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Grandmaternal perinatal serum DDT in relation to granddaughter early menarche and adult obesity: Three generations in the Child Health and Development Studies cohort

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Abstract

Background: Serum DDTs during or just after pregnancy were associated with breast cancer in mothers (F0), and with breast cancer, mammographic density and obesity in adult daughters (F1) in the Child Health and Development Studies multi-generational cohort in prior publications. Here we investigate F0 perinatal serum DDT associations with granddaughters' (F2) measured obesity at a median age of 26 and self-reported age at menarche.

Methods: F2 weight, height and waist circumference were measured by trained examiners. *o,p'*-DDT, *p,p'*-DDT and *p,p'*-DDE were measured in archived F0 perinatal serum. F0 DDT associations with F2 outcomes, accounting for F1 characteristics, were estimated in log-linear models adjusted for F0 and F1 body mass index (BMI), race, and menarche timing (N=258 triads for obesity; N=235 triads for early menarche). Interactions between F0 BMI and DDTs were estimated.

Results: F0 *o,p'*-DDT was associated with F2 obesity (Odds ratio, OR, 2.6; 95% Confidence Interval (CI), 1.3, 6.7, tertile 3 vs. 1), among normal weight F0 (70%), but not among overweight and obese F0 (p-interaction=0.03), independent of other DDTs. F0 *o,p'*-DDT was also associated with F2 early menarche (OR, 2.1; 95% CI, 1.1, 3.9, tertile 3 vs. 1) and this association was not modified by F0 BMI.

Conclusions: Ancestral exposure to environmental chemicals, banned decades ago, may influence the development of earlier menarche and obesity, which are established risk factors for breast cancer and cardiometabolic diseases.

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AUTHOR CONTRIBUTIONS

P. M. Cirillo: Formal analysis, methodology, writing-original draft, writing-review and editing. M. M. La Merrill: Conceptualization, writing-original draft, writing-review and editing, funding acquisition. N.Y. Krigbaum: Data curation, writing-review and editing. B. A. Cohn: Conceptualization, writing-original draft, writing-review and editing, funding acquisition.

CONFLICT OF INTERESTS

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

Impact: Discovery of actionable biomarkers of response to ancestral environmental exposures in young women may provide opportunities for breast cancer prevention.

Keywords

DDT; *o,p'*-DDT; Developmental Origins of Health and Disease; Early Age at Menarche; Obesity; Breast Cancer Risk Factors; Multi-generational; Child Health and Development Studies; CHDS; Prospective; cohort; women's health; environmental chemicals; cardiometabolic risk; pregnancy

INTRODUCTION

Persistent organic pollutants (POPs) were ubiquitous, world-wide chemical exposures during the 1960s (1–3). As such these legacy chemicals, including DDT, remain highly relevant to the health of people born in the 1960s who were exposed during development and to their children born in the 1980s and 1990s who were exposed when they were in ovum.

The commercial product known as “DDT” was composed primarily of *p,p'*-DDT, the active insecticide, but *o,p'*-DDT was an additional low level contaminant (about 15%) of commercial DDT (4). The primary metabolite of *p,p'*-DDT, *p,p'*-DDE, is the most environmentally persistent of these DDT compounds (https://www.cdc.gov/biomonitoring/DDT_BiomonitoringSummary.html). In contrast, *o,p'*-DDT is more quickly metabolized than either *p,p'*-DDT or *p,p'*-DDE and exhibits considerable inter-individual differences in metabolism (5). Importantly, the different chemical structures of these DDTs results in different biological activities. For example, *o,p'*-DDT is known as more estrogenic, with *p,p'*-DDE as largely anti-androgenic (6). During active use of DDT, as during the 1960's when bloods were collected in the current study, these three compounds could all be found in human serum (7–11). However, consistent with more rapid metabolism of *o,p'*-DDT, after DDT was banned in the U.S. in 1972, serum levels of *o,p'*-DDT were detected in only about 1 percent of NHANES II (1976–1980) samples (12). In contrast, exposure to *p,p'*-DDE, the primary metabolite of *p,p'*-DDT, is ongoing worldwide due to its ubiquitous contamination of the food supply (13). These secular changes are reflected in a series of breast cancer studies in human populations conducted with blood samples from 1960s to early 2000s which demonstrate dramatic declines in *p,p'*-DDT that were steeper than for *p,p'*-DDE (7,8). It is important to note that DDT has not been universally banned, thus exposure to DDT remains ongoing among people living where DDT is or has been recently manufactured or used, as well as among migrants from these countries (14–16).

The fetal origins of disease model introduced by Barker (17) and further developed as the Developmental Origins of Health and Disease concept by Gluckman and Hanson (18) extends to chemical exposures (19,20). A meta-analysis of 7 prospective cohort studies across North American, Europe and Asia found that prenatal or early life *p,p'*-DDE exposure was associated with obesity in people up to twenty years old (13). Since the publication of that meta-analysis, it was shown in the Child Health and Development Studies (CHDS) cohort that perinatal *o,p'*-DDT exposure in the 1960s was associated with increased risk of obesity in middle-aged (fifties) daughters (11). Experimental evidence supports these findings; for example mice dosed with a mixture of *p,p'*-DDT and *o,p'*-DDT (replicating their commercial mixture) during pregnancy and early lactation at exposure doses that

produced circulating DDT and DDE levels in the mouse mothers (F0) within the range of the CHDS F0 mothers had obese ‘daughter’ (F1) mice (21). Multiple generations of rats were also vulnerable to *p,p'*-DDT-induced obesity and ovarian diseases, such as polycystic ovarian syndrome and primary ovarian insufficiency (22–24). The multi-generational obesogenic and ovarian effects of environmental DDT and DDE isoform exposures that have been observed experimentally (22–24) remain to be investigated or demonstrated in humans.

Because *p,p'*-DDT commercial formulas were contaminated with *o,p'*-DDT and both are metabolized in living creatures to their DDE isoforms, these chemicals frequently co-exist in human observation and rodent experimental studies. We suspect because *o,p'*-DDT is shorter lived among these POPs, that it serves as the most sensitive biomarker of uniquely perinatal exposures and associated susceptibilities and thus may be the best biomarker for perinatal exposure among these related compounds in humans. This assumption is supported by findings in the CHDS. F0 perinatal *o,p'*-DDT has been shown to be the relevant DDT compound in association with daughters’ (F1) midlife obesity (11), breast cancer (9), breast density (10) and DNA methylation in breast cancer related genes (25). Hence, we focused on F0 perinatal *o,p'*-DDT as the primary potential predictor of granddaughter outcomes given that the granddaughters are exposed in the egg during their mothers’ development *in utero*.

In the present paper we test the hypothesis that perinatal *o,p'*-DDT exposure in the F0 (grandmother) generation of the CHDS cohort is associated with age at menarche and also with obesity and adiposity at age 26 in the F2 (granddaughter) generation of the CHDS cohort. This is the first paper to report on human associations of F0 exposures to POPs with F2 health outcomes.

MATERIALS AND METHODS

Study Population

The CHDS is a population-based, multi-generational cohort with ongoing follow-up for more than 60-years, beginning in the 1960s (26). Archived serum samples drawn during pregnancy and the early postpartum from the original generation (F0) make it possible to examine associations of environmental chemical levels with health outcomes in 3 generations: founding generation of women exposed during pregnancy (F0), the offspring generation exposed *in utero* during development (F1), and the grandchild generation exposed in the egg (F2). The timing of the CHDS F0 pregnancies, in the early 1960s, coincides with high use of legacy pesticides and industrial chemicals (e.g. DDTs, PCBs and PFAS compounds) (7,27,28). The CHDS F0 are now in their late 70s, their offspring (F1) are in their late 50s, and the grandchildren (F2) are in their mid-20s. Thus, the CHDS represents a unique opportunity to test the concept that ancestral exposures during these critical windows impact the health of current human populations. The CHDS recruited more than 98% of women seeking obstetric care at the Kaiser Foundation Health Plan in the San Francisco East Bay Area from 1959–1967 (26). These founding grandmothers (F0) were interviewed in person early in pregnancy, and provided permission to access medical records for themselves and their offspring (F1) and gave blood samples at each trimester and one in the near post-partum, generally within 3 days of delivery.

The current study is based on the daughters (F1) and granddaughters (F2) who participated in the *Three Generations of Breast Cancer (3Gs) Study* conducted from 2010 to 2013. Daughter offspring (F1) born into the CHDS who were surviving at the time of study recruitment and were phone locatable were eligible to participate in the telephone interview phase of the 3Gs Study (N=5,003). Daughters known to be incarcerated, institutionalized, to have a severe mental illness, or to have requested “not to be contacted”; and, who were already eligible for another adult follow-up study in progress were excluded from eligibility. Sixty percent of eligible subjects completed the phone interview (N=3,003). Due to budget constraints, a subset were targeted for participation in a home visit from the following three groups: daughters of mothers with breast cancer, daughters who had participated in an earlier breast density study and a random sample of daughters (29). Of the 1,879 daughters who were eligible, 1,194 completed a home visit (64%). Daughters (F1) were asked to invite their daughters (granddaughters, F2) to participate in the home visit along with them, using the technique of snowball recruitment. In total, 729 F2 granddaughters participated in the home visit and 356 F2 completed an adult questionnaire designed for participants ages 18 years. The distribution of baseline grandmaternal characteristics by participation type for F1 and F2 samples was highly comparable, suggesting that participation bias is unlikely (Supplementary Table 1).

During the home visit, anthropometric and blood pressure measurements were taken from F1 and F2, using standardized protocols. The F1 and adult F2 questionnaires included questions about menarche, menstruation and pregnancy, health behavior and body image.

Analysis Sample.

The present analysis sample is based on the F2 who completed an adult questionnaire as part of the 3Gs Study (n=365), and also participated in a home visit, had available DDT measures from grandmothers’ serum, and had available information on body mass index (BMI) in all three generations (n=258). The analysis sample for age at menarche associations was slightly smaller, requiring information on age at menarche in all three generations (n=235).

The institutional review board of the Public Health Institute approved the study protocols for this research. At enrollment CHDS F0 gave informed oral consent, as was customary in the 1960’s, for themselves and their children (F1). F1 and F2 who participated in the 3Gs Study from 2010–2013 gave full informed verbal consent before completing the study surveys and written consent before participating in the home visits.

Granddaughter (F2) Outcomes.

Weight and height of the F2 were measured during the 3Gs home visit using a detailed, standardized protocol. Examiners were trained to strictly implement the protocol. Standing height, weight and waist circumference was measured using a standardized protocol (11). Only measured F2 height and weight were used as outcomes in this study. Body mass index (*BMI*) was calculated from weight (kg) divided by height (m²), measured during the home visit. Age at menarche was assessed from self-report via the F2 adult survey completed on hard copy or online. Other socio-demographic information such as F2 age, race and

education were assessed from the adult survey. To create outcome variables based on clinically relevant categories, BMI was classified as ≥ 30 vs. <30 kg/m² to represent obesity in the F2 (30) and waist circumference was classified as ≥ 80 cm vs. <80 cm (31). Age at menarche was classified as ≤ 11 vs. >11 years to represent early menarche. Age 11 was selected as the threshold for early menarche because it was the 25th percentile cut-point for the distribution of age at menarche in the F2.

Grandmother (F0) Exposures and Covariates.

We measured *o,p'*-DDT, *p,p'*-DDT, *p,p'*-DDE, and lipids in non-fasting grandmother (F0) perinatal serum samples collected during 1959–1967 preferentially collected in the early postpartum (1–3 days after delivery) as previously described (9,32). Some subjects lacked postpartum samples so alternative samples were used. F0 serum drawn in the early postpartum accounts for 86% of DDT assays in this study, serum from the 3rd trimester was used for 12%, and serum from the 2nd trimester was used for 2% of assays. Adding trimester of draw as a covariate to models did not affect associations observed for F2 obesity and early menarche, nor was trimester of draw, itself, associated with these outcomes.

Previous research has demonstrated measurement reliability over time and correspondence in levels across gestation for these chemicals, supporting the assumption that timing of specimen collection during the perinatal period is representative of pregnancy exposures (33). Samples were measured using gas chromatography with electron capture detection (GC/ECD) fit with a capillary column (18%), GC/ECD fit with dual columns (45%), or GC triple quadrupole mass spectrometry (37%). Companion assays for total cholesterol and triglycerides were measured at the Clinical and Epidemiologic Research Laboratory (CERLab) at Boston Children's Hospital, using methods previously described (34).

DDTs were characterized as continuous variables, as log-transformed variables and as tertiles using 2 dummy variables representing tertile 2 and tertile 3, versus tertile 1 as the reference category. Tertile variables were used to test for patterns of dose response.

Demographics and health-related behavior were collected from grandmothers during in-person interviews at enrollment, including age, race and education. Clinical measures were abstracted from medical records beginning 6 months prior to pregnancy through labor and delivery and are the source of data on baseline weight of grandmothers. Grandmothers' BMI was calculated from *weight* (kg) divided by height (m) squared, measured or reported at interview or first prenatal visit. Weight was adjusted to compensate for variation in the timing of measurement by regressing weight on gestational age using the locally weighted scatterplot smoothing technique (35). Adjusted weight was then imputed as the fitted mean weight at day 101 of gestation (median value for day of interview) plus the residual from the regression procedure. F0 BMI was assessed as a continuous variable and as a binary variable, dichotomized as overweight, BMI ≥ 25 vs. <25 kg/m². Grandmothers' age at menarche was collected from both baseline interview and medical record.

Daughter (F1) Exposures and Covariates.

Weight and height at age 30 years, and age at menarche, as well as age and other socio-demographic information were assessed via self-report during a computer-assisted telephone

interview. F1 BMI at age 30 was calculated from reported weight and height at age 30. F1 BMI at age 30 and F1 age at menarche were implemented as continuous variables.

Statistical Methods.

We implemented log-linear models (PROC GENMOD) executed in SAS 9.4 to estimate associations using the repeated option to adjust the covariance matrix for correlation within family clusters. Models included all three DDTs: *o,p'*-DDT, *p,p'*-DDT and *p,p'*-DDE. Although the DDTs are correlated, each compound represents different exposure sources, metabolic response, and bioactivity. We have recently examined this empirically in a metabolomics analysis that identified different correlated metabolomic pathways for these compounds (36). The decision to mutually adjust for these compounds is based on biological considerations as previously reviewed (8). The three DDTs are different exposures that vary in proportion in individuals and have different toxicities and consequences. *p,p'*-DDE is acquired independently of its parent compound, *p,p'*-DDT, via environmental sources due to extreme persistence and via individual metabolism during active exposure. The relative amount of *o,p'*-DDT compared to *p,p'*-DDT is an indication of recency of exposure to commercial DDT because *o,p'*-DDT is metabolized and excreted more rapidly than *p,p'*-DDT. The biological activities of these compounds differ as well because of their different chemical structures (6). So, one of these compounds cannot be a proxy for the others (8). This is partly empirically revealed by finding that associations can be in opposing directions for these compounds or observed only for one of the three—very likely for complex reasons including metabolic rate, and exposure source and timing. It is essential to consider the compounds together to exclude mutual confounding. In the CHDS, we have previously observed different relationships for these different DDTs with various outcomes in jointly adjusted models (7–9,36–38). We have also observed that individuals in our population have widely varying relative proportions of these compounds – consistent with differences in exposure timing, exposure sources and metabolism that is particular to the active use and intake of commercial DDT during the 1960's (39). These variations suggest that these compounds should be considered in models together to determine differential effects. As noted in the introduction, in this paper we focus on *o,p'*-DDT associations as a marker for recent perinatal DDT exposure and because in utero exposure to *o,p'*-DDT has been associated with F1 outcomes in the CHDS. However, in supplementary tables we present estimated coefficients for all three DDT compounds for completeness.

Models were stratified by whether F0 had a BMI ≥ 25 kg/m² vs. <25 kg/m² at pregnancy interview. We applied a lower BMI threshold for F0 (≥ 25 kg/m²) than for F2 (≥ 30 kg/m²) owing to the different distributions in the two samples. Since only 9% of F0 met the definition of obesity (>30 kg/m²) compared to 32% of F2, we combined the overweight and obese categories to examine BMI in F0. Our rationale for stratifying by grandmothers' (F0) BMI is that *p,p'*-DDT and *p,p'*-DDE are highly lipophilic and grandmothers' BMI could serve to moderate exposures during gestation. Stratified models that estimated F0 *o,p'*-DDT associations with F2 obesity (BMI ≥ 30 kg/m²) included: F0 pregnancy *p,p'*-DDT (continuous), *o,p'*-DDT (continuous), *p,p'*-DDE (continuous), race and self-reported F1 BMI at age 30 years. To support the decision to stratify models by F0 overweight we created a product term between F0 BMI (continuous) X *o,p'*-DDT (continuous) to test whether

associations estimated from stratified models were statistically different. A similar strategy was used for models estimating early menarche (≤ 11 vs. >11 years) which included continuous terms for *p,p'*-DDT, *o,p'*-DDT and *p,p'*-DDE, and were adjusted for F0 age at menarche and race, and F1 age at menarche and self-reported weight at age 30. In a set of secondary analyses designed to evaluate adiposity as distinct from BMI, we also examined F0 *o,p'*-DDT associations with large F2 waist circumference (≥ 80 cm vs. <80 cm) in models as described for F2 obesity. We estimated odds ratios for the difference between high F0 *o,p'*-DDT (median of 3rd tertile) levels vs. low levels (median of 1st tertile) using contrasts calculated from models for each outcome. We chose tertiles based on our prior work which also used this classification (7,9). We identified outlier *o,p'*-DDT values using the exploratory data approach (EDA) described by Tukey (40) and conducted a sensitivity analysis to determine whether results were impacted by extreme values.

RESULTS

Table 1 provides distributions of study variables for three generations, grandmother (F0), daughter (F1) and granddaughter (F2). Birth cohorts for the three generations span more than 60 years, with an approximate 25-year span between the median ages at data collection for successive generations. This study observed grandmothers at a median age of 26 years in 1962 (median year of interview), daughters at a median age of 49 years and granddaughters at a median age of 26 years, the same median age as when their grandmothers donated biospecimens during pregnancy. Consistent with secular changes observed nationally and worldwide (41), weight and BMI increased in consecutive generations. Age at menarche also shows an expected downward trajectory between the F1 and F2 generations, also consistent with international patterns (42–45).

Table 1 also presents distributions for the serum DDT compounds measured in grandmothers (F0) during their pregnancy. High levels of both DDT isomers (*p,p'*-DDT and *o,p'*-DDT) result from the timing of the blood draw. F0 serum was taken when DDT was in active use in the U.S., prior to its ban in 1972 (46).

As shown in Table 1, there was strong representation among African Americans in all three generations that derives from the founding generation which was an archetypal population-based sample of Alameda County, CA with a robust community of African Americans (47). In the ensuing generations, care was taken to ensure continued participation from African Americans by diligently recruiting them to the follow-up 3Gs Study to achieve representation in proportion to the baseline population. Table 1 also demonstrates an upward progression of educational level across F0 and F1 generations. The greater number of women at the highest level of education in the F1 generation compared to the F2 is likely due to the differences in the upper age boundary in the two generations. The F1 are older and have had more opportunity to attain post-graduate credentials.

Figure 1, panel A provides the distributions of grandmothers' (F0) pregnancy serum *o,p'*-DDT (ng/mL) levels by granddaughter (F2) obesity (BMI ≥ 30 vs. <30 kg/m²) within F0 weight sub-groups (BMI ≥ 25 vs. <25 kg/m²). F2 obesity was correlated with higher F0 *o,p'*-DDT for normal weight F0 but this pattern was opposite for overweight F0. Thus, in further

analyses we stratified F2 obesity analyses by F0 BMI (≥ 25 vs. <25 kg/m²). The rationale to stratify by F0 overweight is provided by the statistically significant p-value for the product term, *o,p'*-DDT x F0 BMI ($P_{\text{interaction } (o,p'\text{-DDT} \times \text{F0 BMI})} = 0.0276$) in a model adjusted for all DDT congeners.

Figure 1, Panel B provides the distributions of grandmothers' (F0) pregnancy serum *o,p'*-DDT (ng/mL) levels for early granddaughter (F2) menarche (age at menarche ≤ 11 vs. >11 years) and shows that granddaughters with early menarche had higher *in ovo* exposures to *o,p'*-DDT.

Table 2 provides modeled associations of grandmothers' (F0) perinatal serum *o,p'*-DDT with granddaughter (F2) obesity (BMI ≥ 30 vs. <30 kg/m²) for various levels of model adjustment, stratified by F0 overweight (BMI ≥ 25 vs. <25 kg/m²).

Granddaughter (F2) obesity for normal weight grandmothers (F0).

Risk of obesity in granddaughters was 2 to 3-fold greater when grandmothers' *o,p'*-DDT was in tertile 3 compared to tertile 1 (Table 2). A significant >2 -fold association of F0 *o,p'*-DDT with F2 obesity remained after adjustment for F0 African American race and F1 BMI at age 30 (a proxy for F1 BMI during F2 gestation). The adjusted models in Table 2 show that the F0 *o,p'*-DDT association with F2 obesity persists, even though F0 African American race and F1 BMI at age 30 were positively and significantly correlated with F2 obesity. The F0 *o,p'*-DDT association with F2 obesity was independent of grandmothers' *p,p'*-DDT which showed a significant negative correlation with F2 obesity (Supplementary Table 2A). Supplementary Figure 1 demonstrates that the positive association between grandmothers' *o,p'*-DDT and granddaughter BMI is comparable within tertiles of *p,p'*-DDT, establishing that the positive *o,p'*-DDT effect is not neutralized by the negative *p,p'*-DDT effect. This supports the conjecture that the impact of *o,p'*-DDT exposure per ng/mL on F2 obesity is more potent than that of the *p,p'*-DDT exposure, consistent with the larger absolute value for the *o,p'*-DDT coefficient compared to the coefficient for *p,p'*-DDT (Supplementary Table 2A).

Granddaughter (F2) obesity for overweight/obese grandmothers (BMI ≥ 25 kg/m²).

Risk of obesity in granddaughters was about 70% lower when grandmothers' *o,p'*-DDT was in tertile 3 compared to tertile 1 (Table 2). Adjustment for F0 African American race and F1 BMI at age 30 had little effect on these associations (Table 2). The sample size of overweight/obese grandmothers is small, reflecting the lower incidence of overweight and obesity in the 1960's compared to current obstetric populations. This smaller sample size likely reduced statistical power in the F0 overweight subset.

Granddaughter (F2) abdominal fat.

All but one of the F2 participants who were obese also had waist circumference ≥ 80 cm. For this reason, models that predicted F2 obesity, also predicted F2 waist circumference ≥ 80 cm.

Granddaughter (F2) Age at Menarche.

Unlike F2 obesity, the F0 *o,p'*-DDT association with F2 age at menarche did not vary significantly according to F0 BMI. Therefore, we present results for F2 age at menarche for all F0 (Table 3). Risk of early menarche (age 11 vs >age 11 years) in F2 was estimated to be about 2 times more likely when F0 *o,p'*-DDT levels were in tertile 3 compared to tertile 1. Adjustment by F0 African American race and age at menarche, and by F1 age at menarche and BMI at age 30 did not exert a notable effect on estimated F0 *o,p'*-DDT associations with F2 age at menarche (Table 3). The association of F1 BMI at age 30 with F2 early menarche did not reach statistical significance ($p=.0572$), but still is unlikely to have occurred by chance. *p,p'*-DDT was associated with a lower risk of F2 early menarche in the fully adjusted model (Supplementary Table 2B).

When *o,p'*-DDT is represented categorically (Supplementary Tables 3A and 3B), results are consistent with the linear associations estimated from models in Tables 2 – 3 where *o,p'*-DDT was represented as a continuous variable. As seen in Supplementary Table 2A and Supplementary Figure 1, *p,p'*-DDT and *o,p'*-DDT, which are positively correlated, have associations in opposite directions with F2 obesity and also with F2 age at menarche (Supplementary Table 2B). Thus, we include both in all models due to mutual confounding. Removing *p,p'*-DDE from models (null associations with F2 outcomes, Supplementary Tables 2A and 2B) has no substantial impact on findings, but we include *p,p'*-DDE in final models as it is the DDT compound measured in many studies and readers often wish to see or rule out its impact on health outcomes. Exclusion of outlier F0 *o,p'*-DDT values did not materially affect the associations.

DISCUSSION

Results show that grandmothers' *o,p'*-DDT was associated with an increase in granddaughter obesity and early age at menarche, presenting a consistent pattern of associations that support the concept that in egg exposure to *o,p'*-DDT has potential for altering risk of breast cancer across generations. To the best of our knowledge, such a study has never been previously conducted in humans.

The present study builds on a body of evidence from the CHDS that *o,p'*-DDT exposure during F0 pregnancy or immediately after birth increases risk of breast cancer in F1 daughters (9), and increases the prevalence of breast cancer risk factors among F1 daughters, such as obesity (11) and breast density (10) and altered DNA methylation of breast cancer related genes in midlife (25). The impact of grandmaternal DDT exposure on F2 breast density is unknown but could soon be investigated in this cohort either using new, age-safe techniques for estimating density in young adulthood such as ultrasound (48) or waiting until the F2 achieve the age of mammography screening.

Results in the F1 generation are consistent with another study based in California which found that prenatal *o,p'*-DDT exposure was associated with childhood adiposity (49). We are not aware of previous human studies that have examined associations between perinatal *o,p'*-DDT exposure and age at menarche. The two studies which only examined *p,p'*-DDE or *p,p'*-DDT prenatally, reported conflicting results, one found no association between prenatal

p,p'-DDE or *p,p'*-DDT and the reported timing of menarche (50) while the other study reported prenatal *p,p'*-DDE was associated with earlier menarche (51), consistent in principle with our results. We are also unaware of any studies of age at menarche or obesity after *grandparent o,p'*-DDT exposure. We suspect that perinatal *o,p'*-DDT is most consistently associated with obesity, menarche, and breast cancer across the CHDS generations because *o,p'*-DDT is relatively short lived in humans and thus is the best biomarker for perinatal exposure among these related compounds in humans.

Several experimental studies have reported that both *p,p'*-DDT and *o,p'*-DDT are toxic to ovarian follicles (24,52–54) (those which exist in the ovaries of an F1 fetus at the time of F0 exposure during pregnancy). Although the multigenerational evidence of rat ovarian disease was only demonstrated for *p,p'*-DDT in the F1 and F3 generations, and did not examine F2 rats or *o,p'*-DDT exposure, this body of experimental evidence raises the possibility that *o,p'*-DDT could directly target the F2 follicles within the F1 fetus at the time that F0 serum *o,p'*-DDT is measured. We speculate that changes in DNA methylation could unite the relationship of perinatal *o,p'*-DDT exposure to breast cancer risk. Perinatal *o,p'*-DDT has been associated with altered blood DNA methylation in adult CHDS F1 daughters at loci associated with breast cancer and /or menarche (25), and obesity (11,55–58). In addition, epigenetic changes were observed in F3 rats with heightened obesity and ovary disease prevalence after ancestral *p,p'*-DDT exposure (22–24). The F2 daughters are too young to examine whether F0 *o,p'*-DDT exposure will be associated with increased breast cancer risk. However, if obesity and/or age at menarche are on a causal pathway between *o,p'*-DDT and breast cancer then the impact of grandmaternal *o,p'*-DDT exposure across generations cannot be ruled out here.

Increased grandmothers' BMI appeared to have a protective effect on the association between grandmothers' *o,p'*-DDT and granddaughter obesity. Because DDT and DDE isoforms are lipophilic, excess body fat mass dilutes a given quantity of DDT or DDE exposure. However, if these DDT 'dilution' effects of BMI were the only or primary explanation for the observed interaction, we would expect grandmothers' BMI to also modify the DDT effect on age at menarche, which it did not. Thus, we suspect that the interplay between grandmothers' BMI and *o,p'*-DDT may be more complex than a simple decrease in dose due to 'dilution' and/or 'excretion' effects, possibly linked to activation of differential metabolic pathways in high BMI environments.

Weaknesses of this study include a lack of data on diet and exposures to chemicals at other periods of the lives of these three generations and lack of consideration of other unknown exposures that are correlated with DDTs. The F0 samples have been stored since the 1960's and DDTs were measured about 40 years later. While we cannot rule out some desiccation of the samples, desiccation is likely to apply similarly or at random for the samples in this study. The NIH repository that stores our samples reports when a sample we have requested appears desiccated. This is an infrequent occurrence, and in this case, we do not assay that sample. As this is an observational study, there is uncertainty regarding whether *o,p'*-DDT is a marker of recent exposure or is itself the primary toxic influence among the DDTs measured. We lack data from other geographic locations and our sample has limited representation of Hispanics and Asians. However, our F0 cohort represents in-migration

from East and South geographic areas of the U.S, where DDT was in heavy use (59); and, reflects the racial and ethnic composition of Alameda County California in the 1960's. We do not know how grandfather exposures may have contributed to our findings. The strengths of this study are its prospective design, direct measurement of multiple DDTs in F0 perinatal serum during active exposure, the ability to account for the contribution of grandmother and mother obesity and age at menarche to observed associations of grandmother DDTs with granddaughter obesity and age at menarche.

CONCLUSIONS

Our findings are consistent with the hypothesis that grandmother (F0) exposures to DDT could have contributed the dramatic increases in obesity in current young adult women including the granddaughter (F2) generation in the CHDS cohort. This conclusion is strengthened by our prior observation that F0 exposures to DDT were also associated with daughters' mid-life obesity (F1) in this same cohort. Notably, perinatal exposure to DDT is associated with breast cancer risk factors in subsequent generations.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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ABBREVIATIONS

CHDS	Child Health and Development Studies
DDT	a pesticide widely used worldwide from the 1940's to the 1970's and still used in some countries
DDTs	various chemical compounds related to the commercial insecticide known as DDT
<i>p,p'</i>-DDT	1,1,1-trichloro2,2-bis(<i>p</i> -chlorophenyl) ethane

<i>o,p'</i>-DDT	1,1,1-trichloro2-(<i>o</i> -chlorophenyl)- 2-(<i>p</i> -chlorophenyl)ethane
DDE	1,1-dichloro-2,2-bis (<i>p</i> -chlorophenyl) ethylene
F0	Founding generation in the Child Health and Development Studies refers to mothers pregnant at entry to the study
F1	offspring generation in the Child Health and Development Studies refers to daughters of F0 in this study
F2	grandchildren generation in the Child Health and Development Studies refers to granddaughters of F0 in this study
BMI	body mass index

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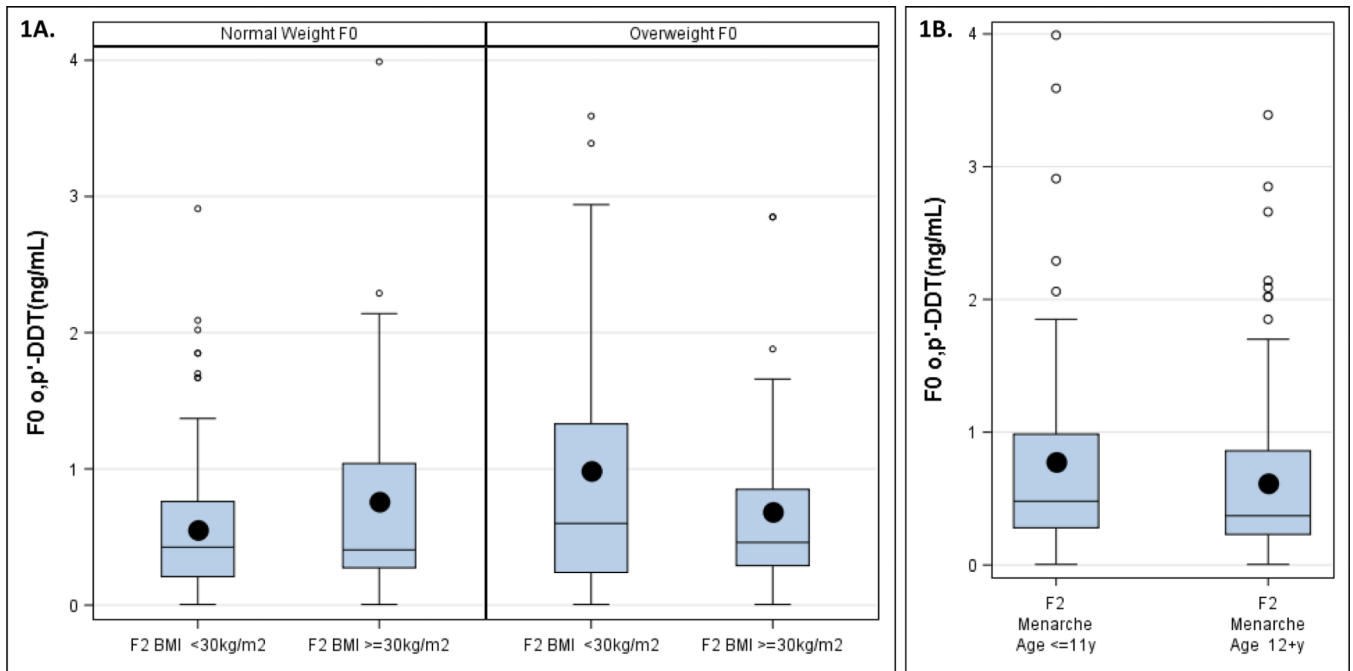


Figure 1. Distribution of grandmothers' (F0) serum *o,p'*-DDT (ng/mL) levels for: **A.** Granddaughter (F2) obesity (BMI ≥ 30 vs. <30 kg/m²) stratified by grandmothers' (F0) overweight (BMI ≥ 25 vs. <25 kg/m²) and **B.** Granddaughter (F2) early menarche (age at menarche ≤ 11 vs >11 years). Overall, F0 serum *o,p'*-DDT ranged from 0.0004 – 3.99 ng/mL.

Table 1. Distribution of study characteristics among three CHDS generations: grandmothers (F0), daughters (F1) and granddaughters (F2)

Study variable	Grandmothers (F0), n=258					Daughters (F1), n=258					Granddaughters (F2), n=258				
	Percentile					Percentile					Percentile				
	Mean (SD)	25 th	50 th	75 th		Mean (SD)	25 th	50 th	75 th		Mean (SD)	25 th	50 th	75 th	
Year of Birth	1936 (6.6)	1931	1937	1941	1961	1962 (1.8)	1961	1962	1964	1989 (3.6)	1987	1990	1993		
Year of Interview	1962 (1.8)	1961	1962	1964	2011	2011 (0.7)	2011	2012	2012	2013 (0.0)	2013	2013	2013		
Age (years)	26 (6.0)	22	25	31	49	49 (2.0)	48	49	51	26 (4.4)	22	26	29		
Weight (kg)	62.9 (14.0)	53.7	59.3	69.0	70.8	70.8 (16.4)	59.0	65.8	77.1	76.9 (23.4)	61.0	70.05	89.0		
Height (m)	1.62 (0.07)	1.57	1.63	1.68	1.66	1.66 (0.07)	1.60	1.65	1.70	1.66 (0.07)	1.62	1.66	1.71		
Body Mass Index (kg/m ²)	23.9 (5.1)	20.9	22.5	25.6	25.8	25.8 (5.6)	22.0	23.9	29.2	28.0 (8.4)	21.8	25.1	32.9		
Age at menarche (years)	12.7 (1.4)	12	13	13	12.8	12.8 (1.8)	12	13	14	12.4 (1.6)	11	12	13		
Parity	1.8 (2.0)	0	1	3											
Year of blood draw	1962 (1.8)	1961	1962	1964											
<i>p,p'</i> -DDT (ng/mL)	14.5 (10.4)	7.1	11.6	19.8											
<i>o,p'</i> -DDT (ng/mL)	0.66 (0.67)	0.24	0.45	0.86											
<i>p,p'</i> -DDE (ng/mL)	48.5 (23.8)	31.6	43.4	60.4											
	N (%)					N (%)					N (%)				N (%)
Race															
African American	78 (30%)					84 (33%)									73 (33%) ¹
Asian	6 (3%)					4 (2%)									12 (5%) ¹
Hispanic	13 (5%)					17 (6%)									17 (8%) ¹
White	153 (59%)					146 (56%)									117 (53%) ¹
Other/Mixed	8 (3%)					7 (3%)									3 (1%) ¹
Education															
<High School	53 (22%) ²					12 (4%)									13 (6%) ³
High School	89 (37%) ²					31 (12%)									48 (22%) ³

Study variable	Grandmothers (F0), n=258				Daughters (F1), n=258				Granddaughters (F2), n=258			
	Mean (SD)	25 th	50 th	75 th	Mean (SD)	25 th	50 th	75 th	Mean (SD)	25 th	50 th	75 th
Some College	52 (22%) ²				105 (41%)				115 (54%) ³			
College	44 (19%) ²				110 (43%)				38 (18%) ³			
BMI												
<25kg/m ²	182 (71%)				153 (59%)				129 (50%)			
25 --<30kg/m ²	52 (20%)				47 (18%)				47 (18%)			
30kg/m ²	24 (9%)				58 (23%)				82 (32%)			

¹ F2 missing on race, n=36.

² F0 missing on education, n=20.

³ F2 missing on education, n=44.

Associations of grandmothers' (F0) serum *o,p'*-DDT with adult F2 Obesity (< 30 v. >30 kg/m²), stratified by grandmothers' BMI², n=258 triads

Table 2.

Model Adjustment Level	Estimated <i>o,p'</i> -DDT association by grand-maternal (F0) BMI ²					
	Grandmothers' (F0) BMI <25 (n=182)		Grandmothers' (F0) BMI ≥ 25 (n=76)		<i>o,p'</i> -DDT Odds Ratio ³	p-value
	<i>o,p'</i> -DDT Odds Ratio ³	(95% CI) ⁴	(95% CI) ⁴	(95% CI) ⁴		
DDT congeners	3.62	(1.37, 9.56)	0.0092	0.37	(0.13, 1.02)	0.0557
+ F1 BMI ⁵ Adjusted	3.43	(1.47, 8.00)	0.0044	0.36	(0.11, 1.13)	0.0809
+ F1 BMI ⁵ + Race ⁶ Adjusted	2.59	(1.00, 6.73)	0.0505	0.31	(0.09, 1.01)	0.0526

¹ Associations are estimated from log-linear models adjusted for family clustering and stratified by F0 BMI (categorized as < 25 kg/m² vs >25 kg/m²). Unadjusted models include linear terms for *p,p'*-DDT, *p,p'*-DDE, *o,p'*-DDT. Models adjusted for F1 BMI added a linear term for F1 BMI reported at age 30 nearest the age when F1 were pregnant with F2 and nearest the ages when F0 and F2 BMI were captured. Models additionally adjusted for race added a dichotomous variable representing African Americans vs. all other races.

² The rationale to stratify by F0 overweight (BMI <25 vs. >25 kg/m²) is based on results from the model estimating F2 obesity that included linear terms for *p,p'*-DDT, *p,p'*-DDE, *o,p'*-DDT and F0 BMI, and the product term between *o,p'*-DDT (continuous) X F0 BMI (continuous). The p-value for the product term, Pinteraction (*o,p'*-DDT X F0 BMI) = 0.0276.

³ Reported *o,p'*-DDT Odds Ratios are estimated for the median of the 3rd vs. the median of the first tertile. The inter-tertile range for F0 *o,p'*-DDT at the medians was 0.86 ng/mL.

⁴ CI=Confidence Limits

⁵ BMI=Body Mass Index, continuous kg/m².

⁶ Race is dichotomized as African American (1) vs. all else (0).

Table 3.

Associations of grandmothers' (F0) serum *o,p'*-DDT with F2 early menarche¹, n=235 triads

Model Level of Adjustment	Estimated <i>o,p'</i> -DDT Association		
	Odds Ratio ²	95% CI ³	p-value
DDT congeners and F0 Age at Menarche Adjusted	2.23	(1.24, 4.01)	0.0076
+ F1 Age at Menarche Adjusted	2.25	(1.24, 4.08)	0.0077
+ Race Adjusted	2.06	(1.10, 3.87)	0.0235
+ F1 BMI Adjusted	2.08	(1.11, 3.90)	0.0222

¹ Early menarche was defined as age 11 vs. >11 years based on the 25th percentile for the distribution of F2 age at menarche. Associations are estimated from log-linear models adjusted for family clustering. Models include linear terms for *p,p'*-DDT, *p,p'*-DDE, *o,p'*-DDT and F0 age at menarche (continuous). Subsequent models were adjusted by cumulatively adding: F1 age at menarche (continuous), race dichotomized as African American (1) vs. all other races (0), and F1 BMI (continuous) reported at age 30 nearest the age when F1 were pregnant with F2 and nearest the ages when F0 and F2 BMI were captured.

² Reported *o,p'*-DDT odds ratios are estimated for the median of the 3rd vs. the median of the first tertile. The inter-tertile range for F0 *o,p'*-DDT at the medians was 0.86 ng/mL.

³ CI=Confidence Interval.