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A systematic review and meta-analysis of the kynurenine pathway of tryptophan metabolism in rheumatic diseases

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There is an increasing interest in the pathophysiological role of the kynurenine pathway of tryptophan metabolism in the regulation of immune function and inflammation. We sought to address the link between this pathway and the presence rheumatic diseases (RD) by conducting a systematic review and metaanalysis of studies reporting the plasma or serum concentrations of tryptophan, kynurenine, and other relevant metabolites in RD patients and healthy controls. We searched electronic databases for relevant articles published between inception and the 30th of June 2023. Risk of bias and certainty of evidence were assessed using the Joanna Briggs Institute Critical Appraisal Checklist and the Grades of Recommendation, Assessment, Development and Evaluation Working Group system. In 24 studies selected for analysis, compared to controls, RD patients had significantly lower tryptophan (standard mean difference, SMD= -0.71, 95% CI -1.03 to -0.39, p<0.001; I² = 93.6%, p<0.001; low certainty of evidence), and higher kynurenine (SMD=0.69, 95% CI 0.35 to 1.02, p<0.001; $l^2 = 93.2\%$, p<0.001; low certainty), kynurenine to tryptophan ratios (SMD=0.88, 95% CI 0.55 to 1.21, p<0.001; I² = 92.9%, p<0.001; moderate certainty), 3-hydroxykynurenine (SMD=0.74, 95% CI 0.30 to 1.18, p=0.001; I^2 = 87.7%, p<0.001; extremely low certainty), and quinolinic acid concentrations (SMD=0.71, 95% CI 0.31 to 1.11, p<0.001; I² = 88.1%, p<0.001; extremely low certainty). By contrast, there were non-significant between-group differences in kynurenic acid, 3-hydroxyanthranilic acid, kynurenic acid to kynurenine ratio, or quinolinic acid to kynurenine acid ratio. In metaregression, the SMD of tryptophan, kynurenine, and kynurenine to tryptophan ratio were not associated with age, publication year, sample size, RD duration, Creactive protein, or use of anti-rheumatic drugs and corticosteroids. In subgroup analysis, the SMD of tryptophan, kynurenine, and kynurenine to tryptophan ratio was significant across different types of RD, barring rheumatoid arthritis. Therefore, we have observed significant alterations in tryptophan, kynurenine, 3hydroxykynurenine, and quinolinic acid concentrations in RD patients. Further research is warranted to determine whether these biomarkers can be useful for diagnosis and management in this patient group. (PROSPERO registration number: CRD CRD42023443718).

Systematic review registration: https://www.crd.york.ac.uk/prospero, identifier CRD CRD42023443718.

KEYWORDS

tryptophan, kynurenine, kynurenine to tryptophan ratio, 3-hydroxykynurenine, quinolinic acid, rheumatic diseases, inflammation, biomarkers

Introduction

The kynurenine pathway is responsible for approximately 95% of the metabolism of the essential amino acid tryptophan (Figure 1) (1). While the majority (90%) of physiological tryptophan degradation through this pathway occurs in the liver, the extrahepatic kynurenine pathway plays a relatively greater role in states

of immune activation (2). The main roles of the kynurenine pathway include the regulation of tryptophan availability for the synthesis of serotonin in the central nervous system, and the synthesis of heme, nicotinic acid, oxidized nicotinamide adenine dinucleotide (phosphate), $NAD^+(P^+)$, and its reduced form NAD (P)H (3). There has been an increasing interest, particularly over the last three decades, in the biological and pathophysiological role of



the kynurenine pathway. Several studies have reported that a) local and systemic alterations in kynurenine metabolites, e.g., tryptophan and kynurenine, resulting from changes in enzyme expression and/ or activity suggest the presence of conditions such as cancer, immune diseases, and neuropsychiatric disorders, and b) individual kynurenine metabolites, e.g., kynurenic acid, 3hydroxyanthranilic acid, 3-hydroxykynurenine, and quinolinic acid, can directly influence various processes, including the redox state, immune function, glutamate neurotransmission in the central nervous system, and carbohydrate metabolism (3–8).

The interplay between the kynurenine pathway, inflammation, and immunity has also stimulated research on the pathophysiological and clinical role of kynurenine metabolites in patients with rheumatic diseases (RD). RD is an umbrella term that includes various chronic systemic conditions affecting the musculoskeletal system with a predominantly autoimmune (e.g., systemic lupus erythematosus, SLE, rheumatoid arthritis, RA, Sjogren's syndrome, SSj, systemic sclerosis, SSc, and progressive systemic sclerosis, pSS), mixedautoimmune-autoinflammatory (e.g., psoriatic arthritis, PsA, ankylosing spondylitis, AS, axial spondylarthritis, axSpA, and Behcet's disease, BD), or autoinflammatory component (e.g., familial Mediterranean fever, FMF) (9-11). Whilst significant progress has been made in the diagnosis and treatment of clinically overt RD, significant challenges remain with the identification of early forms of disease, which supports the search for novel RD biomarkers (12-14).

We sought to address these issues by conducting a systematic review and meta-analysis of published studies investigating the plasma or serum concentrations of metabolites within the kynurenine pathway (Figure 1) in patients with RD and healthy controls. We further investigated the presence of associations between the effect size of between-group differences in individual metabolites and a range of study and patient characteristics, particularly C-reactive protein, type of RD (autoimmune, mixed autoimmune-autoinflammatory, or autoinflammatory disease), and use of disease-modifying anti-rheumatic drugs (DMARDs) and corticosteroids.

Materials and methods

Search strategy and study selection

We systematically searched the electronic databases PubMed, Web of Science, and Scopus for relevant articles published from inception to the 30th of June 2023 using the following terms and their combination: "rheumatic diseases" OR "rheumatoid arthritis" OR "psoriatic arthritis" OR "ankylosing spondylitis" OR "systemic lupus erythematosus" OR "systemic sclerosis" OR "Sjogren's syndrome" OR "connective tissue diseases" OR "vasculitis" OR "Behçet's disease" AND "tryptophan" OR "kynuren*" OR "anthranil*" OR "xanthurenic" OR "cinnabar*" OR "picolinic" OR "quinolinic". Two investigators independently screened each abstract and, if relevant, the full-text articles according to the following inclusion criteria: (i) assessment of tryptophan catabolites and/or their ratio in plasma or serum, (ii) comparison of participants with RD and healthy controls aged \geq 18 years in case-control studies, and (iii) full-text articles available in English language. The references of each article were also searched for additional studies. A third investigator was involved in case of disagreement.

Two investigators independently extracted the following information from the selected manuscripts and transferred them to an electronic spreadsheet for further analysis: first author, year of publication, continent where the study was conducted, number of participants, male to female ratio, tryptophan, kynurenine, kynurenine to tryptophan ratio, kynurenic acid, 3-hydroxyanthranilic acid, 3hydroxykynurenine, quinolinic acid, kynurenine acid to kynurenine ratio, quinolinic acid to kynurenine acid ratio, C-reactive protein (CRP), disease duration, type of RD (autoimmune: SLE, RA, SSj, SSc, pSS; mixed autoimmune-autoinflammatory: PsA, AS, axSpA, and BD; autoinflammatory: FMF), use of DMARDs and corticosteroids, matrix used for analysis (serum or plasma), and analytic method used for the measurement of individual metabolites.

We assessed the risk of bias with the Joanna Briggs Institute Critical Appraisal Checklist for analytical studies (15). Studies addressing \geq 75%, \geq 50% and <75%, and <50% of checklist items were considered as having low, moderate, and high risk, respectively. We also assessed the certainty of evidence using the Grades of Recommendation, Assessment, Development and Evaluation (GRADE) Working Group system (16). We complied with the Preferred Reporting Items for Systematic reviews and Meta-Analyses (PRISMA) 2020 statement (17), and registered our study in the International Prospective Register of Systematic Reviews (PROSPERO registration number: CRD CRD42023443718).

Statistical analysis

We generated forest plots of standardized mean differences (SMDs) and 95% confidence intervals (CIs) to assess for differences in serum or plasma concentrations of individual metabolites between RD patients and healthy controls (a p-value of less than 0.05 was considered statistically significant). Where necessary, means and standard deviations were extrapolated from medians and interquartile ranges or ranges, as previously reported (18, 19), or from graphs using the Graph Data Extractor software (San Diego, CA, USA). Heterogeneity of SMD across studies was assessed using the Q statistic (20, 21). The stability of the results of the meta-analysis was assessed in sensitivity analysis (22). The Begg's and Egger's tests and the "trim-and-fill" procedure were used to assess and eventually correct publication bias (23-25). Univariate meta-regression and subgroup analyses were conducted to investigate associations between the effect size and the following parameters: year of publication, study continent, number of participants, age, male to female ratio, CRP, disease duration, type of RD, use of DMARDs and corticosteroids, sample matrix (serum or plasma) and assay method used to measure analytes.

Statistical analyses were performed using Stata 14 (Stata Corp., College Station, TX, USA).

Results

Screening process

A flow chart of the screening process is described in Figure 2. We initially identified 2,156 studies, of which 2,128 were excluded after the first screening because they were either duplicates or irrelevant. After full-text revision of the remaining 28 articles, two were further excluded because of missing data, one because it was not a case-control study, and one because it was conducted in participants <18 years of age. Thus, 24 studies were selected for analysis (Table 1) (26–49). The risk of bias, presented in Supplementary Table 3, was moderate in 14 studies (26–35, 37, 41, 44, 45, 47), and low in the remaining 10 (34, 36, 38–40, 42, 43,

46, **48**, **49**). The initial certainty of evidence was low for all studies (rating 2, $\oplus \oplus \odot \odot$) given their cross-sectional design.

Tryptophan

We identified 21 studies (23 comparator groups) reporting tryptophan concentrations in a total of 1,526 RD patients (mean age 48 years, 24% males) and 1,404 healthy controls (mean age 43 years, 38% males; Table 1) (26–38, 40–42, 44, 46–49). Ten were conducted in Europe (26–29, 32–37), and 11 in Asia (30, 31, 38, 40–42, 44, 46–49). Six study groups included patients with RA (28, 31, 34, 36, 38, 41), five with pSS (29, 33, 44, 46, 47), four with SLE (27, 30, 32, 35), three with AS (38, 40), two with SSc (26, 37), and one with BD (42), FMF (48), and axSpA (49), respectively. Liquid chromatography was used in 21 study groups (27–38, 40–42, 44, 46, 48, 49), a spectrofluorimetric assay in one (26), and an enzyme-linked immunosorbent assay (ELISA) in the remaining one (47).



TABLE 1 Characteristics of the selected studies.

		Hea	lthy co	ontrols			Patients with	h rheu	imatic disea	ises
Study	n	Age (Mean or median)	M/ F	Trp KynA 3HK 3HAA (Mean <u>+</u> SD)	QuinA Kyn/Trp KynA/ Kyn QuinA/ KynA (Mean <u>+</u> SD)	n	Age (Mean or median)	M/ F	TRP KynA 3HK 3HAA (Mean <u>+</u> SD)	QuinA Kyn/Trp KynA/ Kyn QuinA/ KynA (Mean <u>+</u> SD)
Csipo I et al., 1995, Hungary (26)	46	NR	NR	63.1 ± 11.1 NR NR NR NR	NR NR NR NR	29	50	NR	38.2 ± 9.7 NR NR NR NR	NR NR NR NR
Widner B et al., 2000, Austria (27)	49	38	28/ 21	71.5 ± 12.4 1.84 ± 0.61 NR NR NR	NR 0.026 ± 0.007 NR NR	55	41	10/ 45	54.6 ± 13.6 2.53 ± 1.22 NR NR NR	NR 0.046 ± 0.021 NR NR
Schroecksnadel K et al., 2003, Austria (28)	20	50	NR	64.8 ± 12.8 1.98 ± 0.54 NR NR NR	NR 0.032 ± 0.005 NR NR	38	57	2/36	45.1 ± 7.1 1.9 ± 0.57 NR NR NR	NR 0.043 ± 0.009 NR NR
Pertovaara M et al., 2005, Finland (29)	309	45	170/ 139	79.4 ± 13.7 2.05 ± 0.48 NR NR NR	NR 0.026 ± 0.006 NR NR	103	60	7/96	74 ± 13.2 2.53 ± 1.05 NR NR NR	NR 0.034 ± 0.014 NR NR
Xiang ZY et al., 2010, China (30)	80	34	37/ 42	$\begin{array}{c} 43.4 \pm 4.6 \\ 1.54 \pm 0.36 \\ 25.2 \pm 6.5 \\ \text{NR} \\ \text{NR} \end{array}$	NR 0.037 ± 0.008 16.6 ± 3.2 NR	30	35	5/25	$30.6 \pm 4.1 \\ 3.72 \pm 0.56 \\ 60.3 \pm 5.9 \\ NR \\ NR$	NR 0.1236 ± 0.058 16.2 ± 2.7 NR
Ozkan Y et al., 2012, Turkey (31)	20	62	4/16	42.9 ± 6.6 2.71 ± 0.65 NR NR NR	NR 0.062 ± 0.011 NR NR	32	59	5/27	43.9 ± 15.6 2.56 ± 0.56 NR NR NR	NR 0.062 ± 0.027 NR NR
Lood C et al., 2015, Sweden (32)	79	48	10/ 69	54.1 ± 11.4 1.85 ± 0.61 NR NR NR	NR 0.057 ± 0.0143 NR NR	148	47	22/ 126	49.9 ± 15.2 1.85 ± 0.66 NR NR NR	NR 0.06 ± 0.0181 NR NR
Maria NI et al., 2016, The Netherlands (33)	71	52	5/66	8,201 ± 2,163 584 ± 188 NR NR NR	NR 0.074 ± 0.0215 NR NR	124	58	8/ 116	6,147 ± 1,008 613 ± 219 NR NR NR	NR 0.107 ± 0.035 NR NR
Smolenska Z et al., 2016, Poland (34)	19	41	2/17	48.2 ± 26.8 NR NR NR NR	NR NR NR NR	46	42	4/42	47.7 ± 13.3 NR NR NR NR	NR NR NR NR
Åkesson K et al., 2018, Sweden (35)	30	47	NR	61.8 ± 14 0.712 ± 0.230 NR	0.38 ± 0.15 0.012 ± 0.0048 NR NR	132	48	NR	56.6 ± 23 0.966 ± 0.530 NR	0.546 ± 0.48 0.045 ± 0.02

(Continued)

TABLE 1 Continued

		Неа	lthy co	ontrols			Patients witl	n rheu	matic disea	ises
Study	n	Age (Mean or median)	M/ F	Trp Kyn KynA 3HK 3HAA (Mean <u>+</u> SD)	QuinA Kyn/Trp KynA/ Kyn QuinA/ KynA (Mean <u>+</u> SD)	n	Age (Mean or median)	M/ F	TRP Kyn SHK 3HK 3HAA (Mean <u>+</u> SD)	QuinA Kyn/Trp KynA/ Kyn QuinA/ KynA (Mean <u>+</u> SD)
				NR NR					NR NR	NR NR
Urbaniak B et al., 2019, Poland (36)	51	32	10/ 41	50.3 ± 10.8 NR NR NR NR	NR NR NR NR	50	51	5/45	45.5 ± 10.9 NR NR NR NR	NR NR NR NR
Smolenska Z et al., 2020, Poland (37)	27	NR	7/35	40.8 ± 12.3 NR NR NR NR	NR NR NR NR	42	60	NR	32.5 ± 9.6 NR NR NR NR	NR NR NR NR
Zhou Y et al. (a) 2020, China (38)	30	47	16/ 14	54.6 ± 14.2 NR NR NR NR	NR NR NR NR	30	34	20/ 10	102.5 ± 30.8 NR NR NR NR	NR NR NR NR
Zhou Y et al. (b) 2020, China (38)	30	47	16/ 14	54.6 ± 14.2 NR NR NR NR	NR NR NR NR	32	44	14/ 18	79.5 ± 38.2 NR NR NR NR	NR NR NR NR
Anderson EW et al., 2021, USA (39)	74	36	NR	NR NR NR NR	NR 0.031 ± 0.009 NR 9.13 ± 4.44	74	38	NR	NR NR NR NR	NR 0.045 ± 0.02 NR 18.0 ± 10.9
Eryavuz Onmaz D et al. (a) 2021, Turkey (40)	50	42	27/ 23	$12,215 \pm 4,255$ 317 ± 146 6.07 ± 2.65 2.1 ± 0.81 6.34 ± 3.08	19.3 ± 6.7 0.1125 ± 0.0175 NR NR	35	40	24/ 11	$11,694 \pm 7,076$ $1,217 \pm 918$ 4.39 ± 2.04 2.62 ± 1.65 5.24 ± 2.95	23.2 ± 8.7 0.281 ± 0.230 NR NR
Eryavuz Onmaz D et al. (b) 2021, Turkey (40)	50	42	27/ 23	$\begin{array}{c} 12,215 \pm \\ 4,255 \\ 317 \pm 146 \\ 6.07 \pm 2.65 \\ 2.1 \pm 0.81 \\ 6.34 \pm 3.08 \end{array}$	19.3 ± 6.7 0.1125 ± 0.0175 NR NR	50	40	31/ 19	$\begin{array}{c} 9,165 \pm \\ 4,250 \\ 431 \pm 163 \\ 5.18 \pm 3.00 \\ 2.73 \pm 1.73 \\ 5.24 \pm 2.95 \end{array}$	17.7 ± 4.9 0.0775 ± 0.05 NR NR
Kor A et al., 2022, Turkey (41)	41	54	10/ 31	17,175 ± 18,310 707 ± 932 NR NR NR	NR 0.0333 ± 0.0195 NR NR	50	59	13/ 37	10,244 ± 9,699 669 ± 761 NR NR NR	NR 0.0628 ± 0.0726 NR NR
Eryavuz Onmaz D et al., 2022, Turkey (42)	120	41	51/ 67	$10,908 \pm 4,274 \\ 280 \pm 123 \\ 4.67 \pm 3.11$	17.7 ± 7.4 0.039 ± 0.02 NR NR	120	41	55/ 63	$9,444 \pm 4,334$ 781 ± 520 6.00 ± 3.27	27.1 ± 16.3 0.078 ± 0.0425 NR NR

(Continued)

TABLE 1 Continued

		Неа	lthy co	ontrols		Patients with rheumatic diseases					
Study	n	Age (Mean or median)	M/ F	Trp Kyn KynA 3HK 3HAA (Mean <u>+</u> SD)	QuinA Kyn/Trp KynA/ Kyn QuinA/ KynA (Mean <u>+</u> SD)	n	Age (Mean or median)	M/ F	TRP KynA 3HK 3HAA (Mean <u>+</u> SD)	QuinA Kyn/Trp KynA/ Kyn QuinA/ KynA (Mean <u>+</u> SD)	
				2.25 ± 1.21 5.53 ± 3.15					4.52 ± 2.71 6.67 ± 3.81		
Pellicano C et al., 2022, Italy (43)	20	59	2/18	NR NR 54.2 ± 14.2 NR NR	NR NR NR NR	52	57	5/47	NR NR 68.6 ± 24.8 NR NR	NR NR NR NR	
Apaydın H et al., 2022, Turkey (44)	42	54	3/39	12,804 ± 2,421 413 ± 104 NR NR NR	NR 0.0333 ± 0.0074 NR NR	34	53	2/32	10,700 ± 2,313 488 ± 165 NR NR NR	NR 0.0433 ± 0.0222 NR NR	
Jeon C et al., 2023, Republic of Korea (45)	22	33	22/0	NR 445 ± 91 NR NR NR	NR NR NR NR	87	39	87/0	NR 474 ± 147 NR NR NR	NR NR NR NR	
Eryavuz Onmaz D et al., 2023, Turkey (46)	80	51	4/76	$54 \pm 28 \\ 1.06 \pm 0.33 \\ 19.3 \pm 11.4 \\ 16.6 \pm 15.8 \\ 35.2 \pm 34.3$	$90 \pm 44 \\ 0.0204 \pm \\ 0.0107 \\ 18.3 \pm 12 \\ 5.42 \pm 3.84$	80	52	3/77	$38 \pm 26 \\ 3.49 \pm 2.97 \\ 30.3 \pm 21.7 \\ 21.3 \pm 19.3 \\ 47.2 \pm 27.4$	$\begin{array}{c} 165 \pm 61 \\ 0.0806 \pm \\ 0.0329 \\ 10.1 \pm 5.9 \\ 6.59 \pm 5.01 \end{array}$	
Park Y et al., 2023, Republic of Korea (47)	30	NR	NR	30.5 ± 3.8 19.8 ± 5 NR NR NR	3.29 ± 2.5 0.676 ± 0.202 NR NR	81	53	2/79	18 ± 4.2 17.4 ± 6.8 NR NR NR	10.8 ± 6.7 0.892 ± 0.919 NR NR	
Tezkan D et al., 2023, Turkey (48)	80	35	38/ 42	$10,032 \pm 5,107$ 249 ± 156 4.46 ± 3.34 4.03 ± 1.84 5.85 ± 3.77	10.96 ± 3.82 0.026 ± 0.14 NR NR	81	34	40/ 41	$7,673 \pm 4,400 \\ 406 \pm 288 \\ 2.60 \pm 1.96 \\ 7.39 \pm 2.84 \\ 7.17 \pm 2.89$	21.37 ± 15.78 0.0728 ± 0.0695 NR NR	
Yurt EF et al., 2023, Turkey (49)	50	42	22/ 28	10,958 ± 2,253 366 ± 100 NR NR NR	NR 0.0337 ± 0.093 NR NR	104	45	50/ 54	9,530 ± 1,869 414 ± 86 NR NR NR	NR 0.0432 ± 0.0153 NR NR	

F, female; M, male; NR, not reported; Trp, tryptophan; Kyn, kynurenine; KynA, kynurenine acid; QuinA, quinolinic acid, 3HK, 3-hydroxykynurenine; 3HAA, 3-hydroxyanthranilic acid; Kyn/ Trp, kynurenine to tryptophan ratio; KynA/Kyn, kynurenine acid to kynurenine ratio; QuinA/KynA, quinolinic acid to kynurenine acid ratio. Concentrations are expressed in a) µmol/L or ng/mL for Trp and Kyn, and b) nmol/L or ng/mL for KynA, 3HAA, 3HK, and Quin A.

Among the liquid chromatography-based assays, seven used fluorimetric detection (27–33), and the remaining 14 mass spectrometry (34–38, 40–42, 44, 46, 48, 49). Measurements were conducted in serum in 17 study groups (26–33, 36, 38, 40, 42, 46–48), and plasma in the remaining six (34, 35, 37, 41, 44, 49). The risk of bias (Supplementary Table 3) was moderate in 13 studies (26–33,

35, 37, 41, 44, 47), and low in the remaining eight (34, 36, 38, 40, 42, 46, 48, 49).

The forest plot showed that tryptophan concentrations were significantly lower in RD patients compared to controls (SMD=-0.71, 95% CI -1.03 to -0.39, p<0.001; $I^2 = 93.6\%$, p<0.001; Figure 3). The pooled SMD values were not influenced by individual studies



Forest plot of the standard mean differences in tryptophan concentrations between patients with rheumatic disease and healthy controls

(range between -0.82 and -0.61; Supplementary Figure 1). There was no publication bias (Begg's test, p=0.32; Egger's test, p=0.30). The "trim-and-fill" method did not identify any missing study to be added to the funnel plot to ensure symmetry (Supplementary Figure 2).

In meta-regression, there were non-significant associations between effect size and age (t=-1.60, p=0.13), publication year (t=1.89, p=0.07), sample size (t=0.04, p=0.97), RD duration (t=1.68, p=0.12), CRP (t=1.96, p=0.08), or use of DMARDs (t=0.72, p=0.49) and corticosteroids (t=-0.05, p=0.97). By contrast, the effect size was significantly associated with the (male/female RD patients)/(male/female controls) ratio (t=2.25, p=0.04).

In subgroup analysis, the pooled SMD was statistically significant in studies investigating pSS (SMD=-1.22, 95% CI -1.92 to -0.52, p<0.001; I² = 95.1%, p<0.001), SLE (SMD=-1.15, 95% CI -2.15 to -0.15, p=0.02; $I^2 = 96.1\%$, p<0.001), and SSc patients (SMD=1.55, 95% CI -3.10 to -0.01, p=0.049; $I^2 = 93.6\%$, p<0.001), but not RA (SMD=-0.34, 95% CI -0.99 to 0.31, p=0.31; I² = 90.1%, p<0.001), or AS patients (SMD=0.38, 95% CI -1.02 to 1.77, p=0.60; $I^2 = 96.1\%$, p<0.001; Figure 4). In addition, the pooled SMD was statistically significant in studies of patients with autoimmune disease (SMD=-0.93, 95% CI -1.92 to -0.52, p<0.001; $I^2 = 93.8\%$, p<0.001), but not mixed autoimmune-autoinflammatory disease (SMD=-0.01, 95% CI -0.69 to 0.66, p=0.97; $I^2 = 93.6\%$, p<0.001; Supplementary Figure 3). There were non-significant differences (p=0.46) in pooled SMD between European (SMD=-0.89, 95% CI -1.29 to -0.49, p<0.001; I² = 90.7%, p<0.001) and Asian studies (SMD=-0.56, 95% CI -1.07 to -0.05, p=0.03; $I^2 = 95.0\%$, p<0.001). By contrast, there was a significant difference (p=0.006) between the pooled SMD of studies using liquid chromatography (SMD=-0.53,



FIGURE 4

Forest plot of studies investigating tryptophan concentrations in patients and controls according to the type of rheumatic disease

95% CI -0.82 to -0.24, p<0.001; $I^2 = 91.6\%$, p<0.001) and other assays (SMD=-2.71, 95% CI -3.39 to -2.02, p<0.001; $I^2 = 62.6\%$, p=0.10) with a relatively lower heterogeneity in the latter subgroup. In addition, in studies using liquid chromatography the pooled SMD was statistically significant in those using fluorimetric detection (SMD=-1.15, 95% CI -1.77 to -0.53, p<0.001; $I^2 = 94.8\%$, p<0.001) but not in those using mass spectrometry detection (SMD=-0.24, 95% CI -0.54 to 0.05, p=0.10; $I^2 = 86.6\%$, p<0.001), and the difference between the effect sizes was also statistically significant (p=0.03). Finally, there were nonsignificant differences (p=0.63) in pooled SMD between studies investigating plasma (SMD=-0.53, 95% CI -0.78 to -0.28, p<0.001; $I^2 = 48.1\%$, p=0.09) and those investigating serum (SMD=-0.78, 95% CI -1.20 to -0.36, p<0.001; $I^2 = 93.6\%$, p<0.001), with a lower between-study variance in the former subgroup.

The certainty of evidence remained low (rating 2, $\bigoplus \bigoplus \bigoplus \bigoplus)$ after considering the low-moderate risk of bias in all studies (no rating change), the high but partly explainable heterogeneity (no rating change), the lack of indirectness (no rating change), the relatively low imprecision (confidence intervals not crossing the threshold, no rating change), the moderate effect size (SMD=-0.71, no rating change) (50), and the absence of publication bias (no rating change).

Kynurenine

Seventeen studies (18 comparator groups) investigated kynurenine concentrations in a total of 1,384 RD patients (mean age 48 years, 29% males) and 1,223 healthy controls (mean age 44 years, 40% males; Table 1) (27–33, 35, 40–42, 44–49). Eleven studies were conducted in Asia (30, 31, 40–42, 44–49), and six in Europe (27–29, 32, 33, 35). Five study groups included patients with pSS (29, 33, 44, 46, 47), four with SLE (27, 30, 32, 35), three with AS (40, 45), three with RA (28, 31, 41), and one with BD (42), FMF (48), and axSpA (49), respectively. Liquid chromatography was used in

15 study groups (27–33, 40–42, 44, 46, 48, 49), ELISA in two (45, 47), and gas chromatography in one (35). In liquid chromatography studies, seven used mass spectrometry (40–42, 44, 46, 48, 49), six ultraviolet (27–29, 31–33), and one fluorimetric detection (30). Kynurenine was measured in serum in 14 study groups (27–33, 40, 42, 45–48), and plasma in the remaining four (35, 41, 44, 49). The risk of bias (Supplementary Table 3) was moderate in 12 studies (27–33, 35, 41, 44, 45, 47), and low in the remaining five (40, 42, 46, 48, 49).

The forest plot showed that kynurenine concentrations were significantly higher in RD patients compared to healthy controls (SMD=0.69, 95% CI 0.35 to 1.02, p<0.001; $I^2 = 93.2\%$, p<0.001; Figure 5). Sensitivity analysis showed stability of the results as the effect size ranged between 0.48 and 0.75 (Supplementary Figure 4). There was no publication bias (Begg's test, p=0.88; Egger's test, p=0.46). Although the "trim-and-fill" method failed to identify missing studies to be added to the funnel plot to ensure symmetry (Supplementary Figure 5), it highlighted the distortive effects of the study of Xiang et al. (30). The removal of this study from the analysis attenuated the effect size (SMD=0.48, 95% CI 0.23 to 0.73, p=0.001; $I^2 = 87.3\%$, p<0.001).

In meta-regression, there were non-significant associations between effect size and age (t=-0.53, p=0.61), sex (t=-0.95, p=0.36), publication year (t=-0.44, p=0.67), sample size (t=0.11, p=0.91), CRP (t=-1.68, p=0.14), RD duration (t=0.23, p=0.82), or use of DMARDs (t=-1.17, p=0.27) and corticosteroids (t=0.23, p=0.82).

In subgroup analysis, the pooled SMD was statistically significant in studies of SLE (SMD=1.54, 95% CI 0.13 to 2.95, p=0.03; $I^2 = 97.9\%$, p<0.001), and AS (SMD=0.81, 95% CI 0.12 to 1.51, p=0.02; $I^2 = 85.9\%$, p<0.001), but not RA (SMD=-0.12, 95% CI -0.41 to 0.16, p=0.39; $I^2 = 0.0\%$, p=0.84), or pSS (SMD=0.45, 95% CI -0.02 to 0.92, p=0.06; $I^2 = 90.1\%$, p<0.001; Figure 6), with a virtually absent heterogeneity in the RA subgroup. There were non-significant differences (p=0.82) in pooled SMD between studies in patients with autoimmune disease (SMD=0.62, 95% CI 0.16 to 1.09,

Study Name	Year		SMD (95% CI)	Patients N, mean (SD)	CTRL N, mean (SD)	% Weight
Widner B et al.	2000		0.70 (0.31, 1.10)	55, 2.53 (1.22)	49, 1.84 (.61)	5.62
Schroecksnadel K et al.	2003	_ .	-0.14 (-0.68, 0.40)	38, 1.9 (.57)	20, 1.98 (.54)	5.26
Pertovaara M et al.	2005	∔	0.72 (0.49, 0.95)	103, 2.53 (1.05)	309, 2.05 (.48)	5.93
Xiang ZY et al.	2010		• 5.15 (4.35, 5.96)	30, 3.72 (.56)	80, 1.54 (.36)	4.52
Ozkan Y et al.	2012		-0.25 (-0.81, 0.31)	32, 2.56 (.56)	20, 2.71 (.65)	5.21
Lood C et al.	2015	+ :	0.00 (-0.27, 0.27)	148, 1.85 (.66)	79, 1.85 (.61)	5.86
Maria NI et al.	2016	- - -	0.14 (-0.15, 0.43)	124, 613 (219)	71, 584 (188)	5.83
Åkesson K et al.	2018		0.52 (0.12, 0.92)	132, .966 (.53)	30, .712 (.23)	5.61
Eryavuz Onmaz D et al. (a)	2021		1.50 (1.02, 1.99)	35, 1217 (918)	50, 317 (146)	5.40
Eryavuz Onmaz D et al. (b)	2021	•	0.74 (0.33, 1.14)	50, 431 (163)	50, 317 (146)	5.60
Kor A et al.	2022	-	-0.05 (-0.46, 0.37)	50, 669 (761)	41, 707 (932)	5.58
Eryavuz Onmaz D et al.	2022	-	1.33 (1.05, 1.61)	120, 781 (520)	120, 280 (123)	5.85
Apaydın H et al.	2023	- _	0.56 (0.10, 1.02)	34, 488 (165)	42, 413 (104)	5.47
Jeon C et al.	2023		0.21 (-0.26, 0.68)	87, 474 (147)	22, 445 (91)	5.45
Eryavuz Onmaz D et al.	2023		1.15 (0.82, 1.48)	80, 3.49 (2.97)	80, 1.06 (.33)	5.75
Park Y et al.	2023		-0.38 (-0.80, 0.05)	81, 17.4 (6.8)	30, 19.8 (5)	5.56
Tezkan D et al.	2023		0.68 (0.36, 0.99)	81, 406 (288)	80, 249 (156)	5.78
Yurt EF et al.	2023		0.53 (0.19, 0.87)	104, 414 (86)	50, 366 (100)	5.73
	= 0.000)		0.69 (0.35, 1.02)	1384	1223	100.00

FIGURE 5

Forest plot of the standard mean differences in kynurenine concentrations between patients with rheumatic disease and healthy controls.



Forest plot of studies investigating kynurenine concentrations in patients and controls according to type of rheumatic disease.

p=0.009; $I^2 = 94.5\%$, p<0.001) and mixed autoimmuneautoinflammatory disease (SMD=0.86, 95% CI 0.41 to 1.32, p<0.001; $I^2 = 85.8\%$, p<0.001; Supplementary Figure 6). Furthermore, there were non-significant differences (p=0.34) in pooled SMD between Asian (SMD=0.89, 95% CI 0.38 to 1.40, p=0.001; $I^2 = 94.7\%$, p<0.001) and European studies (SMD=0.34, 95% CI 0.04 to 0.64, p=0.03; $I^2 = 80.0\%$, p<0.001). By contrast, the pooled SMD was statistically significant in studies using liquid chromatography (SMD=0.80, 95% CI 0.42 to 1.18, p<0.001; $I^2 = 93.8\%$, p<0.001) but not in those using other methods (SMD=0.12, 95% CI -0.42 to 0.65, p=0.66; $I^2 = 78.4\%$, p=0.010). In studies using liquid chromatography, the pooled SMD was significantly different with mass spectrometry (SMD=0.81, 95% CI 0.47 to 1.14, p<0.001; $I^2 = 84.6\%$, p<0.001) but not ultraviolet detection (SMD=0.23, 95% CI -0.11 to 0.56, p=0.19; I² = 82.6%, p<0.001), and the difference between the effect sizes was statistically significant (p=0.038). Finally, there were non-significant differences (p=0.56) in pooled SMD between studies in plasma (SMD=0.39, 95% CI 0.11 to 0.67, p=0.006; $I^2 = 48.2\%$, p=0.12) and serum (SMD=0.78, 95% CI 0.36 to 1.20, p<0.001; I² = 94.6%, p<0.001), with a lower between-study variance in the plasma subgroup.

The certainty of evidence remained low (rating 2, $\bigoplus \bigoplus \bigoplus \bigoplus)$ after considering the low-moderate risk of bias in all studies (no rating change), the high but partly explainable heterogeneity (no rating change), the lack of indirectness (no rating change), the relatively low imprecision (confidence intervals not crossing the threshold, no rating change), the moderate effect size (SMD=0.69, no rating change) (50), and the absence of publication bias (no rating change).

Kynurenine to tryptophan ratio

Seventeen studies (18 comparator groups) investigated the kynurenine to tryptophan ratio in a total of 1,371 RD patients (mean age 48 years, 24% males; Table 1) and 1,275 healthy controls

(mean age 44 years, 39% males) (27-33, 35, 39-42, 44, 46-49). Ten studies were conducted in Asia (30, 31, 40-42, 44, 46-49), six in Europe (27-29, 32, 33, 35), and one in America (39). Five study groups included participants with pSS (29, 33, 44, 46, 47), five with SLE (27, 30, 32, 35, 39), three with RA (28, 31, 41), two with AS (40), and one with BD (42), FMF (48), and axSpA (49), respectively. Liquid chromatography was used in 16 study groups (27-33, 39-42, 44, 46, 48, 49), whereas ELISA (47) and gas chromatography (35) were used in the remaining two. Among the studies using liquid chromatography, the detection method included mass spectrometry in nine (39-42, 44, 46, 48, 49), ultraviolet in four (27-29, 32), and fluorimetric detection in the remaining three (30, 31, 33). Serum was the biological matrix in 14 study groups (27-33, 39, 40, 42, 46-48), and plasma in the remaining four (35, 41, 44, 49). The risk of bias was moderate in 11 studies (27-33, 35, 41, 44, 47) and low in the remaining six (39, 40, 42, 46, 48, 49) (Supplementary Table 3).

The forest plot showed that the kynurenine to tryptophan ratio was significantly higher in RD patients compared to controls (SMD=0.88, 95% CI 0.55 to 1.21, p<0.001; $I^2 = 92.9\%$, p<0.001; Figure 7). The effect size ranged between 0.77 and 0.98 in sensitivity analysis, highlighting the stability of the results (Supplementary Figure 7). There was no publication bias (Begg's test, p=1.0; Egger's test, p=0.68). Accordingly, the "trim-and-fill" method did not identify any missing study to be added to the funnel plot to ensure symmetry (Supplementary Figure 8).

No significant associations were observed in meta-regression between effect size and age (t=0.83, p=0.42), sex (t=-1.62, p=0.13), publication year (t=-0.91, p=0.37), sample size (t=0.28, p=0.78), RD duration (t=-0.10, p=0.93), CRP (t=-1.49, p=0.18), or use of DMARDs (t=0.70, p=0.51) and corticosteroids (t=0.19, p=0.85).

In subgroup analysis, the pooled SMD was statistically significant in studies of SLE (SMD=1.10, 95% CI 0.37 to 1.84, p=0.003; $I^2 = 94.7\%$, p<0.001) and pSS (SMD=1.07, 95% CI 0.45 to 1.69, p=0.001; $I^2 = 93.6\%$, p<0.001), but not RA (SMD=-0.63, 95% CI -0.09 to 1.35, p=0.08; $I^2 = 82.3\%$, p<0.001) or AS (SMD=0.10,

Study Name	Year		SMD (95% CI)	N, mean (SD); Treatment	N, mean (SD); Control	% Weight
Widner B et al.	2000		1.25 (0.83, 1.67)	55, .046 (.021)	49, .026 (.007)	5.53
Schroecksnadel K et al.	2003	↓ <u>+ • </u>	1.40 (0.80, 2.00)	38, .043 (.009)	20, .032 (.005)	5.05
Pertovaara M et al.	2005		0.92 (0.69, 1.15)	103, .034 (.014)	309, .026 (.006)	5.91
Xiang ZY et al.	2010		2.81 (2.25, 3.37)	30, .124 (.058)	80, .037 (.008)	5.16
Ozkan Y et al.	2012	 + !	0.00 (-0.56, 0.56)	32, .062 (.027)	20, .062 (.011)	5.17
Lood C et al.	2015	+•- :	0.18 (-0.10, 0.45)	148, .06 (.0181)	79, .057 (.0143)	5.84
Maria NI et al.	2016		1.07 (0.76, 1.38)	124, .107 (.035)	71, .074 (.0215)	5.77
Åkesson K et al.	2018	;	0.51 (0.11, 0.91)	132, .019 (.015)	30, .012 (.0048)	5.58
Anderson EW et al.	2021		0.90 (0.56, 1.24)	74, .045 (.02)	74, .031 (.009)	5.71
Eryavuz Onmaz D et al. (a)	2021		1.14 (0.67, 1.61)	35, .281 (.23)	50, .112 (.0175)	5.42
Eryavuz Onmaz D et al. (b)	2021		-0.93 (-1.35, -0.52)	50, .0775 (.05)	50, .112 (.0175)	5.55
Kor A et al.	2022	i	0.53 (0.11, 0.95)	50, .0628 (.0726)	41, .0333 (.0195)	5.53
Eryavuz Onmaz D et al.	2022		1.17 (0.90, 1.45)	120, .078 (.0425)	120, .039 (.02)	5.84
Apaydın H et al.	2023	.	0.63 (0.17, 1.10)	34, .0433 (.0222)	42, .0333 (.0074)	5.42
Eryavuz Onmaz D et al.	2023		2.46 (2.05, 2.87)	80, .0806 (.0329)	80, .0204 (.0107)	5.55
Park Y et al.	2023	+•	0.27 (-0.15, 0.69)	81, .892 (.919)	30, .676 (.202)	5.53
Tezkan D et al.	2023		0.93 (0.61, 1.26)	81, .0728 (.0695)	80, .026 (.014)	5.74
Yurt EF et al.	2023		0.70 (0.35, 1.04)	104, .0432 (.0153)	50, .0337 (.0093)	5.70
Overall (I-squared = 92.9%, J	o = 0.000)		0.88 (0.55, 1.21)	1371	1275	100.00
NOTE: Weights are from rand	lom effects analysis					

FIGURE 7

Forest plot of studies reporting the kynurenine/tryptophan ratio in patients with rheumatic disease and healthy controls.

95% CI -0.93 to 2.13, p=0.92; $I^2 = 97.7\%$, p<0.001, Figure 8). In addition, the pooled SMD was statistically significant in studies of patients with autoimmune disease (SMD=0.98, 95% CI 0.60 to 1.37, p<0.001; $I^2 = 92.5\%$, p<0.001) but not of patients with autoimmune-autoinflammatory diseases (SMD=0.52, 95% CI -0.39 to 1.43, p=0.26; $I^2 = 96.0\%$, p<0.001; Supplementary Figure 9). There were non-significant differences (p=0.99) in pooled SMD between European (SMD=0.86, 95% CI 0.50 to 1.22, p<0.001; $I^2 = 85.3\%$, p<0.001) and Asian studies (SMD=0.88, 95% CI 0.33 to 1.43, p=0.002; $I^2 = 95.1\%$, p<0.001). Similarly, the pooled SMD was non-significantly different (0.40) between studies using liquid chromatography (SMD=0.94, 95% CI 0.58 to 1.30, p<0.001; $I^2 = 93.4\%$, p<0.001) and other methods (SMD=0.40, 95% CI 0.11 to 0.69, p=0.007; $I^2 = 0.0\%$, p=0.42), with a virtually absent heterogeneity in the latter subgroup. In liquid chromatography

studies, the pooled SMD was statistically significant in studies using mass spectrometry (SMD=0.84, 95% CI 0.32 to 1.36, p=0.002; $I^2 = 94.3\%$, p<0.001) and ultraviolet detection (SMD=0.90, 95% CI 0.37 to 1.43, p=0.001; $I^2 = 89.5\%$, p<0.001), but not in those using fluorimetric detection (SMD=1.29, 95% CI -0.06 to 2.64, p=0.06; $I^2 = 96.0\%$, p<0.001). Finally, there were non-significant differences (p=0.45) in pooled SMD between studies analysing plasma (SMD=0.96, 95% CI 0.55 to 1.37, p<0.001; $I^2 = 94.4\%$, p<0.001) and those on serum (SMD=0.60, 95% CI 0.40 to 0.80, p<0.001; $I^2 = 0.0\%$, p=0.90), with a virtually absent between study-variance in the serum subgroup.

The certainty of evidence was upgraded to moderate (rating 3, $\oplus \oplus \oplus \odot$) after considering the low-moderate risk of bias in all studies (no rating change), the high but partly explainable heterogeneity (no rating change), the lack of indirectness (no

Study Name	Year			SMD (95% CI)	Patients N, mean (SD)	CTRL N, mean (SD)	% Weight
RA Schroecksnadel K et al. Ozkan Y et al. Kor A et al. Subtotal (I-squared = 82.3%, p	2003 2012 2022 p = 0.004)			0.00 (- 0.53 (0	0.80, 2.00) 0.56, 0.56) 0.11, 0.95) 0.09, 1.35)	38, .043 (.009) 32, .062 (.027) 50, .0628 (.0726) 120	20, .032 (.005) 20, .062 (.011) 41, .0333 (.0195) 81	5.05 5.17 5.53 15.76
Pertovaara M et al. Maria NI et al. Apaydin H et al. Eryavuz Onmaz D et al. Park Y et al. Subtotal (I-squared = 93.6%, p	2005 2016 2023 2023 2023 2023 2023 2023	-	-++ +	1.07 (0 0.63 (0 2.46 (2 0.27 (-	0.69, 1.15) 0.76, 1.38) 0.17, 1.10) 0.05, 2.87) 0.15, 0.69) 0.45, 1.69)	103, .034 (.014) 124, .107 (.035) 34, .0433 (.0222) 80, .0806 (.0329) 81, .892 (.919) 422	309, .026 (.006) 71, .074 (.0215) 42, .0333 (.0074) 80, .0204 (.0107) 30, .676 (.202) 532	5.91 5.77 5.42 5.55 5.53 28.18
SLE Widner B et al. Xiang ZY et al. Lood C et al. Anderson EW et al. Subtotal (I-squared = 94.7%, p	2000 2010 2015 2018 2021 0 = 0.000)	+		 2.81 (2 0.18 (- 0.51 (0 0.90 (0 	0.83, 1.67) 2.25, 3.37) 0.10, 0.45) 0.11, 0.91) 0.56, 1.24) 0.37, 1.84)	55, .046 (.021) 30, .124 (.058) 148, .06 (.0181) 132, .019 (.015) 74, .045 (.02) 439	49, .026 (.007) 80, .037 (.008) 79, .057 (.0143) 30, .012 (.0048) 74, .031 (.009) 312	5.53 5.16 5.84 5.58 5.71 27.82
AS Eryavuz Onmaz D et al. (a) Eryavuz Onmaz D et al. (b) Subtotal (I-squared = 97.7%, p	2021 2021 p = 0.000)	-	*	-0.93 ().67, 1.61) -1.35, -0.52) 1.93, 2.13)	35, .281 (.23) 50, .0775 (.05) 85	50, .112 (.0175) 50, .112 (.0175) 100	5.42 5.55 10.97
Eryavuz Onmaz D et al. Tezkan D et al. Yurt EF et al. Subtotal (I-squared = 56.5%, p	2022 2023 2023 5 = 0.100)		++++++++++++++++++++++++++++++++++++	0.93 (0 0.70 (0	0.90, 1.45) 0.61, 1.26) 0.35, 1.04) 0.67, 1.22)	120, .078 (.0425) 81, .0728 (.0695) 104, .0432 (.0153) 305	120, .039 (.02) 80, .026 (.014) 50, .0337 (.0093) 250	5.84 5.74 5.70 17.28
Overall (I-squared = 92.9%, p <u>NOTE: Weights are from rando</u>			-	0.88 (0	0.55, 1.21)	1371	1275	100.00

FIGURE 8

Forest plot of studies reporting the kynurenine/tryptophan ratio in patients and controls according to the type of rheumatic disease.

rating change), the relatively low imprecision (confidence intervals not crossing the threshold, no rating change), the large effect size (SMD=0.88, upgrade one level) (50), and the absence of publication bias (no rating change).

Kynurenic acid

Six studies (seven comparator groups) reported kynurenic acid concentrations in a total of 448 RD patients (mean age 43 years, 37% males) and 480 healthy controls (mean age 41 years, 39% males; Table 1) (30, 40, 42, 43, 46, 48). Five studies were conducted in Asia (30, 40, 42, 46, 48), and the remaining one in Europe (43). Two study groups included individuals with AS (40), and one with SLE (30), SSc (43), pSS (46), BD (42), and FMF (48), respectively. Liquid chromatography was used in six study groups (30, 40, 42, 46, 48), and ELISA in the remaining one (43). In liquid chromatography studies, five study groups using mass spectrometry for detection (40, 42, 46, 48), and the remaining one fluorimetry (30). All studies investigated serum. The risk of bias (Supplementary Table 3)was moderate in one study (30), and low in the remaining five (40, 42, 43, 46, 48).

The forest plot showed that kynurenic acid concentrations were not significantly different between RD patients and controls (SMD=0.72, 95% CI -0.14 to 1.59, p=0.10; $I^2 = 96.7\%$, p<0.001; Figure 9). Sensitivity analysis showed that the study of Xian et al. had a significant effect on the corresponding pooled SMD direction (30) (Supplementary Figure 10). Its removal reduced the effect size (SMD=0.00, 95% CI -0.50 to 0.50, p=1.00).

The limited number of studies prevented the assessment of publication bias and the conduct of meta-regression and subgroup analyses.

The certainty of evidence was downgraded to extremely low (rating $0, \bigcirc \bigcirc \bigcirc \bigcirc$) after considering the low-moderate risk of bias in all studies (no rating change), the high and unexplainable heterogeneity (downgrade one level), the lack of indirectness (no

rating change), and the lack of assessment of publication bias (downgrade one level).

3-hydroxyanthranilic acid

Four studies (five comparator groups), all conducted in Asia, reported 3-hydroxyanthranilic acid concentrations in a total of 366 patients with RD (mean age 42 years, 51% males) and 380 healthy controls (mean age 42 years, 39% males; Table 1) (40, 42, 46, 48). Two study groups included individuals with AS (40), and one with pSS (46), BD (42), and FMF (48), respectively. In all studies, measurements were conducted in serum using liquid chromatography with mass spectrometry detection. The risk of bias (Supplementary Figure 3) was low in all studies (40, 42, 46, 48).

The forest plot showed the absence of significant between-group differences in 3-hydroxyanthranilic acid concentrations (SMD=0.06, 95% CI -0.30 to 0.42, p=0.73; $I^2 = 83.0\%$, p<0.001; Figure 10). The corresponding SMD values were stable in sensitivity analysis (range between -0.03 and 0.22; Supplementary Figure 11).

The limited number of studies prevented the assessment of publication bias and the conduct of meta-regression and subgroup analyses.

The certainty of evidence was downgraded to extremely low (rating $0, \Theta \Theta \Theta \Theta$) after considering the low-moderate risk of bias in all studies (no rating change), the high and unexplainable heterogeneity (downgrade one level), the lack of indirectness (no rating change), and the lack of assessment of publication bias (downgrade one level).

3-hydroxykynurenine

Four studies (five comparator groups) reported 3hydroxykynurenine concentrations in a total of 366 RD patients (mean age 42 years, 51% males) and 380 healthy controls (mean age

Study					Patients	CTRL	%
Name	Year			SMD (95% CI)	N, mean (SD)	N, mean (SD)	Weight
Xiang ZY et al.	2010		_•_	5.53 (4.68, 6.38)	30, 60.3 (5.9)	80, 25.2 (6.5)	13.05
Eryavuz Onmaz D et al. (a)	2021			-0.69 (-1.14, -0.25)	35, 4.39 (2.04)	50, 6.07 (2.65)	14.36
Eryavuz Onmaz D et al. (b)	2021	-•		-0.31 (-0.71, 0.08)	50, 5.18 (3)	50, 6.07 (2.65)	14.47
Eryavuz Onmaz D et al.	2022		*	0.42 (0.16, 0.67)	120, 6 (3.27)	120, 4.67 (3.11)	14.73
Pellicano C et al.	2022			0.64 (0.12, 1.17)	52, 68.6 (24.8)	20, 54.2 (14.2)	14.14
Eryavuz Onmaz D et al.	2023		-	0.63 (0.32, 0.95)	80, 30.3 (21.7)	80, 19.3 (11.4)	14.63
Tezkan D et al.	2023	+		-0.68 (-1.00, -0.36)	81, 2.6 (1.96)	80, 4.46 (3.34)	14.63
Overall (I-squared = 97.2%,	p = 0.000)		\bigcirc	0.72 (-0.14, 1.59)	448	480	100.00
NOTE: Weights are from rar	ndom effects analysis						

FIGURE 9

Forest plot of studies reporting kynurenic acid concentrations in patients with rheumatic disease and healthy controls.



42 years, 39% males; Table 1) (40, 42, 46, 48). Two study groups included individuals with AS (40), and one with pSS (46), BD (42), and FMF (48), respectively. In all studies, measurements were conducted in serum using liquid chromatography with mass spectrometry detection. The risk of bias (Supplementary Table 3) was low in all studies (40, 42, 46, 48).

The forest plot showed that RD patients had significantly higher 3-hydroxykynurenine concentrations compared to healthy controls (SMD=0.74, 95% CI 0.30 to 1.18, p=0.001; $I^2 = 87.7\%$, p<0.001; Figure 11). The effect size was stable in sensitivity analysis, with a range between 0.57 and 0.86 (Supplementary Figure 12).

The limited number of studies prevented the assessment of publication bias and the conduct of meta-regression and subgroup analyses.

The certainty of evidence was downgraded to extremely low (rating $0, \bigcirc \bigcirc \bigcirc \bigcirc \bigcirc$) after considering the low-moderate risk of bias in all studies (no rating change), the high and unexplainable heterogeneity (downgrade one level), the lack of indirectness (no rating change), and the lack of assessment of publication bias (downgrade one level).

Quinolinic acid

Six studies (seven comparator groups) reported the concentrations of quinolinic acid in a total of 579 RD patients (mean age 45 years, 35% males) and 440 controls (mean age 42 years, 39% males; Table 1) (35, 40, 42, 46–48). Five studies were conducted in Asia (40, 42, 46–48), and the remaining one in Europe (35). Two study groups included individuals with AS (40), two with pSS (46, 47), and one with SLE (35), one BD (42), and FMF (48), respectively. The method used was liquid chromatography with mass spectrometry in five study groups (40, 42, 46, 48), gas chromatography with mass spectrometry in one (35), and ELISA in the remaining one (47). All studies assessed serum, barring one which assessed plasma (35). The risk of bias (Supplementary Table 3) was moderate in two studies (35, 47), and low in the remaining four (40, 42, 46, 48).

The forest plot showed that the concentrations of quinolinic acid were significantly higher in RD patients compared to healthy controls (SMD=0.71, 95% CI 0.31 to 1.11, p<0.001; $I^2 = 88.1\%$, p<0.001; Figure 12). Sensitivity analysis showed that the

Study					Patients	CTRL	%
Name	Year			SMD (95% CI)	N, mean (SD)	N, mean (SD)	Weight
Eryavuz Onmaz D et al. (a)	2021			0.42 (-0.01, 0.86)	35, 2.62 (1.65)	50, 2.1 (.81)	18.74
Eryavuz Onmaz D et al. (b)	2021		1 1 1	0.47 (0.07, 0.86)	50, 2.73 (1.73)	50, 2.1 (.81)	19.36
Eryavuz Onmaz D et al.	2022			1.08 (0.81, 1.35)	120, 4.52 (2.71)	120, 2.25 (1.21)	21.15
Eryavuz Onmaz D et al.	2023			0.27 (-0.04, 0.58)	80, 21.3 (19.3)	80, 16.6 (15.8)	20.62
Tezkan D et al.	2023			> 1.40 (1.06, 1.75)	81, 7.39 (2.84)	80, 4.03 (1.84)	20.14
Overall (I-squared = 87.7%, p	= 0.000)			0.74 (0.30, 1.18)	366	380	100.00
NOTE: Weights are from rando	om effects analysis		- - - - - - - - - - - - - - - - - - -				

Forest plot of studies reporting 3-hydroxykynurenine concentrations in patients with rheumatic disease and healthy controls.



corresponding pooled SMD values were not influenced by sequentially removing individual studies (effect size range between 0.59 and 0.87; Supplementary Figure 13).

The limited number of studies prevented the assessment of publication bias and the conduct of meta-regression and subgroup analyses.

The certainty of evidence was downgraded to extremely low (rating $0, \Theta \Theta \Theta \Theta$) after considering the low-moderate risk of bias in all studies (no rating change), the high and unexplainable heterogeneity (downgrade one level), the lack of indirectness (no rating change), and the lack of assessment of publication bias (downgrade one level).

Kynurenic acid to kynurenine ratio

Two Asian studies reported the kynurenine acid to kynurenine ratio in a total of 110 RD patients (mean age 43 years, 7% males) and 160 healthy controls (mean age 47 years, 26% males; Table 1)

(30, 46). One study included individuals with SLE (30), and one with pSS (46). Liquid chromatography was used in both studies, one with mass spectrometry detection (46), and the other with fluorimetric detection (30). Serum was analysed in both studies. The risk of bias (Supplementary Table 3) was moderate in one study (30) and low in the other (46).

The forest plot showed that the kynurenine acid to kynurenine ratio was non-significantly different between RD patients and controls (SMD=-0.51, 95% CI -1.23 to 0.21, p=0.17; $I^2 = 86.5\%$, p=0.006; Figure 13).

The limited number of studies prevented sensitivity analysis, the assessment of publication bias, and the conduct of meta-regression and subgroup analyses.

The certainty of evidence was downgraded to extremely low (rating $0, \Theta \Theta \Theta \Theta$) after considering the low-moderate risk of bias in all studies (no rating change), the high and unexplainable heterogeneity (downgrade one level), the lack of indirectness (no rating change), and the lack of assessment of publication bias (downgrade one level).



Quinolinic acid to kynurenine acid ratio

Two studies reported the quinolinic acid to kynurenine acid ratio in a total of 154 RD patients (mean age 45 years) and 154 healthy controls (mean age 44 years; Table 1) (39, 46). One study was conducted in Asia (46), and the other in America (39). One study included individuals with SLE (39), and the other with pSS (46). In both studies, serum was assessed using liquid chromatography with mass spectrometry detection and the risk of bias was low (Supplementary Figure 3) (39, 46).

The forest plot showed the absence of significant between-group differences in the quinolinic acid to kynurenine acid ratio (SMD=0.66, 95% CI -0.13 to 1.45, p=0.10, $I^2 = 91.3\%$, p=0.001; Figure 14).

The limited number of studies prevented sensitivity analysis, the assessment of publication, and the conduct of meta-regression and subgroup analyses.

The certainty of evidence was downgraded to extremely low (rating $0, \Theta \Theta \Theta \Theta$) after considering the low-moderate risk of bias in all studies (no rating change), the high and unexplainable heterogeneity (downgrade one level), the lack of indirectness (no rating change), and the lack of assessment of publication bias (downgrade one level).

Discussion

In this systematic review and meta-analysis, we have observed significant alterations in metabolites within the kynurenine pathway of tryptophan metabolism in patients with RD. Specifically, RD was associated with significantly lower tryptophan concentrations and higher kynurenine, kynurenine to tryptophan ratios, 3-hydroxykynurenine, and quinolinic acid concentrations. By contrast, there were non-significant betweengroup differences in kynurenic acid, 3-hydroxyanthranilic acid, kynurenic acid to kynurenine ratio, or quinolinic acid to kynurenine acid ratio. In meta-regression, the SMD of tryptophan, kynurenine, and kynurenine to tryptophan ratio were not associated with age, publication year, sample size, RD duration, C-reactive protein, or use of anti-rheumatic drugs and corticosteroids. In subgroup analysis, the SMD of tryptophan, kynurenine, and kynurenine to tryptophan ratio were statistically significant in different types of RD, barring rheumatoid arthritis. Whilst our meta-analysis on tryptophan, kynurenine, and kynurenine to tryptophan ratio included a substantial number of studies, between 17 and 21, caution is required with data interpretation in view of the relatively low certainty of evidence in our analysis, particularly when investigating other metabolites, suggesting the need for additional research to confirm or refute our findings.

The biotransformation of tryptophan to kynurenine via the intermediate N'-formylkynurenine, a critical step within the kynurenine pathway, is regulated by the enzymes tryptophan 2,3dioixygenase (TDO) in the liver and indoleamine 2,3-dioxygenase (IDO) extra-hepatically (Figure 1) (2, 3). A substantial number of studies have demonstrated a distinct regulation of these enzymes. TDO is upregulated by the substrate tryptophan, the cofactor heme, and glucocorticoids, and downregulated by the end-product NAD (P)H (2, 3, 51). By contrast, IDO can be inhibited by excess tryptophan, the endogenous messenger, nitric oxide, and antiinflammatory cytokines, e.g., interleukin-4, interleukin-10, and transforming growth factor beta, and upregulated either directly or indirectly by pro-inflammatory cytokines such as interferon gamma, interferon alpha, interleukin-1B, tumour necrosis factor alpha, and interleukin-2 (2, 3, 51-53). Therefore, the observation of a relative deficiency of tryptophan and excess of kynurenine, with a consequent increase in the kynurenine to tryptophan ratio, suggests the increased contribution of extra-hepatic IDO in tryptophan metabolism in RD, possibly mediated by IDO upregulation by pro-inflammatory cytokines, a common finding in these patients (54-56). However, the putative activation of extra-hepatic IDO could also be secondary to a relative deficiency in systemic nitric oxide, a key endogenous regulator of several biological processes such as vascular homeostasis, immune function, and neuroplasticity (57, 58). This mechanism could be relevant given the evidence of a dysregulation in nitric oxide pathways in several types of RD, and the consequent increased risk of endothelial dysfunction, atherosclerosis, and cardiovascular disease in these patients (59-66). However, it is important to emphasise that the lack of information regarding the measurement of specific cytokines and

Study				Patients	CTRL	%
Name Year			SMD (95% CI)	N, mean (SD)	N, mean (SD)	Weight
Anderson EW et al. 2021			- 1.07 (0.72, 1.41)	74, 18 (10.9)	74, 9.13 (4.44)	49.56
Eryavuz Onmaz D et al. 2023		_	0.26 (-0.05, 0.57)	80, 6.59 (5.01)	80, 5.42 (3.84)	50.44
Overall (I-squared = 91.3%, p = 0.001)			> 0.66 (-0.13, 1.45)	154	154	100.00
		\downarrow	,			
NOTE: Weights are from random effects analysis						
	0	l				

nitric oxide metabolites in the studies selected in our literature search prevented the conduct of further analyses to investigate possible associations between these molecules and the observed between-group differences in tryptophan, kynurenine, and kynurenine to tryptophan ratio. Therefore, further research is warranted to investigate this important issue.

Interestingly, in meta-regression the SMD values of tryptophan, kynurenine, and kynurenine to tryptophan ratios were not significantly associated with CRP, a conventional biomarker of systemic inflammation that is routinely used for the diagnosis and management of patients with RD (67-69). Whilst this observation suggests that alterations in tryptophan and kynurenine are not necessarily correlated with CRP elevations, further research is required to determine whether the measurement of these metabolites significantly enhances diagnostic capacity, particularly in early disease, over and above available criteria for RD (70-74). Another interesting observation, in subgroup analysis, was the different association between the reported alterations in tryptophan and kynurenine concentrations in specific types of RD. For example, the SMD of the kynurenine to tryptophan ratio was statistically significant in SLE, pSS, and autoimmune RD type, but not RA, AS, or mixed autoimmune-autoinflammatory RD type. Once again, future studies should investigate whether kynurenine metabolites are useful for the diagnosis and management of specific types of RD. This proposition is supported by the results of an elegant study reporting the utility of kynurenine pathway metabolomics in discriminating clinical subtypes in patients with multiple sclerosis and identifying those at risk of progression (75). Importantly, the absence of significant differences in the effect size between Asian and European studies downplays the potential role of ethnicity as a factor influencing the link between the kynurenine pathway and RD and confirms the results of previous studies investigating this issue (76).

Other significant RD-associated alterations in the kynurenine pathway observed in our study involved the metabolites 3hydroxykynurenine and quinolinic acid. 3-hydroxykynurenine, derived from kynurenine by kynurenine hydroxylase, has been shown to exert a bi-modal pro-oxidant and antioxidant effect in the central nervous system (77). Further studies have also highlighted the potential for this molecule to bind to α -synuclein and amyloid-beta peptides, with the potential of triggering neuroinflammatory and neurotoxic processes (78). Quinolinic acid has been extensively investigated as an N-methyl-D-aspartate receptor agonist with pro-oxidant and neurotoxic effects and as a biomarker of neurodegenerative and depressive disorders and inflammatory states (79-83). The results of our systematic review and metaanalysis extend the potential role of 3-hydroxykynurenine and quinolinic acid to the pathophysiology of RD and warrant further research to investigate their local and systemic effects in autoimmune and autoinflammatory conditions, and their role in the reported associations between RD and neuropsychiatric disorders (84-86).

An important additional issue, not investigated in our study, is related to the potential role of dietary factors as well as physical activity in modulating the complex interplay between the kynurenine pathway of tryptophan metabolism and RD. Emerging evidence underscores the role of diet in shaping the kynurenine pathway, thereby affecting the metabolism of tryptophan (87-89). A high-fat diet, for instance, has been shown to modify the flux of metabolites along the kynurenine pathway, suggesting that diet could potentially introduce a confounding element in studies comparing metabolite profiles between RD patients and healthy individuals. Additionally, recent studies have reported that downstream metabolites of kynurenine, particularly kynurenic acid, are increased in muscle biopsies of physically active adults, including after an acute endurance exercise, and associated with cardiorespiratory fitness (90-92). Therefore, future studies should also consider dietary habits and patterns of physical activity as potential factors contributing to the observed variations in the concentrations of tryptophan metabolites. Furthermore, the concept of a comorbidome, referring to the collective presence of multiple comorbid conditions in a patient, introduces another layer of complexity. For example, recent studies have reported that comorbid conditions can interact with the kynurenine pathway and potentially alter the concentrations of tryptophan metabolites (93). Considering the intricate interplay between various medical conditions and metabolic pathways, it is plausible that the presence of a distinct comorbidome might confound the association between RD and tryptophan metabolites in individual patients.

Our systematic review and meta-analysis have several strengths, including the comprehensive assessment of the kynurenine pathway in a wide range of RD types, the study of possible associations between the effect size of between-group differences and clinical and demographic characteristics and analytical methods used, and a rigorous assessment of the risk of bias and the certainty of evidence. A significant limitation is represented by the high between-study heterogeneity observed. However, we were able to identify specific sources of heterogeneity in subgroup analyses for tryptophan (analytical method and matrix used), kynurenine (type of RD and matrix used), kynurenine to tryptophan ratio (analytical method and matrix used). Furthermore, sensitivity analysis ruled out the effect of individual studies on the overall effect size.

In conclusion, this systematic review and meta-analysis has demonstrated the presence of significant alterations in metabolites within the kynurenine pathway of tryptophan metabolism in patients with RD, particularly tryptophan, kynurenine, 3hydroxykynureine, and quinolinic acid. Further studies, particularly in view of the relatively low certainty of evidence in our analysis, are required to determine the capacity of kynurenine metabolites to identify early and overt types of RD over and above existing clinical criteria and biomarkers with a view to improve the management and quality of life in this complex patient population.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding author.

Author contributions

AM: Conceptualization, Methodology, Validation, Visualization, Writing – original draft, Writing – review & editing. AZ: Conceptualization, Data curation, Investigation, Methodology, Validation, Writing – review & editing.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

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SUPPLEMENTARY FIGURE 1

Sensitivity analysis of the association between tryptophan concentrations and rheumatic disease.

SUPPLEMENTARY FIGURE 2

Funnel plot of studies investigating associations between tryptophan and rheumatic disease after "trimming-and-filling".

SUPPLEMENTARY FIGURE 3

Forest plot of studies investigating tryptophan concentrations in patients and controls according to the presence of autoimmune, mixed autoimmuneautoinflammatory, or autoinflammatory disease.

SUPPLEMENTARY FIGURE 4

Sensitivity analysis of the association between kynurenine concentrations and rheumatic disease.

SUPPLEMENTARY FIGURE 5

Funnel plot of studies investigating the association between kynurenine and rheumatic disease after "trimming-and-filling".

SUPPLEMENTARY FIGURE 6

Forest plot of studies investigating kynurenine concentrations in patients and controls according to the presence of autoimmune, mixed autoimmuneautoinflammatory, or autoinflammatory disease.

SUPPLEMENTARY FIGURE 7

Sensitivity analysis of the association between the kynurenine/tryptophan ratio and rheumatic disease.

SUPPLEMENTARY FIGURE 8

Funnel plot of studies investigating the association between the kynurenine/ tryptophan ratio and rheumatic disease after "trimming-and-filling".

SUPPLEMENTARY FIGURE 9

Forest plot of studies reporting the kynurenine/tryptophan ratio in patients and controls according to the presence of autoimmune, mixed autoimmuneautoinflammatory, or autoinflammatory diseases.

SUPPLEMENTARY FIGURE 10

Sensitivity analysis of the association between kynurenic acid concentrations and rheumatic disease.

SUPPLEMENTARY FIGURE 11

Sensitivity analysis of the association between 3-hydroxyanthranilic acid concentrations and rheumatic disease.

SUPPLEMENTARY FIGURE 12

Sensitivity analysis of the association between 3-hydroxykynurenine concentrations and rheumatic disease.

SUPPLEMENTARY FIGURE 13

Sensitivity analysis of the association between quinolinic acid concentrations and rheumatic disease.

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