FHF prosjekt 900706:

"Sporing av laks: SNP-tilnærming"

"Tracing the origin of farmed Atlantic salmon escapees by DNA parentage assignment; optimizing methods, and real-life validation studies"





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<u>CIGENE</u>







- Cigene Est. 2003 and funded by the Norwegian Government for:
 - high-throughput genotyping in humans, micro-organisms, plants, animals and fish.
 - integration of experimental and theoretical methods needed to bridge the genotype-phenotype gap.
 - development of a competence base to advance livestock and marine functional genomics



WP1.2 Utvalg av markører samt teknisk og kostnadsmessig optimalisering av SNP-basert genotyping.

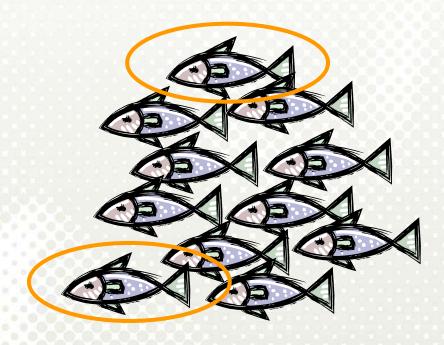
 I denne arbeidspakken vil man arbeide videre med å velge ut antall og finne optimal gjennomføringsmodus for SNP-basert tilordning av foreldrefisk.

Potential Parents

Sample



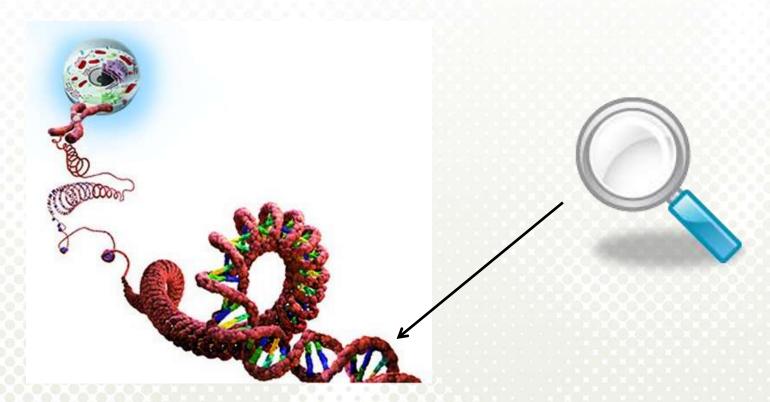






Single Nucleotide Polymorphisms (SNP)

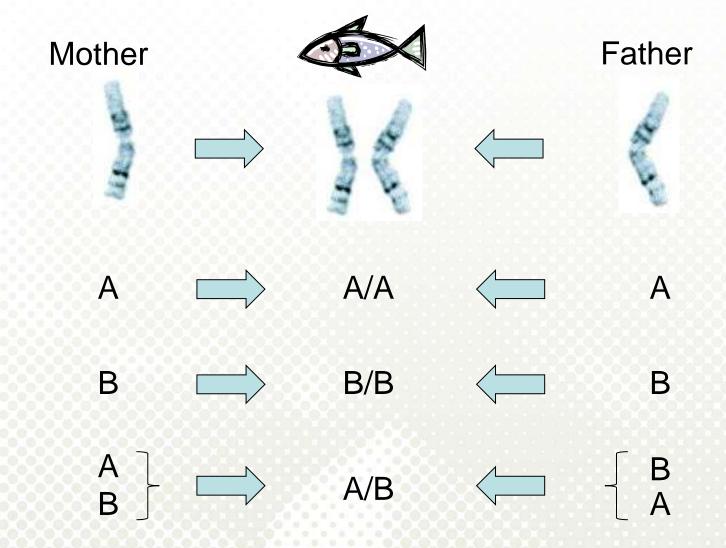
 SNPs are small genetic variations within chromosomes that are passed on through generations



They are easy to assay, and easy to code (binary).



Single Nucleotide Polymorphisms (SNP)





SNP marker panel selection

- 5,500 SNPs tested in 756 samples collected from MOWI, Salmobreed and Aqua Gen.
- A subset of SNPs (n=114) was identified using the following criteria:
 - SNPs must have high minor allele frequency (MAF >0.45)
 - > 3-4 SNPs from each chromosome
 - 3 SNPs from mitochondrial genome to provide extra assurance for female assignment
- Further reduced to <u>59 SNPs</u> compatible for analysis in a single reaction





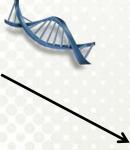
Genotyping protocol



DNA extraction



Sample

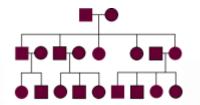




SNP Testing (genotyping)

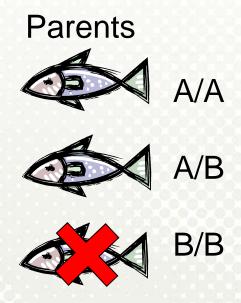


Assignment



Assignment uses the principle of exclusion to identify possible parents:







Development of മ traceability tool using SNPs

Assignment

After identifying candidate parents, exclude impossible parent x parent combinations. For example:

Parents Offspring A/B B/B B/B A/B B/B A/A A/A A/B



Pilot experiment 1





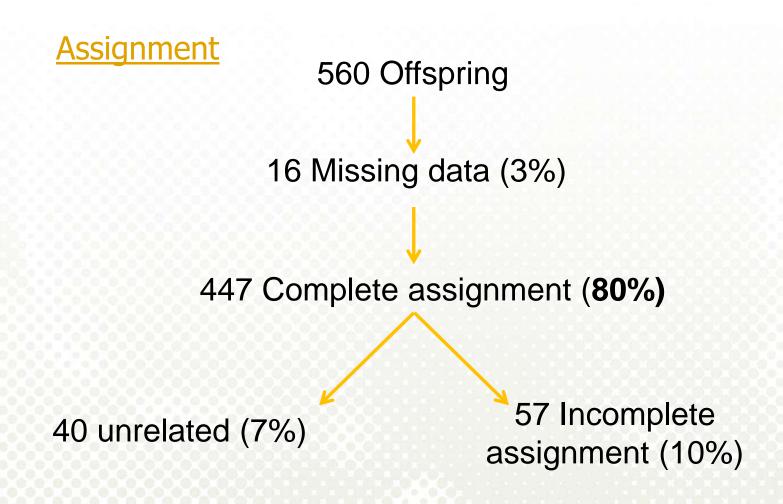


229 Parents

520 Blinded offspring 40 Unrelated fish











<u>Assignment – Continued</u>

- Reduce the percentage of incomplete assignments by
 - including additional information:
 - Sex of parents (utilize mitochondrial SNPs)
 - Parental crossings (reduce possible combinations)
 - include additional SNP sets

Assignment has not been confirmed



Coming activity

- Test assignment with a large excess of potential parents
- Test assignment in blinded material from other breeding companies (Salmobreed, Rauma, Marine Harvest)
- Test ability of assignment panel to differentiate wild and farmed salmon



Thanks

