

26-28 ottobre 2023

### INCREASE MALNUTRITION AWARENESS: CHALLENGE FOR THE FUTURE

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Could disease activity and protein intake affect serum amino acids concentration in adult patients with Crohn's disease?



# BACKGROUND (1)

- Crohn's disease (CD) is an immune-mediated inflammatory bowel disease (IBD) characterized by a remitting-relapsing course.
- Chronic inflammation along with malabsorption and/or inadequate dietary intake can facilitate the development of malnutrition in CD patients, impairing their quality of life.
- Among nutrients, amino acids (AAs), the building blocks of protein macromolecules, may play critical roles in the intestinal manifestations of disease, due to their involvement in many metabolic and immune functions.

# BACKGROUND (2)

- AA concentrations can vary in serum and tissues with developmental stage, diseases, nutritional state, endocrine status, and physical activity.
- Variations in AA concentrations have been observed in patients with IBD and in different other conditions such as liver diseases, heart failure, type 2 diabetes etc.
- Some AAs alterations, including reduced levels of Gln, Trp and His have been found in active IBD patients, suggesting that AA profiles might be altered in this population and might reflect disease activity as well as patients' nutritional status.

### AIMS OF THE STUDY

The present study aimed to explore serum AA concentrations in adult patients with CD, looking into their variations due to disease activity and protein content of diet. Finally, the link between AAs and inflammatory markers was also assessed.

# METHODS (1)

- This is a secondary analysis of the cross-sectional REECD (Resting Energy Expenditure evaluation in subjects with Crohn's Disease) study.
- Adult patients aged 18–65 years with diagnosis of CD were consecutively recruited from July 2016 to March 2018 at the Department of Clinical Medicine and Surgery, Federico II University Hospital, Naples, Italy.
- Exclusion criteria:
- use of corticosteroids in the last 3 months
- history of acute or chronic liver or kidney disease
- current enteral (i.e., tube feeding) or parenteral nutrition
- presence of fistulae, ileostomy, or colostomy; presence of extensive small bowel resections (residual small bowel <2 meters)
- pregnancy or lactation; unstable body weight in the last month; and unable or unwilling to give informed consent.
- Socio-demographic data, disease duration, previous surgery, smoking habits, drug treatment, location and disease behavior (according to Montreal classification) were collected.

# METHODS (2)

- Disease activity was clinically defined using the Crohn's Disease Activity Index (CDAI) score classifying patients in the active (≥150) and quiescent phases (< 150).</li>
- Blood samples were collected to analyze serum AA profiles and inflammatory markers (IL-6, II-1β, TNF-α).

Essential AAs (EAAs)	Non-essential (NEAAs)	Non-proteogenic AAs
Valine (Val)	Arginine (Arg)	Ornithine (Orn)
Isoleucine (Ile)	• Glycine (Gly)	Citrulline (Cit)
Leucine (Leu)	Alanine (Ala)	Taurine (Tau)
Methionine (Met)	Serine (Ser)	
Lysine (Lys)	Tyrosine (Tyr)	
Phenylalanine (Phe)	Cystine (Cys)	
<ul> <li>Tryptophan (Trp)</li> </ul>	Asparagine (Asn)	
Threonine (Thr)	Gutamine (GIn)	
Histidine (His)	Aspartic Acid (Asp)	
	Glutamate (Glu)	

 Participants underwent anthropometry (body weight, height and BMI) and were instructed to fill in a 3-day food record to assess their protein intake.

### **RESULTS – Demographic and clinical characteristics**

TABLE 1 Demographic and clinical characteristics of CD patients.

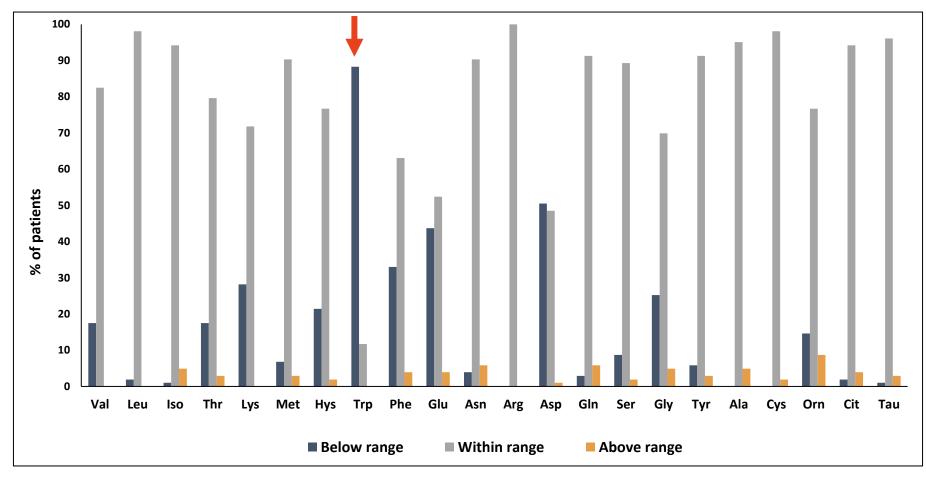
	Total	Men	Women
N, (%)	103 (100)	62 (60.2)	41 (39.8)
Age (years), mean [SD]	39.9 [13.9]	38.7 [13.4]	41.6 [14.5]
Body weight (kg), mean [SD]	65.5 [12]	70.6 [10.5]	57.8 [9.58]
BMI, n (%)	BMI, n (%)		
<18.5 kg/m <sup>2</sup>	6 (5.8)	1 (1.6)	5 (12.2)
18.5-24.9 kg/m <sup>2</sup>	74 (71.8)	48 (77.4)	26 (63.4)
25-29.9 kg/m <sup>2</sup>	18 (17.5)	10 (16.1)	8 (19.5)
>30 kg/m <sup>2</sup>	5 (4.9)	3 (4.8)	2 (4.9)
Previous surgery, n (%)	54 (52.4)	29 (46.8)	25 (61)
Disease duration (years), median [range]	6.50 [1-30]	6.42 [1-30]	6.58 [1-23]
Currently smoking habits n (%)			
Yes	22 (21.4)	11 (17.7)	11 (26.8)
No	63 (61.2)	40 (64.5)	23 (56.1)
Ex-smoker	18 (17.5)	11 (17.7)	7 (17.1)

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Clinical activity, n (%)			
CDAI <150	55 (53.4)	36 (58.1)	19 (46.3)
>150 CDAI <450	48 (46.6)	26 (19.4)	22 (53.7)
Montreal age at diagnosis, n (%)			
A1: < 16 y	18 (17.5)	11 (17.7)	7 (17.1)
A2: 17–40 y	70 (68.0)	43 (69.4)	27 (65.9)
A3: > 40 y	15 (14.6)	8 (12.9)	7 (17.1)
Montreal disease location, n (%)			
L1: Ileum	36 (35)	20 (32.2)	16 (39.0)
L2: Colon	10 (9.7)	10 (16.1)	0
L3: Ileum and colon	54 (52.4)	30 (48.4)	24 (58.5)
L4: Upper GI tract	3 (2.9)	2 (3.2)	1 (2.4)
Montreal disease behavior, n (%)			
B1: Inflammatory 32 (31.1) 24 (38.7) 8 (19.5)			
B2: Stricturing	54 (52.4)	32 (51.6)	22 (53.7)
B3: Penetrating	17 (16.5)	6 (9.7)	11 (26.8)
Perianal disease, n (%)	19 (18.4)	12 (19.4)	7 (17.1)
Medications, n (%)			
None	30 (29.1)	19 (30.6)	11 (26.8)
5-ASA	17 (16.5)	11 (17.7)	6 (14.6)
IMMs	15 (14.6)	8 (12.9)	7 (17.1)
Biologics	41 (39.8)	24 (38.7)	17 (41.5)

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# RESULTS – Percentage of patients with AA levels below, above or within the range



Data are expressed as percentage of patients resulting below (blue column), above (orange column) or within (grey column) the reference range considered by the laboratory.

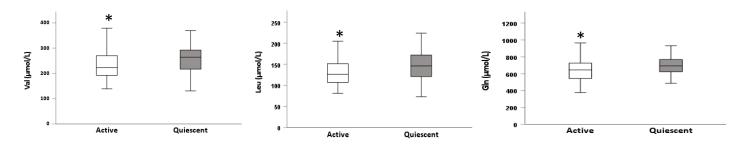
### RESULTS – Differences in AA concentrations between Active vs. Quiescent CD patients

TABLE 2 Serum amino acids concentration in patients with CD according to disease activity.

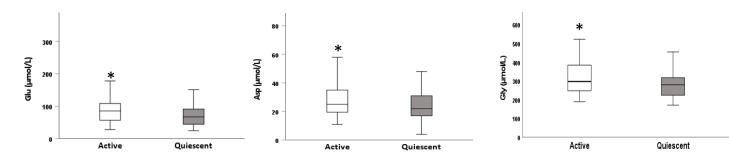
	All	Active	Quiescent	Reference (35, 36
EAA, µM/L				
Valine	233 (79)	223.5 (84)*	264 (76)	187-411
Leucine	139 (57)	127 (59.3)*	146 (53)	82-258
Isoleucine	74 (26)	73.5 (26)	74 (25)	42-111
Threonine	146 (65)	147.5 (64.3)	145 (67)	109-268
Lysine	176 (60)	169.5 (51)	183 (66)	150-286
Methionine	28 (10)	26.5 (10)	29 (10)	21-44
Histidine	88 (22)	84 (24.8)	90 (18)	72-131
Tryptophan	6 (3)	6 (3)	7 (3)	10-30
Phenylalanine	64 (17)	60 (17.3)	67 (14)	59-112
BCAA	445 (149)	421 (156)	484 (154)	-
Total EAAs	908 (264)	862 (253)	957 (263)	-
NEAA, µM/L				
Glutamic acid	73 (45)	85.5 (52.3)*	67 (50)	67-223
Asparagine	60 (15)	59.9 (15.8)	60 (16)	43-87
Arginine	101 (34)	95.5 (37)	107 (28)	36-172
Aspartic acid	23 (16)	25 (15.8)*	22 (14)	24-102
Glutamine	664 (141)	644 (187.5)**	693 (42)	432-871
Serine	147 (42)	147.5 (37.3)	146 (42)	111-297
Glycine	287 (113)	296.5 (139)*	280 (98)	240-460
Tyrosine	62 (20)	61 (21.8)	62 (17)	45-107
Alanine	379 (132)	376.5 (143.5)	397 (129)	193-597
Cysteine	31 (12)	30.5 (12.5)	31 (11)	5-55
Total NEAAs	1864 (417)	1853 (469)	1890 (316)	-
Other metaboli	tes, µM/L			
Ornithine	93 (12)	99.5 (46.8)	91 (44)	63-133
Citrulline	33 (11)	33 (10.8)	33 (14)	15-57
Taurine	151 (70)	150 (57)	155 (78)	50-270

Data are expressed as median and interquartile range. BCAA, branched-chain amino acid; EAAs, essential amino acids; NEAAs, non-essential amino acids. \*p < 0.01; \*p < 0.05.

### Val, Leu and Gln decreased in active vs. quiescent CD patients



### Glu, Asp and Gly increased in active vs. quiescent CD patients



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### RESULTS - Differences in serum AAs according to total intake of protein

Average intake of protein was  $71.2 \pm 22.5$  g (17% of total energy intake).

When expressed as g/kg of body weight, it was  $1.11 \pm 0.36 g/kg/body$  weight.

To assess the effects of protein amount on serum AAs, we used as reference values the protein requirements suggested by the last ESPEN guideline in adults with IBD to split our sample:

- 1.0 g/kg/die in remission phase
- 1.2 g/kg/die in active phase

Using the above values, 47 (46%) out 103 patients did not meet the protein requirements.

Serum levels of Threonine, Lysine and Arginine were reduced in the group with unmet protein intake (UPI) compared to the one meeting the requirements (MPI).

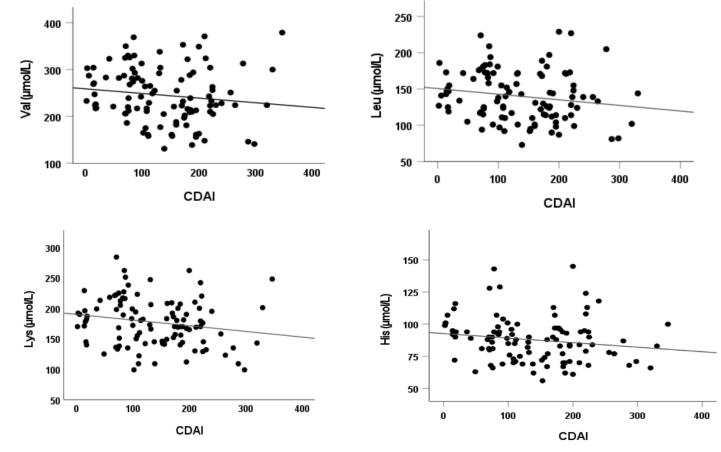
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TABLE 3 Ser	um amino acids concentration according to protein
requirement	s in patients with CD.

	UPI <i>N</i> = 47	MPI <i>N</i> = 56
EAA, µM/L		
Valine	224 (91)	259 (76)
Leucine	131 (48)	140 (55)
Isoleucine	72 (26)	77 (29)
Threonine	134 (61)*	156 (54)
Lysine 🗾	170 (41)**	190 (68)
Methionine	26 (8)*	29 (10)
Histidine	88 (23)	89 (20)
Tryptophan	6 (2.1)	7 (4)
Phenylalanine	61 (15)*	66 (20)
BCAA	424 (157)	484 (144)
Total EAAs	873 (231)	948 (246)
NEAA, µM/L		
Glutamic acid	74 (42)	70.5 (47)
Asparagine	59 (15)	60 (18)
Arginine	96 (43)*	108 (38)
Aspartic acid	23 (14)	24.5 (17)
Glutamine	684 (172)	662 (130)
Serine	145 (42)	149 (40)
Glycine	280 (151)	291 (73)
Tyrosine	61 (20)	62.5 (20)
Alanine	399 (104)	367 (134)
Cysteine	32 (13)	29 (10)
Total NEAAs	1892 (507)	1846 (288)
Other metabolites, µM/L		
Ornithine	92 (44)	93.5 (42)
Citrulline	34 (11)	33 (14)
Taurine	149 (71)	154.5 (72)

BCAA, branched-chain amino acid; EAAs, essential amino acids; IQR, interquartile range MPI, Met protein intake; NEAAs, non-essential amino acids; UPI, Unmet protein intake. \*\*p<0.01; \*p<0.05.</p>

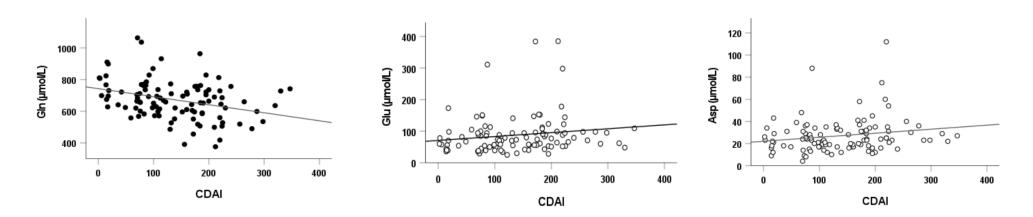
# **RESULTS - Correlation between EAAs and CDAI**



#### FIGURE 3

Correlations between serum essential AA levels and disease activity. Significant inverse Spearman's rank correlation coefficients between the following essential AAs valine (Val), leucine (Leu), lysine (Lys), and histidine (His) and disease activity (CDAI).

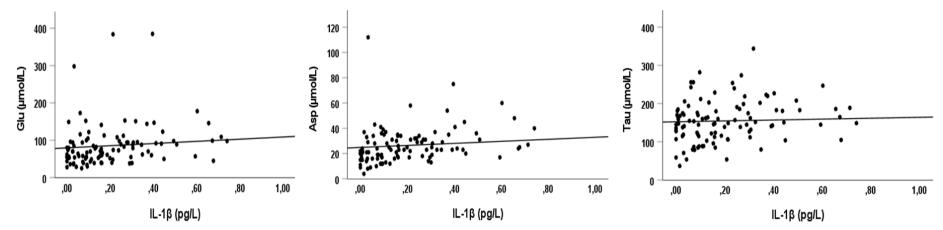
### **RESULTS – Correlation between NEAAs and CDAI**



#### FIGURE 4

Correlations between serum non-essential AA levels and disease activity. Significant Spearman's rank correlation coefficients between the following non-essential AAs: glutamine (Gln) glutamate (Glu), and aspartic acid (Asp) and disease activity (CDAI).

### RESULTS – Correlation between NEAAs and IL-18

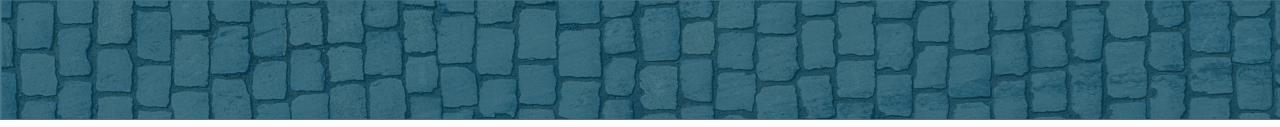


### FIGURE 5

Correlations between Glu, Asp, Tau levels and IL-1β. Significant direct Spearman's rank correlation coefficients between glutamate (Glu), aspartic acid (Asp), and taurine (Tau) and a proinflammatory cytokine (interleukin-1beta).

### CONCLUSIONS

- This secondary analysis showed differences in specific AA concentrations between clinically active and quiescent CD patients as well as according to different protein intakes, especially for some EAAs.
- Levels of serum Trp were reduced in 90% of patients unrelated to CDAI.
- Glu and Asp values were positively correlated to CDAI and serum IL-1β, suggesting a potential link with disease activity.
- In light of these promising results, extensive research is needed to understand the mechanisms underpinning the relationship between AAs, disease activity and protein intake in patients with CD.



### Thanks for your attention!

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