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# Polychlorinated biphenyl and polybrominated diphenyl ether profiles in serum from cattle, sheep, and goats across California

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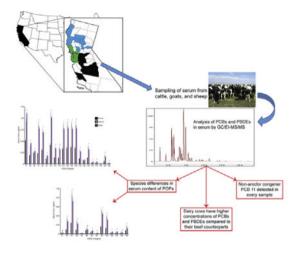
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#### **Abstract**

It has been previously been shown by our lab and others that persistent organic pollutants, such as polychlorinated biphenyls (PCBs) and polybrominated diphenyl ethers (PBDEs), are contaminants in milk produced for human consumption. To further this research we determined the concentration of 21 PCB and 14 PBDE congeners in livestock serum, mainly bovine, across California. Congeners were extracted from serum using solid phase extraction (SPE), cleaned up by silica cartridge and quantified using gas chromatography-triple quadruple mass spectrometry. We detected significant differences among species and the production class of cattle (beef or dairy). The sum of all 21 PCB congeners (ΣPCBs) in caprine and ovine sera had a mean value of 9.26 and 9.13 ng/mL, respectively, compared to 3.98 ng/mL in bovine sera. The mean value for the sum of all 14 PBDE congeners (ΣPBDEs) in caprine and ovine sera was 2.82 and 2.39 ng/mL, respectively, compared to 0.91 ng/mL in bovine sera. Mean ΣPCBs in dairy cattle was 5.92 ng/mL compared to 2.70 ng/mL in beef cattle. Mean ΣPBDEs in dairy cattle was 1.33 ng/mL compared to 0.70 ng/mL in beef cattle. There were no regional differences in the ΣPCBs or ΣPBDEs in cattle distributed across California. These results highlight the fact that livestock are still being exposed to these pollutants yet little is known about where this exposure may be coming from.

#### **Graphical Abstract**

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#### Keywords

Bovine; Caprine; Ovine; Serum; PCB; PBDE

#### 1. Introduction

Polychlorinated biphenyls (PCBs) and polybrominated diphenyl ethers (PBDEs) are synthetic industrial chemicals that have become widespread persistent organic pollutants (POPs) and are resistant to biodegradation leading to their persistence in our environment. Both of these classes of POPs contain 209 different congeners with varying degree of halogenation and position of halogen atoms on their aromatic rings. One of the key characteristics shared amongst POPs are their high lipophilicity leading to their ability to accumulate in fatty animal tissue, thus leading to diet being an important source of exposure to these chemicals for humans (Schecter et al., 2010; Chan-Hon-Tong et al., 2013; Cimenci et al., 2013; Ampleman et al., 2015). Human exposure is a point of concern because epidemiologic studies have implicated PCBs and PBDEs in a variety of adverse health effects including immune system dysfunction (Jusko et al., 2012; Kramer et al., 2012), endocrine disruption (Abdelouahab et al., 2011; Silverstone et al., 2012; Valvi et al., 2012), and deficits in neurodevelopment (Schantz et al., 2003; Korrick and Sagiv, 2008; Herbstman et al., 2010; Winneke, 2011; Gascon et al., 2012; Eskenazi et al., 2013). One form of dietary exposure that has a relatively high fat content is milk. Since these toxic compounds are highly lipophilic, milk has been thought to be a prominent source of exposure for humans (Kim et al., 2008; O'Donovan et al., 2011). Many countries have analyzed bovine milk made within its borders for PCB and PBDE content and have found a multitude of congeners at quantifiable amounts (Focant et al., 2003; Durand et al., 2008; Kim et al., 2013; Lake et al., 2013). To date, there has only been one study within the United States (U.S.) looking at a small subset of PCBs in milk samples collected across the country (Schaum et al., 2003), not allowing for any region specific conclusions on PCB content or detection of PBDEs. To expand on this previous study, and since California produces approximately 20% of the total milk supply in the U.S., we analyzed milk samples made in California and discovered PCBs

and PBDEs at quantifiable levels (Chen et al., 2017). As a follow up we conducted this current study to pursue these same pollutants in bovine serum, a matrix not yet investigated in the U.S in terms of PCB and PBDE contamination. A previous study from Italy has shown a difference in PCB content between ovine and bovine samples (Benedetto et al., 2016), which led us to expand the scope of this study to include ovine and caprine serum samples for assessing possible species differences.

PCBs are categorized into dioxin-like (DL) and non-dioxin-like (NDL) compounds. DL PCBs are potent activators of the aryl hydrocarbon receptor (Vondracek et al., 2005). Previous assessment of POP contamination of dairy products have primarily evaluated DLcompounds (Focant et al., 2003; Durand et al., 2008; Pizarro-Aranguiz et al., 2015) because, in the past, these compounds were considered to be the most toxic. However, NDL-PCBs dominate over DL-PCBs in biological and environmental samples (DeCaprio et al., 2005), and specifically multiple NDL-PCBs have been shown to be potent neurodevelopmental toxicants (Schantz et al., 1997; Howard et al., 2003; Yang et al., 2009; Yang and Lein, 2010; Wayman et al., 2012; Lesiak et al., 2014; Yang et al., 2014). Many of the previous studies with focus on DL-compounds also do not include assessment of PBDEs leaving out an entire class of compounds implicated as neurodevelopmental toxicants (Chen et al., 2012; Bradner et al., 2013; Behl et al., 2015; Jarema et al., 2015). PBDEs have been used extensively in products in California until their proposed ban in 2003. Following the ban in 2004, two commercial formulations, penta-BDE and octa-BDE, were phased out of production in some U.S. states after a voluntary agreement between the U.S. EPA and the sole manufacturer of these products (Dodson et al., 2012a). Despite the phase out of many PBDEs used in industry, these compounds persist in our environment due to their resistance to biodegradation (Dodson et al., 2012b; Bradman et al., 2014; Whitehead et al., 2015). Thus, we focused our study on mainly NDL-PCBs and PBDEs with potential neurodevelopmental toxicity as this is a highly sensitive endpoint of concern for these POPs. In addition, previous work assessing PBDEs and PCBs in bovine milk for human consumption detected a nonlegacy PCB, or a PCB that was never intentionally synthesized for industrial purposes, PCB 11, at quantifiable levels (Chen et al., 2017). PCB 11 has recently emerged as a global pollutant and is currently produced as an unintentional byproduct of paint pigment synthesis (Choi et al., 2008; Du et al., 2008; Hu et al., 2008; Basu et al., 2009; Du et al., 2009; Hu and Hornbuckle, 2010; Heo et al., 2014); thus, we included PCB 11 in the analysis of serum samples collected in this study.

This study was performed to 1) evaluate the presence of PCBs and PBDEs in bovine, caprine and ovine sera and to 2) assess differences in pollutant profiles based on species or cattle production class (beef or dairy).

#### 2. Materials and methods

#### 2.1. Materials

All organic solvents used were of HPLC grade and obtained from Fisher Scientific (Fair Lawn, NJ, USA). The PCB standards (PCB-11, 28, 52, 77, 84, 91, 95, 101, 118, 131, 132, 135, 136, 138, 149, 153, 174, 175, 176, 180, 196) and PBDE standards (BDE-17, 28, 47, 49, 52, 66, 85, 95, 99, 100, 136, 153, 154, 183) were purchased from AccuStandard Inc. (New

Haven, CT, USA). The <sup>13</sup>C12 labeled 2,2′,3′,4,5-pentachlorobiphenyl (<sup>13</sup>C12-PCB-97) and <sup>13</sup>C12 labeled 2,3′,4,4′,5-pentabromodiphenyl ether (<sup>13</sup>C12-BDE-118) were purchased from Cambridge Isotope Laboratories (Tewksbury, MA, USA). Control human serum was purchased from Golden West Biologicals (Temecula, CA, USA). Mirex was purchased from Sigma Aldrich (St. Louis, MO, USA). Solutions were diluted with isooctane to appropriate concentrations.

#### 2.2. Sample collection

Samples were collected from August 2012 to September 2015 from locations depicted in Fig. 1. Counties were grouped into three different areas to identify any regional differences: Northern, Bay Area and Central California. All ovine and caprine serum samples were collected during appointments at the Veterinary Medical Teaching Hospital (VMTH), School of Veterinary Medicine, University of California Davis. Bovine samples were from cattle presented to the VMTH for appointments, and to the California Animal Health and Food Safety Laboratory, University of California Davis for diagnostic work-up. Additional bovine serum samples were collected in the Bay Area region under the supervision of a licensed large animal veterinarian. All samples were stored at -80 °C prior to analysis. In total, 172 serum samples were included in this study (Bovine n = 145, Caprine n = 17, Ovine n = 10). Information on location and production class was also noted when available. Location was designated by county and the various counties sampled were grouped into three regions of California based on geography and anthropogenic activities within the region. Only bovine serum samples were used in regional comparisons (Northern n = 51, Bay Area n = 46, or Central n = 53). Caprine and ovine samples were collected in the northern and bay area regions and were thus excluded from regional analyses since there was uneven regional distribution. The Bay Area was defined by any county touching the San Francisco bay and counties north of the bay were considered to be in the Northern region. All other counties were grouped into the Central region (See Table 1).

#### 2.3. Sample processing and analysis

The serum sample extraction protocol was adapted from our previously published method (Lin et al., 2013). Samples were thawed and briefly vortexed. Aliquots of 0.5 mL of serum were transferred into a disposable glass tube. Serum samples were spiked with 10 µL of an isooctane solution containing 100 ng/mL of <sup>13</sup>C12-PCB-97 and <sup>13</sup>C12-BDE-118. 0.5 mL of formic acid was added and then samples were vortexed for 30 s. Water Oasis HLB SPE cartridges (Milford, MA, USA) were gravimetrically conditioned with 3 mL of methanol and then 3 mL of ultrapure water containing 1% formic acid. Samples were added to the SPE cartridges and the disposable glass tubes were rinsed with 1 mL of ultrapure water with 1% formic acid. This wash was also added to the SPE cartridge and samples filtered gravimetrically. The cartridges were then dried under vacuum (~5 mm Hg) for 300 s. Waters disposable Sep-Pak® Light Silica cartridges (Milford, MA, USA) were placed underneath the SPE cartridges. 3 separate aliquots of 3 mL dichloromethane were added to the SPE cartridges while placed under vacuum (~10 mm Hg) to extract the pollutants of interest into disposable glass tubes containing 100 µL of an isooctane solution containing 100 ng/mL Mirex. The dichloromethane was then evaporated to 1.5 mL under  $N_2$  in a warm water bath (~45 °C). Samples were then vortexed for 30 s and evaporated to dryness under  $N_2$  in a

warm water bath (~45 °C). The residue was reconstituted with 100  $\mu$ L isooctane and vortexed for 30 s. Samples were then placed into an auto-sampler vial for GC/EI-MS/MS analysis.

Seven-point calibration curves at PBDE and PCB concentrations of 0.04, 0.1, 0.2, 0.8, 2, 4, and 10, ng/mL were prepared by adding PBDE and PCB analytical standards to 0.5 mL of control human serum. Calibration samples were processed following the same extraction method as samples.

All samples were analyzed following a previously published method (Lin et al., 2013) using a Bruker Scion TQ triple quadruple mass spectrometer (Bruker, Fremont, CA, USA) equipped with a Bruker 451 GC and CP 8400 auto-sampler and series split/splitless injector set at 280 °C. GC separation was performed by a 15 m BR-5MS, 0.25 mm i.d. column with 0.25 μm film thickness (Bruker, Fremont, CA, USA). The GC oven temperature was as follows: 1) started at 90 °C and held for 1 min; 2) increased to 220 °C at a rate of 50 °C/min and held for 1 min; 3) increased to 260 °C at a rate of 5°C/min and held for 1 min; 4) increased to 300 °C at a rate of 50 °C/min and held for 2 min. The flow rate of the carrier gas, helium, was set at 1.8 mL/min. Source temperature was set at 280 °C and transfer line temperature at 300 °C. PBDE/PCB concentrations were determined in multiple reactions monitoring (MRM) mode. MS/MS was operated in EI positive mode at 70 eV. An aliquot of 2 μL sample was injected by pulsed splitless method (split ratio of 50:1, 50 psi, 0.2 min).

#### 2.4. Quality control

A procedural blank was run in parallel with every batch of samples using isooctane. No PCB or PBDE congeners were detected in the blanks. The accuracy of the method was assessed using three quality control (QC) samples of human control serum fortified with all PBDEs and PCBs at concentrations of 0.1, 0.8 and 4 ng/mL. QC samples were prepared following the same extraction method as described for samples and analyzed in parallel with each group of samples. For each batch of samples processed and analyzed, the determined concentration of each PBDE and PCB congener in the QC samples, as quantified by the standard curves, was required to fall within  $\pm 30\%$  of the known concentration of the individual congener for the data to be included in the final analysis.

#### 2.5. Data and statistical analysis

The GC-MS/MS data was processed by Bruker Mass Spectrometry Working Station version 8.2 (Bruker, Fremont, CA, USA). All analytes were quantified using the 7-point calibration curve. The peak areas were used for quantification following an internal algorithm. The limit of detection (LOD) and limit of quantification (LOQ) were defined based on signal-to-noise (S/N) ratio exceeding three and ten, respectively. Values found below the LOD were reported as "non-detected" (ND) and, for statistical purposes, were assigned a value of half the LOD. The LODs for PBDEs ranged from <0.00001 ng/mL for BDE 154 to <0.094 ng/mL for BDE 66. For PCBs, the LODs ranged from <0.001 ng/mL for PCB 176 to <0.094 ng/mL for PCB 136.

Mann-Whitney Rank Sum Test was conducted using GraphPad Prism v6.07 (San Diego, CA) to compare differences between beef and dairy bovine samples with values of p < 0.05

considered significant. Comparisons between three or more groups used a nonparametric Kruskal Wallis test using an alpha set using the Holm-Bonferroni method using Stata IC/13 (StataCorp LP, College Station, TX).

#### 3. Results and discussion

# 3.1. Caprine and ovine sera contain higher concentrations of PCBs and PBDEs than bovine sera

The sum of the concentrations of PCBs and PBDEs in bovine, ovine, and caprine are depicted in Fig. 2A and B and summarized in the Supplementary Material. Location and production class was not available for every bovine sample, precluding us from using these parameters when evaluating concentrations between species. Total PCBs are significantly greater in caprine and ovine sera than bovine sera with a mean of 9.26 and 9.13 ng/mL compared to 3.98 ng/mL, respectively (Fig. 2A). Total PBDEs are also significantly greater in caprine and ovine sera than bovine sera with a mean of 2.82 and 2.39 ng/mL compared to 0.91 ng/mL, respectively (Fig. 2B).

Individual congener concentrations for each species can be found in Fig. 3 and Supplementary Material. The finding of differing concentrations between species is not surprising because of known differences in biotransformation enzymes and activities between goats, sheep, and cattle (Poppi et al., 1981; Huston et al., 1986; Chen et al., 1990; Szotakova et al., 2004; Girolami et al., 2016). While PCBs and PBDEs are not highly susceptible to metabolic processes, they are subject to cytochrome P450-mediated hydroxylation, primarily performed by the CYP1A and CYP2B family in mammals (Grimm et al., 2015). Cows have higher amounts of CYP1A protein and CYP1A activity compared to sheep and goats, and also have detectable CYP1B mRNA expression and CYP2B activity while goat and sheep lack any CYP2B activity (Szotakova et al., 2004; Girolami et al., 2016). Another species difference in biotransformation mediators is the increased expression of the aryl hydrocarbon receptor (AhR) and a key protein involved in mediating AhR effects, the AhR nuclear translocator (ANTR), in cows compared to sheep (Girolami et al., 2016). The AhR can be activated by coplanar pollutants, such as DL-PCBs, and this activation increases the expression of metabolic enzymes, such as P450s (Denison et al., 2011). Therefore, it is likely that cattle are able to metabolize these pollutants at a higher rate than goats or sheep. Once PCBs are hydroxylated they undergo phase 2 metabolism, attaching even more polar groups leading to their excretion (Grimm et al., 2015). Different digestive processes, such as rumen retention time or microbial communities in the rumen amongst these species can also influence bioavailability of xenobiotics, thereby altering uptake (Agarwal et al., 2015; Ferreira et al., 2016). Other factors, such as body composition, breeding and grazing habits may also contribute to the observed differences. Cattle have a higher mature body fat fraction compared to sheep or goats (Johnson et al., 2012; Maeno et al., 2013). This increase in adipose tissue can serve as a larger reservoir for highly lipophilic POPs allowing them to sequester into fat and removing them from circulation. An additional important physiological consideration to consider in the comparison between bovine, ovine and caprine PCB and PBDE levels is the lactation status of the animals. While about 42% of the analyzed serum samples from cattle came from dairy, none of the ovine and caprine

samples were collected from lactating animals. It is well documented that PCBs and PBDEs partition into bovine milk (Focant et al., 2003; O'Donovan et al., 2011; Lake et al., 2013) presenting a path of elimination of these pollutants.

All livestock, bovine, ovine and caprine, can be exposed to contaminants from air-soil deposition from industrial emissions, and run-off from wastewater treatment plants. In addition, husbandry practices can expose animals to painted fences, barns and hutches, and corral pipes. Husbandry techniques also vary between ruminant species with cattle generally being most closely monitored while small ruminants are allowed more free exploration. Livestock markers are commonly used to temporarily mark livestock after certain procedures including husbandry practices; pigments in these markers including yellow and green may present a potential source for exposure to non-legacy PCBs, such as PCB 11 (Hu and Hornbuckle, 2010). Yellow and green dyes contain the greatest amount of inadvertent PCB contamination providing a potential exposure source for animals through licking and ingesting paint pigments used in markers or paint.

#### 3.2. Dairy cows contain higher serum concentrations of PCBs and PBDEs than beef cattle

The sum of the concentrations of PCBs and PBDEs in beef and dairy bovine samples are depicted in Fig. 2C and D and summarized in the Supplementary Material. Total PCBs are significantly greater in sera from dairy cows compared to beef with a mean of 5.92 ng/mL versus 2.70 ng/mL (Fig. 2C). Total PBDEs are also significantly greater in sera from dairy cows versus beef with a mean of 1.33 ng/mL versus 0.70 ng/mL (Fig. 2D). Individual congener concentrations for each production class can be found in Fig. 4 and in Supplementary Material. The significant difference in body burden of these pollutants and the directionality of the observed difference are surprising. We expected the difference to be in the opposite direction with sera of dairy cows containing lower concentrations of PCBs and PBDEs due to the fact that lactation allows for enhanced elimination in milk.

However, diet is likely a major contributor to the observed differences in PCB and PBDE serum concentrations. One study conducted in Germany strongly supports this hypothesis by finding that 99% of a cow's PCB burden came from its diet with less than 1% coming from the air and water (McLachlan, 1993). U.S. studies confirmed contamination of animal feed with PCBs and other POPs (Sapkota et al., 2007). Dairy cows are usually kept in controlled environments and fed total mixed rations to achieve optimum performance; those rations contain known commodity ingredients which are regularly assessed for quality and nutritional content. The total mixed ration typically contains corn and or corn silage, a feed component documented to contain PCBs and PBDEs (Wang et al., 2011; Brambilla et al., 2015; Hoogenboom et al., 2015; Sun et al., 2016). In contrast, in California, beef cattle tend to be kept on pastures and generally allowed free range, but may be fed hay or other roughage when needed. There are studies reporting PCB levels in fresh grass from the United Kingdom and corn from Italy (Lake et al., 2005; Wang et al., 2011). While the locations for these feed ingredients differed, corn samples have 10-1000 times the PCB content measured in grass. This may be due, in part, to the fact that corn contains a large amount of fatty acids (Gronewald et al., 1982) which would facilitate partitioning of lipophilic compounds into seeds, and/or roots and leaves that are used in grain production

(Sidhu et al., 2000). Various forms of corn that are used in animal feed have lipid percentages ranging from 3.8 to 10% while those among different grasses range from 2.0 to 4.1% (NRC, 2000). Based on these data, rations including corn, such as typical dairy cow rations, may present a greater risk for exposure to PCBs and PBDEs than pastures. However, a recent study in Italy has shown PBDE contamination in pastures and detected its carryover into cow milk (Parolini et al., 2012) showing that pastures are not free of contamination either. While PCB and PBDE concentrations in sera of beef cattle were lower than in dairy cattle, the reported concentrations are still able to make into dietary samples prepared for human consumption. This is illustrated by a study surveying meat samples at a U.S. market detecting PCBs and PBDEs in meat products derived from beef cattle (Huwe and Larsen, 2005) and multiple studies in Europe detecting PCBs and PBDEs in the animal tissues of cattle and goats (Kierkegaard et al., 2007; Ounnas et al., 2010). Unfortunately these past studies mainly looked at dioxin-like POPs, so it is difficult to make comparisons to this study, which focused mainly on the NDL congeners. It would be of great interest to assess how the levels of POPs in the sera of the animal relate to the presence of these pollutants in their meat prepared for human consumption.

With this study we can compare the levels of PCBs and PBDEs present in the sera of the dairy cattle to the results found in a previous study looking at these same congeners in supermarket milk produced in California (Chen et al., 2017) although a limitation is that these dairy cattle were not the producers of the milk analyzed. In the milk study, the most prominent congeners are PCB-138, -118 and -101. These three congeners are present at lower concentrations in bovine sera compared to the other congeners detected in this study, which makes sense and would suggest that there is deposition of specific PCBs from bovine sera to their milk. To further support this hypothesis some of the more prominent congeners in bovine sera PCB-95, -131, -132, -135 and -136 are present at lower concentrations in supermarket milk suggesting an inverse relationship between serum concentration and milk concentration.

# 3.3. There are no regional differences in the sum of PBDEs and PCB congeners in California

The sum of the concentrations of PCBs and PBDEs are illustrated in Fig. 2E and F and summarized in Supplementary Material. There were no statistically significant differences in the  $\Sigma$ PCBs or  $\Sigma$ PBDEs in bovine sera among the three regions of California. There was not a large enough sample size of each type of production cattle in all three regions, and since dairy cattle already have a larger burden of PCBs and PBDEs compared to beef cattle and there were already species differences this made it difficult to draw statistically valid conclusions about this region specific data. Thus we only separated the bovine sera data by samples where location was available (Fig. 5), and although no statistics were run on the individual congeners there appear to be possible regional differences which is not entirely surprising due to the varied anthropogenic activities that occur in each region.

Differences in PCBs may be due to different construction materials used in each region. It is known that the caulking in buildings of many cities (Herrick et al., 2004; Robson et al., 2010; Klosterhaus et al., 2014) contain appreciable amounts of PCBs that can move into the

surrounding environment by either leeching into soil, or volatilizing into the air (Herrick et al., 2007). Studies of buildings from the Bay Area that contain PCBs confirmed a PCB profile similar to Aroclor 1254 (Klosterhaus et al., 2014), a commercial mixture used prior to the ban of PCBs. Unfortunately, there are no data on PCB contamination of older or agricultural buildings in the Northern or Central regions of California. Other regional differences in pollutant profiles may also be due to the proximity of livestock to man-made waste processing facilities (Greichus and Dohman, 1980; Fernandez-Gonzalez et al., 2011).

Due to the high volume of PBDEs used in consumer products and the Bay Area being the most densely populated region of California examined in this study, it is not surprising that the individual PBDE congener concentrations tend to be highest in the Bay Area. Prior to the ban, California mainly used Penta-BDE mixtures, comprised mainly of BDE-47 and –99 (>70%), with smaller contributions from BDE-100, BDE-153, and BDE-154 (La et al., 2006; Stapleton et al., 2012). One epidemiological study assessed sera PBDEs in mothers and their children from the Bay Area of California and found that BDE-47, –99, –100, –153 made up ~90% of serum PBDE content (Eskenazi et al., 2013) consistent with the composition of flame retardants. Our data agrees with these trends with the Bay Area having the highest concentrations of BDE-47, –99, –100 and –153 compared to the Northern and Central regions (Fig. 5).

Another possibility for the regional differences observed in both PCBs and PBDEs levels could be the microbial communities and flora present in the environment. Microbes have been shown to metabolize and dehalogenate POPs, and different microbial communities have higher activities, or preference to degrade specific congeners (Furukawa and Fujihara, 2008; Zanaroli et al., 2015). There are different strains of dehalogenating microbes and each strain prospers under its own specific conditions (Hiraishi, 2008), and it is known that microbial communities vary greatly based on geographic location (Arp et al., 2014), thus these communities may be changing the POP profile in their region. Alfalfa is able to take up PCBs from soil, with enhanced uptake in the presence of nitrogen fixing bacteria (Xu et al., 2010). Maize, or corn has also been shown to uptake both PCBs and PBDEs, with PBDEs primarily staying in the roots, and being more susceptible to metabolic transformations compared to PCBs (Wang et al., 2011). Plant genera and species have different abilities for uptake and metabolism of PCBs that can vary with congeners, as shown in maize, wheat and rice (Sun et al., 2016). Important is also the relationship between plants and the microbial community in the environment as illustrated by the influence of microbial activity and corn on PCB degradation rates (Federici et al., 2012). Collectively, these studies illustrate the effect that varying microbial environments in the environment can have on PCB disposition. In addition, dietary rations change with region based on available commodities.

#### 3.4. PCB 11, a non-legacy PCB, is present in every serum sample collected in California

PCB 11 has recently emerged as a ubiquitous pollutant in multiple regions of the world (Choi et al., 2008; Hu et al., 2008; Du et al., 2009; Heo et al., 2014), yet no study to date has reported serum PCB 11 levels in livestock. PCB 11 was present at quantifiable levels in each serum sample collected for this study. This is surprising because PCB 11 was never a part of

any industrial PCB mixtures, and was not present in the caulking of buildings in the Bay Area (Klosterhaus et al., 2014). It is known that PCB 11 is a byproduct of modern pigment manufacturing processes and detected in consumer goods such as newspapers, plastic bags, magazines and napkins. It is hypothesized that PCB 11 enters the environment by waste release from paint production facilities, or by leaching into the water and soil from consumer products (Hu and Hornbuckle, 2010; Guo et al., 2014). PCB 11 is present at highest concentrations in bright yellow and green dyes, which are both colors used to mark livestock after various procedures. Once marked these dyes can be readily licked or ingested by livestock. PCB 11 can also volatilize and is recognized as a ubiquitous airborne pollutant in multiple regions around the world (Choi et al., 2008; Hu et al., 2008; Heo et al., 2014). Therefore, California livestock are likely being exposed to PCB 11 through inhalation and diet, but it is unknown which route of exposure represents the major contributor to the burdens of PCB 11 we have detected. The fact that our study did not detect geographic differences with this congener suggests that PCB 11 is a ubiquitous pollutant across California without a specific point source. This is of great concern because although PCB 11 is clearly a ubiquitous pollutant present in all forms of environmental media there is currently very little data regarding the potential toxicity of this congener.

While this study provided novel data on the content of PCBs and PBDEs in livestock sera from California there are some limitations. When observing species differences one factor we were unable to control for is obtaining caprine and ovine samples from the same locations as the bovine samples. Assessment of contamination of various feed ingredients was not part of this study; thus we were unable to control for dietary differences. Our analytical method did not include analysis for PCB 209 or BDE-209. In future studies, these congeners should be included because PCB 209 is a prominent paint pigment contaminant (Hu and Hornbuckle, 2010) and BDE-209 occurs at high levels due to firefighting activities (Shaw et al., 2013; Shen et al., 2015).

#### 4. Conclusions

This study is the first to detect PCBs and PBDEs in serum from livestock in the U.S. and also the first study to separate out bovine samples by production class, either beef or dairy. This study is also the first to report the presence of PCB 11 in sera of all livestock evaluated. Sera of goats and sheep have higher concentration of these pollutants than cattle. Dairy cattle have higher serum PCB and PBDE concentrations than beef cattle. This detection of PCBs and PBDEs in sera of livestock in California, along with their detection in commercial milk from California is a point of concern since PCBs and PBDEs are both well-recognized as developmental neurotoxicants. The results of this study bring attention to the lack of knowledge regarding the exposure to these pollutants in livestock and the unknown associated risk factor to humans through the consumption of animal-derived product.

### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

#### **Acknowledgments**

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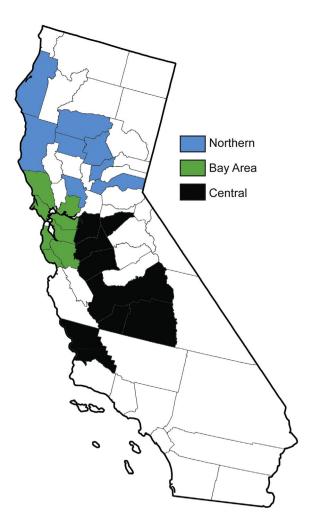
### Appendix A. Supplementary data

Supplementary data related to this article can be found at http://dx.doi.org/10.1016/j.chemosphere.2017.04.059.

#### **HIGHLIGHTS**

• Caprine and ovine sera contain higher levels of PCBs and PBDEs than sera of bovine.

- Dairy cows have higher levels of PCBs and PBDEs than cows used in beef production.
- A non-legacy PCB, PCB 11, was detected at quantifiable levels in all livestock sera.
- There are no significant regional differences in the sum concentration of PCBs and PBDEs in bovine sera in California.



**Fig. 1.**Map of California depicting the counties where serum samples were collected and how those counties were grouped into Northern (blue), Bay Area (green), and Central (black) regions. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

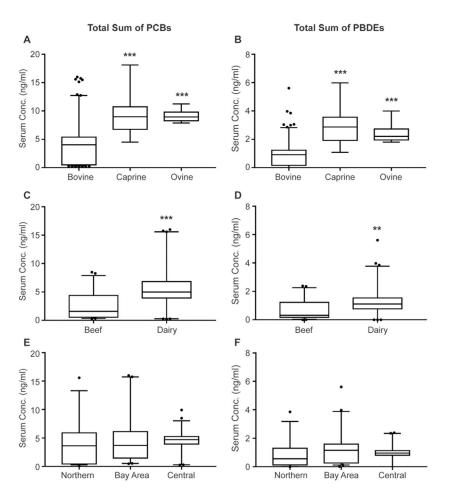


Fig. 2. Species, production, and regional differences in the sum of total PCBs and total PBDEs in serum samples collected across California. Caprine and ovine serum have a greater amount of PCBs (A) and PBDEs (B) compared to bovine serum (n = 145 bovine, n = 17 caprine, n = 10 ovine). Dairy producing cattle have greater amounts of PCBs (C) and PBDEs (D) compared to beef cattle (n = 44 beef, n = 61 dairy). There are no regional differences in the total PCBs (E) and total PBDEs (F) (n = 36 northern, n = 41 bay area, n = 46 central). Plots are showing the  $5^{th}$ – $95^{th}$  percentile. \*Significant differences with p values set at p < 0.05, \*\*p < 0.01, and \*\*\*p < 0.001.

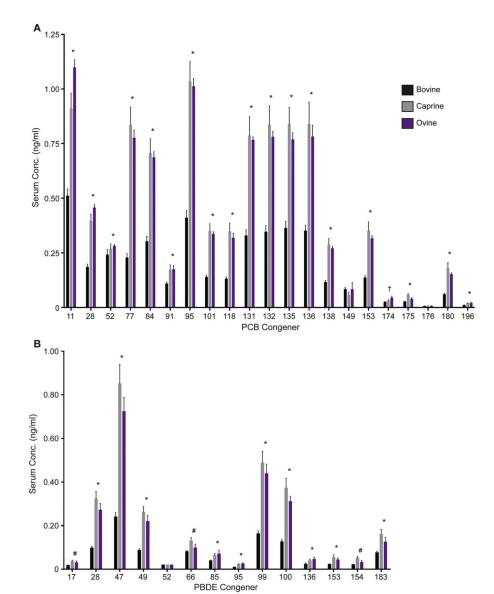


Fig. 3. Species differences in individual PCB (A) and PBDE (B) congeners. Data are presented as mean  $\pm$  standard error of the mean. \* Caprine and Ovine > Bovine, † Ovine > Bovine, # Caprine > Bovine. Significance was set at p < 0.05.

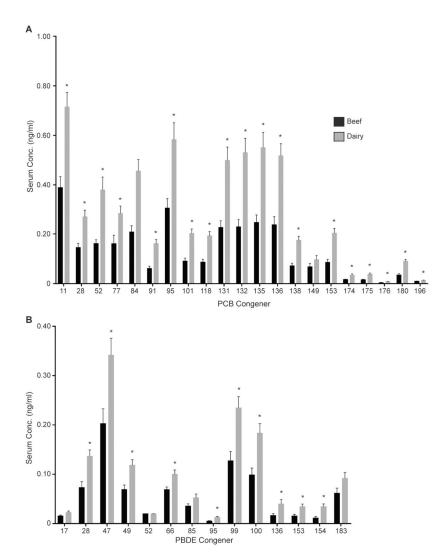
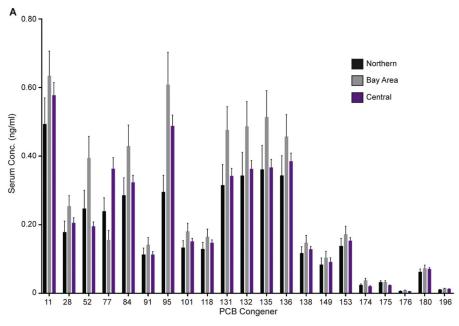


Fig. 4. Production class differences in individual PCB (A) and PBDE (B) congeners. Data are presented as mean  $\pm$  standard error of the mean. \*Significant differences with p values set at p < 0.05.



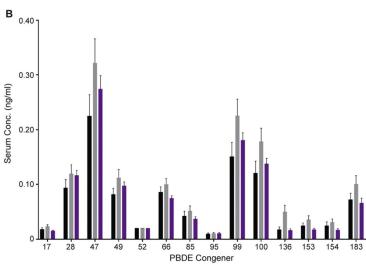


Fig. 5. Regional representation of individual PCB (A) and PBDE (B) congeners in bovine samples including both beef and dairy. Data are presented as mean  $\pm$  standard error of the mean.

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**Table 1** Description of serum samples analyzed in this study.

	Production Class		
	Beef	Dairy	Information Unavailable
Bovine	44 (30%)	61 (42%)	40 (28%)
Caprine	N/A	N/A	17 (100%)
Ovine	N/A	N/A	10 (100%)