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## **1** Benchmarking of techniques used to assess the freeze damage in potatoes

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## 13 Abstract

14 In this study, benchmarking of methods used for assessing freeze damage in potatoes was 15 carried out. Initially, the samples were frozen by subjecting them to three different 16 temperatures (i.e. at -18 °C, -30 °C, and at -74 °C). Then, different analytical techniques 17 comprising of focused methods (i.e. cryo-SEM, confocal laser scanning microscopy-CLSM) 18 and global methods (i.e. texture analysis, low field nuclear magnetic resonance (NMR), 19 exudate loss and colour change) were used to assess the impact of the freezing treatment from 20 the different point of view addressed by each method. As a result, each of these methods were 21 able to distinguish significantly fresh samples from the frozen-thawed samples. Focused 22 methods like cryo-SEM and CLSM methods could differentiate the impact of all three 23 different protocols. Meanwhile, texture analysis (including conventional method and novel 24 method based on a touchless laser puff firmness tester), NMR and exudate loss could only determine the quality difference between -18 °C and -74 °C freezing conditions. Colour 25 26 analysis was found as an inappropriate parameter for comparing the three freezing protocols. 27 Among all analytical techniques, cryo-SEM provides the most authentic information about the 28 product as the analysis is performed in frozen state, while for other techniques the product is 29 thawed prior to analysis.

## 30 **1. Introduction**

Freezing is a preservation technique that has been used for ages for increasing the shelf life of
 food products. Compared to other long-term preservation methods, freezing technique causes

33 less deterioration to nutrients and sensory properties in fruits and vegetables (Barbosa-34 Cánovas, Altunakar & Mejía-Lorío (2005). However, it can cause some irreversible damages such as texture loss, colour change, etc. Three main phenomena may describe freeze damage; 35 36 the primary mechanical effect caused by the transition of water into ice, the biochemical 37 effect caused by cryo-concentration and the secondary mechanical effect eventually caused by 38 ice contraction after freezing (Reid, 1997; Shi, Datta, & Mukherjee, 1999, 1998; Shi, Datta, & 39 Throop, 1998). The extent of freeze damage caused to the food products on thawing is greatly 40 linked to the size and location of the ice crystals, which in turn is related to the applied 41 freezing rate (Chevalier, Le Bail, & Ghoul, 2000). Freeze damage can also be linked to the 42 cryo-concentration effect; the aqueous solution present in the cells is exposed to a progressive 43 concentration in solute caused by ice formation. The remaining aqueous phase which tends to 44 become more and more concentrated can cause denaturation of the proteins and other 45 organized biopolymers (i.e. cellulose based systems for fruits and vegetables), which in turn degrade their mechanical properties and by the way the overall texture of the tissue (Gao & 46 47 Critser, 2000; Lovelock, 1957; Reid, 1997). The mechanisms of freeze damage associated 48 with cryo-concentration effect have been thoroughly discussed in a review article from Gao & 49 Critser (2000). Rapid freezing (high freezing rate) favours the genesis of fine ice crystals 50 distributed uniformly within the product, thus reducing the dislocation of water from intra to 51 the extra cellular spaces, resulting in higher water holding capacity upon thawing. In addition, 52 the time of exposure to the highly concentrated aqueous solution caused by cryo-53 concentration is being reduced during rapid freezing, resulting to a lesser extend of the 54 biochemical degradations (Delgado & Sun, 2001; Fennema, 1966; Orlowska, Havet, & Le-55 Bail, 2009; Sadot, Curet, Rouaud, Le-bail, & Havet, 2017; Shi, Datta, & Mukherjee, 1998; 56 Singh & Heldman, 2009). The result is reduced cell destruction and better quality attributes. 57 However, two main disadvantages lie behind rapid freezing rates : (i) rapid freezing process 58 requires a higher amount of energy, hence increases the consumed energy and the overall 59 operating costs and (ii) the use of extremely high freezing rates such as cryogenic freezing, 60 may lead to crack development in the samples and yield unexpectedly poor quality of the final 61 product (secondary mechanical effect quoted above) (Shi, Datta, & Mukherjee, 1999, 1998; Shi, Datta, & Throop, 1998). On the contrary, at a slow freezing rate, intracellular water move 62 63 to the extracellular domains (from the inside of the cell) and thus can result in greater ice 64 crystal size (in the extracellular spaces) and also in a higher cell dehydration (Gao & Critser, 65 2000; Mazur, 1984). Besides, slow freezing rate will also result in a prolonged exposition of 66 the tissue to the concentrated solution occurred from the cryo-concentration effect (Gao &

Critser, 2000; Mazur, 1977, 1984). In fact, the formation of larger ice crystals along with 67 volume shrinkage and long-term exposure to high solute concentrations would cause a higher 68 69 destruction of cellular structure (high risk of punctured cell membrane/cell wall, collapsed cell 70 structure and cell separation) (Chassagne-Berces et al., 2009; Chassagne-berces, Fonseca, 71 Citeau, & Marin, 2010; Gao & Critser, 2000; Mazur, 1984). As an outcome, the leakage of 72 the fluid from the cell will be higher and a product having a water-soaked appearance and 73 mushy texture will be obtained. The last important parameter that may be responsible for 74 quality losses of frozen foods is the storage conditions; time-temperature parameters, as well 75 as temperature fluctuations, are also responsible for the quality loss.

76 The assessment of the quality of a frozen food product is obtained via analytical methods used 77 to quantify the freeze damage. Broadly, these methods can be categorized as global methods 78 and focused methods. The global methods provide quality information at a macro/meso level. 79 In other terms, the quality information obtained using such methods is the average value from 80 a whole sample or at least from a substantial piece of sample (several grams or several cm<sup>3</sup>). 81 The global methods comprise of texture analysis, exudate loss measurement, solute 82 diffusivity, impedancemetry, colour analysis, etc. While the focused methods provide quality 83 information at a micro level (cell level or even at a lower scale such as ice crystal or water 84 molecule scale). For instance, information about ice crystals size, cellular structure intactness, 85 etc. can be obtained by using focused methods. Microstructures evaluation methods such as 86 Scanning electron microscopy (SEM) (includes conventional SEM, environmental SEM 87 (ESEM), and cryo-SEM), confocal laser scanning microscopy (CLSM), X-rays tomography, 88 etc. fall under focused method. Low field nuclear magnetic resonance relaxometry (NMR) is 89 at the same time a global and focused method as it embraces a sample of *circa*  $1 \text{ cm}^3$  scale 90 and provides info at the level of water molecules (Jha et al., 2018).

91 In the literature, numerous benchmarking studies are available related to freezing effects on 92 the quality parameters of the fruits and vegetables. However, no studies are available in the 93 literature that benchmarks the techniques used to assess the freeze damage in fruits and 94 vegetables.

95 This work was framed under the following objectives

96 i. To introduce new freeze damage assessment methods and check their efficiency in97 quantifying the freeze damage.

98 ii. To benchmark the freeze damage methods (both conventional and novel) used for
99 assessing the freeze damage in plant-based products with the intention to facilitate the
100 researchers and industries to choose the appropriate method to assess the freeze damage.

101 The ultimate objective of this study is to provide a decision table which will allow to 102 categorize the analytical techniques based on sensitivity, efficiency, accuracy, cost of 103 operation, and ease of operation.

## 104 **2.** Materials and methods

## 105 **2.1. Product properties**

106 Potatoes (Solanum tuberosum L. cv. Innovator) were the food matrix chosen for this 107 benchmarking study. McCain Foods (Harnes, France) kindly supplied us potato batch having 108 uniform size and maturity. Two batches of potato were used for this test. The first batch of 109 potato was used for NMR tests. In order to reduce the variability, for each freezing condition, 110 samples intended for NMR test were taken from the same potato. Potatoes from the second 111 batch were used for time-temperature history study and freeze damage assessment tests. The 112 average moisture content of both batches was  $75.70 \pm 1.40\%$  (wet basis) (Note: the values of 113 moisture content are presented here as mean  $\pm$  standard deviation. Similarly, the values for 114 other parameters in the entire manuscript are expressed as mean  $\pm$  standard deviation).

#### 115 **2.2.** Sample preparations

116 The sample size used to study the effect of conventional freezing processes on the quality parameters depended on the freeze damage assessment methods. For instance, to study the 117 118 effect of freezing rates on freezing characteristics, texture, colour and drip loss, cuboids of potato  $(1.3 \times 1.3 \times 1 \text{ cm}^3 - \text{length} \times \text{height} \times \text{width}; \text{ weight } 1.95 \pm 0.5 \text{ g})$  were used. The 119 120 potatoes were cut in cuboid shape using a dicer (Ibili Manage, Inc. Spain). The potato samples 121 used for NMR relaxometry and cryo-SEM analysis were cylindrical in shape ( $\phi = 8 \text{ mm}$  and 122 H = 10 mm). For confocal laser scanning microscopy (CLSM) analysis, cylindrical potato 123 samples ( $\phi = 8$  mm and H = 5 mm) were chosen. The samples were then immediately 124 transferred into the freezer and were cooled from the ambient temperature to the desired 125 temperature.

126 It is worth mentioning that during this study, one of our aims was to analyze the samples in its 127 original state without causing any alterations to them, thus we had to vary sample size for 128 potato according to the freeze damage assessment methods. For instance, NMR tube has an

129 internal diameter of 1 cm and hence required sample having a diameter < 1 cm. Also, another 130 requirement of NMR test is to have the NMR tube covered minimum up to 1 cm in height. 131 The cylindrical sample with a diameter of 8 mm and height of 1cm perfectly matched both the 132 requirements. For cryo-SEM, sample size similar as NMR facilitated easy cutting of sample in 133 the frozen state and allowed the selection of the analyzed sample from nearly the same 134 location for each condition. For CLSM, small sample size helped the proper and fast staining 135 of the sample and facilitated the selection of the analyzed sample from the same location of 136 each matrix.

## 137 **2.3. Freezing apparatus and freezing conditions**

138 Freezing of potatoes (unblanched) was performed at -18 °C in a cold room, at -30 °C and 0.5 m/s air velocity in an industrial batch freezer (MATAL, France), and at -74 °C in an 139 140 ultra-low temperature freezer (TSE240V, Thermo Scientific, Marietta, Georgia, USA). The 141 three conditions will be referred to as slow freezing (SF), intermediate freezing (IF) and fast 142 freezing (FF) in further sections of the manuscript. Once frozen, the samples were packed in the zip-lock and stored at  $-40 \pm 2$  °C ( $\approx$  for 2-3 days) until quality evaluation tests were 143 144 performed. The time-temperature profile during the freezing tests was studied by inserting a 145 K-type thermocouple at a geometric centre of the product. The (thermocouple) was calibrated 146 against a reference platinum probe (Comptoir Lyon Allemand – Lyon-France). During 147 measurements the temperature of the sample was recorded every 2 s with a data logger with an accuracy of  $\pm 0.1$  °C. For comparison purposes, temperature profiles from the initial 148 149 temperature 18 °C to -18 °C were considered. The characteristic freezing time and overall 150 freezing time were determined using time-temperature data. The characteristic freezing time 151 was the measure of local freezing rate and it was defined as the time during which the 152 temperature at a particular point changed from the initial freezing point to a temperature at 153 which 80% of the water (at that point) was converted into ice. The temperature range considered for the characteristic freezing time estimation was from -1 to -7 °C, similar 154 155 temperature range was used by Li & Sun (2002) for the determination of characteristic 156 freezing time during ultrasound assisted freezing of potatoes. The overall freezing time was 157 the time required to lower the temperature of the geometrical centre of the product from the 158 ambient temperature (18 °C) in the present study) to a given temperature (-18 °C). Samples 159 for quality analysis were frozen using the same procedure but without optical fibre inserted. 160 Each experiment was performed at a minimum of individual triplicates.

## 161 **2.4. Thawing protocol**

162 The frozen potato samples were thawed at room temperature  $(20 \pm 1 \,^{\circ}\text{C})$  for 2 h in a zip-lock 163 bag ("static air thawing" method) before performing colour, drip loss, texture and solute 164 diffusion measurements. For NMR analysis the samples were thawed at 4 °C for 4 h in the 165 NMR tube.

## 166 **2.5. Colour analysis**

The colour of the potatoes was measured using a portable and handheld chroma meter CR-167 400 (Konica Minolta, Inc. Japan). Using this equipment, the L<sup>\*</sup>, a<sup>\*</sup>, b<sup>\*</sup> values for samples from 168 each condition were obtained with high accuracy. The maximum value for  $L^*$  is 100 169 170 (represents a perfectly white surface or a perfectly reflecting diffuser) and its minimum value is 0 (represents perfectly black surface). The positive and negative a<sup>\*</sup> corresponds to red and 171 green colour. Similarly, positive and negative b<sup>\*</sup> represents yellow and blue colour, 172 173 respectively. A single value for the colour difference was achieved by calculating the overall 174 colour difference ( $\Delta E$ ) value; it takes into account the differences between L<sup>\*</sup>, a<sup>\*</sup>, b<sup>\*</sup> of the specimen (e.g. frozen-thawed sample) and reference (fresh sample), and it was calculated by 175 176 using Eq. (1) (Anon, 2018). The frozen samples were thawed before the colour measurements. 177 At least eight measurements were recorded for each freezing protocol.

$$\Delta E = \sqrt{\Delta L^{*^2} + \Delta a^{*^2} + \Delta b^{*^2}}$$
(1)

178

## 179 **2.6. Drip loss measurement**

180 The drip loss required great care in terms of manipulation. The mass differences that are being 181 determined are usually small and it can be advantageous to use paper tissue to collect all the 182 drip released by a sample. This requires the weight of the bag containing the sample, the 183 initial mass of the tissue paper and all the final masses. There is also a risk of moisture 184 condensation on the sample during frozen storage (if unpacked) or during the thawing process 185 or even sample handling after thawing if their temperature is lower than the dew point 186 temperature. The drip loss unit is very often given as g drip per g of sample. In most existing 187 studies, drip losses are expressed as free drip; this means that drip the collected from the 188 sample due to gravity forces and possible capillary interactions with the tissue paper. In some 189 studies, the samples are compressed under selected stress (Jahncke, Baker, & Regenstein, 190 1992) or after centrifugation for selected g values under given times (Penny, 1975). In our 191 study, the drip loss was determined based on the weight difference between frozen sample 192  $(W_1)$ , and thawed sample  $(W_2)$ . The drip loss (%) was calculated using Eq. (2). At least 9 193 samples were analyzed for each freezing condition.

Drip Loss (%) = 
$$\frac{W_1 - W_2}{W_1} \times 100$$
 (2)

194

## 195 **2.7. Texture analysis**

196 Texture analysis of food is often carried out using the TPA protocol (Texture Profile 197 Analysis) proposed by Bourne (Bourne, 1968). Quite often, simple compression tests are used 198 for frozen food aiming at determining the Young's Modulus (small deformation, typically 199 10% strain) and eventually the failure strain or stress. In our study, the frozen potato samples 200 were thawed and compressed in a texture analyser (AMETEK, Lloyd Instruments, France) 201 equipped with a 1000 N load cell and operating at a test speed of 50 mm/min beyond failure 202 point which was marked by a significant drop in the force reading (Figure 1). The 203 compression test was performed using a 50 mm compression plate. A similar procedure was 204 used by Khan & Vincent (1993, 1996) for determining the failure stress, Young's Modulus, 205 and failure strain of potato. Based on our preliminary trials and the results from Khan & 206 Vincent (1993) and Alvarez, et al. (2002), compression of the sample to 50% strain was found 207 more than sufficient to cause the failure of our sample (Figure 1). The maximum force exerted 208 during the compression test was recorded as the firmness/hardness (N) of the samples (Figure 209 1). Moreover, the stress vs. strain curves during compression tests were examined and 210 Young's modulus (E) (the slope of the loading curve at the point of its highest gradient) was 211 also acquired (Chassagne-Berces et al., 2009). All measurements were performed at  $21 \pm 2^{\circ}$ C. 212 At least 12 samples were analyzed for each freezing condition.

213

## 2.7.1. Laser-puff firmness tester

A touchless laser-puff firmness tester which was designed and constructed in our lab (GEPEA lab, ONIRIS) was used for non-destructive measuring of the texture of fresh and frozenthawed potatoes (Figure 2). The firmness tester consists of the following components: a mean to generate impulsive jet of air; a nozzle (having a diameter of 0.004 m) to direct the air onto the surface of the specimen under investigation; a deformation measurement unit containing a laser source to generate coherent source of light directed on the surface of the object impacted by the air jet, a detector of the light which is reflected from the specimen surface, and an 221 analyzer to estimate the amount of deformation sustained by the product surface; a control 222 panel to change the pressure value and an appropriate software (developed in our lab) to 223 execute the test. Based on the preliminary trials, for measuring the firmness of frozen-thawed 224 potatoes, the air nozzle was located at 10 mm from the food surface and an air pressure of  $5 \times$ 225  $10^5$  Pa and air jet exposure time of 100 ms were found most appropriate. The estimated pressure exerted onto the food surface was in the range of  $3.32 \times 10^5$  Pa assuming a surface of 226  $1.256 \times 10^{-5}$  m<sup>2</sup> impacted by the air jet. Prussia, Astleford, Hewlett, & Hung (1994) used 227 impact pressure of  $3.10 \times 10^5$  Pa (45 psia) in order to determine the firmness of potato. 228 Similar to the authors, Hung, McWatters, & Prussia (1998) who used the same exposure time 229 230 to determine the firmness of peaches.

231 The sample was placed on the sample platform below the air delivery nozzle and the position 232 of sample was adjusted using a positioner in order to ensure (i) the air jet and the laser hits the 233 same point on the sample and (ii) the distance between end of the nozzle and the sample 234 surface is same for all trials. Subsequently, by using the software air jet was directed on the 235 sample and the deformation was measured. The software yielded deformation results in volts 236 (Figure 3). The maximum deformation can be referred to as the difference between the 237 highest initial value and lowest value recorded by the deformation unit during the test. The 238 conversion of volts to mm (millimeter) based unit was made by multiplying with a conversion 239 factor (in the present case, volt value was multiplied by 2 to obtain final value in mm). The 240 final deformation value was an average of sixteen measurements.

## 241 **2.8. Microstructure examination**

## 242 **2.8.1.** Cryo-SEM analysis

243 Cryo-SEM analysis is a very specific technique which requires very careful and proper 244 sample preparation. A thin specimen of slab shape was cut from a given location of the frozen 245 food using a sharp scalpel. In our case, the central section of the frozen potato samples was 246 selected for collecting the sample. Also, in the case of structured tissue, given shape must be 247 provided to the sample to identify a given orientation; for example, in the case of a muscle, 248 the longer section of sample is usually longitudinal to the fibers in order to be able to do cut 249 view either parallel or perpendicular to the fibers. In the case of a vegetal tissue has no major 250 orientation so special care was taken mainly for the location, shape and size of the sample. It 251 was fixed into the groove notched in a copper sample holder (10 mm diam. 10 mm height) 252 using an OCT (optimal cutting temperature) compound (Tissue-Tek, Sakura, Finetek, USA).

253 An important technical issue lies in preventing any thawing of sample when the sample is 254 contacted onto the sample order. To overcome this difficulty, the following protocol was 255 used. All manipulations were done in a cold box filled with CO<sub>2</sub> sticks (-78.5 °C) in order to 256 condition the sample which was kept in this ambiance. The sample holder was refreshed down 257 to around 0 °C by contacting a CO<sub>2</sub> stick. Then a small amount of OCT was installed in the 258 groove of the sample holder. Then, at the same time, the sample holder was contacted onto a 259 CO<sub>2</sub> stick meanwhile the frozen sample was contacted to the OCT located in the groove. 260 Within less than a few seconds, the sample was fixed onto the sample holder thanks to the 261 freezing of the OCT. The very high thermal diffusivity of the copper material (sample holder) 262 compared to the frozen sample ensures that the sample will not undergo any thawing when the 263 sample is contacted to the sample holder. The sample was left for 2 min in a box filled with 264 dry CO<sub>2</sub> to harden the OCT compound. After that, the sample was loaded onto a precooled 265 copper specimen stub (in liquid N<sub>2</sub>) and was quickly transferred in the cryo-preparation 266 chamber of the cryo-SEM (LS10, ZEISS EVO, Germany) maintained under vacuum at - 80 267 °C, where it was cryo-fractured using a precooled sharp knife mounted inside the chamber. 268 The fractured sample was etched in the preparation chamber for about 5 to 10 min to expose 269 the subsurface information in order to allow a partial sublimation of the ice crystals. The 270 etching time required adjustment depending on the size of the sample, the pressure level and 271 the type of tissue. The sample was finally coated with a thin conducting layer of gold (5 nm) 272 in the cryo-preparation chamber and then was transferred to the cold stage in the cryo-SEM 273 (maintained at - 80 °C) where microstructural observation was performed. The fractured 274 surfaces of potatoes were examined with an accelerating voltage of 11 kV. An illustration of 275 the sample preparation steps is presented in Figure 4. It is worth mentioning that all the 276 samples prepared for cryo-SEM analysis after the different freezing conditions acquired from 277 the same potato.

## 278 **2.8.2.** Confocal laser scanning microscopy (CLSM)

The confocal laser scanning microscopy was also used for potatoes samples. Figure 5 provides a brief description of the protocol followed to prepare potatoes for the CLSM observations. The CLSM was performed using an Eclipse Ti inverted microscope (Nikon Ti A1, Japan). For each freezing condition, independent experiments were performed in triplicates.

It is worth noticing that the acridine orange (used for this study) is well known to be a

metachromic dye, especially for staining DNA and RNA. Such dye is able to give fluorescence with emissions wavelength depended on its interaction with chemical functions, charge or geometry of a compound. The acridine orange was chosen here as a dye able to stain simultaneously the different cell layers of the fruits as it has been widely applied for fluorescent staining of plant tissue with a high fluorescence emission. The maximum emission wavelength of this dye is 500-530 nm and excitation occurs at about 488 nm.

## 291 **2.9. NMR relaxometry**

292 For all the freezing protocols, NMR tests were performed for samples in frozen state (at -20293 °C) and frozen-thawed state (at 4 °C). Initially, the frozen samples were placed in the NMR 294 tube ( $\emptyset = 10$  mm) precooled to -20 °C. Then the tube was quickly placed in the spectrometer (maintained at -20 °C) and left for 10 min to ensure that the samples were at 295 296 -20 °C at the beginning of NMR test at a negative temperature. Upon completion of NMR 297 tests in frozen condition, the same samples were thawed (at 4 °C for 4 h) and then NMR tests 298 on frozen-thawed samples (at 4 °C) were carried out. In order to compare the damage caused 299 by different freezing procedures, the NMR measurements of fresh samples were performed at 300 4 °C and were compared with results obtained for frozen-thawed samples obtained under 301 various freezing protocols. The Minispec mq20 spectrometer (Bruker) at 0.47 T (20 MHz proton resonance frequency) equipped with a thermostated ( $\pm 0.1$  °C) <sup>1</sup>H probe was used for 302 NMR analysis. Triplicates were performed for each freezing condition of potato. 303

In the present study,  $T_2$  (transverse) relaxation time of protons and their respective population in the product were evaluated. The  $T_2$  distributions were determined using a Carr–Purcell–Meiboom–Gill (CPMG) sequence. For the NMR test at frozen and frozenthawed state, the 180° pulse separation was 0.04 and 0.1 ms, 2000 and 10000 even echoes were collected, and the 1024 and 256 scans were acquired with a recycle delay of 1 and 5 s resulting in a total acquisition time of about 20 and 40 min respectively.

310 An inverse Laplace transformation (ILT) was applied to convert the relaxation signal into a 311 continuous distribution of  $T_2$  relaxation components. For this purpose, a numerical 312 optimization method was used by including non-negativity constraints and L1 regularization 313 and by applying a convex optimization solver primal-dual interior method for convex 314 objectives (PDCO).

#### 315 **2.10.** Statistical analysis

- One-way ANOVA (analysis of variance) was used to determine any significant difference (in
  terms of freezing and quality characteristics) among the freezing conditions. Duncan's
- 318 multiple range test was performed to determine differences between the means (p < 0.05).

## 319 **3. Results and discussion**

## **320 3.1. Effect of different freezing rates on temperature history**

321 The representative temperature histories of potatoes frozen under the different freezing 322 conditions are presented in Figure 6. The SF process yielded freezing curves having three 323 stages i.e. the supercooling, nucleation, and phase change. Meanwhile, intermediate freezing 324 (IF) exhibited only two stages (i.e. nucleation and phase change stages) and fast freezing (FF) 325 showed only one stage (i.e. phase change state) in Figure 6. SF condition gave a degree of 326 supercooling of  $0.15 \pm 0.07$  °C, while no supercooling curve was observed for the other two 327 freezing conditions. Similar freezing curves lacking obvious supercooling at higher freezing 328 rates (at - 80 °C and liquid nitrogen immersion freezing) were also obtained by Cao et al. 329 (2018) during freezing of blueberries. It was inferred by them that at a faster freezing rate the 330 supercooled state might be unstable or lack persistence. The initial freezing point for SF and 331 IF condition was recorded as  $-0.3 \pm 0.14$  and  $-0.73 \pm 0.06$  °C (Table 1). It seems that a 332 slight depression in freezing point happened upon increasing the freezing rate. For FF 333 condition, it was hard to detect the initial freezing point due to a rapid decline in the 334 temperature during the freezing process. However, more replications could be helpful to be 335 conducted prior to the confirmation of this outcome. Recently, Cao et al. (2018) also reported 336 slight decrease (but not significant) in the initial freezing temperature of blueberries when the 337 freezing rate was increased from 0.023 °C/s (at - 20 °C) to 0.049 °C/s (at - 40 °C) or 0.11°C/s 338 (at - 80 °C) or 0.76 °C/s (liquid nitrogen immersion freezing). The initial freezing point of blueberries at 0.023, 0.049, 0.11, and 0.76 °C/s rates were recorded as  $-2.67 \pm 0.32$  °C, 339  $-3.23 \pm 0.12$  °C,  $-3.53 \pm 0.35$  °C and  $-3.36 \pm 0.60$  °C respectively (Cao et al., 2018). 340 341 Among other freezing parameters being studied, the characteristic freezing time was found to 342 be the shortest for FF condition  $(8.52 \pm 1.53 \text{ min})$ , followed by IF and SF condition 343  $(17.18 \pm 0.79 \text{ min and } 29.12 \pm 3.94 \text{ min respectively})$  (Table 1). The time spent in this zone 344 is very crucial as it determines the quality of the final product. From the perspective of higher 345 quality preservation, a shorter width of this zone is desired. Similar to characteristic freezing 346 time, the overall freezing time and the overall freezing rate also exhibited a similar trend.

## 347 **3.2. Texture analysis**

#### 348 **3.2.1.** Conventional method

349 The confined compression test was the conventional method used to determine the texture of 350 potatoes. The hardness and Young's modulus values of fresh and thawed sample (from 351 different freezing conditions) are shown in Table 2. The values of fresh samples were 352 significantly different (p < 0.05) compared with those of the frozen-thawed samples. FF 353 process caused less decay in hardness value ( $\approx 50\%$ ) than IF process ( $\approx 62\%$ ) or SF process 354 ( $\approx$  74%). However, significant differences in hardness values were observed only between SF 355 and FF samples. Young's modulus values exhibit a similar tendency as hardness values in 356 Table 2.

357 Khan & Vincent (1996) reported that compressive stiffness of potato less degraded when 358 freezing was performed at a higher freezing rate (10 °C/min) compared to slow freezing rate (1 °C/min). According to Mazur (1984) and Chassagne-Berces, Fonseca, et al. (2010), lower 359 360 dehydration associated with small ice crystals induced less breakage of cell walls, and hence, 361 better texture preservation was achieved at higher freezing rates when compared to the slow 362 freezing rate. The high freezing rates also decrease the collapse of cell walls and generate less 363 intercellular spaces, and hence result in better texture preservation (Chassagne-Berces et al., 364 2009). Besides, Phinney, Frelka, Wickramasinghe, & Heldman (2017) reported that the 365 extent of texture loss of potato depends on the freezing time. It was found that the hardness of 366 thawed potato reached a maximum value when the freezing time decreased dramatically up to 367 1000 s. Further, increase in freezing time beyond 1000 s to around 5000 s did not cause any 368 further texture loss in their case. Van Buggenhout et al. (2006) evaluated the effect of three 369 different freezing conditions i.e. slow freezing (freezing time ( $f_t$ ) = 300 min), rapid freezing ( $f_t$ 370 = 40 min)) and cryogenic freezing ( $f_t = 10$  min)) on the hardness retention of thawed carrots. 371 They reported that rapid and cryogenic freezing condition had a higher hardness retention 372 than the slow freezing condition.

373 **3.2.2.** Laser-Puff firmness tester

Laser-Puff firmness tester allows rapid and non-destructive texture analysis of the food products (Hung et al., 1998; McGlone & Jordan, 2000; Prussia et al., 1994). An attempt was made to use this method, to the best of our knowledge for the first time, for measuring the texture of frozen-thawed fruits and vegetables. In this section, the results from laser-puff firmness analysis of potato will be presented and discussed. The deformation curves and deformation values obtained during laser-puff firmness test of fresh and frozen-thawed 380 potatoes (under different freezing rates) are shown in Figure 7. The fresh samples had 381 significantly (p < 0.05) lower deformation value than all frozen-thawed samples (Figure 7b). 382 As expected, SF samples suffered the highest deformation during the tests. The deformation 383 occurred at IF samples was less than SF samples, but greater than the FF samples. However, 384 the deformation values of IF samples were not significantly different (p > 0.05) from the 385 values of SF and FF samples. The FF sample showed lower deformation than the other 386 conditions, and those values were found to be significantly lower than SF, but not 387 significantly different from IF values. The obtained results (in terms of deformation) are 388 coherent with those obtained by the classical method (discussed above). In conclusion, this 389 method could distinguish fresh and frozen-thawed samples, but had limited capability to 390 differentiate the tested freezing conditions.

## 391 3.3. NMR relaxometry

392 NMR tests were performed for the samples in frozen (at - 20 °C) and frozen-thawed states (at 393 4 °C) after being frozen under the different freezing conditions. Figure 8 shows the results 394 from NMR relaxometry of potato in a frozen state (at - 20 °C). The relaxation peaks,  $T_2^*$ 395 (including magnetic field inhomogeneities) and  $T_2$  of frozen samples provide information about the structure of the samples and about the unfrozen water at – 20 °C (Figure 8).  $T_{2\alpha}^*$  is 396 the relaxation peak of protons associated with the macromolecules.  $T_{2\beta}^*$ ,  $T_{2\gamma}$ , and  $T_{2\delta}$  are the 397 398 relaxation peaks associated with the protons of unfrozen water (Foucat & Lahaye, 2014; 399 Luyts et al., 2013). Results showed that the  $T^*_{2\alpha}$  values (relaxation time and proton population) 400 for all freezing conditions were similar, which indicated that the different freezing protocols didn't influence differently the systems at a macromolecular level (Figure 8a). Similarly,  $T^*_{2\beta}$ 401 402 values were not significantly different among the different freezing conditions (Figure 8a). If 403 we follow the hypothesis that the faster the freezing process is the less is the destruction it imparts, the values of relaxation peak components of non-freezable water ( $T_{2\gamma}$  and  $T_{2\delta}$ ) 404 405 associated with the samples frozen quickly provide an evidence of better preservation of 406 structures. Only samples that were frozen slowly (at - 18 °C) showed different values for these relaxation peak components (Figure 8a and b). It was observed that  $T_{2\gamma}$  component 407 408 values (relaxation time and proton population) for IF and FF conditions were not significantly 409 different (p > 0.05) between each other, meanwhile, these values were significantly different  $(p \le 0.05)$  with that obtained for SF condition. The lowest value of  $T_{2\gamma}$  time observed for slow 410 411 freezing (0.72 ms instead of around 0.80 ms for the other freezing conditions) can be

explained by a greater destruction followed by a diffusion of "solutes" inducing a relative 412 413 increase in viscosity (Lahaye, Falourd, Limami, & Foucat, 2015). The  $T_{2\delta}$  component 414 relaxation times for all freezing conditions were not different from each other. But, the 415  $T_{2\delta}$  component proton population was significantly lower (p < 0.05) for SF compared to the 416 other conditions. This seemed to indicate the loss of fluid from the respective water 417 compartment due to greater damage occurred during SF process. No significant difference (p 418 > 0.05) (with respect to  $T_{2\delta}$  component proton population) was observed among IF and FF 419 samples.

420 Figure 9 represents the  $T_2$  peaks of fresh and frozen-thawed samples at 4 °C. With regard to fresh samples, the distribution of  $T_2$  relaxation peak has five components  $(T_{2a}, T_{2b}, T_{2c}, T_{2d} \&$ 421 422  $T_{2e}$ ), whose values averages are in good agreement with the literature data (Rutledge, Rene, Hills, & Foucat, 1994). Based on these data, an allocation of different components of  $T_2$  have 423 424 been proposed:  $T_{2a}$  and  $T_{2b}$  are the relaxation peaks associated with the water present in the 425 cell walls and the vacuolar membrane.  $T_{2c}$  is the relaxation peak of water in starch grains.  $T_{2d}$ 426 and  $T_{2e}$  are the relaxation peaks of water in the non-starch vacuoles, the nucleus and the 427 cytoplasm (Rutledge et al., 1994). The measurement of the  $T_2$  components values (relaxation 428 time and proton population) of the samples after thawing makes it possible to observe the 429 influence of different freezing protocols on the mobility of the water compared with the fresh 430 samples. It can be seen that the freezing-thawing process affects the resolution of the  $T_2$ 431 distribution peaks (Figure 9). For instance, the  $T_{2c}$  and  $T_{2d}$  components which were 432 distinctively visible in the fresh sample could no longer be differentiated in the frozen-thawed 433 sample. Apart from this,  $T_{2d}$  and  $T_{2e}$  times of fresh potato decreased upon freezing-thawing.

434 Among the different freezing conditions used in this study, only FF (at – 74 °C) preserved  $T_2$ components distributions with good resolution over the entire time range studied. Four  $T_2$ 435 436 components were characterized for FF (against five for fresh samples), meanwhile, for IF and SF conditions, three and two  $T_2$  components could only be characterized. For all freezing 437 438 conditions, the mobility of water associated with non-starch vacuoles, nuclei and cytoplasm 439  $(T_{2d} \text{ and } T_{2e})$  decreased. This reflects a reorganization of the fluids following a partial rupture 440 of the cellular structures (Lahaye et al., 2015) irrespective of the freezing speed. Compared to 441 fresh sample, the decrease in  $T_{2e}$  time was lower for FF process ( $\approx 46\%$ ) and was followed by SF ( $\approx$  53%) and IF ( $\approx$  54%) process. However, no significant difference among freezing 442 443 processes (in terms of  $T_{2e}$  time) was observed.  $T_{2e}$  component proton population data reveal

444 that FF and IF samples had similar values (p > 0.05) as the fresh sample, while it significantly 445 decreased (p < 0.05) in the case of SF samples. This decrease was followed by an increase in 446 proton population of the consecutive peak in the  $T_2$  distribution curve for SF samples (Figure 447 9b), depicting the transfer of water between two compartments which might have probably 448 happened due to the breakdown of vacuolar membrane. However, no such trend was observed 449 for other freezing conditions (Figure 9c and d). The  $T_2$  component values (relaxation time and 450 proton proportion) adjacent to the  $T_{2e}$  for SF samples was significantly different (p < 0.05) 451 from those of FF and IF conditions. The  $T_{2a}$  and  $T_{2b}$  components of FF samples had similar 452 relaxation times as the fresh samples, illustrating the overall better preservation of membranes 453 and walls (despite a reorganization at the level of populations). IF also fairly maintained the 454  $T_{2b}$  component values and they were found to be similar to the fresh sample. Due to poor 455 resolution, it was difficult to extract  $T_{2a}$  component value for IF samples.  $T_{2a}$  and  $T_{2b}$ 456 component values for SF sample also could not be resolved due to the poor resolution of the 457 peak.

458 Cao et al. (2018) used  $T_2$  time of water proton to differentiate fresh blueberries from frozenthawed blueberries. They found that the freezing-thawing process caused a reduction in  $T_2$ 459 460 time of vacuole, cell wall, cytoplasm and extracellular water compared to the fresh sample. 461 With respect to the proton population of different compartments, it was observed that freezing 462 and thawing did not cause any alteration in the proton population of different compartments 463 compared to a fresh samples. Moreover, they also used  $T_2$  relaxation peak data to distinguish 464 different freezing conditions (i.e. freezing at - 20, - 40, - 80 °C and freezing by immersing in 465 liquid nitrogen). It was reported that among all freezing conditions, - 80 °C freezing 466 conditions better maintained  $T_2$  time of vacuole (p < 0.05) depicting better protection to the 467 vacuole membrane. Unlike them, we did not observe any significant difference among the 468 freezing trials with respect to the relaxation times of thawed samples frozen by different 469 methods, however, we could observe a significant differences between FF, IF and SF 470 conditions with respect to the proton population of  $T_2$  peaks. The proton population data (associated with  $T_2$  times) was used by Zhang et al. (2018) to study the effect of state/phase 471 472 transition on water mobility in frozen mango during 4-week storage.

## 473 **3.4. Drip loss**

474 Drip loss is one of the commonly and widely used methods to evaluate the freeze damage in 475 frozen products (especially in meat, fish, fruits and vegetable matrices). This method 476 estimates the freeze damage at a global level, or in other terms, provides an average value of 477 freeze damage of a product. In this section the impact of the studied freezing conditions on the 478 exudate loss from the product will be presented and discussed. Figure 10 shows the 479 dependence of drip loss on freezing rate. The results reveal that drip loss decreased slightly 480 when freezing rate was increased, however, significant difference was observed only between 481 samples that were frozen under FF and SF conditions. IF samples were not significantly 482 different (p > 0.05) compared to the samples of the other two freezing rates. The drip loss 483 results exhibited similar trends to those of  $T_{2e}$  component proton population from NMR that 484 showed that the application of -18 °C freezing protocol maintained less the intracellular 485 water content. This result may be attributed to a better preservation of the intracellular water 486 and to a lower damage of the pectocellulosic walls from the faster freezing rates. As a result, 487 this method was able to detect the differences between the selected freezing rates. The water 488 holding capacity of the frozen sample is linked to the size and location of ice crystals as well 489 as the thawing rate (Van Buggenhout, Messagie, et al., 2006). The formation of large ice 490 crystals in the cellular matrix can affect the water holding capacity of a cellular matrix in two 491 ways: (i) large ice crystals genesis during freezing can damage the cell membrane due to 492 mechanical effects, to cryo-concentration phenomena and shrinkage effects (particularly 493 during slow freezing rates) which in turn, will promote loss of mass during thawing 494 (Bevilacqua, Zaritzky, & Calvelo, 1979; Delgado & Sun, 2001; Sadot et al., 2017) and (ii) 495 higher drip loss may occur due to the formation of bigger ice crystals which correspond to 496 smaller specific surface area. This fact is associated with water re-absorption decrease during 497 thawing (Bevilacqua et al., 1979; Sadot et al., 2017). Charoenrein & Owcharoen (2016) and 498 Fuchigami, Hyakumoto, & Miyazaki (1995) observed decrease in exudate loss with 499 increasing freezing rates in frozen mangoes and carrots.

## 500 **3.5. Colour**

Freezing-thawing process significantly (p < 0.05) affected the colour parameters (L<sup>\*</sup>, a<sup>\*</sup>, and 501 b<sup>\*</sup> values) of the unblanched potatoes Figure 11. The L<sup>\*</sup> value (or lightness) and b<sup>\*</sup> value (or 502 vellowness) decreased, while the a\* value (redness) increased for potatoes after freezing-503 504 thawing (Figure 11a, b and c). These results are in agreement with the previously reported 505 study on freezing-thawing of unblanched potatoes (Koch et al., 1996). The colour change 506 during the freezing-thawing process of unblanched potato has been attributed to the browning 507 reaction that generally happens due to enzymatic activity during thawing process (Cano, 508 1996; Koch et al., 1996). The freezing rates had little effect on the colour parameters of 509 potatoes. Interestingly, it was found that FF process increased the redness value of potatoes 510 significantly (p < 0.05) than compared to SF process. Chassagne-berces, Fonseca, et al. 511 (2010) reported that freezing at – 80 °C increased the redness value of Golden Delicious apple 512 compared to the one frozen at – 20 °C. The redness value of IF samples was not significantly 513 different compared with those frozen under FF and SF conditions. No significant differences 514 for rest of the colour parameters (L<sup>\*</sup> value, b<sup>\*</sup> value and  $\Delta E$ ) were observed among the 515 freezing protocols.

## 516 **3.6. Microstructure analysis**

### 517 **3.6.1.** Cryo-SEM analysis

518 Scanning electron microscopy permits to obtain very high-quality images of a food matrix 519 microstructure. With respect to the frozen cellular matrices, micrographs acquired using SEM 520 (especially the cryo-SEM) provide important details about ice and cell morphology (shape of 521 the cell and the integrity of pectocellulosic walls). To the best of our knowledge, for the first 522 time, cryo-SEM was used to compare the changes occurring to the microstructure of potato at 523 different freezing rates. Figure 12 illustrates the SEM images of fresh and frozen potatoes. 524 The images of fresh and frozen potato were obtained using an environmental SEM (E-SEM) 525 and cryo-SEM, respectively. It can be seen in the figure that the fresh potato has polyhedral 526 shape cells with starch embedded in it. The microstructure morphology upon freezing 527 depended highly on the freezing rate being applied. The SF process not only created bigger 528 ice crystals in the cells but also caused the highest damage to the cellular structure. The cells 529 were highly distorted (deformed cells with broken and irregular cell wall structure) under SF 530 conditions. The IF process produced smaller ice crystals and maintained the cell wall integrity 531 better than the SF conditions. However, some cells lost their polyhedral shape and turned 532 almost into round shape when freezing was performed at IF conditions (cells pointed by the 533 orange arrow in Figure 12e and Figure 12f). The FF process yielded smallest ice crystals 534 compared to other freezing processes. Moreover, it can also be observed that the cellular 535 structure (in terms of cell shape i.e. polyhedral shape and cell wall integrity) was maintained 536 well. Similar to the authors, Bomben & King (1982), Chassagne-Berces et al. (2009) and 537 Chassagne-Berces, Fonseca, et al. (2010) cryo-SEM imaging could differentiate the apples 538 frozen by different freezing conditions based on microstructure. Moreover, they also observed 539 that slow freezing process (e.g. freezing at - 20 °C or 0.4 K/min) altered the shape of the cell 540 more than the fast freezing processes (e.g. freezing at - 80 °C or 450 K/min or by liquid 541 nitrogen immersion freezing). The calculations for ice crystals size were not made as it was

542 difficult to locate the boundary of the ice crystals. Moreover, the ice crystals had a 3D 543 structure and if the calculation were made, we could get only 2D information, this would have 544 led to an inaccurate estimation of the size of the ice crystals. Chassagne-Berces et al. (2009) 545 quantified the size of ice crystals formed at different freezing protocols (at - 20 °C, - 80 °C 546 and by immersion in liquid nitrogen) from the cryo-SEM images using gray level 547 granulometry based on mathematical morphology. They reported that the size of ice crystals was between 10 and 30  $\mu$ m after freezing at – 20 °C and below 5  $\mu$ m for faster freezing at 548 549 - 80 °C and by immersion in liquid nitrogen conditions. Since we were not able to quantify 550 the size of ice crystals, a direct comparison with their results remained difficult.

## 551 **3.6.2.** CLSM analysis

552 CLSM images of fresh and frozen-thawed potatoes are presented in Figure 13. This method 553 provides information about the status of the cell such as the shape of cells and integrity of the 554 cell wall. Compared to the fresh sample, the cells were highly disorganised and distorted in 555 SF samples. The altered shape of the cells and damaged cell wall structures in the slowly 556 frozen sample can be observed in the CLSM images (Figure 13). The buckled and folded cell 557 wall structure in SF samples indicated a major dehydration related damage that generally 558 happens at lower freezing rates (Chassagne-Berces et al., 2009; Gao & Critser, 2000; Mazur, 559 1977, 1984). IF process affected the cell shape as cell walls were found slightly distorted, 560 while IF seemed to have fairly preserved the integrity of the cellular structure and avoid cell 561 wall rupture as observed by SF. Moreover, extracellular spaces that can be generated due to 562 the freezing-thawing process were observed in IF samples (Figure 13c), while it was missing 563 in FF samples (Figure 13d). It can be seen (Figure 13c and Figure 14c) that these gaps were 564 adjacent to the shrunk cells, which clearly indicates the dislocation of water from the inner of 565 the cell to the extracellular space. It was difficult to distinguish freezing-thawing induced gaps 566 between the cells in SF sample as the structure was completely destroyed. FF condition 567 helped to preserve the original shape and integrity of the cellular structure. The results from 568 CLSM (in terms of cell morphology) were coherent with the results from cryo-SEM. 569 Charoenrein & Owcharoen (2016) used CLSM to study the effect of freezing rates and freeze-570 thaw cycles on the cellular structure of mangoes. Using this method, they were able to 571 observe the freezing-thawing related degradation of cellular structure. Moreover, based on 572 CLSM images, they were able to discriminate the different freezing protocols (i.e. freezing at 573 - 80, - 40 and - 20 °C). For instance, CLSM micrographs depicted that cells observed after 574 fast freezing (at - 80 °C) and thawing suffered a minimal amount of degradation; the cells

575 were still round and similar to cells from fresh tissues. The IF frozen (at -40 °C)-thawed 576 sample showed a slightly flat cellular structure, while the slow frozen (at - 20 °C)-thawed 577 mangoes exhibited larger changes. The cells of the slowly frozen samples lacked uniformity and some intercellular spaces were also observed in slowly frozen tissues. Sirijariyawat, 578 579 Charoenrein, & Barrett (2012) used CLSM to study the change in cellular morphology in 580 mangoes upon freezing-thawing. They reported that freezing  $(at - 50 \,^{\circ}C)$  followed by storage 581 (at - 20 °C chest freezer for 14 days) and thawing (at 4 °C for 2 h and kept at 25 °C for 30 582 min prior to analysis) of mango samples transformed the well-defined circular to elliptical 583 regular cells of fresh sample to irregular shaped cells with disintegrated cell wall.

## 584 **4.** Conclusions

585 In this study, potato samples were frozen under different freezing regimes and their quality 586 were evaluated using different techniques. The slowly frozen (-18 °C) potatoes exhibited 587 supercooling during freezing while no supercooling was noticed for other freezing conditions (i.e. -30 °C- intermediate freezing process and -74 °C- fast freezing process). The initial 588 589 freezing point (initial freezing temperature) could be detected for slow freezing and 590 intermediate freezing conditions, while it was hard to detect the initial freezing point for the 591 fast freezing condition. The initial freezing point temperature data for slow freezing and 592 intermediate freezing conditions revealed that a depression in freezing point occurred when 593 freezing rate was increased. The slow freezing process resulted in coarser ice crystals and also 594 caused the highest damage to the cellular structure. The cells were highly distorted (deformed 595 cells with buckled and folded cell wall structure) when the slow freezing condition was used. 596 Intermediate freezing rate process led to relatively fine ice crystals compared to slow freezing 597 process. Although IF process led to greater cell wall structure integrity, however, it was not 598 able to preserve the cell shape. The fast freezing process not only promoted the formation of 599 very fine ice crystals but also preserved the morphology of the cells. The NMR analytical 600 parameters, texture and drip loss showed limitations to differentiate the different freezing 601 protocols. . None of the freezing protocols preserved the colour parameters of the fresh potato.

## 602 Benchmarking of freeze damage assessment methods for vegetables on the basis of 603 efficiency, accuracy, cost-of operation, and ease of operation

In this section, freeze damage assessment methods used during this study were evaluated based on various benchmarking parameters and a decision table (dedicated to benchmarking study) was proposed. Table 3 summarizes the results obtained within this study. Pros and cons 607 related to the different methods considered in this study for assessing the freeze damage in 608 potato were discussed.. These observations would provide useful information about the 609 analytical techniques that can be used to estimate freeze damage efficiently.

The focused freeze damage assessment technologies like CLSM, and global freeze damage assessment techniques like texture analysis (also includes laser-puff texture analysis), low field NMR relaxometry, and colour analysis tests used in this study was found to be relevant methods to distinguish the fresh samples from frozen/thawed sample.

The acquired results suggest that cryo-SEM and CLSM are suitable for validating minor quality changes among the different freezing protocols. Meanwhile, the global methods such as texture, NMR, and drip loss can only reflect larger quality changes.

617 In order to compare freezing protocols, colour analysis was found to be an unsuitable618 parameter.

619 Efficiency and accuracy wise, cryo-SEM and CLSM can be termed as best methods to 620 analyze the freezing injuries.

NMR, cryo-SEM and CLSM techniques are expensive techniques, whereas texture analysis,
drip loss measurements, colour analysis are cost-effective technologies. The analyses time for
NMR is long, while other methods take substantially less time.

624 From a global point of view, a debate could be opened on which is the most relevant 625 technique to assess freeze damage. Two key parameters can be tackled, (i) the size of the field 626 that is embraced by the technique and (ii) the representative size of the technique. For 627 example, NMR will tackle a sample of ca 1 cm and will provide information at the level of a water molecule (2.75 Å). Cryo-SEM will look at field of ca 100 µm with information at 628 629 nanometer scale, even though in this case the freeze damage is observed at the scale of an ice 630 crystal (circa 10 µm). The ratio between the size of field and the representative size of 631 technique could be considered as a kind of "freeze damage assessment index" (FDA Index) to 632 assess the relevance of each technique. The higher the FDA index will be, the useful will be 633 (a priori) the technique. From such point of view NMR looks like the best candidate, even 634 though the interpretation and quantification of the freeze damage based on  $T_1$  and  $T_2$  values 635 are not very well documented in the literature and still remain quite subjective. The major 636 concerns relates to the size of the field. Observation of the freeze damage on a single cell can 637 be very informative and detailed, but ca a hundred of cell should be analyzed to obtain an 638 averaged information, which is out of reach for time reasons. Another aspect relies on the 639 possibility to repeat the analysis and on the time needed for a single measurement. CLSM and 640 Cryo-SEM bring informative images that can help to visualize the defaults and that can 641 support observations done with other techniques at a broader field like NMR, texture, drip 642 losses. The precision of the analytical method is one of the most important factors when choosing the 643 methods for freeze damage assessment. A method that imparts minimum error to the measured result 644 will help in better understanding of the impact of process conditions on the sample. Based on the error 645 calculations made for the quantitative methods (Table 3), the drip loss measurement seemed to be 646 more satisfactory, followed by colour analysis, hardness analysis, deformation analysis by the laser-647 puff tester, NMR analysis and Young's Modulus analysis. Overall, the error imparted by the 648 analytical techniques were quite low in values.

649 The proper assessment of freeze damage remains a challenge and requires a mass of 650 experimental work before drawing any conclusion. However, this comparative study is maybe 651 the first one that proposes a benchmarking of so many different analytical techniques often 652 considered to assess the freeze damage.

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661

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## 784 **Captions for the Figures**

- Figure 1: Load-strain curve providing detail about the firmness of the potato during thecompression tests.
- 788789 Figure 2: Laser-puff firmness testing prototype.

Figure 3: A representative deformation curve obtained while performing the laser puff test.
Conversion of volts to mm: 1 V = 2 mm.

- Figure 4: Illustration of sample preparation step prior to imaging in cryo-SEM.
- Figure 5: Protocol followed for CLSM imaging of unblanched potato.
- Figure 6: Representative freezing curves of potatoes under different freezing methods.
- Figure 7: (a) Deformation curve obtained for fresh and frozen-thawed samples (conversion of volts to mm: 1 V = 2 mm) and (b) deformation values (in mm) for fresh and frozen-thawed potatoes (under different freezing rates) acquired using laser-puff firmness tester. SF (slow freezing at -18 °C), IF (intermediate freezing at -30 °C) and FF (fast freezing at -74 °C).
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- Figure 8: Distributions of relaxation peak components ((a)  $T_2^*$  and (b)  $T_2$ ) at -20 °C of potato samples frozen by various freezing protocols: slow freezing at -18 °C (red lines), intermediate freezing at -30 °C (blue lines), and fast freezing at -74 °C (black lines). (The x-axis correspond to the relaxation times expressed in ms (milliseconds).
- Figure 9:  $T_2$  relaxation peak data of frozen-thawed potatoes at 4 °C: (a) fresh sample, (b) after freezing at -18 °C (c) after freezing at -30 °C and (d) after freezing at -74 °C.
- 811 Figure 10: Effect of different freezing conditions on the drip loss of potato.

812 Figure 11: Effect of freezing protocols on color parameters (a - L<sup>\*</sup> value or Lightness, b - a<sup>\*</sup> 813 value or Redness, c - b<sup>\*</sup> value or Yellowness and d -  $\Delta E$ ) of potato. Mean values of 9 814 repetitions are represented with confidence interval.

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816 Figure 12: Microstructure of potato before and after freezing under the different freezing 817 protocols. (a, b) environmental SEM images of the fresh cell showing the cellular structure and starch granules imbedded into it. Cryo-SEM after freezing at -18 °C-SF (c, d), at -30818 819 °C-IF (e, f) and at -74 °C-FF (g, h), respectively. White colored arrows in images are 820 pointing the cells containing ice crystals. Red arrow showing the area where the breakdown of 821 cell structure happened. Orange arrows indicate the cells that might have transformed from 822 polyhedral to almost round shape. Other abbreviations in the picture are A: air space; S: starch 823 granule; W: cell wall structure.

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Figure 13: Microstructure evaluation using CLSM: (a) fresh potato, (b) frozen-thawed after freezing at -18 °C-SF, (c) at -30 °C-IF and (d) at -74 °C-FF. Other abbreviations in the picture are S: starch granule, W: cell wall, S<sub>p</sub> is probably the gap created when of cell moved apart from each other as a consequence of freezing-thawing. The red arrows indicating the deformed, distorted and shrunk cells. Orange arrows indicating loss of regularity of the cell 830 walls compared to fresh and FF samples. Yellow arrows showing the folded and buckled cell

831 wall structure formed due to SF process.

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Figure 14: Microstructure evaluation using CLSM: (a) single cell of fresh potato, (b) single potato cell (imaged in frozen state) after freezing at -18 °C (SF), (c) frozen-thawed potato structure after frozen at -30 °C (IF) and (d) frozen-thawed potato structure after frozen at -74 °C(FF) (d). S<sub>p</sub> is probably the gap between the cells created due to freezing-thawing process. Black arrow showing a part of broken cell wall. White arrow indicating discontinuity in the cell wall structure. Red arrow evince the deformed cell with irregular cell wall structure (orange arrows).

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850	Captions for Tables
851	Table 1. Effects of different freezing protocols on the freezing properties of potatoes.
852	Table 2. Textural parameters measured for potatoes under different freezing conditions.
853	Table 3. Benchmarking study on freeze damage assessment methods.
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Freezing condition	Initial freezing point (°C)	Characteristic freezing time (min)	Overall freezing time (min)	Overall freezing rate (°C/min)
- 18 °C (SF)	$-0.3 \pm 0.14$	29.12 <u>+</u> 3.94	$72.30 \pm 0.14$	$0.48 \pm 0.00$
– 30 °C (IF)	$-0.73 \pm 0.06$	17.18 <u>+</u> 0.79	26.31 ± 0.62	$1.36 \pm 0.05$
– 74 °C (FF)	n.d.	8.52 <u>+</u> 1.53	14.51 <u>+</u> 1.49	2.51 ± 0.25

Table 1. Effects of different freezing protocols on the freezing properties of potatoes.

	Parameters	Hardness (N)	Young's modulus (MPa)
	Fresh	$190 \pm 19^{a}$	$5.46 \pm 0.44^{a}$
	SF (at – 18 °C)	$47 \pm 11^{\circ}$	$1.37 \pm 0.33^{\circ}$
	IF (at – 30 °C)	$68 \pm 16^{b, c}$	$2.09 \pm 0.41^{b, c}$
	FF (at – 74 °C)	$90 \pm 8^{b}$	$2.60 \pm 0.45^{b}$
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883 Table 2. Textural parameters measured for potatoes under different freezing conditions.

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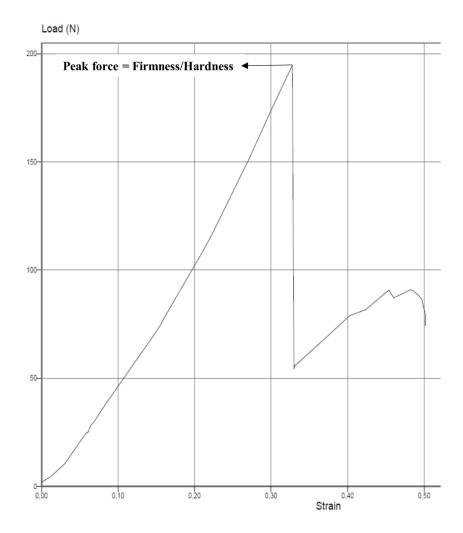
906	Table 3.	Benchmarking study of	n freeze damage assessment	methods.
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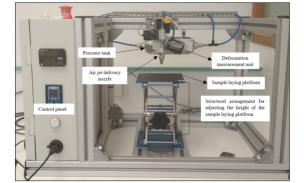
F	Focused Methods		Global methods				
Freeze damage assessment methods for	Cryo-SEM CLS		Texture Analysis		NMR	Drip loss	Colour
fruits and vegetables			Conventional	Laser-Puff	-		
Sample preparation	Difficult	Difficult	Easy	Easy	Easy	Easy	Easy
Ability to detect differences between fresh and	×	++++	++++	++++	++++	++++	++++
frozen/thawed sample							
Ability to distinguish different freezing protocols	++++	++++	++	++	+	++	-
Analysis time (sample preparations + data acquisition and treatment)	+	+	+	+	+++	+	+
Interpretation of measured analytical parameters	Easy	Easy	Easy	Easy	Difficult	Easy	Easy
Nature of sample	F	F/T	F/T	F/T	F & F/T	F/T	F/T
Cost of operation	High	High	Low cost	Low cost	High	Low cost	Low cost
Status of method	OU	NUO	VC	NM	NUO	VC	VC
Error (%)	×	×	Hardness = $\pm 0.5$ YM = $\pm 3.2$	<u>+</u> 0.90	<±1	±0.0016	$L = \pm 0.02$ $a = \pm 0.21$ $b = \pm 0.15$

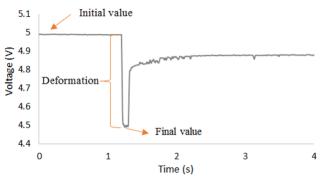
+ = Lowest value; ++++ = Highest value; × = Not applicable; - = No effect.

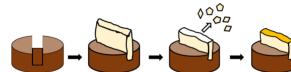
Abbreviations of the words: F = Frozen; F/T = Frozen/Thawed; FD = Freeze dried; OU = Often used; NUO = Not used often; VC = Very common; NM = New method; YM = Young's Modulus.

L, a, b are different colour parameters.









Copper sample holder with its groove (10 mm diam. 10 mm height. Groove 3x3 mm section) Fixing of thin slice of sample in the sample holder

Cryo-fracturing the sample followed by sublimation for 10 min Metal coating (gold coating) followed by imaging  $\frac{Sampling}{Cylindrical shape (diameter (<math>\emptyset$ ) = 8 mm & height (H) = 5 mm); weight = 0.25 g

Sample was stained in Acridine Orange (0.01%) (pH 7.2) for 2 hours at 4 °C

Freezing under different conditions (at - 18 °C, - 30 °C and - 74 °C)

Thawing of sample at room temperature, followed by slicing using a microtome (80 µm slice was used for imaging study).

CLSM imaging was then performed using different resolution objective lenses (for e.g. 20X, 10X, 4X, etc.)

#### Staining the sample



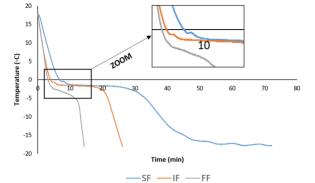
#### Preparation for imaging

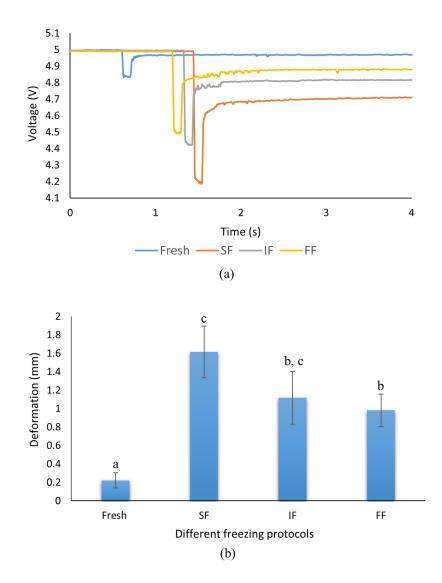


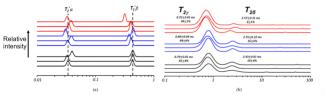
Confocal Laser Scanning Microscope

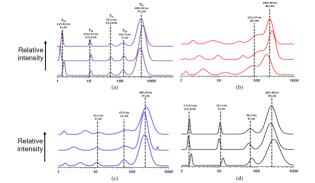


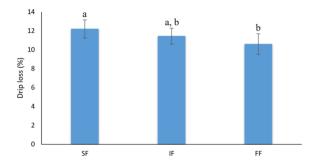
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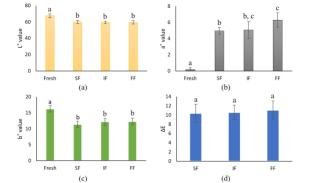


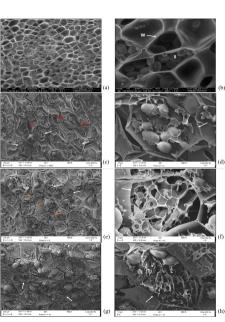


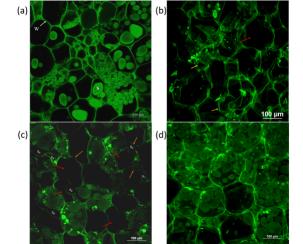


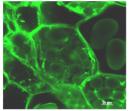


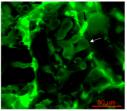






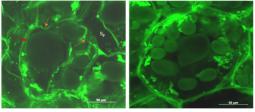












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