**FinDonor 10 000 study: A cohort to identify iron depletion and factors affecting it in Finnish blood donors**

Short title: Finnish blood donor cohort

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## Abstract

### Background and Objectives

There is increasing evidence that frequent blood donation depletes the iron stores of some blood donors. The FinDonor 10 000 study was set up to study iron status and factors affecting iron stores in Finnish blood donors. In Finland, iron supplementation for at-risk groups has been in place since the 1980's.

### Material and Methods

2584 blood donors (N= 8003 samples) were recruited into the study alongside the standard donation at three donation sites in the capital region of Finland between 5/2015 and 12/2017. All participants were asked to fill out a questionnaire about their health and lifestyle. Blood samples were collected from the sample pouch of whole blood collection set, kept in cool temperature and processed centrally. Whole blood count, CRP, ferritin and sTFR were measured from the samples and DNA was isolated for GWAS studies.

### Results

Participant demographics, albeit in general similar to the general blood donor population in Finland, indicated some bias toward older and more frequent donors. Participation in the study increased median donation frequency of the donors. Analysis of the effect of time lag from the sampling to the analysis and the time of day when sample was drawn revealed small but significant time-dependent changes.

### Conclusion

The FinDonor cohort now provides us with tools to identify potential donor groups at increased risk of iron deficiency as factors explaining this risk. The increase in donation frequency during the study suggests that scientific projects can be used to increase the commitment of blood donors.

**Introduction**

Approximately 200 - 250 mg of iron is drawn in a standard blood donation; the amount accounts for 25% of average tissue iron stores in men and up to 75% in women. There is compelling evidence that a portion of blood donors may become iron depleted or deficient. Deferral rates of presenting donors, due to too low level of hemoglobin, vary a lot depending on populations and policies from below 10 % to up to 20 %. In addition, for example, in the U.S. 35% of frequent blood donors were found to be iron deficient [1–6]. Adverse health effects, with various symptoms, such as fatigue, pica, restless legs syndrome and cognitive problems, have been linked to iron deficiency and anemia [7], though the effect of iron deficiency without anemia in otherwise healthy individuals is still unclear [8]. The iron removed by blood donation should be replaced by dietary iron. Another tool to ensure correction of iron stores is the minimum time interval between blood donations, which must be balanced between donor health issues and blood demand. There is evidence [2,4,9] that the current intervals may not be sufficient for iron or Hb recovery, at least for some donors. Genome screening studies [10,11] on various populations, but so far not directly on blood donors, have indicated a number of genes associated with red blood cell indices. The relevance of these results to the response to frequent blood donation is unknown. Identification of donors who are at highest risk for developing iron depletion or anemia as a result of frequent blood donation is warranted as it could give us fact-based tools to steer donor recruitment.

Implementation of iron supplementation to an entire blood donor population to manage iron deficiency is rare, although several countries provide some iron supplementation [12]. The Finnish Red Cross Blood Service has implemented a national risk-group based iron supplementation policy since the 1980's. Iron tablets are provided at donation time to all female donors under the age of 50 and to all other donors donating every 4th month or more frequently.

To complement the above mentioned studies on iron depletion and cohorts collected elsewhere [13–16], we set up a cohort study focusing primarily on the measurement of iron stores, factors affecting them, and, in addition, on the effect of systematic iron supplementation. The study recruited in three blood donation sites in the Helsinki metropolitan area. Any blood donor in these sites was qualified to participate. Based on our sociological study, Finnish blood donors are willing to donate also for research use [17]. In general, blood donors have been regarded as an important resource for biomedical studies [18]. Consent for laboratory and genetics analyses was requested and participants were asked to fill in a health and lifestyle questionnaire during the first donation. We here describe the cohort and give evidence that there are factors in the measurements that may affect the results if not carefully taken into account.

**Materials and methods**

The Finnish Red Cross Blood Service (FRCBS) is a national blood establishment responsible for the collection, processing and distribution of blood products in Finland. An Ethical review and approval for the study was obtained from the Ethical Board of Helsinki University Hospital, Helsinki, Finland. Informed consent was obtained from participants and they were allowed to withdraw from the study at any time.

**Sample collection**

The Fin Donor 10 000 study ran from May 18th 2015 to December 8th 2017. All donors in the participating three fixed donation sites located in the Helsinki metropolitan area (Kivihaka, Sanomatalo, Espoo) were informed about the possibility to join the study with leaflets, posters and Facebook–postings. A permanent deferral was the only criteria for exclusion from the study. The age limit for blood donation and thus for study participation was 18-70 years (18-59 years for first time donors).

All donations and donor registrations were recorded in a single database (e-Progesa, MAK-SYSTEM). Each participant was asked to fill out a questionnaire on health and life style issues using an internet portal provided at the donation site. The questions are provided in Supplementary Table 1 and are similar to those used e.g. in the Danish blood donor studies [19].

Samples of peripheral blood were collected from the diversion pouch (CompoFlow® Quadruple T&B, Fresenius Kabi, Germany). For blood counts and genomic DNA extraction, samples were collected in 3 ml K2-EDTA tubes (Terumo Europe NV, Belgium) and for CRP, ferritin and sTfR measurements in 3 ml Lithium heparin tubes (Terumo Europe NV, Belgium). A cell pellet was stored for genomic DNA preparation. Samples were sent with the signed informed consent to FRCBS headquarters, located in Kivihaka, Helsinki, Finland, where donor identity and coding on the tubes were double-checked and added to the research database. Plasma samples were sent for laboratory measurements (whole blood count, CRP, ferritin and sTfR; see Supplementary Table 2 for further details) in batches twice a day to the clinical laboratory of the University of Helsinki Central Hospital. Blood counts were measured with Sysmex XE (Sysmex, Japan). The measurements are accredited by the local authority FINAS and the laboratory participates in national and international quality assurance rounds. The tubes were kept in cool storage between transportation and handling and time points from blood draw to analysis were recorded.

**Genotyping**

The first 760 enrolled individuals were genotyped using an Illumina HumanCoreExome-24v1-1_A beadchip (Illumina, USA) to estimate ethnic diversity and relatedness. Genotyping of the rest of the participants is under way. Genotyping was performed at the FIMM Technology Centre, Helsinki, Finland. DNA samples from the Finnish cohort were extracted using the QIAamp DNA Blood Mini Kit (Qiagen, Germany). Imputation was carried out as in [20] according to instructions of [21] using plink 1.90b3.29 [22] for quality filtering and IMPUTE2 [23] for the actual imputation with 1000 Genomes Phase 3 [24] as a phased reference. Post-imputation filtering excluded variants having an IMPUTE2 INFO-field <0.5. After post-imputation filtering 11 880 443 variants were included. PCA (Principal Component Analysis) of the imputed and filtered dataset was carried out with eigensoft version 6.1.4 [25]. The datasets analyzed during the current study are not publicly available due to limitations of the ethical permits which do not allow distribution of personal data, including individual genetic and clinical results.

**Blood count data analysis**

The effects of donation time and sampling-to-analysis delay were analyzed using generalized linear multiple regression. Blood count measurement data were used as response variables and donation time, sampling-to-analysis delay, sex and age group (ten year bins) were used as explanatory variables. CRP and ferritin were log2-transformed. Graphical examination of the data suggested interactions between the explanatory factors donation time and sampling-to-analysis delay. Hence, first a model that includes an interaction term for them was tested and if the interaction was not found significant a model without an interaction term was fitted to calculate coefficients of the explanatory variables. The significance of p-values was estimated with Bonferroni correction for multiple testing. The effect of repeated measures was tested by fitting mixed effect models using lme4 R-library [26]. Mixed effect models were identical to the no-interaction models with an additional explanatory variable of individuals as a random effect. Qualitatively similar effect sizes were obtained from the mixed effect models and the no-interaction models. The data were analyzed using R version 3.4.3 [27] and plotted with the ggplot2 R-library [28].

**Results**

In total 2584 individual donors (1015 men, 1569 women) gave consent to participate in the study and donated at least one sample. This represent 6% of all donors who visited the study sites during the study. Of them 2562, (99%; 1004 men, 1558 women) answered the questionnaire. In total, participants donated 8003 samples. None of the participants has withdrawn his or her consent so far (12/2018). 174 were new or reactivated (previous donation more than 2 years before) donors (see Supplementary Table 3 for further details).

Summary statistics of the age, haemoglobin levels, donation interval, count of donations and blood types in the study population were compared to those of the entire national donor population (Group "All" in Table 1 and Figures 1 and 2) and to those of the donor population of the recruitment sites (Group "Study Sites" in Table 1 and Figures 1 and 2). These three groups were found to differ. The median age of study participants in both sexes was higher than both the national and recruitment site populations, but closer to the national population. The age distribution was bimodal and again the study population resembled more the national donor population than the donor population of the study sites (Figure 1A).

**The median donation interval during the study for both sexes was lower than in the national and recruitment site populations.** Accordingly, donation counts were higher (Table 1 and Figure 2). Mean Hb (g/l) was lower in men in the study population than in the national and recruitment site populations (Table 1 and Figure 1B). However, in women, mean Hb (g/l) in the study population was higher than in the recruitment population, which reflects the higher age of the study population. Consequently, the study population consisted of a markedly higher proportion of older donors with frequent donation activity and fewer new donors. The distribution of blood types between different populations resembled each other closely except that the study population has been more comprehensively blood typed due to their frequent donor status (Table 1).

**The during-study donation attempt frequency was found to be higher than the before-study donation attempt frequency for study participants** (median of during-study – before-study frequency = 0.6, p < 2e-44 for men; 0.6, p< 9e-80 for men, Wilcoxon signed rank test, Figure 3). For each donor, the before-study donation frequency was calculated as the annual average count of donations for the two years before study enrollment. The during-study frequency was calculated as the annual average count of donations from the time of enrollment to the study end. Donors who enrolled after 2016.12.08 (less than one year before the end of the study) and new donors were excluded. 2271 (906 men and 1365 women) donors were therefore included in the analysis.

At least two blood samples could be collected from 1976 donors (N= 840 men, N= 1136 women, see Supplementary Table 4 for further details). Multiple samples from the same donors are valuable for longitudinal studies, for example, in serum biomarker or metabolomics studies.

DNA samples from 760 participants were subjected to a genome-wide SNP array screen. To demonstrate the genetic homogeneity of the study population, we performed principal

component analysis (Supplementary Figure 1) that demonstrated that all 760 participants could be classified as Northern Europeans.

For each blood-count test, the sampling time (Figure 4) as well as the time when the analysis result was ready were entered to the laboratory information system. The difference between these two time points was used as the sampling-to-analysis delay time (Figure 5). To assess whether the trends shown in Figures 4 and 5 were significant, a multiple regression model was fitted for each individual blood count measurement including both donation time, sampling-to-analysis delay, an interaction term between these two variables, and sex and age as explanatory factors (Table 2). The interaction term was used to evaluate if the two time variables could be separated from each other in this cohort. If the interaction term was not found to be significant then the coefficient and p-value of the time variables were calculated from a model without the interaction term. While the coefficient from a model without the interaction term is the unit change in an hour, e.g. Hb drops on average 0.2 g/l per hour when going from 8 a.m. to 8 p.m., there is no such intuitive interpretation for coefficients from a model with an interaction term. An effect in same direction and very similar size range has been shown previously from finger prick Hb measurements [29].

**For hematocrit and platelet count there was a separable significant sampling-to-analysis delay time (h) effect and for erythrocyte count, Hb, hematocrit, MCH and leukocyte count there was a separable significant donation time (h) effect.** For ferritin, MCV, MCHC, RDW and sTfR there were significant interactions of donation time (h) and sampling-to-analysis delay time (h) and hence their individual effects could not be separated.

A single elevated CRP measurement, i.e. a value above 10 that may point to inflammation, was found in 1.8 % of women and 0.3 % of men. Respectively, 14.3 % and 7.6 % had a single measurement between 3 and 10, i.e. a low grade inflammation. To investigate if these measurements could point to chronic inflammation we investigated the data graphically (Supplementary Figure 2). No donor was found to have two CRP measurements above 10, hence, they most likely had no chronic inflammation.

**Discussion**

The present cohort was collected primarily for studies investigating the effects of regular blood donation on donors' health, in particular iron stores and factors regulating them in a population with a long-stading systematic iron replacement policy. The cohort complements those collected elsewhere [14–16,19]. We put special emphasis on careful collection and follow-up of plasma and serum samples to understand possible confounding factors in measurements of iron markers. Modelling blood count and related data is not simple as HCT, MCHC, MCV, RDW

and CRP show properties of counts instead of true continuous measurements, hence, defying the assumptions of standard linear regression. To mitigate this, we experimented with various transformations, link functions and error distributions in the modelling process to no avail. The results (Figures 4 and 5, Table 2) clearly demonstrate that it is essential to know the detailed conditions of collected samples if accurate results are assumed. There are interesting studies demonstrating large effects related to the time of drawing the samples [30] and this variation also is relevant in everyday blood donation practice [29]. However, for many screening purposes the variation in measurements found in the present study is apparently not critical.

Another factor that is essential to know in cohort studies is how well the study population represents the overall target population, here blood donors. Based on the demographics, Hb distribution and donation activity, participants (Figures 1 and 2, and Table 1) of the FinDonor 10 000 cohort can be regarded in general to represent Finnish donor pool despite some bias toward frequent and committed donors. As a key question to be clarified with the cohort is related to effects of frequent donation, this bias may not have serious drawbacks in our future studies.

In Finland all blood donations are voluntary and non-remunerated. Such a blood donation system could be sensitive to any additional burden to donors (e.g., participating to research). However, as suggested by two previous studies [15,31], it was found that the donation frequency of study participants, in fact, increased during the present study (Figure 3). It is of note that no additional efforts to remind participants to donate were made. Hence, research projects can be even seen as a way to increase the commitment to donate among blood donors.

The prevalence of CRP between 3 and 10 was very similar to what has been reported in the Danish blood donor population (14.4% women and 6.1% men [32]). The prevalence of CRP > 10 was found to be below 2%. Approximately 7% of the apparently healthy Finnish adults have earlier been reported to have CRP levels over 10 [33,34]. As a minor CRP elevation (between 3 and 10) does not directly point to inflammation but rather to various mild tissue stress or injury [35], the Fin Donor 10 000 cohort appears to have a markedly low level of chronic inflammation. This is in accordance with the healthy donor effect, [36] i.e. healthier individuals are selected as donors and are able to maintain the habit from years to decades

**Conclusions**

Together with similar cohorts from other populations [14–16,19] the FinDonor cohort provides us with tools to identify the potential donor groups at risk for iron deficiency and the genetic and non-genetic factors associated with this increased risk, as well as to study other health effects

of blood donation. These background facts are needed for ensuring safer and more personalized blood donation in the future.

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## Author contributions:

Study concept: JP and JC;

Management and planning of recruitment and sample processing: PM, PN, EP and NN;

Data processing and analysis: AL, ML and MA;

Interpretation of results: PN, ML, EP, JC, MA, JP;

Manuscript preparation: JP and MA.

## Disclosure of conflicts of interest:

Nothing to disclose

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**Figure and Table Legends**

**Table 1 Characteristics of the study population.** Comparison to national (All) and study recruitment site (Study sites) donor populations during the timespan of the study. To facilitate comparison to other cohorts, the count of donations has been normalized from the study period (2.5 years) to 2 years.

**Table 2 Significance of explanatory variables for each measurement.** For each variable a model was fitted with age, sex, sampling-to-analysis delay (h) and donation time (h) as explanatory variables. A model including an interaction term was first fitted ("p-value from a model with interaction"). If the interaction term was not significant, coefficient and p-value is provided for a model without interaction. P-values that are significant after Bonferroni correction of alpha (original alpha 0.05) are marked with an asterix (*).

**Figure 1: Age and Hb distributions of study participants.** Density function of A) age distributions of all donors ("All"), all donors that donated in the collection sites where study took place ("Study sites") and study participants ("Study") during the study period. Respectively the density distribution of B) hemoglobin. Average over all donation events during the study period is used as the value of age and hemoglobin. Mean of hemoglobin was "All": 140 women and 154 men; "Study sites" 138 and 154 and "Study": 138 and 151.

**Figure 2**: **Counts of donations of study participants during study period.** "All" are donations by all donors that donated during the study at any FRCBS site. "Study sites" are donations of all donors in the collection sites where study took place. "Study" are study participants. Y-axis shows the proportion of the donors in the group (All, Study sites, Study) that donated a certain number of donations during the study period (on x-axis).

**Figure 3: Difference of during study donation attempt frequency and before study donation attempt frequency.** Before study donation frequency for a donor was calculated as the annual average count of donations from two years before the study enrollment. During study frequency was calculated as the annual average count of donations from donors who enrolled on 2016.12.08 or earlier from the time of enrollment to the study end.

**Figure 4: Effect of donation time to measurement values.** Data is binned to hexagons and the color of each hexagon shows how many individual measurements are contained in each hexagon. The black line shows a linear regression trend line. See Table 2 for full description of measured variables.

**Figure 5: Effect of sampling-to-analysis delay time (d) to measurement values.** Data is binned to hexagons and the color of each hexagon shows how many individual measurements are contained in each hexagon. The black line shows a linear regression trend line.

## Figures and Tables

**Table 1 Characteristics of the study population**

|  | Men | | | Women | | |
|---|---|---|---|---|---|---|
|  | Study | Study sites | All | Study | Study sites | All |
| n | 1015 | 16492 | 83150 | 1569 | 26390 | 120867 |
| Age (years, median [IQR]) | 49.00 [35.00, 58.00] | 39.00 [28.00, 53.00] | 43.00 [29.00, 55.00] | 43.00 [29.00, 55.00] | 33.00 [25.00, 48.00] | 38.00 [26.00, 52.00] |
| Hb (g/l, mean (sd)) | 151.66 (8.33) | 153.47 (9.85) | 154.39 (11.32) | 137.43 (8.00) | 136.77 (9.53) | 138.76 (10.92) |
| Donation interval (days, median [IQR]) | 107.50 [84.00, 165.00] | 159.00 [85.50, 414.00] | 174.50 [91.00, 465.00] | 162.25 [115.00, 259.75] | 210.50 [107.75, 518.00] | 223.00 [116.00, 572.00] |
| Donations during 2 years (%) |  |  |  |  |  |  |
| 1-3 | 331 (32.6) | 13324 (80.8) | 67091 (80.7) | 900 (57.4) | 23954 (90.8) | 107552 (89.0) |
| 4-6 | 302 (29.8) | 1794 (10.9) | 9738 (11.7) | 523 (33.3) | 2003 ( 7.6) | 11411 ( 9.4) |
| 7-9 | 266 (26.2) | 1001 ( 6.1) | 5005 ( 6.0) | 144 ( 9.2) | 403 ( 1.5) | 1871 ( 1.5) |
| 10+ | 116 (11.4) | 373 ( 2.3) | 1316 ( 1.6) | 2 ( 0.1) | 30 ( 0.1) | 33 ( 0.0) |
| Blood type (%) |  |  |  |  |  |  |
|  | 0 ( 0.0) | 685 ( 4.2) | 3638 ( 4.4) | 2 ( 0.1) | 1575 ( 6.0) | 6598 ( 5.5) |
| A- | 69 ( 6.8) | 981 ( 5.9) | 4719 ( 5.7) | 121 ( 7.7) | 1642 ( 6.2) | 7083 ( 5.9) |
| A+ | 365 (36.0) | 5447 (33.0) | 28205 (33.9) | 546 (34.8) | 8257 (31.3) | 38893 (32.2) |
| AB- | 5 ( 0.5) | 148 ( 0.9) | 901 ( 1.1) | 19 ( 1.2) | 272 ( 1.0) | 1317 ( 1.1) |
| AB+ | 70 ( 6.9) | 926 ( 5.6) | 4792 ( 5.8) | 64 ( 4.1) | 1379 ( 5.2) | 6931 ( 5.7) |

FinDonor study cohort

| | | | | | | |
|---|---|---|---|---|---|---|
| B- | 21 ( 2.1) | 423 ( 2.6) | 1997 ( 2.4) | 49 ( 3.1) | 624 ( 2.4) | 2968 ( 2.5) |
| B+ | 134 (13.2) | 2074 (12.6) | 11059 (13.3) | 191 (12.2) | 3384 (12.8) | 15942 (13.2) |
| O- | 72 ( 7.1) | 1216 ( 7.4) | 5083 ( 6.1) | 132 ( 8.4) | 1917 ( 7.3) | 7808 ( 6.5) |
| O+ | 279 (27.5) | 4592 (27.8) | 22756 (27.4) | 445 (28.4) | 7340 (27.8) | 33327 (27.6) |
| BMI (cm², median [IQR]) | 25.83 [23.91, 28.31] | | | 24.62 [22.31, 28.12] | | |
| Weight (kg, mean (sd)) | 85.66 (14.60) | | | 71.77 (14.70) | | |
| Height (cm, mean (sd)) | 179.82 (6.26) | | | 166.64 (5.92) | | |

**Table 2 Significance of explanatory variables for each measurement.**

| Measurement | Number of measurements | Coefficient from a model without interaction | | p-value from a model without interaction | | p-value from a model with interaction | | |
|---|---|---|---|---|---|---|---|---|
| | | Sampling-to-analysis delay (h) | Donation time (h) | Sampling-to-analysis delay (h) | Donation time (h) | Sampling-to-analysis delay (h) | Donation time (h) | Sampling-to-analysis delay : Donation time |
| log2(CRP mg/l) - C-Reactive Protein | 7906 | 1.000 | 0.999 | 7.2E-01 | 6.2E-01 | 3.1E-01 | 2.8E-01 | 3.2E-01 |
| Eryt (x $10^{12/l}$) -Erythrocyte count | 7909 | -0.001 | -0.011 | 7.9E-03 | 9.7E-11 * | 1.4E-01 | 8.9E-01 | 6.8E-02 |
| log2(Ferrit µg/l) - Ferritin | 7902 | | | | | 2.6E-04 * | 2.1E-04 * | 1.1E-03 * |
| Hb (g/l) - Haemoglobin | 7910 | -0.031 | -0.178 | 2.7E-02 | 7.6E-05 * | 2.9E-01 | 7.8E-01 | 1.9E-01 |
| HKR (%) - Hematocrit | 7919 | 0.028 | -0.083 | 2.1E-14 * | 8.7E-12 * | 2.2E-01 | 1.3E-04 * | 3.0E-02 |
| Leuk (x $10^9$/l) - Leukocyte count | 7909 | 0.004 | 0.108 | 9.3E-02 | 2.6E-41 * | 3.1E-02 | 1.1E-07 * | 4.9E-02 |
| MCH (pg / cell) - Mean corpuscular hemoglobin | 7918 | 0.003 | 0.028 | 2.6E-01 | 1.4E-03 * | 3.4E-01 | 8.0E-01 | 2.7E-01 |
| MCHC (g/l) - Mean corpuscular hemoglobin concentration | 7910 | | | | | 8.4E-06 * | 6.9E-15 * | 2.8E-12 * |
| MCV (fl) - Mean corpuscular volume | 7914 | | | | | 3.0E-04 * | 4.1E-06 * | 2.0E-07 * |
| RDW % - Red cell Distribution Width | 7920 | | | | | 7.8E-03 | 1.3E-04 * | 1.5E-03 * |
| sTfR (mg/l) - soluble Transferrin Receptor | 7646 | | | | | 2.1E-04 * | 5.8E-05 * | 4.8E-05 * |
| Trom (x 109/l) - Thrombocyte count | 7911 | 0.240 | 0.430 | 1.5E-03 * | 8.0E-02 | 9.7E-01 | 9.8E-01 | 6.7E-01 |

**FIGURES**

**Figure 1: Age and Hb distributions of study participants.**

**Figure 2**: **Counts of donations of study participants during study period.**

**Figure 3: Difference of during study donation attempt frequency and before study donation attempt frequency.**

**Figure 4: Effect of donation time to measurement values.**

**Figure 5: Effect of sampling-to-analysis delay time (d) to measurement values.**