



Hva viser genanalyser av muskulatur hos laks med mørke flekker

*Aleksei Krasnov, Hooman Moghadam
Nofima, Ås*

Spørsmål

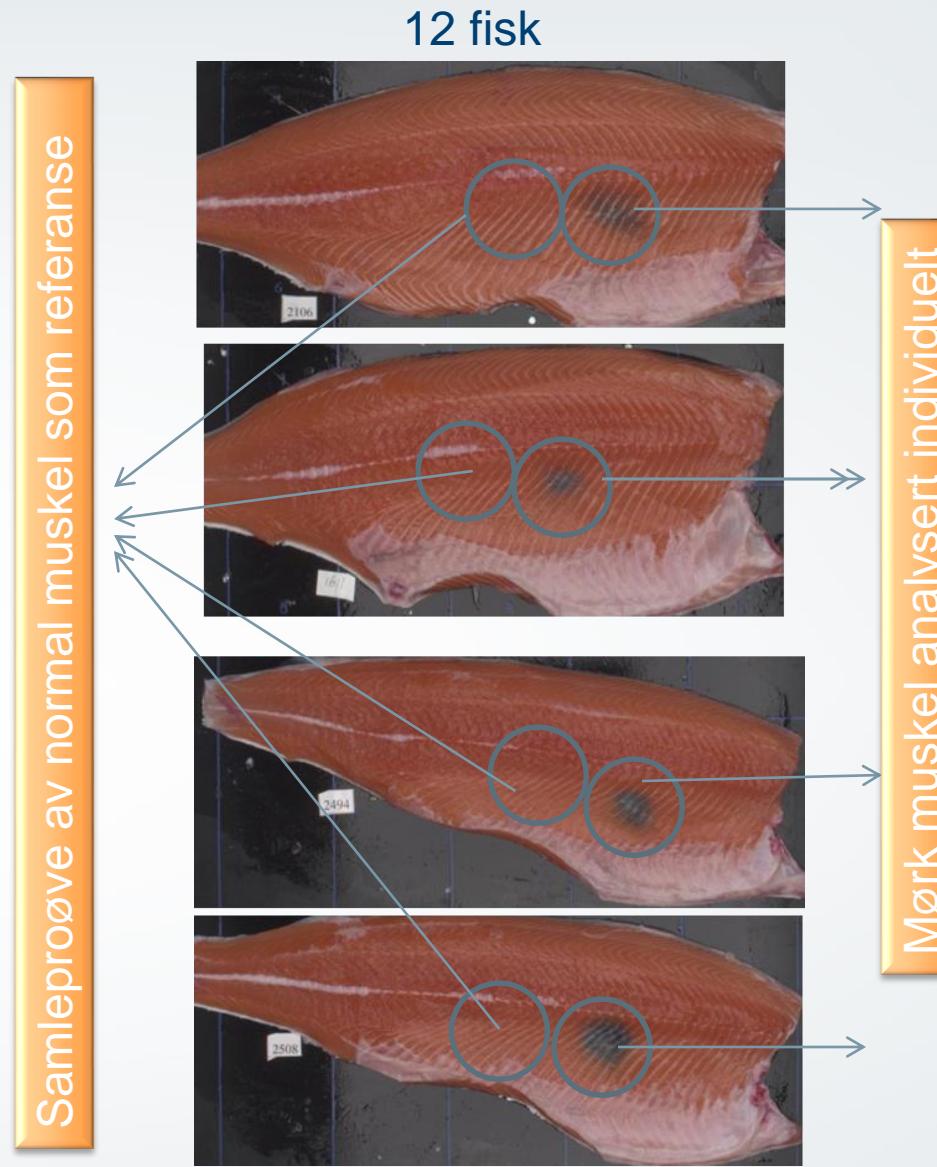
Har mørke flekker lik eller ulik profil?

Vevsskade og nydannelse av vev?

Type betennelse?

Årsak til mørkpigmentering?

Genuttrykk viste vesentlig forskjell mellom normal og mørk muskel



Undersøkt:
Forskjell mellom normal og
mørk muskel

Metode:
Nofima's 15 k Atlantic salmon
oligo chip,
STARS bioinformatics

Resultater genuttrykk:

1570 DE genes (>2-fold, $p<0.01$)

946 høyere i mørk muskel

624 lavere i mørk muskel

Spørsmål

Har mørke flekker lik eller ulik profil?

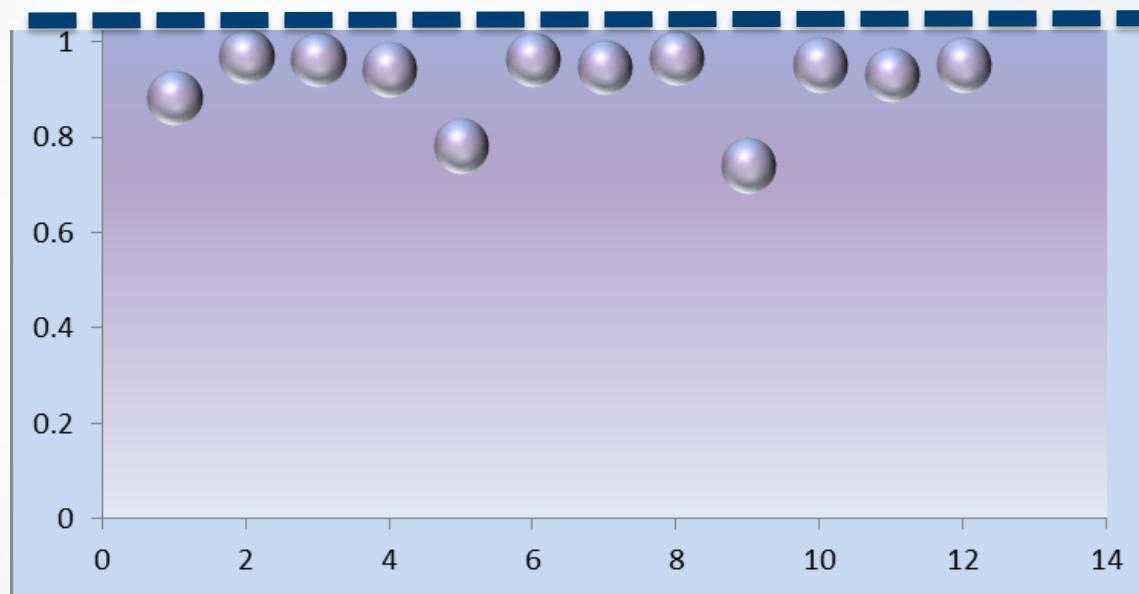
Vevsskade og nydannelse av vev?

Type betennelse?

Årsak til mørkpigmentering?

Tilnærmet likt genuttrykk uavhengig av misfarging og størrelse på flekkene

Korrelasjon, r



Korrelasjon = 1
betyr at genuttrykket er
eksakt det samme. For
vårt materiale var
korrelasjonen 0,92

Flekk fra laks 1 - 12

Resultatene tyder på at vevsskaden hadde samme årsak

Correlation: Pearson r (mean = 0.917)

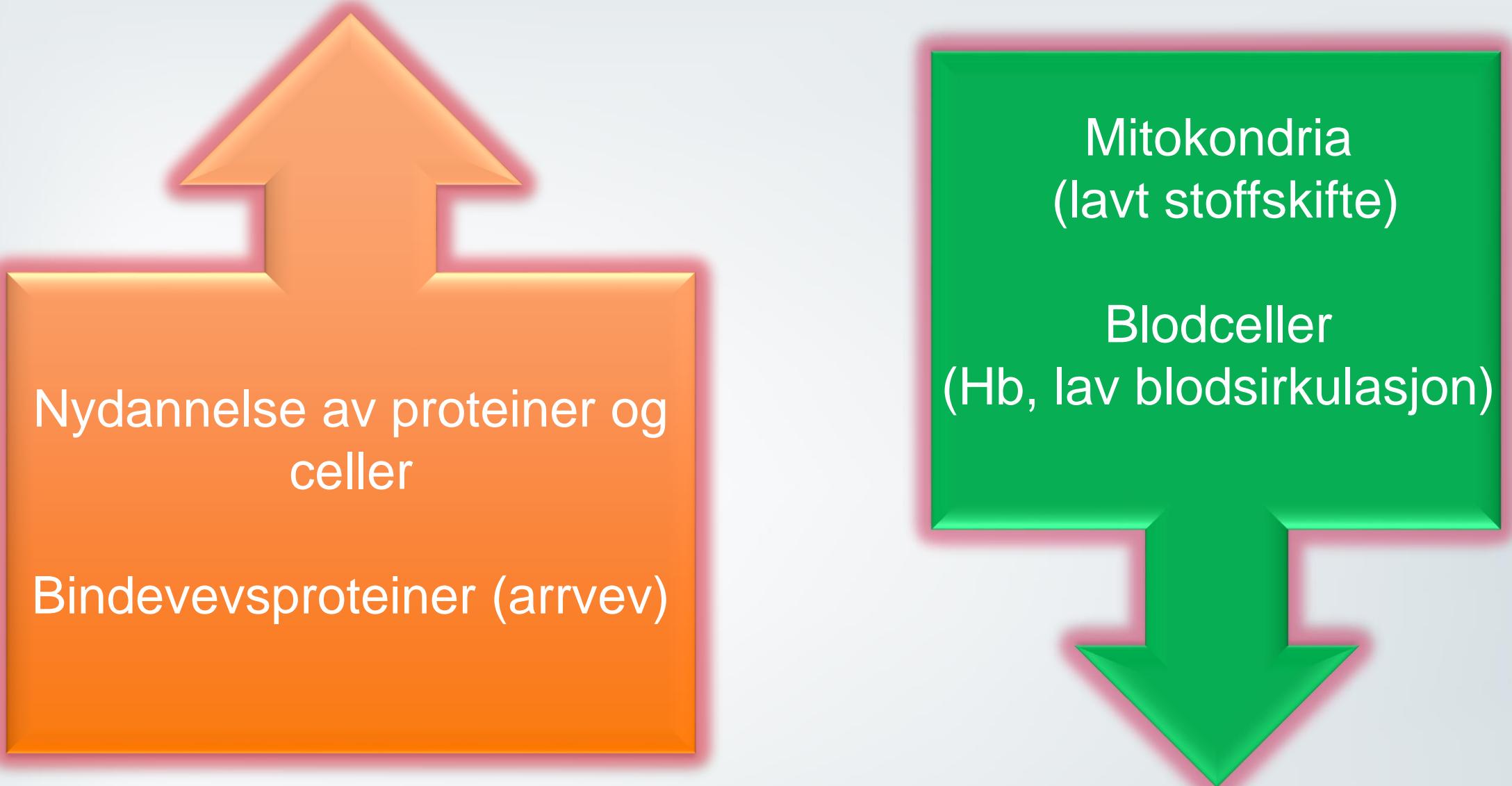
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Nydannelse av proteiner og
celler

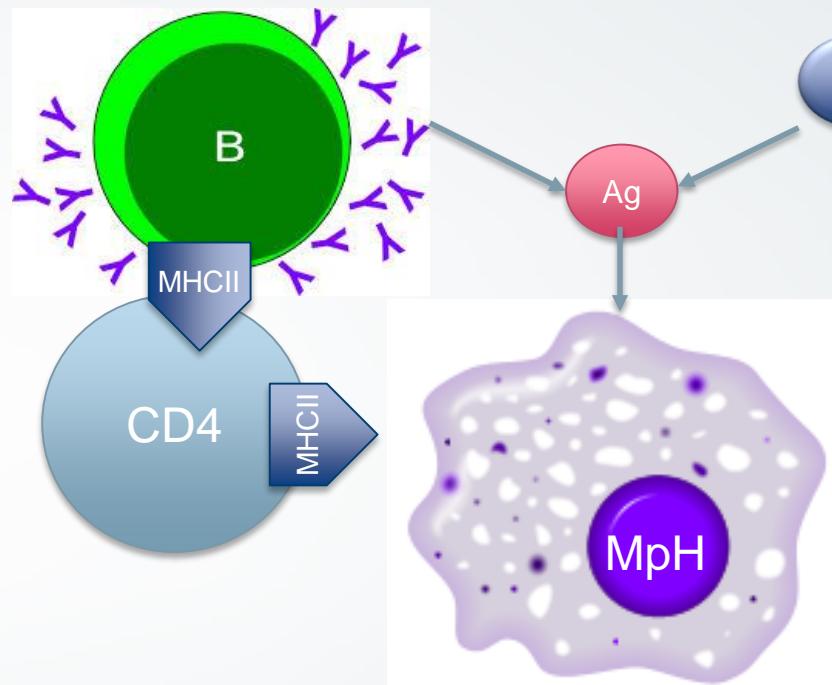
Bindevevsproteiner (arrhev)

Mitokondria
(lavt stoffskifte)

Blodceller
(Hb, lav blodsirkulasjon)

Reparasjon & nydanning, erstatning av muskel med bindevev

CD8



VRG

AFP

Mild kronisk betennelse

Årsak?

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RESEARCH ARTICLE

Open Access

*Piscine orthoreovirus (PRV) in red and melanised foci in white muscle of Atlantic salmon (*Salmo salar*)*



Håvard Bjørgen¹, Øystein Wessel², Per Gunnar Fjelldal³, Tom Hansen³, Harald Sveier⁴, Håkon Rydland Sæbø⁵, Katrine Bones Enger⁶, Eirik Monsen⁷, Agnar Kvellestad¹, Espen Rimstad² and Erling Olaf Koppang^{1*}

Concerns:

Relativt lav interferon respons

CD4 > CD8

MHCII > MHC I

It's like anything known?

Statistisk Meta Analyse: Sammenligning med andre forsøk

FHF Multifactorial disease
Contents lists available at ScienceDirect
Fish & Shellfish Immunology
journal homepage: www.elsevier.com/locate/fsi

Full length article
Comparison of transcriptomic responses to pancreas disease (PD) and heart and skeletal muscle inflammation (HSMI) in heart of Atlantic salmon (*Salmo salar* L)

Lill-Heidi Johansen ^{a,*}, Hanna L. Thim ^b, Sven Martin Jørgensen ^a, Sergey Afanasyev ^{a,c}, Guro Strandskog ^b, Torunn Taksdal ^d, Kjersti Fremmerlid ^b, Marion McLoughlin ^e, Jorunn B. Jørgensen ^b, Aleksei Krasnov ^a

^a Nofima AS, P.O. Box 6122, N-9291 Tromsø, Norway
^b Norwegian College of Fisherries Science, UiT The Arctic University of Norway, N-9037 Tromsø, Norway
^c Sechenov Institute of Evolutionary Physiology and Biochemistry, M. Toreza Av. 44, Saint Petersburg 194223, Russia
^d Norwegian Veterinary Institute, P.O. Box 750, N-0106 Oslo, Norway
^e Fish Vet Group, 22 Carsegar Road, Inverness IV3 8EX, UK

NFR Plattform for Virusvaksiner i fisk, VivaFish
Børre Robertsen (Univ Tromsø)

Muskel injeksjon:

- Placebo (PBS)
- Kontroll plasmid (bakterie DNA)
- Plasmid som produserer IFNa

Condition	Antigen	Trauma	Lesion	Timing
Dark spots	Unknown	Unknown	No	Months?
HSMI	Virus	No	Yes	Weeks
PBS injection	No	Small	No	1-2 weeks
Plasmid	Bacterial DNA	Small	No	1-2 weeks
Plasmid IFNa	Bacterial DNA+IFNa	Small	No	1-2 weeks



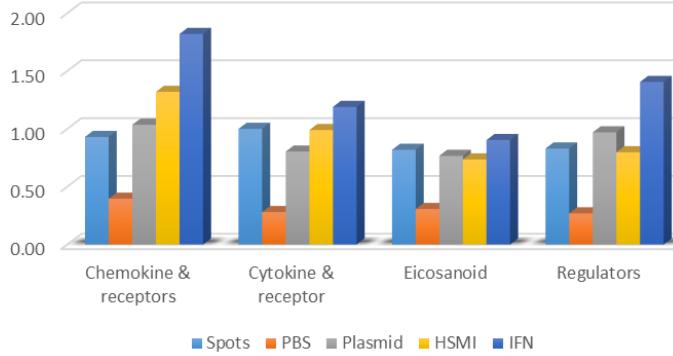
Innate antiviral responses in spots are weak
Correlation is low

TNF α - related responses to treatments are similar

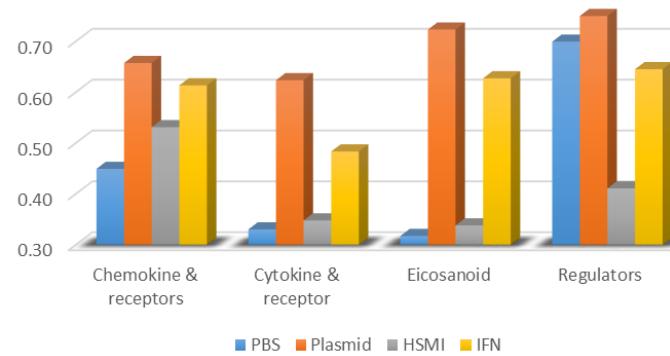
Respons likner ikke
muskel med virusinfeksjon

Gene	Spots	HSMI	PBS	Plasmid	IFN
Barrier-to-autointegration factor	1.30	3.14	0.63	3.81	5.64
Receptor transporting protein 3	0.49	4.80	-1.14	1.57	5.49
Interferon-induced protein 44	0.34	4.14	0.74	0.47	5.15
Very large inducible GTPase 1-1	1.12	4.36	-0.05	3.38	6.67
Sacsin	1.29	5.07	-0.28	3.74	6.12
Gig2-3	0.45	2.38	0.80	3.28	5.15
Mx1	0.49	3.91	0.14	3.90	7.09
Viperin	-0.11	3.67	0.15	2.94	6.39
Ubiquitin-like protein-1	-0.29	4.37	0.01	3.97	5.51
VHSV-inducible protein-4	-1.15	3.10	0.40	2.71	4.57

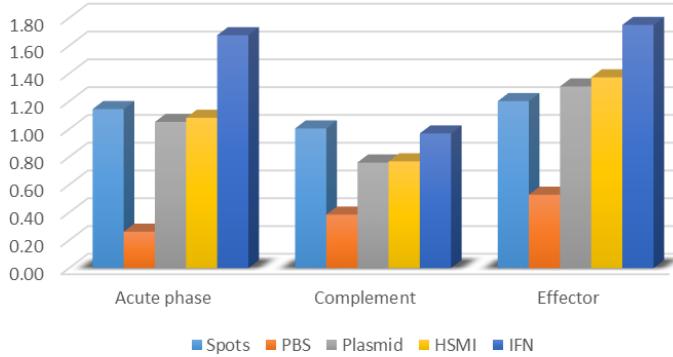
Signaling



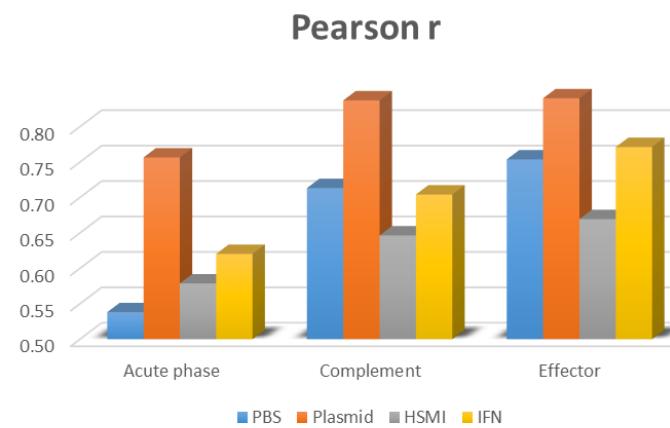
Pearson r



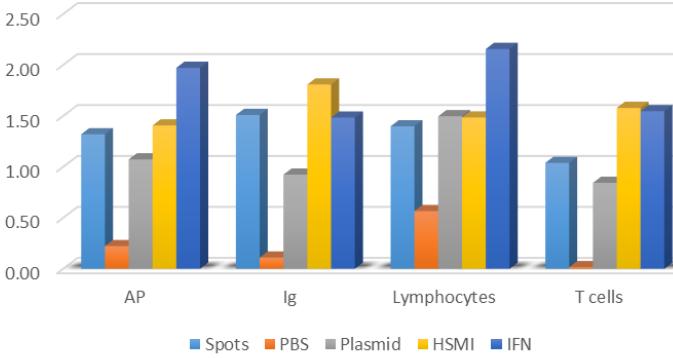
Effectors



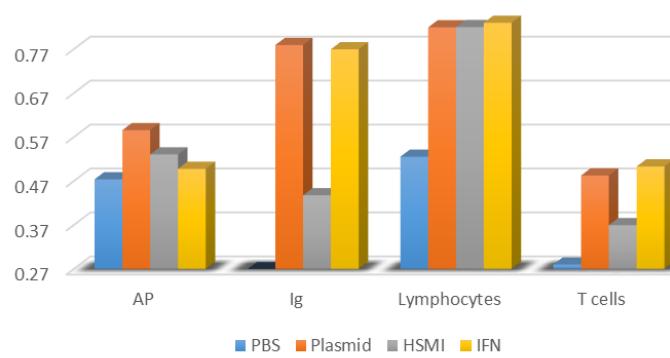
Pearson r



Adaptive



Pearson r



VivaFish fiskemateriale

Immune reactions in spots are similar by magnitude to responses to HSMB and bacterial DNA

Similarity (correlation):

Bacterial DNA > Bacterial DNA + IFNa > HSMI > injection

Least similar: T cells response

Genes	Spots	HSMI	PBS	Plasmid	IFN
CD2	0.69	2.09	1.17	2.19	2.36
CD4-1	3.24	1.11	0.06	-0.43	1.29
CD4-2	2.98	1.95	1.97	0.71	2.59
CD4-3	3.24	0.06	-0.43	1.11	1.29
CD4-4	0.47	1.05	0.58	0.43	1.46
CD8 alpha	1.73	3.30	1.05	0.43	1.17
CD8 beta	0.94	3.57	0.46	-0.88	0.82
CD8 beta	0.85	2.56	0.53	0.32	1.20
CD8 beta	0.41	2.10	-0.70	-0.60	0.31

Mørke flekker samme type
genuttrykk som laks injisert med
bakterie DNA

Immungener som var spesifikke for mørke flekker

- Og uttrykk av de samme genene ved kontrollert «smitte»

Genes	Spots	HSMI	PBS	Plasmid	IFN
Transcription factor PU.1	5.60	2.21	-1.23	1.21	-1.17
Transcription factor SOX-4	2.67	1.10	1.13	-1.12	1.33
CD80-like protein	7.16	1.65	1.00	1.64	2.03
C1q TNF-related protein 5	5.32	1.00	1.19	1.61	1.30
Nattectin C-type lectin	44.53	-1.67	-1.28	2.61	0.97
15-hydroxyprostaglandin dehydrogenase	3.95	-1.88	-1.16	1.21	-1.39
Leukotriene b4 12-hydroxydehydrogenase	11.16	1.22	0.95	-1.08	-1.87
72 kDa type IV collagenase	3.90	1.30	-1.41	-1.28	-2.10

Mulige gen-markører for melanisering

Dual RNA-Seq

- From 6 individuals total RNA was extracted from both normal and adjacent dark spots;
- RNA sequenced using Illumina platform;
- Sequenced reads aligned against the salmon and all viral and bacterial sequence data;

Total RNA (depleted from salmon rRNA)



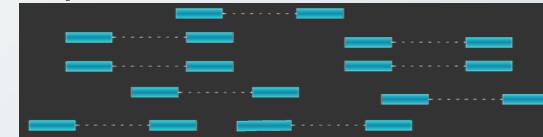
Fragment RNA



Reverse transcribe into cDNA



Sequence cDNA



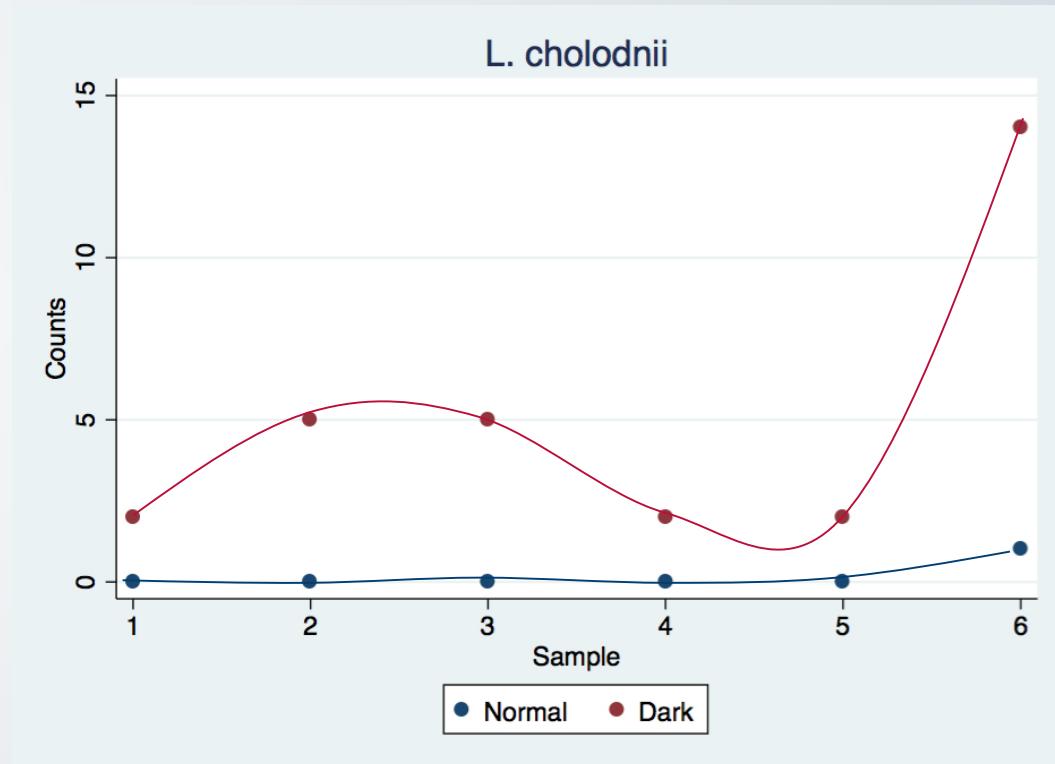
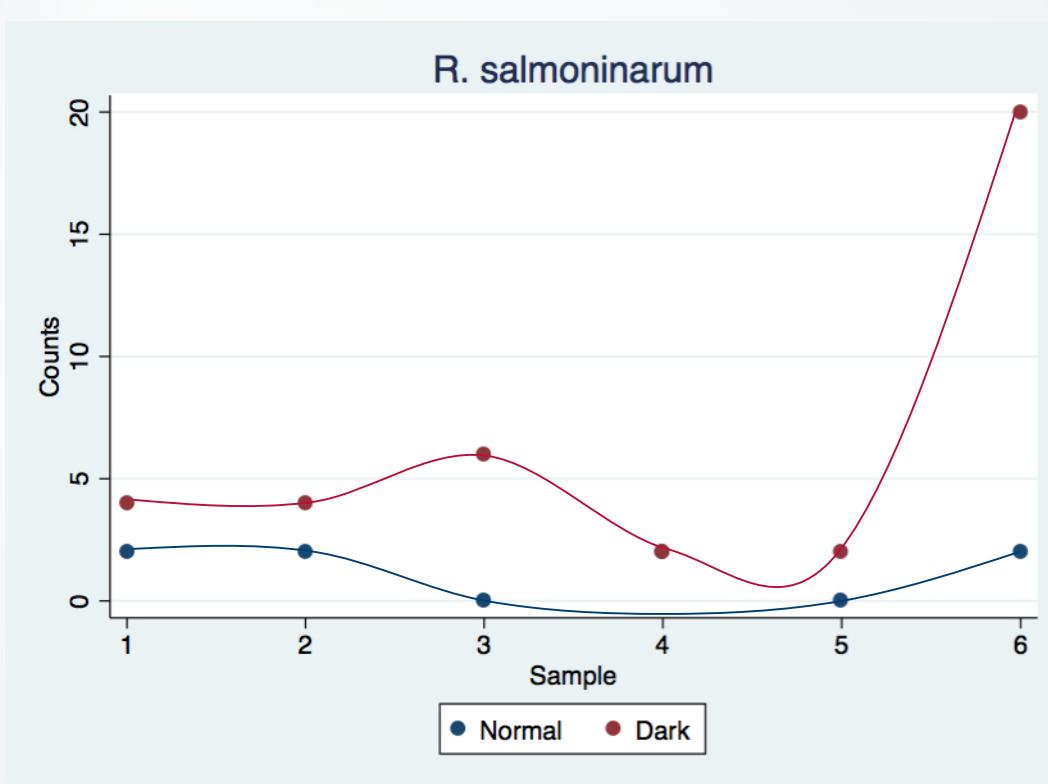
Mapping reads to pathogen genomes



Mapping reads to salmon genome

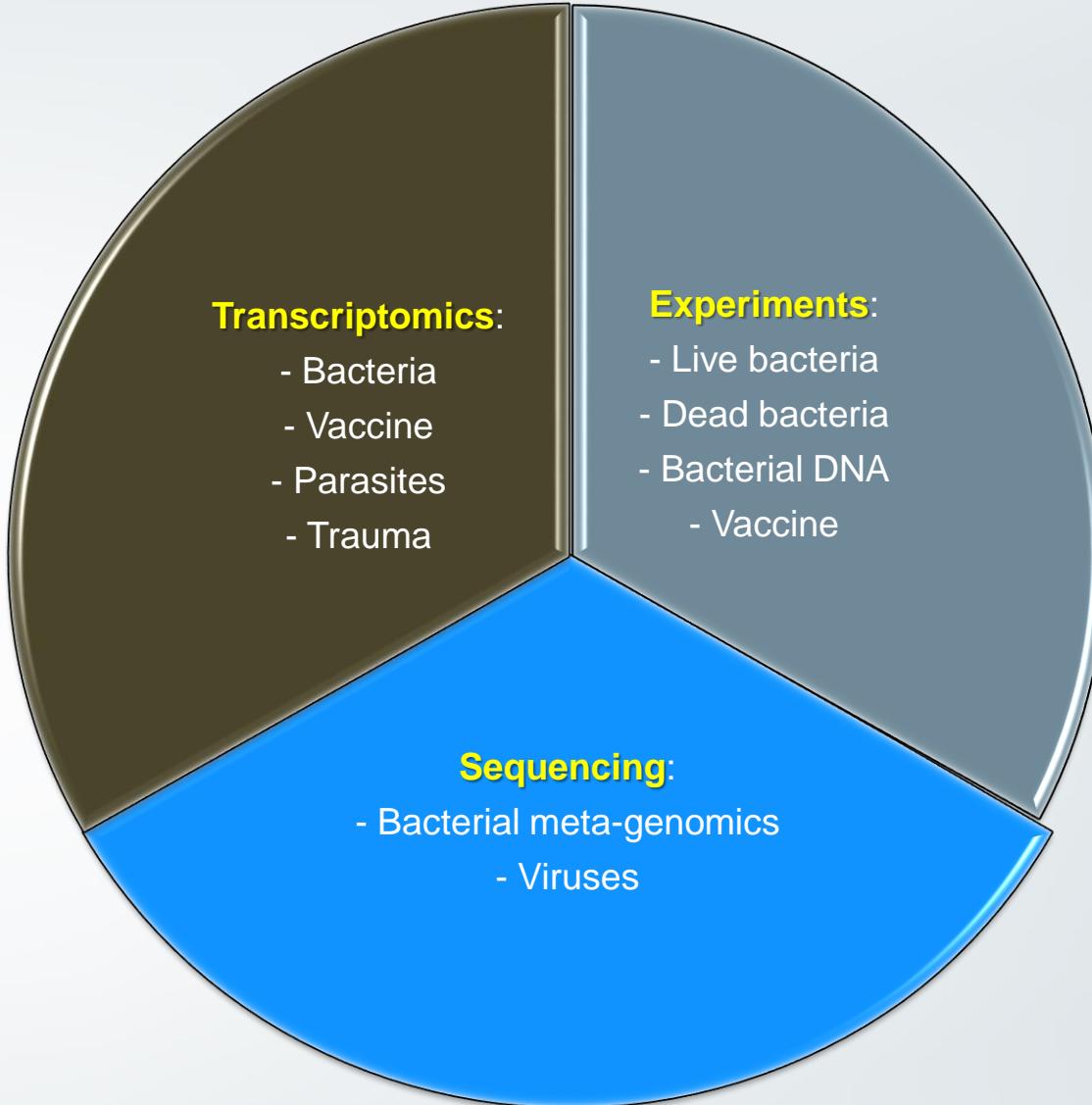
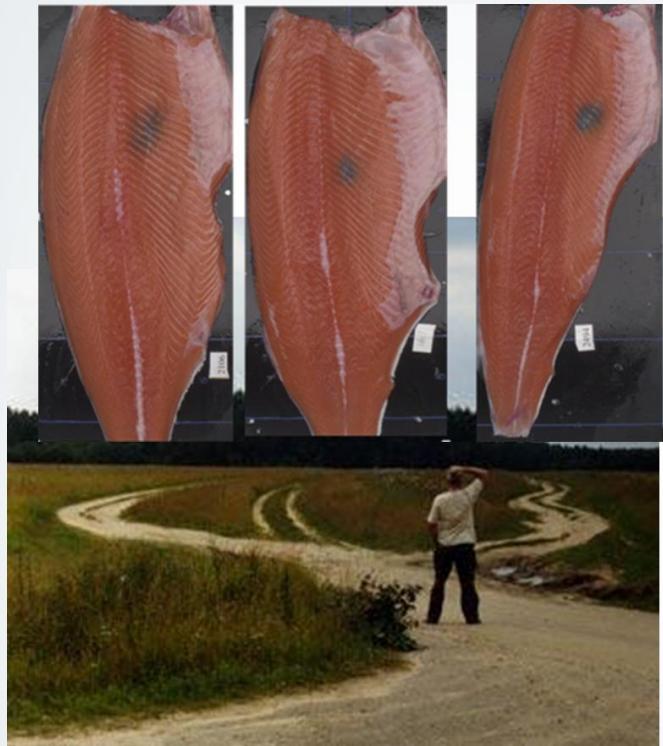


Eksempler på bakterier med ulik konsentrasjon i normal muskel og mørk muskel (loads, rRNA)



Årsak til at flekkene?

Årsak	Sannsynlighet
Virus	Lav
Skader alene	Lav
Patogene bakterier	Lav
Ikke patogene kakterier	Middels høy
Ikke patogene kakterier + skade	Mest sannsynlig
<i>Andre årsaker ???</i>	



Videre forskning
Synes vi bør se
litt nærmere på
denne på
mandag
Bør også
highlighte det
med
genmarkører

Genetic enrichment of functional categories

- Similar to microarray, many immune specific genes are upregulated in dark spots;
- Found evidence many genes involved in muscle development and lipid synthesis and metabolism to also been upregulated;

