



Immunomodulatory effects of dietary seaweeds in LPS challenged Atlantic salmon Salmo salar as determined by deep RNA sequencing of the head kidney transcriptome

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Aim

Results (RNAseq)

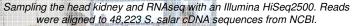
Seaweeds may represent immunostimulants that could be used as health-promoting fish feed components thereby offering an alternative for the use of antibiotics.

This was performed to study gain insights the into immunomodulatory effects of dietary seaweeds in Atlantic salmon. Specifically tested were:

- SW1: 10% inclusion levels of Laminaria digitata;
- SW2: commercial blend of seaweeds (Oceanfeed®);
- against a fishmeal based control diet (FMC).

Differences between groups were assessed in growth, feed conversion ratio (FCR) and blood parameters hematocrit (Hct) and hemoglobin (Hb). After a LPS challenge in fish representing each of the three groups, RNAseg was performed on head kidneys to determine transcriptomic differences in response to the immune activation, to our knowledge for the first time in fish in this context.









Results (growth performance)

No differences in growth performance were detected between FMC, SW1 and SW2 fish. Interestingly, fish fed with seaweed diets showed much less individual variation in growth performance than the FMC fish. SW fish showed higher FCR than fish fed with seaweed diets. No significant differences existed between groups in Hct and Hb.

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RNAseq resulted in ~154 million single-read 1 x 50 nt reads (~8 Gb data). De novo assembly then resulted in 87,600 cDNA contigs

ranging in size from 200 - 14,086 nt. The comparison SW1 vs FMC yielded 28,312 expressed contigs associated with S. salar genes: 59% of the NCBI sequences. 144 of these contigs were differentially expressed: 74 up-regulated (51%) and 70 down-regulated (49%). The comparison SW2 vs FMC yielded 28,189 expressed contigs associated with S. salar genes: 58% of the NCBI sequences. 246 of these contigs were differentially expressed: 136 up-regulated (55%) and 110 down-regulated (45%). SW1 vs. FMC and SW2 vs. FMC revealed 54 common differentially expressed contigs. Expression of all these contigs was in the same direction, either up- or downregulated. These contigs can be considered representing the common genes in the immunomodulatory effects of the tested dietary seaweeds. The immune genes were MHC class I and II and HLA class II antigens, and CD209 antigen-like protein E, all involved in antigen processing and presentation.

specific for SW1 MHC class I (UBA) mRNA, UBA*0901 allele	fc	362			
T-cell receptor alpha chain V region 284 precursor putative mRNA		7.19		/	
MHC class I antigen (UBA) mRNA, UBA*4001 allele		6.61			
Macrophage migration inhibitory factor putative mRNA	Inf	0.01	1		V1 vs FMC SW2 vs FMC
MHC class I antigen (Sasa-UBA) mRNA, Sasa-UBA*0902 allele	Inf		1	28,312 e	28 189 avor senes (58%)
BOLA class I histocompatibility antigen, alpha chain BL3-7 precursor putative mRNA		0.154	1		ff. expressed: 246 diff. expressed: 136 up (55%)
MHC class I (UBA) mRNA, UBA*0501 allele		0.134	•		lown (49%) 110 down (45%)
MHC class I (OBA) III KINA, OBA OSOL allele		0.0467			
MHC class I (UBA) mRNA, UBA*0701 allele		0.00130			
INITE Class I (OBA) IIIKINA, OBA O/OI allele		0.00150			all same direction
specific for SW2	fc		-	commo	n genes in the seaweed effects on immune activation
MHC class I antigen (Sasa-UBA) mRNA, Sasa-UBA*3701 allele		288			
H-2 class II histocompatibility antigen gamma chain putative mRNA		137			The comparisons SW1 vs
MHC class II antigen alpha chain (Sasa-DAA) mRNA		39.7			
MHC class II alpha mRNA		18.6			FMC and SW2 vs FMC
Tumor necrosis factor receptor superfamily member 11B precursor putative mRNA		4.14			
CD209 antigen-like protein E putative mRNA		3.21			revealed 54 common
Ig kappa chain V-IV region Len putative mRNA		2.86	i i		differentially expressed
partial mRNA for MHC class II antigen beta chain (DAB gene)	Inf				, ,
MHC class I antigen (Sasa-UBA) mRNA, Sasa-UBA*1402 allele	Inf				genes all in the same
Class I histocompatibility antigen, F10 alpha chain precursor putative mRNA		0.254			5
MHC class I (UBA) mRNA, UBA*0501 allele		0.131		di	rection. The Table lists the
MHC class I antigen (Sasa-UBA) mRNA, Sasa-UBA*3501 allele		0.125			immune genes that were
Proliferating cell nuclear antigen putative mRNA		0.113			0
MHC class I (UBA) mRNA, UBA*1501 allele		0.00567	1		specific for SW1, for SW2
Anamorsin putative mRNA		C			
					and those in common
common SW1-SW2	SW	1 (fc)		2 (fc)	(fc= fold change; 'inf'
MHC class II antigen beta chain (DAB gene)		309		171	
MHC-Sasa class II (clone c144)		54.7		7.76	indicates expression only
MHC class I (UBA) mRNA, UBA*0201 allele		36.8		35.0	in SW diets and not
MHC class I antigen (Sasa-UBA) mRNA	Inf		Inf		in Svv diets and not
MHC class I antigen (Sasa-UBA) mRNA, Sasa-UBA*0202 allele	Inf		Inf		in FMC)
MHC class II antigen alpha chain (Sasa-DAA) mRNA	Inf		Inf		III I W(O)
HLA class II histocompatibility antigen, DQW1.1 beta chain precursor putative mRNA			Inf		
CD209 antigen-like protein E putative mRNA	Inf		Inf		
MHC class I (UBA) mRNA, UBA*1401 allele		0.350		0.351	
MHC-Sasa class II B (clone c22)		0.276		0.00283	
MHC class I antigen (Sasa-UBA) mRNA, Sasa-UBA*3901 allele		0.00349		0	

Conclusions

- · Seaweed fed fish showed higher FCRs but more homogenous growth performance than the control fish.
- Dietary seaweeds caused important immunomodulatory effects changing antigen processing and presentation.