The ethnicity-specific association of biomarkers with the angiographic severity of coronary artery disease

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Abstract

Background Risk factor burden and clinical characteristics of patients with coronary artery disease (CAD) differ among ethnic groups. We related biomarkers to CAD severity in Caucasians, Chinese, Indians and Malays.

Methods In the Dutch-Singaporean UNICORN coronary angiography cohort (n = 2033) we compared levels of five cardiovascular biomarkers: N-terminal pro-brain natriuretic peptide (NTproBNP), high-sensitivity C-reactive protein (hsCRP), cystatin C (CysC), myeloperoxidase (MPO) and high-sensitivity troponin I (hsTnI). We assessed ethnicity-specific associations of biomarkers with CAD severity, quantified by the SYNTAX score.

Results Adjusted for baseline differences, NTproBNP levels were significantly higher in Malays than in Chinese and Caucasians (72.1 vs. 34.4 and 41.1 pmol/l, \( p < 0.001 \) and \( p = 0.005 \), respectively). MPO levels were higher in Caucasians than in Indians (32.8 vs. 27.2 ng/ml, \( p = 0.026 \)) and hsTnI levels were higher in Chinese than in Caucasians (23.3 vs. 16.4, \( p = 0.031 \)). We found modifying effects of ethnicity on the association of biomarkers with SYNTAX score. NTproBNP associated more strongly with the SYNTAX score in Malays than Caucasians (\( \beta = 0.132 \) vs. \( \beta = 0.020 \) per 100 pmol/l increase in NTproBNP, \( p = 0.032 \)). For MPO levels the association was stronger in Malays than Caucasians (\( \beta = 1.146 \) vs. \( \beta = 0.016 \) per 10 ng/ml increase, \( p = 0.017 \)). Differing biomarker cut-off levels were found for the ethnic groups.

Conclusion When corrected for possible confounders we observe ethnicity-specific differences in biomarker levels. Moreover, biomarkers associated differently with CAD severity, suggesting that ethnicity-specific cut-off values should be considered.

Keywords Ethnicity · Coronary angiography · Biomarkers · Singapore
Introduction

Coronary artery disease (CAD) is highly prevalent worldwide but over the next few decades the majority of deaths due to CAD will occur in Asia [1]. Evidence is accumulating that important differences exist between the Asian ethnic groups and Caucasians, who have been the main focus of cardiovascular research up until now. We know that risk factor levels differ markedly. Specifically, a high prevalence of diabetes has been described for South Asians [2]. Also, the CAD phenotype appears to differ among the ethnic groups. Triple-vessel disease is more common in South Asians than in Caucasians [3], whilst the Chinese suffer from less severe CAD [4].

Blood-derived biomarkers are noninvasive tools that can be indicative of CAD severity. Population means of biomarker levels differ among ethnic groups [5]. However, it is unknown whether biomarkers of CAD report similarly on underlying CAD severity among these groups [6]. In the current study, we investigated patients undergoing coronary angiography for suspected CAD from four globally populous ethnic groups: Caucasians, Chinese, Indians and Malays, enrolled in two countries with high comparable healthcare standards [7]: the Netherlands and Singapore. We evaluated five established biomarkers known to be affected by CAD: N-terminal pro-brain natriuretic peptide (NTproBNP) [8–10], high-sensitivity C-reactive protein (hsCRP) [11], cystatin C (CysC) [12, 13], myeloperoxidase (MPO) [14, 15] and high-sensitivity troponin I (hsTnI) [16, 17]. These biomarkers reflect pivotal CAD aspects and complications: cardiac haemodynamic load (NTproBNP), inflammation (hsCRP and MPO), kidney function (CysC) and cardiomyocyte damage (hsTnI).

In this study we aimed to define inter-ethnic differences in the association of these biomarkers with CAD severity, quantified by the SYNTAX score [18].

Methods

This study was conducted using the parallel United CORo-Nary biobanks, the UNICORN cohort (clinicaltrials.gov ID: NCT02126150), consisting of consecutive patients undergoing coronary angiography recruited from two sites: the University Medical Center Utrecht, the Netherlands and the National University Hospital, Singapore. The institutional review boards of both centres approved of this study, which conforms to the declaration of Helsinki.

Patients were enrolled between September 2010 and March 2013. Patients from four ethnic groups were enrolled, namely Caucasians in the Netherlands and Chinese, Indians and Malays in Singapore. Blood was sampled from the arte-

Clinical and angiographic characteristics

Risk factor profiles were documented at or around the time of coronary angiography. Coronary angiograms were categorised into four groups: no/minor CAD, single-vessel disease, double-vessel disease and triple-vessel disease (defined as number of epicardial vessels with 50% stenosis [19] on visual assessment). In the last three categories, a SYNTAX score was determined [18]. The process of SYNTAX scoring has been described previously [20].

Biomarker assays

Plasma levels of NTproBNP, hsCRP, CysC and MPO were measured at the University Medical Center Utrecht, the Netherlands, using a semi-automated ELISA robot (Freedom EVO, Tecan, Switzerland). Commercial antibody combinations were used to quantify NTproBNP (15 C4 and biotinylated 13G12, Hi-test Finland), hsCRP (Dy1707 duoset, R&D systems), CysC (Dy1196 duoset, R&D systems) and MPO (Dy3667 duoset, R&D systems). In brief, maxisorb plates were coated with mouse anti-human NTproBNP, hsCRP, CysC or MPO. Plates were blocked with 1% bovine serum albumin, and incubated with supernatants. Plates were washed with phosphate buffered saline pH 7.4 with 0.05% Tween 20. Bound factors were detected with biotin coupled detection antibodies. Biotin coupled antibodies were bound with streptavidin horseradish peroxidase (HRP), or goat-anti-human antibodies with rabbit-anti-goat HRP (DAKO, P0449). Detection was performed with SuperSignal West Pico Chemiluminescent substrate, and read with a luminometer. The intra-assay variation coefficient was 10%. Levels of hsTnI were measured using the STAg hsTnI assay on the clinically validated ARCHITECT i2000 analyser (Abbott Laboratories, Lisnamuck, Longford, Ireland).

All samples (Singaporean and Dutch) were randomly distributed across the plates for all analyses.

Statistical analysis

Figures are presented as means with standard deviations for normally distributed variables or medians with interquartile ranges for non-normally distributed variables. Baseline characteristics were compared among the ethnic groups using ANOVA for normally distributed variables, Kruskal-Wallis tests for non-normally distributed data and chi-square tests for categorical data.

We evaluated inter-ethnic differences in biomarker levels, adjusted for baseline differences in age, sex, body...
Table 1  Baseline characteristics of UNICORN participants, stratified by ethnicity

| Characteristic                          | Caucasian | Chinese | Indian | Malay | \( p \)-value |
|-----------------------------------------|-----------|---------|--------|-------|--------------|
| **N**                                   | 1132      | 562     | 158    | 181   |              |
| Males (%)                               | 72.5      | 81.0    | 80.4   | 79.6  | <0.001       |
| Age (mean (SD), years)                  | 64.7 (11.3)| 59.1  (9.9)| 54.4  (9.5)| 55.4 (9.4)| <0.001       |
| BMI (mean (SD), kg/m²)                  | 27.0 (4.3) | 26.2 (4.7)| 27.5 (5.0)| 28.5 (5.4)| <0.001       |
| Diabetes (%)                            | 22.6      | 34.0    | 54.4   | 52.5  | <0.001       |
| Hypertension (%)                        | 56.1      | 65.1    | 63.9   | 64.6  |              |
| Dyslipidaemia (%)                       | 48.2      | 69.6    | 76.6   | 75.1  | <0.001       |
| Smoking, current (%)                    | 23.6      | 29.4    | 42.9   | 39.0  | <0.001       |
| Smoking, ex (%)                        | 31.6      | 19.2    | 11.9   | 16.4  |              |
| Renal failure (%)                       | 3.1       | 6.0     | 3.8    | 5.6   | 0.034        |

**Medications**

| Category                                | Caucasian | Chinese | Indian | Malay | \( p \)-value |
|-----------------------------------------|-----------|---------|--------|-------|--------------|
| Platelet inhibitor (%)                  | 64.6      | 58.4    | 57.6   | 55.8  | 0.016        |
| Statin (%)                              | 58.9      | 44.3    | 43.0   | 43.1  | <0.001       |
| Beta blocker (%)                        | 25.4      | 26.7    | 24.7   | 19.3  | 0.259        |
| RAAS (%)                                | 51.4      | 35.2    | 40.5   | 43.6  | <0.001       |
| Calcium antagonist (%)                  | 64.3      | 61.4    | 61.4   | 55.8  | 0.146        |

**Medical history**

| Category                                | Caucasian | Chinese | Indian | Malay | \( p \)-value |
|-----------------------------------------|-----------|---------|--------|-------|--------------|
| Previous PCI (%)                        | 31.6      | 17.5    | 29.1   | 30.0  | <0.001       |
| Previous CAGB (%)                       | 13.0      | 6.2     | 6.3    | 6.1   | <0.001       |
| Previous ACS (%)                        | 32.6      | 14.8    | 26.8   | 28.5  | <0.001       |
| History of CVA/TIA (%)                  | 10.5      | 5.7     | 8.2    | 6.7   | 0.010        |
| History of PAD (%)                      | 11.5      | 2.8     | 4.4    | 5.0   | <0.001       |

**Indication for coronary angiography**

| Category                                | Caucasian | Chinese | Indian | Malay | \( p \)-value |
|-----------------------------------------|-----------|---------|--------|-------|--------------|
| Stable CAD (%)                          | 67.4      | 65.9    | 58.8   | 54.8  | <0.001       |
| UA/NSTEMI (%)                           | 22.5      | 31.6    | 36.6   | 42.9  |              |
| STEMI (%)                               | 10.1      | 2.4     | 4.6    | 2.3   |              |

**CAD severity**

| Category                                | Caucasian | Chinese | Indian | Malay | \( p \)-value |
|-----------------------------------------|-----------|---------|--------|-------|--------------|
| No/minor CAD (%)                        | 25.0      | 33.8    | 28.7   | 25.1  | <0.001       |
| Single-vessel disease (%)               | 37.3      | 21.7    | 24.0   | 21.1  |              |
| Double-vessel disease (%)               | 25.5      | 23.6    | 25.3   | 27.4  |              |
| Triple-vessel disease (%)               | 12.2      | 20.8    | 22.0   | 26.3  |              |

**Treatment**

| Category                                | Caucasian | Chinese | Indian | Malay | \( p \)-value |
|-----------------------------------------|-----------|---------|--------|-------|--------------|
| Conservative (%)                        | 35.0      | 53.6    | 58.2   | 55.2  | <0.001       |
| PCI (%)                                 | 58.8      | 37.5    | 37.3   | 33.7  |              |
| CAGB (%)                                | 6.2       | 8.9     | 4.4    | 11.0  |              |

**Biomarker levels**

| Biomarker | Median [IQR] | \( p \)-value |
|-----------|--------------|---------------|
| NT-proBNP | 43.1 [10.5, 138.0] | 0.049         |
| hsCRP     | 1.5 [0.6, 3.8] | <0.001        |
| CysC      | 0.8 [0.7, 1.1] | <0.001        |
| MPO       | 28.1 [20.7, 44.8] | 0.001         |
| hsTnI     | 7.8 [3.8, 28.4] | 0.001         |

**Outcome**

| Outcome | Median [IQR] | \( p \)-value |
|---------|--------------|---------------|
| SYNTAX score | 11.6 [8.2] | <0.001        |
| FU time | 823 [653, 988] | <0.001        |

Baseline characteristics of UNICORN patients stratified by ethnicity. Non-normally distributed continuous variables were compared using ANOVA across ethnic groups. Biomarker levels were compared by Kruskal-Wallis testing. Categorical data were compared using chi-square testing. Reported \( p \)-values refer to overall differences across the ethnic groups. Significant \( p \)-values are printed in bold.
ethnicity, defined as the biomarker level at which sensitivity + specificity for a high SYNTAX score was largest. Also, areas under curves were calculated for association with a high SYNTAX score for each biomarker in each ethnic group [21]. The outcome SYNTAX score was dichotomised on a previously reported [22] cut-off of 18 points. All analyses were performed using the R software package [23]. A two-tailed \( \alpha \) of: 0.05 was considered statistically significant.

Results

Baseline characteristics

Marked baseline differences were observed among the ethnic groups (Table 1). Indian and Malay patients were younger (54.4 and 55.4 years, respectively) than the Chinese and Caucasian patients (59.1 and 64.7 years, respectively, \( p<0.001 \)). Among Indian and Malay patients, diabetes (54.4 and 52.5%, respectively) and dyslipidaemia (76.6 and 75.1%, respectively) were strikingly more common than mass index, diabetes, hypertension, dyslipidaemia, smoking, indication for angiography, CAD severity, antiplatelet medication use, beta blocker use, calcium antagonist use, RAAS inhibiting medication use and statin use. Using ANCOVA, adjusted biomarker levels were calculated for each ethnic group. Biomarker levels were positively skewed and therefore log-transformed for analysis when used as the dependent variable. The presented adjusted biomarker level means are antilogs (back-transformed after analysis). Biomarker levels were compared among the ethnic groups through Tukey post-hoc testing, thus adjusting for multiple testing.

Also, we tested for interactions between ethnicity and biomarker levels for the SYNTAX score in univariable and multivariable regression models. Direct comparison of biomarker coefficients between ethnicities was performed by testing an interaction term of biomarker level with ethnicity; Caucasian ethnicity was the reference group. A conservative \( p \)-value of 0.05 for the interaction terms was deemed significant.

Next, using receiver operating characteristic (ROC) analysis, we compared the optimal cut-off biomarker values by ethnicity, defined as the biomarker level at which sensitivity + specificity for a high SYNTAX score was largest. Also, areas under curves were calculated for association with a high SYNTAX score for each biomarker in each ethnic group [21]. The outcome SYNTAX score was dichotomised on a previously reported [22] cut-off of 18 points. All analyses were performed using the R software package [23]. A two-tailed \( \alpha \) of: 0.05 was considered statistically significant.

Results

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in Caucasians (diabetes 22.6%, \( p < 0.001 \), dyslipidaemia 48.2%, \( p < 0.001 \)). The prevalence of current smokers was highest among Indians (42.9%).

The indication for angiography differed among the ethnic groups (\( p < 0.001 \)). Stable CAD was slightly less common among Indians and Malays (58.8 and 54.8%, respectively, vs. 67.4% in Caucasians and 65.9% in Chinese). Unstable angina or non-ST-elevated myocardial infarction was more frequently observed in all three Asian groups than in Caucasians (Chinese 31.6%, Indians 36.6%, Malays 42.9 vs. 25.5% in Caucasians, \( p < 0.001 \)). Also, triple-vessel disease was more often diagnosed in the Asian ethnic groups than in Caucasian (Chinese 31.6%, Indians 36.6%, Malays 42.9 vs. 67.4% in Caucasians and 65.9% in Chinese).
Caucasians (Chinese 20.8%, Indian 22.0%, Malay 26.3 vs. 12.2% in Caucasians, p < 0.001).

Percutaneous coronary intervention (PCI) was performed more often on Caucasians than in the other ethnic groups, in whom a conservative strategy was more frequently opted for (p < 0.001). Coronary artery bypass graft (CABG) surgery was performed most often in Malays (11.0 vs. 4.4% in Indians, 6.2% in Caucasians and 8.9% in Chinese).

### Biomarker levels

Crude and multivariable-adjusted biomarker levels are displayed in Fig. 1, stratified by ethnicity. Crude biomarker levels differed significantly among the ethnic groups for all of the examined biomarkers (Table 1). In Table 2, the crude biomarker levels are displayed for each ethnic group, stratified by indication for angiography and angiographic severity of CAD. All biomarkers differed among the ethnic groups in at least one indication group. High-sensitivity TnI did not differ in any of the CAD severity groups, while the other biomarkers did differ significantly in at least one CAD severity group.

When adjusted for baseline characteristics, certain differences in biomarker levels remained statistically significant. Post-hoc testing revealed that multivariable-adjusted NTproBNP levels were significantly higher in Malays than in Chinese and Caucasians (72.1 vs. 34.4 and 41.1 pmol/l, p < 0.001 and p = 0.005, respectively). MPO levels were higher in Caucasians than in Indians (32.8 vs. 27.2 ng/ml, p = 0.026), hsTnI levels were higher in Malays than in Caucasians and Indians (33.3 vs. 16.4 and 17.8 ng/l, p < 0.001 and p = 0.029) and hsTnI levels were higher in Chinese than in Caucasians (23.3 vs. 16.4, p = 0.031). High-sensitivity CRP levels and CysC levels (additionally adjusted for renal failure) did not differ among the ethnic groups.

### Table 3 Regression coefficients (betas with 95% confidence intervals) of biomarker levels for SYNTAX score

| Biomarker   | Ethnicity | Beta (CI, univariable) | p-value beta (univ.) | Beta (CI, multivariable) | p-value beta (multiv.) | p-value interaction (univ.) | p-value interaction (multiv.) |
|-------------|-----------|------------------------|----------------------|--------------------------|------------------------|-----------------------------|-------------------------------|
| NTproBNP (pmol/l) | Caucasian | 0.056 (−0.002–0.114) | 0.059                | 0.020 (−0.047–0.087)    | 0.560                  | Ref.                        | Ref.                          |
|             | Chinese   | 0.153 (0.053–0.254)   |                      | 0.014 (−0.013–0.221)   | 0.081                  | 0.094                       | 0.115                         |
|             | Indian    | 0.088 (−0.065–0.241)  |                      | 0.020 (−0.119–0.160)   | 0.771                  | 0.669                       | 0.361                         |
|             | Malay     | 0.152 (0.011–0.292)   |                      | 0.132 (−0.020–0.284)   | 0.088                  | 0.157                       | 0.032                         |
| hsCRP (μg/ml) | Caucasian | 0.041 (0.009–0.074)   | 0.012                | 0.030 (−0.006–0.066)   | 0.103                  | Ref.                        | Ref.                          |
|             | Chinese   | 0.029 (−0.020–0.077)  |                      | 0.035 (−0.025–0.095)   | 0.255                  | 0.663                       | 0.709                         |
|             | Indian    | 0.000 (−0.095–0.112)  |                      | 0.033 (−0.084–0.150)   | 0.578                  | 0.505                       | 0.627                         |
|             | Malay     | −0.025 (−0.117–0.066) | 0.584                | −0.040 (−0.148–0.069)  | 0.471                  | 0.113                       | 0.621                         |
| CysC (μg/ml) | Caucasian | 0.162 (−0.015–0.340)  | 0.073                | 0.081 (−0.131–0.293)   | 0.454                  | Ref.                        | Ref.                          |
|             | Chinese   | 0.235 (−0.015–0.485)  | 0.066                | 0.154 (−0.194–0.503)   | 0.383                  | 0.642                       | 0.371                         |
|             | Indian    | 0.022 (−0.449–0.492)  | 0.928                | −0.172 (−0.700–0.356)  | 0.519                  | 0.541                       | 0.753                         |
|             | Malay     | −0.168 (−0.923–0.586) | 0.660                | −0.165 (−0.935–0.605)  | 0.671                  | 0.314                       | 0.457                         |
| MPO (ng/ml) | Caucasian | 0.101 (0.015–0.187)   | 0.022                | 0.016 (−0.095–0.127)   | 0.777                  | Ref.                        | Ref.                          |
|             | Chinese   | −0.008 (−0.485–0.469) | 0.974                | −0.079 (−0.650–0.472)  | 0.778                  | 0.633                       | 0.885                         |
|             | Indian    | −0.077 (−1.314–1.160) | 0.902                | 0.108 (−1.099–1.314)   | 0.859                  | 0.743                       | 0.555                         |
|             | Malay     | 1.517 (0.451–2.583)   | 0.006                | 1.146 (0.023–2.269)    | 0.046                  | 0.002                       | 0.017                         |
| hsTnI (ng/l) | Caucasian | 0.044 (0.026–0.062)   | <0.001               | 0.032 (0.012–0.052)    | 0.002                  | Ref.                        | Ref.                          |
|             | Chinese   | 0.035 (0.008–0.061)   | 0.010                | 0.044 (0.013–0.074)    | 0.006                  | 0.557                       | 0.164                         |
|             | Indian    | 0.022 (−0.023–0.067)  | 0.331                | 0.050 (−0.004–0.104)   | 0.067                  | 0.328                       | 0.958                         |
|             | Malay     | 0.003 (−0.027–0.033)  | 0.858                | 0.003 (−0.027–0.033)   | 0.855                  | 0.010                       | 0.351                         |

The multivariable model adjusts for: age, sex, BMI, diabetes, hypertension, hyperlipidaemia, smoking status, indication for coronary angiography, antiplatelet medication use, beta blocker use, calcium antagonist use, RAAS medication use and statin use. Interaction terms were tested in the univariable and the multivariable model. Ref. = reference group for interaction analysis.

 CI 95% confidence interval, univar. univariable, multivar. multivariable, NTproBNP N-terminal pro brain natriuretic peptide, hsCRP highsensitivity C-reactive protein, CysC Cystatin C, MPO myeloperoxidase, hsTnI high-sensitivity troponin I. Significant p-values are printed in bold.

aBetas are given for a 10-unit increase in biomarker level.
bBetas are given for a 100-unit increase in biomarker level.

When adjusted for baseline characteristics, certain differences in biomarker levels remained statistically significant. Post-hoc testing revealed that multivariable-adjusted NTproBNP levels were significantly higher in Malays than in Chinese and Caucasians (72.1 vs. 34.4 and 41.1 pmol/l, p < 0.001 and p = 0.005, respectively). MPO levels were higher in Caucasians than in Indians (32.8 vs. 27.2 ng/ml, p = 0.026), hsTnI levels were higher in Malays than in Caucasians and Indians (33.3 vs. 16.4 and 17.8 ng/l, p < 0.001 and p = 0.029) and hsTnI levels were higher in Chinese than in Caucasians (23.3 vs. 16.4, p = 0.031). High-sensitivity CRP levels and CysC levels (additionally adjusted for renal failure) did not differ among the ethnic groups.

### Modifying effect of ethnicity on association between biomarker levels and SYNTAX score

We tested interactions of ethnicity with biomarker levels for SYNTAX score (univariable and multivariable, Table 3 and Fig. 2). We found a significantly higher beta for NTproBNP levels in Malays than in Caucasians (only in the multivariable model, β 0.132 vs. β 0.020 p = 0.032), indicating a steeper
increase in SYNTAX score with every 100-unit increase of NTproBNP in Malays than in Caucasians. Also, we found a significantly higher beta, in both the univariable (β 1.517 vs. β 0.101, \( p=0.002 \)) and the multivariable (β 1.146 vs. β 0.016, \( p=0.017 \)) model, for MPO levels in Malays than in Caucasians, indicating that with a 10-unit increase of MPO, the SYNTAX score increases more steeply in Malays than in Caucasians. For hsTnI levels (per 100-unit increase) we found a lower beta for Malays than for Caucasians (β 0.003 vs. β 0.044, \( p=0.010 \)); however, this difference was abolished when adjusting for baseline differences.

Biomarker discrimination of high SYNTAX score

From ROC analysis (Fig. 3) we determined ethnicity-specific optimal biomarker cut-off values for the association with a SYNTAX score of \( \geq 18 \). These cut-offs (Table 4) correspond with the biomarker level at which the sum of sensitivity + specificity is largest. We found markedly different biomarker cut-off values across the ethnic groups. The optimal NTproBNP cut-off for Indians, for example, was 5-fold higher than for Caucasians: 255.77 pmol/l vs. 48.42 pmol/l. For hsCRP the cut-off was highest among Indians (8.95 μg/ml) and lowest in Chinese (1.69 μg/ml). The cut-offs for CysC and MPO were in the same order of magnitude among the ethnic groups.

The optimal cut-off for hsTnI was strikingly higher for Caucasians than the other ethnic groups (116.65 ng/l vs. 60.75 in Chinese, 6.65 in Indians and 17.15 in Malays). This indicates that especially in Indians and Malays, lower levels of hsTnI concur with more severe CAD.

Discussion

We observed inter-ethnic differences in biomarkers related to CAD. Also, the optimal cut-off levels at which these biomarkers offered discrimination of severe CAD varied substantially between ethnicities.
Differences in NTproBNP levels in the general population have mainly been reported between Blacks and Caucasians, with lower levels reported for Blacks [24]. In our cohort we find the highest fully adjusted levels for Malays and very high cut-off levels were found for Indians and Malays, indicating that in those ethnic groups NTproBNP levels correspond to less severe CAD than in Chinese and Caucasians.

High-sensitivity CRP levels are known to differ markedly among the ethnic groups in the general population with very low levels reported for Chinese and Japanese people [5, 25]. In a Singaporean cohort of individuals visiting the hospital for regular health checks, higher hsCRP levels were found in Indians as compared with Chinese, also when adjusted for confounders [26]. Notably, within the general US population, hsCRP levels were not able to differentiate between US Chinese with and without a future cardiovascular event, while some predictive power was observed for Caucasians, indicating different discriminating properties [25]. Similar to the general population cohorts, high adjusted hsCRP levels were found for Indians and Malays in the current study of CAD patients. In contrast to the general population cohorts, however, we also found higher hsCRP levels in Chinese than in Caucasians in our cohort (albeit not significant after adjustment for confounders). While cut-offs for Caucasians, Chinese and Malays were quite comparable,
Atherosclerosis in Blacks, but not in Whites or Hispanics. The severity of CAD is unresolved. In our study, we found a significantly stronger association of MPO levels with SYNTAX score among Malays than among Caucasians. However, comparable Cut-off levels for all ethnic groups were calculated. The association of MPO with severity of CAD among Malay patients deserves the attention of further research.

A striking difference was found for hsTnI levels, which were much higher in Malays (multivariably adjusted) than in Caucasians. Population levels have been reported to be comparable among Chinese, Indians and Malays in a Malaysian study [31], thus the high levels we observe in Malays can probably not be explained by higher baseline levels. High-sensitivity TnI levels can be elevated without actual myocardial necrosis [32]. Experimental data suggest that hsTnI can be released from cardiomyocytes by release of proteolytic degradation products, which can already occur with mild ischaemia. Mechanical stretch and ischaemia have also been demonstrated to increase cellular wall permeability, leading to leakage of troponins from the cytosol. Lastly, cardiomyocyte-derived vesicles, shed upon ischaemia, might be involved in the release of hsTnI without cardiomyocyte necrosis. These phenomena might explain the high hsTnI levels in stable Malay patients, in the absence of necrosis (Table 2).

Cut-off levels of hsTnI for high SYNTAX score were much higher in Caucasians than in the other ethnic groups, indicating that cardiomyocyte damage is greater in Caucasians than in the other ethnic groups at a SYNTAX score of 18 points. One of the explanations for this feature could be preconditioning occurring in more severe, unstable CAD resulting in resilient myocardium releasing less troponin upon prolonged ischaemia [33]. The Caucasian ethnic group most often presented with stable CAD, thus cardiomyocyte preconditioning would have occurred least in this group.

In summary, our results suggest that inter-ethnic differences in biomarker levels exist and that the association of these biomarkers with the extent of disease differs by ethnicity. Prior to implementation of biomarkers into clinical practice, their performance should be evaluated in an ethnicity-specific manner [34].

**Limitations**

Biomarker levels in our study were measured in arterial blood; these levels may differ from venous levels. However, Martin et al. showed that arterial and venous levels of hsCRP and NTproBNP did not differ [35]. Also, levels of NTproBNP, hsCRP, CysC and MPO were measured using an in-house ELISA method instead of clinically standardised

| Biomarker     | Ethnicity | Optimal cut-off | AUC (95% CI) |
|---------------|-----------|----------------|--------------|
| NTproBNP (pmol/l) | Caucasian | 48.42          | **0.594 (0.543–0.645)** |
|               | Chinese   | 120.92         | **0.596 (0.532–0.659)** |
|               | Indian    | 255.77         | 0.584 (0.455–0.712)    |
|               | Malay     | 154.41         | **0.621 (0.515–0.726)** |
| hsCRP (μg/ml)  | Caucasian | 2.40           | **0.589 (0.538–0.641)** |
|               | Chinese   | 1.69           | 0.547 (0.481–0.613)    |
|               | Indian    | 8.95           | 0.550 (0.425–0.675)    |
|               | Malay     | 3.04           | 0.502 (0.395–0.608)    |
| CysC (μg/ml)   | Caucasian | 0.74           | **0.555 (0.501–0.609)** |
|               | Chinese   | 0.86           | 0.527 (0.460–0.594)    |
|               | Indian    | 0.85           | 0.602 (0.475–0.728)    |
|               | Malay     | 0.60           | 0.571 (0.466–0.677)    |
| MPO (ng/ml)    | Caucasian | 30.17          | **0.601 (0.549–0.653)** |
|               | Chinese   | 36.84          | 0.532 (0.467–0.598)    |
|               | Indian    | 22.11          | 0.525 (0.405–0.644)    |
|               | Malay     | 33.40          | **0.611 (0.508–0.715)** |
| hsTnI (ng/l)   | Caucasian | 116.65         | **0.618 (0.566–0.671)** |
|               | Chinese   | 60.75          | **0.605 (0.540–0.669)** |
|               | Indian    | 6.65           | 0.614 (0.499–0.729)    |
|               | Malay     | 17.15          | **0.711 (0.619–0.804)** |

Results from ROC analysis, stratified by ethnicity. The optimal cut-off corresponds to the biomarker level at which the largest sum of sensitivity + specificity is found. The AUC is presented with its 95% confidence interval. When the confidence interval does not contain 0.5 (printed in bold), the cut-off of that biomarker is significantly associated with a SYNTAX score ≥ 18 points.

ROC receiver operating characteristics, AUC area under the curve, NTproBNP N-terminal brain natriuretic peptide, hsCRP high-sensitivity C-reactive protein, CysC cystatin C, MPO myeloperoxidase, hsTnI high-sensitivity troponin I.
assays. Therefore, generalisability of the biomarker levels described in our cohort is limited. Future studies should focus on examining inter-ethnic differences in biomarker levels measured in venous blood, measured by clinically validated assays.

Although follow-up data for all-cause death were available, an ethnicity-specific mortality analysis was not possible due to small numbers of fatalities in certain ethnic groups (e.g. n = 3 in the Indian ethnic group). Follow-up data on other adverse cardiovascular events were unfortunately not available.

We have adjusted our results for baseline differences among the ethnic groups as much as possible. However, profound differences were observed at baseline, which might not have been completely abolished after adjustment. In larger multi-ethnic cohorts propensity-matched [36] analysis might be considered in order to further eliminate confounding. Also, details on nutrition, lifestyle and socio-economic factors were not available in this cohort; therefore we could not consider these (possibly confounding) factors in the analyses.

Conclusion

In a patient group undergoing coronary angiography we found inter-ethnic differences in levels of biomarkers, related to CAD. These differences persisted after correcting for baseline differences among ethnicities. Furthermore, certain biomarkers displayed inter-ethnic differences in the relationship of biomarkers to severity of CAD, with cut-off values from ROC analyses varying over a five-fold range between ethnicities.

Future research should focus on ethnicity-specific cut-off values of established CAD biomarkers.

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Conflict of Interest None declared.

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