

1 **Se improved indole glucosinolate hydrolysis products content, Se-**
2 **methylselenocysteine content, antioxidant capacity and potential**
3 **anti-inflammatory properties of sauerkraut.**

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20 ABSTRACT

21 Selenium (Se) has a well-known role in prevention of chronic diseases
22 associated with oxidative stress and inflammation. The objective of this
23 study was the production of Se-enriched sauerkraut and to know the
24 effect of selenite addition on indole glucosinolate (GLS) hydrolysis
25 products, vitamin C, Se biotransformation and microbial quality as well as
26 the antioxidant and anti-inflammatory properties of sauerkraut. White
27 cabbage was naturally fermented with 0.3 mg Na₂SeO₃/Kg fresh
28 cabbage (NFSe) or without Se addition (NF). Chemical analyses were
29 performed by LC-PAD, CE and LC-ICP-MS. Antioxidant capacity was
30 determined using the oxygen radical antioxidant capacity (ORAC)
31 method. Anti-inflammatory activity was measured as inhibition of nitric
32 oxide (NO) production in LPS-induced macrophages. Total Se content
33 reached up to 1.29 µg/g dry matter (0.11 µg/g fresh weight) and
34 selenomethylselenocysteine was the major Se form found in NFSe
35 cabbage. Se addition caused a slight reduction of ascorbigeno (6%) and
36 vitamin C (5%) content in sauerkraut ($P \leq 0.05$); however, LAB increased
37 (3%), and the formation of indole-3-carbinol (74%) and indole-3-
38 acetonitrile (13%) were markedly enhanced ($P \leq 0.05$). NFSe cabbage
39 extracts showed higher ($P \leq 0.05$) antioxidant activity (163 µmol Trolox/g
40 d.m.) and anti-inflammatory potency ($IC_{50} = 44.01 \mu\text{g/mL}$) compared to
41 NF cabbage extracts. Consequently, Se-enriched sauerkraut can be
42 considered as health-promoting food.

43 **KEYWORDS:** Fermented cabbage, selenium, glucosinolate breakdown
44 products, antioxidant activity, anti-inflammatory activity

45 INTRODUCTION

46 Selenium (Se), as component of selenoamino acids and
47 selenoproteins, is essential in important physiological functions such as
48 redox homeostasis,¹ thyroid hormone metabolism and stimulation of the
49 immune system to increase antibody production.² Recently, Se has
50 attracted tremendous interest because of intensive investigation showing
51 the potential of Se to protect against oxidative stress and chronic
52 inflammation, conditions commonly associated to several chronic
53 diseases.³ Indeed, several human studies suggested that optimal Se
54 status could prevent cancer, cardiovascular disease and type 2
55 diabetes.^{4,5}

56 Recommended Se dietary intake is not well-standardized among
57 different countries. The recommended dietary allowance (RDA) for Se in
58 USA is 55 $\mu\text{g}/\text{day}$ for men and women⁶ while the WHO recommends a
59 Se intake of 40 and 30 $\mu\text{g}/\text{day}$ for men and women, respectively.⁷ Se
60 intake especially in South and Eastern European countries is below
61 RDA⁸ and consumption of Se supplements has been the most
62 widespread approach to prevent Se deficiency. Consumption of Se-
63 enriched foods is increasing markedly in recent years as an alternative to
64 increase Se status and promote health of Se-deficient populations.⁹
65 Several authors have reported preparation of Se-enriched foods by
66 means of fermentation processes.¹⁰ Yeasts and lactic acid bacteria have
67 shown its ability to accumulate and transform inorganic Se to organoSe
68 compounds.^{10,11} For instance, selenomethionine was found to be the
69 main Se-specie in selenised white wine produced by *Saccharomyces*

70 *cerevisiae* and *Saccharomyces bayanus*.¹² Therefore, Se-enriched
71 fermented foods could become a dietary source of bioavailable and
72 physiologically relevant Se forms.

73 Sauerkraut, a popular product resulting from the lactic acid
74 fermentation of white cabbage, is a valuable vegetable plant-food due to
75 its nutritional and health-promoting properties, mainly attributed to its
76 high content of antioxidant compounds such as vitamin C and phenolic
77 compounds^{13,14} which prevent cell damage caused by free radicals.
78 Besides that, sauerkraut presents high levels of glucosinolate (GLS)
79 breakdown products¹⁵ among them, indole-3-carbinol (I3C), indole-3-
80 acetonitrile (I3ACN) and ascorbigen (ABG) are the most abundant.^{14,15,16}
81 The GLS-breakdown products have been linked with a reduction of
82 cancer risk by inhibiting phase I enzymes, involved in carcinogen
83 activation and inducing phase II enzymes, involved in the detoxification
84 of xenobiotics.^{17,18} These compounds also inhibit tumor cell growth and
85 stimulate apoptosis.¹⁹ Different authors described the relationship
86 between inflammation and cancer.²⁰

87 There is no literature information on the effect of the Se addition
88 during cabbage fermentation on the bioactive compounds and biological
89 properties of sauerkraut. Therefore, the objective of this study was the
90 production of Se-enriched sauerkraut and to know the effect of selenite
91 addition on indole GLS hydrolysis products, vitamin C, Se
92 biotransformation and microbial quality as well as the antioxidant
93 capacity and anti-inflammatory properties of sauerkraut.

94

95 **MATERIAL AND METHODS**

96 **Plant material.** White cabbages (*Brassica oleracea* L. var.
97 *capitata* cv. Megaton) grown in the North region of Spain (La Rioja)
98 during winter season 2008 were selected among five Spanish cultivars,
99 based on their glucobrassicin content.²¹ Fresh cabbages were provided
100 by Bejo Iberica S. L. (Madrid, Spain) and fermented immediately upon
101 reception.

102 **Preparation of Se-enriched sauerkraut.** Cabbage heads were
103 prepared by removing the outer leaves and their central core. The edible
104 part of cabbages was then shredded into about 2 mm thick strips using
105 a shredder (Moka Express, Barcelona, Spain). Subsequently, 0.5%
106 NaCl and 0.3 mg of sodium selenite/Kg of fresh cabbage (1.6 mg Se/Kg
107 of dry matter) were added to achieve Se RDA and a high selenite
108 biotransformation in selenoaminoacids. Shredded cabbage and brine
109 were mixed thoroughly, transferred to sterile polyethylene vessels (8 L)
110 and tightly pressed to exclude air. Fermentations were performed
111 spontaneously by the indigenous microbiota present on raw cabbage
112 (NFSe). Sauerkraut without addition of Se were also prepared and
113 considered as control (NF). Fermentations were carried out in 3 batches
114 (4 Kg per batch) at room temperature (22-25 °C) for 7 days. On the third
115 day, cabbage was pricked to remove releasing gases. Raw, NF and
116 NFSe cabbages were freeze-dried, milled and stored at -20 °C under
117 vacuum until their analysis.

118 **Determination of pH during fermentation.** Brine from each
119 fermentation batch (2 mL) was collected at 0, 3 and 7 days of

120 fermentation and their pH was measured in a pH meter Basic 20 (Crison,
121 Barcelona, Spain).

122 **Analysis of indole GLS-derived compounds.** The content of
123 ABG, I3C and I3ACN was determined by HPLC-PAD in raw and
124 fermented cabbage as in Peñas et al.¹⁴

125 **Determination of vitamin C.** The quantification of vitamin C
126 content in raw and fermented cabbages was performed by capillary
127 electrophoresis (CE) as in Frias et al.²² CE analysis were performed in a
128 P/ACE system 2050 (Beckman Instruments, Fullerton, CA, USA) and UV
129 detector at 254 nm. Separation was done using a 47 cm x 75 μm i.d.
130 fused silica capillary TSP075375 column (Composite Metal Services
131 LTD, Worcester, UK) at room temperature.

132 **Oxygen Radical Antioxidant Capacity (ORAC).** ORAC was
133 determined in aqueous extracts from raw and fermented cabbage
134 extracts. Extracts were obtained by suspension of 1 g of freeze-dried
135 sample in 10 mL of phosphate buffer (PBS, pH 7.4) and stirring for 1 h at
136 room temperature. Homogenates were filtered using Whatman No.1 filter
137 paper. The ORAC assay was determined as described by Dávalos et
138 al.²³ Results were expressed as μmol Trolox equivalents (TE) per gram
139 of dry matter ($\mu\text{mol TE/g d.m.}$).

140 **Determination of Se-derived compounds.** Total selenium
141 concentration was determined in raw and fermented cabbages by ICP-
142 MS (Thermo-X Series) after acid digestion in an analytical microwave
143 oven (CEM MSP 1000, Matheus, NC) by following conditions described
144 previously.^{10,11} Selenium speciation was carried out by HPLC-ICP-MS

145 previous enzymatic probe sonication.^{10,11} Enzymatic hydrolysis was
146 performed by 2 minutes of sonication (Sonoplus ultrasonic homogenizer
147 Bandenlin, Germany) after addition of 20 mg of Protease XIV (Sigma-
148 Aldrich, Steinheim, Germany) and 3 mL of Milli-Q water to 100 mg of
149 dried sample. The obtained extracts were centrifuged at at 15,557 $\times g$ for
150 30 min (4 °C) using 10 KDa cut-off filters (Millipore, USA). The ICPMS
151 instrument was coupled with a Hamilton PRP-X100 (250 \times 4.1 mm, 10
152 μ m) chromatographic column. The mobile phase was 10 mM ammonium
153 citrate, pH 5.0, in 2% methanol as mobile phase at a flow rate of
154 1 mL/min. Identification and quantification of selenium species was done
155 by retention time and spiking experiments. Standard stock solutions of
156 1000 mg/L of selenomethionine (SeMet), selenomethylselenocysteine
157 (SeMeSeCys) and selenocystine (SeCys₂) (Sigma) were prepared in
158 ultra-pure Milli-Q water (Millipore, MA, USA), and 3% hydrochloric acid
159 was added for better dissolution of SeCys₂ and SeMeSeCys. Inorganic
160 selenium solutions were prepared by dissolving sodium selenite
161 (Na₂SeO₃) and selenate (Na₂SeO₄) (Merck) in Milli-Q water.

162 **Preparation of extracts for cell treatment.** Extracts were
163 prepared by homogenization of 500 mg freeze-dried sample in 20 mL
164 acetone:water solution (1/1) using an Ultra Turrax homogenizer T-25
165 Digital (Ika Werke GMBH & Co., Staufen, Germany), and centrifugation
166 for 7 min at 3,024 $\times g$ and 5 °C. Supernatant was collected, and the pellet
167 was extracted twice with 10 mL of acetone. Further, supernatants were
168 combined, filtered using Whatman No. 1 paper and concentrated to 7 mL
169 final volume. The concentrate was extracted twice with 15 mL of ethyl

170 acetate. The combined organic layers were dried over anhydrous sodium
171 sulfate, filtered, and evaporated under vacuum to dryness. Finally, the
172 residue was dissolved in 0.1% dimethylsulfoxide (DMSO) (Sigma).

173 **Cell viability assay.** Macrophages cell line RAW 264.7 (ATCC,
174 Manassas, VA, USA) were cultured in Dubelcco's modified Eagle
175 Medium (DMEM; from ATCC) containing 1% penicillin/streptomycin
176 (Sigma), and 10% fetal bovine serum (ATCC) at 37 °C in 5% CO₂
177 atmosphere. The cell proliferation assay was conducted using the
178 CellTiter 96 Aqueous One Solution Proliferation assay kit using the
179 novel tetrazolium compound, 3-(4,5-dimethylthiazol-2-yl)-5-(3-
180 carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium, inner salt
181 (MTS), and an electron coupling reagent, phenazine ethosulfate (PES)
182 (Promega Biotech Iberica, Madrid, Spain). Briefly, 5 x 10⁴ cells/well were
183 seeded in a 96-well plate and the total volume was adjusted to 200 μL
184 with DMEM. The cells were allowed to grow for 24 h at 37 °C in 5% CO₂.
185 After 24 h incubation, they were treated with different concentrations of
186 raw and fermented cabbage extracts (0-150 μg/mL), ABG (0-1000 μM)
187 and SeMeSeCys (0-10 μM) for 24 h. After treatment, DMEM was
188 replaced by 100 μL fresh medium and 20 μL MTS/PES was added to
189 each well. The plate was incubated for 2 h at 37 °C and the absorbance
190 read at 490 nm. The percentage of viable cells was calculated with
191 respect to cells treated with vehicle (0.1% DMSO) as follows: $A_{\text{treatment } 490 \text{ nm}}/A_{\text{control } 490 \text{ nm}} * 100 = \% \text{ cell viability}$
192

193 **Measurement of nitric oxide (NO).** Approximately 5 x 10⁴
194 cells/well were seeded in a 96-well plate and allowed to grow to its 80-

195 90% confluency. The cells were treated with 1 $\mu\text{g}/\text{mL}$ lipopolysaccharide
196 (LPS) from *Escherichia coli* O55:B5 (Sigma) with or without different
197 concentrations of raw and fermented cabbage extracts (0-150 $\mu\text{g}/\text{mL}$),
198 ABG (0-1000 μM) and SeMeSeCys (0-10 μM) for 24 h. After this
199 treatment, medium was collected and NO production analyzed. Nitrite
200 accumulation, and indicator of NO synthesis, was measured in the
201 culture medium by Griess reaction.²⁴ Briefly, 100 μL of DMEM were
202 plated in 96-well plate and an equal amount of Griess reagent constituted
203 by 1% (w/v) sulfanilamide and 0.1% (w/v) N-1-
204 (naphthyl)ethylenediamine-diHCl in 2.5% (v/v) H_3PO_4 , was added. The
205 plate was incubated for 5 min and the absorbance measured at 550 nm
206 in a microplate reader (Biotek, Winooski, VT, USA). The amount of NO
207 was calculated using a sodium nitrite standard curve. Potency was
208 determined by dose–response curves in which the range of
209 concentrations was distributed in a logarithmic scale and the IC_{50} values
210 were calculated using non-linear regression sigmoidal curve fit functions
211 in GraphPad Prism 4.00 (Graphpad Software Inc., San Diego, CA, USA).

212 **Microbiological analysis.** Microbiological analyses were
213 performed on raw and fermented cabbage as described by Peñas et
214 al.²⁵ Briefly, five grams of each sample were aseptically diluted in
215 buffered peptone water (Scharlau Chemie, Spain) in a sterile Stomacher
216 bag and homogenised for 1 min in a Stomacher laboratory blender (IUL
217 Masticator, Barcelona, Spain). Further serial dilutions were made for
218 plating. The pour plate technique was employed to determine the
219 microbial counts. Total aerobic and anaerobic mesophilic bacteria were

220 enumerated on Tryptone Soya Agar (TSA) after incubation in aerobic
221 and anaerobic conditions, respectively, at 30 °C for 72 h; total and faecal
222 coliforms on Violet Red Bile Agar (VRBA) containing lactose as
223 carbohydrate source, after incubation at 37 °C and 44 °C, respectively,
224 for 24 h; moulds and yeasts on Sabouraud-Chloramphenicol Agar, after
225 incubation at 23 °C for 96 h; and LAB on MRS Agar after incubation in
226 anaerobic conditions at 30 °C for 24-48 h.

227 **Statistical analysis.** Data were expressed as means of three
228 experiments. Results were compared by one-way analysis of variance
229 (ANOVA) using the least significant differences ($P \leq 0.05$) (Statgraphic
230 5.0 software, Statistical Graphics Corporation, Rockville, MD, USA).

231

232 **RESULTS AND DISCUSSION**

233 **Evolution of pH during cabbage fermentation.** Figure 1 shows
234 pH evolution during cabbage fermentation in the presence (NFSe) or
235 absence (NF) of selenite solution. Measurement of pH can be considered
236 as an indicator of the success of fermentation processes.²⁶ Raw cabbage
237 pH (6.2) was found similar to the one observed previously by our group in
238 white cabbage cv. Bronco.¹⁴ A rapid pH decrease took place up to 3 days
239 of fermentation as a result of organic acid production by indigenous LAB
240 in NF (pH 3.6) and NFSe (pH 3.7) cabbages, and it did not change up to
241 7 days of fermentation (Figure 1). Our results are consistent with
242 previous studies in cabbages submitted to natural or induced
243 fermentation using LAB starter cultures.^{14,27,28} Moreover, selenite addition
244 had no significant influence ($P \leq 0.05$) on pH evolution during sauerkraut

245 production (Figure 1). Therefore, Se addition seemed to have no adverse
246 effect on LAB fermentative metabolism during sauerkraut production.

247 **Effect of Se-enrichment on the content of indole GLS**
248 **breakdown products in natural fermented cabbage.** Raw cabbage
249 showed small concentrations of indole GLS hydrolysis products such as
250 ABG (~16 $\mu\text{mol}/100\text{g d.m.}$), I3C (0.10 $\mu\text{mol}/100\text{g d.m.}$) and I3ACN (0.24
251 $\mu\text{mol}/100\text{g d.m.}$) since cabbage was shredded immediately before
252 analysis (Table 1). ABG concentration found in raw cabbage was within
253 the range reported in previous studies.^{14,16} However, no literature data
254 has been found about I3C and I3ACN content in raw white cabbage.

255 During natural fermentation, the content of indole GLS hydrolysis
256 products markedly increased ($P \leq 0.05$), being ABG (218.2 $\mu\text{mol}/100\text{g}$
257 d.m.) the major indole GLS-derived compound found in NF cabbage
258 (Table 1), followed by I3C (9.6 $\mu\text{mol}/100\text{g d.m.}$), and I3ACN (2.7
259 $\mu\text{mol}/100\text{g d.m.}$). The content of indole GLS-derived products in NF
260 cabbage differed from previous studies^{14-16,29} which could be attributed to
261 differences in glucobrassicin concentration and myrosinase activity as
262 they depend on genetic and environmental factors that vary among
263 growing locations and year of cultivation.^{21,30}

264 NFSe cabbages showed significantly lower ABG content but
265 higher I3C and I3ACN concentrations ($P \leq 0.05$), compared to NF
266 cabbages (Table 1). No previous studies have been found on the effect
267 of Se on the formation of GLS hydrolysis products during cabbage
268 fermentation. Indole GLS breakdown products are considered
269 anticarcinogenic agents that are associated to lower risk of breast³¹ and

270 colon cancer.³² Furthermore, ABG exert important immunomodulating
271 actions³³ while I3C promote cell cycle arrest in endometrial³⁴ and
272 prostate tumors.³⁵

273 **Effect of Se-enrichment on ascorbic acid content of natural**
274 **fermented cabbage.** As it can be observed in Table 2, raw cabbage
275 presented a high vitamin C content (~329 mg/100 g d.m.). These results
276 are in agreement with those previously reported for different white
277 cabbage cultivars.^{14,16,21} Ascorbic acid content decreased more than
278 25% in NF cabbage compared to raw cabbage. This reduction could be
279 explained by ABG formation which results from the hydrolysis of
280 glucobrassicin by myrosinase enzyme and the further reaction with L-
281 ascorbic acid at low pH,³⁶ and the oxidation of ascorbic acid during
282 cabbage fermentation. The level of vitamin C found in the present work
283 was consistent with our previous studies.^{14,16} Fermentation in the
284 presence of Se caused a small but significant ($P \leq 0.05$) decrease of
285 vitamin C in NFSe cabbage. No information was found about the effect of
286 Se on ascorbic acid content in sauerkraut.

287 **Biotransformation of selenite during natural fermentation of**
288 **white cabbage.** Table 2 collects the content of Se compounds in raw,
289 NF and NFSe cabbages. Total Se in raw and NF cabbage was 0.04 and
290 0.07 $\mu\text{g/g}$ d.m. respectively, and Se species concentration were lower
291 than the quantification limit. Selenite addition into cabbage caused a
292 19.5-fold increase of total Se concentration up to 1.29 $\mu\text{g/g}$ d.m.
293 (equivalent to 11.62 $\mu\text{g}/100$ g f.w.) and Se-methylselenocysteine
294 (SeMeSeCys) (0.74 $\mu\text{g/g}$ d.m.) was the major Se specie found in Se-

295 enriched sauerkraut. Therefore, the consumption of a serving of 150 g
296 f.w. of Se-enriched sauerkraut would contribute to cover a relevant
297 percentage of Se RDA (43.6% for men and 58.1% for women). With
298 regard to safety, upper limit of the safe adult population mean intake of
299 selenium have been established in 400 μg Se/d.⁷ A Se-enriched
300 sauerkraut serving of 150 g f.w. contains 17.43 μg Se, levels that are in
301 the safe range of Se intake.

302 After enzymatic hydrolysis using protease XIV, samples were
303 ultrafiltrated through 10 kDa cutoff membranes. The Se recovery in
304 ultrafiltrated cabbage hydrolysates was 90 \pm 3%. The same extraction
305 procedure was applied without protease (aqueous extraction) providing
306 recovery values of 60 \pm 3%. These results suggested that most of Se in
307 samples was not bound to proteins. Figure 2 shows the chromatographic
308 profiles of selenium species standards (Figure 2A) and the raw, NF,
309 NFSe cabbage hydrolysates (Figure 2B). It is noteworthy the production
310 of SeMeSeCys during fermentation in the presence of selenite.
311 SeMeSeCys was also the main Se-compound found when samples were
312 treated by aqueous extraction. SeMeSeCys is a non-proteinogenic
313 selenoamino acid which is metabolized by lyase to methylselenol *in*
314 *vivo*.³⁷ This fact is very important because there is evidence that ability of
315 endogenous production of monomethylated selenium is a critical factor in
316 Se chemoprotection. Recently, Cuello et al.³⁸ have reported that
317 SeMeSeCys protects human hepatoma cells against oxidative stress.

318 The ability of some microorganisms to biotransform inorganic Se
319 has been widely reported in the literature, especially *Sacharomyces*

320 *cerevisiae*. Yeasts are being used as Se supplement because its ability
321 to accumulate and biotransform high concentrations of inorganic
322 selenium (3000 µg/g) mainly into selenomethionine (SeMet).^{11,12} Thus,
323 several authors have reported preparation of selenized foodstuffs
324 mediated by a fermentation process.^{10,12} Alzate et al.,¹⁰ compared the
325 different Se species that are produced when lactic fermentation in
326 presence of two different types of microorganisms, LAB (*Lactobacillus*)
327 and yeast (*Saccharomyces*) take place to produce yogurt and kefir,
328 respectively. Se organic species formed depended on the type of
329 microorganism involved in the fermentation process, being SeCys₂ and
330 SeMeSeCys the main Se species generated by *Lactobacillus* and SeMet
331 by *Saccharomyces*. These results are in agreement with those reported
332 in the present work, and others previously published supporting the
333 different behaviour of microorganisms *Lactobacillus*^{39,40} and
334 *Sacharomyces* when exposed to inorganic Se.¹¹

335 Results illustrated in Table 2 evidence that up to 50% of the total
336 Se content found in Se-enriched sauerkraut was identified as
337 SeMeSeCys. This Se specie was not found in raw cabbage which again
338 suggests that its production is derived from the action of the
339 microorganisms during the fermentation process in presence of inorganic
340 Se.

341 **Effect of Se-enrichment on the antioxidant capacity of natural**
342 **fermented cabbage.** The antioxidant activity of raw cabbage was ~75
343 µmol TE/g d.m. and natural fermentation (NF) led to a sharp increase
344 (131.5 µmol TE/g d.m.) (Table 3). This effect was also observed by other

345 authors in spontaneously fermented cabbage.⁴¹ These results can be
346 attributed to the formation of GLS-derivatives during fermentation that act
347 as direct antioxidant compounds.⁴² GLS hydrolysis products may
348 enhance antioxidant potential indirectly by the action of antioxidant
349 responsive element which induced gene expression of phase II enzymes
350 in murine models.⁴³ Other factors also contribute to the enhanced
351 antioxidant activity found in sauerkraut compared to raw cabbage.
352 According to Reyes et al.,⁴⁴ the antioxidant activity of cabbage increases
353 during wounding, effect that takes place after shredded during cabbage
354 fermentation. Additionally, LAB may also affect the antioxidant activity
355 exhibited by the fermented cabbage, as has been previously suggested
356 by Kusznierevicz et al.⁴⁵

357 Fermentation in the presence of Se significantly improved ($P \leq$
358 0.05) antioxidant activity in sauerkraut (Table 3). The highest antioxidant
359 activity found in NFSe cabbage compared to NF cabbage may be related
360 to the biotransformation of selenite in Se-organic species which are able
361 to scavenge free radicals.⁴⁶

362 **Effect of Se-enrichment on microbiological quality of**
363 **sauerkraut.** Microbial counts of raw, NF and NFSe cabbage are shown
364 in Table 4. The dominating microbial population of raw cabbage was
365 aerobic mesophilic bacteria (~5 cfu/g), followed by anaerobic bacteria (~4
366 cfu/g), LAB (~2 cfu/g) and total coliforms (~1 cfu/g). Faecal coliforms,
367 moulds and yeasts were not detected in raw cabbage. These results are
368 consistent with those recently reported for other *Brassica* vegetables⁴⁷
369 and white cabbage cv. Bronco.²⁵

370 Fermentation caused a significant ($P \leq 0.05$) increase of aerobic
371 and anaerobic mesophilic bacteria (6.8 cfu/g f.w.) as well as LAB (6.7
372 cfu/g f.w.) in NF cabbage; however, microbial counts of faecal coliforms,
373 moulds and yeasts did not change after fermentation (Table 4). LAB grew
374 in greater extent than other microbial populations, due to the favourable
375 ecological conditions (acidic pH and low oxygen concentration) that take
376 place during cabbage fermentation, enhancing their multiplication and
377 inhibiting the growth of other microorganisms such as aerobic mesophilic
378 bacteria. Slightly lower counts of aerobic mesophilic bacteria and LAB
379 were found in sauerkraut obtained by spontaneous fermentation from
380 white cabbage cv. Bronco,²⁵ probably due to differences in endogenous
381 microflora.

382 The addition of Se during cabbage fermentation caused
383 significantly ($P \leq 0.05$) lower aerobic and anaerobic mesophilic bacteria
384 counts and significantly ($P \leq 0.05$) higher LAB counts. These findings
385 suggest that Se-enrichment enhanced the growth of LAB and,
386 consequently, other microbial populations decreased most likely due to a
387 competition phenomenon. Recently, Molan et al.⁴⁸ reported that inorganic
388 forms of Se (selenate and selenite) exert a prebiotic effect as evidenced
389 by their ability to promote the growth of *Lactobacillus rhamnosus* and
390 *Bifidobacterium breve in vitro*. Furthermore, these authors found that Se-
391 enriched green tea enhanced lactobacilli growth compared to
392 conventional green tea.

393 **Effect of raw and fermented cabbage extracts on**
394 **inflammatory response of LPS-induced RAW 264.7 macrophages.**

395 The cytotoxicity of raw and NF and NFSe cabbage extracts in LPS-
396 induced macrophages was evaluated at a range 0-150 μg extract/mL
397 using MTS reduction assay after 24 h incubation. Macrophages exhibited
398 a survival percentage $> 87\%$ when treated with raw and fermented
399 cabbage extracts at concentrations $\leq 150 \mu\text{g/mL}$. Therefore, these results
400 indicated that range of concentrations used in this study to treat the cells
401 did not exert any cytotoxic effect. In activated macrophages, NO
402 production noticeably increased in response to LPS as shown in Figure
403 3. Excessive production of NO in macrophages represent a potentially
404 toxic effect, which if not counteracted, causes the onset or/and
405 progression of many disease pathologies.⁴⁹ Therefore, the effect of raw,
406 NF and NFSe cabbage extracts on inhibition of NO production was
407 examined in order to study their potential anti-inflammatory effect. Raw,
408 NF and NFSe cabbage extracts induced a significant ($P \leq 0.05$) dose-
409 dependent suppression of NO production (Figure 3A). Table 3 shows the
410 calculated concentrations of raw, NF and NFSe cabbage extracts that
411 resulted in 50% inhibition of NO production in LPS-induced macrophages
412 (IC_{50}). NF cabbage extract exhibited higher ($P \leq 0.05$) potency ($\text{IC}_{50} =$
413 $83.96 \mu\text{g/mL}$) compared to raw cabbage extract ($\text{IC}_{50} = 167.93 \mu\text{g/mL}$).
414 This effect could be due to formation of bioactive compounds such as
415 GLS breakdown products exhibiting anti-inflammatory properties during
416 sauerkraut manufacture. ABG was the major GLS hydrolysis compound
417 in NF cabbage (Table 1), therefore, we further investigated the potential
418 anti-inflammatory activity of synthetic ABG in LPS-induced macrophages.
419 The cytotoxicity of ABG was firstly examined after 24 h incubation.

420 Macrophages exhibited a viability > 92% which indicated that ABG did
421 not induce any cytotoxic effect at concentrations $\leq 1000 \mu\text{M}$. ABG was
422 able to significantly ($P \leq 0.05$) reduce NO production in a dose-
423 dependent manner (Figure 3B); however, ABG was a weak inhibitor of
424 NO production in LPS-activated macrophages ($\text{IC}_{50} = 970.54 \mu\text{M}$). These
425 results suggest that ABG have a small contribution (<10%) to the NO
426 production inhibitory potency indicating that other bioactive compounds
427 would be responsible for the observed potential anti-inflammatory activity
428 of fermented cabbage extracts. A small body of literature suggests that
429 I3C and sulforaphane (SF) may protect against inflammation, inhibiting
430 cytokine production (TNF- α , IL-1, IL-6, IL-8) and expression of
431 proinflammatory enzymes such inducible nitric oxide synthase
432 (iNOS).^{50,51} Therefore, it will be interesting to see if ABG, SF and I3C
433 have synergistic effects in fighting inflammation.

434 Selenite addition (1.6 mg Se/Kg d.m.) markedly improved the NO
435 production inhibitory potency of fermented cabbage ($\text{IC}_{50} = 44 \mu\text{g}$
436 extract/mL). To further confirm the role of Se in the modulation of
437 inflammatory response we tested the effect of SeMeSeCys, the major Se
438 specie found in NFSe cabbage, on NO production in LPS-activated
439 macrophages. Treatment with SeMeSeCys (concentration range 0-10
440 μM) caused a dose-dependent inhibition of NO production in LPS-
441 activated macrophages (Figure 3C) and showed no cytotoxicity at
442 concentrations $\leq 10 \mu\text{M}$. SeMeSeCys potency to inhibit 50% NO
443 production in LPS-activated macrophages (IC_{50}) was $25.20 \mu\text{M}$
444 (equivalent to $10.93 \mu\text{M}$ of Se). This level of Se represents a serum-

445 achievable concentration of total Se and is less than the lower 95%
446 confidence limit of the non-observed adverse effect level (NOAEL).⁵²
447 Taking into consideration SeMeSeCys content as well as NO production
448 inhibitory potency (IC₅₀) of Se-enriched sauerkraut it was estimated that
449 this compound have a relative contribution of 38% to the IC₅₀ value.
450 These results suggest that SeMeSeCys could be acting synergistically
451 with other anti-inflammatory compounds present in Se-enriched
452 fermented cabbage. Our results agree with reports demonstrating the
453 effect of Se on attenuation of proinflammatory response in various cell
454 lines.⁵³ Several studies have been consistent showing the ability of Se in
455 the regulation of expression of the proinflammatory enzymes iNOS and
456 cyclooxygenase 2 (COX-2) through inactivation of the nuclear transcription
457 factor B (NF- κ B).^{54,55} Therefore, downregulation of proinflammatory
458 gene expression by Se explains the inhibition of NO production in LPS-
459 induced macrophages showed in the present study.

460 In conclusion, the addition of sodium selenite during natural
461 cabbage fermentation enhanced the formation of some GLS breakdown
462 compounds that exhibits health promoting properties, increased the
463 antioxidant and potential anti-inflammatory activities of sauerkraut. On
464 the other hand, SeMeSeCys was the primary selenocompound observed
465 in the resulting Se-enriched sauerkraut. Consequently, the consumption
466 of Se-enriched sauerkraut will contribute the Se dietary intake and it can
467 be considered as a good source of health-promoting compounds.

468

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651

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662

663 **FIGURE CAPTIONS**

664 **Figure 1.** Evolution of pH during cabbage fermentation in absence (NF)
665 or presence of Se (NFSe).

666

667 **Figure 2.** Chromatographic profile obtained by anion-exchange LC-ICP-
668 MS of A) a mixture of Se standards containing 100 $\mu\text{g/L}$ of each Se
669 species; and B) enzymatic extraction of raw cabbage (—), NF cabbage
670 (—) and NFSe cabbage (- - -).

671

672 **Figure 3.** Effect of different concentrations of raw, NF and NFSe
673 cabbage extracts (A); ABG (B); and SeMeSeCys (C) on NO production in
674 LPS-induced macrophages RAW 264.7 cells. Data represent the mean \pm
675 standard deviation of a triplicate from three independent experiments.
676 The same letter in the same bar indicates not significant difference ($P \leq$
677 0.05).

Table 1. Effect of Se enrichment on ascorbigen (ABG), indol-3-carbinol (I3C) and indol-3-acetonitrile (I3ACN) content in natural fermented cabbage (*Brassica oleracea* var. *capitata* cv. Megaton)*

Cabbage	ABG ($\mu\text{mol}/100\text{g d.m.}$)	I3C ($\mu\text{mol}/100\text{g d.m.}$)	I3ACN ($\mu\text{mol}/100\text{g d.m.}$)
Raw cabbage	16.43 \pm 1.76a	0.10 \pm 0.04a	0.24 \pm 0.07a
<i>Fermented cabbages</i>			
NF	218.18 \pm 4.21c	9.60 \pm 0.76b	2.66 \pm 0.17b
NFSe	205.26 \pm 3.16b	16.70 \pm 0.99c	3.00 \pm 0.17c

*) Mean value \pm standard deviation of three experiments. The same letter in the same column indicates no significant difference ($P \leq 0.05$)

Table 2. Effect of Se enrichment on ascorbic acid, Se biotransformation and water content in natural fermented cabbage (*Brassica oleracea* var. *capitata* cv. Megaton)*

Cabbages	Ascorbic acid (mg/100g d.m.)	Se ($\mu\text{g/g d.m}$)	SeMeSeCys ($\mu\text{g/g d.m}$)	Water (%)
Raw cabbage	329.45 \pm 8.95c	0.04 \pm 0.01a	NDa	91.6
<i>Fermented cabbages</i>				
NF	242.37 \pm 8.4b	0.07 \pm 0.01b	NDa	91.5
NFSe	229.86 \pm 8.50a	1.29 \pm 0.04c	0.74 \pm 0.02b	91.4

*) Mean value \pm standard deviation of three experiments. The same letter in the same column indicates no significant difference ($P \leq 0.05$).

Table 3. Effect of Se enrichment on antioxidant capacity and anti-inflammatory activity of natural fermented cabbage (*Brassica oleracea* var. *capitata* cv. Megaton)*

Cabbage	Antioxidant capacity ($\mu\text{mol Trolox/g d.m.}$)	Inhibitory potency (IC_{50}) of NO production ($\mu\text{g extract/mL}$)
Raw cabbage	74.78 \pm 0.28a	167.93 \pm 16.09c
<i>Fermented cabbages</i>		
NF	131.54 \pm 13.95b	83.96 \pm 8.50b
NFSe	162.96 \pm 3.71c	44.01 \pm 1.39a

*) Mean value \pm standard deviation of three experiments. The same letter in the same column indicates no significant difference ($P \leq 0.05$). IC_{50} is the concentration of cabbage extract ($\mu\text{g /mL}$) that resulted in 50% inhibition of NO production.

Table 4. Effect of Se enrichment on microbiological quality (cfu/g f.w.) of natural fermented cabbage (*Brassica oleracea* var. *capitata* cv. Megaton)*

Cabbage	Aerobic mesophilic bacteria	Anaerobic mesophilic bacteria	Lactic acid bacteria	Total Coliforms	Faecal Coliforms	Yeasts and moulds
Raw cabbage	5.19±0.11a	4.38± 0.11a	2.42± 0.11a	1.17± 0.18b	<1a	<1 ^a
<i>Fermented cabbages</i>						
NF	6.78± 0.10c	6.75± 0.11c	6.70± 0.14b	<1a	<1a	<1 ^a
NFSe	6.25±0.11b	6.28±0.13b	6.92±0.10c	<1a	<1a	<1 ^a

*) Mean value ± standard deviation of three experiments. The same letter in the same column indicates no significant difference ($P \leq 0.05$)

Figure 1

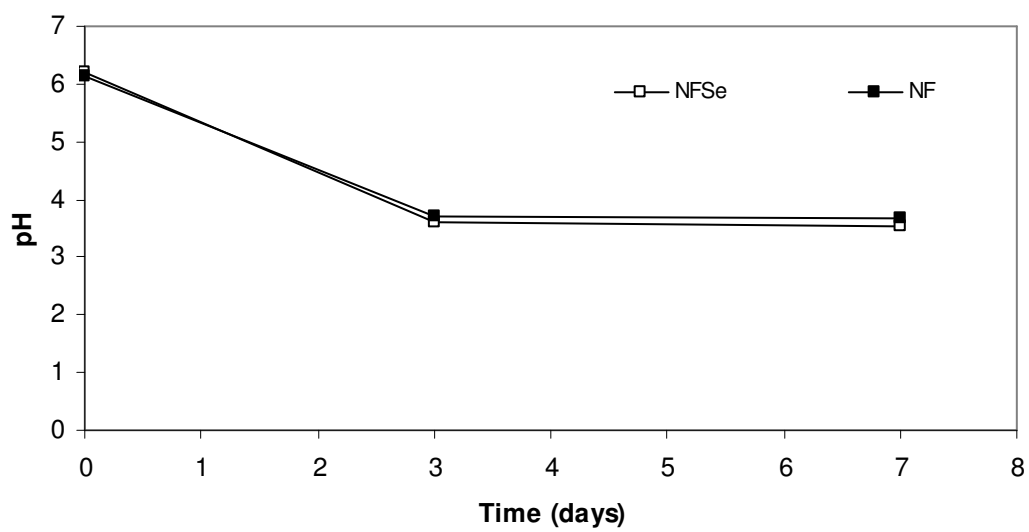


Figure 2

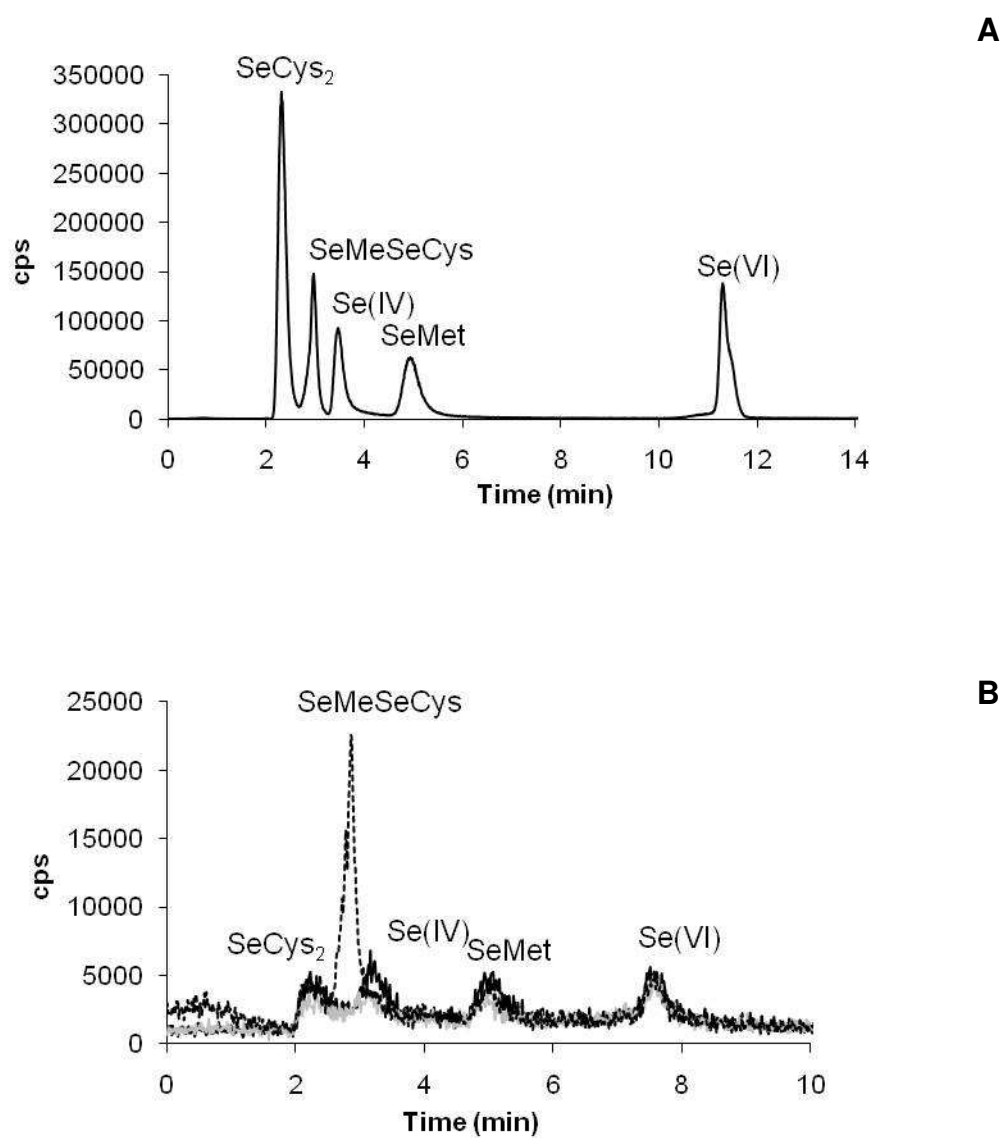
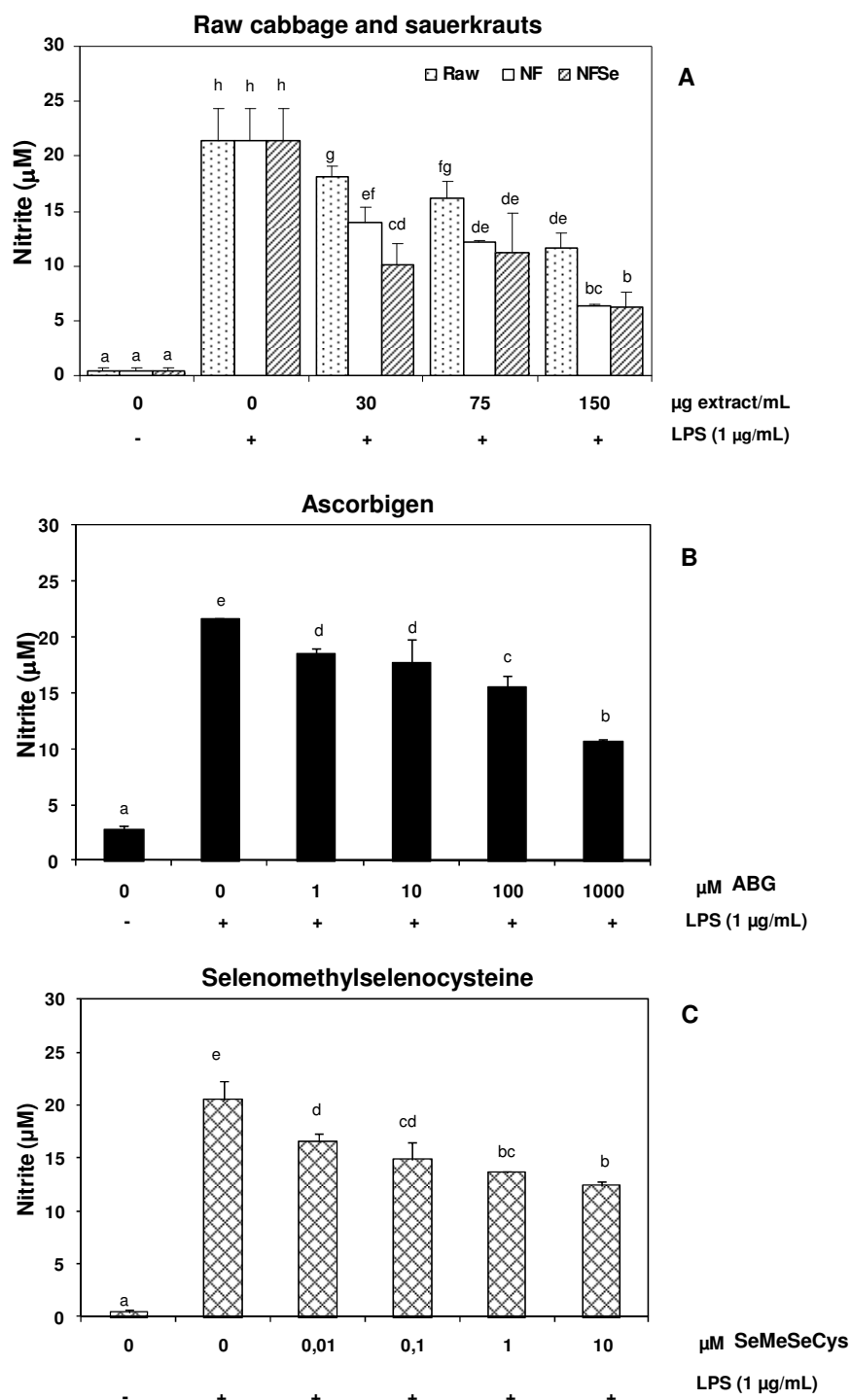


Figure 3



TOC

Selenium species in cabbage fermented with selenite solution

