

## **Changes in Plasma Tryptophan are Inversely Associated with Incident Cardiovascular Disease in the PREDIMED Study\***

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**RUNNING TITLE:** tryptophan and cardiovascular disease

**TOTAL WORD COUNT:** 5250

**NUMBER OF FIGURES:** 1

**NUMBER OF TABLES:** 4

**OSM:** 5 Supplemental Tables and 1 Supplemental Figures

**\*FINANCIAL SUPPORT:** This study was funded, by the Spanish Ministry of Health (Instituto de Salud Carlos III) and the Ministerio de Economía y Competitividad-Fondo Europeo de Desarrollo Regional (Projects CNIC-06/2007, RTIC G03/140, CIBER 06/03, PI06-1326, PI07-0954, PI11/02505, SAF2009-12304 and AGL2010-22319-C03-03 ) and by the Generalitat Valenciana (ACOMP2010-181, AP111/10, AP-042/11, ACOM2011/145, ACOMP/2012/190, ACOMP/2013/159 and ACOMP/213/165).

**POTENTIAL CONFLICTS OF INTEREST:** JSS has received grants from the International Nut and Dried Fruit Foundation and is a nonpaid member of the scientific advisory board of the International Nut and Dried Fruit Foundation. ER has received grants from the California Walnut Commission and is a nonpaid member of its scientific advisory committee. No other authors declare any conflicts of interest.

**AUTHOR LAST NAMES (FOR PUBMED INDEXING):** Yu, Ruiz-Canela, Guasch-Ferré, Zheng, Toledo, Clish, Salas-Salvadó, Liang, Wang, Corella, Fitó, Gómez-Gracia, Lapetra, Estruch, Ros, Cofán, Arós, Romaguera, Serra-Majem, Sorlí, Hu, Martinez-Gonzalez

1 **ABSTRACT**

2

3 **BACKGROUND**

4 During development of cardiovascular disease (CVD), interferon- $\gamma$ -mediated  
5 inflammation accelerates degradation of tryptophan into downstream metabolites. A  
6 Mediterranean Diet (MedDiet) consisting of high intake of extra virgin olive oil, nuts,  
7 fruits, vegetables, and cereals has been demonstrated to lower risk of CVD. The  
8 longitudinal relationship between these metabolites and CVD in the context of a  
9 MedDiet is unstudied.

10

11 **OBJECTIVE**

12 We sought to investigate the relationship between metabolites in the tryptophan-  
13 kynurenine pathway and CVD in the context of diet.

14

15 **METHODS**

16 We employed a case-cohort design nested in the PREDIMED randomized controlled  
17 trial. There were 231 CVD cases (stroke, myocardial infarction, vascular death) among  
18 985 participants over a median of 4.7 y of follow-up [mean  $\pm$  SD age: 67.6  $\pm$  6.1 y; 53.7%  
19 women; mean  $\pm$  SD body mass index: 29.7  $\pm$  3.7 kg/m<sup>2</sup>]. We assessed plasma  
20 tryptophan, kynurenine, kynurenic acid, 3-hydroxyanthranilic acid, and quinolinic acid  
21 levels at baseline and after 1 year of intervention with MedDiet. We combined these  
22 metabolites in a kynurenine risk score (KRS) by weighting each metabolite by the

23 adjusted coefficient of its associations with CVD. Cox models were used in the primary  
24 analysis.

25

## 26 **RESULTS**

27 Changes in tryptophan after 1 year were associated with lower risk of composite CVD  
28 (hazard ratio [HR] per SD: 0.79; 95% CI: 0.63, 0.98). Baseline kynurenic acid was  
29 associated with higher risk of myocardial infarction and CHD death, but not stroke. A  
30 higher KRS was more strongly associated with CVD in the control group than in the 2  
31 intervention groups (*P*-interaction: 0.003). Adjustment for tryptophan changes  
32 attenuated the inverse association between MedDiet+EVOO and CVD.

33

## 34 **CONCLUSIONS**

35 Changes in tryptophan were significantly associated with decreased risk of CVD. A  
36 MedDiet may counteract the deleterious effect of an unfavorable tryptophan-kynurenine  
37 metabolite profile.

38

39 **KEYWORDS:** Metabolomics, Cardiovascular disease, Mediterranean diet, tryptophan

40 **ABBREVIATIONS**

41

42 3-HAA, 3-hydroxyanthranilic acid

43 ANOVA, analysis of variance

44 BCAA, branched chain amino acids

45 BMI, body mass index

46 CHD, coronary heart disease

47 CI, confidence interval

48 CVD, cardiovascular disease

49 EVOO, extra-virgin olive oil

50 HR, hazard ratio

51 ICAM-1, Intercellular adhesion molecule 1

52 IDO, indoleamine 2,3-dioxygenase

53 IFN- $\gamma$ , interferon- $\gamma$

54 IL-6, interleukin 6

55 KA, kynurenic acid

56 KRS, kynurenine risk score

57 Kyn, kynurenine

58 LDL, low density lipoprotein

59 MedDiet, Mediterranean Diet

60 NAD, nicotinamide adenine dinucleotide

61 *P*, probability

62 PREDIMED, Prevención con Dieta Mediterránea

- 63 Q, quartile
- 64 QA, quinolinic acid
- 65 SD, standard deviation
- 66 Tryp, tryptophan
- 67 VCAM-1, Vascular cell adhesion protein 1

## 68 INTRODUCTION

69

70 Inflammation has long been identified as an important component in the  
71 pathophysiology of cardiovascular disease (CVD) (1). Interferon- $\gamma$  (IFN- $\gamma$ ) is a key  
72 molecule that orchestrates many pathways involved in inflammation by acting as a  
73 transcriptional regulator for immune-related genes (2). Its excessive release by  
74 activated T-lymphocytes has been implicated in the pathogenesis of chronic  
75 inflammation and autoimmune disorders (3). IFN- $\gamma$  has also been shown to activate  
76 macrophages, which are intimately involved in plaque formation, and can also trigger  
77 the release of reactive oxygen species, a hallmark of subsequent atherogenesis and  
78 progression of CVD(4). Another well-known consequence of IFN- $\gamma$  release is the  
79 activation of the enzyme indoleamine 2,3-dioxygenase (IDO), which catabolizes  
80 tryptophan into kynurenine (5). The classic kynurenine pathway, involves the  
81 breakdown of tryptophan into downstream products such as kynurenine, kynurenic acid  
82 and 3-hydroxyanthranilic acid, the latter of which is eventually converted into quinolinic  
83 acid and picolinic acid (6). The final result is the production of one equivalent of  
84 nicotinamide adenine dinucleotide (NAD) for every processed tryptophan molecule (7).  
85 To date, most scientific literature regarding this pathway has focused on its relationship  
86 to neurological disorders such as Alzheimer's disease, Huntington's disease, and  
87 depression (8). These publications stem from the early finding that tryptophan is an  
88 essential precursor to brain serotonin and melatonin (9). Some studies relating  
89 tryptophan metabolites to CVD have reported inverse associations with tryptophan and  
90 the risk of CVD (10-16). However, to our knowledge, no studies have examined whether



91 changes in tryptophan levels are prospectively associated with the incidence of CVD.  
92 Furthermore, the effects of diet on metabolites of this pathway in relation to CVD have  
93 not been investigated.

94 In the present case-cohort study nested in the PREvención con Dieta  
95 MEDiterránea (PREDIMED) trial, we used repeated measurements of these metabolites  
96 to longitudinally assess: a) whether baseline and 1-year changes in tryptophan,  
97 kynurenine, kynurenic acid, 3-hydroxyanthranilic acid (3-HAA), and quinolinic acid were  
98 associated with future risk of CVD; b) whether these associations differed by the type of  
99 CVD events (stroke vs. non-stroke events [i.e. coronary heart disease and non-stroke  
100 related cardiac death]). In light of previous research on cognitive diseases, we  
101 hypothesized that a favorable tryptophan profile would be more beneficial in preventing  
102 stroke, as there may be common risk factors for both stroke and cognitive disorders,  
103 such as inflammation or preclinical cerebrovascular disease; c) whether significant  
104 associations could be counteracted by a Mediterranean diet (MedDiet); and d) whether  
105 the cardioprotective effect of a MedDiet is attenuated after adjusting for changes in  
106 these metabolites.

107

## 108 **METHODS**

109

### 110 *Study population and design*

111

112 The PREDIMED trial is a multicenter, randomized trial of dietary interventions  
113 with a MedDiet supplemented with either nuts or extra-virgin olive oil (EVOO) for the

114 primary prevention of CVD compared to a low-fat control group (17). Protocol, design,  
115 and primary results are detailed elsewhere (18, 19). Briefly, 7447 eligible men and  
116 women at high risk of CVD were randomly assigned to a MedDiet supplemented with a  
117 free provision for the family of 1 L/week EVOO (MedDiet + EVOO), a MedDiet  
118 supplemented with 30 g/day mixed nuts (15 g walnuts, 7.5 g hazelnuts, and 7.5 g  
119 almonds) (MedDiet + nuts), or a control diet consisting of advice to reduce the intake of  
120 all types of fat (control group). The primary endpoint was a composite CVD, defined as  
121 stroke, myocardial infarction, or death from cardiovascular causes. Information on  
122 primary end points was collected by physicians and from other sources of information,  
123 such as the National Death Index. Only end points that were confirmed by the  
124 adjudication committee and that occurred between October 2003, and December 2010,  
125 were included in the analyses.

126         For the present study, we sampled approximately 10% of all PREDIMED  
127 participants in a case-cohort design. A random selection of 791 participants was chosen  
128 at baseline as subcohort. From the full cohort (including the subcohort), there were 231  
129 CVD cases during follow-up and were included in the analysis. Of the CVD cases, 118  
130 experienced a stroke (113 ischemic and 5 hemorrhagic), and 113 non-stroke events.  
131 We excluded individuals from specific analyses if they were missing metabolite values  
132 for that analysis. At baseline and at 1-year of follow-up, 896 and 806 participants had  
133 valid measurements for all five metabolites, respectively.

134

135 *Quantification of Metabolites*

136

137 Fasting plasma EDTA tubes were collected for all participants, and aliquots were  
138 coded and kept refrigerated until they were stored at  $-80^{\circ}\text{C}$ . Pairs of samples (baseline  
139 and first-year visits from each participant) were randomly ordered and shipped to the  
140 Broad Institute of Harvard and MIT (Cambridge, MA, USA) for metabolomics analyses.  
141 Liquid chromatography tandem mass spectrometry to quantitatively profile metabolites  
142 in fasting plasma collected at baseline and year 1 of the intervention. Reference  
143 standards were used to confirm metabolite identities. The details of this procedure are  
144 described elsewhere (20, 21).

#### 145 *Statistical Analyses*

146  
147 Rank-based inverse normal transformations were used to transform the non-  
148 normal distributions of the five metabolites under study (22). We calculated a  
149 kynurenine risk score (KRS) using as weights the adjusted coefficients obtained in the  
150 assessment of the association between the respective metabolite and the risk of CVD.  
151 We first regressed composite CVD on each individual metabolite in a fully adjusted Cox  
152 model, then multiplied the resulting beta by the normalized values for the corresponding  
153 metabolite. These five products were then summed to produce the KRS. If any of the  
154 individuals were missing any metabolite values, we did not calculate a KRS for that  
155 individual. Each normalized metabolite was also analyzed according to its quartile  
156 distribution. Quartile cutpoints were generated on the basis of the distributions of  
157 metabolites and score among non-cases. Scores for acylcarnitines (20), BCAAs (21),  
158 and ceramides were also calculated in this fashion. We conducted tests of linear trend

159 by examining an ordinal score on the basis of the median value in each quartile of  
160 metabolites and score in the multivariable models.

161 Baseline data by quartiles of the KRS were presented as means ( $\pm$  standard  
162 deviations) for continuous variables and N and percentages for categorical variables.  
163 Baseline characteristics were compared across quartiles of the KRS using chi-squared  
164 tests for categorical variables and one-way analysis of variance (ANOVA) for  
165 continuous variables.

166 We used weighted Cox regression models with non-cases up-weighted by their  
167 sampling fraction (22), and used robust variance to account for correlation between  
168 observations (23). We calculated hazard ratios (HRs) and their 95% confidence  
169 intervals (95% CIs) for the composite CVD end point, and also separately for stroke and  
170 non-stroke cases (myocardial infarction and non-stroke CVD mortality). Follow-up time  
171 was calculated from the date of enrollment to the date of diagnosis of CVD for cases,  
172 and to the date of the last visit or the end of the follow-up period (December 1, 2010) for  
173 non-cases. In model 1, we adjusted for age, sex, family history of CHD, smoking status,  
174 and body mass index, and stratified by intervention group. In model 2, we additionally  
175 adjusted for model 1 plus baseline hypertension, dyslipidemia, diabetes, and scores for  
176 branched chain amino acids (BCAAs), acylcarnitines, and ceramides (previously  
177 reported as associated with CVD in our cohort). For models including 1-year change  
178 variables, we also adjusted for continuous baseline levels of that metabolite. We  
179 additionally conducted subgroup analyses by restricting the analysis to the group of  
180 interest. To test for effect modification, we used a likelihood ratio test to compare the

181 model without interaction terms vs. model with interaction terms (indicator variable for  
182 subgroup x KRS) among all participants.

183 For the mediation analysis of the MedDiet+EVOO, we report the coefficient for  
184 the MedDiet+EVOO vs. control (excluding individuals allocated to the MedDiet+nuts  
185 group) with and without adjustment for both baseline and change in tryptophan. We  
186 repeated this analyses for MedDiet+nuts excluding the MedDiet+EVOO participants.

187 To calculate 1-year changes in metabolites, we performed an ANOVA for each  
188 metabolites and calculated adjusted means of metabolite values at baseline and at 1  
189 year, stratified by intervention group. We used paired t-tests to test if changes in  
190 metabolites were significant within each arm of the trial.

191 For correlation among metabolites and scores, we used Spearman correlation  
192 coefficients and p-values.

193 All statistical analyses were performed with SAS (v9.4, SAS Institute, Cary, NC)  
194 and R (v2.13.0, R Foundation, Vienna, Austria).

## 195 **RESULTS**

196

### 197 *Descriptive Results*

198

199 Demographic and behavioral characteristics of the participants in the subcohort  
200 with valid measurements of all five metabolites at baseline (n=724) are classified  
201 according to quartiles of KRS in **Table 1**. Increasing quartiles of KRS were significantly  
202 associated with age, % female sex, % hypertension, and % of current smokers. The  
203 median follow-up time was 4.7 years.

204 1-year changes in tryptophan metabolites according to intervention group are  
205 shown in **Supplemental Figure 1**. Metabolites of those randomized to the MedDiet with  
206 EVOO group uniformly decreased after 1-year. Using paired t-tests for the non-  
207 transformed metabolite quantifications at baseline and 1-year, we observed significant  
208 changes in: tryptophan and 3-HAA in the MedDiet with EVOO group, and kynurenic acid  
209 and quinolinic acid in the MedDiet with nuts group.

210 Spearman correlation coefficients of tryptophan metabolites, branched chain  
211 amino acids, ceramides, and acylcarnitine score are depicted in **Supplemental Table 1**.  
212 In general, tryptophan metabolites had low to moderate correlations with each other ( $r =$   
213  $0.00$  to  $0.48$ ), low to moderate correlation with branched chain amino acid score ( $r =$   
214  $0.03$  to  $0.49$ ), and modest correlations with ceramides and acylcarnitine scores ( $r = -$   
215  $0.29$  to  $0.14$ ).

216

217 *Individual Metabolites, Risk Score, and Composite CVD*

218

219 **Table 2** details the associations of baseline and 1-year changes of kynurenine  
220 pathway metabolites with composite CVD incidence. Only baseline kynurenic acid was  
221 associated with composite CVD risk in the fully adjusted model (HR per SD: 1.23; 95%  
222 CI: 1.02, 1.48;  $P < 0.05$ ). Using 1-year changes, an increase in tryptophan was  
223 associated with a significantly reduced risk in the categorical analysis (HR for Q4 vs. Q1:  
224  $0.49$ ; 95% CI: 0.26, 0.95;  $P$ -trend  $< 0.05$ ) models, as well as in the continuous analysis  
225 (HR per SD: 0.79; 95% CI: 0.63, 0.98;  $P < 0.05$ )

226 Since the kynurenine pathway has been examined most often in relation to brain  
227 disorders, we hypothesized that associations of these metabolites would be stronger for  
228 stroke than for non-stroke events. Using stroke as the outcome in **Table 3**, we found no  
229 significant associations of any baseline metabolite concentration with stroke risk.  
230 Furthermore, the inverse association observed for 1-year changes in tryptophan was  
231 not present (HR per SD: 0.91; 95% CI: 0.70, 1.19;  $P = 0.49$ ) when using stroke alone as  
232 the outcome. In **Table 3** when using only non-stroke events as the outcome, kynurenic  
233 acid (HR per SD: 1.46; 95% CI: 1.12, 1.92;  $P < 0.05$ ) was associated with higher  
234 subsequent incidence of outcomes different from stroke (myocardial infarction or any  
235 other cause of vascular death). Furthermore, the relationship between 1-year increase  
236 in tryptophan and non-stroke outcomes was significantly associated with a lower risk of  
237 events in both the continuous models (HR per SD: 0.65; 95% CI: 0.46, 0.91;  $P < 0.05$ )  
238 and categorical models (HR for Q4 vs. Q1: 0.19; 95% CI: 0.06, 0.61;  $P$ -trend  $< 0.05$ ). 1-  
239 year changes in kynurenine were also inversely associated with non-stroke event risk in  
240 the categorical analysis (HR for Q4 vs. Q1: 0.62; 95% CI: 0.28, 1.36;  $P$ -trend  $< 0.05$ ).

241 In **Table 4**, using the KRS as exposure, we observed a similar pattern of  
242 elevated incidence of the composite CVD outcome (HR per SD: 1.41; 95% CI: 1.14,  
243 =1.75;  $P < 0.05$ ), and a strengthened association with non-stroke events (HR per SD:  
244 1.70; 95% CI: 1.27, 2.29;  $P < 0.001$ ). We also considered the ratio of kynurenine to  
245 tryptophan in **Supplemental Table 2**, but found no significant associations of this ratio  
246 with any CVD outcome. Thus, we proceeded with the KRS to test effect modification by  
247 a Mediterranean diet.

248

249 *Effect Modification by a Mediterranean Diet*

250

251 In **Supplemental Table 3**, we sought to address whether several baseline  
252 variables potentially modified the positive association of the KRS with CVD. We only  
253 found significant interaction according to intervention group ( $P$ -interaction = 0.003), with  
254 those in the control group experiencing a greater elevation in risk per SD increase (HR  
255 per SD: 2.02; 95% CI: 1.31, 3.13) compared to the MedDiet + EVOO group (HR per SD:  
256 1.27; 95% CI: 0.83, 1.94) and the MedDiet + nuts group (HR per SD: 1.23; 95% CI: 0.79,  
257 1.92).

258 **Figure 1** depicts the fully adjusted HRs stratified by intervention (combined  
259 MedDiet vs. control) and quartiles of the baseline KRS. Compared to those randomized  
260 to a MedDiet and in the lowest quartile of baseline KRS, we found a significant positive  
261 association for individuals randomized to the control group and quartiles 2 – 4 of the  
262 KRS, but not for those randomized to the MedDiet and quartiles 2 – 4 of the KRS.

263

264 *Attenuation of the Benefits of a Mediterranean Diet by Tryptophan Changes*

265

266 In **Supplemental Tables 4 and 5** we explore the strength of the protective  
267 association of a MedDiet before and after adjustment of tryptophan. When comparing  
268 MedDiet + EVOO vs. control, adjustment for baseline and 1-year changes in tryptophan  
269 attenuated the inverse association from HR per SD: 0.62; 95% CI: 0.38, 1.01 to HR per  
270 SD: 0.95; 95% CI: 0.66, 1.36. This attenuation appeared to be weaker for stroke and  
271 greater for non-stroke outcomes. However, when comparing MedDiet + nuts vs. control,



272 we did not observe these patterns of attenuation. After controlling for baseline and 1-  
273 year changes in tryptophan, the inverse association for the MedDiet + nuts only slightly  
274 changed from HR per SD: 0.61; 95% CI: 0.37, 1.00 to HR per SD: 0.60; 95% CI: 0.35,  
275 1.02. Similarly, effect size was lower for stroke events and higher for non-stroke events.

276

## 277 **DISCUSSION**

278

279 In the present case-cohort study, we report that lower baseline kynurenic acid  
280 and higher baseline tryptophan, were associated with lower risk of non-stroke events,  
281 and 1-year increase in tryptophan levels was strongly associated with lower risk of CVD  
282 and non-stroke events. A combined score of five plasma metabolites in this pathway  
283 (tryptophan, kynurenine, kynurenic acid, 3-hydroxyanthranilic acid, and quinolinic acid),  
284 designed to summarize the pathway in a single parameter by maximizing predictiveness  
285 of outcome within the present study, was significantly associated with the risk of hard  
286 clinical events of CVD. Furthermore, consuming a MedDiet counteracted the harmful  
287 effect of an unfavorable metabolite profile in the tryptophan-kynurenine pathway. Lastly,  
288 changes in tryptophan may be involved in the cardioprotective benefits of the MedDiet.

289 The finding that tryptophan is inversely associated with CVD incidence is in line  
290 with previous publications (24-26). However, we report no significant associations of  
291 kynurenine/tryptophan ratio with any CVD outcome. Clinical studies have reported that  
292 tryptophan degradation, as operationalized by low blood tryptophan levels or a high  
293 kynurenine/tryptophan ratio, were predictive of coronary heart disease status (15),  
294 elevated oxidative stress in patients with kidney dysfunction (13), and greater mean

295 carotid artery intima-media thickness (27). Large scale epidemiologic studies have  
296 reported positive associations of tryptophan catabolism (i.e. lower levels of tryptophan  
297 and higher levels of downstream metabolites in the tryptophan-kynurenine pathway)  
298 with primary acute coronary events (28), worse prognosis after diagnosis of coronary  
299 artery disease (14), and greater risk of mortality from CVD(16).

300       Using repeated measurements of metabolites, our observation that baseline and  
301 1-year increases in tryptophan are associated with lower risk of CVD events (especially  
302 when stroke was excluded) needs to be compared with the existing literature which has  
303 suggested that tryptophan levels are decreased in stroke patients compared to controls  
304 (12), and that lower blood tryptophan concentrations are related to greater infarct  
305 volume in stroke patients (11) and worse prognosis (29) and faster cognitive decline (30)  
306 after stroke. Our results are consistent with an inverse association between tryptophan  
307 and CVD, but they may differ from those of previous studies according to the subtype of  
308 CVD event, given that we have not found a stronger effect for stroke than for other  
309 events. A potential explanation might be that existing publications largely used cross-  
310 sectional designs examining stroke patients at baseline, whereas we used a longitudinal  
311 design for both the exposure and the outcome and followed disease-free individuals for  
312 repeated measurements of metabolites after an intervention and also for incident  
313 endpoints. Mangge et al. (31) suggested that decreases in tryptophan levels may be a  
314 consequence of chronic low-grade inflammation, rather than a cause of disease. Our  
315 findings of an inverse association between changes in tryptophan and non-stroke  
316 endpoints would support the inflammation hypothesis for non-stroke events, given the

317 anti-inflammatory properties of the MedDiet. However, further studies are needed to  
318 elucidate the differing effects of tryptophan on stroke vs. non-stroke CVD events.

319         The causal role of tryptophan and kynurenine pathway metabolites in CVD  
320 remains poorly understood. It is thought that IFN- $\gamma$  plays a central role in the activation  
321 of IDO and subsequent degradation of tryptophan (4); however, activation of the  
322 kynurenine pathway has also been shown to have anti-inflammatory effects (3).  
323 Treatment of human peripheral blood mononuclear cells and monocyte-derived  
324 macrophages with IFN- $\gamma$  attenuated the extent of low density lipoprotein (LDL) oxidation,  
325 and tryptophan degradation in concert with 3-HAA formation was instrumental in this  
326 inhibitory effect (32). 3-HAA has also been independently identified as having anti-  
327 atherogenic properties by regulating lipid metabolism and inflammation (33, 34). Other  
328 experimental studies suggest a beneficial effect of IDO on the vasculature. IDO-  
329 deficient mice fed high-fat diets showed marked increases in F4/80 and TNF mRNA  
330 levels, as well as greater hepatic inflammation compared to controls (35). IDO inhibition  
331 also blunted the protective effects of eicosapentaenoic acid in LDLr<sup>-/-</sup> mice (31). In light  
332 of the prevailing theme in experimental studies that IDO-mediated degradation of  
333 tryptophan is beneficial, we speculate that activation of the tryptophan-kynurenine  
334 pathway may be a compensatory mechanism to, rather than a cause of, inflammation  
335 and cardiovascular dysfunction. Furthermore, since most individuals consume adequate  
336 amounts of tryptophan (i.e. do not suffer from tryptophan deficiency), the protection  
337 conferred by the MedDiet is unlikely to be related to greater availability of tryptophan  
338 (36).

339           The novel finding that a MedDiet may offset the deleterious effects of a high-risk  
340 profile in metabolites of the tryptophan-kynurenine pathway also warrants discussion.  
341 To our knowledge, this the first report of an association of tryptophan metabolites with  
342 CVD in the context of a nutritional intervention in a large randomized trial. Previous  
343 clinical trials have concluded that close adherence to a MedDiet has a beneficial effect  
344 on inflammatory markers (37, 38), and other PREDIMED reports have noted that  
345 participants randomized to the MedDiet interventions had lower incidence of CVD  
346 events compared to those in the control group, even among those with comparable  
347 levels of plasma branched chain amino acids (21) or acylcarnitines (20). These  
348 publications, in addition to support the favorable interaction of the MedDiet with various  
349 CVD biomarkers also point to the need of future experimental studies to clarify the  
350 biological mechanisms underlying this effect.

351           The randomization of dietary interventions at baseline allowed us to study  
352 possible mediating effects of tryptophan in the association between MedDiet and the  
353 risk of CVD. We found that among the MedDiet + EVOO arm, adjustment for changes in  
354 tryptophan attenuated the HRs for CVD. These results suggest a possible role of  
355 tryptophan degradation (or preservation) as a mediator in the causal pathway of EVOO  
356 consumption and CVD prevention. We acknowledge that our statistical power for  
357 detecting interactions is limited. Consumption of both EVOO (39) and nuts (40, 41) has  
358 been associated with reduced circulating levels of inflammatory biomarkers, although  
359 the specific roles of IFN- $\gamma$  and tryptophan degradation in diets enriched with these foods  
360 are not well characterized. Primary results from the PREDIMED trial also reported no  
361 striking differences between the two MedDiet groups in relation to composite CVD or

362 secondary outcomes (18). However, dietary exposure to walnuts has been  
363 characterized by changes in various metabolites, including intermediate metabolites of  
364 the tryptophan pathway (42). Future lines of inquiry should investigate the biological  
365 roles of IFN- $\gamma$ , tryptophan, and related kynurenine metabolites in diets involving EVOO  
366 or nuts.

367         Strengths of the present study include blood draws at repeated intervals to  
368 assess changes in metabolites, the prospective design of the cohort, as well as  
369 adjustment for potential confounders related to CVD. Our study also has limitations.  
370 First, we did not measure all metabolites on interest in the tryptophan-kynurenine  
371 pathway, such as picolinic acid or 3-hydroxykynurenine. Second, we cannot rule out the  
372 possibility that concentrations of the metabolites were different between missing and  
373 non-missing cases. Lastly, results from our study among a population of high-risk  
374 participants living in the Mediterranean region may not be generalizable to individuals of  
375 different demographics.

376         Our results indicate that 1-year changes in tryptophan are predictive of lower  
377 CVD incidence and, especially, non-stroke incidence and that a score combining five  
378 metabolites in the tryptophan-kynurenine pathway is also prospectively associated with  
379 clinical cases of CVD. The harmful effects of an unfavorable tryptophan metabolite  
380 profile were in part mitigated by consuming a MedDiet. The cardioprotective effect of  
381 MedDiets supplemented with EVOO or nuts could be mediated in part by processes  
382 associated with changes in tryptophan.

383

384 **ACKNOWLEDGMENTS**

385 *Sources of Funding*

386 This study was funded by the National Institutes of Health of the United States  
387 (1R01HL118264), Spanish Ministry of Health (Instituto de Salud Carlos III) and the  
388 Ministerio de Economía y Competitividad-Fondo Europeo de Desarrollo Regional  
389 (Projects CNIC-06/2007, RTIC G03/140, CIBER 06/03, PI06-1326, PI07-0954,  
390 PI11/02505, SAF2009-12304 and AGL2010-22319-C03-03 ) and by the Generalitat  
391 Valenciana (ACOMP2010-181, AP111/10, AP-042/11, ACOM2011/145,  
392 ACOMP/2012/190, ACOMP/2013/159 and ACOMP/213/165).

393

394 *Disclosures*

395 JSS has received grants from the International Nut and Dried Fruit Foundation and is a  
396 nonpaid member of the scientific advisory board of the International Nut and Dried Fruit  
397 Foundation. ER has received grants from the California Walnut Commission and is a  
398 nonpaid member of its scientific advisory committee.

399

400 *Author Contributions*

401 EY conducted the analysis and wrote the manuscript. MRC, MGF, YZ, and DDW  
402 provided code and guidance needed to complete the analysis. ET, CBC, JSS, LL, DC,  
403 MF, EGG, JL, RE, ER, MC, FA, DR, LSM, JVS, FBH, and MAM designed the research  
404 plan and oversaw the study. All authors read and approved the final version of the  
405 manuscript.

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Table 1. Baseline characteristics of the full sub-cohort according to quartiles of kynurenine risk score.

	Kynurenine risk score				P value
	Q1	Q2	Q3	Q4	
n	180	181	179	188	
Median Score	-0.43	-0.14	0.08	0.41	
Range	-1.06 to -0.24	-0.24 to -0.03	-0.03 to 0.18	0.18 to 1.08	
Age (years) <sup>1</sup>	66.2 ± 5.8	66.8 ± 6.0	66.8 ± 5.8	69.0 ± 5.7	<0.0001
Sex (% Women)	46.1	57.5	60.3	62.2	0.01
Body mass index, kg/m <sup>2</sup>	29.5 ± 3.6	29.5 ± 3.4	29.8 ± 3.7	30.3 ± 3.5	0.09
Intervention group, %					0.21
MedDiet+EVOO	33.3	32.6	40.2	44.2	
MedDiet+nuts	34.4	37.0	29.1	30.3	
Control	32.2	30.4	30.7	25.5	
Family history of CHD, %	21.7	27.6	22.9	27.7	0.41
Hypertension, %	78.3	84.5	82.1	87.8	0.10
Dyslipidemia, %	71.1	74.6	73.2	73.4	0.90
Diabetes, %	47.2	40.3	53.6	47.9	0.09
Obesity (BMI ≥ 30), %	43.3	40.3	46.9	53.2	0.08
Smoking, %					<0.0001
Never	52.2	65.8	62.6	69.7	
Former	22.2	23.2	26.8	26.6	
Current	25.6	11.1	10.6	3.7	

<sup>1</sup>Data are expressed as means ± SD or percentage. P value for comparisons between cases and controls across quartiles (Pearson  $\chi^2$  test for categorical variables or one way ANOVA for continuous variables) as appropriate.

Table 2. Hazard ratios (95% CI) for composite CVD (stroke, non-stroke, death from vascular causes) by baseline and 1-year changes in plasma metabolites (tryptophan, kynurenine, kynurenic acid, 3-hydroxyanthranilic acid, quinolinic acid) as continuous and categorical variables. Number of participants will vary according to availability of plasma metabolite quantifications.

Baseline <sup>1</sup>					
	Tryptophan	Kynurenine	Kynurenic Acid	3-HAA	Quinolinic Acid
Model 1, Metabolite as continuous variable, per SD					
HR (95% CI)	0.88 (0.74 – 1.04)	1.00 (0.86 – 1.16)	1.14 (0.99 – 1.31)	0.82 (0.69 – 0.98)	1.03 (0.86 – 1.22)
<i>P</i>	0.13	0.99	0.08	0.03	0.77
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	0.91 (0.59 – 1.40)	0.86 (0.55 – 1.35)	1.49 (0.91 – 2.42)	0.83 (0.53 – 1.30)	0.93 (0.59 – 1.47)
Q3	0.67 (0.42 – 1.05)	0.93 (0.61 – 1.44)	1.39 (0.86 – 2.24)	0.62 (0.38 – 1.03)	0.99 (0.64 – 1.54)
Q4	0.75 (0.48 – 1.19)	0.91 (0.59 – 1.40)	1.49 (0.92 – 2.40)	0.69 (0.43 – 1.11)	1.17 (0.75 – 1.83)
<i>P</i> -trend	0.12	0.76	0.14	0.07	0.46
Model 2, Metabolite as continuous variable, per SD					
HR (95% CI)	0.87 (0.67 – 1.11)	1.06 (0.89 – 1.26)	1.23 (1.02 – 1.48)	0.82 (0.67 – 1.01)	1.15 (0.94 – 1.41)
<i>P</i>	0.28	0.52	0.03	0.06	0.19
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	0.95 (0.58 – 1.57)	1.27 (0.75 – 2.13)	1.58 (0.91 – 2.76)	0.72 (0.42 – 1.23)	1.00 (0.60 – 1.67)
Q3	0.58 (0.33 – 1.01)	1.17 (0.72 – 1.92)	1.62 (0.92 – 2.85)	0.84 (0.46 – 1.50)	1.08 (0.65 – 1.82)
Q4	0.75 (0.40 – 1.44)	1.21 (0.73 – 1.99)	1.73 (0.97 – 3.10)	0.64 (0.36 – 1.16)	1.53 (0.91 – 2.57)
<i>P</i> -trend	0.21	0.55	0.08	0.22	0.11
1-Year Changes <sup>2</sup>					
	Tryptophan	Kynurenine	Kynurenic Acid	3-HAA	Quinolinic Acid
Model 1, Metabolite as continuous variable, per SD					
HR (95% CI)	0.77 (0.64 – 0.94)	0.86 (0.69 – 1.06)	0.87 (0.73 – 1.04)	1.15 (0.91 – 1.45)	0.96 (0.80 – 1.15)
<i>P</i>	0.01	0.15	0.13	0.26	0.64
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	1.18 (0.72 – 1.94)	0.79 (0.48 – 1.30)	0.95 (0.56 – 1.61)	1.59 (0.79 – 3.19)	0.84 (0.48 – 1.47)
Q3	0.75 (0.44 – 1.27)	0.37 (0.20 – 0.70)	0.88 (0.50 – 1.53)	1.91 (0.91 – 3.99)	0.95 (0.54 – 1.66)

Q4	0.49 (0.27 – 0.87)	0.68 (0.41 – 1.13)	0.69 (0.40 – 1.20)	1.46 (0.70 – 3.05)	0.89 (0.52 – 1.51)
<i>P</i> -trend	0.005	0.02	0.18	0.27	0.78
Model 2, Metabolite as continuous variable, per SD					
HR (95% CI)	0.79 (0.63 – 0.98)	0.85 (0.67 – 1.06)	0.93 (0.75 – 1.15)	1.15 (0.90 – 1.49)	0.92 (0.75 – 1.13)
<i>P</i>	0.03	0.15	0.52	0.27	0.42
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	1.19 (0.70 – 2.02)	0.78 (0.45 – 1.34)	1.00 (0.53 – 1.87)	1.26 (0.56 – 2.84)	0.82 (0.43 – 1.57)
Q3	0.72 (0.40 – 1.28)	0.39 (0.20 – 0.77)	1.00 (0.52 – 1.93)	1.37 (0.55 – 3.43)	0.97 (0.51 – 1.85)
Q4	0.49 (0.26 – 0.95)	0.60 (0.34 – 1.07)	0.84 (0.44 – 1.58)	1.32 (0.57 – 3.03)	0.78 (0.43 – 1.42)
<i>P</i> -trend	0.01	0.02	0.60	0.50	0.55

*Model 1 was adjusted for age, sex, family history of CHD, smoking status, and body mass index, and was stratified by intervention group.*

*Model 2 was adjusted as for covariates in model 1, plus baseline hypertension, dyslipidemia, diabetes, and scores for BCAAs, acylcarnitines, and ceramides. For change analyses, metabolites were adjusted for baseline values.*

<sup>1</sup>*Baseline analysis consisted of n = 986 participants and n = 231 cases for tryptophan, kynurenine, KA, and quinolinic acid, and n = 904 participants and n = 204 cases for 3-HAA.*

<sup>2</sup>*1-year change analysis consisted of n = 908 participants and n = 162 cases for tryptophan, kynurenine, KA, and quinolinic acid, and n = 750 participants and n = 122 cases for 3-HAA.*

Table 3. Hazard ratios (95% CI) for stroke and non-stroke cases by baseline and 1-year changes in plasma metabolites (tryptophan, kynurenine, kynurenic acid, 3-hydroxyanthranilic acid, quinolinic acid) as continuous and categorical variables. Number of participants will vary according to availability of plasma metabolite quantifications.

STROKE					
Baseline <sup>1</sup>					
	Tryptophan	Kynurenine	Kynurenic Acid	3-HAA	Quinolinic Acid
Model 1, Metabolite as continuous variable, per SD					
HR (95% CI)	0.97 (0.76 – 1.24)	0.97 (0.79 – 1.20)	1.05 (0.87 – 1.26)	0.78 (0.61 – 0.98)	0.91 (0.72 – 1.15)
<i>P</i>	0.81	0.79	0.65	0.04	0.42
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	0.68 (0.38 – 1.22)	0.87 (0.49 – 1.54)	1.43 (0.79 – 2.61)	0.67 (0.37 – 1.21)	0.81 (0.45 – 1.46)
Q3	0.61 (0.34 – 1.10)	0.80 (0.46 – 1.40)	1.00 (0.54 – 1.86)	0.69 (0.37 – 1.30)	0.95 (0.55 – 1.65)
Q4	0.95 (0.54 – 1.66)	0.89 (0.50 – 1.57)	1.16 (0.61 – 2.18)	0.63 (0.34 – 1.17)	0.96 (0.54 – 1.72)
<i>P</i> -trend	0.77	0.62	0.94	0.18	0.97
Model 2, Metabolite as continuous variable, per SD					
HR (95% CI)	0.98 (0.71 – 1.35)	1.02 (0.82 – 1.26)	1.06 (0.83 – 1.37)	0.76 (0.57 – 1.01)	1.05 (0.81 – 1.37)
<i>P</i>	0.89	0.89	0.63	0.06	0.72
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	0.73 (0.38 – 1.41)	1.22 (0.65 – 2.30)	1.43 (0.72 – 2.82)	0.56 (0.28 – 1.12)	0.93 (0.49 – 1.75)
Q3	0.52 (0.25 – 1.09)	0.90 (0.48 – 1.67)	1.07 (0.52 – 2.18)	0.87 (0.42 – 1.80)	1.09 (0.58 – 2.05)
Q4	0.96 (0.43 – 2.16)	1.13 (0.60 – 2.12)	1.19 (0.55 – 2.61)	0.52 (0.24 – 1.12)	1.26 (0.65 – 2.47)
<i>P</i> -trend	0.74	0.96	0.85	0.23	0.43
1-Year Changes <sup>2</sup>					
	Tryptophan	Kynurenine	Kynurenic Acid	3-HAA	Quinolinic Acid
Model 1, Metabolite as continuous variable, per SD					
HR (95% CI)	0.89 (0.70 – 1.12)	0.88 (0.66 – 1.17)	0.90 (0.71 – 1.15)	1.27 (0.93 – 1.72)	0.87 (0.69 – 1.10)
<i>P</i>	0.32	0.37	0.41	0.14	0.24
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	1.51 (0.79 – 2.90)	0.64 (0.33 – 1.26)	0.46 (0.22 – 0.99)	1.65 (0.63 – 4.31)	1.11 (0.54 – 2.27)

Q3	0.88 (0.43 – 1.80)	0.44 (0.20 – 0.98)	0.83 (0.41 – 1.68)	1.90 (0.68 – 5.31)	1.10 (0.54 – 2.25)
Q4	0.84 (0.41 – 1.70)	0.67 (0.34 – 1.31)	0.59 (0.28 – 1.23)	1.79 (0.65 – 4.92)	0.85 (0.43 – 1.70)
<i>P</i> -trend	0.35	0.16	0.40	0.25	0.66
Model 2, Metabolite as continuous variable, per SD					
HR (95% CI)	0.91 (0.70 – 1.19)	0.85 (0.63 – 1.16)	0.97 (0.72 – 1.29)	1.24 (0.90 – 1.72)	0.85 (0.66 – 1.09)
<i>P</i>	0.49	0.31	0.81	0.19	0.20
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	1.60 (0.82 – 3.16)	0.59 (0.28 – 1.25)	0.44 (0.18 – 1.04)	1.30 (0.46 – 3.74)	1.03 (0.46 – 2.29)
Q3	0.90 (0.42 – 1.92)	0.51 (0.23 – 1.14)	0.86 (0.38 – 1.92)	1.42 (0.41 – 4.85)	1.08 (0.49 – 2.36)
Q4	0.89 (0.42 – 1.92)	0.56 (0.26 – 1.18)	0.71 (0.31 – 1.58)	1.55 (0.53 – 4.56)	0.78 (0.37 – 1.63)
<i>P</i> -trend	0.46	0.12	0.78	0.43	0.54
NON-STROKE					
Baseline <sup>3</sup>					
	Tryptophan	Kynurenine	Kynurenic Acid	3-HAA	Quinolinic Acid
Model 1 <sup>2</sup> , Metabolite as continuous variable, per SD					
HR (95% CI)	0.78 (0.63 – 0.97)	1.03 (0.85 – 1.26)	1.24 (1.02 – 1.51)	0.85 (0.69 – 1.06)	1.17 (0.92 – 1.47)
<i>P</i>	0.03	0.74	0.03	0.14	0.20
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	1.20 (0.67 – 2.16)	0.80 (0.43 – 1.50)	1.50 (0.73 – 3.09)	1.04 (0.56 – 1.91)	1.05 (0.57 – 1.95)
Q3	0.78 (0.42 – 1.44)	1.03 (0.58 – 1.85)	1.90 (0.96 – 3.76)	0.53 (0.26 – 1.10)	1.00 (0.54 – 1.86)
Q4	0.58 (0.30 – 1.11)	0.94 (0.52 – 1.71)	2.02 (1.02 – 3.97)	0.74 (0.40 – 1.40)	1.48 (0.80 – 2.73)
<i>P</i> -trend	0.05	0.94	0.03	0.13	0.26
Model 2, Metabolite as continuous variable, per SD					
HR (95% CI)	0.76 (0.54 – 1.05)	1.10 (0.87 – 1.39)	1.46 (1.12 – 1.92)	0.85 (0.66 – 1.10)	1.30 (0.97 – 1.73)
<i>P</i>	0.09	0.42	0.01	0.22	0.08
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	1.16 (0.60 – 2.26)	1.26 (0.60 – 2.65)	1.77 (0.77 – 4.08)	0.99 (0.49 – 2.00)	1.12 (0.54 – 2.35)
Q3	0.69 (0.33 – 1.43)	1.49 (0.75 – 2.94)	2.49 (1.10 – 5.66)	0.81 (0.36 – 1.82)	1.10 (0.53 – 2.30)
Q4	0.52 (0.21 – 1.29)	1.33 (0.65 – 2.74)	2.89 (1.26 – 6.65)	0.74 (0.35 – 1.60)	2.05 (1.00 – 4.21)
<i>P</i> -trend	0.10	0.39	0.01	0.39	0.07
1-Year Changes <sup>4</sup>					

	Tryptophan	Kynurenine	Kynurenic Acid	3-HAA	Quinolinic Acid
Model 1, Metabolite as continuous variable, per SD					
HR (95% CI)	0.64 (0.48 – 0.86)	0.82 (0.62 – 1.07)	0.82 (0.65 – 1.05)	1.04 (0.76 – 1.42)	1.03 (0.79 – 1.33)
<i>P</i>	0.003	0.15	0.11	0.82	0.84
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	0.93 (0.47 – 1.84)	0.94 (0.48 – 1.81)	1.69 (0.85 – 3.36)	1.59 (0.62 – 4.11)	0.59 (0.26 – 1.37)
Q3	0.63 (0.31 – 1.30)	0.27 (0.11 – 0.68)	0.79 (0.34 – 1.86)	2.03 (0.76 – 5.45)	0.81 (0.36 – 1.83)
Q4	0.22 (0.08 – 0.56)	0.64 (0.32 – 1.28)	0.79 (0.37 – 1.70)	1.19 (0.44 – 3.21)	0.90 (0.43 – 1.88)
<i>P</i> -trend	0.001	0.03	0.24	0.64	0.99
Model 2, Metabolite as continuous variable, per SD					
HR (95% CI)	0.65 (0.46 – 0.90)	0.81 (0.60 – 1.09)	0.85 (0.64 – 1.14)	1.09 (0.78 – 1.53)	0.96 (0.71 – 1.30)
<i>P</i>	0.01	0.17	0.29	0.61	0.81
Metabolite in quartile categories, as compared to Q1 (reference)					
Q2	0.84 (0.39 – 1.84)	0.98 (0.48 – 2.00)	2.14 (0.91 – 5.04)	1.30 (0.39 – 4.36)	0.59 (0.22 – 1.57)
Q3	0.55 (0.24 – 1.27)	0.22 (0.07 – 0.64)	1.08 (0.39 – 2.97)	1.45 (0.40 – 5.28)	0.84 (0.30 – 2.35)
Q4	0.19 (0.06 – 0.61)	0.62 (0.28 – 1.36)	0.98 (0.38 – 2.51)	1.20 (0.37 – 3.88)	0.74 (0.31 – 1.77)
<i>P</i> -trend	0.003	0.02	0.62	0.73	0.70

Model 1 was adjusted for age, sex, family history of CHD, smoking status, and body mass index, and was stratified by intervention group.

Model 2 was adjusted as for model 1, plus baseline hypertension, dyslipidemia, diabetes, and scores for BCAAs, acylcarnitines, and ceramides. For change analyses, metabolites were adjusted for baseline values.

<sup>1</sup>Baseline analysis for stroke consisted of  $n = 872$  participants and  $n = 118$  cases for tryptophan, kynurenine, KA and quinolinic acid, and  $n = 804$  participants and  $n = 103$  cases for 3-HAA. Non-stroke cases are excluded.

<sup>2</sup>1-year change analysis for stroke consisted of  $n = 835$  participants and  $n = 86$  cases for tryptophan, kynurenine, KA, and quinolinic acid, and  $n = 692$  participants and  $n = 63$  cases for 3-HAA. Non-stroke cases are excluded.

<sup>3</sup>Baseline analysis for non-stroke consisted of  $n = 868$  participants and  $n = 113$  cases for tryptophan, kynurenine, KA and quinolinic acid, and  $n = 800$  participants and  $n = 100$  cases for 3-HAA. Stroke cases are excluded.

<sup>4</sup>1-year change analysis for non-stroke consisted of  $n = 825$  participants and  $n = 76$  cases for tryptophan, kynurenine, KA and quinolinic acid,  $n = 687$  participants and  $n = 59$  cases for 3-HAA. Stroke cases are excluded.



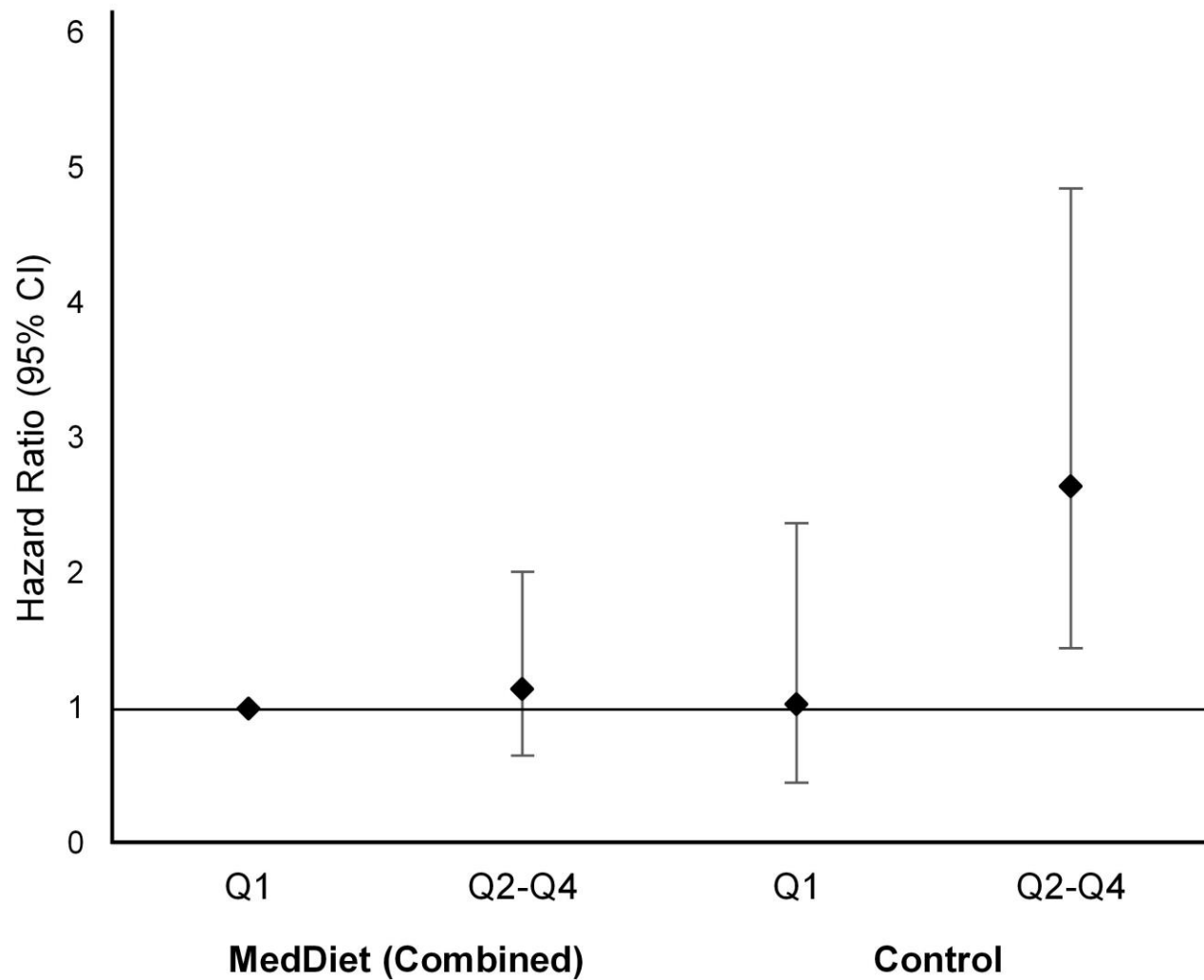
Table 4. Hazard ratios (95% CI) for composite CVD, stroke only, and non-stroke only by baseline kynurenine risk score among participants with available data for all five metabolites under study.

	Outcome		
	Composite CVD	Stroke Only	Non-Stroke Only
n	896	797	793
cases	202	103	99
Model 1, Metabolite as continuous variable, per SD			
HR (95% CI)	1.33 (1.11 – 1.60)	1.17 (0.91 – 1.50)	1.56 (1.22 – 2.00)
<i>P</i>	0.002	0.22	<0.001
Metabolite in quartile categories, as compared to Q1 (reference)			
Q2	1.49 (0.88 – 2.50)	1.17 (0.62 – 2.20)	1.97 (0.92 – 4.22)
Q3	1.03 (0.60 – 1.78)	0.75 (0.38 – 1.47)	1.48 (0.68 – 3.23)
Q4	2.11 (1.26 – 3.52)	1.37 (0.70 – 2.65)	3.72 (1.80 – 7.71)
<i>P</i> -trend	0.02	0.64	0.002
Model 2, Metabolite as continuous variable, per SD			
HR (95% CI)	1.41 (1.14 – 1.75)	1.27 (0.94 – 1.71)	1.70 (1.27 – 2.29)
<i>P</i>	0.002	0.12	<0.001
Metabolite in quartile categories, as compared to Q1 (reference)			
Q2	1.60 (0.89 – 2.86)	1.34 (0.66 – 2.76)	2.07 (0.86 – 5.01)
Q3	1.00 (0.55 – 1.82)	0.71 (0.33 – 1.55)	1.59 (0.67 – 3.78)
Q4	2.41 (1.36 – 4.27)	1.76 (0.80 – 3.88)	4.11 (1.84 – 9.19)
<i>P</i> -trend	0.02	0.40	0.002

*Model 1 was adjusted for age, sex, family history of CHD, smoking status, and body mass index, and was stratified by intervention group.*

*Model 2 was adjusted as for covariates in model 1, plus baseline hypertension, dyslipidemia, diabetes, and scores for BCAAs, acylcarnitines, and ceramides. Kynurenine risk score was built by multiplying normalized individual metabolites (tryptophan, kynurenine, KA, 3-HAA, quinolinic acid) by their beta coefficient in a fully adjusted model with that metabolite alone and then summing up the products for each metabolite x coefficient value. The weights were -0.13474 for tryptophan, 0.0558 for kynurenine, 0.20359 for kynurenic acid, -0.20066 for 3-hydroxyanthranilic acid, and 0.13657 for quinolinic acid.*

Figure 1. Multivariate adjusted HRs (95% CI) of composite CVD by quartiles of baseline kynurenine risk score stratified by intervention group (Mediterranean interventions combined versus control group) among participants with available data for all five metabolites under study (n=896).



Online Supporting Material

Supplemental Table 1. Spearman correlation heatmap of baseline plasma metabolites under study (tryptophan, kynurenine, kynurenic acid, 3-hydroxyanthranilic acid, quinolinic acid) and scores of plasma metabolites related to CVD (branched chain amino acid, ceramides, and acylcarnitine score).

	Tryp	KA	3-HAA	Kyn	QA	CS	SAS	MAS	LAS	BCAAS
Tryp	1	0.13***	0.29***	0.22***	0.00	-0.04	-0.17***	-0.08**	0.13***	0.49***
KA	0.13***	1	0.28***	0.29***	0.31***	-0.09**	-0.24***	0.14***	0.13***	0.30***
3-HAA	0.29***	0.28***	1	0.20***	0.18***	0.00	-0.18***	-0.05	0.01	0.41***
Kyn	0.22***	0.29***	0.20***	1	0.48***	0.10**	-0.16***	0.14***	0.07*	0.06
QA	0.00	0.31***	0.18***	0.48***	1	0.00	-0.29***	0.06*	-0.04	0.03
CS	-0.04	-0.09**	0.00	0.10**	0.00	1	0.04	0.07*	0.18***	0.04
SAS	-0.17***	-0.24***	-0.18***	-0.16***	-0.29***	0.04	1	0.08*	0.05	-0.14***
MAS	-0.08**	0.14***	-0.05	0.14***	0.06*	0.07*	0.08*	1	0.49***	0.07*
LAS	0.13***	0.13***	0.01	0.07*	-0.04	0.18***	0.05	0.49***	1	0.17***
BCAAS	0.49***	0.30***	0.41***	0.06	0.03	0.04	-0.14***	0.07*	0.17***	1

\* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$

Abbreviations: 3-HAA, 3-hydroxyanthranilic acid; BCAAS, branched chain amino acid score; CS, ceramide score; KA, kynurenic acid; Kyn, kynurenine; LAS, long acylcarnitine score; MAS, medium acylcarnitine score; QA, quinolinic acid; SAS, short acylcarnitine score; Tryp, tryptophan.

Online Supporting Material

Supplemental Table 2. Association of baseline plasma kynurenine/tryptophan ratio on composite CVD, stroke only, and non-stroke only among participants with available data for both kynurenine and tryptophan under study.

	Baseline		
	Composite CVD	Stroke Only	Non-Stroke Only
n	984	866	871
cases	231	118	113
Model 1, Metabolite as continuous variable, per SD			
HR (95% CI)	1.07 (0.91 – 1.26)	0.99 (0.79 – 1.24)	1.17 (0.94 – 1.45)
<i>P</i>	0.44	0.89	0.15
Metabolite in quartile categories, as compared to Q1 (reference)			
Q2	1.39 (0.89 – 2.19)	1.13 (0.64 – 1.99)	1.70 (0.91 – 3.17)
Q3	1.12 (0.71 – 1.76)	0.96 (0.54 – 1.68)	1.34 (0.70 – 2.57)
Q4	1.32 (0.84 – 2.07)	1.02 (0.57 – 1.81)	1.80 (0.97 – 3.37)
<i>P</i> -trend	0.42	0.90	0.14
Model 2, Metabolite as continuous variable, per SD			
HR (95% CI)	1.10 (0.91 – 1.37)	1.01 (0.79 – 1.30)	1.22 (0.95 – 1.57)
<i>P</i>	0.32	0.93	0.11
Metabolite in quartile categories, as compared to Q1 (reference)			
Q2	1.67 (0.81 – 3.41)	1.16 (0.48 – 2.82)	2.60 (0.95 – 7.12)
Q3	1.39 (0.54 – 3.55)	1.14 (0.36 – 3.65)	2.07 (0.56 – 7.60)
Q4	1.57 (0.41 – 5.93)	1.10 (0.20 – 6.10)	2.96 (0.50 – 17.47)
<i>P</i> -trend	0.60	0.93	0.30
	1-Year Change		
	Composite CVD	Stroke Only	Non-Stroke Only
n	904	829	818
cases	161	86	75
Model 1, Metabolite as continuous variable, per SD			
HR (95% CI)	1.05 (0.86 – 1.29)	0.89 (0.67 – 1.17)	1.27 (0.96 – 1.68)

Online Supporting Material

<i>P</i>	0.64	0.38	0.10
Metabolite in quartile categories, as compared to Q1 (reference)			
Q2	0.75 (0.44 – 1.28)	0.61 (0.32 – 1.20)	1.05 (0.47 – 2.37)
Q3	0.89 (0.52 – 1.52)	0.44 (0.21 – 0.93)	1.87 (0.88 – 3.96)
Q4	1.03 (0.61 – 1.74)	0.69 (0.36 – 1.31)	1.77 (0.80 – 3.93)
<i>P</i> -trend	0.77	0.17	0.07
Model 2, Metabolite as continuous variable, per SD			
HR (95% CI)	1.03 (0.83 – 1.30)	0.85 (0.63 – 1.15)	1.28 (0.92 – 1.77)
<i>P</i>	0.79	0.30	0.14
Metabolite in quartile categories, as compared to Q1 (reference)			
Q2	0.71 (0.39 – 1.29)	0.57 (0.28 – 1.18)	0.95 (0.38 – 2.41)
Q3	0.76 (0.42 – 1.38)	0.34 (0.15 – 0.80)	1.69 (0.73 – 3.95)
Q4	1.00 (0.56 – 1.80)	0.63 (0.31 – 1.30)	1.78 (0.71 – 4.42)
<i>P</i> -trend	0.95	0.12	0.10

*Cases occurring within one year of follow-up were excluded. For analyses using stroke as the endpoint, non-stroke cases were excluded, and for analyses using non-stroke as the endpoint, stroke cases were excluded.*

*Model 1 was adjusted for age, sex, family history of CHD, smoking status, and body mass index.*

*Model 2 was adjusted as for model 1, plus baseline hypertension, dyslipidemia, diabetes, and scores for BCAAs, acylcarnitines, and ceramides.*

Online Supporting Material

Supplemental Table 3. Subgroup analysis for the associations between baseline kynurenine risk score and risk of composite CVD among participants with available data for all five metabolites under study (n=896).

Characteristic		Continuous HR <sup>1</sup> per 1 SD (95% CI)	P-interaction
Sex	Men (n=419)	1.59 (1.19 – 2.13)	0.23
	Women (n=477)	1.26 (0.90 – 1.75)	
Age	< 65 years (n=312)	1.58 (0.91 – 2.77)	0.73
	≥ 65 years (n=584)	1.39 (1.09 – 1.79)	
Intervention group	Control group (n=281)	2.02 (1.31 – 3.13)	0.003
	Mediterranean diet + EVOO (n=330)	1.27 (0.83 – 1.94)	
	Mediterranean diet + nuts (n=285)	1.23 (0.79 – 1.92)	
Obesity	≥ 30 kg/m <sup>2</sup> (n=406)	1.32 (0.93 – 1.88)	0.07
	< 30 kg/m <sup>2</sup> (n=490)	1.68 (1.18 – 2.39)	
Smoking status	Current/former smoking (n=365)	1.79 (1.21 – 2.64)	0.81
	Never smoking (n=531)	1.46 (1.05 – 2.02)	
Family history of CHD	Yes (n=215)	1.18 (0.72 – 1.93)	0.68
	No (n=681)	1.56 (1.21 – 2.02)	
Baseline T2D	Yes (n=450)	1.26 (0.95 – 1.66)	0.06
	No (n=446)	1.83 (1.26 – 2.67)	
Baseline Hypertension	Yes (n=749)	1.40 (1.10 – 1.79)	0.88
	No (n=147)	1.42 (0.64 – 3.16)	
Baseline Dyslipidemia	Yes (n=631)	1.50 (1.14 – 1.98)	0.51
	No (n=265)	1.40 (0.91 – 2.16)	

<sup>1</sup>Hazard ratios were adjusted for age, sex, family history of CHD, smoking status, body mass index, baseline hypertension, dyslipidemia, diabetes, and scores for BCAAs, acylcarnitines, and ceramides and were stratified by intervention group.

Supplemental Table 4. Analysis for mediation by plasma tryptophan change for the Mediterranean diet with extra virgin olive oil vs. control.

	Outcome		
	Composite CVD	Stroke Only	Non-Stroke Only
n	630	574	556
cases	130	74	56
Model 1			
MedDiet+EVOO vs. Control			
HR (95% CI)	0.66 (0.44 – 1.00)	0.63 (0.38 – 1.06)	0.69 (0.37 – 1.26)
<i>P</i>	0.05	0.08	0.22
Adjusted for 1-year tryptophan change and baseline tryptophan			
HR (95% CI)	0.91 (0.72 – 1.15)	0.68 (0.39 – 1.19)	0.88 (0.43 – 1.83)
<i>P</i>	0.42	0.18	0.74
Model 2			
MedDiet+EVOO vs. Control			
HR (95% CI)	0.62 (0.38 – 1.01)	0.56 (0.30 – 1.02)	0.69 (0.33 – 1.44)
<i>P</i>	0.05	0.06	0.33
Adjusted for 1-year tryptophan change			
HR (95% CI)	0.95 (0.66 – 1.36)	0.62 (0.68 – 1.26)	0.93 (0.40 – 2.17)
<i>P</i>	0.77	0.60	0.86

Cases occurring within one year of follow-up were excluded. For analyses using stroke as the endpoint, non-stroke cases were excluded, and for analyses using non-stroke as the endpoint, stroke cases were excluded.

Model 1 was adjusted for age, sex, family history of CHD, smoking status, and body mass index.

Model 2 was adjusted as for model 1, plus baseline hypertension, dyslipidemia, diabetes, and scores for BCAAs, acylcarnitines, and ceramides.

Supplemental Table 5. Analysis for mediation by plasma tryptophan change for the Mediterranean diet with nuts vs. control.

	Baseline		
	Composite CVD	Stroke Only	Non-Stroke Only
n	575	520	523
cases	107	55	52
Model 1			
MedDiet+nuts vs. Control			
HR (95% CI)	0.62 (0.40 – 0.95)	0.48 (0.27 – 0.86)	0.78 (0.43 – 1.40)
P	0.03	0.01	0.40
Adjusted for 1-year tryptophan change and baseline tryptophan			
HR (95% CI)	0.57 (0.35 – 0.93)	0.47 (0.24 – 0.90)	0.70 (0.34 – 1.44)
P	0.02	0.02	0.33
Model 2			
MedDiet+nuts vs. Control			
HR (95% CI)	0.61 (0.37 – 1.00)	0.43 (0.22 – 0.84)	0.89 (0.45 – 1.77)
P	0.05	0.01	0.73
Adjusted for 1-year tryptophan change			
HR (95% CI)	0.60 (0.35 – 1.02)	0.47 (0.23 – 0.98)	0.83 (0.36 – 1.88)
P	0.06	0.04	0.65

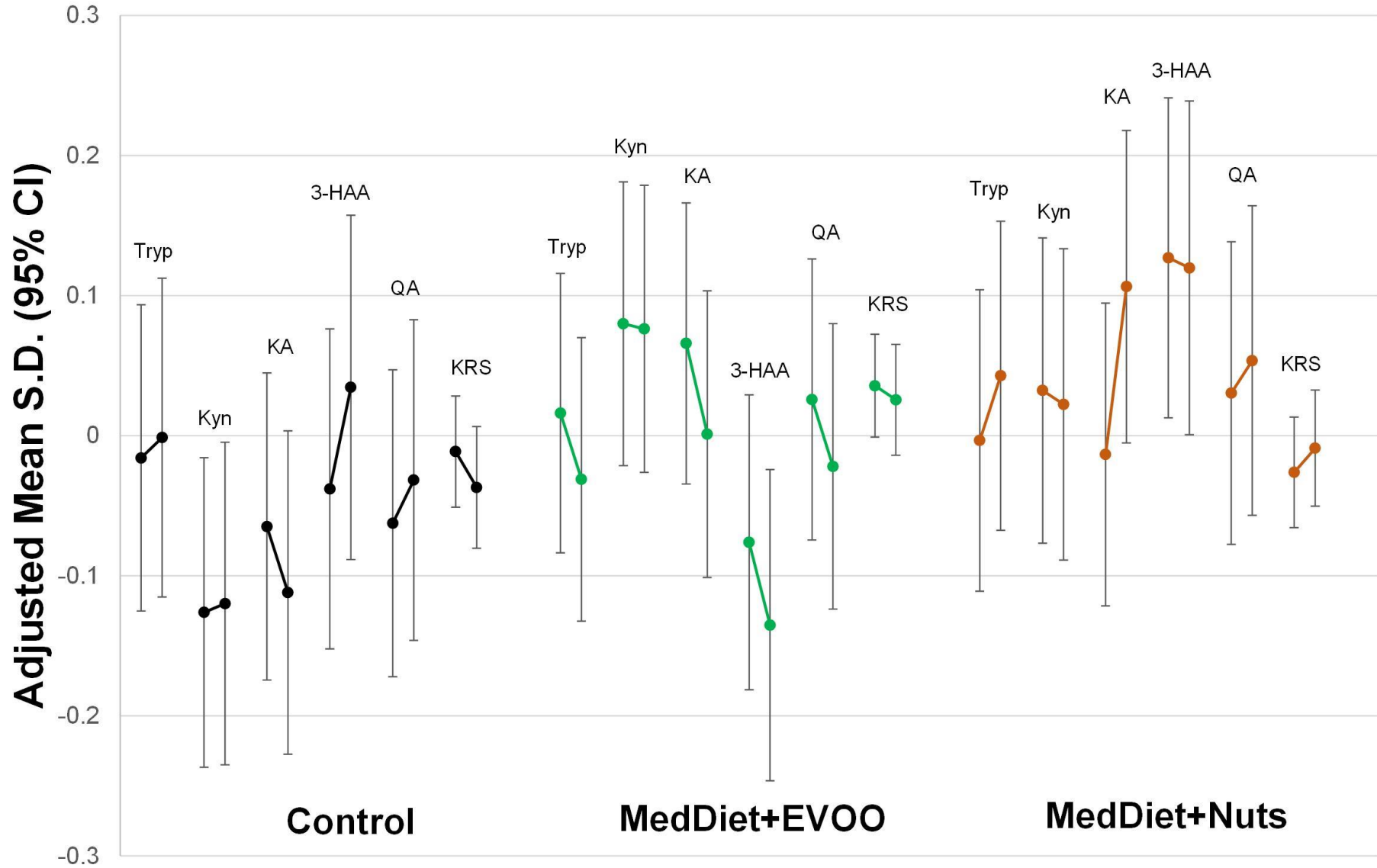
Cases occurring within one year of follow-up were excluded. For analyses using stroke as the endpoint, non-stroke cases were excluded, and for analyses using non-stroke as the endpoint, stroke cases were excluded.

Model 1 was adjusted for age, sex, family history of CHD, smoking status, and body mass index.

Model 2 was adjusted as for model 1, plus baseline hypertension, dyslipidemia, diabetes, and scores for BCAAs, acylcarnitines, and ceramides.



Supplemental Figure 1. Changes in plasma tryptophan, kynurenine, kynurenic acid, 3-hydroxyanthranilic acid, quinolinic acid, and kynurenine risk score from baseline to 1 year (of intervention), stratified by intervention group.



\*Abbreviations: 3-HAA, 3-hydroxyanthranilic acid; KA, kynurenic acid; KRS, kynurenine risk score; Kyn, kynurenine; QA, quinolinic acid; Tryp, tryptophan.