

Elevated Transferrin Saturation and Risk of Diabetes

Three population-based studies

CHRISTINA ELLERVIK, MD, PHD^{1,2,3}
 THOMAS MANDRUP-POULSEN, MD, DMSCI^{2,4,5,6}
 HENRIK ULLITS ANDERSEN, MD, DMSCI⁶
 ANNE TYBJÆRG-HANSEN, MD, DMSCI^{2,7,8,9}

MERETE FRANDBEN, BMLT⁶
 HENRIK BIRGENS, MD, DMSCI^{2,10}
 BØRGE G. NORDESTGAARD, MD, DMSCI^{1,2}

RESEARCH DESIGN AND METHODS

—Details are provided in the Supplementary Data online.

OBJECTIVE—We tested the hypothesis that elevated transferrin saturation is associated with an increased risk of any form of diabetes, as well as type 1 or type 2 diabetes separately.

RESEARCH DESIGN AND METHODS—We used two general population studies, The Copenhagen City Heart Study (CCHS, $N = 9,121$) and The Copenhagen General Population Study (CGPS, $N = 24,195$), as well as a 1:1 age- and sex-matched population-based case-control study with 6,129 patients with diabetes from the Steno Diabetes Centre and 6,129 control subjects, totaling 8,535 patients with diabetes and 37,039 control subjects.

RESULTS—In the combined studies, odds ratios in those with transferrin saturation $\geq 50\%$ vs. $< 50\%$ were 2.1 (95% CI 1.3–3.4; $P = 0.003$) for any form of diabetes; 2.6 (1.2–5.6; $P = 0.01$) for type 1 diabetes; and 1.7 (1.4–2.1; $P = 0.001$) for type 2 diabetes.

CONCLUSIONS—Elevated transferrin saturation confers a two- to threefold increased risk of developing any form of diabetes, as well as type 1 and type 2 diabetes separately.

Diabetes Care 34:2256–2258, 2011

Diabetes is a common late complication of hereditary hemochromatosis, an autosomal-recessive disease with lifelong iron accumulation in metabolically highly active tissues, primarily in the endocrine pancreas, the myocardium, the synovial membrane, the anterior pituitary lobe, and the liver (1). Diabetes as a consequence of hemochromatosis is characterized by insulin resistance and β -cell destruction (2), and the metabolic phenotype can therefore mimic type 2 diabetes and nonautoimmune type 1 diabetes (3).

The defects in insulin-producing and insulin-sensitive tissues are most likely caused by iron-dependent catalysis via the Fenton reaction of reactive oxygen radical species, which impair insulin signaling in skeletal muscle and liver and cause β -cell destruction due to an insufficient β -cell-deficient antioxidant defense (4). It is unclear, however, whether elevated transferrin saturation confers an increased risk of diabetes in the general population. Our aim was to test this hypothesis in three large populations (Supplementary Data online).

Participants

The studies were approved by the appropriate institutional review boards and ethical committees. The investigation was conducted according to the principles outlined in the Declaration of Helsinki. Written informed consent was obtained from participants. All participants were white and of Danish descent. There was no overlap of individuals between the three studies, thus allowing independent confirmation of findings. We included all attending individuals with eligible transferrin saturation.

General population studies

On the basis of the Danish Central Person Registration number, The Copenhagen City Heart Study (CCHS, $N = 9,121$; recruited 1991–1994) (5) and The Copenhagen General Population Study (CGPS, $N = 24,195$; recruited 2003–2007) (6) recruited participants randomly from the general population of Copenhagen, Denmark. The studies included a self-administered questionnaire, a physical examination, and blood samples.

Population-based case-control study

We recruited (2001–2007) consecutively 6,129 population-based patients with diabetes from Copenhagen County who attended the Steno Diabetes Centre.

Transferrin saturation

Transferrin saturation (percentage) was determined as iron levels (in micromoles per liter) divided by $2 \times$ transferrin levels (in micromoles per liter) $\times 100$. Transferrin was measured by turbidimetry, and iron levels were measured by colorimetry using a Konelab autoanalyzer (Thermo Fisher Scientific, Waltham, MA) and Hitachi 912 (Roche Diagnostics, Indianapolis, IN). Transferrin saturation $> 50\%$ was chosen as suggestive of increased transferrin saturation, in accordance with accepted clinical practice (7–9). Transferrin saturation range was 2.8–236%

From the ¹Department of Clinical Biochemistry, Herlev Hospital, Copenhagen University Hospital, Herlev, Denmark; the ²Faculty of Health Sciences, University of Copenhagen, Copenhagen, Denmark; the ³Department of Clinical Biochemistry, Næstved Hospital, Copenhagen University Hospital, Næstved, Denmark; the ⁴Center for Medical Research Methodology, University of Copenhagen, Copenhagen, Denmark; the ⁵Department of Medicine and Surgery, Karolinska Institute, Stockholm, Sweden; the ⁶Steno Diabetes Centre and Hagedorn Research Institute, Gentofte, Denmark; the ⁷Department of Clinical Biochemistry, Rigshospitalet, Copenhagen University Hospital, Copenhagen, Denmark; the ⁸Copenhagen City Heart Study, Bispebjerg Hospital, Copenhagen University Hospital, Copenhagen, Denmark; the ⁹Copenhagen General Population Study, Herlev Hospital, Copenhagen University Hospital, Herlev, Denmark; and the ¹⁰Department of Haematology L, Herlev Hospital, Copenhagen University Hospital, Herlev, Denmark.

Corresponding author: Børge G. Nordestgaard, brno@heh.regionh.dk.

Received 2 March 2011 and accepted 19 July 2011.

DOI: 10.2337/dc11-0416

This article contains Supplementary Data online at <http://care.diabetesjournals.org/lookup/suppl/doi:10.2337/dc11-0416/-/DC1>

© 2011 by the American Diabetes Association. Readers may use this article as long as the work is properly cited, the use is educational and not for profit, and the work is not altered. See <http://creativecommons.org/licenses/by-nc-nd/3.0/> for details.

(CCHS), 0.9–130% (CGPS), and 0.1–97% (population-based case-control study). Increased transferrin saturation was detected in 355, 346, and 361 individuals in the CCHS, CGPS, and in the population-based case-control study, respectively.

Diabetes end point definitions

A combination of ICD codes, self-reported diabetes (yes/no), information

on antidiabetic medication, and a non-fasting glucose >11 mmol/L was used.

Statistics

Cox proportional hazards regression with age as timescale (i.e., left truncation) was used to estimate hazard ratios (HRs) with 95% CIs. The assumption of proportional hazards was tested with the use of Schoenfeld residuals, and no violations

were observed. Analyses not stratified for sex were adjusted for sex. In the case-control study, conditional logistic regression models were used to estimate odds ratios (ORs) with 95% CI.

RESULTS—Details are provided in the Supplementary Data online. The HRs (95% CI) for any form of diabetes in individuals with transferrin saturation of ≥50% vs.

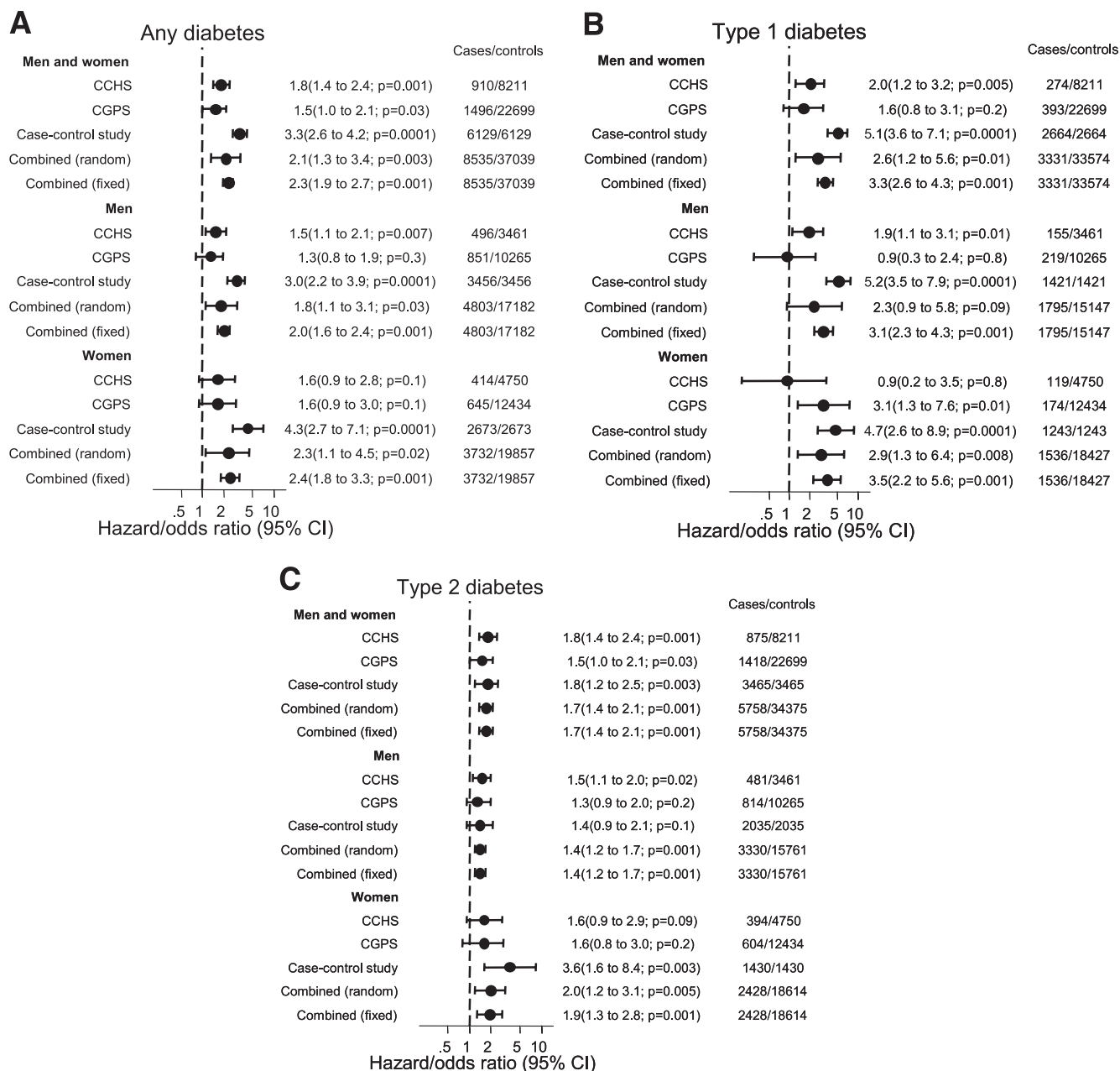


Figure 1—Risk of diabetes according to transferrin saturation ≥50% vs. <50% in the CCHS, the CGPS, a case-control study, and combined. A: Any form of diabetes. B: Type 1 diabetes. C: Type 2 diabetes. The case-control study comprised patients with diabetes from the Steno Diabetes Centre in Copenhagen, and control subjects were ascertained as in the CGPS but from a different sample than those included in the CGPS-only study. Combined, there was no overlap of individuals among these three studies, thus allowing independent confirmation of findings. 95% CIs are shown. Heterogeneity for the combined results for men and women was Q = 17 and P = 0.001 for any form of diabetes, Q = 15 and P = 0.001 for type 1 diabetes, and Q = 0.7 and P = 0.7 for type 2 diabetes.

<50% were 1.8 (1.4–2.4; $P = 0.001$) in CCHS and 1.5 (1.0–2.1; $P = 0.03$) in CGPS (Fig. 1A). In the case-control study, the OR (95% CI) for any form of diabetes in individuals with transferrin saturation of $\geq 50\%$ vs. $< 50\%$ was 3.3 (2.6–4.2; $P = 0.0001$). In the combined studies, OR in the meta-analysis under the random-effects model was 2.1 (1.3–3.4; $P = 0.003$) for any form of diabetes in those with transferrin saturation $\geq 50\%$ vs. $< 50\%$ (Fig. 1A); statistical heterogeneity was $Q = 17$ and $P = 0.001$. The ORs for type 1 and type 2 diabetes in the combined studies in the meta-analysis under the random-effects model were 2.6 (1.2–5.6; $P = 0.01$) and 1.7 (1.4–2.1; $P = 0.001$), respectively, in those with transferrin saturation $\geq 50\%$ vs. $< 50\%$ (Fig. 1B and C); statistical heterogeneity was $Q = 15$ and $P = 0.001$ for type 1 diabetes and $Q = 0.7$ and $P = 0.7$ for type 2 diabetes.

CONCLUSIONS—Details are provided in the Supplementary Data online. Here we demonstrated that transferrin saturation $\geq 50\%$ was associated with a two- to threefold increased risk of developing any form of diabetes, as well as type 1 diabetes and type 2 diabetes separately. This is the first and largest population-based study that consistently demonstrates transferrin saturation as a risk marker of any form of diabetes, and of type 1 and type 2 diabetes separately, in three independent studies. There was no overlap of individuals between the three studies, thus allowing independent confirmation of findings.

Because the risk of developing diabetes observed in this study increased with

the degree of iron overload for which there is a simple treatment (phlebotomy), 1–3% of diabetes cases in the general population and 7% of cases in a diabetes population likely could be resolved if patients received such a treatment earlier in life, equal to ~ 300 patients/million in Denmark. These findings reinforce the importance of investigating iron overload in the differential diagnosis of secondary causes of diabetes and support arguments in favor of screening for iron overload in the general population. A health-economic analysis of the cost/benefit ratio of screening for elevated iron saturation in the general or in diabetic populations may help to determine the public health consequences of this study (e.g., advice against iron supplements), and whether large prospective intervention studies should be initiated to show causality between elevated transferrin saturation and diabetes.

Acknowledgments—No potential conflicts of interest relevant to this article were reported.

C.E. contributed to the literature search and the study design, analyzed and interpreted data, wrote and edited the manuscript, and created the figures and tables. T.M.-P. contributed to the literature search and the study design, collected and interpreted data, and edited the manuscript. H.U.A. contributed to the study design, collected and interpreted the data, and edited the manuscript. A.T.-H. contributed to the study design, collected and interpreted the data, and edited the manuscript. M.F. collected the data and edited the manuscript. H.B. contributed to the study design, interpreted the data, and edited the manuscript. B.G.N. contributed to the literature search and the study design, collected the data,

analyzed and interpreted the data, edited the manuscript, and created the figures and tables.

References

1. Pietrangelo A. Hereditary hemochromatosis—a new look at an old disease. *N Engl J Med* 2004;350:2383–2397
2. Hanson EH, Imperatore G, Burke W. HFE gene and hereditary hemochromatosis: a HuGE review. *Human Genome Epidemiology*. *Am J Epidemiol* 2001;154:193–206
3. Ellervik C, Mandrup-Poulsen T, Nordestgaard BG, et al. Prevalence of hereditary haemochromatosis in late-onset type 1 diabetes mellitus: a retrospective study. *Lancet* 2001;358:1405–1409
4. Oberley LW. Free radicals and diabetes. *Free Radic Biol Med* 1988;5:113–124
5. Ellervik C, Tybjærg-Hansen A, Grande P, Appleyard M, Nordestgaard BG. Hereditary hemochromatosis and risk of ischemic heart disease: a prospective study and a case-control study. *Circulation* 2005;112:185–193
6. Ellervik C, Tybjærg-Hansen A, Nordestgaard BG. Total mortality by transferrin saturation levels: two general population studies and a metaanalysis. *Clin Chem* 2011;57:459–466
7. Cartwright GE, Edwards CQ, Kravitz K, et al. Hereditary hemochromatosis. Phenotypic expression of the disease. *N Engl J Med* 1979;301:175–179
8. Edwards CQ, Griffen LM, Goldgar D, Drummond C, Skolnick MH, Kushner JP. Prevalence of hemochromatosis among 11,065 presumably healthy blood donors. *N Engl J Med* 1988;318:1355–1362
9. Olynyk JK, Cullen DJ, Aquilia S, Rossi E, Summerville L, Powell LW. A population-based study of the clinical expression of the hemochromatosis gene. *N Engl J Med* 1999;341:718–724