



# Dietary patterns and biomarkers of oxidative stress and inflammation: A systematic review of observational and intervention studies

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## ABSTRACT

**Introduction:** Oxidative stress and inflammation are known to play a critical role in ageing and chronic disease development and could therefore represent important targets for developing dietary strategies for disease prevention. We aimed to systematically review the results from observational studies and intervention trials published in the last 5 years on the associations between dietary patterns and biomarkers of oxidative stress and inflammation.

**Methods:** A systematic search of the PubMed, MEDLINE and Web of Science (January 2015 to October 2020) was conducted following the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines. Methodological quality of selected studies was evaluated based on the NUTRIGRADE and BIOCROSS assessment tools.

**Results:** In total, 29 studies among which 16 observational studies and 13 intervention studies were found eligible for review. Overall, results indicated an inverse association between plant-based diets - the Mediterranean and Dietary Approaches to Stop Hypertension (DASH) diet - and oxidative stress and proinflammatory biomarkers. In observational studies, inverse associations were further revealed for the vegetarian diet, the USDA Healthy Eating Index (HEI) - based diet and the paleolithic diet, whereas a positive association was seen for western and fast food diets. Quality assessment suggested that majority of dietary intervention studies (n = 12) were of low to moderate quality.

**Conclusions:** This study provides evidence that the plant-based dietary patterns are associated with lowered levels of oxidative stress and inflammation and may provide valid means for chronic disease prevention. Future large-scale intervention trials using validated biomarkers are warranted to confirm these findings.

## 1. Introduction

Oxidative stress is characterized by an imbalance between production and accumulation of oxygen reactive species (ROS) in cells and tissues and the natural ability of organisms to detoxify abundant reactive species, leading to global oxidative damage and cellular aging [1,2]. ROS represent partially reduced metabolites of molecular oxygen generated as products of metabolic reactions or as by-products of various cellular processes, including inflammation [3–5]. Oxidative stress and inflammation are two closely interrelated and interdependent pathophysiological processes. On one hand, ROS initiate intracellular signaling cascade that enhances proinflammatory gene expression [6];

on the other hand, inflammatory cells secrete ROS and immune mediators (i.e. cytokines and chemokines) leading to induced oxidative stress and tissue damage at the site of inflammation [7]. Oxidative stress and inflammation are known to play an important role in ageing and age-related diseases, including cardiovascular diseases (CVDs), neurodegenerative diseases and cancer [6,8–13]. Understanding the role of nutrition as modifiable determinant of these pathophysiological pathways may therefore hold the key to age-related disease prevention.

Paradoxically, so far, the majority of trials aimed to assess the role of antioxidant nutrients – minerals and vitamins – as preventive targets for chronic diseases, such as CVD and cancer, have not been successful [14, 15]. A potential explanation for these disappointing results could be that those studies were based on measurements of individual pro- and

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## Abbreviations

ABTs	2,2'-azino-bis-3-ethylbenzthiazoline-6-sulphonic acid	MDA	Malondialdehyde
AGEs	Advanced glycation end products	MetS	Metabolic syndrome
AHEI	Alternative Healthy Eating Index	MI	Myocardial infarction
aMED	Alternate Mediterranean diet	MIP-1 $\beta$	Macrophage inflammatory protein-1 beta
AOPPs	Advanced oxidation protein products	MMP-7	Matrix metalloproteinase-7
ARE	Antioxidant response elements	MPO	Myeloperoxidase
CEACAM8	Carcinoembryonic antigen-related cell adhesion molecule 8	MUFA	Monounsaturated fatty acids
CRP	C-reactive protein	NAFLD	Non-alcoholic fatty liver disease
CML	N(6)-carboxymethyllysine	NO	Nitric oxide
COWs	Coke oven workers	Ox-LDL	Oxidated low-density lipoprotein
CRP	C-reactive protein	PCA	Principal component analysis
CVD	Cardiovascular disease	PON-3	Paraoxonase 3
DASH	Dietary Approaches to Stop Hypertension	PRISMA	Preferred Reporting of Systematic Reviews and Meta-Analyses
DPPH	1,1-diphenyl-2-picrylhydrazyl	PUFA	Polyunsaturated fatty acids
FFQ	Food frequency questionnaire	Q	Quartile
FLOP	Fluorescent oxidation product	RCT	Randomized-controlled trial
FRAP	Ferric reducing ability of plasma	ROS	Reactive oxygen species
Gal-4	Galectin-4	SFA	Saturated fatty acids
GSH	Glutathione	sNox2-dp	Soluble Nox2-derived peptide
HEI	Healthy Eating Index	SOD	Superoxide dismutase
HT	Hashimoto thyroiditis	TAC	Total antioxidant capacity
IL	Interleukin	TBARS	Thiobarbituric acid reactive substances
IL1RT1	Interleukin-1 receptor type 1	TNF- $\alpha$	Tumor necrosis factor alpha
IL2RA	Interleukin-2 receptor subunit alpha	TRAIL-R2	Tumor necrosis factor-related apoptosis inducing ligand receptor 2
MCP-1	Monocyte chemoattractant protein-1	UPAR	Urokinase plasminogen activator surface receptor
		8-OH-dG	8-hydroxy-2-deoxyguanosine

anti-oxidant markers and thus may not provide a comprehensive assessment of both the nutritional exposure and the systemic redox status [14]. While previous research has been mostly focused on individual nutrients or single foods, the last decade has been marked by an increased recognition of the importance of the whole diet as opposed to specific nutrients [16]. The dietary pattern approach recognizes that foods are composed by various nutrients and bioactive constituents and are consumed in combinations and may interact with each other in complex ways [17]. Due to this complexity, the associations between single nutrients and age-related diseases and phenotypes may be difficult to explore and interpret. Dietary patterns allow examining combinations of various food components and their complex interactions [18]. Dietary patterns can be defined by *posteriori* (data-driven or empirical approaches) and *a priori* guidelines and recommendations (diet quality indices). The *posteriori* approaches include statistical modeling techniques aimed to disentangle intercorrelated structures dietary items (principal component analysis) or to group individuals into patterns based on their reported mean intakes of foods (cluster analysis). On the other hand, the *a priori* diet quality indices are useful to assess adherence to dietary guidelines, i.e. Healthy Eating Index [19] or to a particular types of diet such as the Dietary Approaches to Stop Hypertension (DASH) [20] or Mediterranean diet [21]. A number of studies suggested that dietary patterns are associated with cardiometabolic risk factors, inflammatory levels and all-cause mortality in human research [21–23].

The field of biomarker discovery has been booming in recent years leading to new advances in the identification of novel molecules characterizing oxidative stress and inflammation in human research [24]. Likewise, in nutrition research the dietary pattern assessment has been in focus over the last years [23]. So far, the link between dietary patterns in relation to oxidative stress was only partly addressed in a previous systematic review [25]. The newly emerging evidence on link between dietary patterns and various biomarkers representing these pathways has not been evaluated.

We therefore aimed to systematically review and synthesize the

results from observational studies and dietary intervention trials published in the last 5 years to clarify the association between dietary patterns and biomarkers characterizing oxidative stress and inflammation.

## 2. Methods

### 2.1. Protocol registration

The systematic review followed the requirements of the Preferred Reporting of Systematic Reviews and Meta Analyses (PRISMA) statement [26] and was registered at the International Prospective Register of Systematic Reviews (PROSPERO, Reference number: 212315; <http://www.crd.york.ac.uk/prospero/212315>).

### 2.2. Search strategy

A systematic search of the PubMed, MEDLINE and Web of Science (January 2015 to October 2020) was conducted using the following combination of Medical Subject Heading (MeSH) terms and text words, with limitation to English language ("Eat-Lancet diet" [All Fields] OR "planetary health diet" [All Fields] OR "portfolio diet" [All Fields] OR "DASH" [All Fields] OR "Dietary Approaches to Stop Hypertension" [All Fields] OR "Dietary Inflammatory Index" [All Fields] OR "nordic diet" [All Fields] OR "paleolithic diet" [All Fields] OR "plant-based diet" [All Fields] OR "vegetarian diet" [All Fields] OR "vegan diet" [All Fields] OR "Mediterranean diet" [All Fields] OR "dietary pattern\*" [All Fields] OR "eating pattern\*" [All Fields] OR "food pattern\*" [All Fields] OR "diet index" [All Fields] OR "dietary index" [All Fields] OR "diet score" [All Fields] OR "dietary score" [All Fields]) AND ("oxidative damage" [All Fields] OR "oxidative stress" [All Fields] OR "immun\*" [All Fields] OR "inflammat\*" [All Fields] OR "CRP" [All Fields] OR "C-reactive protein" [All Fields] OR "IL" [All Fields] OR "interleukin\*" [All Fields] OR "TNF" [All Fields] OR "tumor necrosis factor" [All Fields] OR "acute-phase protein\*" [All Fields] OR "adipokin\*" [All Fields] OR "cytokine\*" [All

Fields]). The PICOs (Population, Intervention, Comparator, Outcome, Study Design) criteria used to define our research question are listed in [Supplementary Table 1](#).

### 2.3. Eligibility criteria

Studies were included if they reported on the association (observational studies) or effect (intervention studies) of dietary patterns (as exposure) with biomarkers of oxidative stress and inflammation (as outcome). A primary focus was put on studies reporting results on biomarkers representing pathways denoting oxidative stress and wherever available simultaneous measurements of inflammatory biomarkers.

The inclusion criteria were as follows: a) assessment of dietary patterns (based on whole foods) as main exposure; b) plasma/serum/urine measurements of biomarkers as main outcome measures; c) enrolled humans at adult and old age; d) analytical epidemiological studies, i.e. observational studies (cross-sectional, case-control or prospective cohort studies) and intervention studies (non-randomized trials, i.e. pre-post studies, and randomized control trials); and e) studies written in English and published in peer-reviewed journals.

The exclusion criteria were: (a) no original research (e.g. reviews, editorials, non-research letters); (b) case reports or case series; (c) ecological studies; (d) lack of data on dietary patterns (e.g., examined only individual nutrients or did not examine all dietary components); (e) no biomarker measurement reported; (f) studies not conducted in humans; (g) studies not conducted in adult population (<18 years old); and (h) studies without reported effect estimates. Additionally, intervention studies were excluded if: a) they used lifestyle interventions in conjunction with diet intervention (e.g., exercise or behavioral management); b) postprandial studies; or c) studies with intervention duration of less than 4 weeks. The latter criteria were applied to allow characterizing potential effect of habitual diet on changes of biomarker concentrations at a longer run rather than acute effects of initial drastic dietary change.

### 2.4. Selection of studies

Identified records were imported in EndNote referencing software (version X7, 2013; Thomson Reuters) and their titles and abstracts were screened by two independent reviewers (LK and TH). Full-text articles were retrieved if the article was considered eligible, and subjected to a second evaluation by two other independent reviewers (CER and KA). Any discrepancies and disagreements were discussed and resolved by consensus among reviewers. After retrieval of full-text articles, the reference lists of selected articles and other reviews were checked to identify additional potentially relevant articles. Where results from the same study were reported in multiple articles, the most recent article was included to avoid duplication of results.

### 2.5. Data extraction

Data extraction was performed by two independent reviewers (LK and TH) using a predefined data extraction form and extracted data was verified by a third reviewer (CER). The following information was extracted: first author, publication year and country, study design, sample size, dietary patterns identified (incl. dietary assessment tool and method of identifying dietary patterns), details of intervention and control groups (intervention studies), biomarker measured, and main findings. In addition, data on analytical methods, sample type and participants' fasting status was recorded for each of the selected studies. When a study provided several estimates with adjustment for different confounders, results were reported based on the one adjusting for the largest number of factors (see [Supplementary Table 2](#) for adjustment models). Discrepancies in data extraction were discussed and resolved by consensus among the reviewers.

### 2.6. Assessment of quality and risk of bias of included studies

The assessment of the quality of studies was performed by three independent reviewers (CER, LK and TH) using a combination of tools specifically developed for the assessment of nutritional exposures (NUTRIGRADE) [27] and biomarker outcomes (BIOCROSS) [28]. These tools were additionally adapted to consider features of cross-sectional and intervention study designs of the included studies. The specific questions and assigned points of the combined tool for observational and intervention studies are presented in [Supplementary Tables 3 and 4](#), respectively. Each study was graded for a) overall risk of bias pertaining to how the study was conducted and reported, incl. study quality, study limitations, statistical analysis, data interpretation and funding bias (based on NUTRIGRADE) and b) risk of bias related to biomarker assessment and reporting, incl. biomarker measurement, specimen characteristics and assay methods, and laboratory measurements (based on BIOCROSS). The total scores of the study quality assessment tool for cross-sectional studies and intervention studies were 14.5 and 13, respectively. Studies scoring 0–5 were considered low quality, studies scoring 5–10 were considered moderate quality and studies scoring  $\geq 10$  points were considered high quality.

## 3. Results

### 3.1. Search results

The process of study selection, including identification, screening, eligibility, and inclusion is illustrated in a flowchart presented in [Fig. 1](#). The search strategy retrieved 2534 unique records. Initial screening of title and abstract excluded 2304 citations. 230 records were then selected for full-text detailed evaluation. Following the screening of the full-texts, further 203 articles were excluded because: a) studies did not report on measurements of oxidative stress biomarkers ( $n = 165$ ); b) did not include information on dietary patterns ( $n = 20$ ); c) the study included inappropriate study population ( $n = 5$ ); d) had inappropriate intervention duration (<2 weeks) ( $n = 4$ ); and e) other reasons ( $n = 7$ ). Among these, 27 articles were identified as meeting the eligibility criteria. In addition to that, 2 studies were found through manual search of reference lists of selected studies and review articles. Finally, a total of 29 articles have been included in this current systematic review among which 16 observational studies and 13 intervention studies.

### 3.2. Study characteristics

The study characteristics of the selected observational and intervention studies are presented in [Tables 1 and 2](#), respectively. Overall, there were 16 cross-sectional studies [29–44], 5 non-randomized intervention trials [45–49] and 8 RCTs [50–57]. None of the studies applied a prospective observational design to investigate associations between food patterns and changes in biomarker concentrations. Of the 8 RCTs, 6 used a parallel design and 2 used a crossover design [50–57]. The 29 studies were conducted in the following countries: Australia [53, 54], Brazil [29], Chile [47], China [43], Cyprus [38], Greece [42,45], Germany [49], Italy [30,31,37,44,46,51], Korea [57], Spain [39,50,56], Sweden [35], United States of America [34,40,41,48,52] and Iran [32, 33,36,55]. The study sample sizes ranged from 40 to 2240 participants for observational studies [29–44], and 20 to 805 participants for intervention studies [45–57]. The duration of intervention studies ranged from 2 weeks to 5 years with a mean duration of 10 weeks [45–57]. Participants' ages ranged from 16 to 80 years at the time of study in the observational studies [29–44] and 20–80 years in the intervention studies [45–57]. 5 studies were restricted to women [33,38,41,47,57] and 4 studies included only men [29,37,43,49]. The remaining 20 studies examined both sexes. Most observational studies recruited healthy community-dwelling adults and several studies included also participants with underlying health condition, i.e. metabolic syndrome

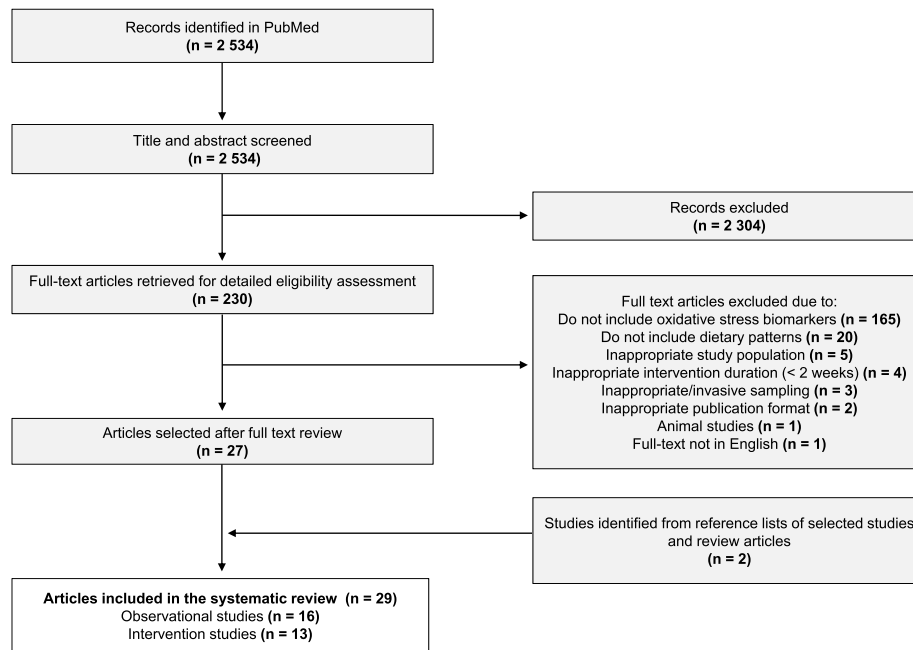


Fig. 1. Flowchart of study selection, including identification, screening, eligibility, and inclusion of studies.

(MetS) [36], non-alcoholic fatty liver disease (NAFLD) [30], euthyroid Hashimoto thyroiditis (HT) [31] obesity [34] or atrial fibrillation patients [44]. Conversely, majority of intervention studies were conducted in participants with existing health problem such as obesity and MetS [46,47,52,55–57], CVD [50], NAFLD [45,55], whereas five of the 13 studies were conducted in predominantly healthy individuals [48,49,51, 53,54].

### 3.3. Dietary patterns

In observational studies, dietary assessment was performed predominantly using validated food frequency questionnaires (FFQs) ( $n = 11$  [31–33,35–38,40–43]), whereas the remaining studies used dietary records ( $n = 4$  [29,30,39,44]) and repeated 24-h dietary recalls ( $n = 1$  [34]). Majority of observational reports were based on *a priori* approaches (diet quality scores or indexes) to define dietary patterns ( $n = 9$  [30,31,34,38–42,44]). These included the Mediterranean diet score [30, 31,38–42,44], USDA Healthy Eating Index (HEI) [34,41] and various other dietary patterns such as paleolithic diet [40] and DASH diet [41]. Overall, 5 studies used *a posteriori* approaches, which included factor/principal component analysis (PCA) to identify data-driven dietary patterns [32,33,35,36,43]. Two studies included participants who were vegetarian, vegan or omnivorous by choice so they did not include a score [29,37]. The dietary components of the identified dietary patterns are summarized in Supplementary Table 5. In intervention studies, Mediterranean diet was again the most commonly used main intervention approach (9 studies [45–47,49–52,54,56]), among which 5 RCTs [50–52,54,56]). Further dietary interventions included the DASH diet [55,57], the lacto-ovo vegetarian low-calorie diet [51] and the traditional Daniel Fast diet [48]. 2 studies explored variations of Mediterranean diet whereby specific food components were additionally included [56] or excluded [47]. While most studies focused on evaluation of dietary patterns generally considered healthy, one study explored effect of a western diet (high in red and processed meat and refined grains) so far described to exert unfavorable health effects [53]. In the majority of the RCTs, dietary recommendations were provided for both the intervention and control groups. Most commonly, control groups followed a habitual diet [52,54], habitual diet with dietary counselling [57], or a low-fat/low-calorie diet [50,55,56]. The most common

methods for evaluating dietary intervention compliance were dietary questionnaires and diet records.

### 3.4. Biomarkers evaluated

A summary of evaluated biomarkers for oxidative stress and inflammation used in the selected studies according to their main characteristics and pathophysiological actions is provided in Table 3. A short description of methods for measurement and sample type used in the observational and intervention studies has been summarized in the Supplementary Tables 6 and 7, respectively. Overall, various biomarkers were used as study endpoints to reflect different pathways involved in oxidative stress including both pro-oxidative and antioxidant defense mechanisms [29–57]. In addition to oxidative stress biomarkers, 14 studies addressed biomarkers of specific immune-inflammatory activation, with C-reactive protein (CRP), interleukin-6 (IL-6), and tumor necrosis factor alpha (TNF- $\alpha$ ) being most commonly assessed [29,33–35, 40,42,45,46,48,50,53–55].

### 3.5. Associations between dietary patterns and biomarkers of interest

An overview of studies reporting results on significant differences between biomarker levels in participants according to identified dietary patterns in observational and intervention studies is presented in Table 4. Overall, reduced concentrations of oxidative stress and pro-inflammatory biomarkers and increased concentrations of antioxidant and anti-inflammatory biomarkers were reported for the Mediterranean diet, the vegetarian diet, the DASH diet, the USDA HEI diet and the paleolithic diet. Vice versa, studies that explored Western and fast-food diets (based on data from observational studies) reported higher levels of oxidative stress and inflammation biomarkers. Of note, only for the Mediterranean and DASH diet significant results could be seen in both observational and intervention studies. The associations described for the other dietary patterns were based just on data from observational studies [29–57].

In observational studies, adherence to the Mediterranean dietary pattern compared to reference diets (i.e., habitual or Western diet) was associated with significant differences in levels of oxidative stress biomarkers in 6 out of 9 studies with no differences reported in 3 out of 9

**Table 1**

Summary of observational studies investigating the associations between dietary patterns and biomarkers of oxidative stress and inflammation.

Authors, Year	Country	Study design, duration	Participants	Dietary pattern/ dietary assessment method	Biomarkers	Results <sup>a</sup> Positive association (↑); Inverse association (↓); No association (↔)
Cinegaglia et al. [29], 2020	Brazil	Cross-sectional	2 groups: 1) N = 44, mean age (sem): 46.8 (1.4) years, healthy omnivorous adults 2) N = 44, mean age (sem): 45.5 (1.2) years; healthy vegetarian adults 100% male	1) Omnivorous diet 2) Vegetarian diet (reference)	Plasma: hs-CRP; heme oxygenase-1 levels	Omnivorous diet vs vegetarian diet: ↑ heme oxygenase-1 levels; ↔ hs-CRP
Baratta et al. [30], 2020	Italy	Cross-sectional/ PLINIO Study	N = 238, mean age (sd): 53.1 (12.4) years, 57% male, patients with NAFLD	Mediterranean diet /validated dietary questionnaire	Serum: sNox2-dp	↓ sNox2-dp
Ruggeri et al. [31], 2020	Italy	Cross-sectional	2 groups (pooled together): 1) N = 81, age range: 18–66 years, 12% male, patients with euthyroid HT 2) N = 119, age range: 18–65 years, 14% male, without euthyroid HT	Mediterranean diet /FFQ	Serum: AGEs; AOPPs Plasma: SOD; Glutathione reductase activity; Glutathione peroxidase activity; Thioredoxin reductase activity; TAC	↔ AGEs; ↔ AOPPs; ↔ SOD; ↔ Glutathione reductase activity; ↔ Glutathione peroxidase activity; ↔ Thioredoxin reductase activity; ↔ TAC
Seyedi et al. [32], 2020	Iran	Cross-sectional/ Tehran Lipid and Glucose Study	N = 470, age range: 40–70 years, 25% male	1) Western pattern 2) Semi-Mediterranean pattern Healthy pattern (reference) /FFQ	Lipoprotein-associated phospholipase A2	1) Western pattern vs healthy pattern: ↑ Lipoprotein-associated phospholipase A2 (univariate and multivariate models) 2) Semi-Mediterranean pattern vs healthy pattern: ↓ Lipoprotein-associated phospholipase A2 (univariate model); ↔ Lipoprotein-associated phospholipase A2 (multivariate model)
Abashzadeh et al. [33], 2020	Iran	Cross-sectional	N = 320, age range: 20–45 years, 0% male, healthy female nurses	1) Healthy pattern 2) Traditional pattern 3) Unhealthy pattern /FFQ	Serum: Ceruloplasmin; protein carbonyl; TAC	1) Healthy pattern: ↔ Ceruloplasmin; ↔ protein carbonyl; ↔ TAC 2) Traditional pattern: ↔ Ceruloplasmin; ↔ protein carbonyl; ↔ TAC 3) Unhealthy pattern: ↓ Ceruloplasmin; ↓ protein carbonyl; ↔ TAC
Crowe-White et al. [34], 2019	USA	Cross-sectional	N = 133, mean age (sd): 70.4 (4.8) years, 40% male, obese older adults	HEI /3 × 24h dietary recalls	Serum: Hydrophilic antioxidant capacity; lipophilic antioxidant capacity; TAC; hs-CRP; TNF-α; IL-6	↔ Hydrophilic antioxidant capacity; ↔ lipophilic antioxidant capacity; ↔ TAC; ↔ hs-CRP; ↔ TNF-α; ↔ IL-6
Lemming et al. [35], 2019	Sweden	Cross-sectional/ EpiHealth	N = 2240, mean age (sd): 61 (8.4) years, 50% male	1) Healthy pattern 2) Western pattern 3) Dairy and sandwich pattern 4) Fast food and alcohol pattern /FFQ	Plasma: MPO protein level; resistin; spondin-2; follistatin; MMP-7; PON-3; Gal-4; TRAIL-R2; IL1RT1; UPAR; CEACAM8; IL2RA	1) Healthy pattern: ↑ PON-3; ↓ MPO protein level; ↓ spondin-2; ↓ follistatin; ↓ MMP-7; ↓ TRAIL-R2; ↔ resistin; ↔ Gal 4; ↔ IL1RT1; ↔ UPAR; ↔ CEACAM8; ↔ IL2RA 2) Western pattern: ↑ MPO protein level; ↑ resistin; ↑ follistatin; ↑ CEACAM8; ↑ IL1RT1; ↑ TRAIL-R2; ↑ IL2RA; ↑ UPAR; ↓ Gal-4; ↔ spondin-2; ↔ PON-3; ↔ MMP-7; ↔ resistin 3) Dairy and sandwich pattern: ↑ CEACAM8; ↔ spondin-2; ↔ follistatin; ↔ MMP-7; ↔ Gal-4; ↔ TRAIL-R2; ↔ IL1RT1; ↔ UPAR; ↔ IL2RA; ↔ PON-3; ↔ MPO protein level; ↔ resistin 4) Fast food and alcohol pattern:

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Table 1 (continued)

Authors, Year	Country	Study design, duration	Participants	Dietary pattern/ dietary assessment method	Biomarkers	Results <sup>a</sup> Positive association (↑); Inverse association (↓); No association (↔)
Mirmiran et al. [36], 2018	Iran	Cross-sectional/ Tehran Lipid and Glucose Study	N = 400, age range: 20–60 years, adults with MetS	1) Healthy pattern (high in fruits and vegetables) 2) Unhealthy pattern (high in soft drinks, fast foods, organ meats) /FFQ	Plasma: TAC; MDA	↑ PON-3; ↓ MPO protein level; ↓ spondin-2; ↓ Gal-4; ↓ IL1RT1; ↓ CEACAM8; ↔ resistin; ↔ follistatin; ↔ TRAIL-R2; ↔ UPAR; ↔ IL2RA; ↔ MMP-7 1) Healthy pattern: ↑ TAC; ↓ MDA 2) Unhealthy pattern: ↓ TAC; ↑ MDA
Vanacore et al. [37], 2018	Italy	Cross-sectional	N = 30 (N = 10 omnivore diet; N = 10 vegan diet; N = 10 vegetarian diet), age range: 20–30 years, 100% male, metabolically healthy adults	1) Omnivore diet 2) Vegetarian diet 3) Vegan diet /FFQ	Serum: DPPH; FRAP; total phenol; ABTS; TBARS; nitrite	1) Omnivore diet vs vegan: ↑ Total phenol; ↔ FRAP; ↔ DPPH; ↔ ABTS; ↔ TBARS; ↔ nitrite 2) Vegetarian diet vs vegan: ↑ Total phenol; ↑ FRAP; ↔ DPPH; ↔ ABTS; ↔ TBARS; ↔ nitrite (↑ nitrite compared to omnivore diet) 3) Vegan diet vs the other diets: ↑ TBARS; ↑ nitrite compared to omnivore diet, ↔ nitrite compared to vegan diet; ↔ DPPH; ↔ ABTS; ↔ total phenol; ↔ FRAP
Kakkoura et al. [38], 2017	Cyprus	Cross-sectional	N = 564, mean age (sd): 55.3 (7.3), 0% male, women without breast cancer	Mediterranean diet /FFQ	Serum: GSH; Flavin mononucleotide; methionine sulfoxide; homocysteine; cystathionine; total cysteine	↔ GSH; ↔ flavin mononucleotide; ↔ methionine sulfoxide; ↔ homocysteine; ↔ cystathionine; ↔ total cysteine
Aranada et al. [39], 2017	Spain	Cross-sectional	N = 81, mean age (95% CI): 43.6 (40.1–47.1) years, 43% male, healthy adults	Mediterranean diet with various types of seafood /3-day dietary record	Plasma: Ox-LDL; F <sub>2</sub> -isoprostane	1) Mediterranean diet + white fish (adjusted): ↓ F <sub>2</sub> -isoprostane; ↔ Ox-LDL 2) Mediterranean diet + oily fish (adjusted): ↔ Ox-LDL; ↔ F <sub>2</sub> -isoprostane 3) Mediterranean diet + shellfish (adjusted): ↑ Ox-LDL; ↔ F <sub>2</sub> -isoprostane
Whalen et al. [40], 2016	USA	Cross-sectional/ pooled MAPI and MAPII studies	N = 646 (N = 558 with hs-CRP samples; N = 434 with F <sub>2</sub> - isoprostane samples), age range: 30–74 years	1) Paleolithic diet 2) Mediterranean diet /FFQ	Plasma: F <sub>2</sub> -isoprostane; hs-CRP	1) Paleolithic diet ↓ F <sub>2</sub> -isoprostane; ↓ hs-CRP 2) Mediterranean diet: ↓ F <sub>2</sub> -isoprostane; ↓ hs-CRP
Jung et al. [41], 2016	USA	Cross-sectional/ Nurses' Health Study	N = 1688, age range: 30–55 years, 0% male, women free of cancer and MI	1) AHEI 2) DASH 3) aMED /FFQ	Plasma: Fluorescent oxidation products (FLOP_320; FLOP_360; FLOP_400)	1) AHEI: ↑ FLOP_320; ↑ FLOP_360; ↔ FLOP_400 2) DASH ↑ FLOP_320; ↑ FLOP_360; ↔ FLOP_400 3) aMED: ↑ FLOP_320; ↑ FLOP_360; ↔ FLOP_400
Koloverou et al. [42], 2016	Greece	Cross-sectional/ ATTICA study	N = 191	Mediterranean diet /FFQ	Serum: TAC; ox-LDL; CRP; IL-6; TNF-α; serum amyloid A; homocysteine	↑ TAC; ↓ ox-LDL; ↓ CRP; ↓ IL-6; ↓ TNF-α; ↓ serum amyloid A; ↓ homocysteine
Xie et al. [43], 2015	China	Cross-sectional	N = 51 topside COWs, mean age (sd): 35.5 (8.3) years, 100% male, adults exposed to high levels of toxic chemicals N = 79 other COWs, mean age (sd): 37.8 (8.2) years, 100% male, adults exposed to high levels of toxic chemicals	1) Rice-noodle pattern 2) Fruit-vegetable pattern 3) High-protein food pattern 4) Snack-sugar pattern /FFQ	Serum: MDA; SOD protein level; glutathione peroxidase protein level	In all COWs (Q4 vs Q1 of pattern): 1) Rice-noodle pattern: ↔ MDA; ↔ SOD protein level; ↔ glutathione peroxidase protein level 2) Fruit-vegetable pattern: ↑ SOD protein level; ↑ glutathione peroxidase protein level; ↓ MDA

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Table 1 (continued)

Authors, Year	Country	Study design, duration	Participants	Dietary pattern/ dietary assessment method	Biomarkers	Results <sup>a</sup> Positive association (↑); Inverse association (↓); No association (↔)
Pastori et al. [44], 2015	Italy	Cross-sectional	N = 709, median age (IQR): 72.6 (8.7) years, 56% male, atrial fibrillation patients	Mediterranean diet /validated short dietary questionnaire	Serum: sNOX2-dp	3) High-protein pattern: ↔ MDA; ↔ SOD protein level; ↔ glutathione peroxidase protein level 4) Snack-sugar pattern: ↔ MDA; ↔ SOD protein level; ↔ glutathione peroxidase protein level ↓ sNOX2-dp

Results ordered according to type of biomarker (oxidative stress, immune-inflammatory) and according to significance.

Abbreviations: ABTs, 2,2'-azino-bis-3-ethylbenzthiazoline-6-sulphonic acid; AGEs, advanced glycation end products; AHEI, Alternative Healthy Eating Index; aMED, Alternate Mediterranean diet; AOPPs, advanced oxidation protein products; ARE, antioxidant response elements; CEACAM8, carcinoembryonic antigen-related cell adhesion molecule 8; COWs, coke oven workers; CRP, C-reactive protein; DASH, Dietary Approach to Stop Hypertension; DPPH, 1,1-diphenyl-2-picrylhydrazyl; FFQ, food frequency questionnaire; FLOP, fluorescent oxidation product; FRAP, ferric reducing ability of plasma; Gal-4, Galectin-4; HEI, Healthy Eating Index; hs, high sensitivity; HT, hashimoto thyroiditis; IQR, interquartile range; IL, interleukin; ILTR1, interleukin-1 receptor type 1; IL2RA, interleukin-2 receptor subunit alpha; MDA, malondialdehyde; MetS, metabolic syndrome; MI, myocardial infarction; MMP-7, matrix metalloproteinase-7; MPO, myeloperoxidase; NAFLD, non-alcohol fatty liver disease; NO, nitric oxide; ox-LDL, oxidized low density lipoprotein; PON-3, paraoxonase 3; Q, quartile; ROS, reactive oxygen species; sd, standard deviation; sem, standard error of mean; sNOX2-dp, soluble Nox2-derived peptide; SOD, superoxide dismutase; TAC, total antioxidant capacity; TBARS, thiobarbituric acid reactive substances; tGSH, total glutathione; TNF- $\alpha$ , tumor necrosis factor alpha; TRAIL-R2, tumor necrosis factor-related apoptosis inducing ligand receptor 2; UPAR, urokinase plasminogen activator surface receptor.

<sup>a</sup>  $P < 0.05$ .

studies [30–32,38–42,44]. Levels were lower for the soluble Nox2-derived peptide (sNOx2-dp) [30,44] and F<sub>2</sub>-Isoprostane [40] and higher for fluorescent oxidation product 320 (FLOP\_320), FLOP\_360 [41] and total anti-oxidant capacity (TAC) [42]. Contradicting results for oxidized low-density lipoprotein (ox-LDL) were reported in two observational studies, with one study showing elevated levels following Mediterranean diet enriched with higher intakes of fish and shellfish [39], whereas the second study reported lower ox-LDL levels related to consumption of traditional Mediterranean diet [42].

In intervention studies, significantly different biomarker levels in Mediterranean diet groups compared to control groups/baseline levels were reported in 7 out of 8 studies [45–47,49–52,54]. Following Mediterranean diet intervention led to significantly reduced levels of a wide range of biomarkers reflecting different aspects of oxidative stress, including biomarkers of lipid peroxidation (F<sub>2</sub>-isoprostanes, ox-LDL, malondialdehyde (MDA), thiobarbituric acid reactive substances (TBARS)), oxidative DNA damage (8-hydroxydeoxyguanosine (8-OH-dG)) [45,46,51,54], reactive metabolic products and byproducts (methylglyoxal and N (6)-carboxymethyllysine (CML)) [47,50], and biomarkers representing endogenous immune-inflammatory activation as sources of oxidative stress (myeloperoxidase (MPO) activity) [46,52]. Conversely, higher levels were observed for of biomarkers of antioxidant defense and ROS detoxification such as superoxide dismutase (SOD) activity and protein level, catalase activity, xanthine oxidase activity [56]. Moreover, in addition to oxidative stress biomarkers, adherence to Mediterranean diet was related to lower concentrations of pro-inflammatory biomarkers - CRP, IL-6, TNF- $\alpha$ , serum amyloid A and homocysteine in observational studies [40,42]. Consistently, in intervention studies, the results for CRP, IL-6 and TNF- $\alpha$  were further confirmed and additionally increased levels of anti-inflammatory biomarkers - IL-10 and adiponectin were demonstrated [45,46,50].

Vegetarian dietary patterns were evaluated in six observational studies and in one intervention study. Compared to reference diets consumption of vegetarian diet was associated with significant differences in levels of oxidative stress biomarkers in 4 out of 6 observational studies. Levels were lower for oxidative stress biomarkers reflecting lipid peroxidation and immune-inflammatory activation as sources of oxidative stress MDA and MPO [35,36,43], whereas levels for

biomarkers of antioxidant defense and ROS detoxification (ferric reducing ability of plasma (FRAP), TAC, nitrite, SOD protein level, glutathione (GSH) peroxidase protein level) were higher [36,37,43]. In the intervention study, reduced levels of L-derived ROS and TBARS were additionally reported [51]. One observational study explored adherence to vegan diet in relation to oxidative stress biomarkers and showed that the levels of TBARS and nitrite were elevated compared to an omnivorous diet [37]. The DASH diet was evaluated in one observational and two intervention studies. Unexpectedly, in the observational study, DASH diet was positively, albeit weakly, associated with FLOP\_320 and FLOP\_360 [41], whereas lower levels of MDA, TBARS and CRP and higher levels of nitric oxide (NO) and GSH were seen in the intervention studies [55,57]. Adherence to the USDA HEI diet and the paleolithic diet were shown to be associated (also weakly) with higher levels of FLOP\_320 and FLOP\_360 [41] and lower levels of F<sub>2</sub>-isoprostane and CRP [40], respectively.

Finally, results from observational studies have reported that consumption of Western/fast-food diets was associated with elevated levels of a range of biomarkers reflecting different aspects of oxidative stress (e.g., MDA, heme oxygenase-1 levels, paraoxonase 3 (PON-3)) and inflammation (e.g., resistin, tumor necrosis factor-related apoptosis inducing ligand receptor 2 (TRAIL-R2), urokinase plasminogen activator surface receptor (UPAR), interleukin-2 receptor subunit alpha (IL2RA)) and with decreased levels of biomarkers reflecting antioxidant defense (e.g., TAC, MPO) [29,32,33,35,37]. The direction of association in Western and fast-food diets differed for MPO and interleukin-1 receptor type 1 (IL1RT1), showing to be elevated in a Western diet pattern while decreased in a fast-food pattern [35]. In the same study, galectin-4 (Gal-4) was inversely associated with both Western and fast-food patterns [35].

### 3.6. Risk of bias and study quality assessment

The study quality and risk of bias assessments for each observational and intervention study included in the systematic review are presented in Supplementary Tables 8 and 9, respectively. The total assessment points ranged from 4.5. to 12 for observational studies (median = 9.75) and from 4 to 11.5 for intervention studies (median = 7). Among the 16

Table 2

Summary of intervention studies investigating the associations between dietary patterns and biomarkers of oxidative stress and inflammation.

Authors, Year	Country	Study design, duration	Participants	Dietary pattern	Biomarkers	Results <sup>a</sup> Increase in biomarker concentration (↑); Decrease in biomarker concentration (↓); No change (↔)
<b>Randomized controlled trials</b>						
Yubero-Serrano et al. [50], 2020	Spain	Parallel RCT, 1 year	Intervention: N = 418, mean (sd) age: 60.4 (0.5) years, 91% male, coronary heart disease patients Control: N = 387, mean (sd) age: 59.9 (0.5) years, 94% male, coronary heart disease patients	Intervention: Mediterranean diet Control: Low-fat, high-complex carbohydrate diet	Serum: Methylglyoxal; hs-CRP	↓ Methylglyoxal; ↓ hs-CRP
Sofi et al. [51], 2018	Italy	Crossover RCT, 3 months each	Intervention (2 arms): 1) N = 60, age range: 24–70 years, 18% male, healthy adults 2) N = 58, age range: 21–75 years, 26% male, healthy adults	Intervention (2-arms): 1) Lacto-ovo vegetarian low-calorie diet 2) Mediterranean low-calorie diet	Plasma: TBARS; TAC; L-derived ROS; M-derived ROS; G-derived ROS	1) Vegetarian diet: ↓ TBARS; ↓ L-derived ROS; ↔ M-derived ROS; ↔ G-derived ROS; ↔ TAC 2) Mediterranean diet: ↓ TBARS; ↔ L-derived ROS; ↔ M-derived ROS; ↔ G-derived ROS; ↔ TAC
Jaacks et al. [52], 2018	United States	Parallel RCT, 8 weeks	Intervention: N = 11, overweight or obese adults Control: N = 9, overweight or obese adults Mean (sd) age: 51.4 (6.6) years, 27% male	Intervention: Mediterranean diet Control: Habitual US diet	Plasma: Cysteine (reduced form); cystine (oxidized form); GSH	↓ Cystine; ↔ cysteine; ↔ GSH
Kim et al. [53], 2017	Australia	Crossover RCT, 4 weeks each	All participants: N = 51, mean (sd) age: 35.1 (15.6) years, 42% male, people without diabetes	Intervention (2 arms): 1) Diet high in red and processed meat and refined grains 2) Diet high in whole grains, nuts, legumes, dairy, and devoid of red and processed meat	Plasma: Fluorescent AGEs; CML; IL-6; hs-CRP	Post-intervention levels of diet 1 vs diet 2: ↔ Fluorescent AGEs; ↔ CML; ↔ IL-6; ↔ hs-CRP
Davis et al. [54], 2017	Australia	Parallel RCT, 24 weeks	Intervention: N = 80, mean (sd) age: 71.0 (4.9) years, 42% male, healthy adults Control: N = 72, mean (sd) age: 70.8 (4.7) years, 46% male, healthy adults	Intervention: Mediterranean diet Control: Habitual Australian diet	Plasma: F <sub>2</sub> -Isoprostanes; hs-CRP	↓ F <sub>2</sub> -Isoprostanes; ↔ hs-CRP
Zade et al. [55], 2016	Iran	Parallel RCT, 8 weeks	Intervention: N = 30, mean (sd) age: 39.7 (7.3) years, 50% male Control: N = 30, mean (sd) age: 42.8 (10.6) years, 50% male Overweight and obese patients with NAFLD	Intervention: DASH diet Control: Calorie restricted diet	Plasma: MDA; NO; GSH; TAC Serum: hs-CRP	↑ NO; ↑ GSH; ↓ MDA; ↔ TAC; ↓ hs-CRP
Sureda et al. [56], 2016	Spain	Parallel RCT, 5 years	Intervention (2 arms): 1) N = 25, 46% male 2) N = 25, 45% male Control: N = 25, 48% male Age range: 55–80 years, people with metabolic syndrome	Intervention (2 arms): 1) Mediterranean diet + olive oil 2) Mediterranean diet + nuts Control: Low-fat diet	Total blood: SOD activity Plasma: Catalase activity; MPO activity; MPO protein level; xanthine oxidase activity; xanthine oxidase protein level; SOD protein level; nitrate; nitrite; nitrotyrosine index; carbonylated proteins	1) Mediterranean diet + olive oil vs control: ↑ SOD activity; ↑ SOD protein level; ↑ catalase activity; ↑ nitrate; ↓ xanthine oxidase activity; ↔ xanthine oxidase protein level; ↔ MPO activity; ↔ MPO protein level; ↔ nitrite; ↔ nitrotyrosine index; ↔ carbonylated proteins 2) Mediterranean diet + nuts vs control: ↑ SOD activity; ↑ SOD protein level; ↑ catalase activity; ↑ nitrate; ↓ xanthine oxidase activity; ↔ xanthine oxidase protein level; ↔ MPO activity; ↔ MPO protein level; ↔ nitrite; ↔ nitrotyrosine index; ↔ carbonylated proteins
Choi et al. [57], 2015	Korea	Parallel RCT, 8 weeks	Intervention: N = 21, Mean age: 73.0 (3.9) years	Intervention: DASH diet Control: Dietary counselling	Plasma: TBARS; FRAP	↓ TBARS; ↔ FRAP

(continued on next page)



Table 2 (continued)

Authors, Year	Country	Study design, duration	Participants	Dietary pattern	Biomarkers	Results <sup>a</sup> Increase in biomarker concentration (↑); Decrease in biomarker concentration (↓); No change (↔)
			Control: N = 18, Mean age: 73.8 (5.8) years 0% male, women with abdominal obesity			
<b>Intervention trials (no comparison group)</b>						
Kaliora et al. [45], 2019	Greece	24 weeks	N = 44, mean (sd) age: 50.4 (10.2) years, 40.9% male, patients with nonfibrotic NAFLD	Intervention: Mediterranean diet	Serum: Ox-LDL; CRP; TNF-α; IL-6; visfatin; leptin	↓ Ox-LDL; ↓ CRP; ↓ visfatin; ↔ TNF-α; ↔ IL-6; ↔ leptin
Luisi et al. [46], 2019	Italy	3 months	Two groups: 1) Overweight or obese: N = 18, age range: 20–61 years, 61% male 2) Normal weight: N = 18, age range: 24–71 years, 33% male	Intervention: Mediterranean diet	MPO activity; MDA; 8-OH-dG; TNF-α; IL-6; IL-10; adiponectin	1) Overweight or obese: ↓ MPO activity; ↓ MDA; ↓ 8-OH-dG; ↑ IL-10; ↑ adiponectin; ↓ TNF-α; ↓ IL-6 2) Normal weight: ↓ MPO activity; ↓ MDA; ↓ 8-OH-dG; ↑ adiponectin; ↓ TNF-α; ↓ IL-6; ↔ IL-10
Rodríguez et al. [47], 2015	Chile	3 months	N = 47, age range: 25–45 years, 0% male, overweight and obese premenopausal women	Intervention: Mediterranean diet (excluding wine)	Serum: CML	↓ CML
Bloomer et al. [48], 2015	USA	21 days	Intervention (3 arms): 1) N = 12, mean (sd) age: 31.1 (4.7) years 2) N = 12, mean (sd) age: 27.9 (3.8) years 3) N = 11, mean (sd) age: 31.5 (4.5) years 16% male, healthy adults	Intervention (3 arms): 1) Traditional Daniel Fast diet <sup>b</sup> 2) Modified Daniel Fast diet <sup>c</sup> 3) Unrestricted vegan diet	Plasma: MDA; nitrate/nitrite; AOPP; hs-CRP	1) Traditional Daniel Fast diet: ↔ MDA; ↔ nitrate/nitrite; ↔ AOPP; ↔ hs-CRP 2) Modified Daniel Fast diet: ↔ MDA; ↔ nitrate/nitrite; ↔ AOPP; ↔ hs-CRP 3) Unrestricted vegan diet: ↔ MDA; ↔ nitrate/nitrite; ↔ AOPP; ↔ hs-CRP
Parcina et al. [49], 2015	Germany	2 weeks	Intervention (3 arms): 1) N = 14, mean (sd) age: 31.9 (6.3) years 2) N = 13, mean (sd) age: 29.1 (5.8) years 3) N = 12, mean (sd) age: 27.4 (5.7) years 100% male, healthy adults	Intervention (3 arms): 1) Mediterranean diet 2) Habitual German diet 3) Fast food diet	Serum: 8-OH-dG; MDA; methylglyoxal; homocysteine	1) Mediterranean diet: ↔ 8-OH-dG; ↔ MDA; ↔ methylglyoxal; ↔ homocysteine 2) Habitual German diet: ↔ 8-OH-dG; ↔ MDA; ↔ methylglyoxal; ↔ homocysteine 3) Fast food diet: ↔ 8-OH-dG; ↔ MDA; ↔ methylglyoxal; ↔ homocysteine

Abbreviations: AOPP, advanced oxidation protein products; CML, N(6)-carboxymethyllysine; CPT1, carnitine palmitoyl transferase 1; CRP, C-reactive protein; DASH, Dietary Approaches to Stop Hypertension; ELISA, enzyme-linked immunosorbent assay; FRAP, ferric reducing ability of plasma; GSH, glutathione; hs, high-sensitivity; IFN-γ, interferon gamma; IL, interleukin; IP-10, interferon gamma induced protein-10; ox-LDL, oxidized low density lipoprotein; MCP-1, monocyte chemoattractant protein-1; MDA, malonyldialdehyde; MIP-1β, macrophage inflammatory protein-1 beta; MPO, myeloperoxidase; NO, nitric oxide; RCT, randomized control trial; ROS, reactive oxygen species; SD, standard deviation; SOD, Superoxide dismutase; TAC, total antioxidant capacity; TBARS, thiobarbituric acid reactive substances; TNF-α, tumor necrosis factor alpha; VEGF, vascular endothelial growth factor; 8-OH-dG, 8-hydroxy-2-deoxyguanosine.

<sup>a</sup> P < 0.05.

<sup>b</sup> Traditional Daniel Fats Diet eliminate all processed foods, white flour products, additives, preservatives, sweeteners, flavorings, caffeine, alcohol.

<sup>c</sup> Animal products, including lean meat and milk.

evaluated observational studies, 8 studies scored as high quality, 6 as moderate quality and one as low quality. Among the 13 included intervention studies, one study scored as high quality, 8 as moderate quality and 4 as low quality.

#### 4. Discussion

This systematic review provides a comprehensive summary and evaluation of the recent evidence from human studies on the association between dietary patterns and biomarkers of oxidative stress and inflammation. Overall results from both observational and intervention studies indicated an inverse association between plant-based diets - the Mediterranean and DASH diet - and oxidative stress and proinflammatory biomarkers. In addition, the vegetarian diet, the USDA HEI diet and the paleolithic diet were associated with lower levels of oxidative stress and inflammation, whereas Western and fast-food diets were positively associated with oxidative stress and inflammation biomarkers based on evidence from observational studies. To our knowledge this is the first systematic review to provide an updated summary

and evaluation of various dietary patterns and biomarkers of oxidative stress and inflammation depicting novel trends in human research within the last few years.

##### 4.1. Dietary patterns and oxidative stress biomarkers

The studies in the current systematic review explored a number of dietary patterns among which the most commonly assessed was the Mediterranean diet. This may not be surprising due to the increased popularity of the health beneficial properties of the Mediterranean diet and its components in the recent years. Greater adherence to the Mediterranean diet was consistently associated with a lower risk of cardiovascular disease, diabetes, cancer and neurodegenerative diseases [81, 82], as well as with reduced overall mortality [83]. To explain these favorable associations, the potential ability of the Mediterranean diet to decrease oxidative stress due to its high antioxidant capacity emerged as one of the leading candidate hypotheses [84–87].

The Mediterranean diet is characterized by high intakes of fruit, vegetables, cereals, legumes, nuts, and seeds; a low-to-moderate intake

**Table 3**

Classification of commonly assessed biomarkers of oxidative stress and inflammation in the studies included in the systematic review.

Biological process	Biomarker	Short description (mechanistic pathway)	Interpretation	References <sup>a</sup>
<b>Reactive oxygen and nitrogen species (RONS)</b> RONS (e.g., ROMs, NO)		Products and byproducts of biological processes that mediate signal transduction and induce oxidative damage.	↑Oxidative stress	[58,59]
<b>Biomarkers of oxidative damage</b>				
Lipid peroxidation	Isoprostanes (F <sub>2</sub> -Isoprostanes) Ox-LDL	Formed by free radical-catalyzed oxidation of arachidonic acid. Originated from oxidative modification hypothesis of atherosclerosis.	↑Oxidative stress ↑Oxidative stress	[60] [60]
	Malondialdehyde (MDA)	Reactive aldehyde derived from lipid peroxidation of various polyunsaturated fatty acids that can form DNA- and protein adducts.	↑Oxidative stress	[61,62]
	TBARS	TBA-reactive substances (reactive carbonyl groups-containing compounds) measured as proxy of MDA levels. <sup>b</sup>	↑Oxidative stress	[61,62]
Oxidative damage to proteins/amino acids	Protein carbonyls Nitrotyrosine	Result of oxidative cleavage of protein backbones. Nitration of tyrosine (free amino acid or within a peptide) induced by RNS.	↑Oxidative stress ↑Oxidative stress	[60] [60]
Oxidative DNA damage	8-OH-dG	Oxidative stress induced base modification. If not repaired, 8-OH-dG can lead to GC-TA transversion (mutation).	↑Oxidative stress	[60]
<b>Reactive metabolic products and byproducts</b>				
Metabolic byproducts	Methylglyoxal	Highly reactive dicarbonyl compound that is a by-product of glycolysis and major cell-permeant precursor of AGEs.	↑Oxidative stress	[63]
Glycooxidation - Protein or lipid become glycation	AGEs (e.g., CML)	Heterogeneous group of molecules formed in a nonenzymatic reaction between reducing sugars and amino groups (lipids, DNA, proteins) during normal metabolism.	↑Oxidative stress	[60]
<b>Antioxidant defense mechanisms (detoxification of ROS)</b>				
Superoxide dismutases	SOD1, 2 and 3	A family of metalloenzymes that catalyze the dismutation of two molecules of superoxide anion to hydrogen peroxide and molecular oxygen, functioning as powerful antioxidant in the cells.	↓Oxidative stress	[64]
Catalase system	Catalase	Catalyzes the detoxification of hydrogen peroxide.	↓Oxidative stress	[64]
Glutathione system	GSH Glutathione reductase	The reduced form of the most important low molecular weight antioxidant synthesized in cells involved in detoxification of reactive substances. Detoxifies GSSG, a potentially toxic product of the oxidation of GSH.	↓Oxidative stress ↓Oxidative stress	[65] [65]
	Glutathione peroxidase	Catalyzes the detoxification of hydrogen peroxide to water, and lipid peroxides to their corresponding alcohols.	↓Oxidative stress	[64]
Thioredoxin system	Thioredoxin reductase	NADPH-dependent reducing enzyme in reactions involving thioredoxin.	↓Oxidative stress	[66]
Paraoxonases	PON1, 2 and 3	Reflect antioxidant activity and play a fundamental role in detoxification of many compounds (e.g., early oxidative products).	↓Oxidative stress	[67]
Heme oxygenases	Heme oxygenase-1	Antioxidant and anti-inflammatory enzyme that is usually expressed at low levels, but that can be highly upregulated in response to oxidative stress (Nrf2-regulated)	↑Oxidative stress	[68]
Antioxidant capacity	TAC FRAP	General term that depicts total antioxidant capacity which can be assessed following various approaches. Result of a reaction of sample antioxidants with inorganic oxidants (e.g., Fe <sup>3+</sup> or Cu <sup>2+</sup> ), representing rather the reducing capacity.	↓Oxidative stress ↓Oxidative stress	[69] [69]
<b>Immune-inflammatory activation as sources of oxidative stress</b>				
Immune-system machinery for generating ROS	MPO AOPP sNOX2-dp	Enzyme stored in the granules of neutrophils that catalyzes the formation of hypochlorous acid from hydrogen peroxide. Formed mainly by chlorinated oxidants as a result from activity of myeloperoxidase. Product of NOX2 activation, an enzyme that plays a role in ROS generation by phagocytic leukocytes.	↑Oxidative stress ↑Oxidative stress ↑Oxidative stress	[70,71] [72] [73,74]
	Lipoprotein-associated phospholipase A2	Enzyme secreted from immune cells regulated by cytokines and steroid hormones that can release isoprostanes from esterified phospholipids.	↑Oxidative stress	[75]
<b>Biomarkers of immune-inflammatory pathways</b>				
Acute phase reactant	hsCRP	Produced mainly in the liver in response to increase of proinflammatory cytokines that represents a sensitive and nonspecific marker of systemic low-grade inflammation.	↑Inflammation	[76]
Cytokines	IL-6 TNF-α IL-10	Produced in response to infections, tissue injuries, hematopoiesis, and other immune reactions. Its dysregulation plays a role in on chronic inflammatory states and autoimmunity. Induces inflammation, activation of vascular endothelium, recruitment of immune cells, and tissue destruction and plays a role in chronic inflammation. Anti-inflammatory function, with a central role in preventing inflammatory and autoimmune diseases.	↑Immune activation ↑Immune activation ↓Immune activation	[77] [78] [79]
Adipokines	Adiponectin Leptin	Plays a role in various aspects of metabolism and exerts anti-inflammatory activity (e.g., inhibiting phagocytic activity and IL-6 and TNF production). Plays a role in various aspects of metabolism and exerts pro-inflammatory and immune activation properties.	↓Inflammation ↑Inflammation	[80] [80]

Abbreviations: AOPP, advanced oxidation protein products; CML, N(6)-carboxymethyllysine; FRAP, ferric reducing ability of plasma; GSSG, glutathione (oxidized); GSH, glutathione (reduced); hsCRP, high-sensitivity C-reactive protein; IL, interleukin; MDA, malondialdehyde; MPO, myeloperoxidase; NADPH, nicotinamide adenine dinucleotide phosphate; NO, nitric oxide; NOX2, NADPH oxidase 2; ox-LDL, oxidized low-density lipoprotein; PON, paraoxonase; ROMs, reactive oxygen

metabolites; RONS, reactive oxygen and nitrogen species; sNOX2-dp, soluble NOX2-derived peptide; SOD, superoxide dismutase; TAC, total antioxidant capacity; TBARS, thiobarbituric acid reactive substances; TNF- $\alpha$ , tumor necrosis factor- $\alpha$ ; 8-OH-dG; 8-oxo-2'-deoxyguanosine.

<sup>a</sup> Selected publications describing mechanistic pathways of biomarkers.

<sup>b</sup> TBARS Assay is characterized by low sensitivity and specificity.

**Table 4**

Overview of results on significant differences between biomarker levels in participants according to dietary patterns reported in observational and intervention studies.

Dietary patterns	Observational studies		Intervention studies	
	Oxidative stress biomarkers	Inflammatory biomarkers	Oxidative stress biomarkers	Inflammatory biomarkers
Mediterranean diet	N = 6/NS = 3 [31,32,38] ↓ F <sub>2</sub> -isoprostane [39,40] ↓ sNox2-dp [30,44] ↓ Ox-LDL [42] ↑ FLOP <sub>320</sub> ; ↑ FLOP <sub>360</sub> [41] ↑ TAC [42] ↑ Ox-LDL [39]	N = 2/NS = 0 ↓ hs-CRP [40] ↓ CRP [42] ↓ IL-6; ↓ TNF- $\alpha$ ; ↓ serum amyloid A; ↓ homocysteine [42]	N = 8/NS = 1 [49] ↓ F <sub>2</sub> -Isoprostanes [54] ↓ Ox-LDL [45] ↓ MDA (measured via TBARS) [46] ↓ TBARS [51] ↓ 8-OH-dG [46] ↓ Methylglyoxal [50] ↓ CML [47] ↓ MPO activity ↓ Cystine [52] ↑ nitrate; ↓ xanthine oxidase activity; ↑ SOD activity; ↑ SOD protein level; ↑ catalase activity [56]	N = 3/NS = 2 [49,54] ↓ hs-CRP [50], ↓ CRP [45] ↓ visfatin [45] ↓ TNF- $\alpha$ ; ↓ IL-6 [46] ↑ IL-10; ↑ adiponectin [46]
Vegetarian Diet	N = 4/NS = 2 [33] ↓ MDA [36,43] ↓ MPO protein level; ↓ spondin-2; ↓ follistatin; ↓ MMP-7; ↓ TRAIL-R2 [35] ↑ PON-3 [35]; ↑ TAC [36] ↑ Total phenol; ↑ FRAP, ↑ nitrite [37] ↑ SOD protein level; ↑ glutathione peroxidase protein level [43]	N = 0/NS = 1 [35]	N = 1/NS = 0 ↓ L-derived ROS [51] ↓ TBARS [51]	–
Vegan Diet	N = 1/NS = 0 ↑ TBARS; ↑ nitrite [37]	–	N = 0/NS = 1 [48]	N = 0/NS = 1 [48]
DASH Diet	N = 1/NS = 0 ↑ FLOP <sub>320</sub> ; ↑ FLOP <sub>360</sub> [41]	–	N = 2/NS = 0 ↓ MDA [55] ↓ TBARS [57] ↑ NO; ↑ GSH [55]	N = 1/NS = 0 ↓ hs-CRP [55]
USDA HEI diet	N = 1/NS = 1 [34] ↑ FLOP <sub>320</sub> ; ↑ FLOP <sub>360</sub> [41]	N = 0/NS = 1 [34]	–	–
Paleolithic diet	N = 1/NS = 0 ↓ F <sub>2</sub> -isoprostane [40]	N = 1/NS = 0 ↓ hs-CRP [40]	–	–
Western diet/Fast food diet	N = 6/NS = 1 [43] ↑ heme oxygenase-1 [29] ↑ Lipoprotein-associated phospholipase A2 [32] ↑ PON-3 [35] ↑ Total phenol [37] ↓ protein carbonyl [33] ↓ MMP-7; ↓ MPO protein level [35] ↓ TAC; ↑ MDA [36] ↓ Ceruloplasmin [33]	N = 1/NS = 1 [29] ↑ resistin ↑ TRAIL-R2; ↑ UPAR; ↑ IL2RA [35] ↑ IL1RT1 ↑ follistatin; ↑ CEACAM8 (Western diet); ↓ IL1RT1 (Fast food diet) [35] ↓ Gal-4 [35]	N = 0/NS = 2 [49,53]	N = 0/NS = 1 [49]

NS = number of studies that found non-significant findings of measured biomarkers in corresponding category ( $P > 0.05$ ).

Abbreviations: CEACAM8, carcinoembryonic antigen-related cell adhesion molecule 8; CML, N(6)-carboxymethyllysine; CRP, C-reactive protein; DASH diet, Dietary Approaches to Stop Hypertension diet; FLOP, fluorescent oxidation product; FRAP, ferric reducing ability of plasma; Gal-4, Galectin-4; GSH, glutathione; HEI, Healthy Eating Index; hs, high-sensitivity; IL, interleukin; IL1TR1, interleukin-1 receptor type 1; IL2RA, interleukin-2 receptor subunit alpha; ox-LDL, oxidized low density lipoprotein; MDA, malonyldialdehyde; MMP-7, matrix metalloproteinase-7; MPO, myeloperoxidase; N, number; NO, nitric oxide; PON-3, paraoxonase 3; ROS, reactive oxygen species; sNOX2-dp, soluble Nox2-derived peptide; SOD, Superoxide dismutase; TAC, total antioxidant capacity; TBARS, Thiobarbituric acid reactive substances; tGSH, total glutathione; TNF- $\alpha$ , tumor necrosis factor alpha; TRAIL-R2, tumor necrosis factor-related apoptosis inducing ligand receptor 2; UPAR, urokinase plasminogen activator surface receptor; 8-OH-dG, 8-hydroxy-2-deoxyguanosine.

of dairy products, fish, poultry and wine; and low intakes of red meat and eggs; with olive oil used as a main source of fat [88]. The synergistic effects of the various plant-based foods with antioxidant potential may explain why the overall quality of the diet could be more valuable compared to single food components. In this vein, the observational studies that explored associations between Mediterranean diet and oxidative stress biomarkers provided promising results reporting inverse associations for circulating levels of ox-LDL [89] and MDA [90] and positive associations with biomarkers that reflect antioxidant defense

mechanisms such as SOD and glutathione peroxidase activity [90], and plasma ratio of reduced to oxidized glutathione (GSH/GSSG ratio) [91]. However, intervention trials yielded inconsistent results with some studies showing that Mediterranean diet was associated with decreased blood levels of MDA, TBARS [92], ox-LDL [93] and urine levels of F<sub>2</sub>-isoprostane and 8-oxo-dG [94], others have reported no change in oxidative stress biomarkers such as blood MDA [95] and TBARS [96] levels, and urinary F<sub>2</sub>-isoprostanes [97]. A previous systematic review that summarized results from four intervention studies published up to

2012 further concluded that the evidence was inconsistent [25]. Potential explanations of these inconsistencies could include the differences in study designs, the lack of standard definition of Mediterranean diet and the variety of biomarkers and analytical techniques used for their measurement. Compared to previous work, the current systematic review included a larger number of studies of both observational ( $n = 6$ ) and intervention design ( $n = 8$ ). Consistently, negative associations between Mediterranean diet and blood biomarkers of lipid peroxidation were shown in both observational and intervention studies for  $F_2$ -isoprostanes [39,40,54] and in intervention studies for MDA (measured via TBARS or as a proxy from TBARS) [46,51]. Both,  $F_2$ -isoprostanes and MDA are considered to be among the most reliable available markers to assess oxidative stress [98]. Both higher levels of  $F_2$ -isoprostanes, a lipid peroxidation product of arachidonic acid, and MDA, a reactive aldehyde derived from lipid peroxidation of various polyunsaturated fatty acids that can form DNA adducts [60–62,99], have shown to be associated with several chronic diseases, e.g., cardiovascular diseases, type 2 diabetes, neurodegenerative diseases and cancer [60,100].

The oxidation of lipids, particularly of LDL cholesterol, has been long suggested to predispose the atherosclerotic lesion formation [101,102]. Thus, ox-LDL has been associated with several cardio-metabolic diseases, including cardiovascular diseases and type 2 diabetes [103], however, it is important to note that ox-LDL is a non-specific measure of oxidative stress [60]. Overall, the extent of lipid peroxidation largely depends on the generation of oxygen free radicals, the presence of lipid substrates, and the activity of antioxidants [99]. Thus, increased consumption of dietary antioxidants through diet high in fruits and vegetables and low in saturated fat, total fat, and cholesterol such as the Mediterranean diet may protect against oxidative stress-mediated lipid peroxidation via decreasing lipid substrate available for peroxidation and increasing the concentration of antioxidants [104]. In fact, the current review has shown that Mediterranean diet has been associated with increased TAC in an observational study [42], while showing to increase SOD and catalase activity (enzymes involved in the detoxification of ROS) and decrease the activity of xanthine oxidase [56] and MPO [46] in intervention studies. The antioxidant properties of the Mediterranean diet itself might also be involved in mechanisms of ROS detoxification [56]. Furthermore, the current review suggested that Mediterranean diet leads to a decrease in 8-OH-dG levels, one of the best known and widely used biomarker of oxidative DNA damage [46]. 8-OH-dG is an oxidative DNA base lesion that is increased in oxidative stress conditions and that, if not repaired, can lead to GC  $\rightarrow$  TA transversion – a point mutation [105]. The interaction of ROS with DNA along with of membrane lipid peroxidation products such as MDA lead to formation of numerous DNA adducts [99]. It is well known that DNA stability is essential for the maintenance of normal cell functions and the damaged DNA promotes a spectrum of acute and chronic disorders [106, 107]. These results are in line with a recent review of human intervention studies ( $n = 8$ ) on the effect of Mediterranean diet on markers of DNA damage and DNA repair that suggested that intervention with Mediterranean diet alone or in combination with bioactive-rich foods is protective against DNA damage [108]. One plausible mechanism explaining this effect may involve the balance between polyunsaturated fatty acids (PUFA) and saturated fatty acids (SFA) [108]. For example, inverse associations have been reported between monounsaturated fatty acids (MUFAs) and DNA damage, whereas positive associations were seen between intake of SFAs and DNA damage [109].

Our review also suggested that the DASH diet is associated with lower levels of oxidative stress biomarkers. The DASH diet is characterized by consumption of high amount of fruit and vegetables, low sodium intake and low-fat milk and dairy products [20]. This type of diet not only effectively reduced blood pressure in intervention studies [110], but was also shown to lead to a lower risk of chronic disease incidence and mortality [111,112]. Previous research has shown inconclusive results for the association between this type of diet and oxidative stress biomarkers [25]. Despite we could only identify a few

studies on the DASH diet, the results from intervention studies support its potential to reduce oxidative stress and inflammation levels. In particular, the DASH diet was shown to lower biomarkers of lipid peroxidation, i.e., MDA and TBARS [55,57], and to increase antioxidant status biomarkers, i.e., GSH levels. Similar to the Mediterranean diet, the DASH diet contains plant-based foods that are rich in antioxidants which regular intakes could favor a better balance between cellular oxidant and antioxidant systems and support the regulation of the oxidizing (redox) mechanisms in health and disease states [113]. Furthermore, results suggested that dietary intervention based on the DASH diet led to increased levels of nitric oxide, known as a potent vasodilator [55]. This may be reasonable since this type of diet is composed by many vegetables with high nitrate content that can especially promote the formation of nitric oxide and protect against the initiation of salt-induced hypertension and associated cardio-vascular complications [113]. This finding may be of special interest in designing prevention strategies in high-risk populations characterized by high salt intakes [114]. Our review included studies that evaluated different types of commonly assessed plant-based diets such as different vegetarian diets, USDA HEI diet and the Paleolithic diet [34,37,41]. However, these were mainly explored in observational research setting and no intervention studies were available to evaluate effects of these diets in modulating oxidative stress biomarkers.

In addition to favorable dietary patterns, our review identified a number of observational studies that explored unhealthy dietary patterns classified as ‘Western diet’ and ‘fast-food diet’ in relation to oxidative stress biomarkers [29,32,33,35–37]. Empirically derived dietary patterns characterized by low consumption of fruits, vegetables, legumes and fiber-rich foods, but with high amounts of refined grains, sugar-sweetened beverages, red and processed meat, were associated with higher levels of biomarkers of lipid peroxidation, i.e., MDA [36] and lipoprotein-associated phospholipase A2 [32] and lower levels of biomarkers of antioxidant defense, i.e., TAC [36] and MPO protein levels [35]. Western diets have been associated with obesity and metabolic dysfunction [115]. Adipose tissue is an active endocrine organ releasing a variety of biologically active molecules known as adipokines. The prolonged adipokine secretion in obesity leads to chronic low-grade inflammation and oxidative stress, thereby potentially predisposing chronic disease development [116]. However, due to the cross-sectional study designs employed so far, causal inference on the link between Western diet and oxidative stress mediators is limited. To better understand the potentially detrimental role of Western diets in modulating various aspects of oxidative stress and inflammation, further studies with repeated assessment of dietary patterns and measurements of changes in biomarker concentrations over time are highly warranted.

#### 4.2. Dietary patterns and inflammatory biomarkers

Having the close interdependence between oxidative stress and inflammation, the present systematic review also included studies that simultaneously assessed dietary patterns in relation to inflammatory biomarkers. It is known that inflammatory cells can produce large amounts of ROS as part of an immunological defense mechanism to protect human organisms against invading pathogens [117]. In line with previously discussed improvement in oxidative stress markers, results from observational studies [29–44] and intervention studies [45–57] collectively suggested a corresponding reduction in levels of common biomarkers of inflammation and immune response – CRP, IL-6 and TNF- $\alpha$  as well as favorable modulation of specific adipokines – visfatin and adiponectin following Mediterranean and DASH diets [40,42,45,46, 50,55]. These results extend on our previous work on a systematic review and meta-analysis of intervention trials that evaluated different types of plant-based diets which revealed that the Mediterranean diet and the DASH led to the largest reduction in inflammatory biomarkers among different evaluated diet types [118]. These associations could be partially accounted for by the array of phytochemicals and other

compounds present in plant-based diets i.e. carotenoids and flavonoids that may directly or indirectly modulate inflammatory and immunological processes [119].

#### 4.3. Strengths and limitations

Strengths of the present study include the application of state-of-the-art approaches of conducting systematic search process conducted by several investigators, the variety of dietary patterns and spectrum of biomarkers that allowed making a comprehensive overview of research evidence published on this topic. The review is also based on most recently published studies that allow depicting novel tendencies in research.

Limitations of the present study also warrant consideration. Due to the large heterogeneity of evaluated biomarkers used in studies as proxies of oxidative stress and inflammation and the various analytical techniques used for measuring biomarker concentrations, the quantification of effect size for individual biomarkers by means of meta-analysis was not feasible in this systematic review. The numerous biomarkers of oxidative stress can bear own advantages and limitations, including the lack of tissue and signaling pathway specificity in target tissues [120]. Therefore, some of the non-significant results may be explained by inability of measured biomarkers to reflect a true change rather than by the inability of diet to modulate oxidative stress levels. The various biomarkers represent different pathways and possibly complementing pathways, therefore assessing single biomarkers may not be representative of the level of oxidative stress, however so far none of the studies attempted to evaluate a combination of multiple oxidative stress biomarkers as endpoint in their analyses.

The observational studies had cross-sectional design which does not allow making inferences on the causal effects of dietary patterns in modulating oxidative stress and inflammation and may be influenced by potential bias and confounding. Despite of the prospective study design and randomization that address these limitations of observational studies, the quality of intervention studies ranged mostly from low to moderate and the majority of studies included low number of participants and were of short duration. Differences in the study populations and methods used to evaluate the adherence or compliance to dietary patterns may have also contributed to differences in reported associations with biomarkers.

#### 4.4. Implications for future research

Approaches for reducing the generation of oxidative stress are increasingly deemed important in chronic disease risk prevention, especially in adult and older age, when most of the endogenous antioxidant defense systems fail to offer appropriate protection against elevated oxidative stress and the human organism is exposed to increased ROS formation in ageing cells. While intervention trials on synthetic antioxidants failed to support the beneficial effects of these compounds in preventing age-related diseases [15], the current results suggest that following a balanced plant-based diet, such as the Mediterranean diet, may represent an important alternative to targeted disease prevention. Thus, a healthy adult person may not need additional vitamin and mineral supplements if he follows a balanced food pattern that includes diversity of plant sources. However, to further strengthen this evidence a number of gaps in research are still to be filled.

Our review revealed that studies used to employ a variety of biomarkers and analytical techniques for their quantification with often poorly reported biomarker measurement information. Although numerous analytical methods for assessing oxidative stress have been developed in the recent years, so far there have not been studies to validate biomarker use in epidemiological research in a systematic manner [60]. This largely restricts validity of conclusions regarding observed effects particularly in a prospective study setting where biomarkers serve as endpoints and assessment of exposure-related

long-term effects are of major interest. Reliability assessment studies, measurements of panels of biomarkers and improved reporting of biomarker measurements could prove useful in improving the quality of the studies and the understanding of how diets might affect different aspects of oxidative stress and inflammatory pathways.

Individual genetic variation is another factor to be taken into account in future research when studying the effect of diet on biomarkers of oxidative stress. Genetic variation in antioxidant enzymes may influence the susceptibility of oxidative stress and may be a potential modifier when studying the responses to dietary interventions rich in antioxidants [121]. As an example, nutrigenetic studies suggested that fruit and vegetable consumption from diet can modify the association between polymorphisms in genes coding antioxidant enzymes and breast cancer [122]. So far, few studies have utilized biomarkers of oxidative stress in understanding the link between diet, genetic variation and diseases associated with oxidative stress [121].

With regards to dietary pattern evaluation, most of the studies used *a priori* dietary indices and no study was identified to apply innovative approaches in nutritional epidemiological data modeling of developing *a posteriori* patterns, i.e., identifying patterns that explain largest variation in oxidative stress biomarkers. In observational studies, future investigation should therefore include methodological studies based on population cohorts to identify dietary patterns best suited to reduce oxidative stress levels. Further methodological modeling is also needed to determine whether specific dietary components or combinations of components could play more protective role than others research. In this context, novel statistical approaches for complex data modeling, incl. bioinformatics and machine learning techniques could be employed to establish novel food and biomarker combinations. Prospective cohort studies with available biosample collections could be designed to explore mediating effects of biomarkers of oxidative stress and inflammation on observed associations with incident disease outcomes. For example, based on data from a large prospective cohort study we previously reported that the association between obesity and colorectal cancer was partly mediated by biomarkers of oxidative stress and inflammation [123]. Additionally, studies with repeated measurements over longer periods of time in healthy individuals will be valuable to assess whether dietary patterns limit the onset of chronic diseases, when corrected for confounding factors such as environment and lifestyle.

In intervention research, future randomized trials should include larger and longer trials conducted in heterogeneous populations to assess adherence, efficacy, and effect of dietary patterns on a range of oxidative stress and inflammatory biomarkers assessed as individual and combined exposure. In addition, randomized trials are also warranted to assess the relative effectiveness of specific dietary patterns, i.e., Mediterranean diet compared with other healthy diets, i.e., the DASH or healthy diet. Finally, further work from well conducted randomized trials is needed to identify novel dietary patterns with high antioxidant potential (potentially generated by observational studies) able to counteract systemic as well as mitochondrial-derived oxidative stress, enhance the endogenous antioxidant defenses, and alleviate symptoms or prevent complications of oxidative-stress associated diseases.

## 5. Conclusion

In conclusion, the current systematic review of observational and intervention studies suggested that the plant-based diets, including the Mediterranean and DASH diet, bear potential in reducing concentrations of various biomarkers of oxidative stress and inflammation. These findings are consistent with observed beneficial effects of plant-based dietary patterns on age-related pathologies. Nevertheless, due to the modest quality of evidence, future well-designed dietary trials using validated biomarkers are needed to corroborate the evidence highlighting the beneficial effects of dietary patterns on oxidative stress and inflammation.



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## 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 A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.redox.2021.101869>.

## Author contributions

Conceptualisation, KA (Krasimira Aleksandrova); writing—original draft preparation, KA (Krasimira Aleksandrova); writing—review and editing, KA, LK, CER (Krasimira Aleksandrova, Liselot Koelman, Caue Egea Rodrigues); supervision, KA (Krasimira Aleksandrova). All authors have read and agreed to the published version of the manuscript.

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