Erythrocyte membrane trans-fatty acid index is positively associated with a 10-year CHD risk probability

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Abstract

Industry-generated trans-fatty acids (TFA) are detrimental to risk of CHD, but ruminant-originated TFA have been reported as neutral or equivocal. Therefore, the total TFA amount should not be the only factor considered when measuring the effects of TFA. In the present study, we addressed whether a version of the TFA index that unifies the effects of different TFA isomers into one equation could be used to reflect CHD risk probability (RP). The present cross-sectional study involved 2713 individuals divided into four groups that represented different pathological severities and potential risks of CHD: acute coronary syndrome (ACS, n 581); chronic coronary artery disease (CCAD, n 631); high-risk population (HRP, n 659); healthy volunteers (HV, n 842). A 10-year CHD RP was calculated. Meanwhile, the equation of the TFA index was derived using five TFA isomers (trans-16:1n-7, trans-16:1n-9, trans-18:1n-7, trans-18:1n-9 and trans-18:2n-6n-9), which were detected in the whole blood, serum and erythrocyte membranes of each subject. The TFA index and the 10-year CHD RP were compared by linear models. It was shown that only in the erythrocyte membrane, the TFA isomers were significantly different between the groups. In the ACS group, industry-generated TFA (trans-16:1n-9, trans-18:1n-9 and trans-18:2n-6n-9) were the highest, whereas ruminant-originated TFA (trans-16:1n-7 and trans-18:1n-7), which manifested an inverse relationship with CHD, were the lowest, and vice versa in the HV group. The TFA index decreased progressively from 7·12 to 6·90 (𝑝< 0·001) and manifested a strong linear correlation, which might reflect the true pathological severity of CHD.

Key words: Trans-fatty: acid index: CHD: Erythrocyte membranes

The effect of blood trans-fatty acid (TFA) levels on human diseases has recently aroused considerable attention(1–3). Chavarro et al.(4) reported that the whole-blood TFA level was associated with an increased risk of non-aggressive prostate tumour. Chajes et al.(5) showed that a high serum level of TFA contributed to the risk of invasive breast cancer in women. Benatar et al.(6) proposed that plasma total TFA may be associated with vascular disease and increased C-reactive protein in patients with severe coronary artery disease. Meanwhile, Lemaitre et al.(7) found that a high erythrocyte TFA index level was correlated with an increased risk of sudden cardiac arrest.

It is well accepted that TFA are highly associated with CHD risk, and different kinds of TFA isomers play different roles in CHD events(8,9). Industry-originated TFA, such as trans-16:1n-9, trans-18:1n-9 and trans-18:2n-6n-9, are considered to have deleterious effects on cardiovascular health(2,10,11), while TFA from ruminant sources are associated with a slightly neutral risk(12), because trans-16:1n-7 and trans-18:1n-7, the dominant TFA isomers in milk, can be biotransformed to conjugated linoleic acid (CLA) such as 9-cis, 11-trans-18:2n-6 CLA by Δ9-desaturase, which is conducive to anticancer, anti-diabetic and anti-CHD effects(13). Therefore, the effects of TFA on CHD vary in differently originated TFA, and it may be irrational to estimate the effects of TFA simply by considering the total amount of TFA levels. However, if there were combined parameters, taking into account both industry-originated TFA and ruminant-sourced TFA, the effects

Abbreviations: ACS, acute coronary syndrome; CCAD, chronic coronary artery disease; CLA, conjugated linoleic acid; HRP, high-risk population; HV, healthy volunteers; RP, risk probability; TFA, trans-fatty acids.

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of TFA on CHD could be estimated according to different TFA isomers and their own physico-chemical characteristics.

In the present study, we explored a TFA index that unifies the disparate impacts of different TFA isomers on CHD by discriminating the different effects into one equation. Meanwhile, a 10-year CHD risk probability (RP) was also calculated in the present cross-sectional study to represent different pathological severities and potential risks of CHD. The index may be used to confirm the presence of a true association between TFA and CHD. The aims of the present study were to verify whether the TFA index was associated reliably with the 10-year CHD RP and to identify which study sample (whole blood, serum or erythrocyte membrane) could truly reflect the change in body TFA levels.

Methods

Ethical statement

The present study was conducted according to the guidelines laid down in the Declaration of Helsinki, and all procedures involving human subjects/patients were approved by the Institutional Ethics Committee, the First and Second Affiliated Hospital, Nanchang University, China. Written informed consent was obtained from all subjects/patients.

Patients

In the present study, we divided CHD into two relatively independent clinical pathological forms: a severely harmful acute coronary syndrome (ACS), which is manifested by myocardial infarction and unstable angina, and a chronic/stable form known as chronic coronary artery disease (CCAD), which has a phenotype of stable angina and ischaemic cardiomyopathy. The ACS and CCAD patients were selected to represent an acute and a chronic CHD pathology, respectively. A high-risk population (HRP) and healthy volunteers (HV) served as controls for CHD patients in the ACS and CCAD groups. Thus, the four groups represented different pathological severities and potential risks of CHD.

A cross-sectional survey on six large physical populations of 12,880 individuals was conducted at the First and Second Affiliated Hospitals, Nanchang University between 2007 and 2010. Excluding confounding factors and other diseases, 2713 individuals were screened in the ACS (n=81), CCAD (n=659), HRP (n=659) and HV (n=842) groups to represent different pathological severities and potential risks of CHD. The ACS was diagnosed on the basis of pre-specified criteria (3) atrial fibrillation; (4) malignant disease; (5) dialysis; (6) diabetes mellitus; (7) renal insufficiency.

Blood sample preparation

After an overnight fasting of the participants, blood (5 ml) was drawn from the antecubital vein into vacutainer tubes containing ethylene diamine tetraacetic acid. Serum was separated from erythrocytes by centrifugation at 2000 g at 4°C for 10 min. Erythrocytes were washed three times with ice-cold isotonic saline to remove the buffy coat. Membranes were isolated by a modification of Burton's method. Briefly, packed cells were lysed with cold distilled water, centrifuged at 20,000 g at 10°C for 20 min, and washed several times to eliminate Hb residues. Whole-blood, serum and erythrocyte membrane samples from each participant were stored at −80°C under liquid N2 until lipid extraction.

Extraction and analysis of fatty acids

Lipids were extracted by chloroform–methanol (1:1) and methylated by sodium methoxide as described previously. Fatty acid methyl esters were analysed by a gas chromatograph (GC 6890 N; Agilent) equipped with a flame ionisation detector, an autosampler injector (7683-B; Agilent) and a fused silica capillary column (CP-Sil 88, 100 m x 0.25 mm inner diameter, 0.20 μm film thickness; Varian). The temperature was held at 45°C for 4 min, and then ramped to 175°C at a flow rate of 13°C/min, held for 27 min, and, finally, increased to 215°C at a flow rate of 4°C/min, held for 35 min.

The levels of five TFA (trans-16:1n-7, trans-16:1n-9, trans-18:1n-7, trans-18:1n-9 and trans-18:2n-6n-9) were calculated from the GC results using normalisation and internal standard methods, as described previously. The results were compared with standard fatty acid methyl esters (GLC-465; Nu-Chek Prep, Inc.), with 21:0 fatty acid methyl esters added. Fatty acid profiles were expressed as the percentage of TFA. The TFA index was defined by the following equation:

\[
\text{TFA index} = \exp \left[ \frac{\sum \text{trans-16:1n-9} + \text{trans-18:1n-9} + \text{trans-18:2n-6n-9}}{\text{trans-16:1n-7} + \text{trans-18:1n-7}} \right].
\]

In this equation, the numerator adds the levels of industrial TFA (trans-16:1n-9, trans-18:1n-9 and trans-18:2n-6n-9) together, and the denominator adds the levels of ruminant TFA (trans-16:1n-7 and trans-18:1n-7) together. An exponential function was used to make the index stronger and more obvious as the TFA index.

Characteristic data

Serum lipids (TAG, total cholesterol and HDL), apoA (mainly included apoA1), apoB (included both apoB100 and apoB48) and lipoprotein α were measured in the hospital clinical laboratory with an automatic biochemistry analyser (Beckman...
Table 1. Characteristics of study participants (n 2713)
(Mean values and standard deviations; number of patients and percentages)

<table>
<thead>
<tr>
<th></th>
<th>ACS (n 581)</th>
<th>CCAD (n 631)</th>
<th>HRP (n 659)</th>
<th>HV (n 842)</th>
<th>P</th>
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</thead>
<tbody>
<tr>
<td><strong>Age (years)</strong></td>
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<td>56·78</td>
<td>48·21a,b</td>
<td>52·99b</td>
<td>46·93a</td>
</tr>
<tr>
<td></td>
<td>sd</td>
<td>19·78</td>
<td>8·23</td>
<td>12·11</td>
<td>9·34</td>
</tr>
<tr>
<td><strong>Male</strong></td>
<td>n</td>
<td>371</td>
<td>63·86a</td>
<td>412</td>
<td>65·29b</td>
</tr>
<tr>
<td></td>
<td>%</td>
<td>48·21a,b</td>
<td>8·23</td>
<td>12·11</td>
<td>9·34</td>
</tr>
<tr>
<td><strong>Tobacco use</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Current (%)</td>
<td>Mean</td>
<td>20a</td>
<td></td>
<td>23a</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Ex (%)</td>
<td>69c</td>
<td>31a,b</td>
<td>40b</td>
<td></td>
</tr>
<tr>
<td><strong>BMI (kg/m²)</strong></td>
<td>Mean</td>
<td>26·83c</td>
<td>25·30b,c</td>
<td>23·01b</td>
<td>21·81a</td>
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<td></td>
<td>sd</td>
<td>2·11</td>
<td>1·98</td>
<td>0·95</td>
<td></td>
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<tr>
<td><strong>Weight (kg)</strong></td>
<td>Mean</td>
<td>80·30</td>
<td>75·08</td>
<td>68·87</td>
<td>69·79</td>
</tr>
<tr>
<td></td>
<td>sd</td>
<td>2·98</td>
<td>3·31</td>
<td>4·20</td>
<td>9·18</td>
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<td><strong>Waist circumference (cm)</strong></td>
<td>Mean</td>
<td>90·12</td>
<td>89·21</td>
<td>80·23</td>
<td>78·60</td>
</tr>
<tr>
<td></td>
<td></td>
<td>0·92</td>
<td>0·32</td>
<td>1·21</td>
<td></td>
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<td><strong>College education</strong></td>
<td>Mean</td>
<td>120</td>
<td>210</td>
<td>298</td>
<td>321</td>
</tr>
<tr>
<td></td>
<td></td>
<td>20·65</td>
<td>33·28</td>
<td>45·22</td>
<td></td>
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<tr>
<td><strong>Blood lipids</strong></td>
<td></td>
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<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>TAG (mmol/l)</td>
<td>Mean</td>
<td>2·19b</td>
<td>1·85b</td>
<td>1·62b</td>
<td>1·12a</td>
</tr>
<tr>
<td></td>
<td>sd</td>
<td>0·92</td>
<td>1·03</td>
<td>0·91</td>
<td>1·09</td>
</tr>
<tr>
<td>TC (mmol/l)</td>
<td>Mean</td>
<td>6·61c</td>
<td>5·45b</td>
<td>6·14a</td>
<td>4·24a</td>
</tr>
<tr>
<td></td>
<td>sd</td>
<td>1·57</td>
<td>2·31</td>
<td>1·22</td>
<td>1·22</td>
</tr>
<tr>
<td>HDL (mmol/l)</td>
<td>Mean</td>
<td>0·90</td>
<td>1·01</td>
<td>1·19</td>
<td>1·08</td>
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<tr>
<td></td>
<td>sd</td>
<td>0·19</td>
<td>1·01</td>
<td>0·91</td>
<td>0·53</td>
</tr>
<tr>
<td>LDL (mmol/l)</td>
<td>Mean</td>
<td>3·58b</td>
<td>3·51b</td>
<td>3·62b</td>
<td>2·35a</td>
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<tr>
<td></td>
<td>sd</td>
<td>2·11</td>
<td>1·33</td>
<td>0·89</td>
<td>0·86</td>
</tr>
<tr>
<td>ApoA (g/l)</td>
<td>Mean</td>
<td>1·87</td>
<td>1·76</td>
<td>1·29</td>
<td>1·30</td>
</tr>
<tr>
<td></td>
<td>sd</td>
<td>0·58</td>
<td>0·19</td>
<td>0·21</td>
<td>0·28</td>
</tr>
<tr>
<td>ApoB (g/l)</td>
<td>Mean</td>
<td>1·26</td>
<td>1·01</td>
<td>1·29</td>
<td>0·99</td>
</tr>
<tr>
<td></td>
<td>sd</td>
<td>0·02</td>
<td>0·71</td>
<td>0·23</td>
<td>0·07</td>
</tr>
<tr>
<td>Lp-a (mg/l)</td>
<td>Mean</td>
<td>178·69</td>
<td>150·18</td>
<td>134·19</td>
<td>140·31</td>
</tr>
<tr>
<td></td>
<td>sd</td>
<td>9·81</td>
<td>8·12</td>
<td>7·31</td>
<td>31·23</td>
</tr>
<tr>
<td><strong>Blood pressure (mmHg)</strong></td>
<td>Systolic</td>
<td>129·21</td>
<td>139·12</td>
<td>131·11</td>
<td>113·98</td>
</tr>
<tr>
<td></td>
<td>Diastolic</td>
<td>86·31</td>
<td>82·21</td>
<td>80·98</td>
<td>81·56</td>
</tr>
<tr>
<td></td>
<td>Glucose (mmol/l)</td>
<td>6·21</td>
<td>5·91</td>
<td>6·10</td>
<td>5·27</td>
</tr>
</tbody>
</table>

ACS, acute coronary syndrome; CCAD, chronic coronary artery disease; HRP, high-risk population; HV, healthy volunteers; TC, total cholesterol; Lp-a, lipoprotein a.

a,b Values within a row with unlike superscript letters were significantly different (P<0·05 or P<0·001).

* Blood lipids tested include: TAG; TC; HDL-cholesterol (HDL); LDL-cholesterol (LDL); apoA (mainly including apoAI); apoB (including both apoB100 and apoB48); Lp-a.
The 10-year RP of CHD was calculated on the basis of the Cox proportional hazards multivariate model formulation and the Framingham risk equation\(^{22}\). The risk score was defined as

\[
 f_{\text{group}}(X, M) = \beta_1(X_1 - M_1) + \beta_2(X_2 - M_2) + \cdots + \beta_i(X_i - M_i).
\]  

(2)

Furthermore, the 10-year CHD RP was calculated from the following equation:

\[
 \text{RP}_{\text{group}} = 1 - S_t(t)^{\exp{f_{\text{group}}(X, M)}},
\]  

(3)

where \(\text{RP}_{\text{group}}\) is the CHD probability within the next 10 years of each group (ACS, CCAD, HRP and HV); \(S_t(t)\) is the hazard function at time \(t\); \(X_1, X_2, X_3, \ldots, X_i\) are independent risk factor variables for each individual; \(M_1, M_2, M_3, \ldots, M_i\) are the average levels of risk factors in each group; \(\beta_1, \beta_2, \beta_3, \ldots, \beta_i\) are the partial regression coefficients of different risk factors.

The estimated partial regression coefficients, hazard ratio and their corresponding 95% CI are shown in Table 2; the descriptive characteristics are shown in Table 1. These parameters and data were substituted into equation 2 and the risk score for the ACS group was calculated as \(f_{\text{ACS}}(X, M)\), which, in turn, was substituted into equation 3 and the average 10-year \(\text{RP}_{\text{ACS}}\) was calculated as \(1 - 0.9876^{\exp{f_{\text{ACS}}(X, M)}}\). In the same way, the average 10-year \(\text{RP}_{\text{CCAD}}\) of the CCAD group was calculated as \(1 - 0.9821^{\exp{f_{\text{CCAD}}(X, M)}}\), the average 10-year \(\text{RP}_{\text{HRP}}\) of the HRP group as \(1 - 0.9781^{\exp{f_{\text{HRP}}(X, M)}}\), and the average 10-year \(\text{RP}_{\text{HV}}\) of the HV group as \(1 - 0.9753^{\exp{f_{\text{HV}}(X, M)}}\) (For the equation calculation process, see the Supplementary material, available online).

The assumption of proportional hazards and the calibration were adjusted for confounding factors such as baseline age, sex, smoking and LDL, and verified by the Hosmer and Lemeshow test\(^{24,25}\). The proportional hazards assumption was considered to be valid when the difference in the \(P\) value was <0.05.

SPSS for Windows version 18.0 was used to calculate the regression equations and correlation coefficients between the TFA index and the 10-year CHD-RP for the different groups. Bonferroni correction was performed for ANOVA and correlation analysis to verify statistically significant differences with a \(P\) value <0.05.

### Results

**Trans-fatty acid profile and index**

The erythrocyte membrane, serum and whole-blood TFA profiles are listed in Table 3. Only in erythrocyte membranes were
Table 3. Erythrocyte membrane, serum and whole-blood trans-fatty acid (TFA) profiles in each group*  
(Mean values and standard deviations)

<table>
<thead>
<tr>
<th></th>
<th>Trans-16:1n-7</th>
<th>Trans-16:1n-9</th>
<th>Trans-18:1n-7</th>
<th>Trans-18:1n-9</th>
<th>Trans-18:2n6n-9</th>
<th>Trans-18:1n-9</th>
<th>Trans-18:2n6n-9</th>
<th>Total TFA</th>
<th>TFA index</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>SD</td>
<td>Mean</td>
<td>SD</td>
<td>Mean</td>
<td>SD</td>
<td>Mean</td>
<td>SD</td>
<td>Mean</td>
</tr>
<tr>
<td>Erythrocyte membrane</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>ACS</td>
<td>0.19</td>
<td>0.01</td>
<td>0.21</td>
<td>0.09</td>
<td>0.39</td>
<td>0.04</td>
<td>0.43</td>
<td>0.11</td>
<td>0.42</td>
</tr>
<tr>
<td>CCAD</td>
<td>0.17</td>
<td>0.06</td>
<td>0.18</td>
<td>0.04</td>
<td>0.41</td>
<td>0.08</td>
<td>0.38</td>
<td>0.12</td>
<td>0.38</td>
</tr>
<tr>
<td>HRP</td>
<td>0.16</td>
<td>0.03</td>
<td>0.23</td>
<td>0.08</td>
<td>0.51</td>
<td>0.19</td>
<td>0.25</td>
<td>0.10</td>
<td>0.29</td>
</tr>
<tr>
<td>HV</td>
<td>0.26</td>
<td>0.11</td>
<td>0.14</td>
<td>0.01</td>
<td>0.59</td>
<td>0.21</td>
<td>0.21</td>
<td>0.10</td>
<td>0.18</td>
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<tr>
<td>HRP</td>
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<td>0.57</td>
<td>7.12</td>
<td>1.23</td>
<td>1.52</td>
<td>0.09</td>
<td>5.06</td>
<td>1.03</td>
<td>3.11</td>
</tr>
<tr>
<td>Serum</td>
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<td>0.01</td>
<td>0.21</td>
<td>0.09</td>
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<td>0.01</td>
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<td>0.13</td>
</tr>
<tr>
<td>CCAD</td>
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<td>0.03</td>
<td>0.11</td>
<td>0.02</td>
<td>0.31</td>
<td>0.02</td>
<td>0.08</td>
<td>0.01</td>
<td>0.12</td>
</tr>
<tr>
<td>HRP</td>
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<td>0.01</td>
<td>0.08</td>
<td>0.01</td>
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<td>0.02</td>
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<td>0.02</td>
</tr>
<tr>
<td>HV</td>
<td>0.04</td>
<td>0.01</td>
<td>0.17</td>
<td>0.07</td>
<td>0.38</td>
<td>0.03</td>
<td>0.05</td>
<td>0.02</td>
<td>0.08</td>
</tr>
<tr>
<td>Whole blood</td>
<td></td>
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<td>ACS</td>
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<td>HRP</td>
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<td>0.08</td>
<td>0.01</td>
<td>0.08</td>
</tr>
</tbody>
</table>

* Values are given as the percentages of individual fatty acids relative to the total chromatogram area (FA%).

**Correlation between trans-fatty acid index and 10-year CHD risk probability**

**CHD risk probability**

The results of the trans-fatty acid (TFA) and 10-year CHD risk probability regression analysis in each group are described in Table 1. TFA index was negatively correlated with BMI and waist circumference, while smoking status and BMI were positively correlated with the TFA index. Partial regression coefficients and their corresponding 95% CI are shown in Table 2. The estimated partial regression coefficients were -0.003, -0.004, -0.005, and -0.006, respectively, for the HS, HS-, CCAD, and CCAD- groups, respectively. The TFA index value represents a coincidental trend with the regression line.
generally ruminant-sourced (29), although on CHD events from the Nurses’ Health Study in 32 826 participants which is consistent with the proven detrimental impact of in the membrane (28). Therefore, erythrocyte membrane TFA no fatty acid synthesis, chain elongation or desaturation

Discussion

Erythrocyte membrane trans-fatty acids reflect the change in body trans-fatty acids in CHD

The levels of all of the erythrocyte membrane TFA isomers differed significantly between the groups. In contrast, only trans-18:1n-9 and trans-18:2n-6n-9 differed significantly between the whole-blood and serum samples. The probable reasons for this difference were as follows: on the one hand, the serum fatty acid level reflected fat intake over the past few days due to the immediate penetration of dietary fat into the serum(26), and it is hard to get much change in fatty acid profile during a few days, especially TFA; on the other hand, the fatty acid level of erythrocyte membrane reflected chronic lipid storage, as well as the average level of body lipids over the previous 4–12 weeks(27), and there was no fatty acid synthesis, chain elongation or desaturation in the membrane(28). Therefore, erythrocyte membrane TFA reflected the change in body TFA levels in CHD(29).

Trans-fatty acid index reflects trans-fatty acid hazard on CHD

The five TFA isomers in the erythrocyte membrane included two n-9 TFA (trans-16:1n-9 and trans-18:1n-9), two n-7 TFA (trans-16:1n-7 and trans-18:1n-7) and one double-bonded TFA (trans-18:2n-6n-9). The two n-9 TFA and trans-18:2n-6n-9 primarily originate from partially hydrogenated vegetable oils (‘industrial’ TFA)(29). The two n-7 TFA are generally ruminant-sourced(20), although trans-18:1n-7 may also come from an industrial source. The levels of n-9 TFA and trans-18:2n-6n-9 were significantly higher in ACS, which is consistent with the proven detrimental impact of industrial TFA in the promotion and induction of CHD events from the Nurses’ Health Study in 32 826 participants and 6 years of follow-up(30). Another study investigating TFA and sudden cardiac death among 86 762 women has shown that the levels of trans-18:2 and trans-16:1n-9 were positively associated with myocardial infarction (P<0.001) after adjusting for established risk factors and other confounders, and that the trans-18:2 isomer may play a greater role in sudden cardiac death among individuals with clinically manifest atherosclerosis(30).

Remarkably, in contrast, n-7 TFA levels were high in the HV group while low in the ACS group, which may result from the ruminant-sourced generation. Ruminant trans-16:1n-7 was associated with trans-18:1n-7 and may be converted into trans-18:1n-7 by the carboxyl chain increase(31). As a major TFA of ruminant fat, trans-18:1n-7 is produced in the rumen and converted in tissues to 9-cis, 11-trans-18:2n-6 CLA by Δ9-desaturase with an average conversion rate of 19%(32). 9-cis, 11-trans-18:2n-6 CLA could positively modulate HDL-cholesterol metabolism and enhance reverse cholesterol transport, and prevent the progression of atherosclerosis in humans(33). Hence, the effect of ruminant TFA on CHD may be neutral or somewhat favourable due to the indirect benefit of 9-cis, 11-trans-18:2n-6 CLA(34). However, the industrial TFA are harmful. It has been reported that the consumption of industrial trans-18:1n-9 by LDL receptor-deficient (LDL−/−) mice stimulated atherosclerotic development(35), while consumption of a diet enriched in trans-18:1n-7 reduced cholesterol-induced hyperlipidaemia and atherosclerosis and thus protected against atherosclerotic lesions(36). Chronic trans-18:1n-7 supplementation also significantly abated dyslipidaemia in both the food-deprived and postprandial states in JCR:LA-cp rats due to reductions in intestinal chylomicrons and hepatic de novo lipogenesis pathways(37). Another study investigating the effects of ruminant-derived TFA on immune function in a model of the metabolic syndrome (JCR:LA-cp rats) has shown that vaccenic acid might protect from CVD due to an anti-inflammatory action(38).

Fig. 1. Linear correlation between the erythrocyte membrane trans-fatty acid (TFA) index and the 10-year CHD risk probability (RP) \(y = 17.122x + 1.1365\) and \(R^2 = 0.9881\). Values (Δ) are average TFA index and RP in the healthy volunteers (HV, \(n = 842\)), high-risk population (HRP, \(n = 659\)), chronic coronary artery disease (CCAD, \(n = 631\)) and acute coronary syndrome (ACS, \(n = 581\)) groups, respectively, with standard deviations represented by vertical bars.

Fig. 2. Linear correlation between the whole-blood trans-fatty acid (TFA) index and the 10-year CHD risk probability (RP) \(y = 14.365x + 21.141\) and \(R^2 = 0.9607\). Values (Δ) are average TFA index and RP in the healthy volunteers (HV, \(n = 842\)), high-risk population (HRP, \(n = 659\)), chronic coronary artery disease (CCAD, \(n = 631\)) and acute coronary syndrome (ACS, \(n = 581\)) groups, respectively, with standard deviations represented by vertical bars.
Although ruminant TFA have been shown to elicit neutral or less detrimental effects, further studies in human subjects are still needed to prove the protective effects.

To unify the inverse trend of decreased ruminant TFA and increased industrial TFA from the HV to the ACS group, we divided the total industrial TFA levels by the total ruminant TFA levels. This index value represents a coincidental trend of TFA change. An exponential function was used to make the index stronger and more obvious as the TFA index. The TFA index could estimate the true objective change in TFA levels comprehensively by unifying the disparate effects of different TFA on the body; accordingly, the hazardous level of all TFA isomers on CHD could be evaluated. In addition, it was interesting to note that none of the five TFA isomers showed any significant differences except the TFA index \((P<0.001)\) between the ACS and CCAD groups, indicating that the TFA index could discriminate the acute CHD symptoms of the ACS from the chronic CHD symptoms of CCAD.

**Trans-fatty acid index is associated positively with 10-year CHD risk probability**

The average TFA index in erythrocyte membranes differed between the groups and increased progressively from the HV to the ACS group. The 10-year CHD RP were 4, 13, 21 and 31% in the HV, HRP, CCAD and ACS groups, respectively. These values generally reflect the potential probability and pathological severity of CHD. A strong linear correlation was seen between the average TFA index and the 10-year CHD RP \((R^2=0.9981; \text{Fig. 1})\), which suggests that as the erythrocyte membrane TFA level goes up, CHD may get worse.

CHD incidence could be influenced directly or indirectly by TFA through TAG accumulation, vasodilation, inflammation, PG translation and/or platelet aggregation. Trans-fatty acids elicit an unfavourable effect on the lipoprotein profile by stimulating cholesteryl ester transfer protein activity \((r=0.58, P<0.005)\), increasing the LDL level and decreasing the HDL level \((r=-0.57, P<0.01)\). These changes may contribute to a more atherogenic lipoprotein profile. A high intake of TFA could adversely affect endothelial function; this would partially explain why the positive relationship between trans-fats and cardiovascular health took precedence over the adverse effects of trans-fats on lipids and lipoproteins. In addition, a recent study has suggested that trans-18:1n-9 maintains the levels of vascular cell adhesion molecule-1 and intercellular adhesion molecule-1 up-regulated by TNF-α or lipase. This kept the human brain microvascular endothelial function at the stimulated phenotype, which could promote CHD.

In summary, the erythrocyte membrane TFA index was proposed as a method to unify the content changes in different TFA isomers and their effects on CHD in one equation. The TFA index manifested a strong and positive linear correlation with the 10-year CHD RP. Although the present study might be limited to the variety of TFA isomers analysed, the results should contribute to further studies on the relationship between TFA and CHD.

**Supplementary material**

To view supplementary material for this article, please visit http://dx.doi.org/10.1017/S0007114513000196

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