

Supplementary Material for

Fish sidestream-derived protein hydrolysates suppress DSS-induced colitis by modulating intestinal inflammation in mice

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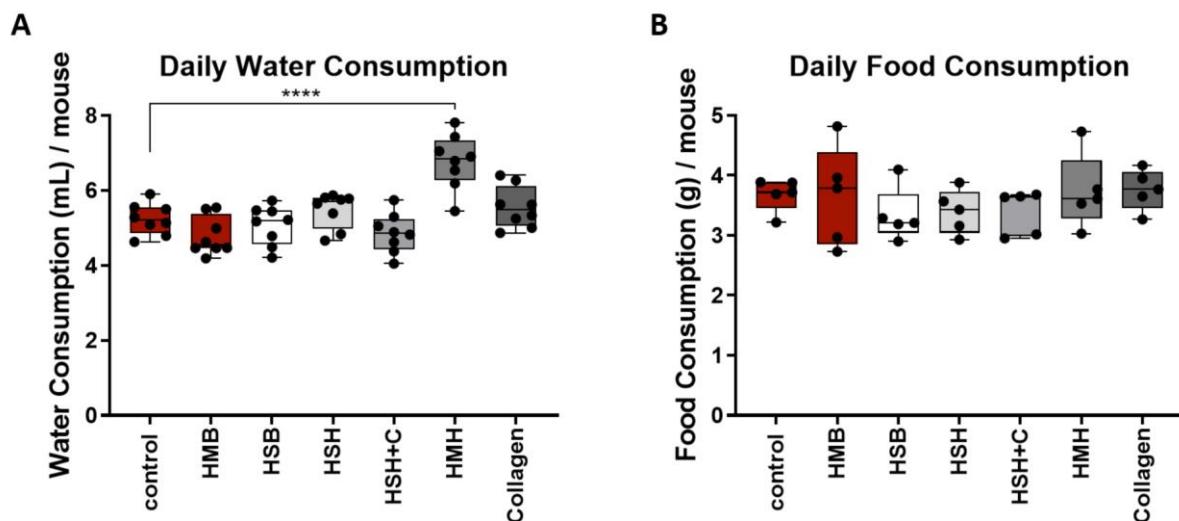


Figure S1. Daily A. water and B. food consumption of animals fed diets containing the different supplements. Graphs represent median \pm SD and 2-way ANOVA statistical analysis was performed. *** p <0.0001.

Table S1. Composition of the different fish sidestream-derived supplements used in the cutaneous model.

Chemical composition		HMB	HMH	HSB	HSH	Collagen
Crude protein Kjeldahl (N*6.25)	%	82.4	59.3	89.8	89.3	>90
Total dry matter	%	96.3	96.4	96.3	98	96.7
Ash	%	15.7	38.3	9.3	12.6	3.1
Watersoluble crude protein	g/100g sample	82	58.5	88.7	89.2	>95
Peptide size distribution						
Mw-peptide > 20000	% of water sol. peptides	<0.1	<0,.1	<0,.1	<0,.1	<0,.1
Mw-peptide 20000-15000	% of water sol. peptides	<0,.1	<0,.1	<0,.1	<0,.1	<0,.1
Mw-peptide 15000-10000	% of water sol. peptides	0.1	0.1	0.1	0.1	0.1
Mw-peptide 10000-8000	% of water sol. peptides	0.1	0.3	0.2	0.2	0.5
Mw-peptide 8000-6000	% of water sol. peptides	0.5	1	0.9	1	2.4
Mw-peptide 6000-4000	% of water sol. peptides	1.9	3.7	3.3	3.6	9.4
Mw-peptide 4000-2000	% of water sol. peptides	8.2	13.9	12.9	15.9	28.1
Mw-peptide 2000-1000	% of water sol. peptides	14.5	17.7	18.4	22.2	28
Mw-peptide 1000-500	% of water sol. peptides	17.7	16.6	18.9	19	17.6
Mw-peptide 500-200	% of water sol. peptides	19.7	16.9	20.3	17.3	9.3
Mw-peptide 200-	% of water sol. peptides	37.3	29.7	24.9	20.7	4.6
Total amino acids						
Aspartic acid	g/100g sample	6.5	4.4	7	6.8	6.4

Glutamic acid	g/100g sample	10.7	7.4	10.9	10.8	10
Hydroksyproline	g/100g sample	1.4	1.9	3	4.1	9.8
Serine	g/100g sample	3.1	2.5	3.4	3.8	6.1
Glycine	g/100g sample	5.3	6.2	8.8	12.1	26
Histidine	g/100g sample	5	1.8	1.8	1.7	1.1
Arginine	g/100g sample	4.5	3.7	5.3	5.6	8.9
Threonine	g/100g sample	2.9	2	3.2	2.9	2.8
Alanine	g/100g sample	4.6	4.2	5.6	6	9.8
Proline	g/100g sample	3	3.1	4.5	6.1	11.7
Tyrosine	g/100g sample	1.7	1.1	1.7	1.4	0.37
Valine	g/100g sample	3.1	2	3.2	2.8	1.9
Methionine	g/100g sample	1.9	1.4	2.3	2.4	2.4
Isoleucine	g/100g sample	2.4	1.5	2.6	2.1	1.1
Leucine	g/100g sample	5	3.2	5	4.3	2.4
Phenylalanine	g/100g sample	1.9	1.5	2.3	2.3	2.1
Lysine	g/100g sample	6.9	4.3	6.5	5.5	3.7

Quality parameters

Putrescine	mg/kg	63	44	110	180	41
Cadaverine	mg/kg	<20	<20	<20	<20	<20
Histamine	mg/kg	74	38	67	68	23
Trimethylamin-N	mg N/100 gram	17	15	14	<1	66
Trimethylaminoxide-N	mg N/100 gram	162	53	125	<1	5

Free amino acids

Aspartic acid	g/100 g sample	0.06	0.07	0.09	0.1	0
Glutamic acid	g/100 g sample	0.39	0.27	0.36	0.32	0.01
Hydroksyproline	g/100 g sample	0.01	0.01	0.02	0.03	0
Serine	g/100 g sample	0.09	0.12	0.12	0.16	0.02

Asparagine	g/100 g sample	0.02	0.01	0.02	0.02	0.01
Glycine	g/100 g sample	0.15	0.21	0.15	0.21	0.05
Glutamine	g/100 g sample	0.21	0.43	0.43	0.45	0
Histidine	g/100 g sample	3.2	0.81	0.22	0.2	0
Threonine	g/100 g sample	0.1	0.1	0.14	0.14	0
Alanine	g/100 g sample	0.29	0.27	0.44	0.4	0.03
Arginine	g/100 g sample	0.25	0.34	0.19	0.21	0.02
Proline	g/100 g sample	0.06	0.09	0.1	0.07	0
Tyrosine	g/100 g sample	0.18	0.26	0.18	0.14	0.02
Valine	g/100 g sample	0.1	0.14	0.23	0.19	0.01
Methionine	g/100 g sample	0.29	0.36	0.47	0.37	0
Cysteine	g/100 g sample	0	0	0	0	0
Isoleucine	g/100 g sample	0.08	0.13	0.24	0.19	0.01
Leucine	g/100 g sample	0.42	0.78	0.83	0.72	0.04
Phenylalanine	g/100 g sample	0.3	0.43	0.48	0.67	0.07
Tryptophane	g/100 g sample	0.07	0.07	0.12	0.1	0
Lysine	g/100 g sample	0.56	0.39	0.4	0.3	0
Creatinine	g/100 g sample	0.94	0.15	0.4	0.34	0.07
β-alanine	g/100 g sample	0	0	0.14	0.21	0
Taurine	g/100 g sample	0.85	1.4	0.46	1.3	0.03
4-aminobutanoic acid	g/100 g sample	0	0	0	0.01	0
Citrulline	g/100 g sample	0	0	0	0.01	0
Carnosine	g/100 g sample	0.06	0.03	0.08	0.03	0.01
Anserine	g/100 g sample	0.11	0.05	2.3	0.64	0
L-Ornithine	g/100 g sample	0.02	0.02	0.02	0.02	0

Table S2. List of oligonucleotides used in real time PCR reactions.

Primer	Sequence
Actin-Fwd	CATTGCTGACAGGGATGCAGAAGG
Actin-Rev	TGCTGGAAGGTGGACAGTGAGG
TNF α -Fwd	GCCCCACGTCGTAGCAAACCACC
TNF α -Rev	CGGGGCAGCCTGTCCCTTG
IL-6-Fwd	CAAAGCCAGAGTCCTTCAGAG
IL-6-Rev	CACTCCTCTGTGACTCCAGC
IL-1 β -Fwd	CGGACCCC AAAAGATGAAGGGCTG
IL-1 β - Rev	GCTCTTGTGATGTGCTGCTGCGAG
Cxcl1- Fwd	CCCAAACCGAAGTCATAGCCA
Cxcl1- Rev	CTCCGTTACTTGGGGACACC
Cxcl2-Fwd	CGCCCAGACAGAAGTCATAGCCAC
Cxcl2-Rev	CGTTGAGGGACAGCAGCCCAG
Ccl2-Fwd	GGCTGGAGCATCCACGTGTTGG
Ccl2-Rev	TTGGGGTCAGCACAGACCTCTCTC
Ccl3-Fwd	GAAGGATAACAAGCAGCAGCG
Ccl3-Rev	TTCTCTTAGTCAGGAAAATGACACC
IL-10 Fwd	GCGCTGTCATCGATTCTCCCCCTG
IL-10-Rev	GGCCTTGTAGACACCTTGGTCTTGG
TGF-b-Fwd	GACACACAGTACAGCAAGGTCC
TGF-b-Rev	CGACCCACGTAGTAGACGATG