



**UvA-DARE (Digital Academic Repository)**

**Results from a family and DNA based active identification programme for familial hypercholesterolaemia**

ten Asbroek, A.H.A.; Marang-van de Mheen, P.J.; Defesche, J.C.; Kastelein, J.J.P.; Gunning-Schepers, L.J.

*Published in:*  
Journal of Epidemiology and Community Health

*DOI:*  
[10.1136/jech.55.7.500](https://doi.org/10.1136/jech.55.7.500)

[Link to publication](#)

*Citation for published version (APA):*

ten Asbroek, A. H. A., Marang-van de Mheen, P. J., Defesche, J. C., Kastelein, J. J. P., & Gunning-Schepers, L. J. (2001). Results from a family and DNA based active identification programme for familial hypercholesterolaemia. *Journal of Epidemiology and Community Health*, 55(7), 500-502.  
<https://doi.org/10.1136/jech.55.7.500>

**General rights**

It is not permitted to download or to forward/distribute the text or part of it without the consent of the author(s) and/or copyright holder(s), other than for strictly personal, individual use, unless the work is under an open content license (like Creative Commons).

**Disclaimer/Complaints regulations**

If you believe that digital publication of certain material infringes any of your rights or (privacy) interests, please let the Library know, stating your reasons. In case of a legitimate complaint, the Library will make the material inaccessible and/or remove it from the website. Please Ask the Library: <https://uba.uva.nl/en/contact>, or a letter to: Library of the University of Amsterdam, Secretariat, Singel 425, 1012 WP Amsterdam, The Netherlands. You will be contacted as soon as possible.



## Results from a family and DNA based active identification programme for familial hypercholesterolaemia

A H A ten Asbroek, P J Marang-van de Mheen, J C Defesche, J J P Kastelein and L J Gunning-Schepers

*J. Epidemiol. Community Health* 2001;55;500-502  
doi:10.1136/jech.55.7.500

---

Updated information and services can be found at:  
<http://jech.bmjournals.com/cgi/content/full/55/7/500>

---

*These include:*

### References

This article cites 7 articles, 4 of which can be accessed free at:  
<http://jech.bmjournals.com/cgi/content/full/55/7/500#BIBL>

2 online articles that cite this article can be accessed at:  
<http://jech.bmjournals.com/cgi/content/full/55/7/500#otherarticles>

### Rapid responses

You can respond to this article at:  
<http://jech.bmjournals.com/cgi/eletter-submit/55/7/500>

### Email alerting service

Receive free email alerts when new articles cite this article - sign up in the box at the top right corner of the article

---

### Topic collections

Articles on similar topics can be found in the following collections

[Other Cardiovascular Medicine](#) (2039 articles)  
[Screening](#) (718 articles)

---

### Notes

---

To order reprints of this article go to:  
<http://www.bmjournals.com/cgi/reprintform>

To subscribe to *Journal of Epidemiology and Community Health* go to:  
<http://www.bmjournals.com/subscriptions/>

# Results from a family and DNA based active identification programme for familial hypercholesterolaemia

A H A ten Asbroek, P J Marang-van de Mheen, J C Defesche, J J P Kastelein, L J Gunning-Schepers

Heterozygous familial hypercholesterolaemia (FH) is a common inborn error of lipoprotein metabolism, which strongly predisposes for coronary artery disease and premature cardiac death.<sup>1</sup> In 1994, a family based active identification programme for FH was implemented in the Netherlands.<sup>2</sup> It is based on DNA diagnosis of the LDL-receptor gene mutation, which enables us to search selectively for patients in a high risk population. The programme initially targets first and second degree relatives of FH probands (diagnosed at Lipid Research Clinics throughout the country) and extends further into the family only when new patients are identified. The programme aims to identify mutation carriers and to refer them to Lipid Research Clinics for extensive individual risk assessment and, if necessary, treatment. As no carefully collected data are available for cholesterol levels among the general population of LDL-receptor gene mutation carriers, the large majority of whom are asymptomatic, we studied the prevalence of hypercholesterolaemia among screenees with a proved LDL-receptor gene mutation.

## Methods and Results

Between 1994 and 1998 2814 adults were screened. The estimated response rate was constant over the years at 90%.<sup>3–4</sup> For reasons of comparison with available population data for total serum cholesterol levels,<sup>5</sup> we selected those who were between 20 and 60 years of age (1856 screenees). Depending on the available funds in the screening programme, which were lacking in certain periods, single cholesterol measurements were taken at the time of screening. Therefore, we analysed the data of all 1005 persons who had DNA test results as well as cholesterol measurements. These were a non-selective sample of the 1856 screenees between 20 and 60 years of age. Cholesterol was measured using commercially available kits (Boehringer Mannheim, Mannheim, Germany). Genomic DNA was isolated from the leucocyte fraction of 10 ml of freshly collected blood, followed by polymerase chain reaction and restriction enzyme analysis.

From the perspective of the screening programme, the screenees that are already treated cannot be considered as new cases and they do not benefit from the screening programme in the same manner as newly identified cases. Therefore, we present the prevalence of hypercholesterolaemia among all screenees as well as the prevalence of hypercholesterolaemia among those not yet treated with HMG-CoA reductase inhibitors.

Hypercholesterolaemia was defined as either an untreated total cholesterol (TC) level above the 95th centile for age and sex in the Dutch population (table 1),<sup>5</sup> or as receiving HMG Co-A reductase inhibitors. We also show the total cholesterol distribution for the untreated screenees using conventional cut off points (<6.5, 6.5–7.9, ≥ 8 mmol/l). All LDL-receptor gene mutation carriers were heterozygotes. None of the screenees had been tested for a LDL-receptor gene mutation before.

Table 2 shows the results for the screened population.

It is evident that each age category contains LDL-receptor gene mutation carriers who do not have hypercholesterolaemia: 19.8% in all men, 32.3% in untreated men and 16.7% in women, 28.7% in untreated women. Furthermore, it is shown that the prevalence of mutation carriers among all screenees tends to be lower in the older age groups. This is probably the result of selective mortality. However, the prevalence of mutation carriers among untreated screenees is also lower in the older age groups. This is not purely the result of selective mortality but it is mainly attributable to the fact that an increasing proportion of those screened in the older age groups is already treated with cholesterol lowering drugs, and more in mutation carriers than in those without a mutation as they have generally higher cholesterol levels. This might also explain why the prevalence of hypercholesterolaemia in untreated female mutation carriers is lower in the older age groups.

Department of Social  
Medicine, Academic  
Medical Centre,  
University of  
Amsterdam, P O Box  
22660, 1100 DD  
Amsterdam, the  
Netherlands

A H A ten Asbroek  
P J Marang-van de  
Mheen  
L J Gunning-Schepers

Department of  
Vascular Medicine,  
Academic Medical  
Centre, University of  
Amsterdam  
J C Defesche  
J J P Kastelein

Correspondence to:  
Dr ten Asbroek  
([g.tenasbroek@amc.uva.nl](mailto:g.tenasbroek@amc.uva.nl))

Accepted for publication  
18 January 2001

Table 1 MORGEN Project data 1996–1997: mean (SD) total serum cholesterol (TC) as well as 95th centile (C95) in the general Dutch population by sex and age group

| Age group (y) | Men       |           |           |           | Women     |           |           |           |
|---------------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
|               | 20–29     | 30–39     | 40–49     | 50–59     | 20–29     | 30–39     | 40–49     | 50–59     |
| Number        | 589       | 890       | 1210      | 1030      | 843       | 1140      | 1450      | 1139      |
| Mean TC (SD)  | 4.4 (0.9) | 5.0 (1.0) | 5.5 (1.0) | 5.5 (1.0) | 4.6 (0.8) | 4.8 (0.8) | 5.1 (0.9) | 5.7 (1.0) |
| C95           | 5.8       | 6.7       | 7.2       | 7.1       | 6.1       | 6.3       | 6.7       | 7.4       |

Table 2 Prevalence of hypercholesterolaemia\* (HC) by sex, age group and DNA test result in all screenees and prevalence of hypercholesterolaemia, mean total serum cholesterol (TC), standard deviation (SD) and total serum cholesterol distribution, using conventional cut off values (6.5 and 8.0 mmol/l) as well as the 95th centile (C95)† in untreated screenees

|                    | 20–29 years |           |           | 30–39 years |           |            | 40–49 years |           |            | 50–59 years |           |           | All ages (20–59 years) |            |            |
|--------------------|-------------|-----------|-----------|-------------|-----------|------------|-------------|-----------|------------|-------------|-----------|-----------|------------------------|------------|------------|
|                    | All         | FH+       | FH–       | All         | FH+       | FH–        | All         | FH+       | FH–        | All         | FH+       | FH–       | All                    | FH+        | FH–        |
| <b>Men</b>         |             |           |           |             |           |            |             |           |            |             |           |           |                        |            |            |
| All‡               | 105 (100)   | 35 (33.3) | 70 (66.7) | 141 (100)   | 63 (44.7) | 78 (55.3)  | 134 (100)   | 38 (28.4) | 96 (71.6)  | 98 (100)    | 26 (26.5) | 72 (73.5) | 478 (100)              | 162 (33.9) | 316 (66.1) |
| % HC               | 39.0        | 82.9      | 17.1      | 46.1        | 76.2      | 21.8       | 35.8        | 76.3      | 19.8       | 32.7        | 92.3      | 11.1      | 38.9                   | 80.2       | 17.7       |
| Untreated          | 101 (100)   | 31 (30.7) | 70 (69.3) | 116 (100)   | 41 (35.3) | 75 (64.7)  | 109 (100)   | 17 (15.6) | 92 (84.4)  | 79 (100)    | 10 (12.7) | 69 (87.3) | 405 (100)              | 99 (24.4)  | 306 (75.6) |
| Number (%)         | 5.5 (1.4)   | 6.9 (1.3) | 4.9 (0.9) | 6.3 (1.5)   | 7.5 (1.4) | 5.7 (1.1)  | 6.2 (1.2)   | 7.4 (1.0) | 6.0 (1.1)  | 6.3 (1.0)   | 7.7 (0.7) | 6.1 (0.9) | 6.1 (1.3)              | 7.3 (1.3)  | 5.7 (1.1)  |
| Mean TC (SD)       | 81 (80.2)   | 13 (41.9) | 68 (97.1) | 68 (58.6)   | 10 (24.4) | 58 (77.3)  | 71 (65.1)   | 4 (23.5)  | 67 (72.8)  | 52 (65.8)   | 0         | 52 (75.4) | 272 (67.2)             | 27 (27.3)  | 245 (80.1) |
| TC < 6.5 (%)       | 11 (10.9)   | 9 (29.0)  | 2 (2.9)   | 34 (29.3)   | 19 (36.3) | 15 (20.0)  | 30 (27.5)   | 8 (47.1)  | 22 (23.9)  | 19 (24.1)   | 6 (60.0)  | 13 (18.8) | 94 (23.2)              | 42 (42.4)  | 52 (17.0)  |
| 6.5 ≤ TC < 8.0 (%) | 9 (8.9)     | 9 (29.0)  | 0         | 14 (12.1)   | 12 (29.3) | 2 (2.7)    | 8 (7.3)     | 5 (29.4)  | 3 (3.3)    | 8 (10.1)    | 4 (40.0)  | 4 (5.8)   | 39 (9.6)               | 30 (30.3)  | 9 (2.9)    |
| TC ≥ 8.0 (%)       | 36.6        | 80.6      | 17.1      | 34.5        | 63.4      | 18.7       | 21.1        | 47.1      | 16.3       | 16.5        | 80.0      | 7.2       | 27.9                   | 67.7       | 15.0       |
| % > C95            |             |           |           |             |           |            |             |           |            |             |           |           |                        |            |            |
| <b>Women</b>       |             |           |           |             |           |            |             |           |            |             |           |           |                        |            |            |
| All‡               | 103 (100)   | 40 (38.8) | 63 (61.2) | 168 (100)   | 60 (35.7) | 108 (64.3) | 146 (100)   | 43 (29.5) | 103 (70.5) | 110 (100)   | 31 (28.2) | 79 (71.8) | 527 (100)              | 174 (33.0) | 353 (67.0) |
| % HC               | 44.7        | 90.0      | 15.9      | 39.3        | 83.5      | 14.8       | 34.2        | 83.7      | 13.6       | 31.8        | 74.2      | 15.2      | 37.4                   | 83.3       | 14.7       |
| Untreated          | 89 (100)    | 26 (29.2) | 63 (70.8) | 146 (100)   | 40 (27.4) | 106 (72.6) | 124 (100)   | 22 (17.7) | 102 (82.3) | 89 (100)    | 13 (14.6) | 76 (85.4) | 448 (100)              | 101 (22.5) | 347 (77.5) |
| Number (%)         | 5.7 (1.7)   | 7.5 (1.5) | 5.0 (1.1) | 5.7 (1.7)   | 7.3 (1.4) | 5.1 (1.1)  | 5.9 (1.2)   | 7.3 (1.5) | 5.6 (0.9)  | 6.5 (1.1)   | 7.8 (1.4) | 6.3 (0.9) | 5.9 (1.4)              | 7.4 (1.4)  | 5.5 (1.1)  |
| Mean TC (SD)       | 65 (73.0)   | 8 (30.8)  | 57 (90.5) | 106 (72.6)  | 12 (30.0) | 94 (88.7)  | 93 (75.0)   | 7 (31.8)  | 86 (84.3)  | 45 (50.6)   | 1 (7.7)   | 44 (57.9) | 309 (69.0)             | 28 (27.7)  | 281 (81.0) |
| TC < 6.5 (%)       | 12 (13.5)   | 8 (30.8)  | 4 (6.3)   | 29 (19.9)   | 18 (45.0) | 11 (10.4)  | 26 (21.0)   | 10 (45.5) | 16 (15.7)  | 37 (41.6)   | 8 (61.5)  | 29 (38.2) | 104 (23.2)             | 44 (43.6)  | 60 (17.3)  |
| 6.5 ≤ TC < 8.0 (%) | 12 (13.5)   | 10 (38.5) | 2 (3.2)   | 11 (7.5)    | 10 (25.0) | 1 (0.9)    | 5 (4.0)     | 5 (22.7)  | 0          | 7 (7.9)     | 4 (30.8)  | 3 (3.9)   | 35 (7.3)               | 29 (28.7)  | 6 (1.7)    |
| TC ≥ 8.0 (%)       | 36.0        | 84.6      | 15.9      | 30.1        | 75.0      | 13.2       | 22.6        | 68.2      | 12.7       | 15.7        | 38.5      | 11.8      | 26.3                   | 71.3       | 13.3       |
| % > C95            |             |           |           |             |           |            |             |           |            |             |           |           |                        |            |            |

\*Hypercholesterolaemia was defined as either an untreated total cholesterol (TC) level above the 95th centile for age and sex in the Dutch population (table 1), or as receiving HMGCo-A reductase inhibitors. †95th centile for total cholesterol in the Dutch general population as observed in the MORGES study (Source: RIVM Bilthoven The Netherlands) (see table 1). ‡All = treated and untreated screenees.

## Discussion

These data have not been shown before in such a large and well defined cohort, and have important consequences for case finding strategies for inherited disorders of lipoprotein metabolism with a known molecular basis. Our study shows that if in a high risk population of yet untreated, mainly asymptomatic mutation carriers, a single TC level would be used for the diagnosis of FH rather than the current gold standard—that is, the presence of a LDL-receptor gene mutation—the diagnosis would be missed in more than a quarter of the FH patients.

As is shown by others, FH is not fully penetrant from birth onwards.<sup>6</sup> It has not been reported before, however, to which extent the genetic disorder causes hypercholesterolaemia in a population of mainly asymptomatic adult relatives of genetically diagnosed patients as is shown by this study.

The high prevalence of “lower” TC levels in our cohort may well reflect a combination of factors like patients’ adherence to low calorie diets for weight loss, intercurrent illness of infectious nature, a better general health of participants in a screening programme, or a protective genetic constitution. In addition, because cholesterol levels vary with LDL-receptor mutation,<sup>7</sup> we cannot exclude that our findings of normal cholesterol levels in LDL-receptor gene mutation carriers are the result of screening for mutations that may result in a milder than expected phenotype as reported elsewhere, albeit very unlikely. Nevertheless, even though a single measurement of cholesterol is not very reliable when assessing the individual cardiovascular disease risk, for the purpose of assessing the cholesterol values on a population level a single measurement can be used.<sup>8</sup>

The prevalence of hypercholesterolaemia in those without a LDL-receptor gene mutation is higher than in the general population, which indicates that in this high risk population probably also other factors than the LDL-receptor gene mutation contribute to the prevalence of hypercholesterolaemia.

The importance of our findings depends largely on whether patients with a LDL-receptor gene mutation but without hypercholesterolaemia, experience an increased risk of coronary heart disease and whether they need the same rigorous treatment as other FH patients. This is currently unknown. As damage to the vascular wall in FH patients is probably the result of the number of cholesterol years,<sup>9</sup> a single TC level below the 95th centile could be falsely reassuring. Comprehensive appraisal of cardiovascular disease risk and cholesterol screening at regular intervals is advised. In a follow up study, we plan to further explore explanations for our findings and to assess whether the risk of coronary heart disease is increased in this group of mutation carriers in whom FH does not seem to be fully penetrant.

A H A ten Asbroek was the main author and carried out the analyses and interpreted data. P J Marang-van de Mheen

formulated the research questions, assisted in writing, analyses and interpreting results. J C Defesche participated in data screening and cleaning, assisted in the analyses as well as in the interpretation of the data and assisted in writing the report. J J P Kastelein assisted in writing and interpreting results. L J Gunning-Schepers was principal investigator, formulated the research questions, assisted in writing the report and is head of the study group.

Funding: this study is funded by the Health Research and Development Council (grant number 28-2751).  
Conflicts of interest: none.

- 1 Anonimus. Risk of fatal coronary heart disease in familial hypercholesterolaemia. Scientific Steering Committee on behalf of the Simon Broome Register Group. *BMJ* 1991;**303**:893-6.
- 2 Umans-Eckenhausen MA, Defesche JC, Scheerder RL, et al. [Tracing of patients with familial hypercholesterolemia in the Netherlands]. [In Dutch]. *Ned Tijdschr Geneesk* 1999;**143**:1157-61.

- 3 Stichting Opsporing Erfelijke Hypercholesterolemie SrOEH [Foundation for Tracing Hereditary Hypercholesterolemia]. Annual report 1997. Amsterdam: SrOEH, 1998.
- 4 Stichting Opsporing Erfelijke Hypercholesterolemie SrOEH [Foundation for Tracing Hereditary Hypercholesterolemia]. Annual report 1998. Amsterdam: SrOEH, 1999.
- 5 Blokstra A. [The Project Monitoring Risk Factors and Health, The Netherlands] (MORGEN-project). Annual report 1996. Report no 263200 006. Bilthoven: Rijksinstituut voor Volksgezondheid en Milieu, 1997.
- 6 Leonard JV, Whitelaw AG, Wolff OH, et al. Diagnosing familial hypercholesterolaemia in childhood by measuring serum cholesterol. *BMJ* 1977;**1**:1566-8.
- 7 Humphries SE, Galton D, Nicholls P. Genetic testing for familial hypercholesterolaemia: practical and ethical issues. *QJM* 1997;**90**:169-81.
- 8 Verschuren WMM. Blood cholesterol: a public health perspective. Wageningen: Ponsen and Looijen, 1995.
- 9 Hoeg JM, Feuerstein IM, Tucker EE. Detection and quantitation of calcific atherosclerosis by ultrafast computed tomography in children and young adults with homozygous familial hypercholesterolemia. *Arterioscler Thromb* 1994;**14**:1066-74.



Want to know more?

### Data supplements

Limited space in printed journals means that interesting data and other material are often edited out of articles; however, limitless cyberspace means that we can include this information online.

Look out for additional tables, references, illustrations.

[www.jech.com](http://www.jech.com)