# Non-invasive Assessment of Packaged Cod Freeze-Thaw History by Hyperspectral Imaging

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#### 9 Abstract

10 Freezing, storage and thawing all significantly affect the quality of seafood products. In this article, we explore the potential for estimating these parameters using online optical spectroscopic 11 12 measurements. Fillets of cod were vacuum packed and underwent different programs of freezing, thawing and storage. Hyperspectral imaging of the fillets was performed at each program stage. Both 13 14 single and double-frozen samples are studied. Different freezing and thawing methods are also 15 investigated. Multivariate analysis of the hyperspectral data show that freezing history can clearly be 16 determined on samples in the frozen state and to some degree on samples in the thawed state. Analysis 17 of the data shows a less clear distinction between methods of thawing. Measurement on samples after 18 a year of storage still showed the ability to classify samples based on their freezing history. These results 19 show that hyperspectral technology can be used to evaluate the freeze-thaw history and potentially 20 estimate quality of cod products in a manner that meets the speed and non-invasive requirements of 21 an industrial setting.

22 Keywords: hyperspectral imaging; fish, cod; storage; freezing; classification

## 23 1 INTRODUCTION

24

25 Freezing and thawing can lead to changes in seafood that affect both their sensory properties and 26 market value. Because consumers believe fresh fish to be a superior product (Peavy et al. 1994), it 27 commands a better market price than previously frozen fish. For frozen seafood, the freezing 28 procedure may affect the product; both the rate of freezing (Pan and Yeh, 1993; Chen and Pan, 1997) 29 and the freezing temperature (Mørkøre and Lilleholt, 2007) have been shown to influence changes 30 that occur to fish during freezing, such as water loss, gaping, or toughness. Such changes can alter the 31 perceived desirability and market price of the product. Refreezing of previously frozen samples has 32 been shown to negatively affect sensory properties of fish such as toughness, dryness, fibrousness and 33 fishy smell (Kent et al. 2004). Fish that have been twice-frozen deteriorate more rapidly in sensory 34 quality under long term storage compared to fish that have been frozen only once (MacCallum et al. 35 1966; Desrosier and Tressler, 1977). In particular, if the initial freezing of the twice-frozen samples was 36 performed using poor freezing conditions, this led to a significantly shorter shelf life of the fish 37 (Desrosier and Tressler, 1977). All these reasons highlight the importance for companies to have38 control over the freeze-thaw history of their seafood products.

39 Ensuring the product delivered to the consumer is the product promised is vital for companies to 40 protect brand reputation. A challenge to the fish processing industry is that a delivery may not consist 41 of fish that have experienced identical handling. A delivery may be comprised of catches from several 42 different vessels, each with different equipment capabilities and handling procedures. Furthermore, 43 previously frozen fish may be purposely mixed in with fresh fish due to the higher market price. These 44 issues make detection of fish of unacceptable quality by spot checks challenging. Therefore, there is a 45 demand for evaluation methods rapid enough to be performed on every fish. Such measurements can 46 potentially be used to detect fraud (e.g. frozen-thawed fish sold as fresh) and measuring if a product 47 has been handled according to the standard set by the buyer. Investigation of double freezing is 48 especially relevant for products based on fish frozen at sea. Here, the fish is thawed before processing 49 (e.g. filleting) and subsequently frozen again as a finished product and proper freezing and thawing 50 parameters are vital if high quality is to be maintained. In this work, we investigate if measurements 51 can be made to determine the full freezing history of double-frozen products.

52 Previous research has investigated a variety of techniques to distinguish between fresh, once-frozen 53 and twice-frozen seafood. Differential scanning calorimetry found a difference in the denaturation 54 temperatures between fresh, once-frozen and twice-frozen gilthead seabream (Matos et al. 2011). 55 Time domain reflectometry to estimate dielectric constants of minced cod was able to differentiate 56 between once-frozen and twice-frozen samples with good accuracy (Kent et al. 2004). Similarly, 57 dielectric measurements in the frequency domain showed the ability to distinguish between fresh, 58 once-frozen and twice-frozen sea bass (Vidacek et al., 2008). Raman spectroscopy was able to reliably 59 identify fresh, once-frozen, and twice-frozen cod by measuring on fat extracted from thawed samples. 60 (H.M. Velioğlu et al., 2015). While all these techniques have proven successful in a lab environment, 61 they take from minutes to hours to perform and frequently require alteration of the sample (e.g., 62 mincing, extraction, etc.). Therefore, the measurement time and extensive sample handling they 63 require make them not suitable for high-throughput industrial screening, where approximately one 64 fillet per second needs to be scanned.

65 In this article, hyperspectral imaging was assessed as a method to evaluate the freeze-thaw history of 66 seafood. This technology has been applied previously to perform many different types of quality 67 control in a wide range of foods (Sun 2010, Siche 2016). Hyperspectral measurement is appealing 68 compared to other forms of spectroscopy, such as Raman or nuclear magnetic resonance, because it 69 is both non-invasive and rapid. Optical spectroscopy is a well-established method for material 70 characterization. When light interacts with a material, some wavelengths will be absorbed while others 71 will pass through the material. The amount of light absorbed by or transmitted through a sample 72 depends on both its structure and its composition. Therefore, by analyzing how a sample absorbs light 73 at different wavelengths, information can be gained on its properties. Hyperspectral imaging is a form 74 of spatially resolved optical spectroscopy. In traditional imaging, each pixel has an intensity associated 75 with a single wavelength for gray scale images or intensities at three different wavelengths for color 76 images. In hyperspectral imaging, every pixel is associated with a plurality of wavelengths that can be 77 used to evaluate the structure and composition of that location in the sample.

The hyperspectral images analyzed in the present study were acquired using the interactance acquisition mode (Schaare and Fraser, 2000). Here, light travels through the sample and the reemitted light is measured to determine which wavelengths have been absorbed. Interactance spectroscopy differs from transmission spectroscopy in that it measures light that has traveled back to the illuminated surface as opposed to light that has passed through the sample. The method is useful for

- 83 samples where complete transmission is not possible, for example if the sample is very thick or where
- 84 opaque constituents are present. Another practical advantage is that one-sided measurement enables
- 85 imaging on conveyor belts where only one side of the sample is accessible. Interactance is also useful
- 86 when the sample thickness is variable and would lead to an inconsistent transmission path length.
- 87 This work builds upon previous research where it was observed that fresh and previously frozen
- 88 samples could be clearly distinguished using hyperspectral measurements on thawed samples
- 89 (Sivertsen et al. 2011, Zhu et al. 2013). There are several aims to this study. First, the scope of
- 90 investigation is expanded to evaluate whether these techniques can distinguish between fresh, once
- 91 freeze-thawed and twice freeze-thawed products. Secondly, we evaluate if the freeze-thaw history
- can be estimated based on measurements on samples in the frozen state. Furthermore, this study
   also seeks to determine whether hyperspectral measurements can identify the freezing and thawing
- also seeks to determine whether hyperspectral measurements can identify the freezing and thawingprocedures the samples had experienced.

# 95 2 MATERIALS AND METHODS

- 96 2.1 HYPERSPECTRAL IMAGING EQUIPMENT
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98 Figure 1 shows a diagram of the hyperspectral imaging setup.

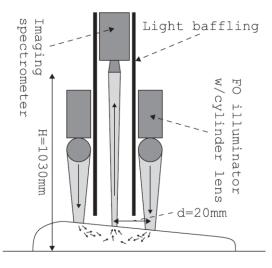


Figure 1: Diagram of the hyperspectral imaging setup. Note that to save space, the vertical scale of the figure has been
 compressed

102 The illumination source was a pair of fiber optic line lights, each 200 mm wide and powered by three 103 150 W halogen lamps. To maximize the amount of NIR light directed into the fiber optic system, lamps 104 with a special aluminum coating were used (International Light Technologies, lamp model L1090). 105 Cylindrical lenses mounted in front of the line lights focused the light into two 10 mm thick parallel 106 lines, 40 mm apart. The camera line of view was placed between the light lines, resulting in a 20 mm 107 distance to each of the lines, as shown in Fig. 1. Previous work has shown that dual illumination of the 108 sample improves signal to noise compared to a single illumination source, as well as resolves the 109 problem of being unable to obtain a signal at the edges of the sample (Sivertsen et al. 2009; Wold et 110 al. 2006). Two aluminium light baffles painted black were used to block light which would otherwise 111 illuminate the sample directly in the camera line of view. The fiber optic line lights and detector were 112 mounted at heights of 150mm and 1030 mm respectively above the conveyor belt. The detector used was a VNIR-1024 from Norsk Elektro Optikk (Skedsmokorset, Norway). The hyperspectral cameraoperates in the VIS-NIR range from 430 to 1000 nm.

115 The hyperspectral camera field of view is 0.56 mm x 300 mm. Each pixel is 0.28 mm x 0.56mm in size and light from that area is measured in the 430-1000 nm spectral region with approximately a 5.4 nm 116 117 resolution. The region of detection is centered parallel between the two illumination lines. Imaging of 118 the samples was performed on a conveyor belt traveling at 40 cm/s, a rate that meets the industrial 119 production requirement of approximately one fillet per second. Each sample was scanned line by line 120 at 400 frames per second as it moved through the hyperspectral camera field of view. The data is 121 stored as a hyperspectral image  $R_i(\lambda, x, y)$  consisting of successive frames  $F(\lambda, x)$ . Calibration of the 122 system was performed using a 300 mm square Teflon target of 25mm thickness. The average reference 123 frame  $R_a(\lambda, x)$  was calculated from one hundred successive frames of the Teflon target. The absorption of the interactance images are calculated as  $I(\lambda, x, y) = -\ln(R_i(\lambda, x, y)/R_a(\lambda, x))$  where 124 125  $R_i(\lambda, x, y)$  is the hyperspectral image of the packaged cod loin.

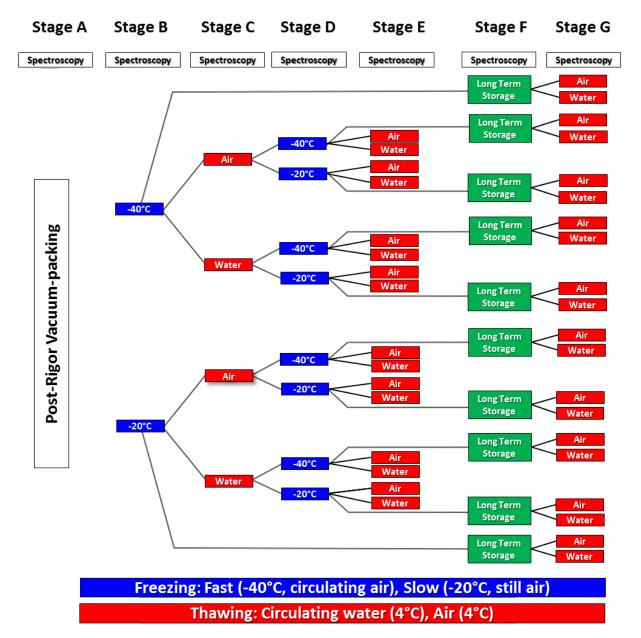
#### 126 2.2 FISH SAMPLES

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128 Atlantic cod for the study were procured by capture based aquaculture. Here, wild cod are captured 129 with seine nets in the spring and then held in sea cages for later slaughter. This both minimizes stress 130 on the fish and allows for better control of the slaughter process, enabling the supply of high quality 131 fish in the months traditionally out of season by conventional fishing. The cod were stored and fed at 132 the Tromsø Aquaculture Research Station, Norway and were killed by a blow to the head and 133 immediately gutted. They were bled for 30 minutes, iced and transported to Nofima, where they 134 were kept on ice for two days prior to filleting. These procedures helped ensure that uniformly high 135 quality fish were used in the study to minimize potential confounding effects resulting from defects (e.g. inadequate bleeding) caused by poor sample handling. The back loin was used for the 136 137 experiments. Each loin was cut in 2 or 3 pieces  $(179 \pm 32 \text{ g})$ , depending on the size of the fish. The 138 loin-pieces were then vacuum-packed (pressure 10 millibar) in plastic pouches (20 µm polyamide 139 inside layer and 70 µm polyethylene outside layer) and kept on ice until the first imaging and 140 subsequent freezing and frozen storage. 141 The two hundred vacuum packed loin pieces of cod were divided into forty groups (n=5). The groups 142 each underwent a different program of freezing, thawing and measurement; the samples were kept 143 vacuum packed for the entire program to minimize the influence of additional factors besides freezing, 144 thawing and frozen storage. Two modes of sample freezing were used: blast freezing at -40° C (fast) 145 and freezing in still air at -20°C (slow). Two modes of sample thawing were used: in 4° C circulating 146 water (fast) and at 4°C by gently circulating air in a climate controlled cabinet (slow). The different 147 freeze-thaw procedures and measurement stages are shown in Figure 2. Every sample in a group

148 underwent the same sequence of freezing, thawing and measurement as the others in the group. All 149 frozen samples were brought to equilibrium at -20° C before hyperspectral measurement to avoid 150 temperature effects producing a bias in the measured spectra.

151 In stage A, samples were vacuum-packed and hyperspectral measurements were performed on all 152 samples. In stage B, the samples were frozen using one of the two different freezing protocols. After 153 freezing, samples were all kept at -20 °C for two days and then hyperspectral measurements were 154 repeated on all the samples in the frozen state. Some samples were placed in long term storage at -155 20°C (Stage F) and the remaining samples moved onto Stage C. Here, they were thawed using the two 156 different thawing protocols and hyperspectral measurement performed. In stage D, the remaining 157 samples were refrozen. Some samples maintained the same freezing protocol for the second freezing 158 as the first freezing. Others changed freezing protocols between the two freezings -- i.e. samples first 159 frozen using the fast protocol were refrozen using the slow freezing protocol and vice versa. Again, the 160 samples were kept at -20°C for two days and then the twice-frozen samples underwent hyperspectral 161 measurement in the frozen state. Again, a set of samples were removed for long term storage. The 162 rest of the samples moved onto Stage E, where they were thawed for hyperspectral measurement. In 163 Stage E, there existed a sample group that covered every permutation of fast and slow freezing and 164 fast and slow thawing for twice-frozen samples. For Stage F, hyperspectral measurements were made 165 on the samples in the frozen state at the end of storage. The long term storage samples were stored 166 in a freezer at -20 °C for 12 months. Finally in Stage G, the samples were thawed and the final 167 hyperspectral measurements were performed. As with Stages C and E, in Stage G a complete set of samples existed that covered all the possible permutations of freezing and thawing protocols for both 168 169 once and twice-frozen samples.



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Figure 2: Program of Analysis. Stage A - vacuuming packing and measurement on the samples in the fresh state. Stage B first freezing of samples and hyperspectral measurement on the samples in the frozen state. Stage C - first thawing of samples
 and hyperspectral measurements of the samples in the thawed state. Stage D - second freezing of samples and hyperspectral
 measurement on the samples in the frozen state. Stage E - second thawing of samples and hyperspectral measurements of

175the samples in the thawed state. Stage F – hyperspectral measurement of samples in the frozen state that have undergone176long term storage frozen Stage G – hyperspectral measurement of samples in the thawed state that have undergone long177term frozen storage.

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#### 179 2.3 DATA ANALYSIS

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181 Extraction of hyperspectral data for analysis was performed using IDL (Exelis Visual Information Solutions, Bracknell, United Kingdom). Regions of interest were manually selected for each sample. 182 183 Regions were selected in the center of the sample, avoiding areas with sample defects (e.g. blood clots) 184 or image aberrations (e.g. reflections, distortions, etc.). All spectra from the region were exported from 185 IDL in the raw form and further analysis was performed using the R programming language (Free 186 Software Foundation Inc., Boston, MA). Spectra were area normalized, averaged to a single spectrum 187 for each sample and the first derivative taken. Taking the first derivative makes subsequent analyses 188 more sensitive to small differences between the spectra as well as to correct for baseline effects. No 189 smoothing was applied. Principal component analysis (PCA) (Wold et al. 1987) was then performed 190 using three different approaches: examining the entire spectrum, examining only the region of 450-191 600nm, which is associated with blood in the fish (Olsen and Elvevoll, 2011), and examining the region 192 of 900-990 nm, which is associated with water (Hale and Query, 1973; Pope and Fry, 1997). The spectra 193 were analysed using the different region selections in order to evaluate whether a particular sample 194 constituent was predominant in the ability to assess a samples freeze-thaw history. As using 195 extraneous information in multivariate analysis typically leads to worse predictions (Anderssen et al., 196 2006), it is considered good practice to limit analysis to wavelengths known to be associated with the 197 effect to be measured. For example, previous research found that limiting analysis of VIS-NIR 198 measurements on cod to the visual region (450-700 nm) improved prediction results for refrigerated 199 storage time (Nilsen et al. 2002) and differentiation between fresh and once-frozen samples in the 200 thawed state (Sivertsen et al. 2011).

201 The spectra were scaled and centered during principal component analysis. After principal component 202 analysis, classification was performed using k-nearest neighbors classification (Cover and Hart, 1967). 203 The k-nearest neighbors (kNN) algorithm is a form of supervised learning, where a set of samples with 204 known classifications are used to predict the classification of new samples. New samples are predicted 205 by assigning a classification based on the classifications of the nearest samples to the new sample in 206 the variable space. K-nearest neighbors cross-validation was performed here on the full set of principal 207 components using 3 nearest neighbors by the "Leave One Out" method; one sample was left out of 208 the data set and the remaining samples were used to predict its classification. This was repeated for 209 all samples.

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# 211 **3 Results**

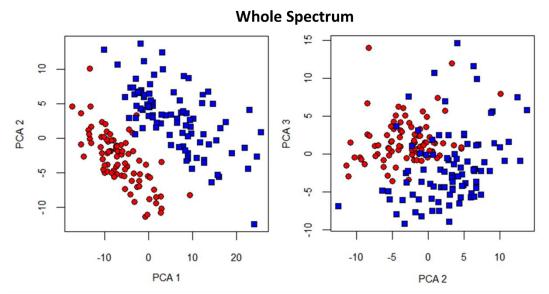
212 **3.1** DIFFERENTIATION OF FREEZING METHOD

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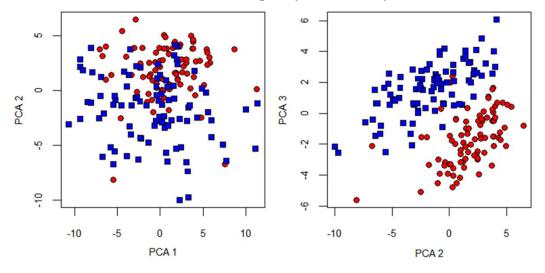
Figure 3 shows the result of principal component analysis on the once-frozen samples measured in

the frozen state. Table 1 contains information on k-nearest neighbors accuracies. Analysis on the

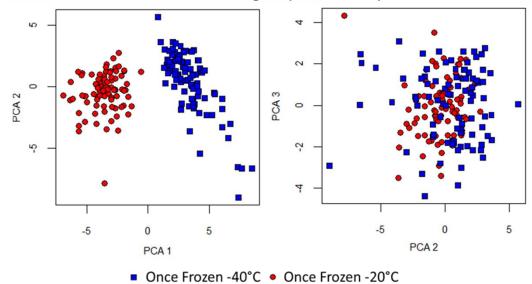
- 216 entire spectrum shows good separation between the two freezing methods. For analysis performed
- 217 using the 450-600 nm region, while clustering is less prominent in the first principal component, the
- two groups can be distinguished visually when examined using the second and third principal
- components. Analysis using the water region of the spectra showed the best separation; the two
- 220 freezing methods show clear separation in the first principal component. K-nearest neighbors
- analysis showed the two groups could be classified with excellent accuracy. The kNN prediction for
- the entire spectrum identified both freezing protocols with 100% accuracy. Using the 450-600 nm
- region, the -40°C samples were predicted with 98% accuracy and the -20°C samples with 99%
- accuracy. For the analysis using only the water region of the spectrum, again both freezing protocols
- could be predicted with 100% accuracy.



Blood Region (450-600 nm)



Water Region (900-990 nm)



226 Figure 3 Principal component analysis on once-frozen samples measured in the frozen state

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230 Table 1: K-nearest neighbors classification accuracies of freezing protocol for once-frozen samples in the frozen state

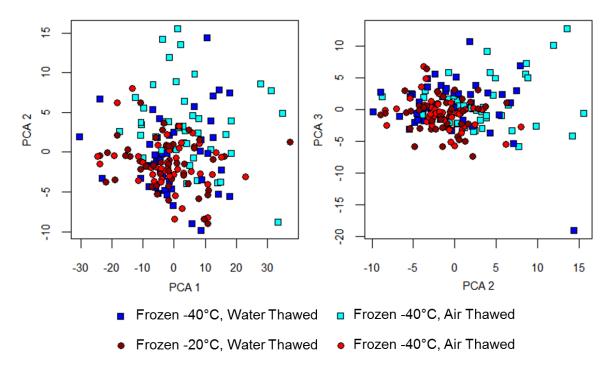
Sample Type	Full Spectrum	450-600 nm	900-990 nm
Frozen -40°C	100%	98%	100%
Frozen -20°C	100%	99%	100%

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#### 3.2 INFLUENCE OF FREEZING AND THAWING METHOD IN ONCE-THAWED SAMPLES

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234 Figure 4 shows the PCA scores plot of the spectra on the once-frozen samples in the thawed state using 235 the entire spectrum for analysis. Table 2 contains the k-nearest neighbor classification accuracy for the 236 thawed samples by freezing protocol, by thawing protocol, and by both freezing and thawing protocol. 237 Regardless of the region of the spectrum used for analysis, there appeared to be no visually consistent 238 grouping of the samples by either freezing method or thawing method used. K-nearest neighbor 239 classification of the samples by freezing protocol showed poor classification, ranging from 38% to 58% 240 regardless of which region of the spectrum had been used for analysis. K-nearest neighbor 241 classification by thawing method performed marginally better, with prediction accuracies ranging from 242 45% to 64%. Classification by both freezing protocol and thawing method simultaneously performed 243 very poorly, with accuracies ranging from 9% to 53%.





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247 Table 2: K-nearest neighbors classification accuracies for freezing protocol for once-thawed samples measured in the thawed

248 state, for thawing protocols of once-thawed samples measured in the thawed state, and for both freezing and thawing

249 protocol for once-thawed samples measured in the thawed state

Sample Type	Full Spectrum	450-600 nm	900-990 nm
Frozen -40°C	58%	56%	53%
Frozen -20°C	38%	38%	47%
Thawed Air	60%	64%	57%
Thawed Water	45%	51%	50%
Frozen -40°C, Thawed Air	51%	53%	42%
Frozen -40°C, Thawed Water	20%	20%	26%
Frozen -20°C, Thawed Air	22%	22%	9%
Frozen -20°C, Thawed Water	31%	27%	36%

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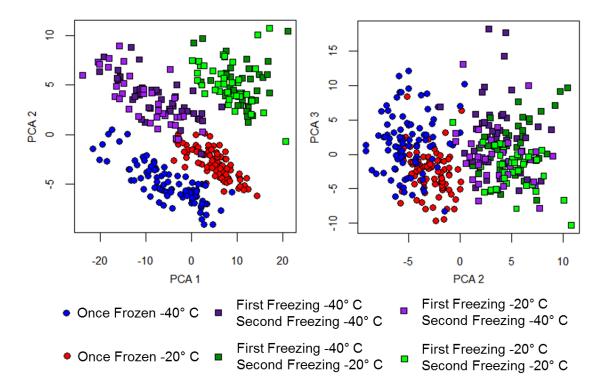
#### 251 3.3 DIFFERENTIATION OF ONCE AND TWICE-FROZEN SAMPLES

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253 Figure 5 shows the PCA of the once and twice-frozen samples in the frozen state for the entire 254 spectrum. Table 3 shows the k-nearest neighbor classification accuracies for once and twice-frozen 255 samples, for freezing protocol in once and twice-frozen samples evaluating only the last freezing for 256 the twice-frozen samples, and for freezing protocol in once and twice-frozen samples evaluating both 257 freezings for the twice-frozen samples. The samples could be classified into once-frozen and twice-258 frozen with good accuracy, with success rates ranging from 97%-100%. Similarly, classification of the 259 samples by freezing protocol produced very good results. K-nearest neighbor prediction using the 260 entire spectrum had almost perfect classification. Analysis using the blood region of the spectrum was still very good, but had slightly more misclassified samples. Analysis using the water region alone 261 accurately identified the freezing protocol for the once-frozen samples, but had a few 262 263 misclassifications in the twice-frozen samples.

For all approaches, the second freezing in the twice-frozen samples appeared to be the dominant effect. Reliable separation of twice-frozen samples by their initial freezing protocols did not appear to be possible. Accuracy of prediction for first freezing protocol on the twice-frozen samples ranged from 40% to 67%. For context, completely random prediction of the four different possible freezing histories would produce results with roughly 25% accuracy.

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271 Figure 5: Principal component analysis for once and twice-frozen samples measured in the frozen state

272 Table 3: K-nearest neighbors classification accuracies for once and twice-frozen samples measured in the frozen state, for

freezing protocol in once and twice-frozen samples measured in the frozen state, and for freezing protocol in once and twice-

274 frozen samples. For twice-frozen samples, both first and second freezing protocol were evaluated for classification

Sample Type	Full Spectrum	450-600 nm	900-990 nm
Once-frozen	100%	99%	100%
Twice-frozen	98%	97%	95%
Once-frozen -40°C	100%	99%	100%
Once-frozen -20°C	100%	97%	100%
Twice-frozen, last freezing -40°C	95%	86%	92%
Twice-frozen, last freezing -20°C	100%	95%	97%
Once-frozen -40°C	100%	99%	100%
Once-frozen -20°C	100%	97%	100%
Twice-frozen -40°C, -40°C	50%	40%	63%
Twice-frozen -40°C, -20°C	54%	56%	59%
Twice-frozen -20°C, -40°C	60%	50%	53%
Twice-frozen -20°C, -20°C	67%	59%	62%

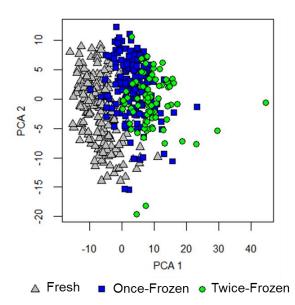
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#### 276 3.4 DIFFERENTIATION OF FRESH, ONCE-THAWED AND TWICE-THAWED SAMPLES

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Figure 6 shows the analysis of fresh, once-thawed and twice-thawed samples. Table 4 contains the knearest neighbor classification accuracies for fresh, once-thawed and twice-thawed samples. These groups contain all possible permutations of fast and slow thawing parameters for the once and twice thawed samples. As was seen previously (Sivertsen et al., 2011) the fresh and thawed samples can be readily distinguished from one another. In all classifications, fresh samples could be reliably distinguished from thawed samples. Analysis of the entire spectrum produced the best classification of the thawed samples, with 98% and 93% correct classification for the once and twice-thawed

- samples. Analysis using the water region produced the worst classification, with 91% and 75% accuracy
- 286 for once and twice-thawed samples respectively.



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288 Figure 6 Principal component analysis for fresh, once-frozen and twice-frozen samples in the thawed state

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Table 4: K-nearest neighbors classification accuracies for fresh, once-thawed and twice-thawed samples measured in the
 thawed state

Sample Type	Full Spectrum	450-600 nm	900-990 nm
Fresh	100%	100%	99.5%
Once-thawed	98%	96%	91%
Twice-thawed	93%	81%	75%

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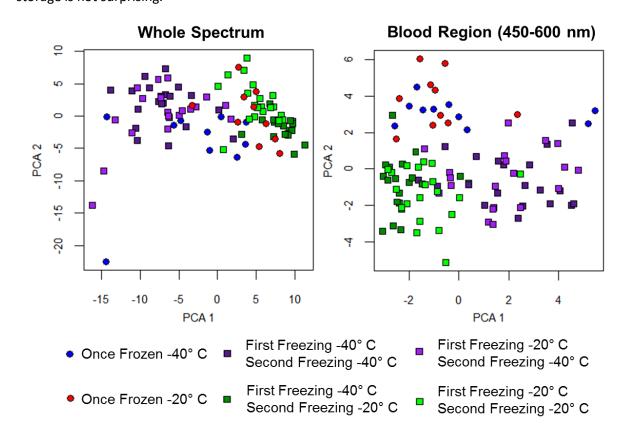
# 293 3.5 EFFECTS AFTER LONG TERM FROZEN STORAGE

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295 For the samples that underwent long term frozen storage, due to changes in the illumination source 296 between the initial measurements and the measurements made after the storage period, the data 297 from the two cannot be directly compared. Therefore, measurements of the long-term storage 298 samples were only compared to one another, and not compared to measurements performed before 299 the long-term frozen storage. Figure 7 shows the PCA analysis for the stored samples measured in the 300 frozen state using the entire spectrum; Table 5 contains the k-nearest neighbors prediction accuracies 301 for the stored samples in the frozen state by number of freezing cycles and by freezing protocol. Figure 302 8 shows the PCA analysis of thawing history for the stored samples measured in the thawed state, 303 Table 6 contains the k-nearest neighbor prediction accuracies for the stored samples in the thawed 304 state by number of freeze-thaw cycles and by freezing parameters.

305 Compared to the results on samples measured before frozen storage, there is a less clear distinction 306 between the different freezing protocols, but clustering can still be observed in frozen samples that 307 have undergone long term frozen storage. Once and twice-frozen samples can be differentiated 308 between with 85-95% success rate when either the entire spectrum or the 450-600 nm region is used. 309 Analysis on the water region alone produced significantly worse classification for the once-frozen samples, but this may be an artifact of the small sample set. The last freezing protocol could also be identified using the entire spectrum with good results on the frozen samples after long term frozen storage. Analysis using the blood region or the water region alone produced significantly worse classification results for the freezing protocol, particularly in the once-frozen samples. As with the samples that had not undergone long term frozen storage, identification of the first freezing protocol for twice-frozen samples was not possible.

316 Analysis of the thawed samples showed the ability to predict the freezing history of the samples 317 using the entire spectrum. Classification of once-thawed samples using the blood or water regions 318 alone performed poorly. The classification of the twice-thawed samples generally performed well, 319 but the classification of the once-thawed samples performed significantly worse. This may simply be statistical noise due to the low number of once-thawed samples; there were eighty twice-frozen 320 321 samples but only twenty once-frozen samples that were placed in long term storage. The technique 322 also produced poor classification of samples by freezing protocol. Given the inability to determine 323 freezing protocol on the thawed samples before storage, the inability to do so after long term frozen 324 storage is not surprising.



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326 Figure 7 Principal component analysis using the entire spectrum of freezing history for samples after long term storage 327 measured in the frozen state.

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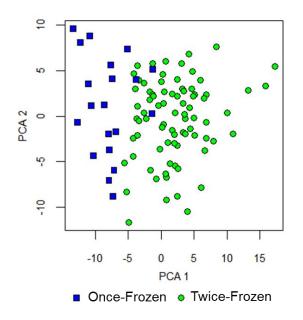
333 Table 5: K-nearest neighbors classification accuracies for the number of freezings and for freezing protocol for once and twice-

frozen samples after long term frozen storage measured in the frozen state. For twice-frozen samples, only the freezing

335 protocol for the second freezing was evaluated for classification

Sample Type	Full Spectrum	450-600 nm	900-990 nm
Once-frozen	80%	85%	15%
Twice-frozen	96%	95%	80%
Once-frozen -40°C	70%	50%	10%
Once-frozen -20°C	90%	80%	10%
Twice-frozen, last freezing -40°C	85%	73%	68%
Twice-frozen, last freezing -20°C	98%	90%	85%

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Figure 8: Principal component analysis using the entire spectrum for the freezing history of samples after long term frozen
 storage measured in the thawed state.

340 Table 6: K-nearest neighbors classification accuracies for the number of thawings and freezing protocol for once and twice-

341 frozen samples after long term frozen storage measured in the thawed state. For twice-frozen samples, only the freezing 342 protocol for the second freezing was evaluated for classification

Sample Type	Full Spectrum	450-600 nm	900-990 nm
Once-thawed	75%	50%	45%
Twice-thawed	93%	96%	86%
Once-frozen -40°C	40%	30%	50%
Once-frozen -20°C	50%	30%	10%
Twice-frozen, last freezing -40°C	45%	50%	50%
Twice-frozen, last freezing -20°C	40%	43%	43%

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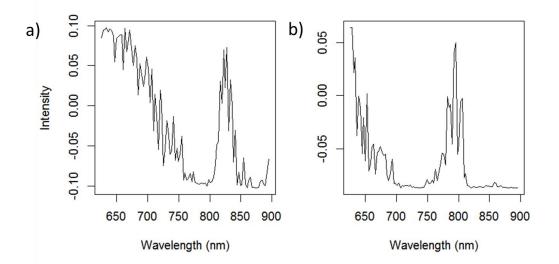
# 344 **4 DISCUSSION**

The results show that hyperspectral imaging can be used to reliably predict some aspects of the freezethaw history of a packaged fish sample. An unexpected result was the ability to much better identify the freeze-thaw history of the samples in the frozen state than in the thawed state. Analysis of hyperspectral images can accurately predict, in the frozen state, whether a sample is once or twicefrozen. Reliable discrimination of whether the frozen samples were frozen quickly at -40°C or slowly at -20°C is also possible, but only for the last freezing if the sample is twice-frozen. For samples in the non-frozen state, it appears possible to determine whether the samples are fresh, once-thawed or twice-thawed. These capabilities appear to hold even after twelve months of storage at -20°C. However, estimation of the reliability of prediction for the samples that underwent long term storage was hampered by the small sample set; a larger study will need to be performed to better understand the accuracy and limits of the technique on samples that have undergone long term frozen storage.

Given the approach used here, it does not appear possible to determine the first freezing protocol for twice-frozen samples. Similarly, information regarding the freezing protocol used appeared to be lost during the thawing process, as analysis on data from the thawed samples could not accurately discriminate those that had been frozen quickly at -40°C from those that had been frozen slowly at -20°C. Discrimination of the thawing protocol used also did not appear to be possible.

361 For this study, while separating the spectra into different regions appeared useful for visual analysis of 362 the sample clustering and to help understand the physical processes behind the analysis, using the 363 entire spectrum for classification appears to be the superior approach. Curiously, analysis performed 364 using both the blood and water regions of the spectrum simultaneously typically performed marginally 365 worse than the classifications performed on the regions individually. Inspection of the full-spectrum 366 loading values for the first principal component for the frozen samples, shown in Figure 9a, shows a 367 peak at the 800-840 nm region that is not known to be associated with any particular sample 368 constituent. In the loading values for the thawed samples, shown in Figure 9b, the peak is shifted to 369 the 750-800 nm region. Previous researchers have seen the 760 nm OH stretch in water shifts towards 370 800 nm with increased ice fraction (Ottestad et al. 2009), which is what we believe is occurring here. 371 We expect this behavior is the reason the full spectrum analyses generally were superior to the blood 372 or water region analyses.

373



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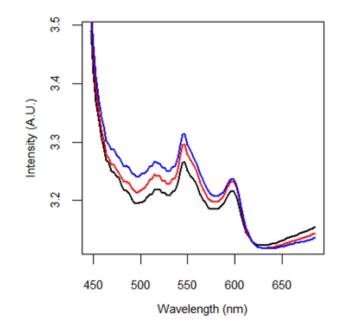
Based on what we are and are not able to predict, we believe that the structure and quantity of ice in

378 the frozen samples plays an important role in the ability to determine the last freezing protocol used.

379 Despite the sample being frozen, some water in the tissue may remain in liquid form due to freezing 380 point depression. For cod, it has been found that the quantity of unfrozen water decreases down to -381 33°C, where it remains stable at approximately 5% (Tolestorebrov et al. 2014). This super-cooled 382 water, once-frozen, will remain frozen even when heated back to -20 °C, though structural changes to 383 the ice may occur (Syamaladevi et al. 2012). Previous research on frozen salmon has shown that 384 different freezing temperatures will also result in different size and shapes of ice crystals in the fish 385 flesh (Syamaladevi et al. 2012). Scattering of light as it travels through the sample is affected by 386 crystalline structure, which would then be detectable by the hyperspectral measurement. This would 387 also explain why analysis on the thawed samples was unable to classify samples by their freezing 388 temperature history.

389 Even though classification using the spectral range associated with blood was less accurate than using 390 the full spectrum or the water region of the spectrum, good classification of freezing protocol was still 391 possible for the frozen samples. If this was due to the different freezing processes causing chemical 392 changes to the blood, such as oxidation, these effects should still be detectable in the thawed state, 393 which they were not. Therefore, it is believed the spectral changes associated with freezing protocol 394 in the blood region mainly arise from light scattering effects, which by nature span the whole spectral 395 range measured (Jacques 2013). Much of this likely is associated with the ice formation. There may 396 also be an additional scattering effect that arises from the blood itself due to the hindered molecular 397 rotation during freezing. Future work is planned to test these theories and better understand the 398 changes that occur to the tissue during the freezing processes.

399 Averaged spectra for the fresh, once thawed and twice thawed samples are shown in Figure 10. In 400 Sivertsen et al. (2011), the results indicated that changes in the oxidation state of blood allowed the 401 discrimination between fresh and once-frozen samples. While the thawed spectra in this study also 402 show indications of oxidation, the changes are less than might be expected given the duration of the 403 program. This is attributed to the vacuum packing of the samples, such that the available oxygen is 404 minimal. In this study, the ability to discriminate between the fresh, once thawed and twice thawed 405 samples appears to depend on increased light scattering with each freeze-thaw cycle. This effect is 406 prominently observed by the increased wavelength dependent baseline shift in the spectra in the 450 407 to 600 nm region with each freeze-thaw cycle. The increased light scattering is attributed to 408 denaturation of proteins during freezing and thawing (Love 1962).



#### 410 Figure 10: Averaged absorbance spectra for fresh (black), once thawed (red) and twice thawed (blue) samples

411 Examination of the plot of the first and second components of the once and twice-frozen samples (Fig. 412 5) shows an interesting trend. All the once-frozen -40°C samples cluster in the lower left quadrant. 413 Moving up and to the right, we see the clusters of the once-frozen at -20°C and twice-frozen with the 414 second freezing at -40°C. Finally, the twice-frozen samples with the second freezing at -20°C cluster in 415 the upper right quadrant. We speculate that movement along this axis from lower left to upper right 416 quadrant may be related to fish quality attributes. Samples frozen at a lower temperature have been 417 shown to have characteristics associated with higher quality (e.g. less water loss, gaping, etc.). 418 Similarly, twice-frozen samples have been shown by previous researchers (Kent et al. 2004) to have 419 more undesirable characteristics such as toughness, dryness and fishy smell. How characteristics of 420 samples once-frozen at -20°C compare to twice-frozen at -40°C is unknown; to the authors' knowledge, 421 no studies in the literature exist that compare these situations. Future work is planned to compare 422 sample location on this axis to other quantifiable properties that previous research (Bonilla et al. 2007, 423 Warm et al. 1998) has shown to be associated with perceived high and low quality in cod such as 424 appearance, texture, odor and flavor.

425 While the results here are promising, there are important differences in the study compared to real 426 world conditions. It is unrealistic that a sample's entire freeze-thaw history would occur in the vacuum 427 packed, filleted state. For twice-frozen products, the first freezing would occur at sea with the fish 428 being headed, gutted and frozen whole. The fish would then be thawed ashore for filleting, and then 429 refrozen. Also, measurements were performed on a section of cod loin in samples that had been 430 handled to produce high quality fillets. How robust the results are in other sections of fish, species of 431 fish or in fish that have had less than ideal handling are unknown. Therefore, generalizations of results 432 of this study to other situations need to be treated with caution.

433 With these considerations in mind, there exist potential applications of the results at the industrial 434 scale. There are multiple stages at which freezing procedure control could be relevant. For the fish 435 processing industry, the speed and non-invasive nature of hyperspectral imaging make it an excellent 436 technology for online quality control when buying deliveries of fish; samples can easily be measured 437 at a rate compatible with online sorting equipment. While region selection was performed manually 438 here, this type of task could be easily automated using standard image analysis methods. This means 439 that fish could be automatically sorted by different handling protocols as they can be currently sorted 440 by size today. A delivery could be sorted into different batches based on freezing protocol used or 441 number of freezings undergone. By sorting the fish by handling procedure, and by proxy the perceived 442 quality, this would enable the maximum price to be obtained for a delivery as quality is currently 443 assigned to a whole batch by the lowest quality specimens. For buyers of finished frozen seafood 444 products, such as a supermarket chain, the technology would enable them to confirm the product has 445 been handled as stated and avoid overpaying for lower quality product. Similarly, there may exist 446 situations where the end buyer also wishes to evaluate the freeze-thaw history of seafood products, 447 for example to confirm that the products have not been allowed to thaw during transport and then 448 refrozen later.

449

## 450 5 **CONCLUSIONS**

451 We demonstrate here that hyperspectral imaging has the potential for use as an online method for 452 evaluation of sample freeze-thaw history. It is possible to differentiate between the freezing protocols 453 on packaged samples in the frozen state. In the thawed state, differentiation between fresh and 454 frozen-thawed samples is straightforward, as is differentiation between thawed samples subjected to one or two freeze-thaw cycles. On samples frozen for the second time, it is possible to differentiate 455 456 between the freezing protocols of the second freezing but not the first freezing. Identification of 457 applied thawing protocols did not appear to be possible with the approach used in this study. Differentiation between freezing protocols is still possible after approximately 12 months of storage. 458 459 We also find a possible correlation between sample clustering and characteristics relating to perceived 460 quality. These findings could enable the development of technologies for online quality screening of 461 frozen seafood products.

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