Insulin-like growth factor-I, its binding proteins (IGFBP-1 and IGFBP-3), and growth hormone and breast cancer risk in The Nurses Health Study II

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Abstract

Earlier data suggest that the relationship between circulating insulin-like growth factor I (IGF-I) levels and breast cancer risk differs according to menopausal status. We evaluated the association between IGF levels as well as the primary regulator of IGF-I production, growth hormone (GH), and breast cancer risk in the Nurses' Health Study II (NHS II) cohort, a large cohort of primarily premenopausal women. We conducted a case-control study nested within the prospective NHS II cohort. Plasma concentrations of IGF-I, IGF binding protein (IGFBP)-3, IGFBP-1, and GH were measured in blood samples collected between 1996 and 1999. Totally 317 women were identified who had a diagnosis of invasive or *in situ* breast cancer between the date of blood collection and June 1 2003; 75% of these women were premenopausal at blood collection. To each of the 317 women, two controls were age-matched for a total of 634 controls. We used conditional logistic regression models to estimate the relative risk of breast cancer. Overall, plasma IGF-I, IGFBP-1, IGFBP-3, and GH levels were not associated with breast cancer risk (relative risks, top vs bottom quartile; IGF-I, 0.98, 95% confidence interval (CI), 0.69–1.39; IGFBP-1, 0.95, 95% CI, 0.63–1.41; IGFBP-3, 1.10, 95% CI, 0.78–1.54; GH, 1.09, 95% CI, 0.82–1.46). These risks were similar for premenopausal women of age 45 years or less. Further adjustment for additional breast cancer risk factors did not change these estimates. In conclusion, circulating IGF-I, IGFBP-1, IGFBP-3, and GH levels appear to have no important association with breast cancer risk in a large cohort of premenopausal women.

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Introduction

Insulin-like growth factor I (IGF-I) is a peptide hormone that, with IGF-II, IGF binding proteins (IGFBPs), and cell-surface receptors, is integral to the regulation of cell survival and death. IGF-I appears to be associated with breast cancer risk, particularly among premenopausal women (Hankinson & Schernhammer 2003). However, results from observational studies are not entirely consistent (Kaaks *et al.* 2002, Hankinson & Schernhammer 2003), and uncertainty exists about the true association between IGF-I and breast cancer risk. IGF-I production in the liver, the primary source of circulating IGF-I (Jones & Clemmons 1995), is regulated primarily by human growth hormone (GH), which is produced by the pituitary gland. To our knowledge, no study has evaluated the association between circulating GH levels and breast cancer risk. IGFBP-3, along with IGFBP-1, are two of six

currently identified IGFBPs that, by binding IGF peptides, prolong their half-lives and may alter the interaction of IGFs with their cell surface receptors (Shimasaki & Ling 1991).

Using a nested case-control design within the large, prospective Nurses' Health Study II (NHS II) cohort of primarily premenopausal women, we investigated an association between plasma IGF-I, its binding proteins IGFBP-1 and IGFBP-3, and GH and breast cancer risk. C-peptide was measured for the purpose of conducting analyses stratified by C-peptide as an index of insulin resistance.

Materials and methods

Study population

The NHS II is a prospective cohort study that started in 1989 when 116 671 registered female US nurses aged 25 to 42 from 14 US states were enrolled. The NHS II was designed akin to the Nurses' Health Study (NHS), an earlier, independent cohort study of similar size that was initiated in 1976 (Colditz & Hankinson 2005). The baseline questionnaire sought hormone use, reproductive history, current medication, history of disease, and a number of life-style factors. Since then, women have been followed biennially by mailed questionnaires, ascertaining any diagnosis of breast cancer, including date of diagnosis. For women who reported a diagnosis of breast cancer, we requested permission to review relevant medical records, all of which were reviewed by trained physicians. For deceased participants, we sought permission from next-of-kin to review these records. More than 99% of reported breast cancers were confirmed by medical record review. Further details of the cohort have been published (Rockhill *et al.* 1998).

Women who had not previously reported a diagnosis of cancer were eligible for sample collections; in total, 29 611 women in the NHS II cohort participated in our blood collection study from 1996 to 1999. We provided blood collection kits and advised each participant to have blood samples drawn by a local laboratory or colleague. First samples were drawn in the follicular phase of the menstrual cycle; second samples were collected in the luteal phase. Samples were returned to our laboratory via overnight courier, with a frozen water sample to keep them cool. To time samples within the menstrual cycle, a postcard was included in the blood kit on which nurses indicated the first day of their next menstrual cycle after blood collection. Of the 29 611 participants, 18 521 provided

detailed information on the onset of their menstrual cycle, and 11 090 more women provided single, untimed blood samples. A brief questionnaire was included with the blood kit, asking the specific date and time when blood samples were drawn, the first day of the nurse's current menstrual cycle, the number of hours since she had last eaten, her current weight and medication use, and any changes in her menstrual cycle characteristics. For women who gave both follicular and luteal samples, we used luteal samples in this study, because cyclic variations of IGF are only modest (Juul *et al.* 1997, Helle *et al.* 1998).

Cases in this analysis are women with no cancer diagnosis (with the exception of non-melanoma skin cancer) prior to blood collection and with breast cancer diagnosis between the date of blood collection and June 1 2003. In all, 317 cases of breast cancer ($n = 80$ *in situ*) were confirmed by medical record review (two controls selected before 2001 became cases in 2003, we kept them as controls only, leaving 317 cases of the initial 319 cases). Although the 317 cases included 11 women whose pathology reports have not yet been obtained, we based our analyses on the total, because the accuracy of self-reporting was extremely high (Rockhill *et al.* 1998). Estrogen receptor (ER) status was also abstracted from medical records (168 of the invasive cancers were ER+). To each case, we matched two controls on year of birth, menopausal status at blood draw and at diagnosis of breast cancer (postmenopausal vs premenopausal vs unknown), time of day and month of blood draw, fasting status at blood draw (≥ 8 h since a meal vs not), luteal day (number of days before start of next cycle), and ethnicity (African–American, Asian, Hispanic, Southern European/Scandinavian/other Caucasian, and other). In sum, a total of 317 cases and 634 controls formed the study population for the current analyses. The study was approved by the Committee on the Use of Human Subjects in Research at the Brigham and Women's Hospital and the Harvard School of Public Health.

We defined menopausal status at the time of blood collection in our main analyses. Women who provided a timed sample were considered to be premenopausal. Women providing a random sample were considered premenopausal if they (a) reported that periods had not ceased or (b) had a hysterectomy but had at least one ovary remaining and were ≤ 45 (for nonsmokers) or ≤ 47 (for smokers) years of age – at these ages fewer than 10% of the cohort had had a natural menopause. Women were

considered postmenopausal if they (a) reported that natural menstrual periods had ceased permanently, (b) had a bilateral oophorectomy, or (c) had a hysterectomy but had at least one ovary remaining, and were at least 56 (for nonsmokers) or 54 (for smokers) years of age – by these ages natural menopause had occurred in 90% of the cohort. All other women were considered to be of unknown menopausal status (<1%). Of the 239 women who were premenopausal at blood collection, all but 14 were still premenopausal at diagnosis.

For IGFBP-1 analyses, only fasting blood samples were used. Total IGF-I, IGFBP-1, IGFBP-3, GH, and C-peptide levels were assayed by ELISA after acid extraction, using reagents from Diagnostic Systems Laboratory (DSL, Webster, TX, USA). To test whether the IGFBP-3 assay itself could account for variability in the results of associations between IGFBP-3 and breast cancer risk, IGFBP-3 levels for a portion of the data set (cases up to 2001 and their matched controls) were also assayed in Professor Holly's laboratory in Bristol, UK by a previously validated RIA, using an in-house polyclonal antibody and calibrated against recombinant human glycosylated IGFBP-3 (Cheetham *et al.* 1998). For a comparison of the ELISA results with this RIA see Table 2. As they were highly correlated, we used results obtained from the ELISA assays for IGFBP-3 in our final analyses. For that data set we also had IGF-I and IGFBP-3 levels assayed with an alternative ELISA, using the chemiluminescent immunoassay system (Immulate, Diagnostic Products Coorporation, CA, USA). When plotting the differences against the averages of the two measurements (ELISA versus alternative ELISA), the two different methods demonstrated good agreement between the two assays (Spearman $r = 0.94$) for IGF-I. For IGFBP-3, the plot suggested that the variation of at least one ELISA measurement depended on the magnitude of the measurements (Spearman $r = 0.90$). We therefore compared the standard DSL ELISA with IGFBP-3 measurements obtained by RIA, which is currently considered the gold standard for measuring IGFBP-3. The Bland–Altman plot (Bland & Altman, 1986) demonstrated good agreement between these two measurements. Moreover, IGFBP-3 RIA measurements were highly correlated with those obtained by standard ELISA (see Table 2) and will therefore not be used for this report. All samples were masked to case or control status. Masked split specimens included within each batch were used to calculate the coefficient of variation within batches; for IGF-I

these were 6.8%, for IGFBP-3 4.2%, for IGFBP-1 1.6%, for GH 11.3%, and for C-peptide 4.3%.

Evidence that a single plasma measurement of IGF-I and IGFBP-3 reflects longer term circulating levels is limited, but correlations ranged from 0.94 to 0.97 for samples measured over 8 weeks (Goodman-Gruen & Barrett-Connor 1997), 0.81 for IGF-I and 0.60 for IGFBP-3 measured over 1 year (Muti *et al.* 2002), suggesting that a single measure reflects average levels over at least a 1-year period. We also found good reproducibility of IGF-I and IGFBP-3 in NHS II (Missmer *et al.* 2006) over time (over 3 years, premenopausal women only, $r = 0.83$ for IGF-I, and intra-class correlation (ICC) $= 0.76$ for IGFBP-3). This level of reproducibility is similar for other biological variables, such as blood pressure and serum cholesterol measurements (ICC, 0.6–0.8 over several years), parameters considered reasonably well measured and consistent predictors of disease in epidemiological studies. In the NHS, we had conducted similar analyses for GH, with reproducibility over 3 years (ICC $= 0.44$).

Statistical analyses

We identified statistical outliers based on the generalized extreme studentized deviate (ESD) many-outlier detection approach (Rosner 1983); two women with improbable IGFBP-1 concentrations were identified as outliers and excluded from analyses that included IGFBP-1. Overall, we had fewer women with IGFBP-1 levels (191 cases and 378 controls) available for our analyses because the set of women who had provided fasting blood samples was smaller. To test for differences in hormone levels between cases and controls, we used mixed-effects regression models for clustered data to adjust for possible confounding due to the matching factors and for any residual correlation between case and control subjects within the matched set (Zeger *et al.* 1988). Quartiles of IGF levels were defined on the basis of plasma levels of all controls for the overall analyses and of IGF levels of all premenopausal controls for the analyses restricted to various subgroups of premenopausal women. In sub-analyses, to better classify individuals with respect to their individual insulin levels, we stratified by the plasma C-peptide median (using C-peptide as a marker of insulin production). As more than 30% of all GH values, which we analyzed in two batches, were below the detection limit of the GH assay (0.14 ng/ml and 0.21 ng/ml

respectively), all women with values ≤ 0.21 ng/ml constituted the reference group; for the remaining women tertiles were defined according to plasma levels of all controls, for a total of four categories.

To estimate the relative risks (odds ratios, ORs) and 95% confidence intervals (95% CIs), we used conditional logistic regression models, adjusting for the matching factors. In sub-analyses, we additionally adjusted for other breast cancer risk factors (age at menarche (less than 12 years, 12 years, 13 years, and 14+ years), parity (0, 1–2 births, 3–4 births, more than 4 births), age at birth of first child (0, less than 25 years, 25–29 years, 30+ years), family history of breast cancer (in mother or sister, yes/no), and body mass index (BMI) (<21 kg/m^2, 21–22.9 kg/m^2, 23–24.9 kg/m^2, 25–28.9 kg/m^2, and 29+ kg/m^2)). We tested for trends by calculating the Wald statistics for a continuous distribution of IGF values, using the square-root of IGF-I because of its skewed distribution. For analyses stratifying on number of years diagnosed after blood collection, we also estimated ORs for continuous measures of IGF-I and IGFBP-3 transformed on the log$_2$ scale [$\log_2 x = \log(x)/\log(2)$].

In contrast to our previous publication (Hankinson et al. 1998), we included in situ breast cancer cases in the current analyses, since our results, which overall are similar to our recent update in the NHS (Schernhammer et al. 2005) remained essentially unchanged after exclusion of in situ cases (but not their controls).

Results

Table 1 shows baseline characteristics of the 317 cases and 634 controls, by subgroups. The mean time between blood collection and diagnosis was 31 months (S.D. 20) with a range of 1–88 months.

Table 1 Baseline characteristics

	Cases	Controls
All women	$n = 317$	$n = 634$
‡Age (mean (S.D.))	45.3 (4.3)	45.1 (4.3)
Age at menarche (mean (S.D.))	12.4 (1.4)	12.4 (1.4)
Parity* (mean (S.D.))	2.2 (0.8)	2.3 (1.0)
Family history of breast cancer (%)	16.4	10.4
Oral contraceptive use† (%)	2.8	2.2
BMI (mean (S.D.))	25.3 (5.3)	25.7 (6.0)
Premenopausal women	$n = 239$	$n = 478$
‡Age (mean (S.D.))	44.1 (4.0)	43.8 (3.9)
Age at menarche (mean (S.D.))	12.5 (1.4)	12.4 (1.4)
Parity* (mean (S.D.))	2.1 (0.8)	2.3 (1.0)
Family history of breast cancer (%)	15.5	9.8
Oral contraceptive use† (%)	2.9	2.3
BMI (mean (S.D.))	24.9 (5.0)	25.1 (5.4)
Premenopausal women age $\leq$45‡	$n = 129$	$n = 275$
‡Age (mean (S.D.))	41.1 (2.8)	41.1 (2.8)
Age at menarche (mean (S.D.))	12.4 (1.5)	12.4 (1.4)
Parity* (mean (S.D.))	2.1 (0.8)	2.3 (0.9)
Family history of breast cancer (%)	13.2	8.0
Oral contraceptive use† (%)	3.9	1.8
BMI (mean (S.D.))	24.8 (5.7)	24.7 (5.0)

* Among parous women only. † Current (1997). ‡ Age at blood draw.

Total IGF-I and IGFBP-3 were positively correlated with each other, whereas IGF-I and age were inversely correlated, whereas height was not correlated with either IGF-I or GH (Table 2). IGFBP-1 was weakly and inversely correlated with both IGF-I and IGFBP-3 and weakly positively associated with GH. The correlation between GH and IGF-I, finally, was $r = -0.09$ ($P = 0.03$). The two measures for IGF-I and IGFBP-3 that we had available for our analyses in a portion of the data set (ELISA versus alternative ELISA) were well correlated with each other (IGF-I: Spearman $r = 0.94$, $P < 0.001$; IGFBP-3: $r = 0.87$, $P < 0.001$).

Table 2 GH and IGFs correlation matrix, all women combined (controls only)

	Spearman correlation coefficient (r) between the untransformed continuous measures of GH and various IGFs						
	GH	IGF-I	IGFBP-1	IGFBP-3 (ELISA)	IGFBP-3 (RIA)	Age	Height
GH	1.0	−0.09	0.16	−0.02	−0.06	−0.0006	−0.01
IGF-I		1.0	−0.16	0.51	0.39	−0.25	0.05
IGFBP-1			1.0	−0.16	−0.17	0.19	−0.03
IGFBP-3 (ELISA)				1.0	0.85	−0.13	0.01
IGFBP-3 (RIA)					1.0	−0.02	0.01
Age						1.0	−0.06
Height							1.0

Table 3 Plasma GH, IGF-I, IGFBP-1 and IGFBP-3 concentrations by case or control status

	Cases		Controls		
	n	Median (range*)	*n*	Median (range*)	*P*†
IGF-I (ng/ml)					
All cases or controls	317	230 (153–346)	634	239 (135–341)	0.83
Premenopausal	239	242 (164–352)	478	249 (150–350)	0.85
Premenopausal, age ≤45	129	260 (167–367)	275	258 (151–350)	0.40
IGFBP-3 (ng/ml) (ELISA)					
All cases or controls	317	4864 (4068–5967)	634	4881 (4000–5911)	0.72
Premenopausal	239	4918 (4137–5967)	478	4936 (4000–6030)	0.81
Premenopausal, age ≤45	129	5139 (4267–6002)	275	5021 (4092–5929)	0.27
IGFBP-1 (ng/ml)					
All cases or controls	191	38.7 (10.6–73.1)	378	35.3 (10.3–75.8)	0.55
Premenopausal	129	36.4 (11.1–66.8)	255	33.6 (11.4–67.9)	0.61
GH (ng/ml)					
All cases or controls	317	0.24 (0.14–5.27)	634	0.25 (0.14–4.03)	0.45
Premenopausal	239	0.23 (0.14–5.39)	478	0.24 (0.14–4.10)	0.47
Premenopausal, age ≤45	129	0.23 (0.14–5.39)	275	0.21 (0.14–4.03)	0.43

* Range of plasma values, 10th to 90th percentile.
† *P*-values for comparison of mean natural IGF plasma levels between cases and controls, based on mixed-effects regression models with adjustment for the matching variables.

Neither total IGF-I, nor IGFBP-1, IGFBP-3, or GH levels varied significantly between cases and controls in the whole cohort, nor did they vary among premenopausal women or premenopausal women of age 45 years or less (Table 3).

Throughout the study population, circulating IGF-I levels were not associated with breast cancer risk; nor did additional adjustment for IGFBP-3 or breast cancer risk factors such as BMI and family history of breast cancer alter these estimates (Table 4). The risks remained essentially unchanged after the exclusion of *in situ* breast cancer cases (relative risk (RR), top vs bottom quartile, 1.03; 95% CI, 0.69–1.54), as they did after exclusion of current or recent users of either hormone-replacement therapy or oral contraceptives (OC) (RR, top vs bottom quartile, 1.01; 95% and 5% CI, 0.68–1.51). Moreover, there was no marked difference in risks stratified along the median (34 months) of duration of OC use, as assessed in 1997 (≥34 months of OC use: RR, top vs bottom quartile, 0.81; 95% CI, 0.51–1.30; <34 months of OC use: RR, top vs bottom quartile, 1.20; 95% CI, 0.62–2.32).

IGFBP-3 and IGFBP-1 were similarly not associated with breast cancer risk and further adjustment for IGF-I did not alter these estimates (Table 4). For GH, the relative risk, comparing top and bottom quartiles was 1.09 (95% CI, 0.82–1.46).

In sub-analyses, when we restricted our data set to women who were premenopausal at blood collection (Table 4) or to premenopausal women of age 45 years or less at blood collection, the lack of an association between IGF-I and breast cancer risk remained essentially unchanged. Furthermore, among invasive cases, we observed no association of plasma IGF levels stratified by hormone receptor status of the tumor: IGF-I was not associated with either ER-negative (RR, top vs bottom tertile, 1.25, 95% CI, 0.49 to 3.18) or ER-positive tumors (RR, top vs bottom tertile, 1.14, 95% CI, 0.70 to 1.85) among women who were premenopausal at blood collection. A previous report (Bruning *et al.* 1995) indicated decreased IGFBP-3 levels in early-stage premenopausal breast cancer. We therefore evaluated the association between circulating IGFBP-3 levels and breast cancer risk in smaller tumors only (tumor size at diagnosis ≤2 cm), but could not confirm those findings (RR top vs bottom IGFBP-3 quartile, 1.08, 95% CI, 0.66 to 1.77).

To further explore differences in the relationship between IGF-I and IGFBP-3 and breast cancer risk depending on the interval between blood donation and tumor diagnosis, we stratified on the number of years since blood collection and evaluated the RR associated with a doubling of levels (i.e. a unit increase on the $\log_2$ scale). The risks

Table 4 Relative risk‡ of breast cancer by plasma IGF and GH quartiles, 1996–2003 for all women combined

	Q1	Q2	Q3	Q4	*P* for trend
IGF-I					
Cases/controls	78/157	90/160	70/159	79/158	
IGF-I	1.0	1.09 (0.80–1.47)	0.92 (0.67–1.27)	1.00 (0.73–1.37)	0.77
IGF-I†	1.0	1.07 (0.79–1.46)	0.91 (0.65–1.27)	0.98 (0.69–1.39)	0.77
IGF-I¶	1.0	1.14 (0.83–1.56)	0.92 (0.65–1.29)	0.96 (0.67–1.37)	0.91
IGFBP-3 (ELISA)					
IGFBP-3	1.0	1.04 (0.76–1.41)	0.92 (0.67–1.27)	1.07 (0.79–1.45)	0.76
IGFBP-3†	1.0	1.05 (0.77–1.45)	0.94 (0.67–1.32)	1.10 (0.78–1.54)	0.64
IGFBP-1					
IGFBP-1‡	1.0	0.73 (0.47–1.14)	1.16 (0.80–1.69)	0.95 (0.63–1.41)	0.50
GH					
Cases/controls	145/284	58/115	46/119	68/116	
GH‡	1.0	0.99 (0.73–1.35)	0.83 (0.59–1.15)	1.09 (0.82–1.46)	0.51
Premenopausal women					
IGF-I					
Cases/controls	47/94	69/119	53/130	70/135	
IGF-I	1.0	1.10 (0.76–1.60)	0.87 (0.59–1.29)	1.02 (0.71–1.48)	0.87
IGF-I†	1.0	1.07 (0.74–1.56)	0.83 (0.55–1.25)	0.94 (0.63–1.42)	0.57
IGF-I¶	1.0	1.13 (0.77–1.65)	0.83 (0.55–1.25)	0.92 (0.61–1.41)	0.48
IGFBP-3 (ELISA)					
IGFBP-3	1.0	1.07 (0.74–1.54)	0.98 (0.67–1.42)	1.17 (0.82–1.66)	0.90
IGFBP-3†	1.0	1.09 (0.75–1.58)	1.02 (0.68–1.51)	1.23 (0.83–1.82)	0.79
IGFBP-1					
IGFBP-1‡	1.0	0.81 (0.48–1.37)	1.07 (0.66–1.73)	0.68 (0.37–1.24)	0.60
GH					
Cases/controls	112/223	41/76	33/89	53/90	
GH‡	1.0	1.05 (0.73–1.50)	0.81 (0.55–1.19)	1.11 (0.80–1.54)	0.50
Premenopausal women age ≤45					
IGF-I					
Cases/controls	21/45	25/57	34/87	49/86	
IGF-I	1.0	0.96 (0.54–1.71)	0.88 (0.51–1.52)	1.14 (0.68–1.90)	0.70
IGF-I†	1.0	0.89 (0.49–1.60)	0.80 (0.45–1.40)	0.97 (0.55–1.70)	0.85
IGF-I¶	1.0	0.85 (0.46–1.56)	0.76 (0.43–1.37)	0.90 (0.51–1.62)	0.85
IGFBP-3 (ELISA)					
IGFBP-3	1.0	1.25 (0.71–2.19)	1.30 (0.76–2.22)	1.49 (0.88–2.51)	0.36
IGFBP-3†	1.0	1.27 (0.72–2.25)	1.31 (0.75–2.27)	1.49 (0.84–2.65)	0.49
GH					
Cases/controls	60/137	25/36	15/51	29/51	
GH‡	1.0	1.35 (0.84–2.15)	0.75 (0.42–1.32)	1.19 (0.76–1.85)	0.54

† Relative risks were, in addition to matching variables, further adjusted for plasma IGF-I or IGFBP-3.
¶ Relative risks were, in addition to matching variables and IGF-I or IGFBP-3, further adjusted for the following breast cancer risk factors: age at menarche (less than 12 years, 12 years, 13 years, and 14+ years), parity (0, 1–2 births, 3–4 births, more than 4 births), age at birth of first child (0, less than 25 years, 25–29 years, 30+ years), family history of breast cancer (yes/no), and BMI ($<21\,kg/m^2$, 21–$22.9\,kg/m^2$, 23–$24.9\,kg/m^2$, 25–$28.9\,kg/m^2$, and $29+\,kg/m^2$).
‡ Analyses based on conditional logistic regression models.

were independent of time since collection (women with a diagnosis of breast cancer within 2 years following blood collection: IGF-I, RR, 0.99, 95% CI, 0.69–1.43; IGFBP-3, RR, 1.09, 95% CI, 0.49–2.42; women with a diagnosis of breast cancer more than two years after blood collection: IGF-I, RR, 1.06, 95% CI, 0.74–1.53; IGFBP-3, RR, 1.24, 95% CI, 0.56–2.74).

Finally, to explore the possible influence of insulin resistance on the associations (using C-peptide as a

marker of insulin production), we stratified our data by the median C-peptide level. We would expect women with lower C-peptide levels (i.e. not resistant to the effects of higher IGF-I levels) to be at higher risk of breast cancer due to IGF-I than those with higher C-peptide levels. However, we were unable to confirm this hypothesis: in unconditional logistic regression models (adjusting for the matching factors and additional breast cancer risk factors), we observed an RR of 1.32 (95% CI, 0.57–3.05) associated with the top quartile of IGF-I (compared with the bottom quartile of IGF-I) in the stratum of women with C-peptide levels above the median compared with 0.83 (95% CI, 0.35–1.93) in women with C-peptide levels below the median.

Discussion

In a matched, nested case-control study of breast cancer, we found no association between IGF levels and premenopausal breast cancer risk and no association between plasma GH, IGFBP-1, or IGFBP-3 and breast cancer risk.

Persuasive basic science suggests that the proliferative effects of IGF-I influence both normal and transformed breast epithelial cells (Bates *et al.* 1995, Yang *et al.* 1996, Ng *et al.* 1997) and increase the frequency of breast tumors (Medical Intelligence Unit 2003). The expression of IGFBP-3 in many tissues, on the other hand, suggests that it locally modulates the action of IGF peptides. IGFBP-3 may have other, not yet fully understood, physiological roles. Finally, IGFBP-1, another binding protein, also regulates IGF actions (Medical Intelligence Unit 2003). GH, a hormone produced by the pituitary gland, is the primary regulator of hepatic IGF-I production, which, in turn, is the main source of circulating IGF-I (Jones & Clemmons 1995).

To date, seven prospective analyses have evaluated associations between IGFs and premenopausal breast cancer risk (Hankinson *et al.* 1998, Toniolo *et al.* 2000, Kaaks *et al.* 2002, Krajcik *et al.* 2002, Muti *et al.* 2002, Allen *et al.* 2005, Rinaldi 2005, Schernhammer *et al.* 2005). Three nested case-control studies among premenopausal and post-menopausal women (Toniolo *et al.* 2000, Krajcik *et al.* 2002, Muti *et al.* 2002, Allen *et al.* 2005) reported a positive association among premeno-pausal women. In all three, women with high premenopausal levels of IGF-I had a two- to three-fold higher risk of breast cancer, while levels in postmenopausal women were unrelated to risk. Recently published extensions of two of these

studies found generally similar results (Rinaldi *et al.* 2005, Schernhammer *et al.* 2005). In the largest prior prospective evaluation, data were analyzed from a total of 513 incident breast cancer cases and 987 matched controls, a study nested within two large Swedish cohorts (Kaaks *et al.* 2002). In contrast to the previous studies, the authors observed no noteworthy associations between plasma IGF levels and breast cancer risk among the 116 pre-menopausal cases and 330 controls (RR, top vs bottom quartile, 0.6, 95% CI, 0.3–1.4).

Many studies have investigated the relationship between IGFBP-3 and breast cancer risk. The first of the prospective studies published (Toniolo *et al.* 2000) noted no significant association for IGFBP-3 among either premenopausal or postmenopausal women. Subsequent cohort studies tended to confirm these findings in premenopausal women, with one exception (Muti *et al.* 2002), and with more inconsistent reports about the association between circulating IGFBP-3 levels and breast cancer risk among postmenopausal women (Kaaks *et al.* 2002, Keinan-Boker *et al.* 2002, Krajcik *et al.* 2002, Muti *et al.* 2002).

A few prospective studies also examined associations of IGFBP-1 with breast cancer risk (Kaaks *et al.* 2002, Krajcik *et al.* 2002, Schernhammer *et al.* 2005). All three studies, including our own results from the NHS cohort (Schernhammer *et al.* 2005), report no association of IGFBP-1 with breast cancer. The three retrospective studies that examined the associations of IGFBP-1 with breast cancer (Favoni *et al.* 1995, Del Giudice *et al.* 1998, Goodwin *et al.* 2002) overall confirm this absence of an important association. To our knowledge, no previous study has evaluated associations of circulating GH levels with breast cancer risk. Previous studies report strong correlations between GH and IGF-I, particularly *in utero* (r = 0.42) (Chellakooty *et al.* 2004). Although the reproducibility of GH appeared to be reasonable in a comparable data set (the NHS, ICC = 0.44), it was still lower than for IGF-I, which may explain the lack of correlation between GH and IGF-I and between circulating GH levels and breast cancer risk in this study. More-over, the quickly changing, pulsatile excretion pattern of GH makes this hormone difficult to measure, and our results must, therefore, be interpreted with great caution.

More recent studies, including our own update (Schernhammer *et al.* 2005) of previously published data (Hankinson *et al.* 1998) and updated data from the New York University Women's Health Study

cohort (Rinaldi *et al*. 2005), have observed weaker associations between circulating IGF-I levels and breast cancer risk among premenopausal women than initially reported. Similarly, the findings from our current study of primarily premenopausal women show no association at all. What could account for these puzzling differences? The most likely explanations are either methodological issues or secular changes over time, particularly if linked to a woman's hormonal status, given the lack of similar changes in results over time with colon cancer (Giovannucci *et al*. 2000, Wei *et al*. 2005).

With our two data sets (NHS and NHS II) at hand, we therefore considered methodological issues comparing the two cohorts, and evaluated a variety of hormone-related breast cancer risk factors that could account for the discrepancies, potentially reflecting a secular change in lifestyle factors.

First, blood sampling, transportation, and storage methods are identical between the two cohorts. Technical issues related to the assays used for IGF-I measurements, although they admittedly exist (Rinaldi *et al*. 2005), also appear to be an unlikely explanation, considering the high correlation between two different ELISA assays for IGF-I in our study. In addition, given that the positive association between IGF-I and colon cancer risk in the NHS cohort remained in a more recent update (Wei *et al*. 2005), whatever affects the changes seen in risk may only relate to breast cancer. We therefore compared some of the baseline breast cancer risk factors between the two cohorts but were unable to detect important differences. For example, the average body mass index of premenopausal women was comparable between cohorts (mean, NHS: cases 25.4; controls 25.7; NHS II: cases 24.9; controls 25.1), as were IGF-I and IGFBP-3 ranges, although levels of both IGF-I and IGFBP-3 were slightly higher in the NHS II (as expected in this younger group) than in the NHS. Correlations between IGF-I and IGFBP-3 (NHS: $r = 0.57$ vs NHS II; $r = 0.51$) and IGF-I and age (NHS: $r = -0.33$ vs NHS II, $r = -0.25$) were also comparable.

With no obvious explanation at hand, there is room for speculation. One obvious hypothesis to explain the discrepancies in IGF studies conducted over the past decade is that circulating IGF-I levels, as measured in epidemiological studies, do not accurately reflect long-term IGF-I levels. However, while no studies have reported the ICC of IGF-I measures taken many years apart, studies have shown a reasonable ICC for shorter intervals.

Also, there is no reason to believe that the misclassification resulting from the use of a single IGF measure would vary substantially between studies. Alternatively, one could hypothesize that, in the past, there were environmental interactions with IGF levels that vanished over time. However, all studies on IGF-I and breast cancer risk collected their blood samples at various time points, and it would be hard (if not impossible) to pin such an event to a certain period. Another theory might support the relevance of much earlier exposures to IGF-I (i.e. in adolescence); it is conceivable that, while adult levels (particularly when measured before menopause) may have reflected adolescence levels in the past, they no longer do so because of additional, differential exposure to IGF-I due to dietary milk (Holmes *et al*. 2002) or other unknown influences. However, we measured IGF earlier in a woman's life than in our previous study (Hankinson *et al*. 1998).

Speculations about a secular trend in nutritional exposures, which may have led not only to growing obesity rates but also to an increase in insulin resistance, thereby possibly negating the impact of IGF-I, could not be corroborated in our analyses stratified on C-peptide levels as a marker for insulin resistance.

Another mechanistic interpretation (Pollak *et al*. 2004) of previous studies showing a higher breast cancer risk to be associated with higher IGF-I levels among premenopausal women was that higher IGF-I levels are associated with higher levels of IGF-I receptor activation in at-risk mammary epithelial cells. This was postulated to increase survival of cells with accumulating DNA damage, which would facilitate stepwise carcinogenesis and/or lead to a higher proliferation rate of early cancers, as suggested by experimental data (Ng *et al*. 1997). However, it is plausible that signaling at and downstream of the IGF-I receptor is a function not only of ligand levels, but also of polymorphic variation in genes encoding key signaling proteins. Thus, risk may vary in a complex fashion involving interactions between polymorphic variation of genes encoding signaling molecules and ligand levels. Ongoing studies will explore the possibility that the relationship between circulating IGF-I levels and risk is confined to subpopulations that can be genetically identified, and that discrepancies between population studies may relate, in part, to differences in genotype distribution.

Our study is prospective and of fairly large size, adding to the strength of our findings. With a large

proportion of premenopausal women, we were able to address associations by menopausal status, with sufficiently large numbers of cases in each quartile. However, the youngest of our premenopausal women is more than 42 years old, which compromises the ability to assess risks among very young women even in this, to date one of the largest cohort studies among premenopausal women. Thus, larger cohorts of young women with a lower median age are needed to address whether IGF-I associations with breast cancer risk vary by age, and whether IGF levels in young women better reflect the exposure period of importance.

An advantage of our prospective design is that blood samples were collected before the occurrence of breast cancer, thus allowing us to evaluate circulating IGF-I levels as potential predictors for breast cancer rather than as tumor markers. However, with an average of only 31 months between blood collection and tumor diagnosis, the influence of a tumor that was not yet diagnosed cannot be completely ruled out, given the long latency period of breast cancer. We did exclude cases that occurred within the first year after blood collection, and breast cancer risks associated with IGF-I remained essentially unchanged; the modest increase of risk associated with IGFBP-3, however, suggests that studies with longer periods between blood collection and tumor diagnosis, particularly among premenopausal women, are needed to rule out a possible influence of a preclinical tumor on circulating IGF levels.

In summary, our study does not support findings from previous studies suggesting that the relation between IGF-I and breast cancer risk differs by menopausal status. While most but not all previous prospective studies reported a relatively strong, positive association between plasma IGF-I and breast cancer risk among premenopausal women, we could not corroborate these findings. Cohort effects may contribute complexity to these associations, which will need to be addressed in future cohort studies. Future studies will also need to further evaluate the usefulness of IGF-I as a prognostic marker for breast cancer.

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