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Longitudinal Analysis Reveals Early-Pregnancy Associations between Perfluoroalkyl Sulfonates and Thyroid Hormone Status in a Canadian Prospective Birth Cohort

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Abstract

Serum perfluoroalkyl acids (PFAAs) have been linked to disruption of maternal thyroid hormone homeostasis, but results have varied between studies which we hypothesized was due to timing of the thyroid hormone measurements, variability in PFAA isomer patterns, or presence of other stressors.

In a longitudinal study design, we investigated the time-dependency of associations between PFAA isomers and thyroid hormones during pregnancy and post-partum while considering thyroid peroxidase antibody (TPOAb) status and mercury (Hg) co-exposure. In participants of a prospective Canadian birth cohort (n=494), free thyroxine (FT4), free triiodothyronine (FT3), thyroid stimulating hormone (TSH) and TPOAb were quantified in maternal plasma collected in each trimester and 3-months postpartum, and 25 PFAAs (15 linear and 10 branched) and Hg were quantified in samples collected during the second trimester.

Perfluorohexane sulfonate (PFHxS) and total branched isomers of perfluorooctane sulfonate (PFOS) were positively associated with TSH in mixed-effect models, with strongest associations early in gestation. Throughout pregnancy and post-partum, PFHxS was inversely associated with FT4, consistent with elevated TSH, while Hg was inversely associated with FT3. In TPOAb-

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positive women, negative associations were found between PFAA and FT4, and 1*m*-PFOS and TSH, supporting previous studies that thyroid disorder increases susceptibility to PFAA-mediated hormone dysregulation. Hg did not confound associations but was a significant interaction term, revealing further positive associations between PFOS isomers (3*m*+4*m*-PFOS) and TSH.

Higher perfluoroalkyl sulfonate exposures were associated with higher TSH and/or lower FT4, strongly suggestive that PFHxS and branched PFOS isomers are risk factors for subclinical maternal hypothyroidism. Isomer-specific analysis is important in future studies, as crude measures of ‘total-PFOS’ masked the associations of branched isomers. A concerning result was for PFHxS which had consistent negative associations with FT4 at all time points and a positive association with TSH in early pregnancy when fetal development is most sensitive to disruption.

Keywords

Perfluoroalkyl acids; Perfluoroalkyl sulfonates; Perfluoroalkyl carboxylates; Thyroid hormones; Pregnancy; Longitudinal study design

1. Introduction

Thyroid hormones are important in critical periods of neurodevelopment, including neurogenesis, neuronal migration, proliferation, and myelination (Howdeshell, 2002; Préau et al., 2015). Maintenance of an adequate maternal concentration of thyroid hormones is, therefore, essential for healthy fetal and postnatal neurodevelopment (Williams, 2008). In humans, fetal production of thyroxine (T4) and triiodothyronine (T3) is not established until late in the first trimester (Burrow et al., 1994), and until this point the fetus relies on the maternal supply (Fitzpatrick and Russell, 2010). Thus, early stages of pregnancy exert stress on the hypothalamic–pituitary–thyroid axis (Glinioer, 1999). Alterations of maternal thyroid hormone status are linked to adverse birth outcomes and child development (Negro and Stagnaro-Green, 2014). Maternal hypothyroidism, defined by elevated thyroid stimulating hormone (TSH) with free T4 (FT4) in the reference range, has been associated with spontaneous abortion, preterm birth, placental abruption, low birth weight (Abalovich et al., 2002; Casey et al., 2005; Leung et al., 1993), and lower scores on neuropsychological tests in children (Haddow et al., 1999). Maternal hypothyroxinemia, defined as FT4 in the lowest 10th percentile without a compensatory increase in TSH, has been associated with lower psychomotor development and delayed mental and motor function in infants (Pop et al., 2003, 1999; Smit et al., 2000).

Perfluoroalkyl acids (PFAAs) are among the most prominent organic contaminants in human blood, with perfluorohexane sulfonate (PFHxS), perfluorooctanoate (PFOA) and perfluorooctane sulfonate (PFOS) present at highest concentrations in Canadian (Haines et al., 2017) and American (Olsen et al., 2017) populations. Historic and ongoing production of these compounds, combined with their environmental persistence and bioaccumulation potential has led to their global distribution and accumulation in people and wildlife (Reiner and Place, 2015). Dietary intake is a major pathway of PFAA exposure (Fromme et al., 2007; Haug et al., 2011; Rylander et al., 2010), particularly where fish and seafood are major dietary items (Berger et al., 2009; Haug et al., 2010), but PFAAs or related precursors

may also be present in carpeting, textiles, indoor air and household dust (De Silva et al., 2012; Kato et al., 2009; Kubwabo et al., 2005; Makey et al., 2017; Shoeib et al., 2011).

In rodent developmental toxicology studies, PFOS generally elicits effects on thyroid hormone metabolism that are consistent with hypothyroxinemia (Lau et al., 2003; Luebker et al., 2005; Thibodeaux et al., 2003; Yu and Liu, 2009). In monkeys exposed to PFOS (Seacat et al., 2002) or PFOA (i.e., ammonium perfluorooctanoate) (Butenhoff et al., 2002), subtle alterations in T3 and T4 homeostasis were induced. There have been fewer studies of PFHxS, but one early rat study reported no developmental, nor reproductive toxicity, even at high doses (10 mg PFHxS/kg-day) (Butenhoff et al., 2009). More recent studies have reported that PFHxS-exposed pregnant rats and their offspring had reduced serum T4 measured post-pregnancy (Ramhøj et al., 2018), while in mice, dams had increased liver weight and slight reductions in mean live litter sizes, but no development effects were noted in pups (Chang et al., 2018).

Epidemiological studies of maternal thyroid disruption by PFAAs have been conducted in Asia, Europe and North America, but with differences in experimental design, including differences in measurement timing for thyroid hormones or PFAAs (Berg et al., 2015; Chan et al., 2011; de Cock et al., 2014; Kim et al., 2011; Lopez-Espinosa et al., 2012b; Wang et al., 2014, 2013; Webster et al., 2014). Positive associations are often reported between certain PFAAs and TSH (Berg et al., 2015; Wang et al., 2014, 2013; Webster et al., 2014), suggesting that PFAA exposure may be a risk factor for maternal hypothyroidism. However, as described in a recent review, associations of PFAAs with TSH were not always significant ($p < 0.05$); moreover, previously reported associations of PFAAs with T3 and T4 between studies have been inconsistent, and future well-designed investigations are suggested to confirm the nature of these relationships (Ballesteros et al., 2017). Due to the dynamic nature of thyroid hormone concentrations throughout normal pregnancy (Glinioer, 1997, 1999), we propose a longitudinal design with repeated measurements of thyroid hormones across the gestational period. A previous study made repeated measurement of thyroid hormones to assess relationship with PFAAs, but this included only one timepoint during pregnancy, with all other measurements after birth (Berg et al., 2015).

Additional stressors to the thyroid system may also contribute to variability between studies. In the US general population, PFAAs were associated with changes in thyroid hormone regulation, but only in a subset of participants considered both iodine deficient and had tested positive for thyroid peroxidase antibody (TPOAb) (Webster et al., 2016); a marker of autoimmune hypothyroidism (i.e., Hashimoto's disease). Similarly, in a Canadian birth cohort, PFAAs (PFNA, PFOA and PFOS) were positively associated with TSH, but only in women with high TPOAb (Webster et al., 2014).

TPOAb has since been incorporated as a covariate into models of PFAAs and other contaminants to evaluate their effects on maternal thyroid hormone status (Berg et al., 2014; Preston et al., 2018), considered important since a high proportion of pregnant women with hypothyroidism (e.g., 31–77 %) have been identified to have elevated TPOAb (Abbassi-Ghanavati et al., 2010; Haddow et al., 1999). In the current study we tested a new hypothesis that co-exposure to mercury (Hg) may confound the effects of PFAAs. Maternal exposure to

Hg has deleterious effects on cognition and motor development of offspring, and has been associated with changes in maternal thyroid hormone status, particularly T3 (Chen et al., 2013; Ursinyova et al., 2012). Moreover, like PFAAs, dietary intake of fish is a major source of Hg exposure (i.e., methylmercury). Previous epidemiological studies have not considered co-exposure of PFAAs with Hg, but for pregnant rats the combined exposure of PFOA and Hg caused non-additive changes in gene expression in brain regions of the offspring (Cheng et al., 2013).

The aim of the current investigation was to examine the longitudinal association between maternal PFAA exposure and thyroid hormone status in the prospective Canadian pregnancy cohort study known as APrON (Alberta Pregnancy Outcomes and Nutrition). With measurements of thyroid hormone status at three timepoints during pregnancy, and once in the post-partum period, while also considering Hg co-exposure and TPOAb status; this is a large (n=494) and highly detailed investigation of risk factors for maternal thyroid disruption. This is also the first such study to use an isomer-specific analysis for the PFAAs, which can have different pharmacokinetics and placental transfer (Beesoon et al., 2011; Benskin et al., 2009; De Silva et al., 2009), and isomer-specific associations have already been noted for birth weight and gestational age (Li et al., 2017).

2. Materials and Methods

2.1 Study Participants and Blood Sample Collection

Recruitment to the APrON longitudinal Canadian pregnancy cohort was between March 2009 and July 2012, including 2140 women from the Edmonton and Calgary metropolitan regions in Alberta, Canada. The study protocols were approved by the University of Calgary Health Research Ethics Board, and the University of Alberta's Human Ethics Research Board. Participants provided written informed consent prior to sample or data collection. A full description of recruitment methods, rationale for APrON and a detailed description of the cohort is published elsewhere (Kaplan et al., 2014; Leung et al., 2016).

For the current study, women were included if residing in Calgary, recruited prior to 18 weeks of gestation, providing a blood sample during second trimester for PFAA analysis, and conceived naturally without the use of fertility hormones or assisted reproductive technique) were considered. Inclusion was further limited to non-smokers, women quitting smoking during pregnancy or quit when finding out they were pregnant (i.e., have a history of smoking), resulting in 494 eligible participants. Data was collected through in-person interviews, administration of a first-visit questionnaire, and follow-ups at each timepoint. Details of the assessments have been published (Kaplan et al., 2014), but included data collection on diet, physical activity, mental health, medical history, and demographics. For this investigation, potential covariates and confounders included maternal age, education, household income, ethnicity, parity, medical conditions, as well as a history of smoking, alcohol, and recreational drug use. A subset of participants (n=25) self-reported to be taking medication (e.g., Levothyroxine) for hypothyroidism were included, and adjusted for in final statistical models. Participants missing demographic data (n=16) were not included in final statistical models, and a flowchart and detailed description of the sample population and selected covariates is in the supplementary information (Fig. A1 and Table A1).

Except for the first trimester, plasma or sera were available for most women in each trimester of pregnancy: Timepoint 1 (< 13 weeks gestation, n=167), Timepoint 2 (14 to 27 weeks gestation, n=487), and Timepoint 3 (27 to 40 weeks gestation, n=465). Additional samples were available from 3 months post-partum, Timepoint 4 (n=479). Although the majority of PFAA analyses were measured in plasma, serum was utilized when plasma was not available (14 % of all samples), as the use of either plasma or sera as comparable matrices representing human exposure to PFAAs has been previously validated (Manzano-salgado et al., 2015). All collection and storage materials were tested for background contamination using HPLC-grade water as a surrogate matrix, and no contamination was detected in any materials. A complete description of blood collection, QA/QC protocols, and analyte recoveries are in the supplementary information (Appendix A).

2.2 Thyroid Hormone Analysis

Free triiodothyronine (FT3), free thyroxine (FT4), thyroid stimulating hormone (TSH), and thyroid peroxidase antibodies (TPOAb) were measured at all time points (Timepoints 1 to 4) in plasma of participants. Chemiluminescent microparticle immunoassay kits were used on the Architect System (Abbott Diagnostics, Santa Clara, CA). Commercially available controls (Abbott Diagnostics) were included with each run (every 20 samples) to verify uniform precision between runs. In addition to commercial kits, an in-house reference sample (fasting plasma) was also included with each new lot number to ensure instrumental accuracy and consistency of calibration curves with a coefficient of variation < 1 % for thyroid hormones. Six sets of duplicates within each run were included as a QC check with an acceptable coefficient of variation of < 10 %.

2.3 Isomer-Specific PFAA Analysis.

Extraction of PFAAs was from maternal plasma collected at Timepoint 2 using a modified method from Glynn et al. (2012). Due to their long half-lives (Olsen et al., 2007), PFAA exposure is highly correlated across trimesters (Fisher et al., 2016), making single observations a robust measurement of PFAA exposure. An aliquot of 0.5 mL of plasma was placed in a 15 mL conical polypropylene centrifuge tube containing 1 ng of 8 isotopically labeled internal standards in methanol (MPFAC-MXA, Wellington Laboratories, Guelph, ON) (listed, Table. A2). Plasma extraction was by protein precipitation with 4 mL of acetonitrile (ACN) and sonication at room temperature for 10 min. The plasma/ACN mixture was centrifuged at 2000 rpm for 5 min in an Eppendorf Sorvall ST-40R tabletop centrifuge (Thermo-Fisher Scientific), the supernatant was transferred to a new 15 mL tube, and the pellet was discarded. The supernatant was then evaporated with nitrogen gas at 40 °C to a volume of 0.3 mL and reconstituted in 50:50 methanol:water to a final volume of 1 mL. This extract underwent dispersive cleanup (Powley et al., 2005) by transferring to a 1.7 mL Eppendorf tube containing 0.025 g of bulk graphitized carbon (Supelclean ENVI-Carb, Sigma Aldrich), that had been acidified with 50 µL of glacial acetic acid and vortexed for 10 s. The sample was centrifuged for 10 min at 10,000 rpm (Sorvall Legend Micro 21R, Thermo Scientific) and the top 0.5 mL was transferred to a glass auto-sampler vial.

A total of 25 PFAA analytes, including 16 perfluoroalkyl carboxylates (11 linear and 5 branched isomers) and 9 perfluoroalkyl sulfonates (4 linear and 5 branched isomers) were

monitored in all maternal plasma samples at Timepoint 2. Analysis was by HPLC-MS/MS with a UFLC-XR Shimadzu HPLC coupled to an API 5000 triple quadrupole mass spectrometer (Applied Biosystems Sciex, Concord, ON) operating in negative ion mode with multiple reaction monitoring. Instrument parameters were modified from a previous method (Benskin et al., 2012). A detailed description of instrument parameters, median concentration, and detection limits for all PFAAs are provided in the supplementary material (Tables A2 and A3). An external solvent-based calibration curve was utilized and each linear and branched PFAA was quantified using the appropriate mass labeled internal standard. A 5 μL sample was injected onto an Ascentis Express F5 PFP analytical column (2.7 μm , 90 \AA , 10 cm \times 2.1 mm, Sigma-Aldrich) equipped with an Ascentis Express F5 PFP guard column (2.7 μm , 5.0 mm \times 2.1 mm) at 40 $^{\circ}\text{C}$. Upstream of the injector, two XTerra C18 columns (5 μm , 30 mm \times 2.6 mm each, Waters) were in place to separate instrumental background PFAAs from PFAAs in the sample injected to the analytical column. A binary gradient elution was used, including (A) 5 mM aqueous formic acid and 5 mM ammonium formate, and (B) 100 % methanol at 0.2 mL/min. The elution gradient was initially 10 % B, 60 % B by 3 min, 88 % B by 14 min, and 100 % B by 14.5 min, held until 15 min and returned to initial conditions by 16 min with a further 5 min equilibration.

2.4 Total Hg Analysis

A 100 μL aliquot of maternal blood cell fraction from Timepoint 2 was diluted with 100 μL of deionized water and diluted with a basic solution containing 25 $\mu\text{g/L}$ of iridium as internal standard, 10 $\mu\text{g/L}$ gold, 0.5 g of EDTA in 1% v/v ammonia hydroxide, 2.5% butanol, and 0.05% v/v Triton X100. The resultant 50-fold dilution was then analyzed for total Hg using inductively coupled plasma mass-spectrometry (Agilent 8800 ICP-MS/MS). Helium was used as collision gas to remove interferences. Two sources of external quality controls were used (SeronormTM and ClinchekTM) at three levels and injected after every 10 samples. The analytical acceptability range was defined as within 20 % of the reference value according to manufacturer guidelines. Duplicates of each sample were run, and for every 10 samples a sample was randomly selected and spiked with the analytes as an additional QC check. Complete details of quality assurance and quality controls, including instrumental limits of detection and quantitation (LOD, LOQ), and Hg recovery are described in the supplementary material (Tables A6, A7, and A8).

2.5 Statistical Analysis

Thyroid hormones were measured over time for the same subject. These observations were not independent, violating the assumption of ordinal regression modeling, thus mixed effect modelling was employed to accommodate the correlation structure within observations of the same subjects. All analyses were performed using R.3.3.2. Log-transformation of both PFAAs and maternal thyroid hormones were considered, but the model fit was best when only the outcomes (thyroid hormones) were log-transformed. Plots of the residuals supported the normality of the residuals and confirmed the goodness of fit (data not shown). The effect of each potential covariate was evaluated in the separated mixed effect model in the presence of the main predictor, either PFAAs or Hg. Covariates with $p < 0.2$ were nominated to enroll in a multiple regression model. The fixed parameters of the multiple mixed models included the main predictor, the covariates from the first step ($p < 0.2$) and a

time variable indicating Timepoints. We also estimated subject-specific trajectories by considering a time variable in the random part of the model. For all mixed effect modeling, three correlation structures (unstructured, autoregressive and compound symmetry) were attempted and the structure with the lowest Akaike information criterion – a method of assessing the quality of a model – was used for final model selection. Multicollinearity between covariates was tested and the goodness of fit for all the models was evaluated exploring the models' residuals. Evaluation in statistical models was restricted to PFAAs that were detected in > 80 % of samples in the overall population. Values below detection limits were replaced with an imputed value of LOD/ 2 (Hornung and Reed, 1990).

For each model of maternal PFAA or Hg exposure with thyroid hormone, in addition to the main effect, the outcome at each timepoint considered the significance of a time interaction ($p < 0.05$), accounting for the effect over each sequential Timepoint (e.g., the change in the effect of each PFAA on modeled thyroid hormone from Timepoint 1 to Timepoint 2, Timepoint 2 to Timepoint 3, etc.). At Timepoint 4, the significance of an additional interaction ($p < 0.05$) was considered to account for the change in effect during pregnancy to post-birth. The main effect, and the outcome at each timepoint were adjusted for all significant covariates for each thyroid hormone. Common significant covariates, including maternal age, ethnicity and history of smoking were identified for all measured thyroid hormones (FT3, FT4, and TSH). In addition to these, a diagnosed thyroid condition was included for FT4, and a history of drug and alcohol use was included for TSH.

3.0 Results

3.1 Population Description

Participants had a mean age of 32, the majority were Caucasian (88 %), had completed post-secondary education (99 %), and most (82 %) had annual household incomes above \$77,000 CDN (Table A1). Most participants were either nulliparous or primiparous (91 %). A proportion (23 %) were self-reported to have a history of smoking, defined as having consumed >100 cigarettes over their lifetime. However, the self-reported proportion of current smokers was 2 % among pregnant participants. Overall, the described population had similar characteristics to a Canadian birth cohort study ($n = 152$) in the metropolitan area of Vancouver, Canada, that previously reported on PFAA-thyroid associations.

3.2 PFAA Concentrations in Maternal Plasma

Certain perfluoroalkyl carboxylates (linear-PFOA, perfluorononanoate (PFNA), and perfluorodecanoate (PFDA)) and perfluoroalkyl sulfonates (PFHxS, linear-PFOS, and most branched PFOS isomers (*iso*, *5m*, *3m*, and *4m*)) were detected in > 99 % of plasma samples. Perfluoroheptanoate (PFHpA), perfluoroundecanoate (PFUnA), and perfluorododecanoate (PFDoA) were detected less frequently, in 66, 89, and 55 % of samples, respectively. Total PFOS had the highest median concentration at 4.77 ng/mL, followed in descending order by PFOA (2.11 ng/mL), PFHxS (1.03 ng/mL), PFNA (0.69 ng/mL), PFDA (0.25 ng/mL), PFHpA (0.08 ng/mL), PFUnA (0.06 ng/mL), and PFDoA (0.06 ng/mL) (Table 1). Among the major PFAAs, total PFOS, total PFOA, and PFHxS concentrations were comparable to recent measurements in Canadians (Fisher et al., 2016; Health Canada, 2013).

For isomers of PFOS, the median concentration of linear PFOS was 2.49 ng/mL (69 % of total PFOS) and total branched PFOS was 1.08 ng/mL (31 % of total PFOS) in maternal plasma (Table 1). Among PFOS branched isomers, the highest median concentration was for *iso*-PFOS (0.42 ng/mL, 12.7 % of total PFOS), followed by 5*m*-PFOS (0.33 ng/mL, 9.8 %), 3*m*+4*m*-PFOS (0.23 ng/mL, 6.1 %), and 1*m*-PFOS (0.07 ng/mL, 2.1 %) (Table 1). Although a broad range of PFOA isomers were also monitored in maternal plasma, only linear-PFOA (2.11 ng/mL, 96.8 % of total PFOA) and *iso*-PFOA (0.07 ng/mL, 3.2 %) were detectable. However, due to the low frequency of detection of *iso*-PFOA, it was excluded from models of associations with thyroid hormones and further mention of PFOA refers only to linear PFOA.

Spearman correlations (listed in Table A9 of the supplementary information), indicated moderate correlations between numerous carboxylate PFAAs and sulfonate PFAAs within each family, but weak correlations between carboxylate and sulfonate groups; isomers of PFOS were moderately to strongly correlated (Table A9).

3.3 Total Hg Concentrations in Maternal Red Blood Cells

Total Hg was frequently detectable (98 %) in maternal blood cell fraction (Table 1). After adjusting for relative volumes of plasma and cell fraction (i.e., Hg concentration in RBC fraction were divided by 2), concentrations of total Hg in the current study (median of 0.490 µg/L) were comparable to whole blood concentrations in pregnant women from another recent Canadian cohort (0.491 µg/L, MIREC) (Arbuckle et al., 2016). Canadian cohort values were lower than recent cohort studies conducted in Norway (1.21 µg/L, MISA) (Veyhe et al., 2015) and Korea (3.19 µg/L, MOECH) (Jeong et al., 2017). In the current study, Hg was not strongly correlated with any PFAA analytes or their isomers, but significant low to moderate correlations were observed between Hg and PFDA, and Hg and PFUnA (Table A9).

3.4 Thyroid Hormone Concentrations in Maternal Plasma

Concentrations of thyroid hormones (TSH, FT4, and FT3) at all Timepoints are summarized in Table 2. TSH increased slightly mid-gestation (Timepoint 2), whereas decreasing trends for FT4 and FT3 were observed throughout the pregnancy (Timepoint 1 to 3). FT4 subsequently increased at 3-months postpartum (Timepoint 4). An increase in TSH between the first and second trimester, and parallel patterns of decreasing FT4 and FT3 have been noted previously, but levels generally remain in the reference range (Glinoe, 1997).

TPOAb data have been used as a binary variable (i.e., high/normal) in previous investigations of PFAAs and thyroid hormones. Cutoff values used have ranged from low (i.e., 9 mIU/mL) (Chen et al., 2013; Webster et al., 2016, 2014), to moderate (34 to 50 mIU/mL) (Abbassi-Ghanavati et al., 2010; Berg et al., 2015) to high (90 mIU/mL) (Pop et al., 1995). Here, we used 9 mIU/mL, to be consistent with previous PFAA studies (Chen et al., 2013; Webster et al., 2016, 2014), and because this value has been considered a threshold for eliciting disturbances in immunological function of thyroid tissue (National Center for Health Statistics, 2009). The range of women with elevated TPOAb in the current study was 5 to 15 % (Table 2), which is comparable but slightly wider than the reference range for

euthyroid women (i.e., 6 – 12 %) based on previous studies of thyroid hormone status and TPOAb (Abbassi-Ghanavati et al., 2010; Loh et al., 2016; Meena et al., 2016; Springer et al., 2017). Including, and adjusting for hypothyroidism in models of the current population may have attributed to increased proportion of women with elevated TPOAb. In women categorized as high TPOAb, TSH was significantly higher early in pregnancy (Timepoint 1 and Timepoint 2), and FT4 was significantly higher postpartum (Timepoint 4) compared to normal TPOAb women ($p < 0.05$ using student t-test).

3.5 Associations between PFAA or Hg Exposure and Maternal Thyroid Hormones

Significant associations ($p < 0.05$) between PFAAs and thyroid hormones (TSH, FT4, FT3) are summarized in Table 3 (see Table A5 for all results). TSH was positively associated with several perfluoroalkyl sulfonates (PFHxS, Br-PFOS, and 5*m*-PFOS) during pregnancy, but not with any perfluoroalkyl carboxylates, nor Hg (Table 3). The strength of the significant associations was dependent on time, with all three associations significant in early pregnancy (Timepoint 1), only PFHxS was significant in the second trimester (Timepoint 2), and none were significant late in pregnancy (Timepoint 3). The effect size of PFAAs with TSH were larger in comparison to PFHxS with FT4, suggesting that PFAAs may elicit a stronger influence on specific maternal thyroid hormones. At 3-months postpartum (Timepoint 4), a significant association with linear-PFOS was revealed, and association with 5*m*-PFOS was significant again, but with a decreased coefficient compared to earlier in gestation (Timepoint 1). For PFHxS, Br-PFOS, and 5*m*-PFOS, the relative trends are similar over time (Fig. 1A–C), with strong dose-dependent positive associations at Timepoint 1 (Fig. 1A–C) which weaken by Timepoint 2, disappear by Timepoint 3, and reappear post-partum. It is concerning that the most significant and strongest associations occurred in the earliest stage of pregnancy when the fetus may be most susceptible to subclinical maternal hypothyroidism.

For FT4, the only significant association was a negative association with PFHxS at all Timepoints (Table 3). This can be visualized over time for increasing inter-quartile concentrations of PFHxS (Fig. 2A), demonstrating the consistent dose-dependent association despite the expected decrease in FT4 during the first and second trimester, as well as the subsequent postpartum rise (Table 2). These significant associations for PFHxS and FT4 are consistent with the positive associations between PFHxS and TSH at Timepoints 1 and 2, because the function of elevated TSH is to stimulate the thyroid gland to produce more T4 in response to low circulating T4 and T3.

For FT3, the only significant associations were for Hg and 1*m*-PFOS (Table 3). In the case of Hg, a significant negative association with FT3 was observed consistently across all Timepoints. This can be observed visually as a dose-response decline in FT3 with increasing median quartile concentration of Hg (Fig. 2B). Increasing 1*m*-PFOS concentrations were positively associated with FT3 at 3-months postpartum (Timepoint 4), but not at other Timepoints.

The significant associations between TSH and the branched isomers (Br-PFOS, 5*m*-PFOS), or between FT3 and 1*m*-PFOS, would likely have been missed in this study without the isomer-specific analysis. This is because the branched isomers are relatively minor, and

the major isomer (Linear-PFOS, 69% of total PFOS) was not associated with TSH during pregnancy (Table 3), nor with FT4 or FT3 at any Timepoint. In fact, by summing all linear and branched isomers we confirmed that total PFOS was not associated with TSH, FT4, or FT3 (Table A5). A recent review pointed out that previous studies of PFOS and thyroid disruption have been inconsistent (Ballesteros et al., 2017), and we propose that 'total-PFOS' analytical methods have contributed to this.

Including an interaction term for Hg co-exposure did not influence any of the above noted associations for PFAAs. However, one significant new association was revealed, a positive association between 3*m*+4*m*-PFOS and TSH ($p < 0.05$) (Table 3) when including the interaction of Hg co-exposure. This indicates that increasing concentrations of 3*m*+4*m*-PFOS are associated with TSH, but that increasing concentrations of Hg interact to weaken the association ($\beta = -0.30$, $p = 0.047$). This new relationship agrees with the general association for Br-PFOS and 5*m*-PFOS which were significantly positively associated with TSH in early pregnancy (Timepoint 1) and at 3-months postpartum (Timepoint 4) (Table 3).

3.6 Inclusion of TPOAb status

Including a TPOAb interaction term into the model (Table A5), whereby participants were classified as either normal (< 9 IU/mL) or high (> 9 IU/mL) at each time point, did not significantly alter any of the established main effect associations (Table 3). Nevertheless, new significant associations were revealed after inclusion of TPOAb status (Table 4). PFUnA was positively associated with FT4 in the high TPOAb group only. The association was consistent at all measured timepoints; thus, the result is unlikely spurious because it was consistent throughout pregnancy and at 3-months postpartum. The branched PFOS isomer, 1*m*-PFOS, was also associated with TSH when including TPOAb status, but the result was time-dependent (Table 4). Specifically, in the normal TPOAb group, there was a significant positive association between 1*m*-PFOS and TSH at Timepoints 1 and 2, but not at Timepoints 3 and 4. The positive association of 1*m*-PFOS with TSH is not surprising, as the normal TPOAb group represents 85% of the population, and similar longitudinal trends were observed for the sum of all PFOS branched isomers (Br-PFOS), as well as 5*m*-PFOS. More unexpected was the opposite association in the high TPOAb group, whereby 1*m*-PFOS was inversely associated with TSH (Table 4). This association was not significant early in gestation (Timepoints 1 and 2) but became stronger over subsequent timepoints, with significant negative associations in late pregnancy and at 3-months postpartum (Timepoints 3 and 4). This vulnerable TPOAb subgroup may be more susceptible to the effects of environmental contaminants such as PFAAs, and these results suggest a unique response compared to the larger normal TPOAb population.

4. Discussion

During pregnancy, specific thyroid hormones were influenced by background exposure to PFAA sulfonates (Table 3), as PFHxS, and branched-PFOS isomers were positively associated with TSH ($p < 0.05$) in a dose-dependent manner that were strongest in early pregnancy (1st trimester) and weakened over subsequent trimesters. In addition to associations of TSH, PFHxS was also negatively associated with FT4, and Hg was

negatively associated with FT3. Including additional interactions did not further affect the established associations of PFAAs and thyroid hormones but newly significant associations were revealed when considering effect modification by Hg (e.g., 3m+4m PFOS with TSH, Table 3), and TPOAb (e.g., PFUnA with FT4 and 1*m*-PFOS with TSH, Table 4).

Maternal thyroid hormone homeostasis is under inherent stress during pregnancy. Increasing estrogen in early gestation coincides with increased T4-binding globulin, resulting in increased total T3 and T4 (Glinoe, 1999) which continues throughout pregnancy to meet maternal and fetal demand, even after onset of fetal thyroid function at 16–20 weeks (Moog et al., 2017). Elevated TPOAb, which occurs in 6 – 12 % of pregnant women, is considered a clinical marker of autoimmune thyroiditis (Hashimoto’s disease) and indicates additional stress on the thyroid that increases the risk of adverse birth outcomes (Abbassi-Ghanavati et al., 2010; Li et al., 2010). Thus, pregnant women may be vulnerable to thyroid disruption from the additional stress of environmental contaminants that act as endocrine disruptors, such as PFAAs and Hg. Epidemiological studies of pregnant women are complicated by gradual changes in maternal physiology, and by changing concentration of thyroid hormones throughout the pregnancy (Harada et al., 1979; Mandel et al., 2005; Tulchinsky and Little, 1994; Yazbeck and Sullivan, 2012), also observed here (Table 2). Given the susceptibility of pregnant women to thyroid disruption, while also considering the biological importance of understanding the timing of any disruption, longitudinal studies are particularly well suited to studies of maternal thyroid disruption.

In prospective cohorts, longitudinal approaches have several advantages over other study designs, including identification of the timing, trends, or recurrence of adverse outcomes (Caruana et al., 2015). This study on PFAAs is the first to measure thyroid outcomes at multiple time points during the gestational period, and important time-dependent associations were revealed for the first time (Table 3). Our statistical models using repeat measures accounted for changes in thyroid hormone levels over time, revealing significant relationships between PFHxS and TSH, or PFOS isomers and TSH that were strongest early in pregnancy, gradually disappeared by the third trimester, and in some cases, reappeared post-partum. These trends suggest a window of vulnerability in early pregnancy at a time when the fetus is almost entirely dependent on the maternal supply of thyroid hormones. In other cases, the longitudinal design revealed no trend over time but demonstrated consistent significant associations at each time point. For example, the negative associations between PFHxS and FT4, and between Hg and FT3 (Table 3). Such consistent results provide great internal validity that the associations are not spurious.

In the current study, overall results for PFHxS were particularly strong and compelling. Higher maternal PFHxS exposure was positively associated with TSH (time-dependent) and negatively associated with maternal FT4 (not time-dependent). This is consistent with previous findings of a positive association of PFHxS with TSH (Wang et al., 2014, 2013; Webster et al., 2014) and negative association with FT4I; an index of free T4 in circulation (Preston et al., 2018). Taking this into consideration with the current dataset showing a consistent dose-response between increasing PFHxS concentrations and declining FT4 at all Timepoints, evidence is building for a causal relationship (Bradford-Hill, 1965). Several mechanisms have been proposed to explain the thyroid disrupting effects of PFAAs,

including, increased hepatic clearance and excretion of T4 through glucuronidation (Yu et al., 2009), deiodinase enzyme mediated conversion of T4 to T3 (Yu et al., 2009), and the competitive displacement of T4 from thyroid hormone binding proteins in serum (Jones et al., 2003; Weiss et al., 2009). In fact, compared to other PFAAs, PFHxS has the highest in vitro competitive binding potency to human transthyretin, an important human T4 serum transport protein (Weiss et al., 2009). Our findings are in support of this latter mechanism, as it has been proposed that competitive displacement of T4 by PFAAs leads to a transient increase in FT4, but that the resulting clearance from induction of T4 metabolism ultimately reduces FT4 at steady-state (Kim et al., 2011). Moreover, dose-dependent reductions in plasma FT4 have also been observed in chicken embryos exposed to PFHxS (Cassone et al., 2012). Notwithstanding such in-vitro results, this mechanism required relatively high concentrations of PFAAs, and the validity in-vivo at lower concentrations is questionable (Ren et al., 2016). Nevertheless, observed associations in this study and previous studies are consistent with this mechanism.

According to NHANES data between 1999–2000 (Calafat et al., 2007) and 2007–2008 (Webster et al., 2016), human exposure to PFOS drastically declined (geometric mean declined from 30.4 to 13.5 ng/mL) whereas PFHxS exposure was effectively unchanged (2.1 and 1.9 ng/mL, respectively) over the same time period. More recent monitoring of PFAAs in plasma samples collected from American Red Cross donors in 2015 found further decreased PFOS (4.3 ng/mL), and to a lesser extent, PFHxS (0.9 ng/mL) (Olsen et al., 2017), which was at comparable concentrations to the current study (Table 1). Human PFOS exposure has declined much faster than PFHxS, and PFHxS may soon be the most prominent PFAA in humans if it is not targeted for mitigation.

To our knowledge, this is the first epidemiological study to include an isomer-specific analysis of PFAAs for investigation of maternal thyroid hormone disruption. This is a limitation of previous work because PFOS and PFOA are present in human blood as various linear or branched isomers that are known to have different pharmacokinetics (Benskin et al., 2009; De Silva et al., 2009). Isomer-specific analysis is particularly important for PFOS which was historically manufactured as a mixture of linear (70%) and branched (total 30%) isomers. Populations around the world have wide-ranging proportions of branched PFOS isomers, for example the branched PFOS content in samples from China (52 %) (Zhang et al., 2013), is higher than in Norway (30–47 %) (Haug et al., 2009; Rylander et al., 2010) or Vietnam (17 %) (Rylander et al., 2009). Higher proportions of branched PFOS isomers have been detected in fetal cord blood (24 %, and 36–54 %) compared to paired maternal samples (17 %, and 27–44 %) (Beesoon et al., 2011; Chen et al., 2017) and branched PFOS isomers have higher transplacental transfer efficiencies than linear PFOS (Beesoon and Martin, 2015). Thus, the associations between branched PFOS isomers and maternal thyroid disruption (Table 3) could also be relevant to thyroid disruption in the developing fetus.

Interestingly, the strongest associations with TSH were for PFHxS and branched PFOS isomers, with no corresponding association for linear PFOS (Table 3). We postulate that PFHxS and branched PFOS isomers share optimal physical properties or molecular sizes which allows them to interact most strongly with receptor biomolecules that control thyroid hormone homeostasis. Compared to linear PFOS, PFHxS and branched PFOS molecules are

shorter in length and less hydrophobic, for example eluting earlier than linear PFOS in reversed phase chromatography. PFHxS has already been demonstrated to be more optimal than PFOS for displacing T4 from transthyretin (Weiss et al., 2009), but the same studies have yet to be done for branched PFOS isomers. For all these reasons, future epidemiological and toxicological studies of PFOS should consider isomer-specific analysis and isomer-specific toxicity.

During pregnancy, thyroid hormone concentrations are dynamic but tightly controlled. PFAAs were significantly associated with altered maternal thyroid hormone concentrations (FT4 and TSH), but it is important to note that thyroid hormone levels remained in the reference range. Although reference ranges exist for the general population (e.g., 0.45 to 4.5 mU/L for TSH and 4.5 to 12.5 mg/dL FT4I), there is uncertainty in the value of universal reference ranges of these hormones during pregnancy as concentrations vary by gestational age, number of fetuses, and between populations (Fitzpatrick and Russell, 2010). Furthermore, even subtle alterations in maternal thyroid hormone status may affect fetal development, as children of women with subclinical maternal hypothyroidism (Haddow et al., 1999) or maternal hypothyroxinemia (Pop et al., 2003) had decreased neurodevelopmental indices. Although only subclinical associations in maternal thyroid hormone status during pregnancy were detected here, the public health significance for fetal neurodevelopment can be important due to widespread exposure of the population to PFAAs and presence of cumulative risk factors (e.g., TPOAb).

The “multiple-hit hypothesis” proposed by Webster et al. (2014), states that populations with pre-existing thyroid related conditions may be more susceptible to the effects of thyroid disrupting environmental contaminants, supported by evidence of PFAA-induced thyroid dysregulation in a subset of people with high TPOAb and an iodine deficiency (Webster et al., 2016). Within the small subgroup of women with elevated TPOAb in the current study, PFUnA was negatively associated with FT4, and 1*m*-PFOS was negatively associated with TSH (Table 4). PFOS has previously been reported to have both positive (Webster et al., 2014) and negative (Preston et al., 2018) associations with TSH in high TPOAb status pregnant women. Although the mechanism of interaction between PFAAs, thyroid hormones, and TPOAb is not well understood, these results further support the multiple-hit hypothesis (Webster et al., 2014).

Total Hg analysis was included in the current study to control for a hypothesized confounding variable, but also to consider any relevant mixture effects on thyroid disruption. In the current Canadian cohort, blood Hg was not strongly correlated with serum PFAAs and was not considered a confounder, but Hg should be included in future studies where fish intake is higher. Nevertheless, Hg was negatively associated with free T3, similar to previous associations between Hg and total T3 (FT3 was not analyzed) in pregnant women from Quebec, Canada, (Takser et al., 2005) and Spain (Llop et al., 2015). In Slovakia, Hg in cord blood was inversely associated with total T3 and FT3 in 6-month old infants (Ursinyova et al., 2012). These findings are consistent with the proposed mechanism of action for Hg through enzymatic inhibition of deiodinase activity (enzyme D3, in particular), resulting in reduced T3 production (Mori et al., 2006; Soldin et al., 2008). A possible mixture effect between Hg and 3*m*+4*m*-PFOS was observed in models of TSH, whereby the main effect

of 3*m*-+4*m*-PFOS was positively associated with TSH when including Hg co-exposure, however, the 3*m*-+4*m*-PFOS-Hg interaction term was negative, indicating that at higher Hg concentrations association of these PFOS isomers with Hg became weaker. Thus, it is possible that Hg increases sensitivity to thyroid disruption by PFOS branched isomers but is dependent on the concentration of Hg.

Although this was among the largest and most detailed studies of PFAAs and thyroid disruption, we acknowledge some study limitations. Due to the logistical challenges of recruiting pregnant women early in pregnancy, first trimester (Timepoint 1) samples for thyroid hormone analysis were not always available, resulting in a smaller sample size when compared to the other timepoints. Participants all resided in the metropolitan area of Calgary, Canada, and were from rather narrow demographic (e.g., high socioeconomic status) that may not be representative of the overall population. This may limit the external validity of our findings when considering other global populations, but it does not decrease the internal validity of our findings. Nevertheless, with respect to PFAA exposure, the concentrations of PFAAs in maternal plasma in this study were almost identical to another Canadian birth cohort (Webster et al., 2014), and similar to other cohorts in Korea (Kim et al., 2011), China (Lin et al., 2016), Taiwan (Wang et al., 2014), and Norway (Berg et al., 2014). With respect to participant iodine sufficiency, this was determined using questionnaires, assuming sufficient if taking prenatal supplements that included iodine. In the general US population, thyroid hormone status was particularly susceptible to PFAAs for a subgroup with combined high TPOAb status and an iodine deficiency, (Webster et al., 2016), thus future studies might consider inclusion of urinary iodine. Finally, thyroid hormones were measured by immunoassay. Although this is one of the most commonly employed methods of quantifying thyroid hormones in human samples, radioimmunoassays, such as the one used in this study, are sensitive to serum binding protein concentration, not designed to be used under conditions when protein levels are subject to change (e.g., during pregnancy) (Soldin and Soldin, 2011). Similar problems resulted in reporting bias and underreporting of FT4 values in animal models (Chang et al., 2007), but have been deemed less problematic in human samples (Lopez-Espinosa et al., 2012a). Nonetheless, pregnancy cohort studies might benefit from using LC-MS/MS for its enhanced specificity and precision over immunoassays.

5. Conclusions

The prospective APrON birth cohort study allowed one of the largest and most detailed studies of maternal thyroid hormone disruption by PFAAs. With three sampling times throughout pregnancy and another sampling time after birth, the longitudinal analysis revealed new trends that are of biological significance, and which could explain variability among previous studies. By considering additional thyroid stressors, Hg and TPOAb status, the study also revealed interactions of relevance to the multiple hit hypothesis, and to the toxicology of environmental chemical mixtures in human blood, respectively. By using an isomer-specific method for PFAA analysis, the distinct toxicological behavior and relevance of branched PFOS isomers was revealed, which highlights the importance of isomer-specific PFAA methodologies in future epidemiology and toxicology studies. Building on previous literature, there is now very strong evidence in support of causal relationships between

PFHxS and PFOS exposure and disruption of maternal thyroid homeostasis in early pregnancy. This information should be considered in future international decisions under the Stockholm Convention, where PFHxS has been nominated for inclusion as a Persistent Organic Pollutant (Stockholm Convention, 2018), and PFOS is already included but with many current exemptions under its listing in Annex B.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

ACN	acetonitrile
APrON	Alberta pregnancy outcomes and nutrition
FT3	free triiodothyronine
FT4	free thyroxine
Hg	mercury
HPLC-MS/MS	high performance liquid chromatography mass spectrometry
ICP-MS/MS	inductively coupled plasma mass-spectrometry
LOD	limit of detection
LOQ	limit of quantitation
PFAAs	perfluoroalkyl acids
PFDA	perfluorodecanoate
PFDoA	perfluorododecanoate
PFHpA	perfluoroheptanoate
PFHxS	perfluorohexane sulfonate
PFNA	perfluorononanoate
PFOA	perfluorooctanoate

PFOS	perfluorooctane sulfonate
PFUnA	perfluoroundecanoate
TPOAb	thyroid peroxidase antibody
TSH	thyroid stimulating hormone

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Highlights

- Perfluoroalkyl acids were associated with thyroid hormones during and after pregnancy
- Associations were strongest in early pregnancy, a sensitive stage of neurodevelopment
- Evidence mounting for cause-effect relationship for perfluorohexane sulfonate
- Isomer-specific associations for branched perfluorooctane sulfonate, not linear
- Associations influenced by mercury co-exposure and thyroid peroxidase antibodies

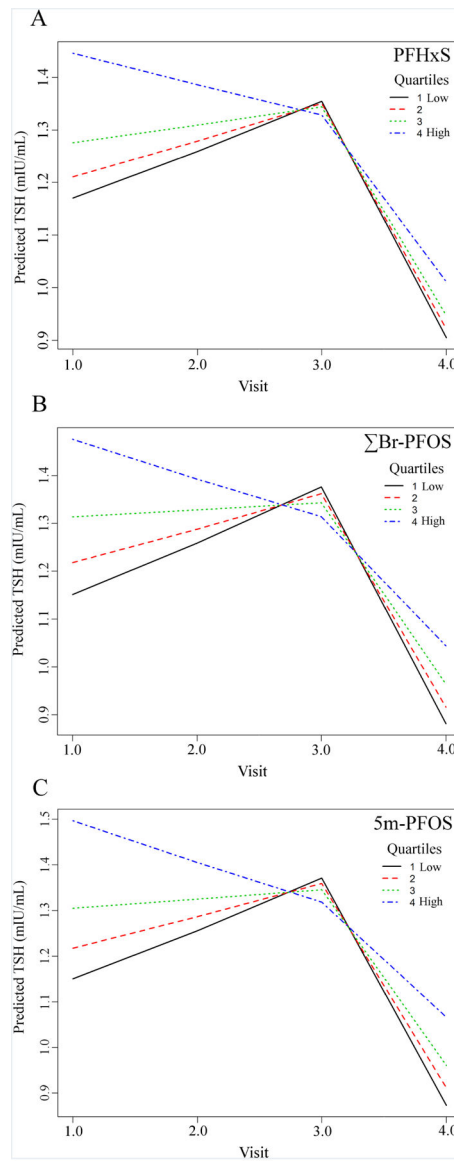


Figure 1. Change in predicted TSH (mIU/L) with each median quartile concentration (ng/mL) of PFHxS (A), Br-PFOS (B), and 5m-PFOS (C), with binary predictors at a fixed reference level and mean maternal age. Includes an interaction term which best fits the directional change from increasing in TSH levels (mIU/mL) during pregnancy (Timepoints 1 – 3) to decreasing TSH postpartum (Timepoint 4).

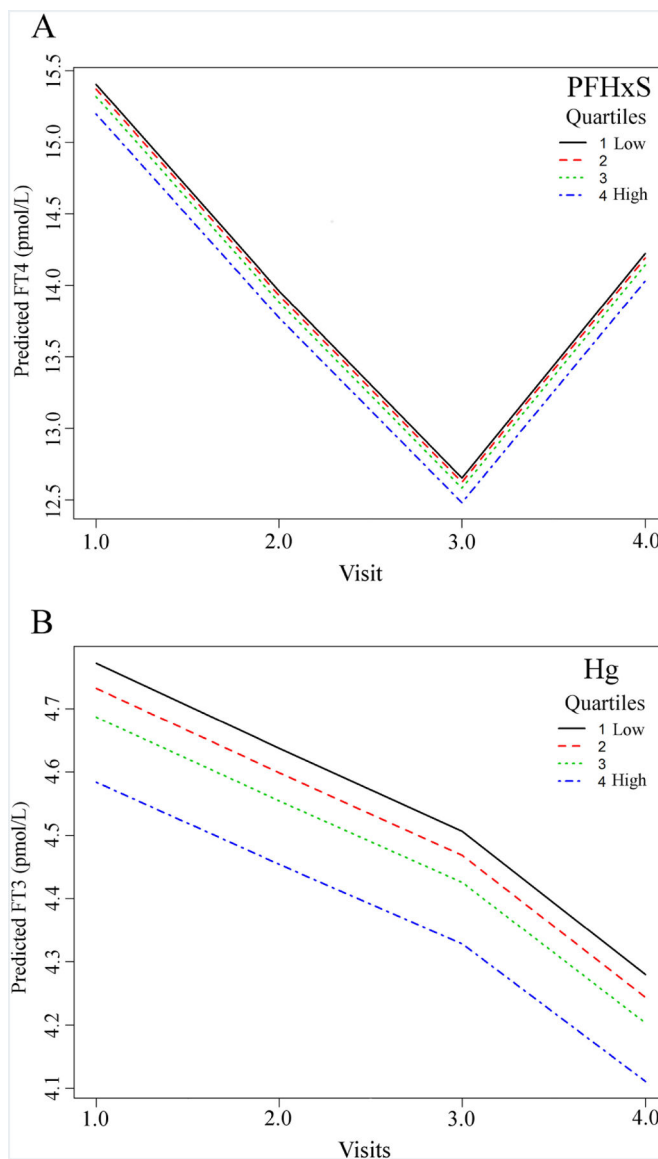


Figure 2. Change in predicted FT4 (pmol/L) with each median quartile concentration (ng/mL) of PFHxS (A), and the change in predicted FT3 (pmol/L) with each median quartile concentration ($\mu\text{g/L}$) of Hg (B), with binary predictors at a fixed reference level and mean maternal age. Includes an interaction term which best fits the directional change of concentration of FT4 (A), and Hg (B) during pregnancy (Timepoints 1 to 3) to post pregnancy (Timepoint 4).

Table 1.

Concentrations, and detection frequency of PFAAs (ng/mL), Hg ($\mu\text{g/L}$), and isomers of PFOS in maternal plasma (n=494), including isomer proportions of total PFOS (%).

	AM (SD)	GM	Min	Median	Max	% > DL	Total PFOS (%) ^b
PFHpA	0.14 (0.33)	0.04	< LOD ^a	0.08	3.87	66.6	
PFOA	2.88 (3.38)	2.12	0.265	2.11	43.3	100	
PFNA	1.05 (1.27)	0.76	< LOD ^a	0.69	15.8	99.0	
PFDA	0.38 (0.51)	0.26	< LOD ^a	0.25	7.01	99.8	
PFUnA	0.21 (0.15)	0.16	< LOD ^a	0.06	1.26	88.5	
PFDoA	0.05 (0.050)	0.03	< LOD ^a	0.06	0.68	55.5	
PFHxS	1.52 (1.76)	1.01	0.03	1.03	15.9	100	
PFOS	5.18 (2.69)	4.54	0.66	4.77	17.9	100	
Hg	1.47 (1.32)	0.98	0.03	1.06	8.68	97.6	
<i>Isomers of PFOS</i>							
Linear-PFOS	2.72 (1.49)	2.34	0.14	2.49	9.15	100	69.0
Br-PFOS	1.22 (0.66)	1.08	0.25	1.08	4.77	100	31.0
1 <i>m</i> -PFOS	0.08 (0.05)	0.06	< LOD ^a	0.06	0.43	96.8	2.1
3 <i>m</i> +4 <i>m</i> -PFOS	0.26 (0.14)	0.23	0.05	0.23	1.19	100	6.1
5 <i>m</i> -PFOS	0.38 (0.23)	0.32	0.06	0.33	1.63	100	9.8
<i>Iso</i> -PFOS	0.50 (0.28)	0.43	0.11	0.42	1.92	99.6	12.7

Note: AM, arithmetic mean; GM, geometric mean; SD, standard deviation; DL, detection limit. Total populations samples were collected at Timepoints 1 (0.6 %), 2 (96 %), 3 (3 %), and 4 (0.4 %)

^aValue below methods quantitation limits (MQL; ng/mL), detailed in supplementary material

^bTotal PFOS (%) applies to isomers of PFOS only

Thyroid hormone concentrations in maternal plasma over the 3 trimesters of pregnancy (Timepoints 1 to 3) and three months postpartum (Timepoint 4).

Table 2.

	Timepoint 1 (n=167)			Timepoint 2 (n=487)			Timepoint 3 (n=465)			Timepoint 4 (n=479)		
	GM (GSD)	Median (range)		GM (GSD)	Median (range)		GM (GSD)	Median (range)		GM (GSD)	Median (range)	
TSH												
All	1.04 (2.65)	1.18 (0.15, 3.21)		1.36 (1.96)	1.38 (0.57, 3.15)		1.20 (1.76)	1.28 (0.52, 2.45)		0.90 (3.20)	1.14 (0.07, 2.28)	
High TPOAb	*2.03 (3.48)	*2.07 (0.48, 8.01)		*1.60 (2.92)	*2.01 (0.59, 4.58)		1.13 (2.53)	1.42 (0.20, 2.57)		0.36 (8.42)	0.53 (0.07, 2.28)	
Normal TPOAb	0.94 (2.41)	1.10 (0.15, 2.52)		1.32 (1.79)	1.33 (0.57, 2.76)		1.22 (1.63)	1.27 (0.54, 2.38)		1.11 (1.74)	1.17 (0.50, 2.28)	
FT4												
All	15.1 (1.18)	14.9 (11.8, 19.6)		14.1 (1.16)	14.0 (11.6, 17.5)		12.9 (1.24)	12.6 (10.0, 16.6)		14.3 (1.18)	14.03 (11.6, 19.4)	
High TPOAb	14.6 (1.24)	14.2 (8.9, 19.6)		14.8 (1.18)	14.4 (11.3, 19.6)		13.7 (1.30)	13.4 (10.6, 18.2)		*16.5 (1.51)	*15.8 (11.6, 28.2)	
Normal TPOAb	15.2 (1.17)	15.1 (12.1, 18.7)		14.0 (1.15)	14.0 (11.6, 17.0)		12.7 (1.23)	12.5 (10.0, 15.9)		14.0 (1.18)	13.9 (11.7, 17.0)	
FT3												
All	4.64 (1.16)	4.62 (3.72, 5.58)		4.54 (1.14)	4.53 (3.69, 5.51)		4.43 (1.23)	4.36 (3.62, 5.41)		4.16 (1.18)	4.13 (3.30, 5.30)	
High TPOAb	4.38 (1.13)	4.38 (3.46, 4.99)		4.53 (1.18)	4.55 (3.61, 5.47)		4.41 (1.13)	4.47 (3.79, 5.11)		4.55 (3.26)	4.49 (3.16, 7.77)	
Normal TPOAb	4.68 (1.16)	4.69 (3.81, 5.58)		4.54 (1.14)	4.53 (3.70, 5.51)		4.43 (1.25)	4.35 (3.61, 5.42)		4.10 (1.14)	4.10 (3.30, 5.05)	
TPOAb ^a	n (%)			73 (15 %)			70 (15 %)			67 (14 %)		

Note: Sample collection: Timepoint 1 (< 13 weeks), Timepoint 2 (14 – 26 weeks), Timepoint 3 (27 – 40 weeks), Timepoint 4 (3 months postpartum) GM, geometric mean; GSD, geometric standard deviation

Range defined as the 5th percentile (lower) and 95th percentile (upper) of the sample population.

* p < 0.05, level of significance (high vs. normal TPOAb)

^aThreshold indicating participants with high TPOAb, n (%) > 9 IU/mL

Table 3.

Mixed effects model of overall associations between exposure to PFAs, Hg or PFAA and Hg co-exposure with maternal thyroid hormones.

PFAs	Main Effect (n=478) β (95 % CI)	Timepoint 1 (n=151) β (95 % CI)	Timepoint 2 (n=471) β (95 % CI)	Timepoint 3 (n=449) β (95 % CI)	Timepoint 4 (n=463) β (95 % CI)
TSH	0.144 (0.039, 0.249) **	0.093 (0.025, 0.160) **	0.042 (0.007, 0.078) *	-0.009 (-0.045, 0.028)	0.049 (-0.001, 0.098)
Br-PFOS ^{a,b,c}	0.286 (0.017, 0.555) *	0.180 (0.007, 0.352) *	0.073 (-0.021, 0.167)	-0.034 (-0.132, 0.065)	0.122 (-0.011, 0.255)
L-PFOS ^{a,c}	0.005 (-0.031, 0.041)	0.005 (-0.031, 0.041)	0.005 (-0.031, 0.041)	0.005 (-0.031, 0.041)	0.062 (0.007, 0.118) *
5m-PFOS ^{a,b,c}	0.851 (0.094, 1.608) *	0.541 (0.055, 1.03) *	0.230 (-0.035, 0.496)	0.080 (-0.357, 0.196)s	0.411 (0.034, 0.787) *
3m+4m PFOS(*Hg) ^{a,d,e}	1.357 (0.060, 2.65) *	0.998 (0.106, 1.89) *	0.639 (0.036, 1.24) *	0.281 (-0.343, 0.905)	0.920 (0.173, 1.67) *
FT4	-0.006 (-0.012, -0.001) *	-0.006 (-0.012, -0.001) *	-0.006 (-0.012, -0.001) *	-0.006 (-0.012, -0.001) *	-0.006 (-0.012, -0.001) *
FT3	-0.122 (-0.302, 0.059)	-0.122 (-0.302, 0.059)	-0.122 (-0.302, 0.059)	-0.122 (-0.302, 0.059)	0.333 (0.082, 0.583) *
Hg	-0.016 (-0.023, -0.008) *	-0.016 (-0.023, -0.008) *	-0.016 (-0.023, -0.008) *	-0.016 (-0.023, -0.008) *	-0.016 (-0.023, -0.008) *

Significance level

* p < 0.05

** p < 0.01

model estimates were calculated using log-values of thyroid hormone concentrations where β = change in log(hormone) per unit (1ng/mL) increase in PFAA.

^aEffect of PFAs on thyroid hormones after adjusted for moderately significant covariates (p < 0.2) in univariate models

^bAssociation includes time as a significant interaction (all Timepoints)

^cAssociation includes birth as a significant interaction (Timepoint 4 only)

^dAssociation includes Hg as a significant interaction

^eInteraction of Hg, effect sizes within the table were calculated when Hg set to a reference value

Table 4.

Mixed effects model coefficients for associations between PFAAs and thyroid hormones in adjusted models after including interaction term for TPOAb status

		Time	Normal TPOAb β (95 % CI)	High TPOAb β (95 % CI)	TPOAb Interaction ^c (p-value)
FT4	PFUnA ^a	all	-0.011 (-0.087, 0.066)	-0.240 (-0.456, -0.025)*	0.049
TSH	1m-PFOS ^{a,b}	1	1.97 (0.198, 3.74)*	-0.819 (-3.13, 1.49)	0.008
		2	1.36 (0.218, 2.51)*	-1.427 (-3.32, 0.468)	
		3	0.754 (-0.273, 1.78)	-2.03 (-3.88, -0.190)*	
		4	0.146 (-1.40, 1.69)	-2.64 (-4.83, -0.453)*	

^aEffect of PFAAs on thyroid hormones after adjusting for significant covariates in univariate models

^bSignificant interaction with time, coefficient calculated for main effect (all) and for each Timepoint (#)

^cLevel of significance (p-value) of TPOAb interaction term of normal vs. high TPOAb of the main effect (all Timepoints) using a threshold of 9 IU/mL

* p < 0.05

** p < 0.01 level of significance