

1 Physiochemical and microbiological quality of lightly processed salmon
2 (*Salmo salar* L.) stored under modified atmosphere.

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16 Abstract

17 Low-temperature cooking such as sous-vide has become a favored method for processing seafood. In
18 order for this method to be applicable for retail products, combinations with other processing steps
19 are needed to keep the products safe and durable while maintaining high quality. The present
20 experiments were designed to investigate the influence of low-temperature treatment (40, 50, or 60
21 °C) in combination with various packaging technologies (modified atmosphere (MA) or soluble gas
22 stabilization (SGS)) on both the microbial growth as well as the physiochemical quality. Salmon loins
23 were either kept natural or inoculated with *Listeria innocua* prior to drying (16-18h) in either 100% CO₂
24 (SGS) or atmospheric air (MA packaging). All samples were sous-vide treated, repackaged in MA and
25 stored at 4 °C for 24 days. The results showed shelf life to be significantly improved with the
26 implementation of SGS, prolonging the of lag-phase and slowing the growth rate of both naturally
27 occurring and inoculated bacteria. Variations in packaging technology did not significantly influence
28 any of the tested quality parameters including drip loss, surface color, and texture. Consumers
29 increasing demand for lightly processed seafood products makes *Listeria spp.* an increased problem,
30 however the present experiment has shown that it is possible to lower processing temperatures to as
31 little as 40 or 50 °C and still obtain inhibition of *Listeria*, but with improved chemical quality compared
32 to traditional processing.

33 Keywords

34 Atlantic salmon, lightly processed, *Listeria spp.*, microbial quality, modified atmosphere packaging,
35 physiochemical quality, soluble gas stabilization, sous-vide.

36 1. Introduction

37 Easy-to-prepare and ready-to-eat meals are increasingly perceived as an optimal solution in a modern
38 lifestyle. This has led to a tremendous increase in the demand for tasty, nutritious, high quality, and
39 yet convenient food products.

40 Seafood is the second largest export sector in Norway, after oil and gas, and salmon make up the
41 largest individual product group within this sector. This has led to Norway being the world's largest
42 producer of farmed salmon (Asche, Roll, & Tveteras, 2009). Currently, Norwegian salmon is mostly sold
43 in whole fresh or frozen form (Straume, 2017) however, contemporary trends for consumption of
44 lightly processed seafood (Speranza, Corbo, Conte, Sinigaglia, & Del Nobile, 2009) have increased the
45 market for valueadded salmon products. This emphasizing the need for devolping and testing of
46 processing methods which allow production of tasty, safe, and durable salmon products with minimal
47 heat treatment.

48 The consumer demands make sous-vide cooking a favored light processing option. By heating in sealed
49 pouches at a lower temperature, sous-vide cooking offers multiple benefits compared to traditional
50 cooking of seafood (Baldwin, 2012). The benefits include reduced heat damage to proteins and lipids
51 and diminishing the loss of liquid, nutrients, and aromatic compounds while improving the perceived
52 texture (Singh et al., 2016). Two factors are influential in relation to the shelf life extension obtain by
53 sous-vide cooking; the intensity of the applied heat and the control of subsequent storage
54 temperatures (García-Linares, Gonzalez-Fandos, García-Fernández, & García-Arias, 2004). Increased
55 temperature or time during heat treatment has the potential to prolong the shelf life further, but at
56 the same time leads to a significant decrease in organoleptic quality. Low storage temperature is
57 necessary to ensure the microbial safety of sous-vide cooked products (García-Linares et al., 2004), yet
58 many food products are subject to temperature abuse during transport, selling, or storage. This
59 emphasizes the need for further preservation steps in a form of hurdle technology (Baldwin, 2012).
60 Multiple technologies are being used for this purpose, and modified atmosphere (MA) packaging has
61 become a well-established method (Bouletis, Arvanitoyannis, & Hadjichristodoulou, 2017; Lambert,
62 Smith, & Dodds, 1991).

63 Depending on species and temperature MA packaging has been found to extend shelf life of seafood
64 products by several days compared to air storage (Powell & Tamplin, 2012; Sivertsvik, Rosnes, &

65 Kleiberg, 2003; Speranza et al., 2009; Torrieri, Cavella, Villani, & Masi, 2006; Tsironi & Taukis, 2010;
66 Özogul, Polat, & Özogul, 2004). The amount of dissolved CO₂ in the foods is proportional to the
67 inhibitory effect of MA packaging (Devlieghere, Debevere, & Van Impe, 1998a, 1998b). Thus
68 constricting the optimal use of MA packaging by the need for a high gas to product ratio to avoid
69 packaging deformation due to CO₂ dissolvment when high CO₂ levels are introduced (Rotabakk,
70 Birkeland, Jeksrud, & Sivertsvik, 2006). Dissolvment of CO₂ prior to retail packaging, a method known
71 as soluble gas stabilization (SGS) (Sivertsvik, 2000) has the ability to overcome this drawback.
72 Regardless of the choice of modified atmosphere applied, the altering of the gas composition in the
73 packages also alters the microbial community (Yesudhason, Lalitha, Gopal, & Ravishankar, 2014). The
74 identified dominant spoilage strains for MA packaged seafood includes lactic acid bacteria (LAB) (Gram
75 & Huss, 1996), *Brochotrix thermosphacta* (Macé et al., 2012; Sivertsvik, 2003) as well as
76 *Photobacterium phosphoreum* (Dalgaard, Mejlholm, Christiansen, & Huss, 1997). Both *B.*
77 *thermosphacta* and *P. phosphorerum* have been shown to be limited by either heat (Gram & Huss,
78 1996) or by CO₂ levels equivalent to those obtained by SGS-treatment (Abel, Rotabakk, & Lerfall, 2019).
79 The processing inhibition of aerobic spoilage microflora has the potential to rendering the food unsafe
80 for consumptions before it appears spoiled (Sivertsvik, Jeksrud, & Rosnes, 2002), thus making the
81 control of pathogens such as *Listeria monocytogenes* and *Clostridium spp.* an even more important
82 task.

83 Multiple studies have been performed on the effect of either heat-treatment or packaging technology
84 on seafood shelf life or product quality; however, research regarding combinations of such
85 technologies on *both* shelf life and product quality are limited. Hence, the aim of this study is to gain
86 knowledge of quality deterioration and microbial development in lightly processed salmon, by studying
87 the effect of combined low heat treatment, MA packaging, and SGS technology on the microbial load
88 as well as perceived product quality parameters.

89 2. Materials and methods

90 A three-factor challenge- storage study was conducted, the factors being degree of heat treatment
91 (core temperature of 40, 50, or 60 °C), packaging technology (MA packaging or SGS followed by MA
92 packaging), and microbial flora (natural or inoculated with *Listeria innocua*) (Table 1). The microbial
93 and physiochemical development was evaluated continuously for a period of 24 days (at day 0, 6, 10,
94 13, 17 or 24). The experiments were executed in two rounds, separated based on choice of packaging
95 technology.

96

97 2.2 Raw material

98 Pre-rigor filleted farmed Atlantic salmon (*Salmo salar L.*) (fillet weight of 1-1.4kg) were obtained from
99 Salmar AS (Frøya, Norway). Fillets were obtained on the day of slaughtering and stored in a fridge
100 (4.2±2.4 °C) on ice for 3 days to ensure post rigor state before processing. Backfins, belly flaps, and
101 tails were trimmed in order to obtain a product with equal height (approx. 3cm). Fillets were portioned
102 into equal size of 79.8±2.3g (Figure 1).

103 2.3 Bacterial strains

104 A pure *L. innocua* culture (-80 °C) (ATCC 33090) were obtained from the culture collection at University
105 of Gothenburg (CCGU). The cultures were thawed and recovered on brain heart infusion (BHI) agar
106 (Oxoid CM1136, Oxoid Ltd., Basingstoke, UK) at 37 °C for 24 hours. Single colonies were inoculated
107 into separate vials of BHI broth (CM1032, Oxoid Ltd., Basingstoke, UK) for enrichment and incubated
108 at 8 °C for 5 days. The procedure resulted in cold-adapted cultures in an early stationary growth phase.
109 Samples were diluted to OD₆₀₀ of approximately 0.1 (0.104-0.110) in order to obtain a cell
110 concentration of approximately 1x10⁵ colony forming units (CFU) x ml⁻¹ (2.7x10⁵ CFU x ml⁻¹).

111 2.4 Inoculation

112 100 µL inoculum were dispersed on the surface of half the samples (estimated 10^4 CFU x surface⁻¹) and
113 all the samples were air-dried for 10 min. The rest of the samples were kept natural, without any
114 inoculation. All the samples were packed in batches (n=13) on trays (C2325-1C, Færch Plast, Holstebro,
115 Denmark) in vacuum pouches (425x650 mm PA/PE sous-vide pouch, Maske AS, Trondheim, Norway,
116 filling degree approx. 17%). The pouches were filled with either atmospheric air (in case of MA
117 packaged samples) or pure CO₂ in excess (in case of SGS samples) using a chamber machine
118 (Webomatic SuperMax s3000, Webomatic, Bochum, Germany). Samples were stored at 3.7 ± 0.5 °C for
119 16-18 h to dry completely and to ensure CO₂ saturation of the SGS-samples. Four replicates were
120 prepared and analysed of each sample at each sampling point, a total of 28 for each treatment group.

121 2.5 Heat treatment and packaging

122 Samples were repacked in vacuum pouches (135x180 mm PA/PE sous-vide pouch, Maske AS,
123 Trondheim, Norway) using a chamber machine (Webomatic SuperMax s3000, Webomatic, Bochum,
124 Germany). A sous vide water bath (Diamond M, Fusionchef by Julaba, Germany) was used for all heat
125 treatments. Temperatures were 45 °C (44.6 ± 0.4 °C), 55 °C (54.5 ± 0.2 °C), or 65 °C (64.6 ± 0.1 °C).
126 Treatment times were 15, 18, or 21 min, respectively. Treatment times were chosen based on pre-
127 experiments conducted to establish time needed to obtain a core temperature 5 °C lower than the
128 water bath temperature (core temperature of 40, 50, or 60 °C, respectively). All sample pouches were
129 cooled in ice water and fish samples repackaed in 300 ml semi-rigid crystalline polyethylene
130 terephthalate (CPET) trays (C2125-1B, Færch Plast, Holstebro, Denmark) using a semi-automatic tray
131 sealing packaging machine (TL250, Webomatic, Bochum, Germany). All trays were equipped with an
132 absorbent. During packaging, the air was evacuated (final vacuum pressure of 25 mbar) and flushed
133 with the pre-set MA gas mixture prior to application of a cover film comprised of a 40 µm combination
134 of polyethylene (PE), ethylene vinyl alcohol (EVOH), polyamide (PA), and polyethylene terephthalate
135 (PET) (Topaz B-440 AF, Plastopil, Almere, The Netherlands). Food grade CO₂ and N₂ were mixed to 60%

136 CO₂ balanced with N₂ (both MA and SGS) using a gas mixer (MAP Mix 9000, Dansensor, Ringsted,
137 Denmark). All handling were done aseptically. Oxygen transmission rate (OTR) was 66-78 cm³ x 25 μm
138 x m⁻² x 24 h¹ x bar¹ at 23 °C for the tray, 2.5 cm³ x 40 μm x m⁻² x 24 h¹ x atm¹ at 23 °C for the cover film,
139 and 50 cm³/m² x 24 h¹ x bar¹ at 23 °C for the vacuum pouches. Packaging resulted in a sample filling
140 degree of approximately 1:3.

141 After packaging, the trays were stored at 2.4±1.0 °C for up to 24 days.

142 2.6 Chemical analysis

143 2.6.1 Headspace gas analysis

144 The headspace gas composition (% O₂ and CO₂) was measured using an oxygen and carbon dioxide
145 analyzer (Checkmate 9900 analyzer, PBI-Dansensor, Ringsted, Denmark) as described by Abel,
146 Rotabakk, Rustad, and Lerfall (2018). The gas compositions were measured at storage day 0, 6, 10, 13,
147 17, and 24.

148 2.6.2 Water-, lipid-, and protein content

149 Water content was determined gravimetrically by drying the samples for 24 hours at 105°C (ISO.6496,
150 1983). Lipids were extracted and the total amount calculated gravimetrically as described by Bligh and
151 Dyer (1959). Protein content was calculated based on the total Kjeldahl nitrogen method, using an
152 automated Kjeldahl digester (KjeldDigester K-449, Büchi, Flawil, Switzerland) and titration-system
153 (KjelMaster K-375, Büchi, Flawil, Switzerland) equipped with an autosampler (KjelSampler K-376,
154 Büchi, Flawil, Switzerland). Only the raw material underwent composition analysis.

155 2.6.3 Drip loss and water holding capacity (WHC)

156 Drip loss was calculated by the difference in weight of the tray plus absorbent between day 0 and days
157 6, 10, 13, 17, and 24. WHC was measured as described by Skipnes, Østby, and Hendrickx (2007) using
158 metal carriers (Part No. 4750, Hettich Lab Technology, Germany) and centrifuged (Rotina 420 R,

159 Hettich centrifuge) for 15 min at 4 °C, using a free swing rotor at RCF = 530×g. The WHC was measured
160 in triplicates of each group on day 6 and 24 of storage to obtain start and end values.

161 2.6.4 Surface color

162 Sample surface color (CIE Lab) was assessed by a digital photo imaging color-measuring system
163 (DigiEye full system, VeriVide Ltd., Leicester, UK). Analysis were carried out in a standardized lightbox
164 (6400 K) using a digital camera (Nikon D7000, 35 mm lens, Nikon Corp. Japan). The pictures were
165 analyzed with DigiPix software ver 2.8.0.2 (VeriVide Ltd., Leicester, UK). Changes in perceived color
166 were calculated as ΔE in accordance with the formula $E = \sqrt{S_L^2 + S_C^2 + S_H^2}$ where $S_L = \Delta L/2$, $S_C =$
167 $\Delta C/(1 + 0.048 * C_1)$, $S_H = \Delta H_{ab}/(1 + 0.014 * C_1)$ and $\Delta H_{ab} = \sqrt{\Delta a^2 + \Delta b^2 - \Delta C^2}$ as described by
168 CIE Int. Commission on Illumination (1994). ΔE values higher than 4 are normally visible to the human
169 eye (Lerfall, 2011).

170 2.6.5 Texture

171 Instrumental textural analyses were performed using a Texture Analyzer TA-XT2 (SMS Ltd., Surrey,
172 England) fitted with a 30 kg load cell and a Warner Bratzler probe (SMS Ltd., Surrey, England). The
173 force-time graph was obtained by the Texture Exponent software for Windows (version 6.1.7.0, SMS
174 Ltd., Surrey, England), which was used for the data analyses. The analyses were performed in four
175 times replicates for each group immediately after processing and cooling. The analysis was done at a
176 speed of 1 mm x s⁻¹, and measurements were performed until 100% penetration was achieved.
177 Portioned raw and treated samples (Figure 1) were placed with the probe adjacent to the mid line, to
178 ensure measures were a result of shearing rather than flaking of the muscle fibers.

179 2.6.6 Degradation products of adenosine triphosphate (ATP)

180 Degradation products of ATP was analysed on a Phenomenex synergi 4u hydro-RP80 A (150×4.6mm,
181 4µm) HPLC column after a method by Sellevold, Jynge, and Aarstad (1986), using an Agilent 1290
182 chromatograph (Agilent technologies, Paolo Alto, CA, USA) (isocratic, flow 1.0 mL/min) connected to

183 an Agilent 1260 diode array UV-VIS detector, as described by Lerfall, Jakobsen, and Bjørge Thomassen
184 (2018). Standard curves of ATP (Sigma, ≥99%, CAS:34369-07-8), ADP (Sigma, ≥95%, CAS:20398-34-9),
185 AMP (Sigma, ≥99%, CAS:149022-20-8), IMP (Sigma, ≥98%, CAS:352195-40-5), HxR (Sigma, ≥99%,
186 CAS:58-63-9) and Hx (Sigma, ≥99.0% CAS:68-94-0) in deionized water were used for identification of
187 quantification.

188 2.7 Microbial analysis

189 Microbial analyses were prepared using 10 g of fish sampled aseptically from the inoculated surface.
190 The fish sample was homogenized in 90ml sterile 0.85% NaCl (w/v) and 0.1% peptone (w/v) water for
191 60 sec. Decimal dilution series were prepared in similar solution in accordance with NMKL-standard 91
192 (NMKL, 2010).

193 Natural bacterial flora (NBF) samples were analyzed for total aerobic plate count, *Clostridium* spores,
194 *Listeria* spp., and LAB. Inoculated samples were analyzed for total aerobic plate count and *Listeria* spp.
195 Negative control samples were tested for total aerobic plate count, *Clostridium* spores, *Listeria* spp.,
196 and LAB.

197 Total aerobic plate count was analyzed as pour plates with a top layer of Lyngby iron (LI)-agar
198 (CM0964, Oxoid Ltd., Basingstoke, UK) prepared as described by the manufacturer, and incubated
199 aerobically at 22 °C (21.6±0.4 °C) for 3 days, in accordance with NMKL-184 (NMKL, 2006). Presence of
200 sulfite-reducing *Clostridium* spores was analyzed in accordance with NMKL-56 (NMKL, 2008) on Shahidi
201 Ferguson Perfringens (SFP) agar base (DIFCO28110, Thermo Fisher Scientific, Waltham, MA, USA)
202 prepared as described by the manufacturer, but without the addition of egg yolk. Dilution of the
203 sample material was heated at 80 °C for 10 minutes prior to plating in order to inhibit any vegetative
204 cells. Samples were incubated anaerobically at 15 °C (15.1±0.6 °C) for 5 days. Presence of *Listeria* spp.
205 were tested on Brilliance™ listeria agar (BLA) containing Brilliance™ listeria selective supplement
206 (Oxoid CM1080 and Oxoid SR0227, Oxoid Ltd., Basingstoke, UK) prepared as described by the
207 manufacturer, and incubated aerobically at 37 °C (37.1±0.2 °C) for 24±2 h. Presence of LAB was tested

208 as described in NMKL-140 (NMKL, 2007) on de Man, Rogosa and Sharpe (MRS)-agar (Oxoid CM0361,
209 Oxoid Ltd., Basingstoke, UK) with 10 mg/l amphotericin B, and inoculated at 25 °C (25.8±0.2 °C) for 3
210 days.

211 Sampling was performed of the raw material, inoculated samples, and on all stored samples after 0, 6,
212 10, 13, 17, and 24 days storage in accordance with experimental design (Table 1).

213 2.8 Statistics

214 Statistical analyses included outlier test (Grubbs outlier test at level $p < 0.05$), analysis of variance
215 (ANOVA) and general linear modeling (GLM, Tukey's HSD test at level $p < 0.05$). All data processing were
216 carried out using Minitab 17.0 (Minitab, Coventry, UK). To meet the requirements of equal variance
217 and normal distribution, all statistical analyses of microbial growth were done on log-transformed
218 data.

219 Data were analysed in 4 time replicate and is presented as mean \pm standard deviation (SD) unless
220 otherwise stated.

221 3. Results and discussion

222 A GLM showed all parameters (core temperature, packaging technology, and storage time) as well as
223 all the interaction effects to be of significant influence on the amount of microbial growth ($p < 0.001$)
224 for both natural- and inoculated samples. No correlation was found between packaging technology
225 and color, WHC, drip loss, or formation of ATP-degradation products. Processing temperature
226 influenced all of the tested parameters ($p < 0.001$)

227 3.1 Quality analyses

228 A concern when it comes to the implementation of SGS is the influence on the quality of the product.
229 The appearance of food products is of major importance to consumers, both with regards to
230 acceptability and preference. When it comes to salmon, the color is generally perceived as one of the
231 most important quality parameters (Anderson, 2000). In the present experiments, the only results that

232 yielded ΔE values above the noticeable limit was comparisons between temperature treatments,
233 showing increased lightness (L^*) and decreased redness (a^*) and yellowness (b^*) with increasing
234 temperatures (Figure 2), as reported by Bhattacharya, Choudhury, and Studebaker (1994). This result
235 can be explained by the increased protein denaturation and coagulation of sarcoplasmic proteins on
236 the surface caused by the increased temperatures. Broadly speaking, fish muscle proteins are
237 separated into three groups, with the more important proteins being myosin, actin and the
238 sarcoplasmic proteins. Multiple studies have demonstrated the temperature stability of these
239 proteins, and it is generally agreed that heat denaturation of myosin in salmon occurs in a range of 43-
240 50 °C, actin around 76-78 °C and the sarcoplasmic proteins, which a more diverse group, in a broad
241 range from 57-67 °C (Ovissipour, Rasco, Tang, & Sablani, 2017). In the present study, the heat
242 treatments were carried out at 45, 55, and 65 °C, respectively. This would indicate that only limited
243 protein denaturation would have taken place at 45 °C, whereas myosin would be completely
244 denatured at 55 °C and at least some of the sarcoplasmic proteins in the 65 °C samples. Temperatures
245 never reach levels of actin denaturation.

246 The results of the color analysis also show that neither choice of packaging technology nor storage
247 time gave rise to any perceivable changes in color (average $a^*=14.5\pm3.4$, $b^*=9.4\pm2.2$, $L^*=67.0\pm2.4$,
248 $\Delta E=2.2\pm1.3$). These findings are in agreement with those by Rotabakk, Birkeland, Lekang, and Sivertsvik
249 (2008) on halibut, or by Mendes and Gonçalves (2008) in sea bream and sea brass. As for the color
250 analysis, WHC was found not to be significantly influenced by treatment temperature, packaging
251 technology, or duration of storage ($p=0.054-0.926$). Average WHC was measured to be 70.9%. The lack
252 of differences in WHC is in agreement with the fact that no significant differences ($p<0.001$) were
253 observed in drip loss as a result of either temperature or packaging technology (Table 2). In contrast,
254 a close relationship has previously been established between protein denaturation and WHC (Kong,
255 Tang, Rasco, Crapo, & Smiley, 2007) and hence drip loss. WHC is highly dependent on the properties
256 of myosin, thus expecting WHC to decrease once myosin denaturation temperatures have been
257 reached (Ofstad, Kidman, Myklebust, & Hermansson, 1993). The reason for the discrepancy is not

258 understood. Moreover, it has been reported that an increase in dissolved CO₂ will alter WHC and
259 increase drip loss (Davis, 1998; Randell et al., 1999) as reported for halibut (Rotabakk et al., 2008) and
260 shrimps (Sivertsvik & Birkeland, 2006). The effect was then ascribed to volume reduction caused by
261 the uptake of CO₂ by the product, an effect which is counteracted by the use of SGS prior to MA
262 packaging (Rotabakk et al., 2008). An alternative explanation could be that the presence of CO₂ during
263 the heat treatment increased the cook loss thereby limiting the drip loss later in the storage; however,
264 cook loss was not measured in the present study. The significant differences between processing
265 temperature in results for all the tested quality parameters show the potential to improve quality by
266 lowering of the processing temperature. At the same time, the lack of significant differences between
267 packaging technology treatments facilitates the lowering of this temperature without affecting the
268 quality by itself.

269 Just as perceived color, tenderness of foods is another important parameters regarding consumer
270 satisfaction (Bhattacharya, Choudhury, & Studebaker, 1993). **Particularly** the shear force has been
271 cited as an influential factor for consumer opinions (Jonsson, Sigurgisladottir, Hafsteinsson, &
272 Kristbergsson, 2001; Sigurgisladottir et al., 1999). Protein denaturation is known to play an important
273 role in the toughening of the texture of muscle products (Hatae, Yoshimatsu, & Matsumoto, 1990).
274 The influence of both treatment temperature and packaging technology were analyzed, but the results
275 were inconclusive as variation in raw material were found to be bigger than variations between
276 treatments (**data not shown**).

277 Fish deterioration is monitored in many different ways, common amongst these are ATP-degradation
278 products, which is considered a good indicator of fish freshness (Shumilina et al., 2016). Post-mortem
279 degradation of ATP in fish muscle occurs due to a combination of endogenous and bacterial enzymes
280 and goes through the intermediate products ADP, AMP, IMP, HxR, and Hx. Most important degradation
281 products include Hx and IMP, which has been associated with the development of unpleasant and
282 enhancing flavors in stored fish, respectively (Mørkøre et al., 2010). Regardless of treatments, all

283 samples showed a significant drop in IMP levels between storage day 0 and 6. Furthermore, at day 0
284 no difference was found between temperature treatments within packaging-groups, however at the
285 end of the storage period, samples treated to 40 °C were significantly lower than those at treated to
286 50, and 60 °C ($p=0.003-0.042$) (Figure 3). This can be explained by the enzymatic nature of the
287 degradation of IMP to HxR. Higher treatment temperatures yield a higher degree of enzyme
288 denaturation, causing the degradation of IMP to slow (Surette, Gill, & LeBlanc, 1988). The temperature
289 dependence is further highlighted by the fact that the levels of HxR were significantly higher in samples
290 heated to 40, followed by 50 and lastly 60 °C. For the duration of the entire storage period, only
291 samples heated to 60 °C with SGS showed any development in HxR levels (Figure 3). **The developmental**
292 **trend indicates** that the conversion rate from IMP to HxR equals that of HxR to Hx. Concerning Hx, no
293 significant initial differences were observed between temperatures within packaging groups. During
294 the storage period levels developed into samples heated to 40 °C having significantly higher levels than
295 those at 50 °C, which in turn was significantly higher than those heated to a core temperature of 60 °C
296 ($p<0.013$). As for the formation of HxR, this development is explained by lower temperatures causing
297 lesser enzyme denaturation. **Further**, the formation of Hx is partly caused by bacterial action (Surette
298 et al., 1988), and as seen from the bacterial counts, bacterial levels were significantly higher with
299 lowering of the core temperature (Table 2), thus further explaining the increased formation of Hx at
300 lower core temperature. Analysis of ATP degradation products in the raw salmon used in the two round
301 of experiments (MA or SGS) showed only HxR levels to be significantly different ($p=0.037$) (data not
302 shown). This could explain the differences observed between packaging groups in HxR- as well as in Hx
303 levels, however, it does not explain why significant differences were observed between packaging
304 technologies for IMP. Due to the differences in initial levels of ATP degradation products observed
305 between the two batches of raw material, it is not possible to distinguish potential effect of variation
306 in packaging technology from the batch variations, rendering comparisons between packaging
307 technologies infeasible.

308 3.2 Microbial community and processing

309 The raw material for the two rounds of experiments was obtained separately in order to obtain equal
310 length from slaughtering to processing and analyzing, and thereby equal rigor-state. Unfortunately,
311 the raw material characteristics were significantly different for the two batches, both regarding protein
312 content ($p=0.003$), lipid content ($p=0.002$), and water content ($p=0.002$). It has previously been shown
313 (Abel et al., 2018) that the important parameter for absorption of CO_2 and consequently the
314 bacteriostatic effect of packaging, is the total content of water and liquid lipid. When combining water-
315 and lipid content from each of the two batches of salmon, no significant differences were seen
316 between them ($p>0.642$) rendering the differences in each compound not important. All other inputs
317 were identical between the two rounds (water bath temperatures $p>0.701$, packaging gas CO_2
318 concentration $p=0.551$, and storage temperature $p>0.921$) and thus the two round is assumed equal.

319 Regarding microbial growth, a clear pattern evolved during the study when examining both the natural
320 flora and the *L. innocua*-inoculated samples. Regardless of packaging technology, no growth was
321 observed on any samples heated to a core temperature of 60°C , the same was the case for SGS treated
322 samples heated to core temperature of 50°C . Temperatures were chosen based on pre-trials, which
323 had shown growth after heat treatment at 40 , 50 , and 60°C (data not shown). The pre-experiments
324 were carried out without the use of modified atmosphere, which might explain the inconsistencies.
325 Furthermore, regardless of packaging technology higher temperature treatment always lead to a lower
326 level of recovered bacteria, as expected (Figure 4 and Figure 5). Headspace gas composition analysis
327 showed that SGS treated samples had significantly higher O_2 levels ($p<0.001$) after equilibrium has
328 been reached (day 6 and onwards). It has previously been shown the presence of CO_2 during the
329 heating step facilitates a higher heat inactivation of the bacteria (Abel et al., 2019; Loss & Hotchkiss,
330 2002), however, this was not observed in the present study (Figure 4 and 5). In the case of *L. innocua*-
331 inoculated samples, heat treatment to a core temperature of 40°C led to an insignificant reduction in
332 CFU ($p>0.072$) unlike that seen for the 50°C samples ($p<0.001$), regardless of presence of CO_2 . This led
333 to no differences in bacterial count at day 0 between samples packed using MA or SGS when heated

334 to 40 °C core temperature. On the other hand, samples heated to a 50 °C core temperature showed
335 significantly higher initial CFU levels ($p < 0.001$) when treated with CO₂ prior to heating (SGS samples).
336 This can be explained by a visible bloating that arose during the heat treatment. The solubility of CO₂
337 is highly temperature dependent (Sivertsvik, Jeksrud, Vågane, & Rosnes, 2004), thus heating at
338 temperatures as high as 55-65 °C for prolonged periods will decrease the solubility of CO₂, causing it
339 to desorb from the salmon and into the headspace of the vacuum pouch. This forms a layer of gas
340 surrounding the sample, **protecting the samples from the heat of the water bath, thus reducing the**
341 **heat load less than anticipated.** Despite the higher initial count from the SGS-treated samples, the
342 bacterial counts are equal already at day 6 and surpassed by the MA packaged samples from day 10
343 and onwards ($p = 0.001-0.035$). This effect is ascribed to the fact that samples treated in 55 °C water
344 and packed using SGS showed no significant bacterial growth throughout the entire 24 days of storage
345 **(Figure 4 and Table 3).** The experiments were performed in two rounds, and no significant differences
346 were observed between bacterial counts in either inoculum or from samples right after inoculations.
347 This means that the outgrowing on the MA packaged samples can be ascribed to the processing and
348 not variations between samples. It has been suggested by multiple studies that increased CO₂ level will
349 increase the inhibitory effect of *Listeria spp.*, e.g. by reducing the growth rate (Augustin & Carlier, 2000;
350 Devlieghere et al., 2001; Farber, Cai, & Ross, 1996; Provincial et al., 2013), **as seen in the present study**
351 **(Table 3).** However, to the best of our knowledge, this is the first time it has been proven that growth
352 of *L. innocua* can be completely inactivated for as long as 24 days of storage, **under the given**
353 **conditions.** Industry practice calls for a heat-treatment at no less than 70 °C for 2 min (concerning
354 *Listeria spp.*; Advisory Committee on the Microbiological Safety of Food (2009)) in order to ensure a
355 safe product, often with unwanted quality deterioration as a result. The present results show that
356 **inhibitory effect on *Listeria spp.* can be reached at** much lower processing temperatures, at least for
357 a refrigerated storage period of up to 24 days. SGS treatment of samples does not only benefit the
358 high-temperature samples. Even when treated to 40 °C, introduction of SGS results in a prolonging of
359 the lag phase of *L. innocua* **(Table 3)** from seeing significant count increase already between day 0 and

360 6, to a lag phase of more than 6 days. This leads to no significant difference at the end of storage
361 between 50 °C treated samples packed using MA packaging and 40 °C treated samples packed using
362 SGS (Figure 4). This highlights that SGS compared to MA packaging makes up for at least a 10 °C
363 difference in core temperatures, when it comes to inhibition of *Listeria*, with potential organoleptic
364 quality improvements as a result. The experiments were only performed under ideal storage
365 conditions e.g. low storage temperature, elimination of cross contamination etc., thus the effect might
366 be different in case of temperature abuse or deviating conditions. However, the comparison between
367 MA packaging and SGS is believed to be true as tests are performed under the same condition.

368 Similar positive results were seen for the natural samples. Unlike *L. innocua* inoculated samples, the
369 bloating of the natural samples during heat treatment was not enough to influence the bacterial
370 inhibition by heat (figure 5). A log reduction in CFU of 0.82, 1.17, 1.14 or 3.14 (MA-40°C, MA-50°C, SGS-
371 40°C or SGS-50°C, respectively) was obtained ($p < 0.003$ for all groups). The difference in heat
372 inactivation can be ascribed to the lower temperature tolerance of natural flora that mainly consists
373 of LAB, as can be seen by the lack of significant differences between bacterial count for natural samples
374 grown on either non-selective or selective LAB growth media (equivalent to Figure 5, with bacterial
375 counts at day 24 for 40 °C core temperature samples: LI $5.8-5.98 \pm 0.35$ log CFU \times g⁻¹ while MRS $5.33-$
376 5.59 ± 0.20 log CFU \times g⁻¹). This further explains why significant bacterial growth was only observed for
377 samples heated to 40 °C, with the exception of growth on samples heated to 50 °C packed in MA.
378 Comparing bacterial growth on samples heated to 40 °C packed using either MA or SGS showed no
379 significant difference in bacterial count immediately after heat treatment. This was in spite of the raw
380 material used for the SGS-treated samples showing a significantly higher bacterial count prior to heat
381 treatment ($p < 0.001$) (Figure 5). At the end of the storage period, no significant differences were
382 observed between bacterial counts from samples packed in MA and SGS ($p > 0.545$), however, at days
383 6 till 17 MA packaged samples had significantly higher bacterial counts compared to SGS ($p < 0.029$)
384 (Figure 5). Manufacturers are not only interested in obtaining the longest possible period below the
385 recommended maximum level of 10^6 CFU \times g⁻¹ (Health Protection Agency, 2009), but they are equally,

386 if not more, interested in prolonging the period with what is perceived as “good quality”, which is what
387 SGS-treatment facilitates **under the conditions** of the present study (Figure 4 and Figure 5). Spoilage
388 of MA packaged seafood has often been associated with the growth of LAB which, due to its
389 proteolytic abilities, can cause serious deterioration of the quality of the products, including increased
390 drip loss and loosening of the texture (Gram & Huss, 1996). Reducing the growth of LAB therefor has
391 the potential to significantly increase the quality of the product. The use of SGS packaging significantly
392 lowered the maximum growth rate by half and increased the lag phase of the natural flora compared
393 to that observed in MA packed samples (Table 3). **The results indicate a potential inhibitory effect on**
394 **the natural flora of salmon, as** introduction of SGS **in the current study** compensates for a 10 °C
395 reduction in processing temperature, as seen for the listeria inoculated samples. **Further certainty**
396 **regarding the effect on the natural flora, can only be achieved through further challenge testing,**
397 **considering both strains and conditions.**

398 **The non-pathogenic *L. innocua* has a high phenotypic similarity to *L. monocytogenes* and is often**
399 **used as surrogate for *L. monocytogenes*. Both species share ecological niches (Hudecova, Buchtova,**
400 **& Steinhauserova, 2010), show no differences in growth patterns (McLaughlin, Casey, Cotter, Gahan,**
401 **& Hill, 2011), and no differences in response to the use of MA packaging (Hugas, Pagés, Garriga, &**
402 **Monfort, 1998). On the other hand, *L. innocua* have been found to be more heat resistant than *L.***
403 ***monocytogenes* under certain conditions (Lorentzen, Ytterstad, Olsen, & Skjerdal, 2010). This makes**
404 ***L. innocua* a suitable, yet more conservative, surrogate for studying effect of processing on *L.***
405 ***monocytogenes* under the conditions in present experiment (Hu & Gurtler, 017).**

406 4 Conclusion

407 In conclusion, SGS has long been expected to have beneficial properties with regard to prolonging shelf
408 life, however, the impact on chemical quality has been questioned. The present study underlines the
409 microbiological benefits of CO₂ in SGS by prolonging the lag phase of both *Listeria innocua* and the
410 naturally occurring flora, slowing the growth rate, and even completely hindering the growth of *L.*

411 *innocua* for 24 days of storage as compared to MA packaging. Furthermore, no negative effect of SGS
412 was observed for any of the chemical parameters tested, including WHC, drip loss, surface color, and
413 texture. Consumers show an increased demand for lightly processed convenient seafood products. The
414 present experiment has shown that it is possible to lower processing temperatures to a little as 40 or
415 50 °C and still obtain an inhibitory effect on *Listeria spp.*, one of the biggest risks regarding food safety,
416 while improving chemical quality compared to traditional processing.

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421 6 Author Contribution

422 The study was designed in cooperation between N. Abel, B.T. Rotabakk, and J. Lerfall. Data was
423 collected by N. Abel, V. B. Ahlsen, and J. Lerfall. Results were interpreted by N. Abel, J. Lerfall, B.T.
424 Rotabakk, and T. Rustad. The manuscript was drafted by N. Abel and revised by B.T. Rotabakk, T.
425 Rustad, V. B. Ahlsen and J. Lerfall.

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427 7 References

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614 Table 1: Experimental design and response variables.
 615 MA=modified atmosphere, SGS=soluble gas stabilization

Design variables	Levels
Core temperature	40, 50, or 60 °C
Packaging method	MA packaging or SGS
Microbial community	Natural or inoculated with <i>Listeria innocua</i>
Storage time	0, 6, 10, 13, 17, or 24 days
Response Variables	Analyses
Quality (only on natural samples)	Color, composition, drip loss, headspace gas composition, metabolites of ATP, texture, water-holding capacity.
Microbiological	Aerobic plate count, H ₂ S-reducing clostridium, Lactic acid bacteria, <i>Listeria spp.</i>
Tested samples	Raw, processed, stored n=4 for each group, at each sampling point

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622 *Table 2: Main effect of treatment temperature and packaging technology on drip loss, WHC, headspace CO₂ concentration,*
 623 *and bacterial count of natural and inoculated samples. Key results are elaborated in Figure 4 and Figure 5*

		Drip loss [%] df=175	WHC [%] df=54	Head space CO₂ [%] df=257	Log CFU x g⁻¹ Inoculated df=141	Log CFU x g⁻¹ Natural df=139
Core temp	40	7.4±2.2 ^a	70.4±5.4 ^a	50.6±7.4 ^a	3.85±0.77 ^a	3.97±1.4 ^a
	50	8.9±1.3 ^a	71.2±4.1 ^a	49.8±6.3 ^a	2.18±1.05 ^b	0.90±1.2 ^b
	60	7.4±1.4 ^a	71.0±5.1 ^a	49.7±6.7 ^a	N.D.	N.D.
		p=0.065	p=0.868	p=0.635	p<0.01	p<0.01
Packaging technology	SGS	8.0±1.9 ^a	72.0±5.6 ^a	54.5±5.8 ^a	2.26±2.01 ^a	1.95±2.05 ^b
	MA	7.8±1.9 ^a	69.7±3.7 ^a	45.4±4.2 ^b	1.69±1.42 ^a	1.17±1.87 ^a
		p=0.329	p=0.087	p<0.01	p=0.053	p=0.020

df = degrees of freedom

N.D.=not detected.

Superscript letters (a-c) indicates significantly mean value differences (p<0.05) according to one-way ANOVA

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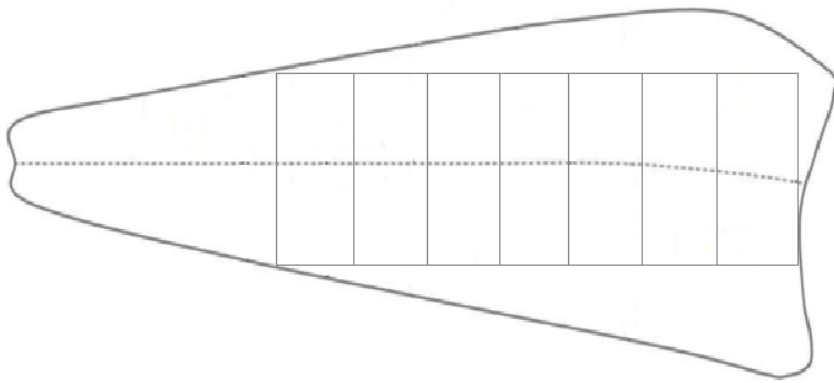
626 Table 3: Growth kinetic parameters (maximum specific growth rate (μ_{max} , day^{-1}) and lag phase (day) for samples subjected to
 627 different heat treatments (40 or 50 °C) and packaging technology (MA and SGS). Samples treated at 60 °C are not included
 628 due to lack of detectable growth. The parameters are estimated from the primary model of Baranyi and Roberts (1994) using
 629 log-transformed bacterial counts.

Treatment	μ_{max} [day^{-1}]	Lag phase [day]	R^2	SE(fit)
<i>Natural flora</i>				
40°C – MA	0.48±0.08	2.8±1.4	0.929	0.515
50°C – MA	0.2±0.05	3.1±2.0	0.827	0.827
40°C – SGS	0.22±0.03	4.0±0.2	0.934	0.336
50°C – SGS	N/A ^a	N/A ^a	N/A ^a	N/A ^a
<i>Inoculated flora</i>				
40°C – MA	0.70±0.06	-	0.951	0.421
50°C – MA	0.38±0.15	1.59±2.4	0.839	0.563
40°C – SGS	0.06±0.03	6.48±0.1	0.643	0.233
50°C – SGS	0.10±0.88	24.0±5.1	-0.0745	0.426

630 R^2 , coefficient of determination, SE (fit), standard error of fit to the model, ^a not estimated due to no detectable growth during storage.

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634 *Figure 1: Schematic illustration showing the sampling of salmon portion after removal of backfin, belly flap, and tail.*

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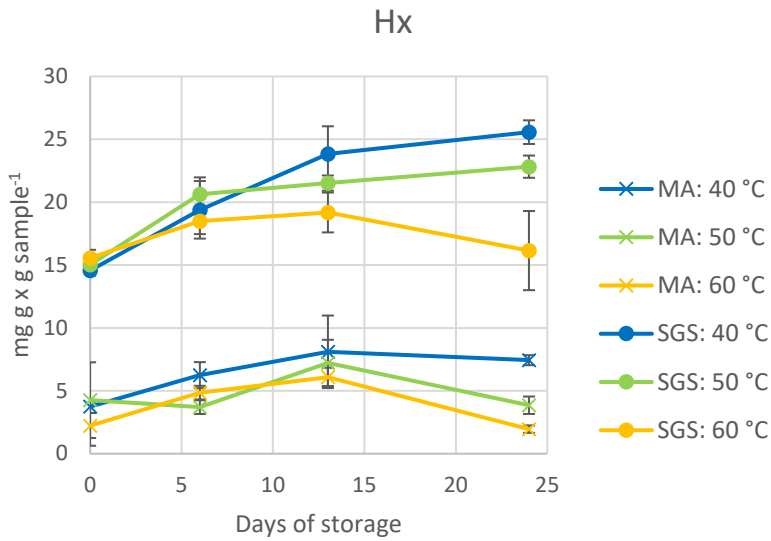
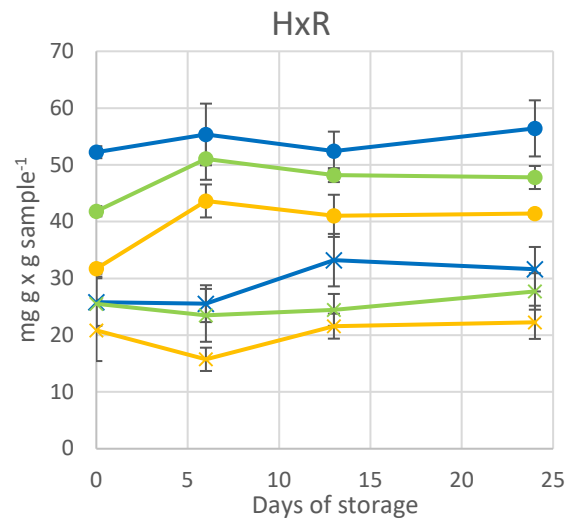
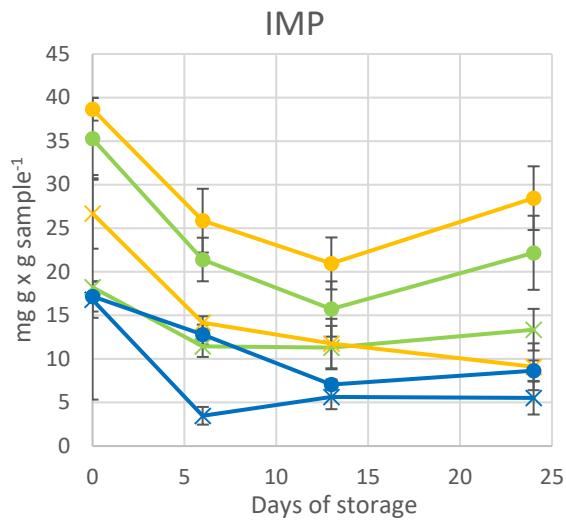
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637 *Figure 2: Image of salmon samples immediate after heat treatment in temperatures of 45 (left), 55 (middle), or 65 °C (right).*

638 *All samples are with no CO₂ exposure prior to heat treatment (MA-samples).*

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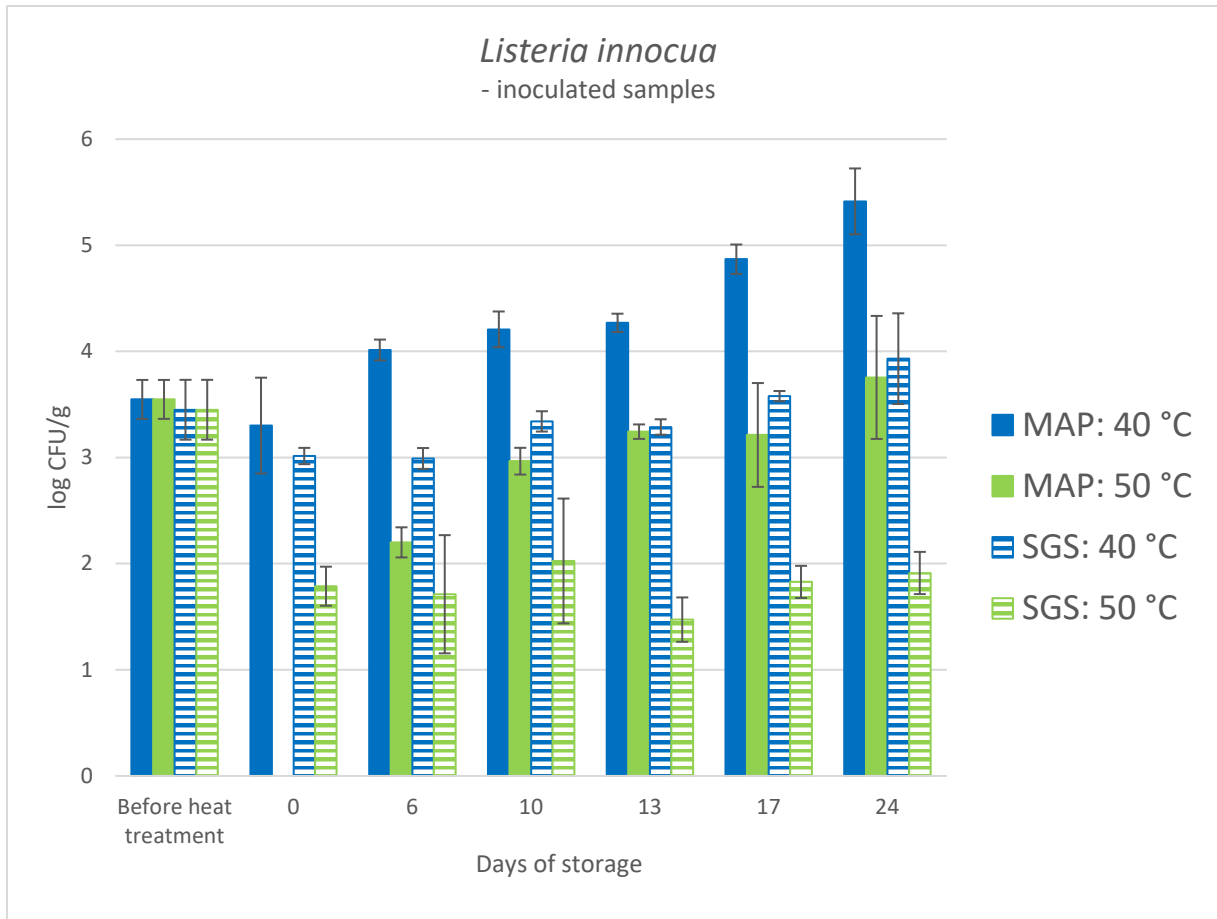


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643 *Figure 3: Development of ATP-degradation products inosine monophosphate (IMP) (upper left), inosine (HxR) (upper right)*
 644 *and hypoxanthine (Hx) (lower left) during 24 days of storage, separated based on treatment temperatures and packaging*
 645 *technology applied. X = MA packaged samples, O = SGS treated samples, blue = 40 °C core temperature samples, green = 50*
 646 *°C core temperature samples, yellow = 60 °C core temperature samples. Error bars indicates mean ±1 standard deviation.*

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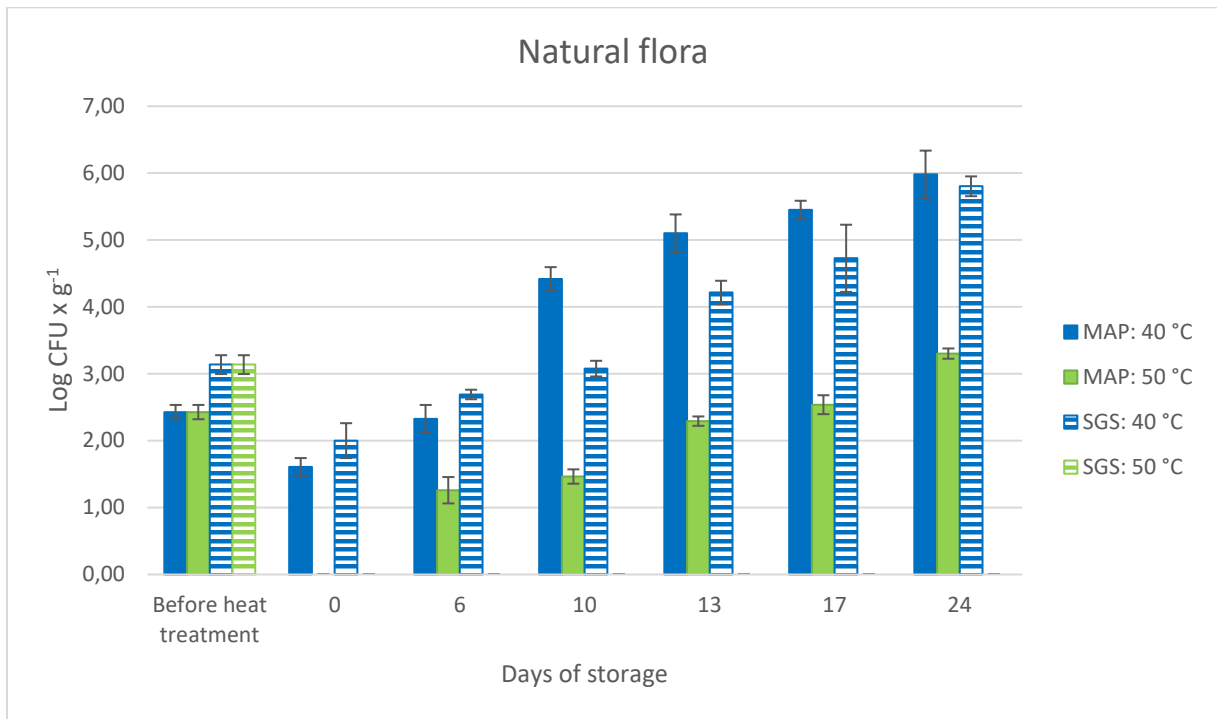
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649 *Figure 4: Growth of Listeria innocua on inoculated samples, separated based on heat treatment temperature and packaging*
 650 *technology applied. 60 °C samples, regardless of packaging technology, showed no growth at any point, and thus have been*
 651 *left out. Solid = MA packaged samples, striped = SGS treated samples, blue = 40 °C core temperature samples, green = 50 °C*
 652 *core temperature samples. Error bars indicates mean ±1 standard deviation.*

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658 *Figure 5: Growth of microbial flora on natural samples, separated based on heat treatment temperature and packaging*
659 *technology applied. 60 °C samples, regardless of packaging technology, showed no growth at any point, and thus have been*
660 *left out. Solid = MA packaged samples, striped = SGS treated samples, blue = 40 °C core temperature samples, green = 50 °C*
661 *core temperature samples. Error bars indicates ±1 standard deviation.*

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