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Pesticides in honey bee colonies: Establishing a baseline for real world exposure over seven years in the USA^{\star}



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ABSTRACT

Honey bees *Apis mellifera* forage in a wide radius around their colony, bringing back contaminated food resources that can function as terrestrial bioindicators of environmental pesticide exposure. Evaluating pesticide exposure risk to pollinators is an ongoing problem. Here we apply five metrics for pesticide exposure risk (prevalence, diversity, concentration, significant pesticide prevalence, and hazard quotient (HQ)) to a nation-wide field study of honey bees, *Apis mellifera* in the United States. We examined samples from 1055 apiaries over seven years for 218 different pesticide residues and metabolites, determining that bees were exposed to 120 different pesticide products with a mean of 2.78 per sample. Pesticides in pollen were highly prevalent and variable across states. While pesticide diversity increased over time, most detections occurred at levels predicted to be of low risk to colonies. Varroacides contributed most to concentration, followed by fungicides, while insecticides contributed most to diversity above a toxicity threshold. High risk samples contained one of 12 different insecticides or varroacides. Exposures predicted to be low-risk were nevertheless associated with colony morbidity, and low-level fungicide exposures were tied to queen loss, *Nosema* infection, and brood diseases.

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1. Introduction

Honey bees forage in a wide radius of approximately 2 km around their colony during resource plentitude and up to 6 km during dearths, collecting both pollen and nectar (Beekman and Ratnieks, 2000; Visscher and Seeley, 1982). Their critical pollination services are valued at \$175 billion worldwide (Gallai et al.,

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2009) and \$17 billion in the US (Calderone, 2012). While foraging in the environment, bees function as living terrestrial bioindicators, picking up traces of heavy metals, pesticides, and other pollutants in the environment (Giglio et al., 2017; Goretti et al., 2020; Smith et al., 2020). Pesticides accumulate in the colony matrix and are often found in the pollen bees consume as their primary protein source (Calatayud-Vernich et al., 2018; de Oliveira et al., 2016). These pesticides play a role in poor bee health (Doublet et al., 2015; Goulson et al., 2015; Sanchez-Bayo and Goka, 2014) and sublethal, chronic pesticide exposures can interact with viruses, parasites, and poor nutrition (Alaux et al., 2010; Poquet et al., 2016; Schmehl et al., 2014; Tosi et al., 2017) leading to decline (Becher et al., 2013;

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Henry et al., 2012; O'Neal et al., 2018; Steinhauer et al., 2018; vanEngelsdorp et al., 2013). The multiplicity of pesticide residues found in the colony complicate analyses (Johnson et al., 2010; Mullin et al., 2010; Traynor et al., 2016a). While current pesticide risk measures focus on individual contaminants (Carnesecchi et al., 2019), different pesticides have non-additive interactions complicating risk assessments in nature.

Honey bees collect and store pollen from plants. It is consumed primarily by nurse bees, which convert it into proteinaceous glandular secretions fed to developing larvae (Crailsheim, 1992). An individual worker will consume over 100 mg of pollen, predominantly when feeding larvae (2012; Crailsheim et al., 1992). Little is known about the actual risk to honey bee health based on the pesticides found in the pollen bees store in their colonies, which is the main protein source nurse bees consume to rear the next generation (Crailsheim et al., 1992; Schmickl and Crailsheim, 2004). Prior surveys suggest high levels of overall pollen contamination in the United States, though we lack scale surveys with samples collected since 2007-2008 (Mullin et al., 2010; Traynor et al., 2016a). We hypothesize that the pesticides found in pollen have changed over the years as pesticide use has shifted, and so seek to establish a baseline of pesticide contamination of pollen. Here we report on the overall pesticide exposure risk from pollen in a subset of colonies randomly surveyed for the National Honey Bee Disease Survey (NHBDS) to determine a baseline of pesticide exposure (for details on sampling see the supplemental information section on sample origin and pollen sampling (Traynor et al., 2016b)) and the potential relationships between pesticides and colony morbidity in the United States.

2. Methods

Freshly stored pollen can easily be identified in a colony by its bright color and matte appearance, indicating the bees collected it recently and thus it is indicative of the pesticides in pollen in the current environment (see supplemental information pollen sampling), whereas honey can be stored for a long time with no change in appearance. We thus focused our survey on freshly stored pollen. Pollen samples were collected from apiaries (n = 1055) in 39 US States and Puerto Rico between 2011 and 2017 (see Table S1 for sampling by state). A subset were simultaneously inspected for overt disease conditions (n = 151, see SI Apiary Inspection Sheet), levels of the ectoparasite Varroa destructor (n = 1048), the spore forming fungal gut parasite *Nosema* spp. (n = 1034), and virus presence (n = 1015). Pesticide contamination of the samples was analyzed by the USDA Gastonia lab for the presence of 218 different pesticide residues using a modified QuEChERS method (Lehotay et al., 2005) that was adapted for 3 g instead of the normal 15 g samples, as amounts greater than 3 g of pollen are hard to obtain from bee colonies (see supplemental information on multiresidue pesticide analysis for details). To understand exposure risk under real world field conditions, we calculated five pesticide risk measures and correlated these with colony morbidity:

- 1) pesticide prevalence (PP): the percentage of samples positive for any pesticide residue
- 2) pesticide diversity (PD): the number of different pesticide residues
- 3) pesticide concentration (in ppb) (PC): the summed concentration of all pesticide residues
- 4) 50+ diversity (50+D): the number of pesticide residues detected, where that residue contributes \geq 50 points to a sample's overall Hazard Quotient score; in particular an HQ score of 50 represents 0.5% of a given pesticide's LD₅₀ consumed over 10 days

5) HQ score (HQ) (Stoner and Eitzer, 2013; Traynor et al., 2016a): an estimate of an adult worker bee's lifetime consumption risk (pesticide residue in ppb/respective LD₅₀ in μg/bee)

Further we summarized exposure patterns by their classification (e.g. fungicides, herbicides, insecticides, and varroacides) and mode of action (Fungicide Resistance Action Committee, 2018; Insecticide Resistance Action Committee) and explored potential relationships between these groups and colony morbidity.

3. Results

The five risk estimates provide different insights into the pesticides bees encounter in pollen (Figs. 1 and 2). The continuous risk variables are all correlated (Table 1). Overall 2933 pesticide detections were made across the 1055 samples, representing 120 of the 218 different active ingredients and metabolites analyzed (Table S2; Table S3). Overall PP was high (Fig. 1A), with 81.9% of all samples contaminated, a stable rate across years (Pearson $X^2 = 9.06$, n = 1,058, df = 6, p = 0.17). When separated by pesticide class (insecticide, fungicide, herbicide, and varroacide), insecticide prevalence decreased over time, while herbicides and fungicides increased (Fig. 2A). Bees were exposed to a diversity of pesticides (Fig. 1B and S1), with up to 21 different residues detected (mean = 2.78 ± 0.09). Varroacides were detected most often, followed by fungicides, and insecticides (Fig. S2A). The varroacides detected with greatest frequency are the currently recommended products for Varroa control (Amitraz metabolite DMPF detected in 45.38% and thymol in 20.63% of tested samples). PD increased over years ($r_{1055} = 0.15$, p < 0.0001; Fig. 2B), with the highest PD occurring in 2014 and 2016. The five pesticide risk values varied by state (Fig. 1, Table S3; **PP**, GLM Binomial $X^2 = 150.18$, df = 39, p < 0.0001; **PD**, GLM Normal X² = 240.56, df = 39, p < 0.0001; **PC**, GLM Poisson X² = 100.50, df = 39, p < 0.0001; **50**+**D**, GLM Normal $X^2 = 143.16$, df = 39, p < 0.0001; HQ, GLM Poisson $X^2 = 220.43$, df = 39, p < 0.0001).

Most pesticides were found at concentrations below 1000 ppb (1 ppm) (Table S2) with some varroacide and fungicide exceptions and a mean concentration of 600.3 ppb \pm 82.0. The concentration did vary across states (Fig. 1C; GLM Normal X² = 56.64, df = 39, p = 0.0336), with NJ (p < 0.001) and NY (p = 0.047) significantly different (Fig. S3). Concentration did not vary across years (Fig. 2C). Several states had mean concentrations \pm SE above our 1000 ppb threshold, a rate previously linked to queen losses for fungicide concentrations (Traynor et al., 2016a) (CA = 1110.0 \pm 228.7; DE = 1228.1 \pm 700.2; IN = 1306.8 \pm 999.7; NJ = 2941.7 \pm 1475.7; NY = 1239.3 \pm 373.1; WV = 1146.1 \pm 753.8).

Many detected products have low recognized toxicity to bees (LD₅₀ in μ g/bee > 100) and so, presumably, pose little risk to bees. To eliminate background noise of low risk exposure, we calculated the diversity of pesticides detected that each contributed 50 or more HQ points (50+D) to a sample's overall score (Fig. 1D). Only 11.9% of detections (n = 349) exceeded the 50+ threshold; the majority (92.0%, n = 321) were insecticides, 7.4% (n = 26) were varroacides, one was a fungicide (THPI), and one was an herbicide (atrazine; Fig. S2B). 50+D differed between sample years (F_{6,1051} = 6.78, p < 0.001) with more detected in 2012 (0.55 ± 0.06 SE) than any other year (0.35 ± 0.04 SE or less).

We classified samples as high risk when they had HQ scores \geq 1000; an amount indicating that honey bees consuming this pollen will ingest 10% or more of their LD₅₀ over their nursing lifetime. Overall 5.4% of samples (n = 57) had HQ scores that exceeded 1000 points (Fig. 1E, Table S2). Fifteen different insecticides and one varroacide (coumaphos) contributed substantially to HQ scores (detected in at least 5 samples and adding more



Scale from 0 (yellow) to 1,000 (red)

Fig. 1. Quantifying pesticide exposure risk in bee bread samples collected as part of the National Honey Bee Disease Survey. (A) Pesticide prevalence: the percentage of samples in each state with one or more pesticide residues. Heat map shows range of positive, from yellow which indicates no samples are positive to dark red where all samples are positive; (B) pesticide diversity: the number of pesticides found per sample, from a scale of 0-10 with yellow indicating an average of no detections and red indicating an average of 10 detections; (C) pesticide concentration: the sum of all detected residues in a sample, with the mean displayed in ppb per state on a scale of 0-1000. We limited the max color scale of deep red to 1000 ppb to illustrate which states regularly meet this threshold concentration of xenobidics, but if increased to 3000 ppb then all states except NJ are shades of yellow to light orange (see Fig. S3); (D) total number of pesticide detections contributing at least 50+ to the hazard quotient, a threshold equivalent to 0.5% of the LD₅₀, used for eliminating trace residues that contribute negligibly to consumption risk; E) mean hazard quotient (HQ) scores per state on a scale of 0-1000, where 1000 is our threshold of high risks. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

than 500 points to at least one HQ score.). Twelve different products (Table S2) in 52 samples had individual pesticide residues that exceeded our 1000 or more point HQ threshold; insecticides: chlorpyrifos (n = 13), clothianidin (n = 8), bifenthrin (n = 6), carbaryl (n = 6), prallethrin (n = 6), thiamethoxam (n = 3), cyfluthrin (n = 2), fenpropathrin (n = 2), permethrin (n = 2), pyridaben (n = 2), imidacloprid (n = 1), and the varroacide coumaphos (n = 1). We analyzed the prevalence and mean HQ contributed by residues that appear in at least 10% of these 57 samples that exceed an HQ of 1,000, contributing at least 100 HQ points. These include chlorpyrifos (n = 33, 57.9%, mean HQ = 877), carbaryl (n = 11, 19.3%, mean HQ = 2931), bifenthrin (n = 10, 17.5%, mean HQ = 1299), clothianidin (n = 8, 14.0%, mean HQ = 2002), thiamethoxam (n = 8, 14.0%, mean HQ = 939), and prallethrin (n = 6, 10.5%, mean HQ = 12,750). Although neonicotinoids were detected only 60 times (2.0% of all pesticide residue detections) throughout our survey, when they are detected they often contribute substantially to the HQ (Table S2). An HQ score above 10,000 indicates that a bee will consume the equivalent of her LD₅₀ over her nursing phase; less than 1% of our samples (n = 5) exceeded this threshold (Table 2). Despite these five extreme risk samples exhibiting high pesticide diversity (mean = 7.4 \pm 0.93 SE), in each of these samples, one insecticide was the main contributor. This pattern of a single insecticide as the predominant contributor to exposure risk is common across our 57 high risk samples (Fig. S3A), except for the five samples from Nebraska where the two neonicotinoids clothianidin and thiamethoxam are frequently co-detected, potentially because the former is the insecticidal metabolite of the latter (Nauen et al., 2003). This single residue as the main contributor to

Fig. 2. Pesticide class trends over time. (A) Pesticide prevalence of positive detections; (B) pesticide diversity; (C) pesticide concentration, (D) 50+ detections by class; and (E) hazard quotient score). Grey = all pesticide detections, yellow = insecticides, orange = fungicides, green = herbicides, red = varroacides. Differences between years (α = 0.05) are marked with different letters. Significant trends over survey years are also indicated by reporting GLM results with binomial distribution for prevalence, normal for diversity and 50+ diversity, and Poisson for concentration and HQ; n. s. indicates that no significant (p > 0.05) trends were found. Trend lines show a linear regression with 95% confidence intervals. The first and last survey years covered 6 non-overlapping months (2011 = July–Nov., 2017 = March–June), so differences from general trends in these years should be viewed cautiously. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

Table 1

Correlations of different risk measures and parasite loads. We report all significant correlations between our four continues estimates of risk (diversity (PD), concentration (PC), 50+D, and HQ, while excluding prevalence as that is binary). We then also compare our continuous risk measures with parasite (*Varroa* and *Nosema*) loads.

Risk Factor	Compared with	Correlation	Count	Lower 95%	Upper 95%	P value
PD	50+D	0.508	1055	0.46	0.55	1.4×10^{-70}
HQ	50+D	0.340	1055	0.29	0.39	$5.9 imes 10^{-30}$
PD	HQ	0.218	1055	0.16	0.27	7.3×10^{-13}
PD	PC	0.163	1055	0.10	0.22	$1.0 imes 10^{-7}$
PC	50+D	0.142	1055	0.08	0.20	$3.6 imes 10^{-6}$
PC	HQ	0.141	1055	0.08	0.20	4.3×10^{-6}
Parasite Loads						
PD	Varroa	-0.153	1048	-0.21	-0.09	6.1×10^{-7}
Nosema	Varroa	-0.087	1034	-0.15	-0.03	0.005
PD	Nosema	0.086	1034	0.03	0.15	0.006
HQ	Varroa	0.065	1048	0.00	0.12	0.036
50+D	Nosema	0.062	1034	0.00	0.12	0.047

HQ risk contrasts with the diversity of residues found in samples with high pesticide concentrations where multiple different products co-occur (Fig. S3B).

Our risk measures varied by pesticide class (Fig. 3). Varroacides, applied by beekeepers directly into colonies to reduce parasite loads, were the main contributor for PP (Fig. 3A), PD (Fig. 3B), and PC (Fig. 3C), though PC decreased significantly over time (Fig. 2C). Varroacides were detected 1226 times (Fig. S2A), with at least one varroacide detected in 65.9% of samples (n = 697), and contributing 0.04 to 1190 points to the HQ score of samples; the majority (99.51%) of varroacide detections contributed less than 50 points to the HQ score. Thymol, considered practically non-toxic to bees ($LD_{50} = 975 \mu g/bee$), contributed little to most HQ scores although it was detected at more than 1000 ppb in 4.1% of positive samples (n = 43) (Table S2) and at over 10,000 pbb in 0.7% of samples (n = 7). Other varroacides detected at over 1000 ppb were the amitraz metabolite DMPF (n = 2), coumaphos (n = 4) and fluvalinate (n = 5).

Insecticides were the second most prevalent pesticide residue class, detected 691 times (Fig. S2A), with at least one insecticide detected in 38.1% of samples (n = 404). Insecticide PP was highest in 2012 (47.5% \pm 0.04% SE) and lowest in 2017 (26.9% \pm 0.05% SE). Insecticide PD varied by year (Fig. 2B). Insecticides contributed the majority of the 50+D detections (Figs. S2B and 3D), and varied

significantly by year. Overall insecticides contributed little to the PC (Fig. 3C), though the concentration increased significantly over time (Fig. 2C). Six different insecticides were found in 14 different samples at concentrations that exceeded 1000 ppb (Table S2). Many insecticides are highly toxic to bees, so it is not surprising that as a pesticide class they contributed the most (95.9%) to the HQ score (Fig. 3E), adding a mean of 646.15 \pm 110.67 SE (range: 0.01 to 29,629.6) points to samples with at least one insecticide detected.

Fungicides were detected 641 times, with at least one detection in 29.5% of samples (n = 312). Fungicide PP increased over years $(r_{1055} = 0.13, p < 0.001, Fig. 2A)$. Fungicide PD also increased $(r_{1055} = 0.19, p < 0.001)$ and varied between years (Fig. 2B). Multiple fungicides were frequently detected; the maximum number of fungicides increased annually from four fungicides in 2011 to ten fungicides in 2017. Altogether 9.4% (n = 30) of the fungicidepositive samples contained five or more fungicides, disproportionately from California (32.5%). After varroacides, fungicides were the most common (22%) (Fig. S2A) and largest contributor to the PC (Fig. 3C). Fungicide PC didn't change over time (Fig. 2C). Eight different fungicides in 25 samples contributed 1000 or more ppb (Table S2; Fig. S4B). Of note is the fungicide tetrahydrophthalimide (THPI) which was detected in 22 samples, contributing a mean of 1526.6 \pm 500.9 SE ppb and 3.14 \pm 0.49 SE HQ points to each sample's concentration (range: 1 to 7060) and HQ score (range: 0.01 to

Table 2

Samples with HQ scores above 10,000 and constituent products

Sample ID	95,350	93,274	93,497	96,361	96,025
State	OR	OR	MD	WV	NJ
Year	2013	2013	2013	2016	2016
HQ Score	19,097.0	29,643.3	11,928.9	11,771.0	16,146.5
Detections	9	6	5	10	7
1	Acephate	Azoxystrobin	Atrazine	DMPF	1-Naphthol
	1.9	0.2	0.1	1.0	226.2
2	Azoxystrobin	Coumaphos	Cyhalothrin total	Atrazine	DMPF
	0.1	10.9	10.0	0.01	0.9
3	Chlorpyrifos	Coumaphos oxon	Fluvalinate	Azoxystrobin	Azoxystrobin
	87.9	0.2	7.4	0.2	0.4
4	Endosulfan II	Fluvalinate	Prallethrin	Boscalid	Carbaryl
	6.6	1.5	11,888.9	0.3	14,932.13
5	Endosulfan sulfate	Prallethrin	Thymol	Captan	Chlorantraniliprole
	0.3	29,629.6	22.6	2.5	0.81
6	Fluvalinate	Thymol		Carbaryl	Chlorothalonil
	0.9	1.0		11,764.7	6.1
7	Methamidophos			Carbendazim	Imidacloprid
	36.1			0.8	979.9
8	Prallethrin			Myclobutanil	
	18,963.0			0.5	
9	Thymol			Pyraclostrobin	
	0.2			0.6	
10				Pyriproxyfen	
				0.4	

Carbaryl is a widely used agricultural insecticide, predominantly used for insect pest control (aphids, ticks, fleas, etc.).

75.16). THPI is the major metabolite of captan, a widely used fungicide that caused brood mortality in cage trials when integrated into larval food (Mussen et al., 2004), but not in field trials (Everich et al., 2009). Fungicides were seldom (<1%) found to contribute 50+ points to the HQ score (Fig. S2B). HQ_{Fung} scores below 5 were implicated in colony losses in a prior study (Traynor et al., 2016a), hence this risk assessment metric may underestimate the number of significant exposures for products with low toxicity and underestimate potential synergies. Fungicides generally contributed little to a sample's overall HQ score (Fig. 2D), so we also calculated which fungicide modes of actions (MOA) were found contributing five or more HQ points (see below).

Herbicides were found in 24% of samples (Fig. S2A). Herbicide PP ($r_{1055} = 0.19$, p < 0.001) and PD ($r_{1055} = 0.20$, p < 0.001) increased over the course of the study (Fig. 2A and B). Herbicide PC was low (Fig. 3C), and did not change over time (Fig. 2C). The herbicides atrazine (n = 1), fluridone (n = 2), metolachlor (n = 2), and propachlor (n = 3) were detected at concentrations above 1000 ppb (Table S2). Only one detection of atrazine added 50+ points to the HQ; overall herbicides added a mean of 1.29 ± 0.38 SE HQ points to samples with at least one herbicide detection (HQ range: 0.01 to 87.32). No herbicide was found at levels which exceeded our HQ safety threshold of 1000 points, though our survey didn't-test for glyphosate, previously linked to disruptions of gut microbiota (Motta et al., 2018), as it requires a separate analysis.

Pesticides have different modes of action (MOA), and some pesticide MOAs have previously been linked to colony morbidity (Böhme et al., 2018; Douglas et al., 2020; Traynor et al., 2016a), hence we grouped detections by MOA and calculated exposure risk (Table S4; Fig. S5). The most frequently detected insecticide and varroacide MOA groups were Group 1 Acetyl Choline Esterases (AChE) and Group 3 Sodium Channel Modulators. These specific MOAs contributed more than 50+ in 324 pollen samples, accounting for 50.1% and 33.9% of 50+ detections in all samples (Table S4). The small contribution fungicides make to HQ scores was mostly made by fungicide products with an MOA defined as M. multisite activity, F. lipid synthesis or transport and C. respiration by the fungicide resistance action committee (Fungicide Resistance)

Action Committee, 2018) (Table S4; Fig. S5).

Varroa destructor (n = 1048) and Nosema spp. (n = 1034) infestation were calculated for most apiaries (Table 3) and correlated with our five measures of risk (Table 1). Overall and varroacide PP, overall and varroacide PD, 50+D, and HQ scores varied across varroa infestation levels (Figs. S5A-F). PP was higher in groups with less than 3 mites, while samples with no mites had more varroacides present than other groups (Fig. S5A, red). Total PD is negatively correlated with Varroa infestation ($r_{1048} = -0.153$, p < 0.0001). Varroacide PD was higher in Varroa free colonies than in samples taken from colonies with detectable Varroa levels (Fig. S5B), suggesting they were being or had recently been treated for this parasite. Overall PC did not vary with Varroa load (Fig. S5C), but varroacide levels were highest in samples with 5 or more mites. When we investigated the major contributors to varroacide PC, we found a surprising inverse relationship between the four most commonly used varroacides, suggesting that amitraz and coumaphos are effective control products, while high levels of thymol and fluvalinate are detected in samples from colonies with high varroa loads (Fig. 4) Interestingly, for both 50+D and total HQ we found the highest scores in the Varroa infestation at zero and at 10+ mites per 100 bees (Figs. S5D and E). Elevated HQ levels in the 10+ Varroa infestation group suggests that either Varroa are more fit in environments of high pesticide exposure (e.g. by perhaps increasing the length of brood development providing Varroa with greater fecundity), or that elevated HQ contamination of pollen may reduce a colony's adult bee population (e.g. by shortening the lifespan of adult bees thus increasing the density of Varroa per bee), resulting in a greater density of Varroa per bee (Gill et al., 2012; Wu et al., 2011).

PD, fungicide PD, fungicide PC, 50+ D and fungicide HQ were significantly higher in *Nosema* positive samples (Fig. 5), a counterintuitive result as *Nosema* is a fungal spore disease. We suspect that just as antibiotics can wipe out beneficial intestinal flora in the gastrointestinal tract, and allow harmful flora to establish (Pamer, 2016), exposure to fungicides may destroy the beneficial fungi in a colony, permitting fungal diseases like *Nosema* to proliferate as seen in a prior study where fungicide PC increased the probability

Fig. 3. Exposure risk assessment by pesticide class, mean ± SE reported for all. (A) Pesticide prevalence of positive detections; (B) pesticide diversity; (C) pesticide concentration; (D) 50+ detections by class; and (E) HQ score; yellow = insecticides, orange = fungicides, green = herbicides, red = varroacides. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

of *Nosema* infection (Pettis et al., 2013). Apiaries with undetectable *Nosema* exhibited 20.6% lower PD compared to apiaries with *Nosema* (Fig. 4B). Samples with *Nosema* had fungicide HQ scores 3x higher than *Nosema* free samples (Fig. 3E). *Nosema* spore load was loosely correlated with PD ($r_{1032} = 0.086$, p = 0.0055), fungicide PD ($r_{1032} = 0.084$, p = 0.0068) and the fungicide HQ score ($r_{1032} = 0.085$, p = 0.0065).

14.3% of apiaries (n = 151) were simultaneously inspected for overt symptoms of honey bee diseases (Table 3). Brood diseases (American foulbrood, European Foulbrood, Sacbrood, Chalkbrood, and/or Snot Brood) were detected in 29.8% of inspected apiaries (n = 45). The fungicide HQ was elevated in samples from apiaries positive for brood disease ($X^2 = 3.98$, df = 1, p = 0.046) with a mean

fungicide HQ = 4.01 \pm 1.81 SE in positive samples compared to fungicide HQ = 1.20 \pm 0.45 SE in colonies free of brood disease. About one third of apiaries (n = 49) experienced queen issues, where at least 1 of 8 colonies inspected was a drone layer, queenless, or the colony had queen cells (Table 3). Colonies experiencing queen issues had elevated fungicide HQ scores (X² = 3.84, df = 1, p = 0.049) and an average fungicide HQ score 3x higher (3.82 \pm 1.73 SE) than colonies without queen issues (1.18 \pm 0.41 SE).

Lastly, we examined how our five exposure risk measures varied with viral prevalence for eight common honey bee viruses (Table 4). Each was associated with either increased or decreased prevelence of at least one of the viruses examined except Acute bee paralysis virus (ABPV) (Table 4).

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Table 3

Prevalence of parasites, pathogens, overt brood disease, and queen conditions in inspected apiaries.

Condition	Sampled Apiaries	Prevalence	Load (mean \pm SE)
Parasites & Pathogens			
Varroa	1048	89%	4.05 ± 2.2 mites per 100 bees
Nosema spores	1034	43.7%	0.32 ± 0.026 million spores per bee
Viruses			
Acute Bee Paralysis virus (ABPV)	993	23.2%	N/A
Black Queen Cell Virus (BQCV)	368	80.4%	N/A
Chronic Bee Paralysis Virus (CBPV)	992	9.6%	N/A
Deformed Wing Virus (DWV)	1002	85.2%	N/A
Israeli Acute Paralysis Virus (IAPV)	991	13.7%	N/A
Kashmir Bee Virus (KBV)	912	9.2%	N/A
Lake Sinai Virus II (LSV II)	629	36.1%	N/A
Varroa Destructor Virus (VDV)	282	56.0%	N/A
Overt Brood Disease			
American Foulbrood	151	<1%	1 colony per infected apiary
European Foulbrood	151	5%	1.3 colonies per infected apiary
Sacbrood	151	7.2%	1.4 colony per infected apiary
Chalkbrood	151	13.2%	1.9 colony per infected apiary
Snot Brood and/or Parasitic Mite Syndrome	151	11.9%	2.5 colony per infected apiary
Deformed wing	151	15.2%	1.4 colony per infected apiary
Black Shiny Bees	151	5.3%	2.3 colony per infected apiary
Queen Issue			
Queen Cells	151	16.6%	1.9 colony per infected apiary
Drone Layer	151	4.0%	1 colony per infected apiary
Queenless	151	8.6%	2.2 colony per infected apiary

Fig. 4. The concentration of the four most commonly used varroacides by varroa infestation, mean \pm SE. The amitraz metabolite DMPF and coumaphos both exhibit high concentrations at samples with no varroa, while fluvalinate and thymol show the inverse relationship with high concentrations appearing in samples with high varroa infestations. This suggests that amitraz and coumaphos may be effective and rapid varroa controls, while fluvalinate and thymol are either ineffective or are applied when beekeepers detect high infestations and dissipate quickly post treatment. Varroa Infestation per 100 adult bees: 0 = 0, <3 = 0 to 3 varroa, <5 = 3 to 5 varroa, < 10 = 5 to 10, and 10+ = more than 10. For each reside, we conducted a GLM analysis with exponential distribution.

4. Discussion

Our national survey is the first to broadly examine environmental pesticide exposure throughout a wide majority of the United States, establishing an important baseline of pesticide pollution of pollen collected by our most important managed pollinator, *Apis mellifera*. Honey bees forage in a wide radius around their colony, functioning as a terrestrial biomonitor that provides critical insight into the contaminants they encounter and bring back to the nest. Here we link field measurements of colony

Fig. 5. Exposure risk assessment by *Nosema* spp. spore level, mean \pm SE. (A) Pesticide prevalence of positive detections; (B) pesticide diversity; (C) pesticide concentration; (D) 50+ detections; and (E) HQ score, with left y-axis for all pesticides and right y-axis for fungicides. Light grey = all pesticides; dark grey = fungicides. Significant differences ($\alpha = 0.05$) marked with different letters. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

morbidity to five different ways of summarizing pesticide exposure and risk in pollen, identifying clear areas of concern. Eleven insecticides and one varroacide were the greatest contributors to HQ consumption risk. Because chlorpyrifos, carbaryl, bifenthrin, clothianidin, thiamethoxam, and prallethrin occurred most frequently in samples we considered high risk, our work raises concern about their current use.

Based on standard lethal effects indexes, most of our samples were not considered risky. Many of our 57 high-risk samples had one main contributor, typically an insecticide. Although fungicides contribute little to direct consumption risk, they are often found at high concentrations and were correlated with increased *Nosema* infections, brood disease and queen issues, reaffirming that a reevaluation of their safety is necessary (Fisher et al., 2017; Simon-Delso et al., 2018; Traynor et al., 2016a; Wade et al., 2019). High levels of varroa were associated with high levels of fluvalinate and thymol, two different varroacide products, suggesting that either beekeepers applied them after noticing high varroa levels and then residues dissipated quickly when varroa mites were controlled or that they are less effective control products and varroa infestation levels escalate despite high dose treatments. In contrast, low levels of varroa were associated with high levels of the amitraz metabilite DMPF and the synthetic varroacide coumaphos, potentially suggesting that these two are effective varroa control products that quickly reduce varroa infestations. Our work thus underlines the complexity of pesticide effects, highlighting that they are not limited to lethal exposures. Colony morbidity was often associated with low-risk exposure suggesting the myriad residues detected may result in poorly understood interaction (i.e. additive, antagonistic, synergistic) effects. While our results provide little evidence of poisoning as a major colony loss driver, we stress the mounting evidence that colonies experience sublethal pesticide exposure and unexpected interactions arise when bees are co-exposed to diseases, as well as chemical mixtures (18) and nutritional stress (9). Our results suggest an urgent need for greater understanding of pesticide impacts on bee losses.

Pesticide residues detected and the amount of points contributed to the HQ score for each of the six samples above 10,000. Residues that contributed more than 1000 points in bold. For 4 of the 5 samples, one residue was the main contributor to the score. The NJ sample had two residues that contributed substantially to the score: carbaryl and imidacloprid.

Table 4

Exposure risk and virus status.

		Negative		Positive		Significance	
		Mean	SE	Mean	SE	t-test	р
РР	ABPV	81.4%	1.4%	82.6%	2.5%	0.42	0.67
	BQCV	68.1%	5.5%	83.4%	2.2%	2.59	0.0110
	CBPV	81.8%	1.3%	80.0%	4.0%	-0.42	0.67
	DWV	84.4%	2.3%	81.3%	1.0%	-0.94	0.35
	IAPV	80.8%	1.3%	86.8%	2.9%	1.85	0.07
	KBV	81.5%	1.0%	86.9%	4.0%	1.37	0.17
	LSV	79.4%	2.0%	88.1%	2.0%	2.96	0.0030
	VDV	83.9%	3.0%	86.7%	3.0%	0.66	0.51
PD	ABPV	2.81	0.10	2.67	0.17	-0.68	0.49
	BQCV	2.10	0.28	2.68	0.15	1.86	0.07
	CBPV	2.81	0.10	2.53	0.26	-1.00	0.32
	DWV	3.27	0.27	2.68	0.09	-2.05	0.0410
	IAPV	2.73	0.10	3.09	0.26	1.30	0.20
	KBV	2.81	0.10	2.88	0.29	0.22	0.82
	LSV	2.53	0.13	3.57	0.22	3.99	< 0.0001
	VDV	2.72	0.24	3.74	0.29	2.67	0.0081
PC	ABPV	527.35	89.26	616.80	137.28	0.55	0.59
	BQCV	414.29	183.02	591.76	156.26	0.74	0.46
	CBPV	540.25	80.33	627.24	221.66	0.37	0.71
	DWV	374.03	78.13	626.43	98.24	2.01	0.0450
	IAPV	511.69	80.47	784.26	219.98	1.16	0.25
	KBV	503.55	57.67	1165.67	677.42	0.97	0.33
	LSV	647.84	165.11	539.15	95.21	-0.57	0.57
	VDV	257.19	57.57	560.72	123.45	2.23	0.0270
50+D	ABPV	0.33	0.02	0.31	0.04	-0.24	0.81
	BQCV	0.31	0.08	0.42	0.04	1.20	0.23
	CBPV	0.34	0.02	0.17	0.04	-3.42	0.0008
	DWV	0.34	0.05	0.32	0.02	-0.50	0.62
	IAPV	0.33	0.02	0.26	0.05	-1.24	0.22
	KBV	0.32	0.02	0.26	0.06	-0.85	0.40
	LSV	0.28	0.03	0.27	0.04	-0.30	0.76
	VDV	0.15	0.04	0.27	0.05	2.05	0.0410
HQ	ABPV	249.42	149.67	315.48	109.22	0.57	0.57
	BQCV	207.98	87.27	244.76	39.09	0.38	0.70
	CBPV	282.96	50.52	91.11	52.79	-2.62	0.0091
	DWV	394.96	205.28	239.62	39.90	-0.74	0.46
	IAPV	240.84	38.09	415.87	235.22	0.73	0.46
	KBV	282.48	54.28	141.12	55.46	-1.82	0.07
	LSV	339.00	103.62	180.21	58.45	-1.33	0.18
	VDV	144.90	96.65	237.10	106.52	0.64	0.52

Imidacloprid is a neonicotinoid, frequently applied as seed treatment where bees can contact the planting dust, but also used as a foliar spray, soil drench, and injected directly into trees, i.e. to protect against citrus greening.

Prallethrin is used primarily for mosquito control and in products to kill wasps/hornets.

For each of our five estimates of exposure risk, we report the mean \pm SE for samples that were negative or positive for eight common honey bee viruses.

Viruses abbreviated: Acute Bee Paralysis Virus (ABPV), Black Queen Cell Virus (BQCV), Chronic Bee Paralysis Virus (CBPV), Deformed Wing Virus (DWV), Israeli Acute Paralysis Virus (IAPV), Kashmir Bee Virus (KBV), Lake Sinai Virus II (LSV II), Varroa Destructor Virus (VDV).

Author statement

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Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

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Further reading

Insecticide Resistance Action Committee, The IRAC Mode of Action Classification Online.

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