cowpea ($y = 3.3 - 0.002x$; $n = 8$; $r^2 = 0.003$; $p = 0.91$).

All watermelon treatments except PPHWHIP + Watermelon + PPHWHIP ($r = 0.35$; $p = 0.41$), showed a negative relationship between yield and number of pollinators (Figure 11). There was a strong, moderate, and weak negative relationship observed in the watermelon control ($r = 0.76$; $p = 0.03$), WHIP + Watermelon + WHIP, ($r = 0.47$; $p = 0.25$) and PPH + Watermelon + PPH ($r = 0.18$; $p = 0.67$) treatments, respectively.

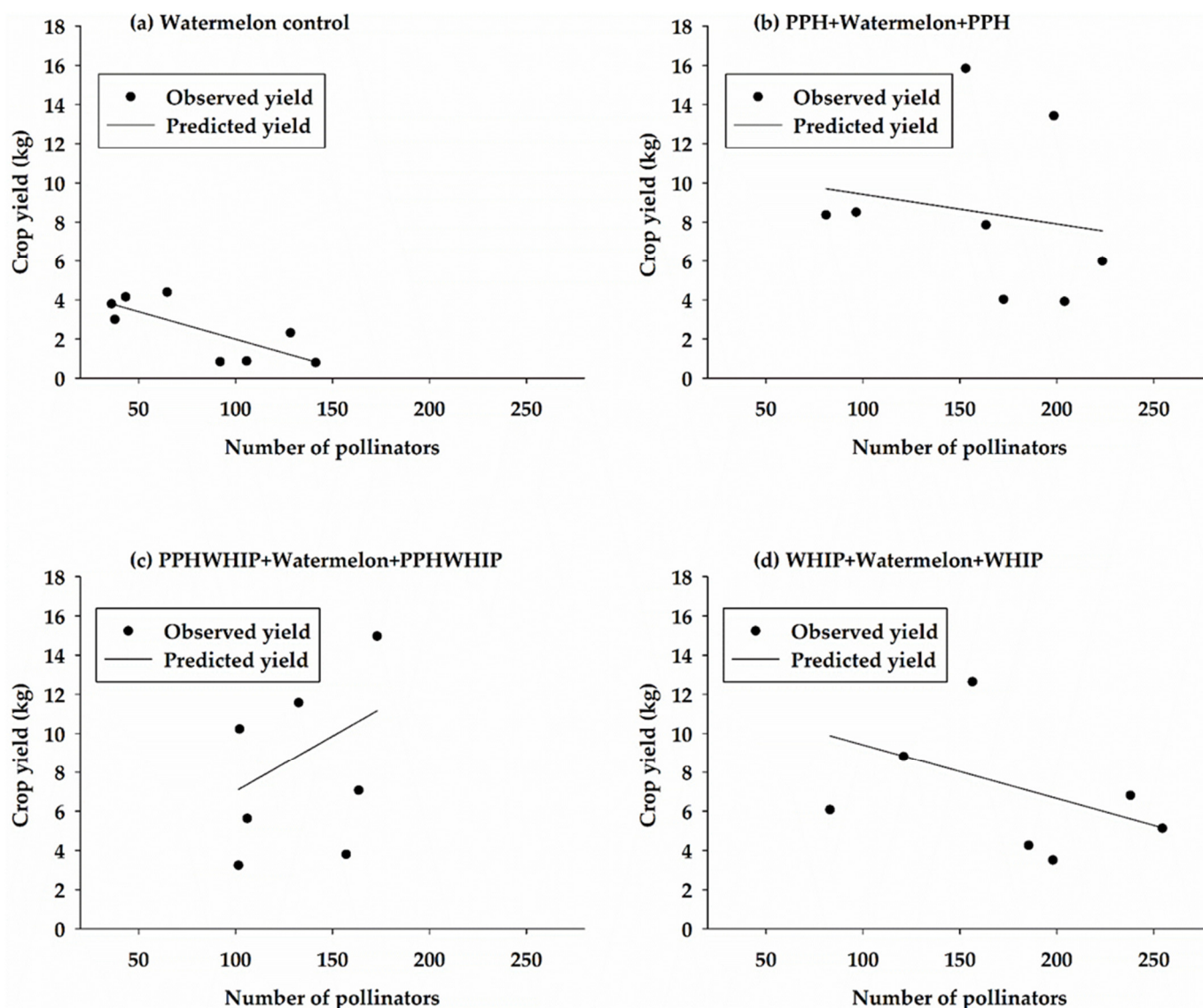


Figure 11. Linear regression showing the relationships between crop yield and number of pollinators recorded on (a) sole watermelon ($y = 4.8 - 0.03x$; $n = 8$; $r^2 = 0.58$; $p = 0.03$), (b) watermelon intercropped with PPH cowpea ($y = 10.9 - 0.02x$; $n = 8$; $r^2 = 0.03$; $p = 0.67$), (c) watermelon intercropped with PPH + WHIP cowpea ($y = 1.4 + 0.06x$; $n = 8$; $r^2 = 0.12$; $p = 0.41$), and (d) watermelon intercropped with WHIP cowpea ($y = 12.2 - 0.03x$; $n = 8$; $r^2 = 0.22$; $p = 0.25$).

4. Discussion

The use of farming systems that encourage the creation of flower-rich habitats to promote ecosystem services such as pollination is crucial at this time, given that global agriculture has expanded and has become increasingly dependent on pollinator services as a result of the increase in the production of pollinator-dependent crops (PDCs). In this study, we evaluated the impact of pollinator abundance and diversity and the presence of other beneficial insects and *H. halys* on crop yield when two cowpea varieties attractive to pollinators were intercropped with three pollinator-dependent crops (squash, okra, and watermelon). This was achieved using three sampling methods (direct visual counts, pan traps, and sticky traps). It has been suggested by several studies that when assessing the abundance and diversity of insect pollinators, the use of multiple sampling methods would provide a better representation of the different types of pollinators in an ecosystem [14,19,39–44]. From our study, diverse pollinator types were recorded. A total of 80,379 pollinators representing seven pollinator families (Andrenidae, Apidae, Crabronidae, Formicidae, Halictidae, Tachinidae, and Vespidae) were captured in pan traps, six families (Apidae, Crabronidae, Halictidae, Pyralidae, Tachinidae, and Vespidae) in sticky

traps, and five pollinator types (bumble bees, carpenter bees, honey bees, butterflies and moths, and wasps) through direct visual counts in both years. This supports the recommendation that different sampling methods are efficient for depicting the different pollinator types, and our findings present a more accurate representation of the diversity of pollinators attracted to cowpea and pollinator-dependent crops. Generally, sticky traps and pan traps catch a proportion of the aerial insect community. Consequently, it is more difficult to attribute the presence and abundance of insects in these traps to specific treatments, especially in small plots like those used in the present study. However, the trapping data support the trends with the direct visual counts of pollinators. The traps also provided an opportunity for measuring diversity within the field.

From each sampling method used in our study, squash, okra, and watermelon intercropped with cowpea resulted in increased pollinator abundance compared to squash, okra, and watermelon planted without cowpea (Tables 1–3). Our findings are similar to other studies that have reported that intercropping with flowering plants or flower strips established adjacent to or within crop fields enhanced pollinator abundance compared to sole cropping. For example, intercropping native and perennial wildflower species adjacent to blueberry farms increased wild bee abundance compared to the control [12]. In another study, flowering basil intercropped with bell peppers resulted in an increased number of pollinators compared to bell peppers without basil [15]. Other studies have suggested that an increase in the number of inflorescences may result in an increase in pollinator visits [45–48]. Pinkeye Purple Hull and Whippoorwill cowpeas produce an average of 49.4 and 56.4 flowers per plant, respectively [19]. These flowers, in addition to those from squash, okra, and watermelon, as well as other floral resources such as nectar and pollen, may have influenced the attraction and abundance of pollinators in the cowpea intercrop.

In most studies, the contribution of the individual crop to the documented outcome has not been investigated. For each treatment and control, pollinator counts were highest on the cowpeas, followed by PDCs intercropped with cowpea and lowest on the control without cowpea (Figure 5). In addition, data from the weekly distribution of pollinators show that among the three cowpea intercrop treatments, there were more pollinators recorded on the cowpeas during each sampling period compared to the PDCs (Figure 6). From our findings, intercropping resulted in more pollinators on the PDCs intercropped with cowpea compared to the control, which had no cowpea (Figure 5). This could be attributed to the consistent high numbers of pollinators on the cowpeas. The flower color and floral resources in cowpea could have contributed to this outcome. It has been reported that colorful flowers induce responses in the form of sensory signals indicating the quantity and quality of floral rewards [20]. In addition to this, the canopy structure of cowpea may have provided more appropriate nesting sites and a suitable microhabitat, as observed on some cowpea varieties [49].

Environmental factors such as temperature, rainfall, and relative humidity can affect pollinator abundance. Data obtained from the State Climate Office of North Carolina [50] indicate that the average relative humidity during the sampling period was 74% in both years; however, the average temperature was 25 °C and 23 °C in 2018 and 2019, respectively. The elevated temperatures in 2018 were accompanied by an increase in pollinator abundance compared to 2019, when temperatures were two degrees lower (Tables 1–3). It is most likely that the temperatures during the sampling period in 2018 could have been optimal for pollinator foraging activity. On the other hand, there was a noticeable decrease in pollinator counts in the third week of sampling (Figure 6). This could be due to the continuous rainfall during this sampling period (22 July to 3 August 2018), where precipitation ranged from 0.05 to 1.11 inches daily, with an average of 0.69 inches [50]. Rainfall has the potential to affect plant–pollinator interactions, as it could have direct physical effects on the floral resources and thus their pollinators.

Creating ways of producing food sustainably requires cropping systems that support biodiversity for increase crop yield [51,52]. From our findings, intercropping with cowpea

resulted in increased pollinator diversity compared to crops without intercropped cowpea. This was accompanied by an increase in crop yield, where okra, squash, and watermelon intercropped with cowpea produced 7%, 27%, and 54% more fruits, respectively, than the controls. Among the insects recorded, pollinators in the Apidae (honey bees and bumble bees) and Halictidae families were the most abundant insects on the intercropped cowpea treatments. Honeybees are essential insect pollinators for all PDCs. Their abundance on the intercrop could have aided pollination efficiency and the observed increase in yield. According to [53], honeybees were the main pollinators of okra flower and resulted in improved yield. The high yield in watermelon and squash could have been due to the combined pollination from honeybees, bumble bees, and halictidae, all of which have been reported to pollinate watermelon and squash and were among the most abundant pollinators recorded on the cowpea intercrops [54–56]. Our findings suggests that pollinators, and subsequently yield, could be increased by intercropping with cowpea varieties that are attractive to pollinators.

On the other hand, pest densities in intercropped systems are reduced as a result of an increase in the density of beneficial insects, resulting in increased crop yield [57,58]. *H. halys* has been reported to be a major pest of cowpea in North Carolina [27]. Even though the number of *H. halys* on the cowpea among the five-cowpea treatments (Figure 7) and the number on each of the PDCs was similar to the control, the yield of each of the PDCs intercropped was higher than the control. This could be due to the presence of more beneficial insects recorded on the three-cowpea intercrop treatments than the control without cowpeas. Intercropping benefitted both the cowpeas and the PDCs in increasing the number of beneficial insects on these crops compared to the controls (Figures 7 and 8). Additionally, the intercropped treatments recorded the highest number of tachinidae (Tables 2 and 3), a family of true flies that are parasitic on some insects. For example, from several cowpea studies, tachinid fauna has been reported to attack cowpea pests such as the cowpea curculio and the pod-sucking bug complex [59]. In addition, in soybean fields, several stink bug species were reported to be parasitized by tachinids [60]. Tachinidae have also been reported to parasitize the leaf-footed bug, the southern green stink bug [61], and the squash bug [62], which are pests of squash and okra. Although all the PDC controls recorded some *H. halys*, watermelon without cowpea recorded no *H. halys*. This could be because in the United States, watermelon is not a host plant of *H. halys*, whereas okra and squash are host plants [63]. This may imply that *H. halys* recorded on the intercropped watermelon probably moved from the cowpeas onto the watermelon.

5. Conclusions

In this study, we demonstrated that the use of Pinkeye Purple Hull (PPH) and Whip-poorwill (WHIP) cowpea varieties in an intercropping system with pollinator-dependent crops (PDCs) squash, okra, and watermelon increased pollinator abundance, diversity, and crop yield. Pollinator counts were higher in all cowpea intercrop treatments than the control; hence, there was no superior cowpea intercrop treatment. Though the number of *H. halys* on the cowpeas and on each of the PDCs was similar to the controls, the yield of each of the PDCs intercropped was higher than the controls, with okra, squash, and watermelon intercrops producing 7%, 27%, and 54% more fruits than the controls, respectively. This suggests that beneficial insects may have kept the pest population below the economic injury level, given that more beneficial insects were recorded on the three-cowpea intercrop treatments than the controls. Overall, an increase in pollinator count resulted in increased crop yield. However, other factors such as pollinator diversity and the presence of pollinators specific to pollinating the PDCs were important. Intercropping with cowpea showed promise in these small field trials. Future research in scaling up the experiment to test intercropping with cowpea on a larger field scale is warranted. We recommend okra, squash, and watermelon intercropped with PPH, WHIP, or both for increase yield.

Supplementary Materials: The following is available online at www.mdpi.com/article/10.3390/su13179612/s1, Table S1: List of the data analyzed using PROC GLIMMIX for specific data types, crop, year, transformation, and distribution and with or without repeated measures.

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