Dietary Inflammatory and Insulinemic Potential and Risk of Type 2 Diabetes: Results From Three Prospective U.S. Cohort Studies

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To examine whether proinflammatory and hyperinsulinemic diets are associated with increased risk of type 2 diabetes.

RESEARCH DESIGN AND METHODS

We prospectively followed 74,767 women from the Nurses' Health Study (1984– 2016), 90,786 women from the Nurses'Health Study II (1989–2017), and 39,442 men from the Health Professionals Follow-up Study (1986–2016). Using repeated measures of food-frequency questionnaires, we calculated empirical dietary inflammatory pattern (EDIP) and empirical dietary index for hyperinsulinemia (EDIH) scores, which are food-based indices that characterize dietary inflammatory or insulinemic potential based on circulating biomarkers of inflammation or C-peptide. Diagnoses of type 2 diabetes were confirmed by validated supplementary questionnaires.

RESULTS

We documented 19,666 incident type 2 diabetes cases over 4.9 million person-years of follow-up. In the pooled multivariable-adjusted analyses, individuals in the highest EDIP or EDIH quintile had 3.11 times (95% CI 2.96–3.27) and 3.40 times (95% CI 3.23–3.58) higher type 2 diabetes risk, respectively, compared with those in the lowest quintile. Additional adjustment for BMI attenuated the associations (hazard ratio 1.95 [95% CI 1.85–2.05] for EDIP and hazard ratio 1.87 [95% CI 1.78–1.98] for EDIH), suggesting adiposity partly mediates the observed associations. Moreover, individuals in both highest EDIP and EDIH quintiles had 2.34 times higher type 2 diabetes risk (95% CI 2.17–2.52), compared with those in both lowest quintiles, after adjustment for BMI.

CONCLUSIONS

Higher dietary inflammatory and insulinemic potential were associated with increased type 2 diabetes incidence. Findings suggest that inflammation and hyperinsulinemia are potential mechanisms linking dietary patterns and type 2 diabetes development.

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Type 2 diabetes is a major cause of morbidity, and the Centers for Disease Control and Prevention ranked diabetes as the seventh leading cause of death in the U.S. in 2017 (1). The estimated prevalence of type 2 diabetes was 12–14% among U.S. adults in 2011–2012 (2), and it is projected to increase to 25–28% by 2050 (3). Type 2 diabetes is characterized by insulin resistance, impaired insulin secretion, and hyperglycemia (4). Moreover, evidence suggests that inflammation playsan important rolein the pathogenesis of insulin resistance and type 2 diabetes (5). It is well documented that major risk factors of type 2 diabetes such as obesity and physical inactivity upregulate inflammation and insulin resistance, which consequently cause hyperinsulinemia (6,7).

Diets modulating these biological pathways of inflammation and insulin response may influence type 2 diabetes incidence. Epidemiologic studies have found healthy dietary patterns indicated by the Alternate Healthy Eating Index (AHEI), alternate Mediterranean Diet (aMED), and Dietary Approaches to Stop Hypertension (DASH) to be inversely associated with risk of type 2 diabetes (8). Also, greater adherence to the prudent pattern and lower adherence to the Western pattern were associated with reduced risk of type 2 diabetes (8), and higher glycemic index (GI) and glycemic load (GL) have been found to be associated with higher risk of developing type 2 diabetes (9). Although these dietary patterns are reported to be associated with inflammation and insulin response (10–12), they may not comprehensively capture one's dietary inflammatory and insulinemic potential, which are important mediators linking diet and type 2 diabetes. Recently, we developed and validated an empirical food-based dietary inflammatory pattern (13) and an empirical food-based dietary index for hyperinsulinemia (14) to assess longterm inflammatory and insulinemic potential of usual diets. These empirical dietary indices showed low-to-moderate correlations with conventional dietary pattern scores (AHEI, aMED, and DASH) $(r = -0.09$ to $-0.45)$ (15). In this study, we investigated the association of dietary inflammatory and insulinemic potential with incidence of type 2 diabetes in three large prospective cohort studies. We also examined whether the association differs by other major risk factors of type 2 diabetes including BMI.

RESEARCH DESIGN AND METHODS

Study Population

The Nurses' Health Study (NHS) cohort was established in 1976 with 121,701 female nurses aged 30–55 years (16), the Nurses'Health Study II (NHSII) cohort was established in 1989 with 116,430 female nurses aged 25–42 years (17), and the Health Professionals Follow-up Study (HPFS) cohort was established in 1986 with 51,529 male health professionals aged 40–75 years (18). In these three cohorts, participants completed questionnaires on medical history and lifestyle at enrollment and every 2 years thereafter. The response rate of each follow-up questionnaire cycle exceeded .90% for all cohorts.

In the current study, we excluded participants who had diabetes, cardiovascular disease, or cancer (except nonmelanoma skin cancer) or who had incomplete dietary information or implausible energy intake ($<$ 600 or $>$ 3,500 kcal/day for women and $<$ 800 or $>$ 4,200 kcal/day for men) at baseline. The final sample included 74,767 women from the NHS, 90,786 women from the NHSII, and 39,442 men from the HPFS. The study protocol was approved by the institutional review boards of the Brigham and Women's Hospital and Harvard T.H. Chan School of Public Health.

Dietary Assessment

Diet was assessed using a validated semiquantitative food-frequency questionnaire (FFQ) (\sim 130 items) every 4 years, starting from 1984 in NHS, 1991 in NHSII, and 1986 in HPFS (19–22). Participants were asked to report how often, on average, they consumed each food item during the previous year, with a standard portion size, using nine response categories ranging from"never or less than once per month" to "six or more times per day."

Assessment of Empirical Dietary Inflammatory Pattern and Empirical Dietary Index for Hyperinsulinemia

Empirical dietary inflammatory pattern (EDIP) and empirical dietary index for hyperinsulinemia (EDIH) scores were developed and validated to capture overall inflammatory and insulinemic potential, respectively, of whole diets (13,14). Briefly, EDIP was derived based on 39 predefined food groups from FFQs using reduced-rank regression followed by stepwise linear regression models to identify a dietary pattern most predictive of three plasma inflammatory biomarkers including interleukin-6, C-reactive protein (CRP), and tumor necrosis factor (TNF) α receptor 2. Similarly, EDIH was derived based on 39 predefined food groups using stepwise regression models to identify a dietary pattern most predictive of fasting plasma C-peptide, an indicator of insulin secretion. Of note, EDIH was predictive of both fasting and nonfasting C-peptide in a previous study (23). EDIP and EDIH are weighted sums of 18 food groups (9 overlapping), and higher (more positive) scores indicate higher inflammatory or insulinemic potential of diets and lower (more negative) scores indicate lower inflammatory or insulinemic potential of diets [\(Supplementary Table 1](https://doi.org/10.2337/figshare.12739601)). Detailed distributions of nutrients and food groups are shown in [Supplementary Table 2.](https://doi.org/10.2337/figshare.12739601) In independent data sets, EDIP and EDIH were validated using biomarkers of inflammation and insulin response (13,14,24). For each participant, we calculated EDIP and EDIH scores using updated FFQ data in each 4-year cycle. We also calculated dietary insulin index (II) and insulin load (IL) for a secondary analysis. Details of II and IL are provided in the [Supplementary](https://doi.org/10.2337/figshare.12739601) [Material.](https://doi.org/10.2337/figshare.12739601)

Assessment of Covariates

Detailed information on age, race, height, body weight, smoking, and physical activity was collected at baseline and updated biennially. Other information such as menopausal status, postmenopausal hormone use, oral contraceptive use, and family history of diabetes was collected from the biennial questionnaires.

Ascertainment of Type 2 Diabetes

Self-report of type 2 diabetes was assessed via questionnaires every 2 years. Diagnoses were confirmed using a supplemental questionnaire, which asked about type 2 diabetes–related symptoms, medication use, and diagnostic tests. In accordance with the National Diabetes Data Group (25), confirmation of diabetes required at least one or more of the following criteria: 1) at least one classic symptom (excessive thirst, polyuria, weight loss, or hunger) plus fasting blood glucose \geq 140 mg/dL (7.8 mmol/L), or random blood glucose \geq 200 mg/dL (11.1 mmol/L); 2) no symptoms but elevated blood glucose on two occasions (fasting blood glucose \geq 140 mg/dL

[7.8 mmol/L] or random blood glucose \geq 200 mg/dL [11.1 mmol/L]), or blood glucose \geq 200 mg/dL after 2-h blood oral glucose tolerance testing); and 3) treatment with hypoglycemic drugs (insulin or oral hypoglycemic agent). The diagnostic criteria for fasting blood glucose was changed to \geq 126 mg/dL (7.0 mmol/L) in 1998 (26), and glycated hemoglobin $(HbA_{1c}) \geq 6.5\%$ was further added in the criteria in 2010 (27). This supplemental questionnaire was validated previously (28,29).

Statistical Analysis

Person-years were calculated from the baseline when EDIP and EDIH scores were first available (1984 for NHS, 1991 for NHSII, and 1986 for HPFS) until type 2 diabetes diagnosis, death, censoring, or the end of follow-up (June 2016 for NHS, June 2017 for NHSII, and January 2016 for HPFS). We used cumulative average of EDIP and EDIH scores calculated from repeated measure of FFQs to capture habitual long-term dietary intake and reduce within-person variation. For example, we calculated the average of 1986 and 1990 scores and then formed quintiles and applied this to 1990–1994 follow-up. Then, we calculated the average of 1986, 1990, and 1994 scores and then formed quintiles and applied this to 1994–1998 follow-up and so on. The scores were adjusted for total energy intake using the residual method (30).

We used Cox proportional hazards regression models to estimate hazard ratios (HRs) and 95% CI of type 2 diabetes associated with quintiles of EDIP and EDIH scores. We also evaluated the doseresponse relationship of EDIP and EDIH with risk of type 2 diabetes using restricted cubic splines. All analyses were stratified by age in months and calendar years. Multivariable models additionally included race, smoking, postmenopausal hormone use, oral contraceptive use, physical activity, and family history of diabetes. We ran an additional multivariable model further adjusting for BMI. To examine the independent association of each empirical dietary index, we conducted a model further mutually adjusting for EDIP and EDIH. As a secondary analysis, we examined the association between other insulin-related indices (II and IL) and type 2 diabetes risk. All analyses were done separately by cohorts (sex), and results were pooled after testing for heterogeneity ($P > 0.05$). Pooled analyses were additionally stratified by cohorts.

To examine whether the association between empirical hypothesis-oriented dietary indices and type 2 diabetes differs by potential effect modifiers, we conducted subgroup analyses by age, BMI, physical activity, smoking status, alcohol intake, and family history of diabetes. The Wald test was used to test for interaction between the empirical dietary indices (continuous) and stratification variables. Lastly, we examined joint associations of EDIP (quintiles) and EDIH (quintiles) and of each empirical hypothesis-oriented index (quintiles) and BMI (five categories) with type 2 diabetes risk. We conducted a sensitivity analysis restricted to symptomatic diabetes cases to address potential bias due to screening. All statistical analyses were performed using SAS 9.4 with two-sided tests, and P value $<$ 0.05 was considered statistically significant.

RESULTS

Participants with higher EDIP or EDIH scores had higher BMI and lower physical activity levels (Table 1). Moreover, women with higher EDIP or EDIH scores were more likely to have family history of diabetes. The Spearman correlation between EDIP and EDIH was 0.65 ([Sup](https://doi.org/10.2337/figshare.12739601)[plementary Table 3](https://doi.org/10.2337/figshare.12739601)).

During 4,949,265 person-years of followup, we documented 19,666 incident type 2 diabetes cases. Diets with higher inflammatory potential were significantly associated with higher type 2 diabetes incidence in all three cohorts (Table 2). In the pooled multivariable-adjusted analyses, compared with individuals in the lowest EDIP quintile, those in the highest EDIP quintile had 3.11 times higher risk of type 2 diabetes (95% CI 2.96–3.27). Additional adjustment for BMI attenuated the magnitude of the association, but the strong positive association remained statistically significant (HR 1.95 [95% CI 1.85–2.05]). Similarly, diets with higher insulinemic potential were associated with increased risk of type 2 diabetes in all three cohorts (Table 2). In the pooled multivariable-adjusted analyses, compared with individuals in the lowest EDIH quintile, those in the highest EDIH quintile had 3.40 times higher risk of type 2 diabetes (95% CI 3.23–3.58). When we further adjusted for BMI, we observed an attenuated but strong positive association (HR 1.87 [95% CI 1.78– 1.98]). In a mutually adjusted model including both EDIP and EDIH, we still observed significant positive associations for both EDIP (HR 1.59 [95% CI 1.49– 1.69]) and EDIH (HR 1.42 [95% CI 1.33– 1.51]) ([Supplementary Table 4](https://doi.org/10.2337/figshare.12739601)). Moreover, additional adjustment for AHEI did not change the results, and the correlations of AHEI were -0.27 for EDIP and -0.47 for EDIH [\(Supplementary Tables 3](https://doi.org/10.2337/figshare.12739601) and [4](https://doi.org/10.2337/figshare.12739601)). Dose-response analyses showed that the associations of EDIP and EDIH with type 2 diabetes deviate from linearity with an accelerated increase in risk at higher scores (P for curvature $<$ 0.001 for both scores) [\(Supplementary Fig. 1\)](https://doi.org/10.2337/figshare.12739601). Restricting the analyses to symptomatic diabetes cases did not change the results (data not shown).

Stratified analyses showed significant positive associations of EDIP and EDIH with type 2 diabetes incidence, regardless of the predefined subgroups (Table 3). However, we observed a stronger positive association between EDIP and type 2 diabetes risk among younger, leaner, or more active adults or those without family history of diabetes ($P_{interaction}$ <0.05 for all). Moreover, the positive association between EDIH and type 2 diabetes risk was stronger among younger, leaner, or more active adults, never smokers, or moderate drinkers ($P_{\text{interaction}}$ < 0.05 for all). In the joint analysis of EDIP and EDIH, individuals in both highest EDIP and EDIH quintiles had 4.62 times (95% CI 4.29–4.97) increased risk of type 2 diabetes, compared with those in both lowest quintiles [\(Supplementary](https://doi.org/10.2337/figshare.12739601) [Fig. 2\)](https://doi.org/10.2337/figshare.12739601). Additional adjustment for BMI attenuated the association, which remained significant (HR 2.34 [95% CI 2.17– 2.52]) (Fig. 1). When we examined the joint association of each empirical hypothesis-oriented dietary index and BMI with type 2 diabetes risk, obese individuals in the highest EDIP or EDIH quintile had 21.8–42.6 times higher risk of type 2 diabetes compared with lean individuals in the lowest quintile ([Supplementary](https://doi.org/10.2337/figshare.12739601) [Figs. 3](https://doi.org/10.2337/figshare.12739601) and [4\)](https://doi.org/10.2337/figshare.12739601).

In a supplementary analysis using II and IL, we observed weak correlations of II and IL with EDIP (0.14 for II and 0.16 for IL) and EDIH (-0.03 for II and -0.02 for IL) ([Supplementary Table 3\)](https://doi.org/10.2337/figshare.12739601). Higher II or IL was inversely associated with risk of type 2 diabetes [\(Supplementary Tables 5](https://doi.org/10.2337/figshare.12739601) and [6\)](https://doi.org/10.2337/figshare.12739601). In the pooled multivariable-adjusted

Table 2—HR (95% CI) of type 2 diabetes according to quintiles of EDIP and EDIH*

Multivariable models adjusted for age (month), race (white or nonwhite), smoking (never, past, or current: 1-14, 15-24, or >24 cigarettes/day), postmenopausal hormone use (women only) (premenopausal, postmenopausal current user, or postmenopausal never/past user), oral contraceptive use (women only) (never, past, or current), physical activity (<3, 3–8.9, 9–17.9, 18–26.9, or \geq 27 MET h/week), and family history of diabetes (yes or no). Multivariable + BMI models additionally adjusted for BMI (<21, 21–22.9, 23–24.9, 25–26.9, 27–29.9, 30–34.9, 35–39.9, or ≥40 kg/m²). ref, reference. *Energy-adjusted dietary pattern. *†P_{trend} was calculated using continuous variables of dietary pattern in the model.*

analyses including BMI, individuals in the highest II or IL quintile had \sim 10% lower risk of type 2 diabetes compared with those in the lowest quintile.

CONCLUSIONS

In three large prospective cohorts, higher dietary inflammatory or insulinemic potential was associated with increased incidence of type 2 diabetes. Notably, the strong positive associations remained significant even after adjustment for BMI and mutual adjustment for the dietary indices including the conventional dietary score (i.e., AHEI). Individuals consuming diets with a highinflammatory and insulinemic potential had a 2.3-fold higher risk of developing type 2 diabetes

than those whose dietary patterns have a low inflammatory and insulinemic potential, after adjustment for BMI.

A number of studies have examined the association of various dietary patterns with incidence of type 2 diabetes. A recent meta-analysis of prospective cohort studies reported that a priori–defined "healthy" dietary patterns including AHEI,

All models were adjusted for age (month), race (White or non-White), smoking (never, past, or current: 1-14, 15-24, or $>$ 24 cigarettes/day), postmenopausal hormone use (women only) (premenopausal, postmenopausal current user, or postmenopausal never/past user), oral contraceptive use (women only) (never, past, or current), physical activity (<3, 3–8.9, 9–17.9, 18–26.9, or ≥27 MET h/week), family history of diabetes (yes or no), and BMI (<21, 21–22.9, 23–24.9, 25–26.9, 27–29.9, 30–34.9, 35–39.9, or \geq 40 kg/m²). ref, reference. *Energy-adjusted dietary pattern. $\dot{P}_{\text{interaction}}$ was calculated using the Wald test by including the interaction term.

aMED, and DASH were associated with 21%, 13%, and 19% decreased risk of type 2 diabetes, respectively, in comparisons of extreme quantiles (8). Similar reduction of type 2 diabetes risk (25– 35%) was shown with restriction to the same study population (i.e., NHS or HPFS) (8). Evidence indicates that the biological mechanisms by which these dietary patterns influence type 2 diabetes are closely related to inflammation and insulin response (10–12). However, studies have not comprehensively assessed dietary inflammatory or insulinemic potential and its relation with type 2 diabetes. The strong positive associations of EDIP and EDIH scores with type 2 diabetes risk remained robust even after adjustment for the conventional dietary pattern score (AHEI), suggesting unique and strong contributions of the empirical dietary indices in predicting type 2 diabetes risk. Previously, one cross-sectional study using a literature-derived nutrient-based index (Dietary Inflammatory Index [DII]) (31) found a positive association between dietary inflammatory potential and type 2 diabetes in 1,174 Mexican adults (32). Given that DII was calculated using both food items and nutrients (also influenced by supplement use), these results are not directly comparable with our findings using EDIP, which is based exclusively on

Figure 1—Joint association of EDIP and EDIH with risk of type 2 diabetes.HRs were calculated in Cox proportional hazards models after adjustment for age (month), race (White or non-White), smoking (never, past, or current: 1-14, 15-24, or >24 cigarettes/day), postmenopausal hormone use (women only) (premenopausal, postmenopausal current user, or postmenopausal never/past user), oral contraceptive use (women only) (never, past, or current), physical activity (<3, 3-8.9, 9-17.9, 18–26.9, or \geq 27 MET h/week), and family history of diabetes (yes or no) and BMI (<21.0, 21.0-22.9, 23.0-24.9, 25.0-26.9, 27.0-29.9, 30.0-34.9, 35.0-39.9, or \geq 40.0 kg/m²).

whole diets. In a previous validation study of 11,053 individuals, EDIP showed greater ability to predict inflammatory markers (i.e., CRP, TNF α receptor 2, and adiponectin) than DII (24). Moreover, the cross-sectional design of that study makes it difficult to assess a temporal relationship between diets and type 2 diabetes.

To our knowledge, studies have not directly examined the association of dietary insulinemic potential with incidence of type 2 diabetes. However, previous studies have used GI and GL as contributors to an immediate insulin response by assessing the influence of carbohydratecontaining foods on postprandial blood glucose (33). In a recent meta-analysis, participants in the highest categories of GI and GL had a 19% and a 16% higher risk of type 2 diabetes, respectively, compared with those in the lowest quintiles (9). Nonetheless, GI and GL have limited capacity to account for noncarbohydrate factors that may affect insulin response, and they do not incorporate effects of diet on insulin resistance, which is a major determinant of insulin response. As an extension of GI and GL, II and IL have been developed to directly quantify the postprandial insulin response to foods independent of insulin resistance (34). However, both GI and II did not predict fasting C-peptide, a marker of β -cell secretory activity (35,36). Moreover, II

and IL did not predict risk of colorectal and pancreatic cancers, which are strongly related with hyperinsulinemia and insulin resistance (37,38). Similarly, in our supplementary analyses, II and IL were associated not positively but inversely with risk of type 2 diabetes-a finding that aligns with previous studies showing that II may be inversely associated with long-term insulin exposure (14,23,35), suggesting that the capacity of foods to induce postprandial insulin secretion may have a role in reducing blood glucose and thus risk of type 2 diabetes.

Interestingly, although EDIP and EDIH were correlated, EDIP and EDIH were independently and jointly associated with type 2 diabetes risk, suggesting two distinct biological mechanisms linking diets and type 2 diabetes. Individuals consuming diets with both higher inflammatory and insulinemic potential had the highest risk of type 2 diabetes. A dietary pattern with higher inflammatory and insulinemic potential has higher intakes of red meat, processed meat, nondark (nonfatty) fish, sugar-sweetened (highenergy) beverages, and refined grains and low intakes of wine, leafy green vegetables, and coffee. Overall, our findings show that EDIP and EDIH may be more comprehensive dietary indices that better capture the biologically relevant aspects of diets for type 2 diabetes development.

Diet and adiposity have a complex relationship (39). The effect of diet can be mediated through adiposity. A previous study has shown that both EDIP and EDIH were associated with substantial longterm weight gain (40); thus, the models adjusting for BMI may highlight the adiposity-mediated influence of EDIP and EDIH on type 2 diabetes risk. As expected, we observed largely attenuated associations when we adjusted for BMI ($-37%$ for EDIP and $-45%$ for EDIH), but the strong positive associations remained. Our study shows that although the association of EDIP and EDIH with type 2 diabetes risk may be partly mediated through adiposity, these empirical hypothesis-oriented dietary indices have direct influences on type 2 diabetes, independent of adiposity. In addition, diet may interact with adiposity. Our findings from the joint analysis provide evidence that the combined influence of the dietary index and adiposity can be substantial. Obesity is a major risk factor for type 2 diabetes that is strongly linked to inflammation and insulin resistance. Thus, proinflammatory and/or hyperinsulinemic diets may exacerbate inflammatory and insulin-related pathological pathways, especially among obese individuals who may already be in a condition of increased inflammation and insulin insensitivity, thus leading to higher risk of developing type 2 diabetes.

Additionally, we observed significant interactions between the empirical hypothesis-oriented dietary indices and major risk factors of type 2 diabetes. EDIP showed a stronger positive association with type 2 diabetes risk in younger, leaner, or active adults or those without family history of diabetes. EDIH also showed a stronger positive association in younger or active adults or never smokers. Although the reasons for these significant interactions are not clear, these findings suggest that dietary inflammatory or insulinemic potential may be a stronger predictor of type 2 diabetes in lower risk groups. More research is needed to replicate our findings in diverse racial and ethnic groups. In addition, future studies incorporating genotype for various type 2 diabetes–associated single nucleotide polymorphisms could provide important insights to understand the interaction between individual genetic variants and proinflammatory/hyperinsulinemic diets in relation to type 2 diabetes. Also,

integration of these dietary indices with new data including biomarkers and metabolomics can provide opportunity to comprehensively elucidate the underlying biological mechanisms linking dietary patterns and type 2 diabetes development.

Our study has considerable strengths. First, we used validated food-based EDIP and EDIH scores that are strongly correlated with systemic inflammatory biomarkers and C-peptide, respectively, which are related to risk of type 2 diabetes. Second, we had a large sample size with $>$ 20 years of follow-up. The large number of type 2 diabetes cases allowed us to obtain precise estimates for stratified and joint analyses. Third, we had detailed prospectively collected information on diets and covariates, which minimizes potential confounding and recall bias. Fourth, repeated measures of diets allowed us to measure long-term effect of diets as well as reduce measurement error due to within-person variation. There are several limitations as well. Self-report diet and other covariates from questionnaires may have measurement errors. However, previous validation studies have shown reasonably good correlation between FFQ and diet records. Moreover, although we thoroughly adjusted for important potential confounding factors in our analyses, we cannot completely rule out the potential for residual confounding by unmeasured variables.

In conclusion, we found that higher dietary inflammatory or insulinemic potential was strongly associated with an increased incidence of type 2 diabetes in U.S. adults. Our study provides strong evidence that inflammation and hyperinsulinemia may mediate the association between diets and type 2 diabetes development. Dietary guidelines and interventions highlighting the importance of reducing or avoiding inflammatory and insulinemic dietary patterns may have great potential for the primary prevention of type 2 diabetes.

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