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Is DNA-testing the Future of Species Control?





Is DNA-testing the future of species control?

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Mixed industrial catches

- Large mixed quantities (>200 tonnes)
- Generally few species
- Uneven distribution of species in catch (hauls/tanks)
- No fixed EU standards (level playing field)
- Current control by manual counting and weighing (buckets)
- Problems with bycatch (e.g. herring)
- Problems with documenting catch
- Minimum reporting 50 kilos

DNA-Mix project, Danish Fisheries Agency, industry and fishermen. Funded by **European** Maritime and Fisheries Fund



Marine Ingredients Denmark









DTU DNA based species-identification –"Barcoding"

- "Barcoding of life" database <u>www.boldsystems.org</u>
- One gene COI (cytochrome c oxidase subunit I)
- 650 bases ACTG.....
- Separates 98% of studied fish species
- 21.073 species (25.05.2022)
- Simple as it relies on categorical differences
- Single species samples (filets, fins) easily identified without taxonomic expertise







Can DNA testing be used for mixed catches?

Challenges:

- -How to sample large inhomogenous mixtures?
- Do some species give more DNA \neq weight?
- Is the precision high enough?

Potential solutions:

- Sample production water on vessel or in factory = more homogenous DNA composition than the fish
- -Calibrate for DNA/weight, with respect to different species
- Test the robustness of inferences with "mock" mixture samples (species proportions and size etc.)





Case 1 Sprat and herring mixed catches

• Experiment:





5 kilos in each bucket





Relationship between weight and DNA fractions (herring)

qPCR-measurements of blood water





qPCR-measurements of discharge water





DNA-fraction measured (μi) = 0.4 (95% CI)

Weight fraction estimated = 0.53 ± 0.07

Weight fraction

Relationship between weight and DNA fractions corrected for relative size of fish (herring larger = less DNA per weight)



qPCR-measurements of discharge water



Fish proportion (based on allometrically scaled weight)

DNA-fraction measured (μ i) = 0.4 (95% Cl) Corrected weight fraction = 0.4 ± 0.05

DTU



Case 2 Bycatch of mackerel in herring fishery





Landing – the unloading process





Mackerel fraction estimated with different methods







Conclusions and what's next

- DNA based species control from production water has high sensitivity and precision = large potential for practical implementation
- Factors like species and relative fish size has to be (and can) be accounted for in relation to DNA proportion
- The process of unloading the fish and reuse of discharge water in the factory is complex and has to be known for DNA testing
- Sampling water from ship tanks before landing may be the best solution
- More industrial scale trials with known weight proportions have to be conducted
- Robustness to factors like maturity, sea-area and time of year should be investigated
- Practical implementation trials can be conducted now using visual and DNA based methods in parallel
- The frequency for updating the relationship between DNA and weight should be assessed
- DNA methods for more complex mixtures are under development



