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Fish Muscle postmortem change and salting performance - Impact of muscle microstructure on salt transfers

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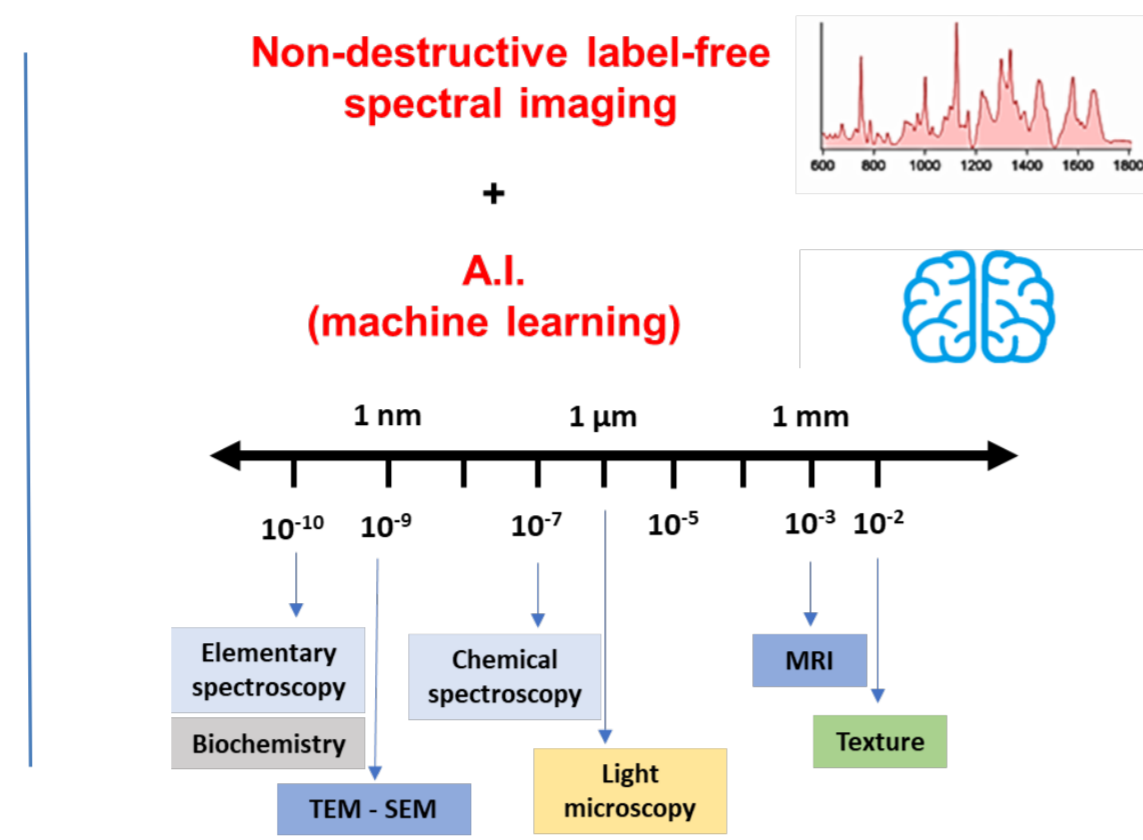
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Objectives

This study aimed to characterize fish muscle change with *postmortem* time, and predict the impact of ageing on salt transfers.

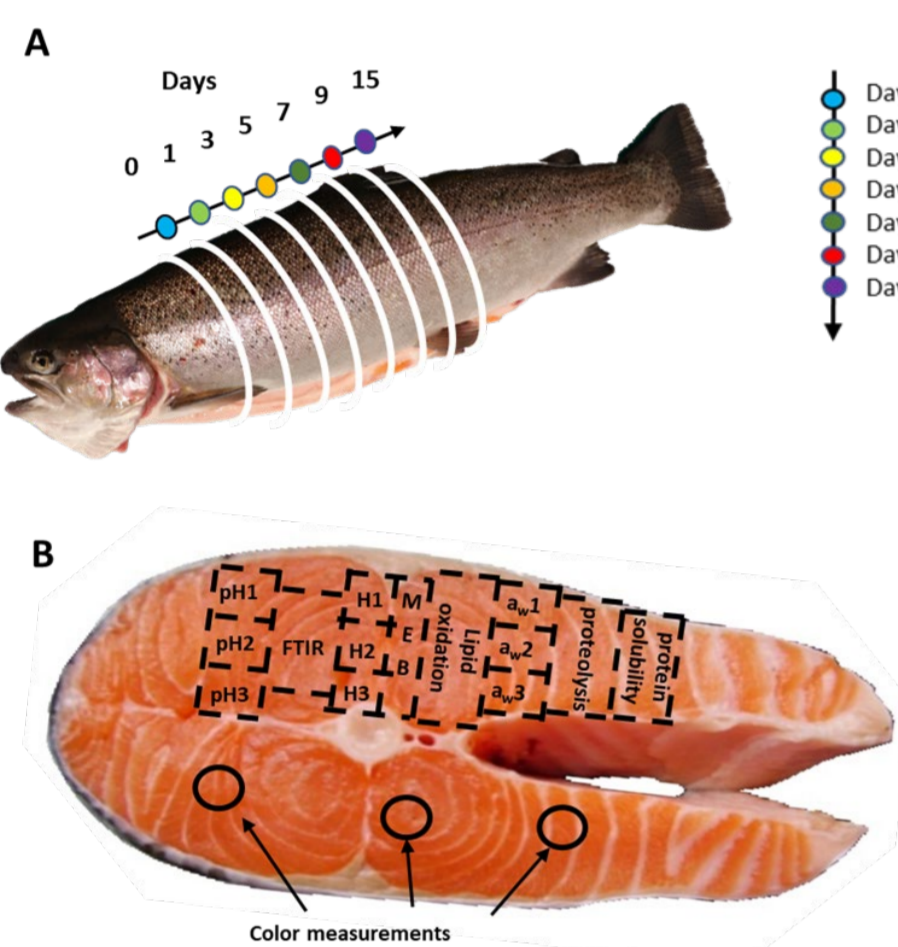


Highlights

- Combinatory approaches of **physicochemical measurements and machine learning** helped characterize the impact of aging and salting in trout muscle.
- Biomarkers accurately **predicted the *postmortem* time** of fish and salt concentrations.
- **Label-free fast characterization** open perspectives to optimize the salting process, improve food quality, and reduce the environmental impacts of salting in the processing industry.

Methods

6 rainbow trouts (3 Kg), from 0 to 15-day *postmortem*. One filet was salted and its contralateral serves as control



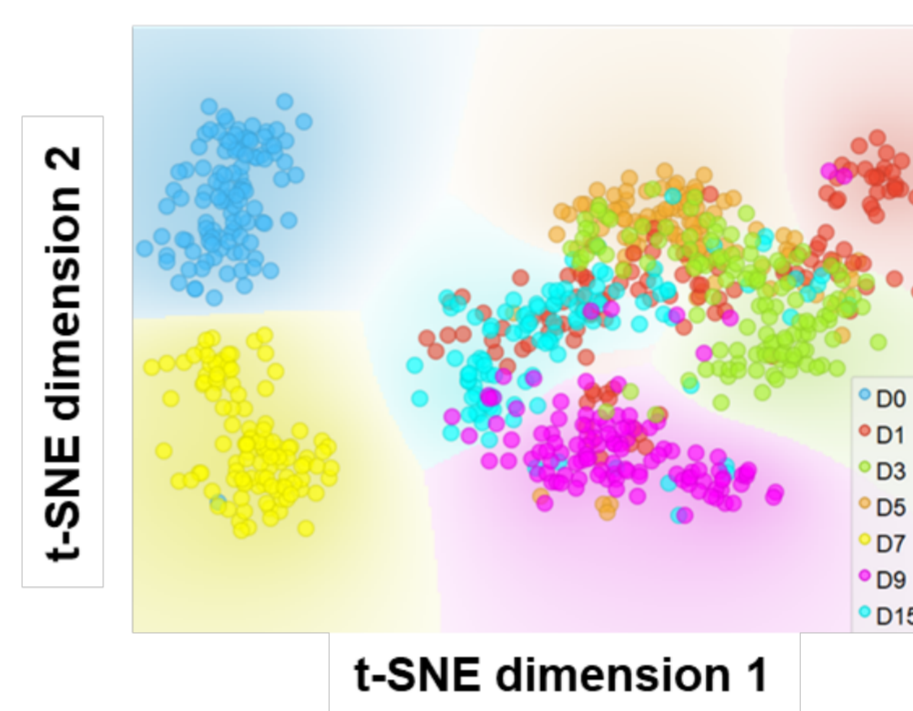
Physico-Chemical Analyses

- pH
- Colour (L, a*, b*)
- Microscopy
- HPLC for Na⁺ & Cl⁻ quantification
- Water activity (a_w)
- Lipid oxidation (T bars)
- Proteolysis
- Solubility

Imaging

- Synchrotron X-ray fluorescence
- MRI of H⁺ et Na²³
- SEM+ EDS
- TEM
- FTIR
- SHG

Machine learning coupled with FTIR revealed biomarkers of *postmortem* time



Wavenumber	Biomarker for	Molecular bond	Biological compounds
1002 cm ⁻¹	D0	Phe, Tyr	aromatic compound, protein
1024 cm ⁻¹	D0	CO	polysaccharides, glycogen
1172-1176 cm ⁻¹	D0	CC of DNA, PO ₂ stretch	DNA, PO ₂ , nucleic acids, carbohydrates
1204-1206 cm ⁻¹	D7	amide III	amide III, collagen
1247 cm ⁻¹	D0	PO ₂	nucleic acids (RNA)
1363 cm ⁻¹	D0	CH ₂ , CC	polysaccharides
1469 cm ⁻¹	D0	CH ₂ bending of lipids	saturated lipids
1490 cm ⁻¹	D7	CC, CH	amide I, fatty acids
1553 cm ⁻¹	D0	amide II	predominantly α-helix of amide II
1635 cm ⁻¹	D0	β-sheet structures of amide I	β-sheets, proportion of secondary protein structures (shoulder peak of amide I)
1670 cm ⁻¹	D7	ν(CC) trans, antiparallel β-sheet structures of amide I	antiparallel β-sheets (shoulder peak of amide I), fatty acids
1712 cm ⁻¹	D0	CO	glycogen, lipid oxidation, unsaturated fatty acids (PUFA)
1760 cm ⁻¹	D0	CO, COO-R	glycogen

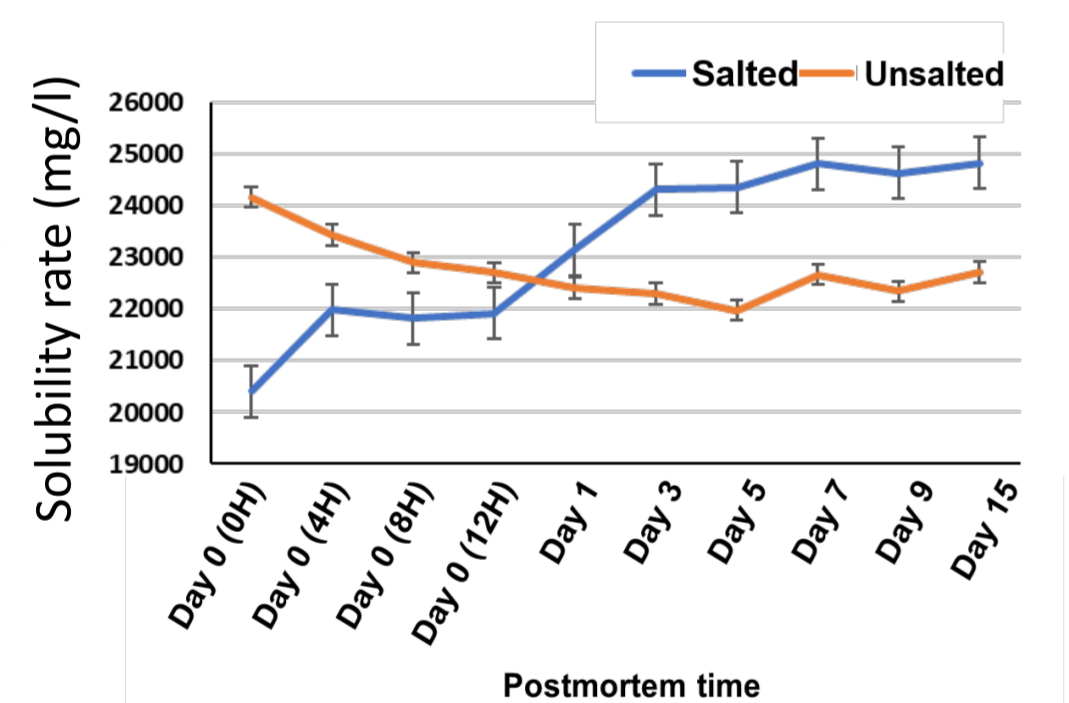
True values	D0	D1	D3	D5	D7	D9	D15	Σ
D0	100.0 %	0.0 %	0.0 %	0.0 %	0.0 %	0.0 %	0.0 %	111
D1	0.0 %	93.1 %	0.0 %	0.7 %	0.0 %	0.0 %	22.1 %	122
D3	0.0 %	0.0 %	82.9 %	27.6 %	0.0 %	0.0 %	0.0 %	129
D5	0.0 %	0.0 %	8.1 %	70.9 %	0.0 %	3.6 %	0.8 %	109
D7	0.0 %	0.0 %	0.0 %	0.0 %	100.0 %	0.0 %	0.0 %	114
D9	0.0 %	3.0 %	0.0 %	0.0 %	0.0 %	91.1 %	6.6 %	113
D15	0.0 %	4.0 %	0.0 %	0.7 %	0.0 %	5.4 %	70.5 %	107
Σ	111	101	111	134	114	112	122	805

Predicted values

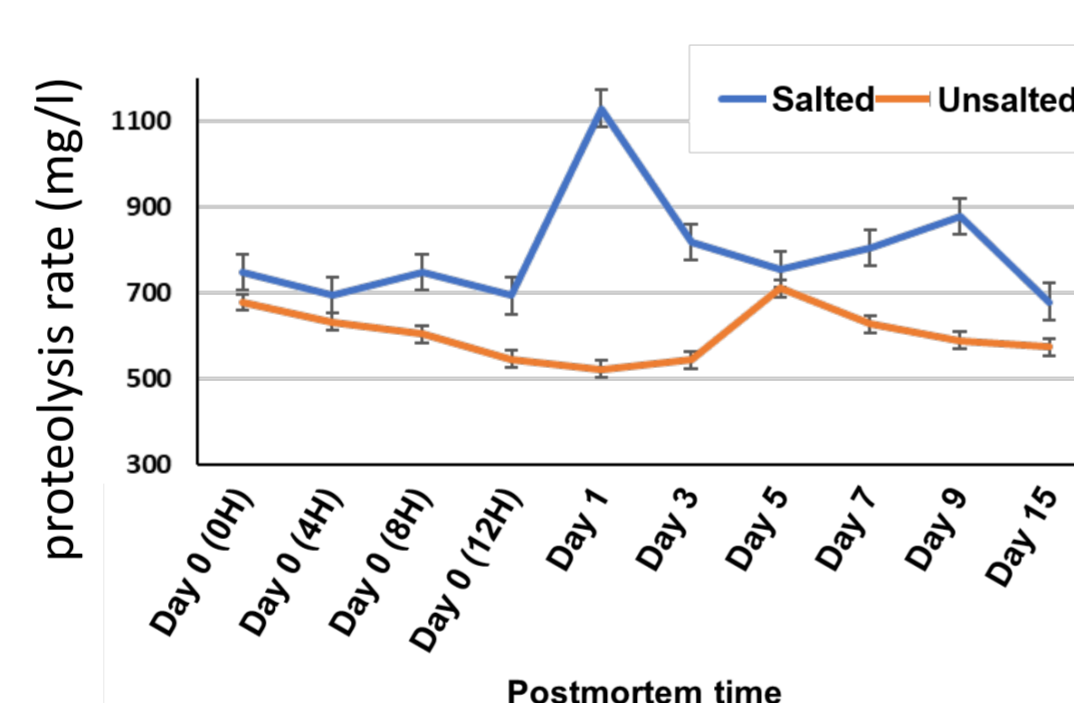
Major physicochemical and structural changes

- In the absence of salt, biochemical analyses revealed little changes over the *postmortem* time. Surprisingly, the proteolysis index and solubility of proteins varied very little during the *postmortem* time. A fast pH drop was observed soon after stunning and bleeding and it stabilized close to 6,5.

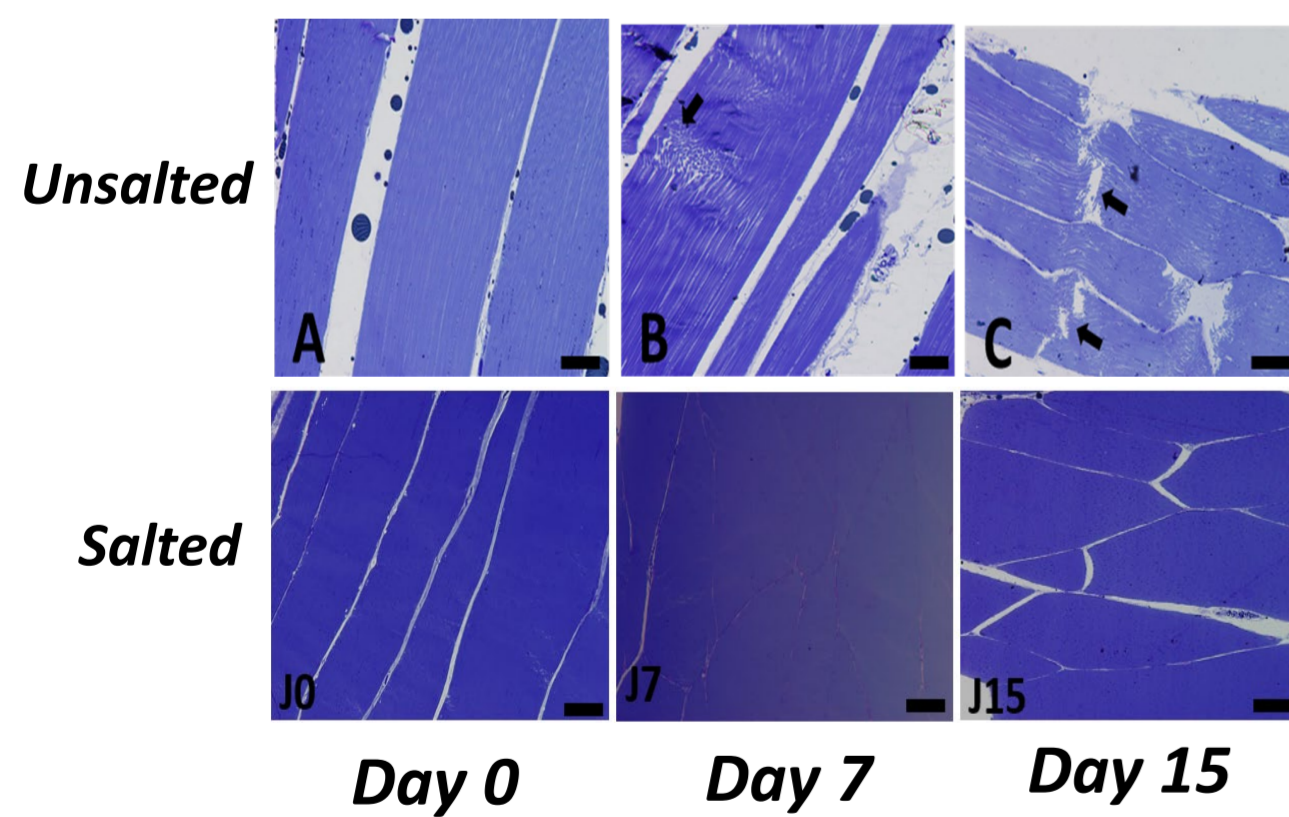
Protein solubility during *postmortem* time



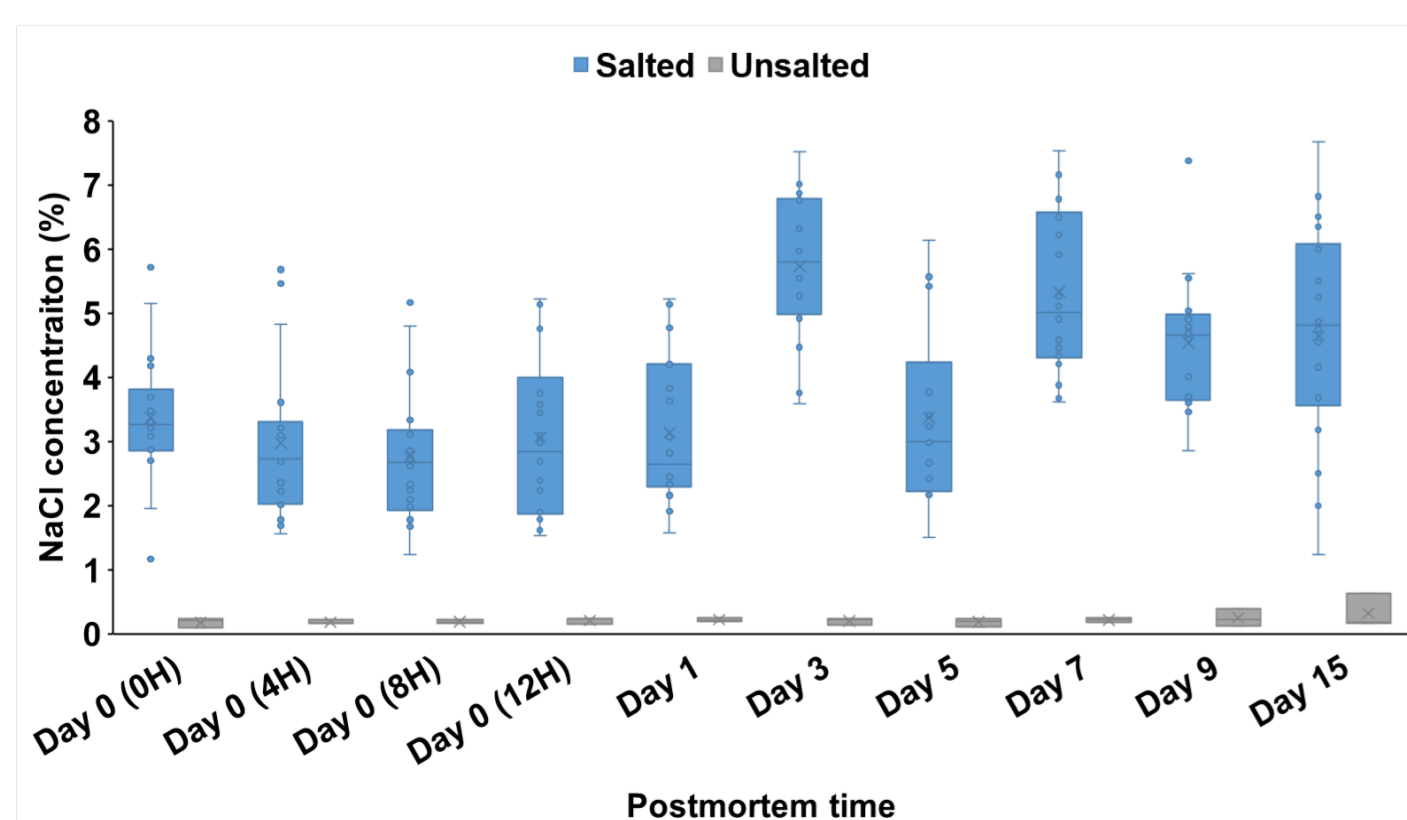
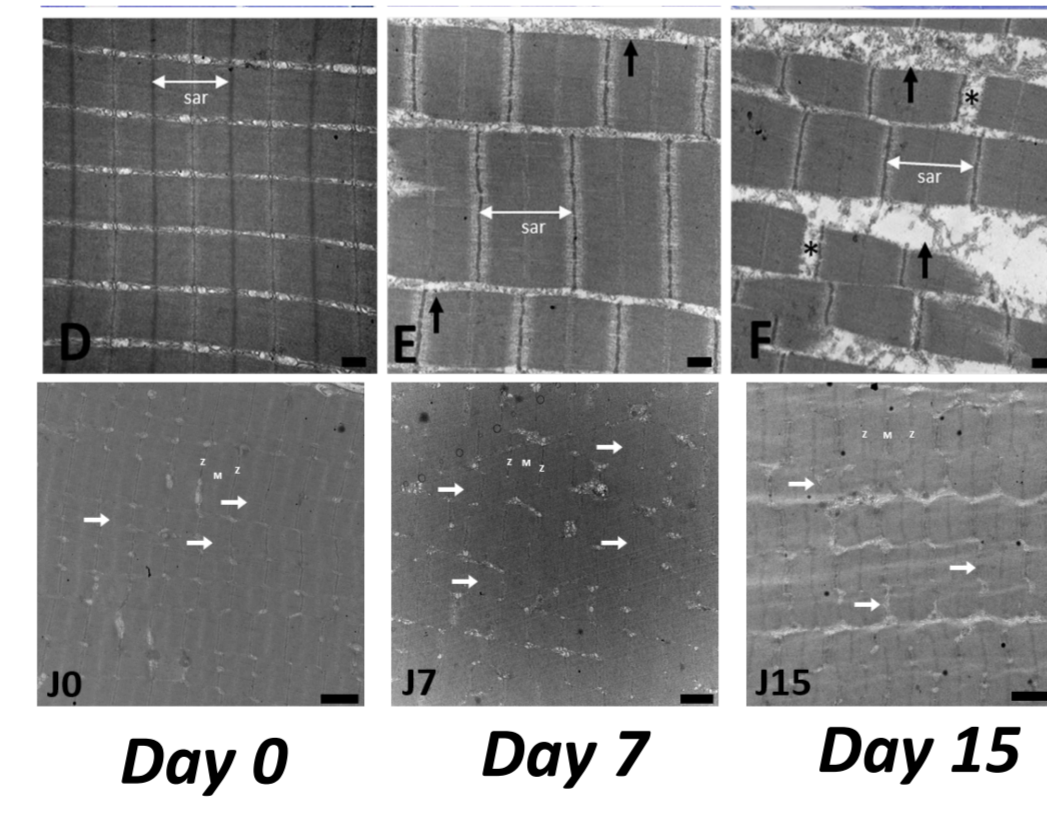
Proteolysis during *postmortem* time



Microstructural changes Light microscopy



Ultrastructural changes Electron microscopy

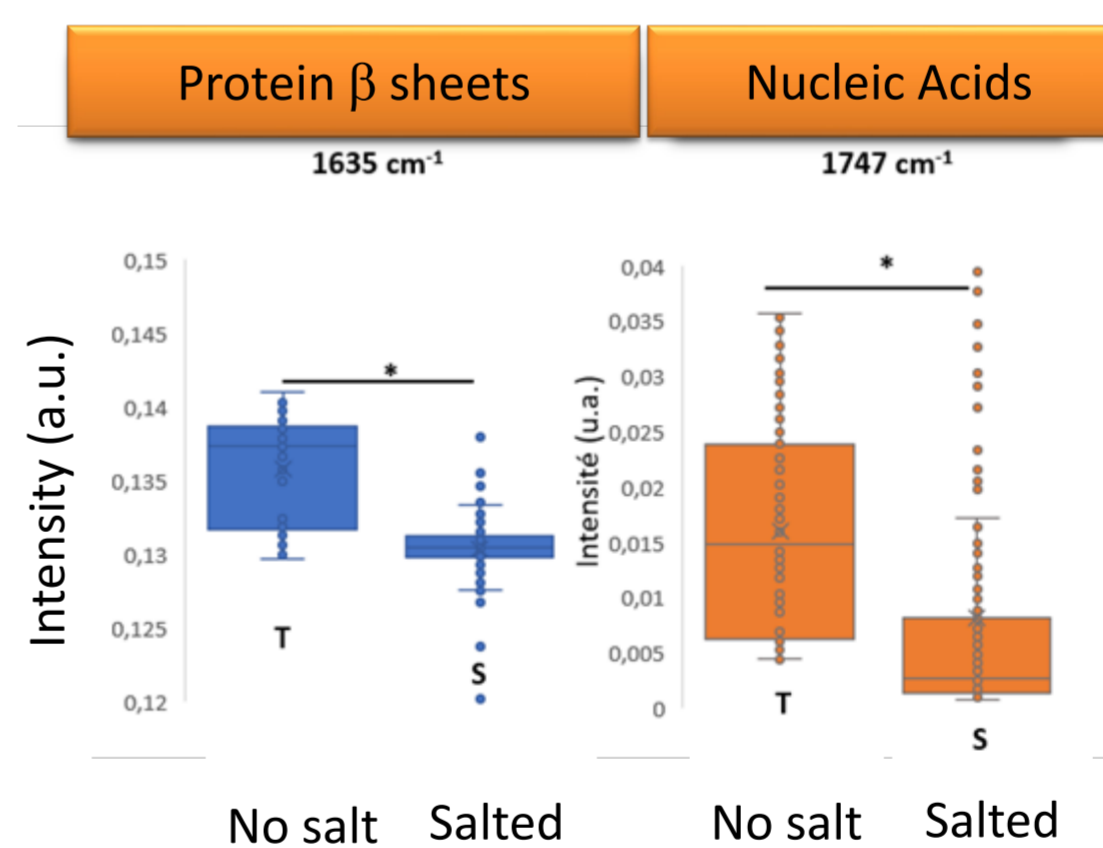


Final salt concentrations in trout muscles increased with advanced *postmortem* times

Salt content increases with *postmortem* time, from 2,5% (salting before 3 days *postmortem*) up to 6% when the filet is salted from 3 days *postmortem*.

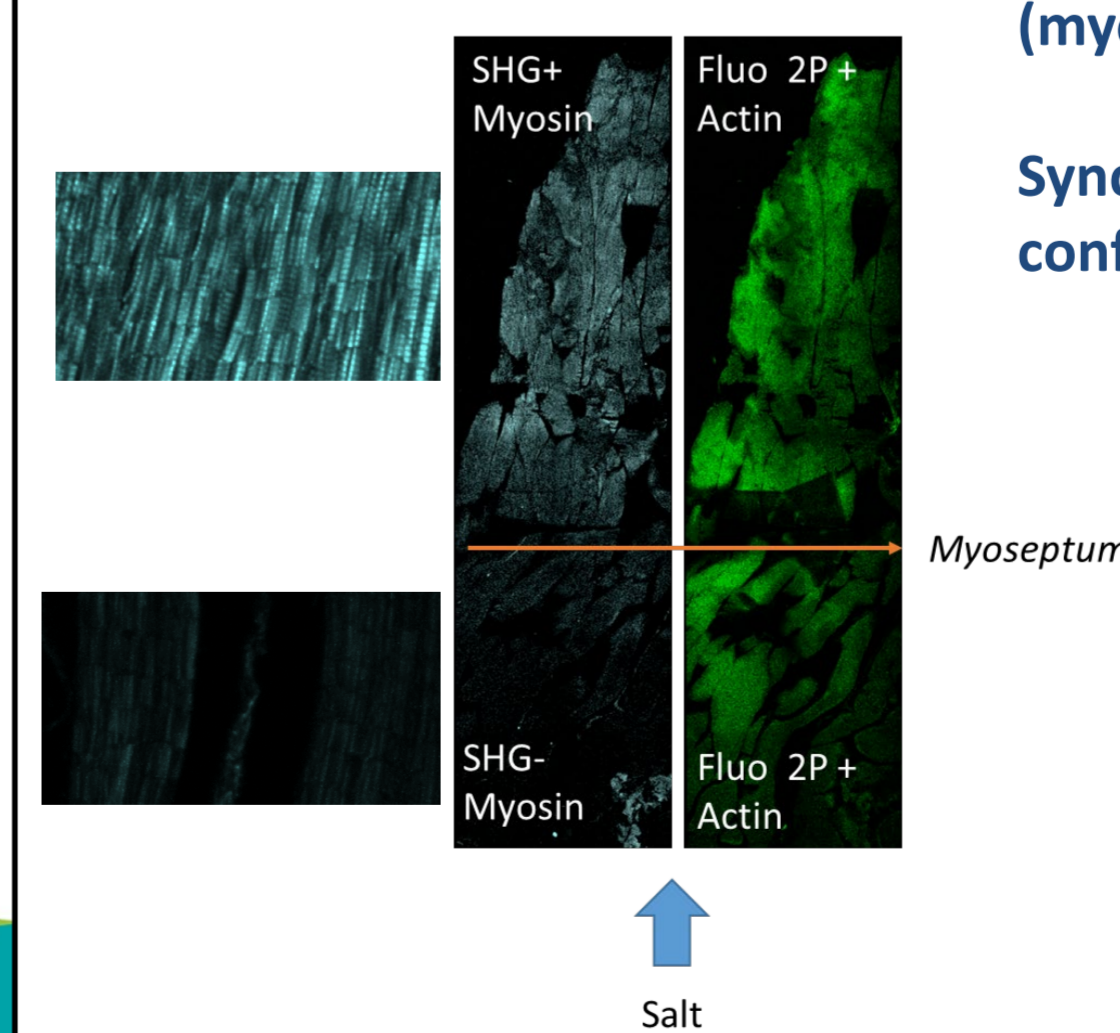
European standard imposes a minimum of 3% final salt concentration

Biomarkers toward the fast characterization of filets



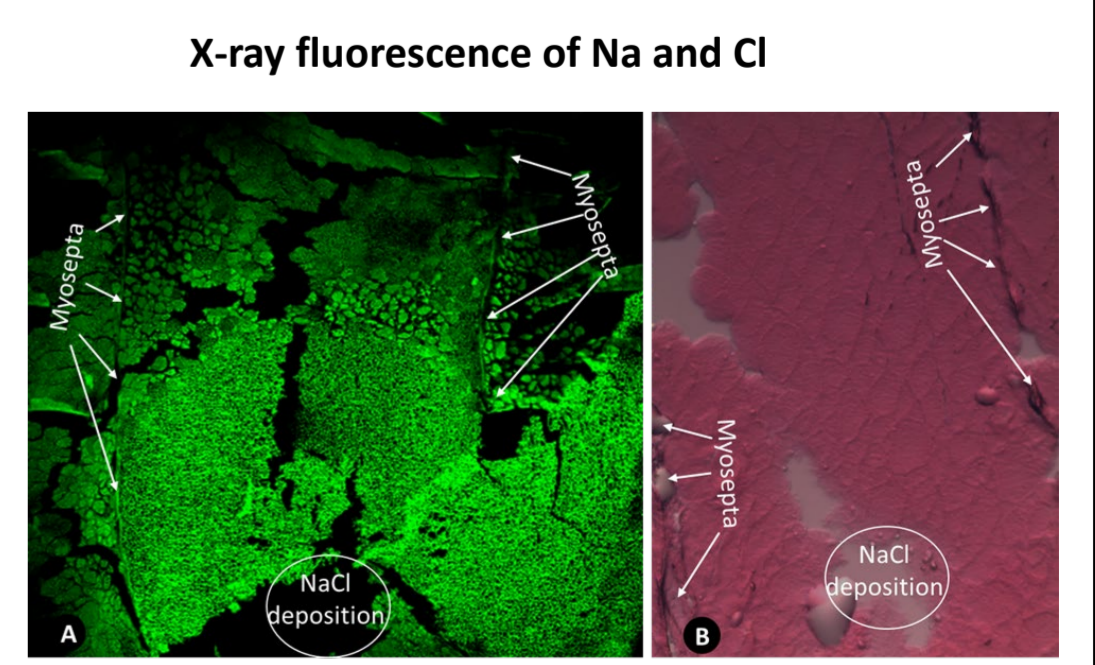
By increasing the ionic strength, salting decreases absorbance at 1635 cm⁻¹ and 1747 cm⁻¹ assigned to β-sheet structure and nucleic acids respectively

SHG microscopy



SHG microscopy suggests that connective tissue (myoseptum) slows down the diffusion of salt.

Synchrotron X-Ray Fluorescence of Na and Cl confirms the barrier effect of myoseptum



Industrial application prospects:

On-line in situ measurements of freshness could be used to adjust salt concentrations to maximize product quality and economic value.