Impact of faecal haemoglobin concentration on colorectal cancer mortality and all-cause death

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ABSTRACT

Objective: To assess the effect of an incremental increase in faecal haemoglobin (f-Hb) concentration on colorectal cancer (CRC) mortality and all-cause death.

Design: We conducted an observational study of cohorts over time based on two population-based CRC screening programmes.

Setting: Two cities of Taiwan.

Participants: 1233 individuals with CRC (217 prevalent cases and 1016 incident cases) and 2640 with colorectal adenoma (1246 prevalent cases and 1394 incident cases) found in the two cohorts of 59 767 and 125 976 apparently healthy individuals, aged 40 years and above, who had been invited to participate in screening since 2001 and 2003, respectively.

Main outcome measures: Death from CRC and all-cause death ascertained by following up from the entire two cohorts over time until 2009.

Results: The effect of an incremental increase in f-Hb on the risk for CRC mortality was noted, increasing from a slightly increased risk for the category of f-Hb of 20–49 ng Hb/mL (adjusted HR (aHR)=1.09; 95% CI 0.68 to 1.75) to 11.67 (95% CI 7.71 to 17.66) for the group with f-Hb of 450 ng Hb/mL as compared with the group considered baseline with f-Hb of 1–19 ng Hb/mL (p<0.001). A similar but less marked increasing trend was found for all-cause mortality, aHR increasing from 1.15 (95% CI 1.07 to 1.24) for the group with f-Hb of 20–49 ng Hb/mL to 11.15 (95% CI 1.07 to 1.24) for the group with f-Hb of 450 ng Hb/mL.

Conclusions: We substantiated the impacts of an incremental increase in f-Hb on the risk for death from CRC and all-cause death, consistently showing a significant gradient relationship. Both discoveries suggest that f-Hb may not only make contribution to facilitating individually tailored screening for CRC but also can be used as a significant predictor for life expectancy.

INTRODUCTION

Use of traditional guaiac faecal occult blood tests (gFOBTs) has been shown to reduce mortality in population-based screening programmes for colorectal cancer (CRC). The newer faecal immunochemical tests (FITs) for haemoglobin (Hb) have many advantages, are recommended in recent guidelines, and have been introduced in many countries. Moreover, there are plans to begin screening using FIT in other countries, and those that still use gFOBT are planning to evolve their programmes and adopt FIT. Quantitative FIT values, generally using automated immunoturbidimetric analytical systems, have the significant advantage that they allow the end-user to select the faecal haemoglobin (f-Hb) to be used as the cut-off for referral for further investigation, usually colonoscopy. Previous study has shown that, as the cut-off concentration is decreased, positivity and sensitivity increase at the expense of specificity and positive predictive value. Moreover, there is some evidence, from data generated in screening programmes using FIT, that f-Hb is directly related to the severity of colorectal disease present in those screened.
f-Hb above a predetermined cut-off concentration chosen to suit the requirements of the particular programme, such as the colonoscopy resource available. However, a recent study has demonstrated the potential usefulness of using baseline f-Hb in predicting the risk of colorectal neoplasia in those who attend screening but who are classified as negative at first screen and are therefore not further investigated at that time. This observed trend for increasing f-Hb to be associated with increasing risk of colorectal neoplasia prompted us to speculate whether the causal effect of f-Hb on CRC mortality might also exist. It also would be of interest to elucidate whether a greater risk of f-Hb can lead to higher all-cause death. To our knowledge, these have never been investigated to date.

The aim of this study was to assess whether the effect of an incremental increase in f-Hb on the risk of death from CRC could be found in a population that has undergone screening using FIT and has been followed up for 3.5 years of median follow-up period. The impact of an incremental increase in f-Hb on all-cause death was also explored.

MATERIALS AND METHODS
Study cohorts
The study population comprised two cohorts who had been offered population-based screening for CRC using FIT: the first were residents of Keelung city, which is situated in the North of Taiwan, and the second resided in Tainan county, which is situated in the far South. Both cohorts underwent screening for CRC as part of community-based integrated screening programmes, in which tests for five neoplastic and three non-neoplastic diseases were performed. The details of study design, screening protocol, policy for referrals and diagnostics, surveillance and the evaluation of outcomes have been described in full elsewhere. Some characteristics of the two cohorts are delineated as follows.

The Keelung cohort included participants in the Keelung community-based integrated screening programme from 1999 until present. The details of the screening protocol for CRC have been presented in previous reports. In brief, the screening population comprised residents aged 40–79 years who were screened at yearly intervals until 2008, and then biennially. There were 59 767 participants and entry since 2001 is ≥100 ng Hb/mL at the first or second screen. Duals for further investigation using colonoscopy was determined at the central laboratory. When samples were assessed with high Hb concentrations (≥1 000 ng Hb/mL) the analyser flagged out of range results for subsequent automated serial dilution for reanalysis to give quantitative f-Hb results ranging from 0 to 2 500 000 ng Hb/mL. The cut-off used to select individuals for further investigation using colonoscopy was 100 ng Hb/mL.

Data collection
Participants with f-Hb ≥100 ng Hb/mL at the first or repeated screens were referred for colonoscopy on each occasion throughout the follow-up period from entry until the end of 2009. For adenoma, we classified abnormal findings by the number, site and size of adenomas found during each colonoscopy. Individuals with more than a single lesion were classified according to the most advanced. Prevalent adenoma and CRC cases were defined as the cases detected at first screen. Incident CRC cases included screen-detected cases (presymptomatic cases) detected by subsequent screens, interval
cancers and cancers from non-participants of subsequent screen (symptomatic CRC). Interval cancers and cancers from non-participants were obtained through the linkage of the entire cohort with the nationwide cancer registry data and the follow-up of the entire cohort until the end of 2009. The national cancer registry is a nationwide resource with high coverage and accuracy (the percentage recorded as ‘death certificate only’ is less than 1% for CRC) but the time to reporting is around 3 years. In the present study, anthropometric measurements (including blood pressure, height, weight and waist circumference), information on demographic characteristics, family history and life-style (obtained from questionnaires) and a series of biochemical variables, including plasma triglycerides, high-density lipoprotein cholesterol and fasting glucose concentrations, were investigated and recorded at each visit.

Note that data derived from both cohorts and used for the following mortality data analysis consisted of the main variable of interest, f-Hb and other confounding factors, all of which were collected at baseline (first screen) and the corresponding variables also collected at subsequent screens, which allowed us to do the following time-independent (baseline) and the time-dependent Cox proportional hazards regression models (see below).

**Statistical analysis**

As far as descriptive analysis is concerned, we reported the detection rate at first screen (prevalence) for colorectal adenoma and CRC. It was calculated as the number of cases found divided by number of participants at first screen. By following up an adenoma-free or a CRC-free cohort (excluding individuals with CRC or colorectal adenoma detected at first screen), the incidence rates of CRC and colorectal adenoma were calculated as the number of incident cases identified during the follow-up divided by the number of person-years at risk.

A Cox proportional hazards regression model was used to investigate the possible causal effect of an incremental increase in f-Hb on time to death from CRC. Time to event for each participant used in the regression analysis was calculated as the time from date of entry to the screening programme until date of death, date of loss to follow-up or the end of the study (whichever came first).

The concentration of f-Hb was classified into eight incremental groups, the undetected, 1–19, 20–49, 50–99, 100–149, 150–249, 250–449 and ≥450 ng Hb/mL following the previous study. In the univariate analysis, we estimated the crude HR (cHR) for each covariate including different categories of f-Hb (main variable) and other confounding factors (including age, gender, smoking, drinking alcohol, body mass index (BMI), metabolic syndrome, diabetes mellitus and hypertension) collected at the same time as f-Hb. In the multivariable regression model, we tested whether each significant confounding factor identified from the univariate analysis was statistically significant when age, gender and f-Hb were retained in the model. The baseline f-Hb was generally used in our time-independent analysis. However, to assess the effect of dynamic changes in f-Hb on the risk for death, by screening round, a time-dependent, Cox proportional hazards regression model was used, treating the repeated data on f-Hb collected from subsequent screens as time-dependent covariates. Note that the last updated f-Hb value near each time to event was used as the covariate of f-Hb included in the time-dependent Cox proportional hazards regression model.

**RESULTS**

**CRC and neoplasia**

Table 1 shows the 217 CRC and 1246 adenoma that were identified at the first (prevalence) screen. For colorectal adenoma, the detection rate at the first screen increased with f-Hb, from 11.2% among those screened with f-Hb of 100 to 149 ng Hb/mL to 15.6% among those with f-Hb≥450 ng Hb/mL. Note that it was not possible to detect adenoma and CRC for f-Hb below 100 ng Hb/mL as these negative cases would not be referred to undergo colonoscopy. A similar increasing trend was found for CRC. Participants who were disease free at first screen were then followed up (the median follow-up time for the two cohorts combined was 3.5 years (IQR 1.4–5.3)) to identify 1016 incident cases of CRC and also 1394 incident cases of adenoma. The incidence rate (per 1000 persons) of colorectal adenoma increased with f-Hb, from 1.17 for f-Hb of 1 to 19 ng Hb/mL to 21.2 for f-Hb≥450 ng Hb/mL (table 1). A similar increasing trend was also noted for CRC.

**CRC mortality**

Table 1 and figure 1 show that the CRC mortality rate (per 1000 person-years) increased as the f-Hb increased from 0.26 for 1 to 19 ng Hb/mL to 3.54 for f-Hb≥450 ng Hb/mL (the far right column of table 1). Table 2 documents the cHR for age, gender, smoking, drinking alcohol, BMI, metabolic syndrome, diabetes mellitus, hypertension and f-Hb based on Cox proportional hazards regression models. In the univariate analysis, we noted a direct effect of f-Hb on the risk for CRC death, with the risk for those in the group with f-Hb of 20–49 ng Hb/mL being only slightly increased (cHR=1.02; 95% CI 0.65 to 1.60) compared with the baseline category (1–19 ng Hb/mL), whereas for f-Hb≥450 ng Hb/mL, the risk increases 15-fold (CI 10.05 to 23.57; trend test, p<0.001). This significant impact persisted after adjustment for age, gender and diabetes mellitus in a multivariable analysis (trend test, p<0.001). Regarding other confounding factors, it should be noted that only diabetes mellitus remained statistically significant when age, gender and f-Hb were retained in the multivariable regression model. Table 2
<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Number of participants</th>
<th>Adenoma Prevalent cases</th>
<th>Detection rate</th>
<th>Incident cases</th>
<th>Incidence rate</th>
<th>CRC Prevalent cases</th>
<th>Detection rates</th>
<th>Incident cases</th>
<th>Incidence rate</th>
<th>Death</th>
<th>Mortality rate</th>
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<tr>
<td>40–49</td>
<td>57 153</td>
<td>123</td>
<td>0.22</td>
<td>147</td>
<td>0.68</td>
<td>32</td>
<td>0.06</td>
<td>89</td>
<td>0.41</td>
<td>13</td>
<td>0.06</td>
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<td>50–59</td>
<td>53 525</td>
<td>454</td>
<td>0.85</td>
<td>397</td>
<td>2.14</td>
<td>41</td>
<td>0.08</td>
<td>180</td>
<td>0.96</td>
<td>32</td>
<td>0.17</td>
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<td>60–69</td>
<td>43 663</td>
<td>514</td>
<td>1.18</td>
<td>699</td>
<td>4.63</td>
<td>66</td>
<td>0.15</td>
<td>348</td>
<td>2.29</td>
<td>57</td>
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<td>≥70</td>
<td>31 402</td>
<td>155</td>
<td>0.49</td>
<td>151</td>
<td>1.22</td>
<td>78</td>
<td>0.25</td>
<td>399</td>
<td>3.20</td>
<td>110</td>
<td>0.88</td>
</tr>
<tr>
<td>Gender</td>
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</tr>
<tr>
<td>Male</td>
<td>75 267</td>
<td>816</td>
<td>1.08</td>
<td>803</td>
<td>3.14</td>
<td>123</td>
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<tr>
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<td>430</td>
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<td>591</td>
<td>1.40</td>
<td>94</td>
<td>0.09</td>
<td>514</td>
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<td>f-Hb (ng Hb/mL)</td>
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<tr>
<td>Undetected</td>
<td>86 574</td>
<td>NK</td>
<td>0.00</td>
<td>394</td>
<td>1.52</td>
<td>NK</td>
<td>0.00</td>
<td>226</td>
<td>0.87</td>
<td>49</td>
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<tr>
<td>1–19</td>
<td>62 235</td>
<td>NK</td>
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<td>300</td>
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<td>NK</td>
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<td>88</td>
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<td>NK</td>
<td>0.00</td>
<td>95</td>
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<td>12</td>
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<td>318</td>
<td>11.16</td>
<td>103</td>
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<td>150–249</td>
<td>2062</td>
<td>258</td>
<td>12.51</td>
<td>92</td>
<td>14.15</td>
<td>20</td>
<td>0.97</td>
<td>36</td>
<td>5.04</td>
<td>8</td>
<td>1.10</td>
</tr>
<tr>
<td>250–449</td>
<td>1522</td>
<td>221</td>
<td>14.52</td>
<td>72</td>
<td>17.39</td>
<td>21</td>
<td>1.38</td>
<td>38</td>
<td>8.05</td>
<td>10</td>
<td>2.09</td>
</tr>
<tr>
<td>≥450</td>
<td>2881</td>
<td>449</td>
<td>15.58</td>
<td>158</td>
<td>21.15</td>
<td>142</td>
<td>4.93</td>
<td>179</td>
<td>22.05</td>
<td>31</td>
<td>3.54</td>
</tr>
<tr>
<td>Total</td>
<td>185 743</td>
<td>1246</td>
<td>0.67</td>
<td>1394</td>
<td>2.06</td>
<td>217</td>
<td>0.12</td>
<td>1016</td>
<td>1.49</td>
<td>212</td>
<td>0.31</td>
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</tbody>
</table>

The baseline f-Hb was used.

f-Hb, faecal haemoglobin; NK, not known.
also shows the adjusted HR (aHR) when the repeated measurements of f-Hb were treated as time-dependent covariates in the multivariable analysis. The aHR for CRC mortality increased from 1.09 (95% CI 0.68 to 1.75) for f-Hb of 20–49 ng Hb/mL to 11.67 (95% CI 7.71 to 17.66) for f-Hb ≥ 450 ng Hb/mL.

**All-cause death**

As shown in figure 2, a similar but less marked increasing trend was observed for all-cause death (per 1000 person-years), being 7.93 for those with the undetectable f-HB, 9.65 for 1–19, 11.06 for 20–49, 13.74 for 50–99, 13.44 for 100–149, 18.21 for 150–249, 21.73 for 250–449 and 21.36 for ≥ 450 ng Hb/mL, respectively.

The effect of f-Hb on all-cause death with the multi-variable Cox proportional hazards regression model is documented in table 3. After adjusting for age and gender, the impact of f-Hb on all-cause death was statistically significant (trend test, p<0.0001) regardless of the model based on baseline or time-dependent f-Hb.

**DISCUSSION**

Our findings have significant implication for CRC screening in many countries with FIT. This is the first population-based study, using data on population-based screening for CRC with FIT, to demonstrate the effect of an incremental increase in f-Hb on CRC mortality and all-cause death. These findings strongly suggest that we could make a much better use of quantitative measures of f-Hb as previously proposed. The increasing f-Hb in groups, which strongly allude to an increase in CRC mortality, can be used to stratify the underlying population into different risk groups in order to consider the possibility of individually tailored screening policy of interscreening interval with reference to the values of f-Hb at baseline to reduce false-negative cases and false-positive cases. To avoid the former, participants with higher f-Hb but not yet classified as worthy of investigation for colorectal neoplasia at first screen may need a shorter interscreening interval with FIT or any further assessment of risk for CRC. On the other hand, the interval between repeated FIT screens could be extended to avoid false-positive cases for those with a lower f-Hb, particularly 50 ng Hb/mL or below. However, further studies are required to elucidate the precise link between f-Hb and objective proposals for individualised screening policies that will be widely adopted. Moreover, the optimal cut-off for individually tailored screening should be considered on the basis of cost-effectiveness analysis. This will become the subject of future research.

It is interesting to note the impact of f-Hb on all-cause death. It could be argued that all-cause death may be mainly attributed to CRC deaths. However, the effect of f-Hb on non-CRC deaths was still statistically significant (p<0.001 for trend test) after excluding all CRC deaths. One biological explanation for this effect might be that an increase in f-Hb leads to a decrease in systemic blood Hb concentration, which has been demonstrated in a previous study to be a strong predictor of all-cause death. However, we found the impact of f-Hb on all-cause death was attenuated, but still an independent predictor (aHR increased from 1.13 (1.04 to 1.22) for 20–49 ng Hb/mL to 1.63 (1.41 to 1.88) for 450 ng Hb/mL or greater; trend test p<0.001) after adjusting for blood Hb concentration(low Hb vs higher Hb (<13 g/dL for men and <12 g/dL for women) with aHR=1.88 (95% CI 1.77 to 1.99)). This finding may reduce this possibility but this postulate still needs to be verified. Another possible explanation is that an increase in f-Hb is a reflection of systemic inflammation, as is the case with...
Table 2  Crude HRs and adjusted HRs (aHR) for CRC death in a CRC screening programme

<table>
<thead>
<tr>
<th>Variables</th>
<th>Univariate analysis</th>
<th>Multivariate analysis</th>
<th>Time-dependent multivariate analysis</th>
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<tr>
<td></td>
<td>HR</td>
<td>95% CI</td>
<td>aHR</td>
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<tr>
<td>Age (years)</td>
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<tr>
<td>Comparison</td>
<td>1.08</td>
<td>(1.07 to 1.09)**</td>
<td>1.08</td>
</tr>
<tr>
<td>Gender</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Male vs female</td>
<td>1.65</td>
<td>(1.26 to 2.16)**</td>
<td>1.30</td>
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<td>Smoking</td>
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<tr>
<td>Yes vs no</td>
<td>1.21</td>
<td>(0.89 to 1.66)</td>
<td>1.30 (0.99 to 1.70)</td>
</tr>
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<td>Drinking alcohol</td>
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<td></td>
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<tr>
<td>Yes vs no</td>
<td>0.91</td>
<td>(0.65 to 1.29)</td>
<td>0.98</td>
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<tr>
<td>BMI (kg/m²)</td>
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<td>≥30 vs &lt;30</td>
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<td>(0.76 to 1.95)</td>
<td>0.98</td>
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<tr>
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<td>(0.72 to 1.33)</td>
<td>1.22 (0.99 to 1.70)</td>
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<tr>
<td>Yes vs no</td>
<td>2.22</td>
<td>(1.59 to 3.08)**</td>
<td>1.69</td>
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<td>Hypertension</td>
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<tr>
<td>Yes vs no</td>
<td>1.49</td>
<td>(1.13 to 1.96)*</td>
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<td>f-Hb (ng Hb/mL)</td>
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<tr>
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<td>(0.57 to 1.20)</td>
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<td>1–19</td>
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<td>(0.67 to 2.27)</td>
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<td>50–99</td>
<td>1.23</td>
<td>(1.34 to 5.81)</td>
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<td>2.79</td>
<td>(2.15 to 9.33)</td>
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<td>(4.58 to 17.31)</td>
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<td>250–449</td>
<td>8.90</td>
<td>(10.05 to 23.57)**</td>
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<td>≥450</td>
<td>15.39</td>
<td>(10.05 to 23.57)**</td>
<td></td>
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<tr>
<td>(trend test)**</td>
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</table>

**p<0.001; *p<0.01.
Smoking status (yes: current or former smokers).
Drinking alcohol status (yes: current or former drinkers).
The metabolic syndrome was defined according to NCEP-modified ATP III criteria with the adjustment of waist circumference for Asian.
The values of f-Hb used in univariate and multivariate analysis were based on baseline (first screen) f-Hb.
The values of f-Hb used in time-dependant multivariate analysis were based on baseline (first screen) f-Hb and also repeated (subsequent screen) f-Hb.
BMI, body mass index; CRC, colorectal cancer; f-Hb, faecal haemoglobin.

Figure 2  Cumulative all-cause death by an incremental increase in baseline faecal haemoglobin concentration.
both chronic diseases (such as type 2 diabetes mellitus and cardiovascular disease) and CRC, in which there is an over expression of a number of inflammatory biomarkers such as tumour necrosis factors α and interleukin I–VI.24–27 We suggest that the impact of f-Hb on all-cause death may be mediated through these inflammatory biomarkers. However, the exact mechanisms require further investigation.

The strengths of this study are that it examines the impact of f-Hb on CRC mortality and all-cause deaths in two large cohorts of individuals over a long period of 8 years, making allowance for other possible confounding correlates (such as diabetes mellitus). As the main focus was the effect of an incremental increase in f-Hb on CRC mortality, we have to control other confounding factors that may affect our main postulate because risk factors that are responsible for the occurrence of colorectal neoplasm may also lead to an increase in CRC mortality that is determined by the incidence and survival of CRC once our denominator is based on participants rather than colorectal neoplasm cases. To enhance the internal validity of our finding on the effect of f-Hb on CRC mortality, we could only measure f-Hb from participants involved in the screening programme. Thus, only measuring f-Hb among participants and identifying the outcomes from these participants may be subject to selection bias if the baseline f-Hb in participants were different from that in non-participant. This may suggest our finding may not be generalised to the entire underlying population but may be only applicable to those who had undertaken the screening programme. We believe such generalisability of our finding was not so serious because the Keelung cohort has almost covered the eligible population and the second cohort covered a large size of population. Moreover, as our community-based screening programme is a multiple service screening programme covering a range of different cancers and chronic diseases it seems unlikely to have the unknown selective factors leading to selection bias if the baseline f-Hb in participants were different from that in non-participant. This may suggest our finding may not be generalised to the entire underlying population but may be only applicable to those who had undertaken the screening programme. We believe such generalisability of our finding was not so serious because the Keelung cohort has almost covered the eligible population and the second cohort covered a large size of population. Moreover, as our community-based screening programme is a multiple service screening programme covering a range of different cancers and chronic diseases it seems unlikely to have the unknown selective factors leading to selection bias if the baseline f-Hb in participants were different from that in non-participant. This may suggest our finding may not be generalised to the entire underlying population but may be only applicable to those who had undertaken the screening programme.

Potential weaknesses are several fold. First, the study was carried out over a period of time when some changes in procedures and practices might have occurred in spite of the ongoing total quality management techniques being applied throughout. Moreover, although it is now widely recognised that f-Hb is dependent on age and gender,9 we did not specifically assess the effect of f-Hb on CRC mortality and all-cause death in potential subgroups, but we undertook Cox proportional hazards’ regression model to explore the effects of a number of variables including age and gender. Second, as our main interest was focused on the effect of f-Hb on CRC mortality, we could only measure f-Hb from participants involved in the screening programme. Thus, only measuring f-Hb among participants and identifying the outcomes from these participants may be subject to selection bias if the baseline f-Hb in participants were different from that in non-participant. This may suggest our finding may not be generalised to the entire underlying population but may be only applicable to those who had undertaken the screening programme. We believe such generalisability of our finding was not so serious because the Keelung cohort has almost covered the eligible population and the second cohort covered a large size of population. Moreover, as our community-based screening programme is a multiple service screening programme covering a range of different cancers and chronic diseases it seems unlikely to have the unknown selective factors leading to selection bias if the baseline f-Hb in participants were different from that in non-participant. This may suggest our finding may not be generalised to the entire underlying population but may be only applicable to those who had undertaken the screening programme.

We substantiated the impact of an incremental increase in f-Hb on CRC mortality, from a slightly increased risk for the category of f-Hb 20–39 ng Hb/mL to approximately 12 times risk for death for f-Hb greater than 450 ng Hb/mL compared with the baseline group of f-Hb 1–19 ng Hb/mL. The similar but less remarkable increasing trend was also noted for all-cause death.
findings suggest that f-Hb can not only be used for risk stratification of the underlying population to facilitate individually tailored screening for CRC but may also be a significant predictor for life-expectancy.

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Contributors
L-SC was responsible for writing up the manuscript, data management and also statistical analysis. AM-FY was responsible for statistical analysis and the interpretation of results. CGF was involved in the interpretation of results, and the edition of the manuscript. SY-HC and JC-YF were in charge of the data collection, linked the data of the two cohorts with mortality data, and did partial data analysis. P-EW and S-CL contributed to the implementation of programme, data collection and the interpretation of findings. D-SL, Y-CL and H-MC were involved in auditing the clinical procedures, including colonoscopy examinations, the pathological confirmation of cases, the interpretation of the findings and discussion, and the revision of manuscript. H-HC provided the main conceptual framework, designed the study, interpreted the results, supervised data analysis and revised the manuscript.

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