Supplemental Method 1. Gut microbiota sequencing protocol, Milieu Intérieur study, France, 2012.

## DNA extraction

DNA was extracted from stool samples using a Repeated Bead Beating Plus Column (RBB+C) method, as previously published (O'Cuiv et al. 2011; Yu et al. 2004). Briefly, cells lysis was achieved by bead beating in the presence of $4 \%(\mathrm{w} / \mathrm{v})$ sodium dodecyl sulfate (SDS), 500 mM NaCl , and 50 mM EDTA. After bead beating, precipitation with ammonium acetate was achieved to remove most of the impurities as well as the SDS; and the nucleic acids were recovered by precipitation with isopropanol. During the last step, RNA and proteins were removed using sequential digestions with RNase and proteinase K respectively, and genomic DNA was purified using QIAamp columns (Qiagen). Finally, DNA quality and quantity were checked using NanoDrop and Qubit together with Broad Range assay.

## Barcoding Polymerase Chain Reaction (PCR)

Barcoding PCR was carried out using indexed primers targeting the V3-V5 of the 16S rRNA gene as described by Kozich et al (Kozich et al. 2013) (forward primer: CCTACGGRAGGCAGCAG; reverse primer: GGACTACHVGGGTWTCTAAT). AccuPrime ${ }^{\text {TM }}$ Pfx SuperMix (Invitrogen - 12344-040) was used to perform the PCR. PCR mix was made up of $18 \mu \mathrm{~L}$ of AccuPrime ${ }^{\text {TM }}$ Pfx SuperMix, $0.5 \mu \mathrm{~L}$ of both V3-340F and V5-926R primers ( $0.2 \mu \mathrm{M}$ ) and $1 \mu \mathrm{~L}$ of DNA (10ng). PCR was carried out as follows: $95^{\circ} \mathrm{C}$ for $2 \mathrm{~min} ; 30$ cycles of $95^{\circ} \mathrm{C}$ for 20sec; $55^{\circ} \mathrm{C}$ for $15 \mathrm{sec} ; 72^{\circ} \mathrm{C}$ for 5 min ; and $72^{\circ} \mathrm{C}$ for 10 min . PCR reaction was quantified using Qubit BR DNA assay kit. Equal amounts of each PCR reaction were pooled and thoroughly mixed. The amplicon pool was then loaded on an agarose gel (1.5\%) and electrophoresis was run for 45 min at 100 V . The band corresponding to the 16 S rDNA V3-V5 amplicon was extracted and purified using Qiaquick gel extraction kit (Qiagen-28704). The amplicon quantity and integrity were checked on agarose gel electrophoresis and Qubit BR DNA assay kit. Amplicon was stored at $-20^{\circ} \mathrm{C}$ until use.

## Sequencing

16S rDNA amplicon library was sequenced at Institut Curie NGS platform on a MiSeq machine using the $2^{*} 300$ bp V3 kit.

## Bioinformatics pipeline

Reads were trimmed using sicle (Joshi et al. 2011) then error corrected using SPAdes (Bankevich et al. 2012), and merged using PEAR (Zhang et al. 2014). Reads were clustered at $97 \%$ of identity using vsearch pipeline (Rognes et al. 2016) to define Operational Taxonomic Units (OTU). Chimeric OTUs were identified using UCHIME (Edgar et al. 2011) and discarded from downstream analysis. Taxonomy of representative OTU sequences was determined using RDP SeqMatch and reference database RDP Release 11.3. OTU sequences were aligned using ssu-align (Nawrocki 2009). The phylogenetic tree was inferred from the OTUs multiple alignments using Fattree2 (Price et al. 2010).

## Computation of microbial diversity indexes (Marcon 2016)

Alpha-diversity indexes - which measure diversity in each sample - and Bray-Curtis dissimilarity matrix - which assesses how samples differ from each other ( $\beta$-diversity) - were computed from the unrarefied OTU count table. Inferred $\alpha$-diversity indexes were observed richness (number of distinct species observed in each sample), Chao1 richness estimate (which estimates the number of unobserved species), and Simpson's diversity index (which corresponds to the probability that two randomly picked sequences belong to the same species).

## Literature Cited

Bankevich A, Nurk S, Antipov D, Gurevich AA, Dvorkin M, Kulikov AS, Lesin VM, Nikolenko SI, Pham S, Prjibelski AD, Pyshkin AV, Sirotkin AV, Vyahhi N, Tesler G, Alekseyev MA, and Pevzner PA. 2012. SPAdes: a new genome assembly algorithm and its applications to single-cell sequencing. J. Comput. Biol. 19 (5): 455-477.

Edgar RC, Haas BJ, Clemente JC, Quince C, and Knight R. 2011. UCHIME improves sensitivity and speed of chimera detection. Bioinformatics. 27 (16): 2194-2200.

Joshi NA, and Fass JN. 2011. Sickle: A sliding-window, adaptive, quality-based trimming tool for FastQ files (Version 1.33) [Software]. Available at https://github.com/najoshi/sickle.

Kozich JJ, Westcott SL, Baxter NT, Highlander SK, and Schloss PD. 2013. Development of a dual-index sequencing strategy and curation pipeline for analyzing amplicon sequence data on the MiSeq Illumina sequencing platform. Appl. Environ. Microbiol. 79 (17): 5112-5120.

Marcon E. 2016. Mesures de la biodiversité. Available at http://www.ecofog.gf/spip.php?article427.

Nawrocki E. 2009. Structural RNA homology search and alignment using covariance models. Structural RNA homology search and alignment using covariance models.

O'Cuiv P, Aguirre de Carcer D, Jones M, Klaassens ES, Worthley DL, Whitehall VL, Kang S, McSweeney CS, Leggett BA, and Morrison M. 2011. The effects from DNA extraction methods on the evaluation of microbial diversity associated with human colonic tissue. Microb. Ecol. 61 (2): 353-362.

Price MN, Dehal PS, and Arkin AP. 2010. FastTree 2--approximately maximum-likelihood trees for large alignments. PLoS. One. 5 (3): e9490.

Rognes T, Flouri T, Nichols B, Quince C, and Mahe F. 2016. VSEARCH: a versatile open source tool for metagenomics. PeerJ. 4: e2584.

Yu Z, and Morrison M. 2004. Improved extraction of PCR-quality community DNA from digesta and fecal samples. Biotechniques 36 (5): 808-812.

Zhang J, Kobert K, Flouri T, and Stamatakis A. 2014. PEAR: a fast and accurate Illumina Paired-End reAd mergeR. Bioinformatics. 30 (5): 614-620.

Supplemental Table 1. Associations between consumption of total, soluble and insoluble dietary fibers, and the risk of breast and prostate cancers from multi-adjusted Cox proportional hazard models, NutriNet-Santé cohort, France, 2009-2019


|  | cancer ${ }^{\text {j }}$ | T2 | 26,581 (295) | 1.10 (0.90-1.34) |  | T2 | 26,581 (279) | 0.99 (0.82-1.21) |  | T2 | 26,581 (304) | 1.13 (0.93-1.36) |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | T3 | 26,581 (317) | 0.95 (0.76-1.19) |  | T3 | 26,581 (337) | 0.95 (0.77-1.17) |  | T3 | 26,581 (293) | 0.91 (0.73-1.13) |  |
|  | Prostate | T1 | 7179 (64) | 1.00 |  | T1 | 7179 (61) | 1.00 |  | T1 | 7179 (73) | 1.00 |  |
|  | cancer ${ }^{\text {k }}$ | T2 | 7179 (128) | 1.17 (0.85-1.60) | 0.8 | T2 | 7179 (120) | 0.99 (0.72-1.36) | 0.8 | T2 | 7179 (129) | 1.21 (0.90-1.64) | 0.6 |
|  |  | T3 | 7179 (131) | 0.98 (0.67-1.42) |  | T3 | 7179 (142) | 0.95 (0.67-1.35) |  | T3 | 7179 (121) | 0.94 (0.66-1.35) |  |
| Model $5^{\text {i }}$ | Breast cancer ${ }^{\text {j }}$ | T1 | 26,580 (171) | 1.00 | 0.2 | T1 | 26,580 (167) | 1.00 | 0.5 | T1 | 26,580 (186) | 1.00 | 0.09 |
|  |  | T2 | 26,581 (295) | 1.03 (0.81-1.31) |  | T2 | 26,581 (279) | 0.93 (0.73-1.18) |  | T2 | 26,581 (304) | 1.03 (0.82-1.30) |  |
|  |  | T3 | 26,581 (317) | 0.88 (0.68-1.14) |  | T3 | 26,581 (337) | 0.92 (0.72-1.17) |  | T3 | 26,581 (293) | 0.82 (0.64-1.06) |  |
|  | Prostate cancer ${ }^{\text {k }}$ | T1 | 7179 (64) | 1.00 | 0.8 | T1 | 7179 (61) | 1.00 | 0.7 | T1 | 7179 (73) | 1.00 | 0.8 |
|  |  | T2 | 7179 (128) | 1.18 (0.80-1.75) |  | T2 | 7179 (120) | 1.02 (0.69-1.53) |  | T2 | 7179 (129) | 1.22 (0.84-1.77) |  |
|  |  | T3 | 7179 (131) | 1.08 (0.70-1.67) |  | T3 | 7179 (142) | 1.09 (0.72-1.64) |  | T3 | 7179 (121) | 1.09 (0.72-1.65) |  |

Cl , confidence interval; HR , hazard ratio
${ }^{\text {a }}$ Cut-offs for tertiles of TDF consumption were $18.2 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $24.2 \mathrm{~g} . \mathrm{d}^{-1}$ for men (prostate cancer model) and $15.5 \mathrm{~g} . \mathrm{d}^{-1}$ and $20.5 \mathrm{~g} . \mathrm{d}^{-1}$ for women (breast cancer model).
${ }^{\mathrm{b}}$ Cut-offs for tertiles of SF consumption were $5.0 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $7.4 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for men (prostate cancer model) and $4.2 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $6.1 \mathrm{~g} . \mathrm{d}^{-1}$ for women (breast cancer model).
${ }^{c}$ Cut-offs for tertiles of IF consumption were $12.8 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $17.1 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for men (prostate cancer model) and $11.0 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $14.5 \mathrm{~g} . \mathrm{d}^{-1}$ for women (breast cancer model).
${ }^{d} \mathrm{HR}(95 \% \mathrm{Cl})$ and $p$-value were computed for Cox proportional hazard model.
${ }^{e}$ Model 1 was adjusted for (timescale), sex, body-mass index, height, physical activity, smoking status, alcohol intake, energy intake, number of 24-hour dietary records, educational level, and family history of the chronic pathology of interest.
${ }^{f}$ Model $2=$ Model 1 further adjusted for dyslipidemia and associated medical treatment, hypertriglyceridemia and associated medical treatment, hypertension and associated medical treatment. CVD and cancer models were also adjusted for T2D and T2D medical treatment.
${ }^{\mathrm{g}}$ Model 3 = Model 1 further adjusted for saturated fat, sodium, and added sugar intakes.
${ }^{h}$ Model $4=$ Model 1 further adjusted for vitamin C, vitamin E, zinc, selenium intakes.
${ }^{i}$ Model 5 = Model 1, carried out by excluding participants whose conditions were diagnosed or whose death occurred within 2 years after inclusion. Cancer models were also adjusted for height.
${ }^{\mathrm{j}}$ Mean follow-up time for breast cancer was 5.8 years, with 435,186 person-years.
${ }^{k}$ Mean follow-up time for prostate cancer was 6.0 y , with 120,200 person-years.

Supplemental Table 2. Associations between consumption of dietary fibers from different sources, and the risk of breast and prostate cancers from multi-adjusted Cox proportional hazard models, NutriNet-Santé cohort, France, 2009-2019

|  | Total dietary fibers from fruits |  |  |  | Total dietary fibers from vegetables |  |  |  | Total dietary fibers from whole grains |  |  |  | Total dietary fibers from legumes |  |  |  | Total dietary fibers from potatoes \& tubers |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Tertile ${ }^{\text {a }}$ | N (of which cases) | $\begin{gathered} \text { HR } \\ (95 \% \mathrm{CI})^{\mathrm{f}} \end{gathered}$ | $\begin{gathered} \mathrm{p}- \\ \text { trend } \end{gathered}$ | Tertile ${ }^{\text {b }}$ | N (of which cases) | $\begin{gathered} \text { HR } \\ (95 \% \mathrm{Cl})^{f} \end{gathered}$ | $\begin{gathered} \mathrm{p}- \\ \text { trend } \end{gathered}$ | Tertile ${ }^{\text {c }}$ | N (of which cases) | $\begin{gathered} \text { HR } \\ (95 \% \mathrm{CI})^{\mathrm{f}} \end{gathered}$ | $\begin{gathered} \mathrm{p}- \\ \text { trend } \end{gathered}$ | Tertile ${ }^{\text {d }}$ | N (of which cases) | $\begin{gathered} \text { HR } \\ (95 \% \mathrm{Cl})^{f} \end{gathered}$ | $\begin{gathered} \mathrm{p}- \\ \text { trend } \end{gathered}$ | Tertile ${ }^{\text {e }}$ | N (of which cases) | HR (95\%CI) ${ }^{\text {f }}$ | $\begin{gathered} \mathrm{p}- \\ \text { trend } \end{gathered}$ |
| Breast cancer ${ }^{\text {g }}$ |  | 26,580 | 1.00 |  |  | 26,580 | 1.00 |  |  | 26,667 | 1.00 |  |  | 44,634 | 1.00 |  |  | 26,580 | 1.00 |  |
|  | T1 | (172) |  |  | T1 | (178) |  |  | T1 | (219) |  |  | T1 | (378) |  |  | T1 | (223) |  |  |
|  |  | 26,581 | 0.92 | 0.8 |  | 26,581 | 1.01 | 0.6 |  | 26,494 | $0.93$ | 0.06 |  | 8154 | 0.99 $(0.79-1.25)$ | 0.9 |  | 26,582 | $1.02$ | 0.5 |
|  | T2 | (261) | (0.76-1.12) | 0.8 | T2 | (290) | (0.84-1.23) | 0.6 | T2 | (283) | (0.77-1.11) |  | T2 | (96) | (0.79-1.25) |  | T2 | (297) | (0.86-1.22) |  |
|  |  | 26,581 | 0.95 |  |  | 26,581 | 0.96 |  |  | 26,581 | 0.84 |  |  | 26,576 | 1.01 |  |  | 26,580 | 0.94 |  |
|  | T3 | (350) | (0.78-1.16) |  | T3 | (315) | (0.79-1.16) |  | T3 | (281) | (0.70-1.01) |  | T3 | (309) | (0.86-1.18) |  | T3 | (263) | (0.78-1.13) |  |
| Prostate cancer ${ }^{\text {h }}$ | T1 | $\begin{aligned} & 7179 \\ & (48) \end{aligned}$ | 1.00 |  | T1 | $\begin{aligned} & 7179 \\ & (65) \end{aligned}$ | 1.00 |  | T1 | $\begin{gathered} 7408 \\ (75) \end{gathered}$ | 1.00 |  | T1 | $\begin{gathered} 11,406 \\ (132) \end{gathered}$ | 1.00 |  | T1 | $7179$ (76) | 1.00 |  |
|  | T2 | $\begin{aligned} & 7179 \\ & (115) \end{aligned}$ | $\begin{gathered} 1.15 \\ (0.81-1.61) \end{gathered}$ | 0.6 | T2 | $\begin{aligned} & 7179 \\ & (127) \end{aligned}$ | $\begin{gathered} 1.14 \\ (0.81-1.60) \end{gathered}$ | 0.9 | T2 | $\begin{gathered} 951 \\ (120) \end{gathered}$ | $\begin{gathered} 1.23 \\ (0.92-1.65) \end{gathered}$ | 0.3 | T2 | $\begin{gathered} 2952 \\ (65) \end{gathered}$ | $\begin{gathered} 1.25 \\ (0.92-1.71) \end{gathered}$ | 0.9 | T2 | $\begin{aligned} & 7179 \\ & (129) \end{aligned}$ | $\begin{gathered} 1.13 \\ (0.85-1.51) \end{gathered}$ | 0.8 |
|  |  | 7179 | 1.14 |  |  | 7179 | 1.11 |  |  | 7178 | 1.21 |  |  | 7179 | 1.02 |  |  | 7179 | 1.00 |  |
|  | T3 | (160) | (0.81-1.60) |  | T3 | (131) | (0.82-1.50) |  | T3 | (128) | (0.90-1.61) |  | T3 | (126) | (0.79-1.31) |  | T3 | (118) | (0.75-1.35) |  |

Cl , confidence interval; HR , hazard ratio
${ }^{\text {a }}$ Cut-offs for sex-specific tertiles of TDF from fruits consumption were $2.1 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $4.6 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for men and $2.0 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $4.1 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for women.
${ }^{\mathrm{b}}$ Cut-offs for sex-specific tertiles of TDF from vegetables consumption were $3.6 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $5.9 \mathrm{~g} . \mathrm{d}^{-1}$ for men and $3.5 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $5.6 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for women.
${ }^{\text {c }}$ Cut-offs for sex-specific tertiles of TDF from whole grains consumption were $0 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $2.3 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for men and $0.3 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $2.1 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for women.
${ }^{d}$ Cut-offs for sex-specific tertiles of TDF from legumes consumption were $0 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $0.6 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for men and $0 \mathrm{~g} \cdot \mathrm{~d}^{-1} 0.4 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for women.
${ }^{e}$ Cut-offs for sex-specific tertiles of TDF from potatoes and tubers consumption were $0.5 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $1.2 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for men and $0.4 \mathrm{~g} \cdot \mathrm{~d}^{-1} 1.0 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for women.
 records, educational level, and family history cancer.
${ }^{\mathrm{g}}$ Mean follow-up time for breast cancer was 5.8 years, with 435,186 person-years.
${ }^{h}$ Mean follow-up times for prostate cancer was $6.0 y$, with 120,200 person-years.

Supplemental Table 3. Sensitivity analyses for the associations between consumption of total, soluble and insoluble dietary fibers, and the risk of overall mortality, mortality attributed to chronic diseases, overall CVD, coronary heart diseases, cerebrovascular diseases, overall cancers, colorectal cancer, and type 2 diabetes from multi-adjusted Cox proportional hazard models, NutriNet-Santé cohort, France, 2009-2019.

| Model $2^{\text {e }}$ | Mortality from chronic diseases ${ }^{\text {' }}$ | Total dietary fibers |  |  |  | Soluble fibers |  |  |  | Insoluble fibers |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Tertile ${ }^{\text {a }}$ | N (of which cases) | HR (95\%CI) ${ }^{\text {d }}$ | $\begin{gathered} \mathrm{p}- \\ \text { trend }^{\mathrm{d}} \end{gathered}$ | Tertile ${ }^{\text {b }}$ | N (of which cases) | HR (95\%CI) ${ }^{\text {d }}$ | $\begin{gathered} \mathrm{p}- \\ \text { trend }^{\mathrm{d}} \end{gathered}$ | Tertile ${ }^{\text {c }}$ | N (of which cases) | HR (95\%CI) ${ }^{\text {d }}$ | p-trend ${ }^{\text {d }}$ |
|  |  | T1 | 35,791 (194) | 1.00 |  | T1 | 35,791 (174) | 1.00 |  | T1 | 35,791 (216) | 1.00 |  |
|  |  | T2 | 35,793 (195) | 0.79 (0.64-0.98) | 0.003 | T2 | 35,793 (204) | 0.83 (0.67-1.02) | 0.003 | T2 | 35,793 (186) | 0.76 (0.62-0.93) | 0.0004 |
|  |  | T3 | 35,793 (191) | 0.70 (0.55-0.88) |  | T3 | 35,793 (202) | 0.71 (0.57-0.89) |  | T3 | 35,793 (178) | 0.66 (0.52-0.83) |  |
|  | Overall cardiovascular diseases ${ }^{\text {j }}$ | T1 | 34,934 (479) | 1.00 |  | T1 | 34,934 (463) | 1.00 |  | T1 | 34,934 (512) | 1.00 |  |
|  |  | T2 | 34,936 (740) | 1.04 (0.92-1.18) | 0.1 | T2 | 34,936 (727) | 0.97 (0.86-1.09) | 0.01 | T2 | 34,936 (710) | 1.05 (0.93-1.18) | 1.0 |
|  |  | T3 | 34,935 (733) | 0.91 (0.80-1.05) |  | T3 | 34,935 (762) | 0.86 (0.75-0.98) |  | T3 | 34,935 (730) | 1.00 (0.88-1.14) |  |
|  | Coronary heart diseases | T1 | 34,934 (302) | 1.00 |  | T1 | 34,934 (296) | 1.00 |  | T1 | 34,934 (314) | 1.00 |  |
|  |  | T2 | 34,936 (455) | 1.05 (0.90-1.22) | 0.5 | T2 | 34,936 (468) | 0.98 (0.84-1.14) | 0.007 | T2 | 34,936 (449) | 1.12 (0.96-1.30) | 0.5 |
|  |  | T3 | 34,935 (462) | 0.96 (0.81-1.14) |  | T3 | 34,935 (455) | 0.81 (0.69-0.96) |  | T3 | 34,935 (456) | 1.07 (0.91-1.27) |  |
|  | Cerebrovascular diseases | T1 | 34,934 (209) | 1.00 |  | T1 | 34,934 (197) | 1.00 |  | T1 | 34,934 (235) | 1.00 |  |
|  |  | T2 | 34,936 (341) | 1.04 (0.87-1.25) | 0.1 | T2 | 34,936 (304) | 0.93 (0.77-1.12) | 0.8 | T2 | 34,936 (315) | 0.96 (0.8-1.14) | 0.3 |
|  |  | T3 | 34,935 (328) | 0.86 (0.70-1.06) |  | T3 | 34,935 (377) | 0.96 (0.80-1.17) |  | T3 | 34,935 (328) | 0.90 (0.74-1.10) |  |
|  | Overall cancers ${ }^{\text {k }}$ | T1 | 33,759 (591) | 1.00 |  | T1 | 33,759 (571) | 1.00 |  | T1 | 33,759 (643) | 1.00 |  |
|  |  | T2 | 33,760 (914) | 1.06 (0.95-1.18) | 1.0 | T2 | 33,760 (883) | 0.97 (0.87-1.08) | 0.6 | T2 | 33,760 (916) | 1.08 (0.97-1.19) | 0.8 |
|  |  | T3 | 33,760 (998) | 1.01 (0.90-1.14) |  | T3 | 33,760 (1049) | 0.97 (0.86-1.08) |  | T3 | 33,760 (944) | 0.99 (0.89-1.12) |  |
|  | Colorectal cancer ${ }^{\text {k }}$ | T1 | 33,759 (44) | 1.00 |  | T1 | 33,759 (45) | 1.00 |  | T1 | 33,759 (52) | 1.00 |  |
|  |  | T2 | 33,760 (77) | 1.12 (0.76-1.65) | 0.1 | T2 | 33,760 (70) | 0.86 (0.59-1.27) | 0.04 | T2 | 33,760 (72) | 1.01 (0.69-1.46) | 0.1 |
|  |  | T3 | 33,760 (61) | 0.74 (0.47-1.16) |  | T3 | 33,760 (67) | 0.66 (0.44-1.00) |  | T3 | 33,760 (58) | 0.71 (0.46-1.08) |  |
|  | Type 2-diabetes' | T1 | 34,901 (226) | 1.00 |  | T1 | 34,901 (210) | 1.00 |  | T1 | 34,901 (258) | 1.00 |  |
|  |  | T2 | 34,903 (292) | 0.86 (0.72-1.03) | 0.02 | T2 | 34,903 (305) | 0.87 (0.73-1.05) | 0.003 | T2 | 34,903 (273) | 0.79 (0.66-0.94) | 0.004 |
|  |  | T3 | 34,903 (303) | 0.79 (0.64-0.96) |  | T3 | 34,903 (306) | 0.75 (0.61-0.91) |  | T3 | 34,903 (290) | 0.74 (0.61-0.91) |  |
| Model $3^{\text {f }}$ | Chronic-disease related mortality ${ }^{\text {i }}$ | T1 | 35,791 (194) | 1.00 |  | T1 | 35,791 (174) | 1.00 |  | T1 | 35,791 (216) | 1.00 |  |
|  |  | T2 | 35,793 (195) | 0.80 (0.64-1.00) | 0.04 | T2 | 35,793 (204) | 0.86 (0.69-1.06) | 0.04 | T2 | 35,793 (186) | 0.76 (0.61-0.94) | 0.004 |
|  |  | T3 | 35,793 (191) | 0.73 (0.55-0.98) |  | T3 | 35,793 (202) | 0.77 (0.59-0.99) |  | T3 | 35,793 (178) | 0.66 (0.50-0.88) |  |


|  | Overall cardiovascular diseases ${ }^{j}$ | T1 | 34,934 (479) | 1.00 |  | T1 | 34,934 (463) | 1.00 |  | T1 | 34,934 (512) | 1.00 |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | T2 | 34,936 (740) | 1.05 (0.93-1.19) | 0.4 | T2 | 34,936 (727) | 0.98 (0.87-1.11) | 0.06 | T2 | 34,936 (710) | 1.06 (0.94-1.20) | 0.6 |
|  |  | T3 | 34,935 (733) | 0.94 (0.81-1.09) |  | T3 | 34,935 (762) | 0.88 (0.76-1.01) |  | T3 | 34,935 (730) | 1.05 (0.91-1.20) |  |
|  | Coronary heart diseases | T1 | 34,934 (302) | 1.00 |  | T1 | 34,934 (296) | 1.00 |  | T1 | 34,934 (314) | 1.00 |  |
|  |  | T2 | 34,936 (455) | 1.06 (0.91-1.24) | 1.0 | T2 | 34,936 (468) | 1.00 (0.85-1.16) | 0.02 | T2 | 34,936 (449) | 1.13 (0.97-1.32) | 0.2 |
|  |  | T3 | 34,935 (462) | 1.01 (0.83-1.21) |  | T3 | 34,935 (455) | 0.82 (0.69-0.99) |  | T3 | 34,935 (456) | 1.13 (0.95-1.35) |  |
|  | Cerebrovascular diseases ${ }^{\text {j }}$ | T1 | 34,934 (209) | 1.00 |  | T1 | 34,934 (197) | 1.00 |  | T1 | 34,934 (235) | 1.00 |  |
|  |  | T2 | 34,936 (341) | 1.05 (0.87-1.26) | 0.2 | T2 | 34,936 (304) | 0.96 (0.79-1.15) | 0.7 | T2 | 34,936 (315) | 0.97 (0.81-1.16) | 0.5 |
|  |  | T3 | 34,935 (328) | 0.89 (0.71-1.11) |  | T3 | 34,935 (377) | 1.03 (0.84-1.28) |  | T3 | 34,935 (328) | 0.94 (0.76-1.15) |  |
|  | Overall cancers ${ }^{\text {k }}$ | T1 | 33,759 (591) | 1.00 |  | T1 | 33,759 (571) | 1.00 |  | T1 | 33,759 (643) | 1.00 |  |
|  |  | T2 | 33,760 (914) | 1.06 (0.95-1.18) | 0.9 | T2 | 33,760 (883) | 0.97 (0.87-1.08) | 0.7 | T2 | 33,760 (916) | 1.07 (0.96-1.19) | 0.7 |
|  |  | T3 | 33,760 (998) | 1.00 (0.88-1.14) |  | T3 | 33,760 (1049) | 0.97 (0.86-1.10) |  | T3 | 33,760 (944) | 0.98 (0.87-1.11) |  |
|  | Colorectal cancer ${ }^{\text {k }}$ | T1 | 33,759 (44) | 1.00 |  | T1 | 33,759 (45) | 1.00 |  | T1 | 33,759 (52) | 1.00.00 |  |
|  |  | T2 | 33,760 (77) | 1.13 (0.76-1.68) | 0.2 | T2 | 33,760 (70) | 0.86 (0.58-1.28) | 0.07 | T2 | 33,760 (72) | 1.02 (0.70-1.49) | 0.2 |
|  |  | T3 | 33,760 (61) | 0.76 (0.46-1.24) |  | T3 | 33,760 (67) | 0.66 (0.42-1.05) |  | T3 | 33,760 (58) | 0.73 (0.46-1.16) |  |
|  | Type 2-diabetes ${ }^{\text {' }}$ | T1 | 34,901 (226) | 1.00 |  | T1 | 34,901 (210) | 1.00 |  | T1 | 34,901 (258) | 1.00 |  |
|  |  | T2 | 34,903 (292) | 0.81 (0.67-0.98) | 0.001 | T2 | 34,903 (305) | 0.83 (0.69-1.00) | <0.0001 | T2 | 34,903 (273) | 0.76 (0.63-0.90) | 0.0003 |
|  |  | T3 | 34,903 (303) | 0.69 (0.55-0.86) |  | T3 | 34,903 (306) | 0.64 (0.52-0.80) |  | T3 | 34,903 (290) | 0.67 (0.55-0.83) |  |
| Model $\mathbf{4}^{\text {g }}$ | Chronic-disease related mortality ${ }^{i}$ | T1 | 35,791 (194) | 1.00 |  | T1 | 35,791 (174) | 1.00 |  | T1 | 35,791 (216) | 1.00 |  |
|  |  | T2 | 35,793 (195) | 0.76 (0.61-0.93) | 0.0002 | T2 | 35,793 (204) | 0.81 (0.65-0.995) | 0.0003 | T2 | 35,793 (186) | 0.72 (0.59-0.89) | <0.0001 |
|  |  | T3 | 35,793 (191) | 0.62 (0.49-0.80) |  | T3 | 35,793 (202) | 0.65 (0.51-0.82) |  | T3 | 35,793 (178) | 0.59 (0.46-0.75) |  |
|  | Overall cardiovascular diseases |  |  |  |  |  | 34,934 (463) |  |  | T1 |  | $1.00$ |  |
|  |  | T2 | 34,936 (740) | $1.04 \text { (0.92-1.17) }$ | 0.1 | T2 | 34,936 (727) | 0.97 (0.86-1.10) | 0.02 | T2 | 34,936 (710) | $1.05 \text { (0.93-1.18) }$ | 1.0 |
|  |  | T3 | 34,935 (733) | 0.90 (0.78-1.05) |  | T3 | 34,935 (762) | 0.85 (0.74-0.98) |  | T3 | 34,935 (730) | 1.00 (0.87-1.15) |  |
|  | Coronary heart diseases | T1 | 34,934 (302) | 1.00 |  | T1 | 34,934 (296) | 1.00 |  | T1 | 34,934 (314) | 1.00 |  |
|  |  | T2 | 34,936 (455) | 1.03 (0.88-1.20) | 0.3 | T2 | 34,936 (468) | 0.92 (0.76-1.10) | 0.003 | T2 | 34,936 (449) | 1.10 (0.95-1.28) | 0.8 |
|  |  | T3 | 34,935 (462) | 0.92 (0.76-1.10) |  | T3 | 34,935 (455) | 0.98 (0.84-1.14) |  | T3 | 34,935 (456) | 1.04 (0.87-1.24) |  |
|  | Cerebrovascular diseases ${ }^{\text {j }}$ | T1 | 34,934 (209) |  |  | T1 | 34,934 (197) |  |  | T1 | 34,934 (235) | $1.00$ |  |
|  |  | T2 | 34,936 (341) | 1.05 (0.88-1.27) | 0.3 | T2 | 34,936 (304) | 0.95 (0.79-1.15) | 0.7 | T2 | 34,936 (315) | $0.97 \text { (0.81-1.16) }$ | 0.6 |
|  |  | T3 | 34,935 (328) | 0.90 (0.72-1.12) |  | T3 | 34,935 (377) | 1.03 (0.83-1.26) |  | T3 | 34,935 (328) | 0.95 (0.77-1.16) |  |
|  | Overall cancers ${ }^{\text {k }}$ | T1 | 33,759 (591) | 1.00 |  | T1 | 33,759 (571) | 1.00 |  | T1 | 33,759 (643) | 1.00 |  |
|  |  | T2 | 33,760 (914) | 1.04 (0.93-1.16) | 0.5 | T2 | 33,760 (883) | 0.96 (0.86-1.07) | 0.3 | T2 | 33,760 (916) | 1.06 (0.95-1.17) | 0.4 |
|  |  | T3 | 33,760 (998) | 0.97 (0.86-1.10) |  | T3 | 33,760 (1049) | 0.93 (0.83-1.05) |  | T3 | 33,760 (944) | 0.96 (0.85-1.09) |  |


|  | Colorectal cancer ${ }^{\text {k }}$ | T1 | 33,759 (44) | 1.00 | 0.3 | T1 | 33,759 (45) | 1.00 |  | T1 | 33,759 (52) | 1.00 |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | T2 | 33,760 (77) | 1.15 (0.77-1.70) |  | T2 | 33,760 (70) | 0.88 (0.60-1.31) | 0.09 | T2 | 33,760 (72) | 1.03 (0.70-1.50) | 0.2 |
|  |  | T3 | 33,760 (61) | 0.78 (0.48-1.27) |  | T3 | 33,760 (67) | 0.69 (0.44-1.08) |  | T3 | 33,760 (58) | 0.74 (0.47-1.17) |  |
|  | Type 2-diabetes ${ }^{\text {' }}$ | T1 | 34,901 (226) | 1.00 | 0.002 | T1 | 34,901 (210) | 1.00 | 0.0009 | T1 | 34,901 (258) | 1.00 | 0.0001 |
|  |  | T2 | 34,903 (292) | 0.82 (0.68-0.99) |  | T2 | 34,903 (305) | 0.88 (0.73-1.05) |  | T2 | 34,903 (273) | 0.75 (0.62-0.89) |  |
|  |  | T3 | 34,903 (303) | 0.71 (0.57-0.88) |  | T3 | 34,903 (306) | 0.71 (0.57-0.87) |  | T3 | 34,903 (290) | 0.66 (0.54-0.81) |  |
| Model $5^{\text {h }}$ | Chronic-disease related mortality ${ }^{\text {i }}$ | T1 | 35,791 (194) | 1.00 | 0.01 | T1 | 35,791 (174) | 1.00 | 0.09 | T1 | 35,791 (216) | 1.00 | 0.0006 |
|  |  | T2 | 35,793 (195) | 0.71 (0.55-0.91) |  | T2 | 35,793 (204) | 0.89 (0.69-1.15) |  | T2 | 35,793 (186) | 0.74 (0.58-0.95) |  |
|  |  | T3 | 35,793 (191) | 0.69 (0.52-0.91) |  | T3 | 35,793 (202) | 0.79 (0.60-1.04) |  | T3 | 35,793 (178) | 0.62 (0.47-0.81) |  |
|  | Overall cardiovascular diseases ${ }^{\text {j }}$ | T1 | 34,934 (479) | 1.00 | 0.04 | T1 | 34,934 (463) | 1.00 | 0.002 | T1 | 34,934 (512) | 1.00 | 0.6 |
|  |  | T2 | 34,936 (740) | 1.04 (0.90-1.19) |  | T2 | 34,936 (727) | $0.91 \text { (0.79-1.04) }$ |  | T2 | 34,936 (710) | $1.04(0.91-1.18)$ |  |
|  |  | T3 | 34,935 (733) | 0.87 (0.75-1.01) |  | T3 | 34,935 (762) | $0.80(0.69-0.92)$ |  | T3 | 34,935 (730) | $0.97 \text { (0.84-1.13) }$ |  |
|  | Coronary heart diseases ${ }^{j}$ | T1 | 34,934 (302) | 1.00 | 0.3 | T1 | 34,934 (296) | 1.00 | 0.0007 | T1 | 34,934 (314) | 1.00 | 0.6 |
|  |  | T2 | 34,936 (455) | 1.08 (0.91-1.29) |  | T2 | 34,936 (468) | 0.89 (0.75-1.06) |  | T2 | 34,936 (449) | 1.14 (0.96-1.35) |  |
|  |  | T3 | 34,935 (462) | 0.92 (0.75-1.11) |  | T3 | 34,935 (455) | 0.73 (0.61-0.88) |  | T3 | 34,935 (456) | 1.07 (0.88-1.29) |  |
|  | Cerebrovascular diseases ${ }^{\text {j }}$ | T1 | 34,934 (209) | 1.00 | 0.09 | T1 | 34,934 (197) | 1.00 | 0.6 | T1 | 34,934 (235) | 1.00 | 0.2 |
|  |  | T2 | 34,936 (341) | 1.02 (0.84-1.24) |  | T2 | 34,936 (304) | 0.91 (0.74-1.11) |  | T2 | 34,936 (315) | 0.96 (0.79-1.16) |  |
|  |  | T3 | 34,935 (328) | 0.84 (0.68-1.05) |  | T3 | 34,935 (377) | 0.94 (0.76-1.15) |  | T3 | 34,935 (328) | 0.88 (0.71-1.09) |  |
|  | Overall cancers ${ }^{\prime}$ | T1 | 33,759 (591) | 1.00 | 0.5 | T1 | 33,759 (571) | 1.00 | 0.7 | T1 | 33,759 (643) | 1.00 | 0.4 |
|  |  | T2 | 33,760 (914) | 1.03 (0.90-1.18) |  | T2 | 33,760 (883) | 0.98 (0.86-1.12) |  | T2 | 33,760 (916) | 1.02 (0.90-1.16) |  |
|  |  | T3 | 33,760 (998) | 0.97 (0.84-1.11) |  | T3 | 33,760 (1049) | 0.97 (0.84-1.11) |  | T3 | 33,760 (944) | 0.95 (0.83-1.10) |  |
|  | Colorectal cancer ${ }^{\prime}$ | T1 | 33,759 (44) |  | 0.06 | T1 | 33,759 (45) | 1.00 | 0.01 | T1 | 33,759 (52) | 1.00 | 0.1 |
|  |  | T2 | 33,760 (77) | 0.95 (0.60-1.51) |  | T2 | 33,760 (70) | 0.76 (0.48-1.19) |  | T2 | 33,760 (72) | 0.86 (0.55-1.35) |  |
|  |  | T3 | 33,760 (61) | 0.62 (0.36-1.05) |  | T3 | 33,760 (67) | 0.55 (0.33-0.90) |  | T3 | 33,760 (58) | 0.66 (0.40-1.10) |  |
|  | Type 2-diabetes' | T1 | 34,901 (226) | 1.00 | 0.0004 | T1 | 34,901 (210) | 1.00 | <0.0001 | T1 | 34,901 (258) | 1.00 | <0.0001 |
|  |  | T2 | 34,903 (292) | 0.85 (0.68-1.06) |  | T2 | 34,903 (305) | 0.83 (0.67-1.04) |  | T2 | 34,903 (273) | 0.77 (0.62-0.96) |  |
|  |  | T3 | 34,903 (303) | 0.64 (0.50-0.82) |  | T3 | 34,903 (306) | 0.61 (0.48-0.78) |  | T3 | 34,903 (290) | 0.58 (0.46-0.75) |  |

[^0]${ }^{\text {a }}$ Cut-offs for sex-specific tertiles of TDF consumption were $18.2 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ and $24.2 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for men and $15.5 \mathrm{~g} . \mathrm{d}^{-1} 20.5 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for women.
${ }^{\text {b }}$ Cut-offs for sex-specific tertiles of SF consumption were $5.0 \mathrm{~g} . \mathrm{d}^{-1}$ and $7.4 \mathrm{~g} . \mathrm{d}^{-1}$ for men and $4.2 \mathrm{~g} \cdot \mathrm{~d}^{-1} 6.1 \mathrm{~g} . \mathrm{d}^{-1}$ for women.
${ }^{\text {c }}$ Cut-offs for sex-specific tertiles of IF consumption were $12.8 \mathrm{~g} . \mathrm{d}^{-1}$ and $17.1 \mathrm{~g} \cdot \mathrm{~d}^{-1}$ for men and $11.0 \mathrm{~g} . \mathrm{d}^{-1} 14.6 \mathrm{~g} . \mathrm{d}^{-1}$ for women.
${ }^{d} \mathrm{HR}(95 \% \mathrm{Cl})$ and $p$-value were computed for Cox proportional hazard model.
${ }^{e}$ Model 2: adjusted for age (timescale), sex, body-mass index, physical activity, smoking status, alcohol intake, energy intake, number of 24-hour dietary records, educational level, and family history of the chronic pathology of interest (CVD, cancer, or T2D) for pathology models or family history of CVD and cancer for mortality models (model 1) + dyslipidemia and associated medical treatment, hypertriglyceridemia and associated medical treatment, hypertension and associated medical treatment. Mortality, CVD, and cancer models were also adjusted for T2D and T2D medical treatment. Cancer models were also adjusted for height.
${ }^{\text {f }}$ Model 3: model 1 + saturated fat, sodium, and added sugar intakes. Cancer models were also adjusted for height.
${ }^{8}$ Model 4: model $1+$ intakes of vitamin C , vitamin E , selenium, and zinc. Cancer models were also adjusted for height.
${ }^{\text {h }}$ Model 5 was adjusted for model 1 +was carried out by excluding participants whose conditions were diagnosed or whose death occurred within 2 years after inclusion. Cancer models were also adjusted for height. 'Mean follow-up time for mortality was 6.3 years, with 616,427 person-years.
${ }^{j}$ Mean follow-up times for overall cardiovascular diseases, coronary heart diseases and cerebrovascular diseases were all equal to 6.0 y . Person-years were $579,099,581,374$, and 582,790 respectively.
${ }^{k}$ Mean follow-up times for overall cancers and colorectal cancer was 6.0y. Person-years was 579,099.
'Mean follow-up times for type 2-diabetes was 6.0y. Person-years was 582,237 .

Supplemental Figure 1. Distributions of the consumption of total dietary fibers (A), insoluble fibers (B), and soluble fibers (C) in $\mathrm{g} \cdot \mathrm{d}^{-1}$ in the study sample ( $\mathrm{N}=107,377$ ), NutriNet-Santé cohort, France, 2009-2019.
(A)

(B)

(C)


Supplemental Figure 2. Scaled Schoenfeld residuals for the DF variables in the (A) mortality, (B) cardiovascular disease, (C) coronary heart disease, (D) cerebrovascular disease, (E) global cancer, (F) colorectal cancer, (G) breast cancer, (H) prostate cancer models, NutriNet-Santé cohort, France, 2009-2019.

The solid black and the dashed black curves represent the smooth curve and its $95 \%$ confidence interval, respectively. For comparison purpose, the estimated coefficient is showed as a solid blue line.
(A)

(B)





Schoenfeld Individual Test p: 0.8866



## (C)





## (D)






Schoenfeld Individual Test p: 0.2112


Schoenfeld Individual Test p: 0.7505


## (E)


(F)





Schoenfeld Individual Test p: 0.148


Schoenfeld Individual Test p: 0.1136

(G)

(H)



[^0]:    Cl, confidence interval; HR, hazard ratio

