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Does allostatic load predict incidental coronary events differently among sexes?

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ARTICLE INFO	A B S T R A C T
Keywords: Allostatic load Biomarkers Sex interaction Cardiovascular diseases Hazard Cox regression Older adults	Background: One measure to quantify the degree of dysregulation is allostatic load (AL). Typically, AL in- corporates information on diverse biomarkers and is associated with health outcomes such as cardiovascular diseases or the incidence of coronary events (C-E). Aims: This study investigates the predictive performance of different AL scoring methods on the incidence of coronary events (C-E). This study also elaborates sex differences in the baseline risks of C-E and the AL associated risks of C-E. Design: Longitudinal data analysis of the Heinz Nixdorf Recall Study (Risk Factors, Evaluation of Coronary Calcification, and Lifestyle) of 4327 participants free of C-E at study baseline aged 45–75. The data contains over 13 biomarkers measuring AL. Methods: After conducting multiple imputations on missing values on AL for 826 participants, the analysis sample consisted of N = 4327 participants. We applied the two most commonly used methods of AL scoring AL (count- based and Z-score) and a recently developed logistic regression weighting method (LRM) approach. Cox regression was used to predict the incidence of C-E for each AL score. Results were estimated without (M0) and with (M1) covariate adjustment, and in a final model (M2), with an interaction between AL and sex. Results: We found no violation of the proportional hazard assumption and significant differences in the survival curves between the sexes for C-E (Log-rank test: prob. > Chi ² = 0.000). In MO, all AL-scoring methods levelled, though the count-based method and LRM performed better than the Z-scoring method. The interaction analysis in M2 showed a significant multiplicative interaction for the count-based method (1.254; [1.066, 1.475]) and for the LRM (1.746; [1.132, 2.692]). The additive relative excess risk due to interaction (RERI) measure was negative for the count-based method (RERI = -1.967 ; [-3.778; -0.156]) and the LRM (RERI = -1.909 [-3.910; 0.091]), indicating subadditivity. Conclusion: AL scores are suitable for predicting C-E. Dif

1. Introduction

McEwen and Stellar [1] introduced the concept of allostatic load (AL) to consider the impact of long-term stress on physiological aspects

of health. AL combines stress theory with the concept of allostasis. The concept of allostasis differentiates between homeostasis and allostasis. Homeostasis describes the regulatory processes of the organism to maintain optimal stability. Allostasis is the reaction to changing

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environments that affects multiple biological systems [2]. The linkage between homeostasis, allostasis and stress theory exists in the stressor evaluation of the organism. McEwen and Gianaros [3] further differentiated the involvement of the brain in the allodynamic processes. Here, allostasis is the short-term adaption, while AL is the long-term maladaptation. More precisely, the primary outcomes of the adaption process are the acute responses of the organism to the stressor, which, e. g., expresses through anti-inflammatory reactions. Secondary outcomes result from chronic exposition to stress, concurrently chronic exposition to primary outcomes. Secondary outcomes induce changes in various regulatory processes, such as the metabolism and the cardiovascular system, that mutually reinforce each other. The adaption process to stress results in pathophysiological adaptions of the organism if exposition to secondary outcomes is permanent and not followed by recovery [4].

AL measures the pathophysiological adaptions of the organism through various biomarkers. Initially, Seeman [21] presented a summary score of ten biomarkers corresponding to different health domains to measure AL. The AL scores have been applied to predict numerous diseases, such as cardiovascular diseases (CVD), mortality, diabetes, depressive symptoms and cognitive limitations (for systematic reviews, see: [5,6]).

1.1. AL - C-E and sex

Regarding AL and its impact on mortality, it is vital to highlight its association with coronary events (C-E) and CVD because these are major causes of mortality [7]. Many empirical studies have confirmed the association of stress, AL and C-E/CVD (e.g., Ref. [8]). Research has shown that the association of AL and C-E/CVD seems sensitive towards stress intersections with race and deprivation (e.g. Refs. [9-11]. Gianaros and Jennings [12] provided biological evidence that further supports stress's impact on the body. They identified neurobiological mechanisms from brain to body and body to brain. Such insights provide a deeper understanding of the reciprocal pathophysiological processes causing C-E/CVD. Especially in the face of the increased prevalence of C-E/CVD among males, such processes can be different between the sexes, biologically and psychologically. McEwen [13] discussed biological sex differences in the autonomic nervous system (ANS) and hypothalamic-pituitary-adrenal (HPA) axis between the sexes, which has been proven to be more robust in females mice than in males mice [14]. Psychologically, gendered coping patterns and stress responses have been associated with increased AL levels [15,16]. However, research so far also found mixed evidence for differences by sex in the effects of AL on health outcomes [17–20]. Still, there is a lack of research on the sex gradient in incidence of C-E/CVD that integrates AL.

1.2. Scoring methods for AL

While the implications of AL on the organism are undoubted, the scoring algorithms of AL are not. Since the AL score's initial presentation by Seeman [21]; different AL scoring methods have evolved with mixed predictive performances (e.g., Refs. [22–24]). Due to AL combining multisystem information (e.g., neuroendocrine, metabolic, inflammatory), some challenges in constructing appear. First, the technical calculation of the index. Second, the relative importance of each biomarker in the AL score. Third, the amount of biomarker information from each domain in the final AL score. Fourth, the standardization of the AL scores within the population.

1.3. The present study

This study utilizes information on a total set of 13 biomarkers. It compares the most commonly used AL-scoring methods (count-based and Z-scoring method) and the recently developed logistic regression weighting method (LRM) by Li et al. [23]. This study also compares the

predictive performance of different AL scoring algorithms on the incidence of cardiovascular events. We emphasize sex differences in the analysis due to the higher prevalence of C-E for males. This study contributes to the methodological discussion of scoring AL and investigates the suitability of the AL concept for predicting C-E in general and between the sexes.

2. Material and methods

2.1. Sample

We used the Heinz Nixdorf Recall (HNR) study (Risk Factors, Evaluation of Coronary Calcification, and Lifestyle) as the database for the analysis. The data consists of randomly selected participants aged 45-75 years from mandatory residence registries of three adjacent cities in the Ruhr area (Bochum, Essen and Mülheim). Between 2000 and 2003, 4814 individuals were enrolled with a recruitment efficacy proportion of 56% [25]. After enrolment, study participants completed questionnaires, blood samples were drawn, and urine was taken. The questionnaires were conducted by computer-assisted personal interviews (CAPI) and contained information on health behaviours, social contexts, attitudes and sociodemographics [26]. Laboratory blood and urine tests provided pathophysiological information. In total, we observed 13 biomarkers relevant for scoring AL. The HNR study aimed to provide information on the first occurrence of nonfatal myocardial infarction and cardiac death. More detailed descriptions of the sampling process of the HNR study are available elsewhere [27,28].

2.2. Study endpoints

The endpoint was defined as incident coronary events (C-E), including unequivocally documented death related to coronary artery disease or incident nonfatal myocardial infarction that met the predefined study criteria [27,28]. The HNR-study asked participants to give medical records information to document emerging conditions during the observation period and validate self-reported diagnoses. Myocardial infarction was validated based on electrocardiographic signs, enzymes (creatinine kinase), troponin T or I and necropsy as nonfatal or fatal myocardial infarction. The outcome measure is binary, with a value of 1 indicating the onset of the condition. The incidence surveillance is ongoing. More details on sample characteristics, outcome and covariate distribution are displayed in Table 1.

2.3. Biomarkers

The 13 biomarkers represent different health domains, dividing into metabolic, immune-inflammatory, cellular, cardiovascular and renal systems. Information on the metabolic system stems from low-density lipoprotein cholesterol (LDL-C) (1) and high-density lipoprotein cholesterol (HDL-C) (2), triglycerides (3), blood glucose (4) and waisthip ratio (5). C-reactive protein (6) and the glycoprotein fibrinogen (7) reflect immune-inflammatory processes. Dehydroepiandrosterone sulphate (DHEA-S) measures cellular ageing (8) and homocysteine vascular cell damages (9). Systolic (10) and diastolic (11) blood pressure measures the cardiovascular system. The renal system contains information on serum albumin (12) and urine creatinine (13). Higher values of HDL-C and DHEA-S do not correspond to higher AL; therefore, we reverse-coded HDL-C and DHEA-S, according to previous studies on AL (e.g., Ref. [23,24]). High values on the biomarkers correspond to high values on the AL scores. More details on origin, related health domain, measurement and distribution of the biomarkers are described in Table 2.

Table 1

Descriptive statistics of outcome and covariates.

Outcome: 100% Incidence of C-E*: 4327 100% Does not apply 4054 93.69% Applies 273 7.95 (4.66) Males: Incidence of C-E*: 2050 100% Does not apply 1864 90.93% Applies 1866 9.07% Years to event 186 9.07% Years to event 87 3.82% Years to event 87 3.82% Years to event 87 3.82% Years to event 87 5.918 (7.74) Born in Germany 4327 100% Yeas 4016 92.81% No 311 7.19% Married/living with partner 3444 79.59% Living alone/divorced 883 20.41% SCED-97 (0-4) 4327 100% Q 47 1.09% I atus 126 2.91% <t< th=""><th>Measures</th><th>Ν</th><th>Mean (SD)/%</th></t<>	Measures	Ν	Mean (SD)/%		
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Does not apply 4021 92.93% Applies 306 7.07% Heart related medication 4327 100% Does not apply 3176 73.40% Applies 1151 26.60%	Diabetes	4327	100%		
Applies 306 7.07% Heart related medication 4327 100% Does not apply 3176 73.40% Applies 1151 26.60%	Does not apply	4021	92.93%		
Heart related medication 4327 100% Does not apply 3176 73.40% Applies 1151 26.60%	Applies	306	7.07%		
Does not apply 3176 73.40% Applies 1151 26.60%	Heart related medication	4327	100%		
Applies 1151 26.60%	Does not apply	3176	73.40%		
	Applies	1151	26.60%		

2.4. Scoring methods

2.4.1. Z-scoring method

The Z-scoring method Z-standardized each biomarker to a mean of 0 and a standard deviation of 1 and then calculates the sum-score (e.g., Ref. [29]). The obtained Z-standardized biomarker values are comparable to one another but not directly interpretable, though higher values on the Z-standardized AL score correspond to higher levels of AL. The Z-scoring method requires continuously measured biomarkers and accounts for variances in the biomarkers. However, each biomarker's relative importance and the number of biomarkers from certain health domains remain unreflected.

2.4.2. Count-based method

The simplest and most applied method to score AL is the count-based method. The count-based method relies on building risk quartiles for the biomarkers in the study sample [21]. If an individual's biomarker value falls into the risk quartile, the individual receives the value "1"; if not, the value is "0". The overall sum of the dichotomized biomarkers represents the AL score. We used the 90th percentile sample distribution of each biomarker as the cut-off values. Most studies so far used the 75th or 90th risk percentile as the cut-off values of the biomarkers [21,23,24]. Subsequently, we summarized the dichotomized biomarkers into one AL score. A value of 13 indicates the highest AL and zero the lowest AL. Another issue of the count-based method is the potential loss of information due to the dichotomization of the biomarkers. The count-based method does also not adjust for the relative importance of each

biomarker for the AL score and the number of biomarkers from each health domain.

2.4.3. The logistic regression method

The LRM addresses the relative importance of each biomarker for the AL score. We applied the LRM according to Li et al. [23]. First, a logistic regression on the outcome (C-E) with all Z-standardized biomarkers as predictors were calculated. Second, the Z-standardized biomarkers with their respective coefficient were multiplied and, third, summarized to an AL score. Thus, the LRM addresses an important issue the two most commonly applied AL-scoring methods do not address.

2.5. Confounding factors and covariate selection

To investigate the AL – C-E/CVD association, we also adjusted for confounding variables. These variables simultaneously affect the exposure (AL) and the outcome (C-E) and can confound any observed association between AL and C-E. We identify four primary confounding sources: (1) socio-demographic, (2) socio-economic, (3) health-behavioral and (4) health-related. Table 1 gives a detailed overview of the distributional properties of all measures.

- Age is an apparent socio-demographic confounder since it increases AL and the probability of experiencing a C-E simultaneously. Living alone at an older age or the experience of a marital disruption have been found to increase AL and the likelihood of cardiovascular diseases [30], especially for males [31, 32]. We used a continuous measure for age and a binary measure for marital status.
- (2) The migration background is often accompanied by lower social status and a higher stress exposition due to racial discrimination [33]. Similarly, socio-economic factors, such as the educational level, can influence the AL and C-E through health-behavioural pathways [34]. We added information on the direct migration background (binary), and we used the International Standard Classification of Education (ISCED) as a proxy for social status (categorical).
- (3) Health-behavioural confounders may affect AL and C-E positively or negatively [35], and therefore confound the association. Hence, we added information on the current smoking status (binary) and the sport participation (binary).
- (4) Health-related conditions, such as certain conditions and medications, could also confound the AL – C-E association. To account for confounding pre-existing health conditions, we added diabetes (binary) and any heart-related medication (binary) as covariates in the models.

2.6. Sample selection and missing values

The sample selection process is shown in Fig. 1. We addressed potential reverse causalities by excluding 432 participants reporting physician-diagnosed coronary artery disease (i.e., history of myocardial infarction or coronary revascularization) at baseline, resulting in a sample of 4482 remaining participants. Only 14 participants were excluded due to missing values on the outcome or control variable (N = 4368).

Due to the nature of sum-scoring, complete information on all biomarkers was required for the AL scores. However, 80.11% had full information, and 16.76% had only one and 2.20% only two missing values on all 13 biomarkers; thus, over 99% of the study participants had at least 11 out of 13 valid entries on the AL score. We imputed missing values on the AL scores for individuals with at least 11 or more valid entries on the biomarker and assumed missingness at random for the remaining biomarker information. In sum, 828 values for the participants with only one or two incomplete entries on the AL score were imputed, resulting in a final analysis sample of 4327 study participants.

Table 2

Biomarker and allostatic load score distribution between sexes.

All								
Domain	Biomarker	Unit	Mean	Std. Dev.	p90	Min	Max	Ν
(1)	LDL-C	mg/dl	147.018	36.123	193	4	360	4315
(1)	HDL-C*	mg/dl	58.765	17.182	81	7	171	4326
(1)	Triglyceride	mg/dl	147.819	102.038	247	13	1846	4324
(1)	Glucose	mg/dl	110.487	26.588	130	48	459	4323
(1)	Waist-Hip-Ratio	cm/cm	0.909	0.092	1.026	0.529	1.281	4315
(2)	C-reactive Protein	g/1	0.299	0.634	0.65	0.01	14	4315
(2)	Fibrinogen	mg/dl	332.834	75.606	434	124	876	4287
(3)	DHEA-S*	µg∕dl	844.911	118.767	953	0	1000	4034
(3)	Homocysteine	µmol/l	11.565	4.288	15.74	4.04	100	3985
(4)	Sys. blood Pressure	mmHg	132.788	20.78	158.5	70	229	4326
(4)	Dias. blood Pressure	mmHg	81.572	10.859	95.5	44	133.5	4326
(5)	Albumin	g/dl	34.438	273.805	44.5	5.4	13100	4128
(5)	Creatinine	mg/dl	0.923	0.22	1.13	0.2	7.12	4323
Raw	Z-Score	AL score	-0.115	5.128	6.05	-14.738	57.757	3499
Imputed	Z-Score	AL score	-0.077	4.796	5.476	-14.738	57.757	4327
Raw	Count score (p90)	AL score	1.255	1.379	3	0	9	3499
Imputed	Count score (p90)	AL score	1.259	1.279	3	0	9	4327
Raw	LRM score	AL score	-0.011	0.65	0.783	-5.007	3.674	3499
Imputed	LRM score	AL score	-0.006	0.618	0.744	-5.007	3.674	4327
Males								
(1)	LDL-C	mg/dl	147.491	35.286	191	38	360	2044
(1)	HDL-C*	mg/dl	51.633	14.496	70	7	155	2049
(1)	Triglyceride	mg/dl	164.938	121.382	275	17	1846	2048
(1)	Glucose	mg/dl	114.336	29.519	136	55	459	2049
(1)	Waist-Hip-Ratio	cm/cm	0.978	0.061	1.055	0.544	1.238	2046
(2)	C-reactive Protein	g/1	0.302	0.657	0.65	0.01	13.7	2043
(2)	Fibrinogen	mg/dl	326.219	75.8	425	124	876	2031
(3)	DHEA-S*	µg∕dl	807.933	130.081	932	0	996	1910
(3)	Homocysteine	µmol/l	12.208	3.891	16.38	4.75	57.72	1894
(4)	Sys. blood Pressure	mmHg	137.971	19.248	163	70	219.5	2049
(4)	Dias. blood Pressure	mmHg	84.441	10.471	98	44	133.5	2049
(5)	Albumin	g/dl	43.581	252.691	58.6	5.4	8480	1946
(5)	Creatinine	mg/dl	1.004	0.189	1.2	0.32	4.46	2049
Raw	Z-Score	AL score	2.068	4.609	7.501	-11.691	28.956	1652
Imputed	Z-Score	AL score	2.107	4.239	6.895	-11.691	28.956	2050
Raw	Count score (p90)	AL score	1.611	1.521	4	0	9	1652
Imputed	Count score (p90)	AL score	1.614	1.393	3	0	9	2050
Raw	LRM score	AL score	0.351	0.536	0.972	-3.996	3.674	1652
Imputed	LRM score	AL score	0.355	0.498	0.936	-3.996	3.674	2050
Females								
(1)	LDL-C	mg/dl	146.592	36.863	194	4	277	2271
(1)	HDL-C*	mg/dl	65.183	16.879	87	8	171	2277
(1)	Triglyceride	mg/dl	132.415	77.629	216	13	1390	2276
(1)	Glucose	mg/dl	107.02	23.102	124	48	380	2274
(1)	Waist-Hip-Ratio	cm/cm	0.848	0.069	0.932	0.529	1.281	2269
(2)	C-reactive Protein	g/1	0.296	0.612	0.66	0.01	14	2272
(2)	Fibrinogen	mg/dl	338.789	74.949	440	128	738	2256
(3)	DHEA-S*	µg∕dl	878.162	96.148	963	0	1000	2124
(3)	Homocysteine	µmol/l	10.983	4.541	14.8	4.04	100	2091
(4)	Sys. blood Pressure	mmHg	128.123	21.005	154	78	229	2277
(4)	Dias. blood Pressure	mmHg	78.99	10.551	92.5	50	121	2277
(5)	Albumin	g/dl	26.285	291.164	31.2	5.9	13100	2182
(5)	Creatinine	mg/dl	0.849	0.22	1.01	0.2	7.12	2274
Raw	Z-Score	AL score	-2.067	4.77	3.565	-14.738	57.757	1847
Imputed	Z-Score	AL score	-2.044	4.401	3.141	-14.738	57.757	2277
Raw	Count score (p90)	AL score	0.936	1.148	3	0	7	1847
Imputed	Counts core (p90)	AL score	0.939	1.07	2	0	7	2277
Raw	LRM score	AL score	-0.334	0.568	0.385	-5.007	2.795	1847
Imputed	LRMs core	AL score	-0.331	0.528	0.335	-5.007	2.795	2277

Note: *Values reversed during calculation of AL scores.

We used the multiple imputations (MI) procedure as an imputation technique in Stata 16.1 (StataCorp LP, College Station, Texas, USA). This MI procedure is based on Rubin [36]; and we included all covariates and the outcome in the imputation model as predictors. We also supplement the imputation model with the incidence of CVD as an auxiliary variable. Auxiliary variables can reduce the bias in imputation. According to the overall missingness of approx. 20% on the AL score, we calculated 20 MI datasets [37]. Robertson et al. [38] conducted a similar MI approach on imputing AL scores.

2.7. Statistical methods

We investigated the AL-associated hazard of C-E with Cox regression modelling [39]. Cox regressions are commonly applied in analyzing events, such as the incidence of C-E (e.g., Ref. [40]). We checked the proportional hazards assumption and the linear relation assumption by inspecting the Kaplan-Meier curve and the Schoenfeld residual plot and test [41]. We provide results before (M0) and after (M1) covariate inclusion for each AL-scoring method.

We evaluated the performance of the AL-scoring techniques by



Fig. 1. Flowchart of sample selection and imputation.

comparing the estimated hazard ratios and the model-fit parameters. As model-fit parameters, we reported Harell's C and Somers' D. The first expresses the balance between concordant and discordant pairs, and the latter is the correlation between predicted risk observed survival. Additionally, we inspected the receiver operator characteristic (ROC). The ROC displays the positive cases (incidence of C-E) that the models correctly classified (sensitivity) and the negative cases that the model correctly classified (specificity). We contrasted the sensitivity and specificity of the scoring methods by testing the equality in the ROCcurves between the scoring methods. Further, we reported pseudo-R² and the Akaike Information Criterion (AIC). Lower AIC values correspond to increases in model fit, and pseudo-R² values can be used to compare the models between one another.

We investigated sex differences through sex separated Kaplan-Meier curves and the log-rank test to compare the survival curves between females and males. In addition, we performed the analysis of the interaction between AL and sex in M2. As demonstrated by Vatcheva et al. [42]; considering interactions in hazard Cox regression can substantially improve the prediction of the main effects. We reported additive and multiplicative interaction measures. As a multiplicative measure, we included the product of sex and AL as a predictor in the hazard Cox regression. For the additive interaction measures, we followed the recommendation by Anderson et al. [43] and constructed a composite variable of sex and AL. For sex, we define female sex as the low-risk category, and for AL, we define having an AL score above the 90th percentile as the high-risk category. The risk combinations are then defined ordinally. The value 1, the reference category, equals the lowest risk combination (being female sex and not having high AL) and the value of 4, the highest risk combination (being male sex and having high AL). As additive interaction measures, we reported the relative excess risk due to interaction (RERI), the synergy index S and the attributable

proportion (AP) due to interaction [43].

3. Results

3.1. Descriptive results

Table 1 presents the sample characteristics of the analysis sample after imposing the selection criterion (Fig. 1) and imputation of the AL score values. The analysis sample consisted of 4327 participants, with 2050 males and 2227 females. The incidence of C-E was observed more often in males (9.07%) than in females (3.82%). On average, it took about eight years after study participation if a C-E occurred. For males, this duration was slightly lower than for females (7.77–8.32). The mean age of the analysis sample was 59.18 (SD = 7.74), and the majority of the analysis sample were married or cohabiting (79.59%), had no direct migration background (92.81%), were currently not smoking (76.50%) and had no diabetes (92.93%). Although we restricted the analysis sample to participants with no baseline C-E or CVD, 26.60% of the analysis sample had heart-related medications prescripted.

Table 2 shows the distribution of the biomarkers and AL scores. A more detailed table on the missing values of the biomarkers and AL score is available in Appendix Tables 1 and 2. Overall, the biomarkers values indicated higher levels of AL for males than for females, except for fibrinogen. Also, the raw HDL-C and DHEA-S values were lower for males than for females; however, higher values for these biomarkers correspond to a lower AL, and the HDL-C and DHEA-S values were reversed during the AL score calculation. Logically, males also had higher values on the AL scores. For the count-based scoring method, the mean AL score was 1.259, whereas the mean value for males was 1.614 and 0.939 for females. Similarly, the signs of the AL scores based on Zstandardization (Z-score and LRM) showed in opposite directions between the sexes, with a negative sign for females and a positive sign for males. We found no significant difference between raw and imputed AL scores in an unpaired t-test for mean differences (see also, Appendix Table 2).

3.2. Hazard Cox regressions on C-E

Fig. 2a shows the Kaplan-Meier curve for the observed (bolt lines) and predicted (dashed lines) values of the C-E between males (black colour) females (grey colour). Fig. 2a clearly illustrates a gap between the sexes in the survival probability for the predicted and observed values. The gap between the sexes further increased with the observation duration. For example, after ten years of observation, males had at least a 0.05 lower survivability than females. The log-rank test for equality of survival curves was also highly significant (prob. $> Chi^2 =$ 0.000), confirming significant differences in the survival function between the sexes. Fig. 2a also visually checked the proportional-hazards assumption and the linear relation. The larger the gap between observed and predicted lines within the sex, the more likely the violation of the proportional-hazards assumption. The Kaplan-Meier curve indicated no violation of the proportional-hazards assumption with a continuously decreasing trend in the survival for both sexes, which plateaued after 17 years of observation.

Table 3 summarizes the hazard Cox regression models for each AL score, where M0 consists only of the AL scores, M1 includes the covariates, and M2 includes the covariates and an interaction term between AL scores and sex. In M0, the AL score of the LRM overall provided the best model fits, with the lowest AIC, the highest pseudo-R² and a ROC-Area of 0.703. Z-scoring method and count-based method performed fairly similar, however the model fit statistics pseudo-R² and AIC indicated a slightly better performance for the count-based method (Z-scoring: pseudo-R² = 0.017; AIC = 4375.6; Count-based: pseudo-R² = 0.020; AIC = 4365.8). The Kendall rank parameters indicated a good fit for all models, with Harrell's C > 0.5 and Somer's D > 0.35. However, the ROC-test for equality between the models revealed significant



Fig. 2. a: Kaplan-Meier survival curve between sexes **b**: Cumulative hazards between sexes at different levels of AL. Note: **a**: relative hazards for males: 1.6167; relative hazards for females = 0.6582; Cox regression-based test for equality of survival: Log-rank Chi²(1) = 51.78; prob. > Chi² = 0.000. Note: **b**: Estimated cumulative hazards at different levels of the count-based AL-scoring method between the sexes. Mean AL = 1.258; SD = 1.278.

differences (prob. > Chi² = 0.012) in the sensitivity and specificity of the predictions between the scoring methods.

Adding covariates in M1 improved all models and levelled most of the initial differences in M0 between the AL scores, and all estimated hazard ratios remained highly significant. After covariate inclusion, the Z-scoring and count-based methods' AIC decreased by nearly 100. Somer's D increased by almost 0.1 to 0.467 for the Z-scoring method and 0.482 for the count-based method. Pseudo-R² values ranged from 0.043 to 0.046. The estimated ROC-Area was at least above 0.720 for all scoring methods. However, the ROC-test of equality between the models remained significant (prob. > $\text{Chi}^2 = 0.046$), indicating significant differences in the ROC-area between the scoring methods. In Appendix Fig. 1, we also present additional diagnostics for the PH assumption of M2 by displaying the Schoenfeld residuals graph. The global test was 0.950, and the single test for the interaction term was 0.644, indicating no violation of the PH assumption in M2.

M2 added the multiplicative interaction between AL score and sex (1 = male; 2 = female) to test if the effect of AL is different across sexes. All model-fit parameters increased in comparison to M1 for all scoring methods. The ROC-test for equality between the models turned insignificant (prob. > $\text{Chi}^2 = 0.077$), indicating no differences in sensitivity and specificity between the scoring methods in M2. The LR-test confirmed the increased fit of M2 over M1, which indicated that M1 is nested in M2 except for the Z-scoring method. Regarding the AL-sex interaction, the multiplicative interaction was insignificant for the Z-scoring method (1.254**: p < 0.01) and LRM (1.746*; p < 0.05). The hazard ratio for the multiplicative interaction effect was >1, while the hazard ratio for sex was <1, indicating that the AL hazard ratio effect is stronger for females than males.

To further illustrate the interaction between AL and sex, we plotted the cumulative hazards in Fig. 2b. It displays the cumulative hazards for males (black lines) and females (grey lines) for the best performing

method in M2 (count-based method). The cumulative hazards on the bolt solid lines were calculated by the mean score of the count-based AL score (1.259). The dotted lines were calculated by the cumulative hazards with two SD units increase from the mean (SD = 1.29; AL score \sim 3.9), reflecting the cumulative hazards for individuals with high levels of AL. The dashed lines are the cumulative hazards where the AL score was set to "0" to reference individuals with no AL. The difference between the bold black solid line (mean AL and males) and the bold grey solid line (mean AL and female) were much larger than the difference between the dotted black (high AL and male) and dotted grey line (high AL and female). The decrease in the differences of the cumulative hazards between the sexes at higher AL scores visualizes that although males have a much higher baseline hazard for experiencing C-E, the relative increase in hazards that is due to increased AL is larger for females than for males. Therefore, the interaction showed that the more elevated the AL score, the lesser the differences between the sexes in the cumulative hazards of C-E.

Following the recommendations of VanderWeele and Knol [44]; the inspection of additive interaction is recommended when the baseline risk for one group is higher than for the other group. We calculated additive interactions measures with the composite variable of AL and sex (Table 3; Model 2b) [43]. The hazard ratios for the composite variable are estimated in reference to the expected lowest risk group, which were females with low AL. For females with high AL, the Z-scoring method predicted a hazard ratio of 1.277, while for the count-based method, the hazard ratio was 3.254***, and for the LRM, the hazard ratio was 2.715***. For the count-based method and the LRM, being male sex and having high AL (highest risk group) had a lower hazard ratio than being female and having high AL. The relative excess risk due to interaction (RERI) is < 0, indicating sub-additivity of the interaction, which means that the increase in hazards is not consistent with the increase in risk groups. The attributable proportion due to interaction (AP) shows that 61% of the observed C-E incidences are due to interaction for

Table 3

Estimates and fit statistics of AL scoring methods on C-E with and without co-variate inclusion.

	-		LIUM
AL HR 1.085*	**	1.450***	3.133***
[1.071]	,1.100]	[1.352,1.556]	[2.630,3.732]
Somers' D 0.378		0.357	0.434
Harrell's C 0.689		0.678	0.717
AIC 4375.6		4365.8	4308.9
Pseudo-R ² 0.017		0.020	0.032
ROC-Area [CI] 0.673	0.644,	0.671 [0.635,	0.703 [0.674,
0.703]		0.697]	0.734]
ROC-M0 equality test: $Chi^2 =$	8.81; prob >	$Chi^2 = 0.012 \\$	
M1 with controls Z-Scor	ing	Count-based	LRM
AL HR 1.047*	**	1.213***	1.939***
[1.020	,1.072]	[1.114,1.321]	[1.499,2.508]
Somers' D 0.467		0.482	0.476
Harrell's C 0.733		0.741	0.738
AIC 4279.5		4271.8	4266.3
Pseudo-R ² 0.043		0.045	0.046
BOC-Area [CI] 0.723		0.731	0.727
[0.694]	,0.753]	[0.702,0.760]	[0.698,0.756]
ROC-M1 equality test $Chi^2 =$	6.32; Prob >	$Chi^2 = 0.042 \\$	
M2 multiplicative Z-Scor	ing	Count-based	LRM
interaction			
AL HR 1.031	[0.997,	1.142** [1.035,	1.568** [1.147,
1.066]		1.260]	2.143]
Sex = Female (2) HR 0.471^*	**	0.330*** [0.220,	0.568***
[0.346	, 0.640]	0.495]	[0.406, 0.794]
AL-sex interaction HR 1.030	0.987,	1.254** [1.066,	1.746* [1.132,
1.0751	-	1.475]	2.6921
Somers' D 0.469		0.485	0.481
Harrell's C 0.742		0.743	0.740
AIC 4279.7		4266.9	4262.1
Pseudo-R ² 0.044		0.046	0.047
BOC-Area [CI] 0.724		0.732	0.730
[0 695	0 753]	[0 703 0 761]	[0 700 0 759]
LR-Test M1 nested in 0.188	,01/00]	0.008	0.012
M2: $Prob > Chi^2$			
ROC-M2 equality test: $Chi^2 =$ Prob $> Chi^2$	5.11; Chi ² =	0.077	
M2 additive 7-Scor	ing	Count-based	IBM
interaction		Sount-Dascu	11111
measures			
Female: low AL (1. 1		1	1
reference)		1	1
Female; high AL (2; 1.277	[0.583;	3.254*** [1.996;	2.715** [1.371;
one risk factor) 2.795]		5.306]	5.377]
Male; low AL (3; the 2.475*	**	2.938*** [2.129;	2.672***
other risk factor) [1.836]	; 3.338]	4.056]	[1.971; 3.622]
Male; high AL (4; all 2.529*	**	3.226*** [2.152;	2.477***
risk factors) [1.686	; 3.792]	4.837]	[1.641; 3.738]
RERI -0.223	3 [-1.541;	-1.967* [-3.778:	-1.909 [-3.910:
1.0951	/	-0.156]	0.091]
AP -0.088	3 [-0.623:	-0.610 [-1.245:	-0.771 [-1.663:
0 4471		0.0261	0.122]
S 0.873 I	0.186:	0.531*** [0.247:	0.436** [0.116:
1.559]	- •	0.815]	0.757]

Note: 95% confidence intervals in brackets * p < 0.05, **p < 0.01, ***p < 0.001.

the count-based method. This amount even increased for the LRM to 77%. The synergy index (S) expresses whether the combinations of the hazard ratios of the risk groups is greater than the separate hazard ratios; in cases where the RERI is < 0, S would indicate no synergy when S = 1 or S = 0. For the count-based method, $S = 0.531^{***}$ and $S = 0.436^{***}$ for the LRM, but S is insignificant S = 0.873 for the Z-scoring method, and its upper C.I. includes values above 1. Overall, the additive interaction analysis confirmed Fig. 2b's illustration; at high levels of AL, the relative increase in hazards is more substantial for females than for males.

4. Discussion

This study investigated the predictive performance of three ALscoring algorithms on the incidence of C-E. using hazard Cox regression models. The data used for analysis was the HNR study, which is an ongoing surveillance of participants aged 45 to 75 living in Germany. The HNR study examined over 13 biomarkers related to the concept of AL, observed any incidence of C-E and CVD and enabled us to apply hazard Cox modelling on the C-E incidence.

This study highlighted sex differences between the incidence of C-E and the effect of AL by providing measures of multiplicative and additive interaction. Irrespective of the scoring algorithms, we found that AL is a solid predictor for the incidence of C-E. This study also contributes to AL being valid for predicting heart-related diseases [45] and to the recent discussions on the differences in performance of various AL-scoring algorithms [22–24].

In the naïve model without covariate inclusion, the LRM outperformed the other methods. With covariates, the AL scores remained highly significant, and the differences between the scoring methods were reduced. In the final adjusted models with an interaction between AL and sex, all scoring methods model fits increased even further, leading to insignificant differences in sensitivity and specificity between the models. Nevertheless, the interaction for the Z-scoring method was insignificant, whereas the interactions for the count-based method and LRM were significant. In covariate-adjusted models, the differences between AL-scoring methods are minor, but the range of the AL score should be reflected when incorporating interaction terms. We also confirmed substantial sex differences in the incidence of C-E, with males having a higher probability of a C-E incidence. However, with increasing levels of AL we found that the relative increase in hazards is larger for females than for males.

4.1. Strengths and limitations

One strength of this study is comparing three different AL scoring algorithms based on the same set of 13 biomarkers. There numerous other scoring AL scoring techiques that either rely on Z-standardization of the biomarkers, such as the factor loading method (e.g. Ref. [46]), or defininig risk-profiles, such as the grade of membership method (e.g., Ref. [47]). However, with increasing amount of specifications in scoring the comprabibility decreases. Recently, McLoughlin et al. [24] presented a genuine discussion and comparison of different scoring algorithms in a nationally representative cohort of older persons living in community-dwellings. They advocated the use of standardized, comparable AL scores across investigations. At the same time, they also recommended using sex-specific AL scores and incorporating medication information either directly in the AL-scoring procedure or as a covariate, as done by Piazza et al. [48]. Recently, Liu et al. [49] even introduced a scoring procedure of AL using the item response theory, which shows that the discussions on optimal AL scoring algorithms are still ongoing. With our chosen scoring methods, we achieved comparability across previous studies, and we also included heart-related medication as a covariate. Though, we did not provide sex-specific AL scores because we investigated sex differences through modelling interactions between AL and sex.

The analysis of interaction is another strength of our study. Interaction analysis allowed us to investigate sex differences in the AL related hazards of C-E incidence. Following the recommendations of Vander-Weeele and Knol [44]; we provided measures of multiplicative and additive interactions, which is important for three reasons. Although multiplicative interactions are most commonly used when assessing interactions, their interpretations can be biased when baseline risks between groups differ largely. The Kaplan-Meier curve revealed that the baseline hazards for experiencing C-E were substantially higher for males; thus, the multiplicative interaction measures could be biased. Secondly, it is possible to observe null or negative multiplicative interactions while also having positive additive interactions. Therefore, relying solely on multiplicative interaction could lead to wrong conclusions [44]. Thirdly, the interpretation of additive interactions is of high public health relevancy. The additive interaction measures identify the sub- or riskgroups for whom an intervention is more effective.

Without the interaction analysis, the large sex differences in the C-E incidence and AL levels could lead to the conclusion that AL is more important in males than in females. Our analysis revealed that for C-E prevention, females could benefit more from a potential allostatic load intervention than males, despite having lower baseline risks than males. Reporting additive and multiplicative interaction measures should become standard practice.

Our study confirms the applicability of the AL concept for predicting coronary events and diseases because our study finds comparable predictive performances to traditional C-E/CVD risk scores. For instance, in the study, Erbel et al. [28] had a similar predictive performance on C-E for the Framingham risk score (FAS) using the same data. Unlike traditional C-E/CVD risk scores such as the FAS, AL is not restricted to and specifically designed to predict C-E/CVD risk. However, we evaluate the similarity in the predictive performance as additional validity for AL predicting C-E and CVD.

To address potential reverse-causality issues between high AL and C-E, we excluded those study participants with a C-E related condition at baseline. Still, we did not exclude individuals that have had heart-related drugs already prescribed at baseline. In this sense, the analysis is not entirely free of reverse-causality issues.

A limitation to this finding is that the Z-scoring method did not provide significant interactions. The Z-score ranges from -14.738 to 57.757, and we assume that the insignificance of the interaction is partly caused by the relatively large range. The LRM is based on the Z-scores of the biomarkers and, overall, provided better performances and significant interactions. We view that the causes of both aspects are the lower range of the LRM based AL score (-3.996 to 3.674) and the LRM AL score weights for the relative importance of each biomarker for the C-E incidence.

Another challenge of our study was the relatively large amount of missing values (n = 869) on the AL scores, which we addressed with multiple imputations. The vast majority of the individuals with missing values on the AL score had only one or two missing values on the AL scores (n = 828); thus, excluding those individuals from the analysis would equal the exclusion of study participants of whom 11 out of 13 valid information on the biomarkers were available. We found no significant mean differences between the imputed and unimputed AL scores (see Appendix; Table 2). Furthermore, missing values were only slightly more common in Albumin, Homocysteine and DHEA-S (see Appendix; Table 1). We can neither empirically nor logically find evidence for these missing values to be systematically correlated with the incidence of C-E; thus, we evaluate the assumption of completely missing at random as justified.

The outcome measure C-E also posits a challenge due to the relative rarity (n = 273). This measure was a predefined study criterion restricted to unequivocally documented death related to coronary artery disease or incident nonfatal myocardial infarction [27,28]. To test the restrictiveness of our findings to C-E as an outcome, we provided the same analysis with a looser outcome that measures any incidence of CVD (n = 655). In Appendix Table 3, we applied the same analysis as in Table 3 for the CVD outcome and found similar results in hazards and interaction, but overall lower model fits. With this comparison, we view our results supported.

Concerning the causal character of our study, a limitation is the missing incorporation of time-varying measures. For instance, observing the AL score on multiple points in time would allow more sophisticated modelling to detect time-varying effects on the C-E incidence. Likewise, time-varying confounding could also change the association of AL and C-E. Although we tried to adjust for socio-demographic/economic, health-behavioural and health-related confounders between AL and C-E, we did

not analyze time-varying measures for these confounders since they were unavailable to us. The issue of time-varying measures also highlights the importance of process-oriented analysis, which would allow investigation of mediation and moderation on the AL – C-E association over time and could also determine the influence of gendered coping responses to stress on the sex differences in C-E incidence and AL [15, 16]. Such process-oriented analysis could also help to explain the substantial higher baseline hazards in C-E for males.

4.2. Conclusion

Our study contributes to three research subjects: First, on the discussion of AL scoring algorithms. Second, the applicability of the AL concept on C-E and related conditions. Third, the sex differences in C-E and AL. We conclude that the commonly used methods perform well. The simplicity of the count-based method has the advantage that it can be easily adapted to clinical practices; thus, we recommend using the count-based method for these settings. Our results give strong support for AL to be a valid predictor of C-E. In all models and irrespective of the scoring methods, AL remained a significant predictor of C-E. Concerning the sex differences in the C-E incidence, we showed with multiplicative and additive interaction analysis that at low levels of AL, the sex differences are larger, and at high levels of AL are lower. The additive interaction indicated that, despite the higher baseline risk for males experiencing C-E, females could benefit more from a potential intervention on AL. However, we would like to emphasize that more research on the physiological and psychological origins of the sex gap in the C-E incidence is needed and would suggest the analysis of time-varying AL levels and intermediate factors such as gendered stress coping patterns.

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Statement

The corresponding author has full access to all data in the study and final responsibility for the submission of the article for publication. Due to data security reasons (i.e., data contain potentially participant identifying information), the HNR Study does not allow sharing data as a public use file. Data requests can also be addressed to: recall@uk-essen. de.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix B. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.cpnec.2021.100089.

Appendix Table 1

Appendix

Vlissing Values in Analysis Sample.				
Biomarker	MV	% of TS (N = 4368)		
LDL-C	21	0.48		
HDL-C*	33	0.76		
Triglyceride	22	0.50		
Glucose	23	0.53		
Waist-Hip-Ratio	12	0.27		
C-reactive Protein	31	0.71		
Fibrinogen	212	4.85		
DHEA-S*	76	1.74		
Homocysteine	330	7.55		
Sys. blood Pressure	378	8.65		
Dias. blood Pressure	4	0.09		
Albumin	4	0.09		
Creatinine	212	4.85		
Allostatic load	869	19.89		

Appendix Table 2. Unpaired t-test of AL scores before and after imputation

Unpaired t-test	Ha: diff <0; Pr(T < t)	Ha: diff ! = ; 0 Pr(T > t)	Ha: diff >0; $Pr(T > t)$
Z-scoring	0.631	0.739	0.369
Count-based	0.555	0.889	0.444
LRM	0.637	0.726	0.363

Appendix Table 3. Estimates and fit statistics of AL scoring methods on CVD with and without covariate inclusion

M0 without controls	Z-Scoring	Count-based	LRM
Hazard Ratio [CI]	1.084*** [1.075,1.094]	1.420*** [1.355,1.489]	2.704*** [2.438,2.999]
Somers' D	0.373	0.314	0.408
Harrell's C	0.686	0.657	0.704
AIC	10475.1	10483.4	10340.8
Pseudo-R ²	0.017	0.017	0.030
ROC-Area [CI]	0.673 [0.644, 0.703]	0.671 [0.635, 0.697]	0.703 [0.660, 0.720]
ROC-M0 equality test:	$\text{Chi}^2 = 3.53; \text{ prob} > \text{Chi}^2 = 0.170$		
M1 with controls	Z-Scoring	Count-based	LRM
Hazard Ratio [CI]	1.055*** [1.039,1.071]	1.206*** [1.141,1.276]	1.814*** [1.559,2.112]
Somers' D	0.453	0.459	0.457
Harrell's C	0.726	0.729	0.728
AIC	10261.5	10259.0	10243.7
Pseudo-R ²	0.039	0.039	0.041
ROC-Area [CI]	0.715 [0.685, 0.745]	0.7237 [0.693, 0.753]	0.714 [0.684, 0.744]
ROC-M1 equality test	$\text{Chi}^2 = 6.37; \text{Prob} > \text{Chi}^2 = 0.041$		
M2 AL-sex interaction	Z-Scoring	Count-based	LRM
AL- Hazard Ratio [CI]	1.044*** [1.022,1.067]	1.155*** [1.083,1.232]	1.643*** [1.369,1.971]
Sex = Female (2) - Hazard Ratio [CI]	0.473*** [0.388,0.576]	0.350*** [0.271,0.453]	0.588*** [0.472,0.731]
AL-sex interaction	1.019 [0.992,1.047]	1.181** [1.060,1.317]	1.314* [1.010,1.710]
Hazard Ratio [CI]			
Somers' D	0.453	0.460	0.457
Harrell's C	0.726	0.730	0.728
			(continued on next page)

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(continued)

M0 without controls	Z-Scoring	Count-based	LRM
AIC	10261.7	10252.4	10241.6
Pseudo-R ²	0.039	0.040	0.041
ROC-Area [CI]	0.724 [0.695,0.753]	0.732 [0.703,0.761]	0.730 [0.700,0.759]
ROC-M2 equality test	$Chi^2 = 5.11$; $Prob > Chi^2 = 0.077$		

Note: 95% confidence intervals in brackets * p < 0.05, **p < 0.01, ***p < 0.001.



Appendix Fig. 1. Schoenfeld residual plot of M2 Table 3. Note: Scaled Schoenfeld residuals for the sex-AL interaction. The dashed line expresses the slope. A nonzero slope would equal a violation of the PH assumption and imply time-dependency.

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