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Detection of peanut (*Arachis hypogaea*) allergens in processed foods by immunoassay:
influence of selected target protein and ELISA format applied.

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Abstract

Direct competitive and sandwich ELISA formats developed to determine Ara h1 and Ara h2 proteins were applied in the detection of peanut in model biscuits prepared with a commercial peanut butter as ingredient. The sandwich format for Ara h2 protein could detect the addition of 2.5% peanut butter, whereas the same format for Ara h1 could not detect 5% added peanut. Direct competitive formats for Ara h1 and Ara h2 proteins could detect the presence of 1% and 0.05% peanut butter, respectively. Therefore, competitive format for Ara h2 was selected to be evaluated by four laboratories, obtaining adequate results in term of repeatability and reproducibility. Results obtained indicate that processing decreased the level of extracted protein and underestimated the amount of Ara h1 and Ara h2 proteins, the effect being more severe for Ara h1. The selection of the target protein and the ELISA format applied greatly influence the detection of peanut in processed foods.

Keywords: Ara h1, Ara h2, allergen, peanut detection, processed foods, ELISA assay.

57 **1. Introduction**

58 Food allergy has emerged as a serious public health problem over recent years and
59 its prevalence is rising, especially in industrialized countries. The reason appears to be
60 related to changes in dietary habits as well as to the use of complex technological
61 processes and ingredients in food industry (Nwaru et al., 2014; Sicherer & Sampson,
62 2010).

63 The estimated prevalence of peanut allergy in developed countries is between 0.6%
64 and 1.0%. Peanut allergy deserves particular attention because very small amounts of
65 peanut proteins can induce severe allergic reactions, it persists throughout life and it
66 accounts for most of food-induced anaphylactic reactions (Al-Muhsen et al., 2003; Wen
67 et al., 2007)

68 Until now, thirteen peanut proteins with allergenic capacity have been identified, and
69 designated as Ara h1 to Ara h13 (Bublin & Breiteneder, 2014; Sáiz et al., 2013). Ara h1
70 and Ara h2 proteins are considered as the major allergens of peanut, more than 65% of
71 peanut allergic individuals have specific IgE to Ara h1 and more than 71% to Ara h2.
72 (Scurlock & Burks, 2004). They are both major proteins in peanut, as they account for
73 12 to 16% and 5.9 to 9.3% of the total seed protein content, respectively (Koppelman et
74 al., 2001).

75 Ara h1 is a seed store glycoprotein that belongs to the vicilin family. It has a
76 molecular mass of 63.5 kDa in its monomer form and an isoelectric point of 5.2. It exists
77 as a trimer formed by three identical monomers stabilized mainly by hydrophobic
78 interactions. Ara h2 is a glycoprotein of the conglutinin family with a molecular mass of
79 17.5 kDa and an isoelectric point of 4.6 (Wen et al., 2007). Both proteins have been
80 found to maintain the IgE binding capacity after being exposed to thermal treatments or
81 *in vitro* digestion with pepsin, chymotrypsin and trypsin (Lehmann et al., 2006; Maleki
82 et al., 2000; Mondoulet et al., 2005).

83 The way to prevent peanut allergy is the strict avoidance of peanut consumption.
84 However, contamination with hidden allergens can occur due to inefficient cleaning
85 procedures of the production equipment or the use of contaminated raw ingredients,
86 among others (Vierk et al., 2002). The implementation of a management plan in the food

87 industry, the enforcement of labeling rules and its control by authorities are important
88 strategies for protecting against allergic reactions.

89 Therefore, reliable methods to detect peanut are required to ensure compliance with
90 the labeling legislation and to assist food manufacturers in order to improve consumer
91 protection. Enzyme-linked immunosassay (ELISA) is the technique most widely used by
92 food industries and official food control agencies for monitoring adventitious
93 contamination of food products by allergenic ingredients because of its sensitivity and
94 specificity (Monaci & Visconti, 2010). Several studies have been performed to develop
95 ELISA techniques to detect peanut in foods. These studies include the design of one
96 ELISA format (sandwich or competitive) and are based on the determination of one
97 selected target (a mixture of peanut proteins or a specific peanut protein) (Holzhauser &
98 Vieths, 1999; Kiening et al., 2005; Pomés et al., 2003; Stephan & Vieths, 2004).

99 It is worthwhile to remark that the determination of peanut proteins in foods can be
100 impaired by their interaction with compounds of the complex food matrix and
101 denaturation during processing. Consequently, protein extraction greatly decreases and
102 protein recognition by antibodies is reduced (Chassaigne et al., 2007; Fu & Maks, 2013;
103 Khuda et al., 2012).

104 Several recent studies have shown that results obtained by different ELISA tests give
105 significantly varying results in quantitative assays when they are used to detect peanut in
106 processed foods (Khuda et al., 2012; Poms et al., 2005). This variability may be
107 explained by the fact that ELISA tests can use different antigens as targets, antibodies
108 for antigen recognition and assay formats (Fu & Maks, 2013; Khuda et al., 2012;
109 Montserrat et al., 2013; van Hengel et al., 2007).

110 In this work, four ELISA assays for the detection of peanut, based on the
111 determination of Ara h1 or Ara h2 proteins (sandwich and direct competitive assay for
112 each protein) have been developed. The performance of the four assays was evaluated
113 using biscuits containing defined concentrations of a commercial peanut butter as
114 ingredient. The ELISA format and the target protein that gave the best sensitivity was
115 selected to determine peanut content in model biscuit the samples in blind duplicate by
116 four laboratories. For clarity and explanation, this part of the study is called
117 interlaboratory study, even though it did not involve the minimum number of

118 laboratories requested by a full interlaboratory study as defined in the ISO 5725 standard
119 (ISO, 1994).

120

121 **2. Materials and methods**

122 *2.1. Materials*

123 Raw peanuts and peanut butter from the Spanish variety was provided by Chocolates
124 Lacasa (Utebo, Spain). Peanut butter was prepared by roasting whole peanuts in a flame
125 oven at 225 °C for 27 min and afterwards, by grinding in a stone mill to obtain an
126 emulsion with dark color. Horseradish peroxidase (HRP, 250-503 units/mg) and goat
127 anti-rabbit IgG antibodies labelled with peroxidase were purchased from Sigma
128 Chemical (Poole, UK). Tetramethylbenzidine (TMB) substrate (Reference ZE/TMB125)
129 was obtained from ZEULAB (Zaragoza, Spain) and Maxisorp microtitration plates from
130 Nunc (Roskilde, Denmark). The bicinchoninic acid (BCA) assay kit was from Pierce
131 (Rockford, IL, USA).

132

133 *2.2. Methods*

134 *2.2.1. Isolation of Ara h1 and Ara h2*

135 Peanut proteins were extracted by stirring 20 g of ground raw peanut with 100 mL of
136 50 mM Tris-HCl buffer, pH 8.2. Proteins precipitated between 40 and 80% ammonium
137 sulphate saturation was collected by centrifugation, suspended in Tris buffer and filtered.
138 The extract was applied onto a Sephacryl S-200 column (90 x 2 cm). Fractions enriched
139 in Ara h1 were applied onto a Q-Sepharose column (15 x 1.5 cm) as previously
140 described (Montserrat et al., 2013) and fractions enriched in Ara h2 protein onto a
141 Sephadex G-50 column (80 x 1 cm). The purity of isolated proteins, determined by SDS-
142 PAGE was higher than 95%.

143

144 *2.2.2. Preparation and conjugation of antibodies to Ara h1 and Ara h2*

145 Antisera to Ara h1 and Ara h2 were obtained by immunization of rabbits as
146 previously described (Wehbi et al., 2005). All procedures were approved by the Ethic
147 Committee for Animal Experiments from the University of Zaragoza (Project Licence PI
148 48/10). The care and use of animals were performed following the Spanish Policy for

149 Animal Protection RD 1201/05, which meets the European Union Directive 86/609 on
150 the protection of animals used for experimental and other scientific purposes. Specificity
151 of antisera against Ara h1 or Ara h2 proteins were assessed by Western blotting analysis
152 (Franco et al., 2010).

153 Specific antibodies to Ara h1 or Ara h2 were purified by affinity chromatography
154 using immunosorbents of the corresponding proteins as described by Montserrat et al.
155 (2013). Antibodies were conjugated with HRP using the periodate method (Nakane &
156 Kawaoi, 1974).

157

158

159 *2.2.3. Sandwich and direct competitive ELISA assays for Ara h1 and Ara h2*

160 For the sandwich ELISA, plates were coated with 120 μ L per well of anti-Ara h1 or
161 anti-Ara h2 antibodies (5 μ g/mL), in 50 mM sodium carbonate buffer, pH 9.6 overnight
162 at 4 °C. Then, wells were blocked with 300 μ L of 2% (w/v) ovalbumin in 8 mM
163 Na_2HPO_4 , 3 mM KCl, 0.14 M NaCl, 1.5 mM KH_2PO_4 buffer, pH 7.4 (PBS) for 2 h at 37
164 °C and washed with PBS containing 0.5% Tween 20 (PBST). Afterwards, 100 μ L of
165 Ara h1 and Ara h2 standards or samples diluted in 0.1 M sodium borate buffer, pH 9.0
166 were added to the wells and incubated for 30 min at 37 °C. Then, wells were incubated
167 with 100 μ L of anti-Ara h1 or anti-Ara h2 antibodies HRP-conjugated diluted 1/6,000
168 and 1/10,000, respectively in the same buffer for 30 min at 37 °C. After washing with
169 PBST, wells were incubated with 100 μ L of TMB substrate for 20 min at room
170 temperature. Finally, the enzymatic reaction was stopped by adding 50 μ L of 2 M H_2SO_4
171 per well, and the absorbance determined at 450 nm using a microplate reader
172 (Labsystem Multiskan, Helsinki, Finland).

173 Calibration curves for the sandwich assay of Ara h1 was obtained by plotting
174 absorbance versus the concentration of standard solutions. For Ara h2, calibration curves
175 were obtained using the relationship between the value of absorbance and the logarithm
176 of the concentration of standard solutions. The concentration of Ara h1 and Ara h2 in the
177 test samples was determined by interpolating absorbance data in the corresponding
178 calibration curves.

179 For the direct competitive ELISA, plates were coated with 120 μ L per well of Ara h1
180 or Ara h2 proteins (5 μ g/mL) in 50 mM sodium carbonate buffer, pH 9.6. After
181 overnight incubation at 4 °C, wells were washed and blocked with ovalbumin as
182 indicated above. After washing with PBST, plates were incubated for 30 min at 37 °C
183 with 50 μ L of protein standards or samples diluted in 0.1 M borate buffer, pH 9.0 and 50
184 μ L of HRP-labeled anti-Ara h1 or anti-Ara h2 antibodies diluted 1/30,000 and 1/40,000,
185 respectively in the same buffer. Finally, after washing wells were incubated with TMB
186 substrate and enzymatic reaction stopped with H₂SO₄ before measuring absorbance at
187 450 nm.

188 Calibration curves for direct competitive assays were obtained using the logit log
189 model (Nix & Wild, 2000). The fraction bound ($r = B / B_0$), where B is the absorbance
190 of each standard and B₀ the absorbance of the blank standard was calculated. A plot of
191 logit (r) of standards against the log₁₀ of the concentration, where logit (r) = ln [(1-r) / r]
192 was obtained. The concentration of Ara h1 and Ara h2 in tests samples was determined
193 from its fraction bound, which is the ratio between absorbance of the sample and
194 absorbance of the blank standard (B₀).

195

196 *2.2.4. Preparation of model biscuits*

197 Biscuits were prepared at the pilot plant of the University of Zaragoza following
198 standard manufacturing processes. They were made by mixing 6 hen eggs (55-65 g), 120
199 g butter, 300 g wheat flour, 150 g sugar and peanut butter to obtain final concentrations
200 of 0, 0.25, 0.5, 1.0, 2.5 and 5.0%, (w/w). The ingredients were kneaded for 30 min using
201 a bread and dough maker (Deluxe: Bread and Dough Maker, Oster, USA) equipped with
202 a blade type "pigtail". Then, 40 g of homogenized material was placed in a baking
203 mould (10 cm diameter) and pressed to obtain round cookies of 1 cm height. Then,
204 biscuits were introduced into an oven and cooked at 160 °C for 12 min.

205

206 *2.2.5. Extraction procedure*

207 Food samples purchased from local retailers and model biscuits were ground into fine
208 powder with a mincer. An amount of 3.00 \pm 0.01 g of ground samples were extracted in
209 30 mL of 0.1 M sodium borate buffer, pH 9.0 and incubated in a shaking water bath at

210 30 °C for 15 min. Extracts were clarified by centrifugation at 3,000 x g for 15 min, and
211 the supernatants stored in aliquots at -20 °C until use. Supernatants were directly assayed
212 in the ELISA plates.

213

214 2.2.6. *Evaluation of direct competitive ELISA for Ara h2*

215 The evaluation study was performed following the procedure previously described
216 (Abbot et al., 2010; AOAC, 2012). Four laboratories with ELISA experience
217 participated in this study to evaluate the direct competitive ELISA for Ara h2 protein to
218 detect peanut in model biscuits. The study was coordinated by the group of the
219 University of Zaragoza.

220 The samples to be sent to the participants were prepared as follows. Biscuits
221 containing 0, 0.25, 0.5, 1.0 and 2.5% peanut were ground and 3.00 ± 0.01 g was
222 weighted into 50 mL plastic tubes. Biscuits with peanut concentrations of 0.01, 0.05 and
223 0.1% were prepared by mixing appropriate quantities of the ground 0.25% samples with
224 the blank sample into plastic tubes to give a total weight of $3.00 + 0.01$ g. Extraction of
225 test samples was performed as indicated above.

226 The coordinator provided two sets of 8 pre-weighed test samples, randomly coded,
227 and ZEULAB provided the ELISA kits containing plates, reagents, standards and
228 instructions. Each set of samples was extracted once in different days and analyzed in
229 triplicate in the ELISA assay. Absorbance data of calibration standards and blind
230 samples of each set were sent to the coordinator. Calibration curves were obtained for
231 each ELISA assay using the logit log model. Determination of repeatability and
232 reproducibility data were calculated according to ISO 5725.

233

234 **3. Results**

235 3.1. *Specificity of antisera to Ara h1 and Ara h2*

236 The specificity of antisera against Ara h1 and Ara h2 proteins were assessed by
237 Western blotting (Figure 1). Results showed that antibodies to Ara h1 only reacts with
238 Ara h1 and antibodies to Ara h2 only bind to Ara h2. In both cases, no reaction was
239 observed with any other protein from crude peanut extract demonstrating that antisera
240 obtained were specific for each protein.

241

242 3.2. *Development of sandwich and direct competitive ELISA for Ara h1 and Ara h2*

243 Immunoassay formats for Ara h1 and Ara h2 were optimized to choose the assay
244 conditions which gave the highest sensitivity, that were chosen for the validation and the
245 interlaboratory study. The relationship found was linear within the range of
246 concentrations between 20 ng/mL and 2 µg/mL for direct competitive assays and for the
247 sandwich format of Ara h2, and curvilinear between 20 ng/mL and 800 µg/mL for the
248 sandwich format of Ara h1 protein. All assays gave regression coefficients $r^2 \geq 0.985$
249 (Figure 2). The detection limit (LOD) of the immunoassays tests was determined as the
250 mean concentration of Ara h1 and Ara h2 corresponding to the absorbance of eight
251 replicates of the blank standard plus 3.3 times the standard deviation (Miller et al., 2006)
252 (Table 1).

253

254 3.3. *Determination of peanut in model biscuits*

255 Results obtained in the analysis of model biscuits which contained different amounts
256 of peanut butter using sandwich and direct competitive assays to determine Ara h1 and
257 Ara h2 proteins are shown in Figure 3. Biscuit samples were extracted in three different
258 days and assayed by triplicate. Previously, a cut-off value was established to consider a
259 sample as positive for peanut addition for each ELISA test. This value was estimated as
260 the average concentration of the blank biscuit plus 3.3 times the value of its standard
261 deviation (Lexmaulová et al., 2013) (Table 1). The assumption of this value ensures that
262 interference caused by the matrix effect in each assay is minimized.

263 In this study, biscuit samples without added peanut gave a concentration value below
264 the cut-off calculated for each format assay. The sandwich format based on Ara h2
265 protein could detect the addition of 2.5% peanut, whereas the same format for Ara h1
266 could not detect samples containing 5.0% peanut. Direct competitive assays for Ara h1
267 and Ara h2 proteins could detect biscuits samples containing 1.0% and 0.05% of peanut
268 addition, respectively. Biscuit samples which contained a lower percentage of peanut
269 than those indicated above gave false-negative results in the corresponding assays and
270 those which contained higher percentages gave a concentration of Ara h1 and Ara h2
271 that increased gradually.

272 On the other hand, the concentration of soluble proteins, estimated by the
273 bicinchoninic acid, and of Ara h1 and Ara h2 was determined in peanut butter and in raw
274 dough of biscuits. The protein concentration in the peanut butter extract was of $8.1 \pm$
275 0.4% (w/w) and the concentration of Ara h1 and Ara h2 proteins, estimated using the
276 direct competitive assays was $1,000 \pm 20$ and $2,750 \pm 13$ mg/kg, respectively. Samples
277 of raw peanut from the same variety were also analyzed and a protein content of $16.2 \pm$
278 0.4% (w/w) and concentrations of Ara h1 and Ara h2 of $20,244 \pm 68$ and $5,873 \pm 87$
279 mg/kg respectively, were obtained. When these proteins were determined in biscuits
280 added with 1.0 and 5.0% peanut butter, the concentration of Ara h1 and Ara h2 was
281 found to be about 1% and 45% of that in the raw dough before the baking treatment.

282

283 *3.4. Cross-reactivity study*

284 The specificity of anti-Ara h1 and Ara h2 antibodies was also examined by testing its
285 cross-reactivity with other food ingredients such as, tree nuts (almond, cashew nut,
286 pistachio, walnut and hazelnut), legumes (chick pea, soya, green pea and lentil), and
287 ingredients used in the elaboration of biscuits (wheat, milk, egg and sugar). Extracts of
288 all ingredients and peanuts were prepared following the extraction protocol and tested
289 undiluted. Protein concentration of extracts assayed ranged from 0 to 32 mg/kg. All
290 ingredients gave a small decrease (in competitive format) or increase (in sandwich
291 format) of the absorbance value compared to the blank standard indicating a certain
292 degree of interference (results not shown). Concentration values of Ara h1 and Ara h2
293 determined in these ingredients were below the cut-off established for each ELISA assay
294 to consider a sample as positive for peanut protein.

295

296 *3.5. Evaluation of direct competitive ELISA for Ara h2*

297 The direct competitive ELISA test to determine Ara h2 protein was evaluated by four
298 laboratories for the detection of peanut in the model biscuits. Concentration of Ara h2 in
299 two set of blind biscuit samples prepared with peanut butter were determined.

300 Using the standards of Ara h2 indicated in Table 1, calibration curves were obtained
301 for every ELISA plate using the logit log model, obtaining regression coefficients higher
302 than 0.976. The concentration of Ara h2 in test samples was calculated as indicated

303 above. The mean concentration of Ara h2 obtained for each set of samples by each
304 laboratory is shown in Table 2.

305 The cut-off value for the interlaboratorial study was determined as 3.3 times the
306 reproducibility (S_R) of the blank biscuit (Lexmaulová et al., 2013), obtaining a value of
307 0.81 mg/Kg.

308 The four laboratories obtained concentrations of Ara h2 in the blank biscuit samples
309 below the cut-off established for interlaboratory study to consider a sample as positive,
310 indicating that no false-positive samples were found. For all laboratories, Ara h2 was
311 detected in samples with a percentage equal or higher than 0.05% of peanut butter. At
312 0.01% of peanut addition, the concentration of Ara h2 was below the cut-off with the
313 exception of one laboratory. At higher percentages, concentration of Ara h2 increased
314 for all laboratories. Results and performance characteristics (repeatability and
315 reproducibility data) of the interlaboratory study are summarized in Table 3. Values of
316 repeatability RSD (RSD_r) ranged between 15.83 and 44.07% and values of
317 reproducibility RSD (RSD_R) between 30.18 and 111.13%.

318

319 **4. Discussion**

320 The search for the selection of an immunoassay format and a target protein to detect
321 peanut in processed foods led us to develop direct competitive and sandwich ELISA
322 formats to determine Ara h1 and Ara h2 proteins, the two major peanut allergens.

323 The optimum conditions led to the development of sandwich and direct competitive
324 ELISA tests with sensitivities comparable to those previously obtained for Ara h1 and
325 Ara h2 proteins (Pomés et al., 2003; Schmitt et al., 2004).

326 Certain degree of interference was observed between Ara h1 and Ara h2 with basic
327 food ingredients when they were analyzed using competitive ELISA tests. The existence
328 of cross-reactivity between Ara h1 and other vicilin storage proteins of legumes such as
329 soya, green pea and beans have been reported (Beardslee et al., 2000; Sicherer et al.,
330 2000). These proteins have some 30-45% of amino acids in common with peanuts and a
331 similar folding. However, homology at surface residues requires a higher degree of
332 amino acid identity (Pomés et al., 2003). In this study, we did not observe a higher level
333 of interference when analyzing legumes compared to other foods. Thus, it is assumed

334 that interference could be produced by non-specific interaction between components of
335 the food matrix and antibodies.

336 Model biscuits containing several different percentages of peanut butter as ingredient
337 were analyzed using developed ELISA assays. We selected this processed material to
338 prepare biscuits because it is commonly used in the elaboration of nougats,
339 confectionery products, seasoning blends, bakery mixes, frostings, fillings, chocolate,
340 creams and cereal bars. Results obtained indicated that the processing of peanut to
341 obtain butter caused a decrease in the level of extracted proteins of about 50% and a loss
342 of immunoreactive proteins of about 95% and 53% for Ara h1 and Ara h2, respectively.

343 Our results are in good agreement with those previously reported on the effect of
344 thermal processing of peanut on protein solubility and detectability by ELISA
345 techniques (Chassaigne et al., 2007; Fu & Maks, 2013; Schmitt et al., 2010). Thus,
346 Chassaigne et al. (2007) found that roasting of peanuts under mild or strong conditions
347 decreased extraction efficiency of proteins by 75% and 82%, respectively. In the same
348 study, the concentration of Ara h1 and Ara h2 proteins under mild and strong roasting of
349 peanuts, determined by ELISA kits, were reported to be about 15% and 8% of that of the
350 raw peanut extract for Ara h1 and 59% and 47% for Ara h2, respectively. Fu & Maks
351 (2013) studied the effect of heat treatment of peanut flour on the solubility of proteins
352 and compared the performance of two commercial ELISA test kits targeting whole
353 peanut proteins or Ara h1 for quantitation of residual peanut. They found that dry
354 heating at 232 and 260 °C for 10 min caused an approximately 49.9% and 85.7%
355 decrease in the amount of proteins extracted, respectively. Likewise, the two ELISA kits
356 underestimated the level of proteins in the samples, the degree of immunoreactivity loss
357 being greater for the kit targeted to Ara h1 than for the kit targeted to whole peanut
358 proteins, about 62.7% and 75.0% at 232 °C and 98.5% and 99.4% at 260 °C for kits
359 targeted whole peanut proteins and Ara h1, respectively.

360 Our study confirms that thermal processing of peanuts decreases solubility of peanut
361 proteins as well as immunoreactivity of Ara h1 and Ara h2 proteins, the effect being
362 more marked for Ara h1. This fact could be attributed to a higher degree of denaturation
363 and/or aggregation of Ara h1 compared to Ara h2, which causes a higher loss of epitopes
364 recognized by antibodies and a higher reduction of its solubility. Our results and those

365 obtained by other authors (Chassaigne et al., 2007; Schmitt et al., 2010) support the
366 previously reported good thermal stability of Ara h2 (Owusu-Apenten, 2002) and
367 suggest that Ara h2 would be a better target than Ara h1 when immunoassays are going
368 to be used for the detection of peanut in processed foods.

369 Results obtained in the analysis of model biscuits which contained different amounts
370 of peanut butter indicate that direct competitive formats have a higher sensitivity to
371 detect added peanut butter than the sandwich formats. Differences in the recognition of
372 antigen by competitive and sandwich ELISAs could be due to the former requires only
373 one site of interaction with the antibodies whereas the later requires two binding sites. It
374 should be also considered that the way that specific antibodies are presented to its target
375 protein is different depending on the ELISA format. In the sandwich format, capture
376 antibodies are coated on the wells whereas in the competitive format antibodies are in
377 solution and thus, the accessibility of adsorbed antibodies may differ from the antibodies
378 in solution.

379 Our results are in accordance with those reported by de Luis et al. (2008) using
380 competitive and sandwich ELISA assays based on the determination of ovomucoid to
381 detect egg in model foods. In that study, both formats performed well to detect egg
382 added to pasteurized sausages and baked bread whereas only the competitive format
383 could detect egg in high heat treated foods such as sterilized pâté.

384 Our results also show that sandwich and direct competitive assays based on the
385 determination of Ara h2 protein are able to detect lower percentages of added peanut
386 compared to their counterparts for Ara h1. These findings can be attributed to a more
387 severe denaturation and/or aggregation for Ara h1 than for Ara h2 induced by the baking
388 process, which result in a lower level of extracted Ara h1 and/or in a lower recognition
389 of this protein by their specific antibodies, as indicated above.

390 Pomés et al. (2003) developed a sandwich ELISA for Ara h1 to monitor peanut
391 allergen in foods that could detect peanut in cookies and pancake mix spiked with 0.2%
392 of ground peanut. They observed that the recovery of Ara h1 progressively decreased
393 when lower amounts of peanut were added to those foods, obtaining recoveries in
394 biscuits of 86% and 6% at spiked levels of 16% and 0.2%, respectively. This fact
395 indicates that compounds of the matrix impaired recognition of Ara h1 by its specific

396 antibodies. Peng et al. (2013) developed a monoclonal-antibody sandwich ELISA for
397 Ara h1 that could detect milk samples spiked with pure Ara h1 at levels between 60 and
398 240 ng/mL, obtaining recoveries ranging from 95.45 to 105.18%.

399 The performance of the assays developed in our work to detect peanut addition is
400 difficult to compare with other studies (Peng et al., 2013; Pomés et al., 2003). Although
401 the standards used are composed in all these studies of Ara h1, we used food samples, in
402 which a commercial peanut butter was added at the ingredient stage and afterwards
403 subjected to processing, whereas in the others, food products analyzed were spiked with
404 pure Ara h1 (Peng et al., 2013) or with a raw peanut extract (Pomés et al., 2003). The
405 use of spiked foods is useful to determine the effect of food matrix but they do not
406 provide information about the effect of processing on assay performance. In the last few
407 years, the potential effects of processing on the quantitation of proteins by ELISA have
408 become recognized. The use of incurred samples, in which the allergenic food is added
409 as ingredient and afterwards, processed in a manner mimicking as closely as possible the
410 actual conditions under which the sample matrix would normally be manufactured,
411 allows evaluating the actual effect of processing on the detection efficiency of an
412 immunoassay (Khuda et al., 2012; Taylor et al., 2009). Although incurred samples are
413 considered difficult and costly to obtain, some regulatory bodies may be unwilling to
414 consider approval of validation data without the inclusion of data generated with
415 incurred samples prepared with material for the allergen being targeted (AOAC, 2012).

416 Recently, Khuda et al. (2012) performed a study to establish the effect of food
417 processing on peanut detection by five commercial ELISA kits using cookie dough
418 prepared with defatted light-roasted peanut flour before baking. These authors obtained
419 that recovery was drastically reduced after baking at 190 °C for 30 min, being less than
420 18% at all added levels.

421 Our study and others demonstrates that ELISA tests could not give accurate results
422 when they are used to determine allergenic proteins present in thermal processed foods
423 due to changes in solubility and immunoreactivity of the target proteins (Fu & Maks,
424 2013; Khuda et al., 2012). Therefore, an understanding of the effects of processing on
425 allergen structure in a specific matrix, as it relates to immunoreactivity and solubility, is
426 necessary to evaluate the performance of ELISA methods to detect allergens in

427 processed foods. The limitations of immunoassays should be considered when they are
428 going to be applied in the evaluation of food allergen control programs.

429 Performance characteristic of direct competitive ELISA for Ara h2 were determined
430 within the interlaboratorial study. This ELISA test could detect percentages of peanut
431 butter addition higher than 0.05% and false-negative results were found at 0.01%
432 addition. It has been shown that relatively low values of RSD_R from 30.18 to 53.47% for
433 model biscuits can be achieved at 0.05-5% peanut addition, obtaining the highest value
434 at the lowest levels of peanut addition (0.01%), in which sample Ara h2 could not be
435 detected.

436 Poms et al. (2005) carried out an interlaboratory validation of five commercial
437 ELISA test kits for the determination of peanut in two food matrices (biscuits and dark
438 chocolate) at four levels of peanut contamination. They found that variance of results
439 between laboratories (RSD_R) for biscuits for the different concentration levels ranged
440 between 23.4 and 127.0%. Matsuda et al. (2006) evaluated the analytical performance of
441 two ELISA kits to detect peanut in an interlaboratory study and found RSD_R values of
442 14% and 9% for cookies added with peanut proteins at a level of 10 $\mu\text{g/g}$ of food.
443 Lexmaulová et al. (2013) performed a collaborative study to validate an ELISA method
444 for the quantitative determination of peanut protein in foods. They used six real foods
445 with peanut declared in the ingredient list and obtained variation coefficient of
446 reproducibility between 31.4 and 59.4% depending on the sample. Thus, RSD_R values
447 obtained in our study are in the range of those reported in other studies.

448

449 **5. Conclusions**

450 In this study, direct competitive and sandwich ELISA formats to determine Ara h1
451 and Ara h2 proteins were developed and assayed in model biscuits prepared with a
452 commercial peanut butter as ingredient. Direct competitive formats could detect lower
453 levels of peanut butter in biscuits compared to sandwich formats. Moreover, ELISA
454 assays based on the determination of Ara h2 protein were able to detect lower
455 percentages of peanut than their counterparts for Ara h1. Therefore, direct competitive
456 format for Ara h2 were selected to be evaluated by four laboratories, obtaining adequate
457 results in term of repeatability and reproducibility.

458 Results obtained revealed that detected levels of Ara h1 and Ara h2 were drastically
459 reduced after the roasting of peanuts to obtain the peanut butter used as ingredient and
460 also after the baking of biscuits, the effect being more marked in the case of Ara h1. This
461 is an important point, as these proteins that are underestimated by ELISA have been
462 reported to retain or even to increase their allergenicity after processing in sensitized
463 individuals.

464 These findings underline the fact that the determination of allergenic proteins is
465 greatly affected by the nature of the immunoassay format, the target protein and the food
466 processing conditions. The limitations of each allergen assay should be considered
467 before applying ELISA assays for evaluation of food allergen control programs and to
468 assess allergen risk management studies.

469

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FIGURE LEGENDS

Figure 1: SDS-PAGE (a) and Western-blotting against rabbit antiserum to Ara h1 (b) and Ara h2 (c) of raw peanut extract.

Figure 2: Calibration curves obtained for sandwich (a, b) and direct competitive (c, d) ELISA formats for determination of Ara h1 (a, c) and Ara h2 (b, d) concentration in standard solutions of pure proteins.

Figure 3: Concentration of immunoreactive Ara h1 (a, c) and Ara h2 (b, d) in model biscuits added with different amounts of peanut butter. Sandwich (a, b) and direct competitive (c, d) ELISA. Values are the mean + SD of three sample extractions assayed by triplicate expressed in mg/kg.

Lines indicate the cut-off value above which biscuits are considered positive for peanut butter addition, and were calculated as the mean value + 3.3 SD of the blank biscuit.

Figure 1

602

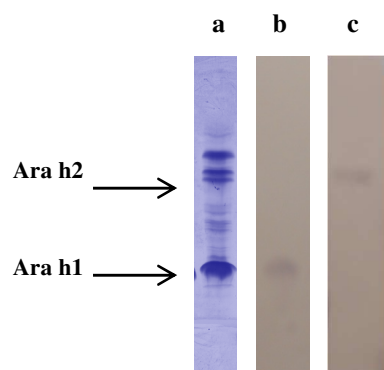


Figure 2

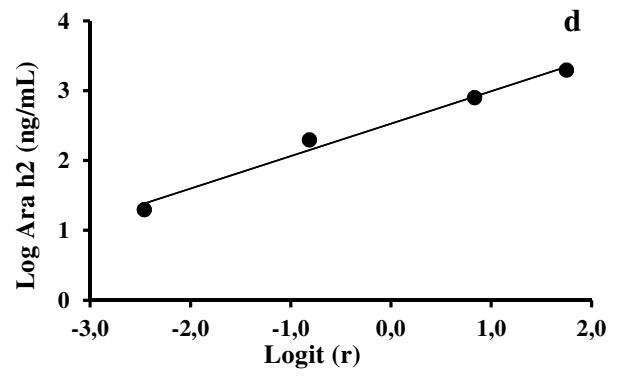
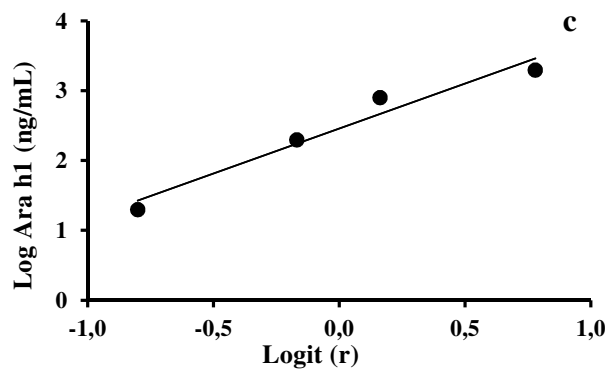
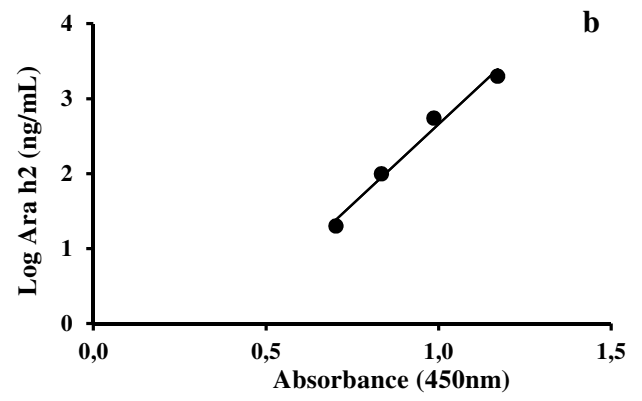
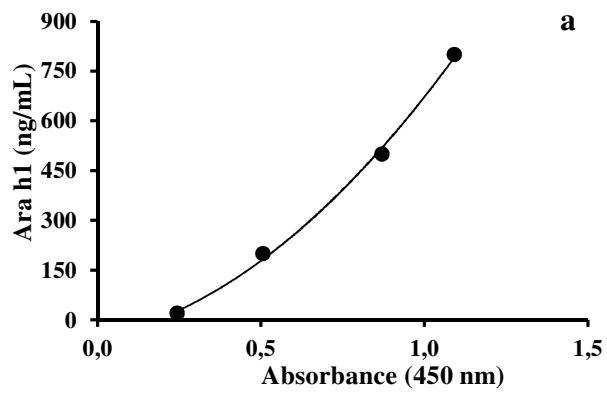


Figure 3:

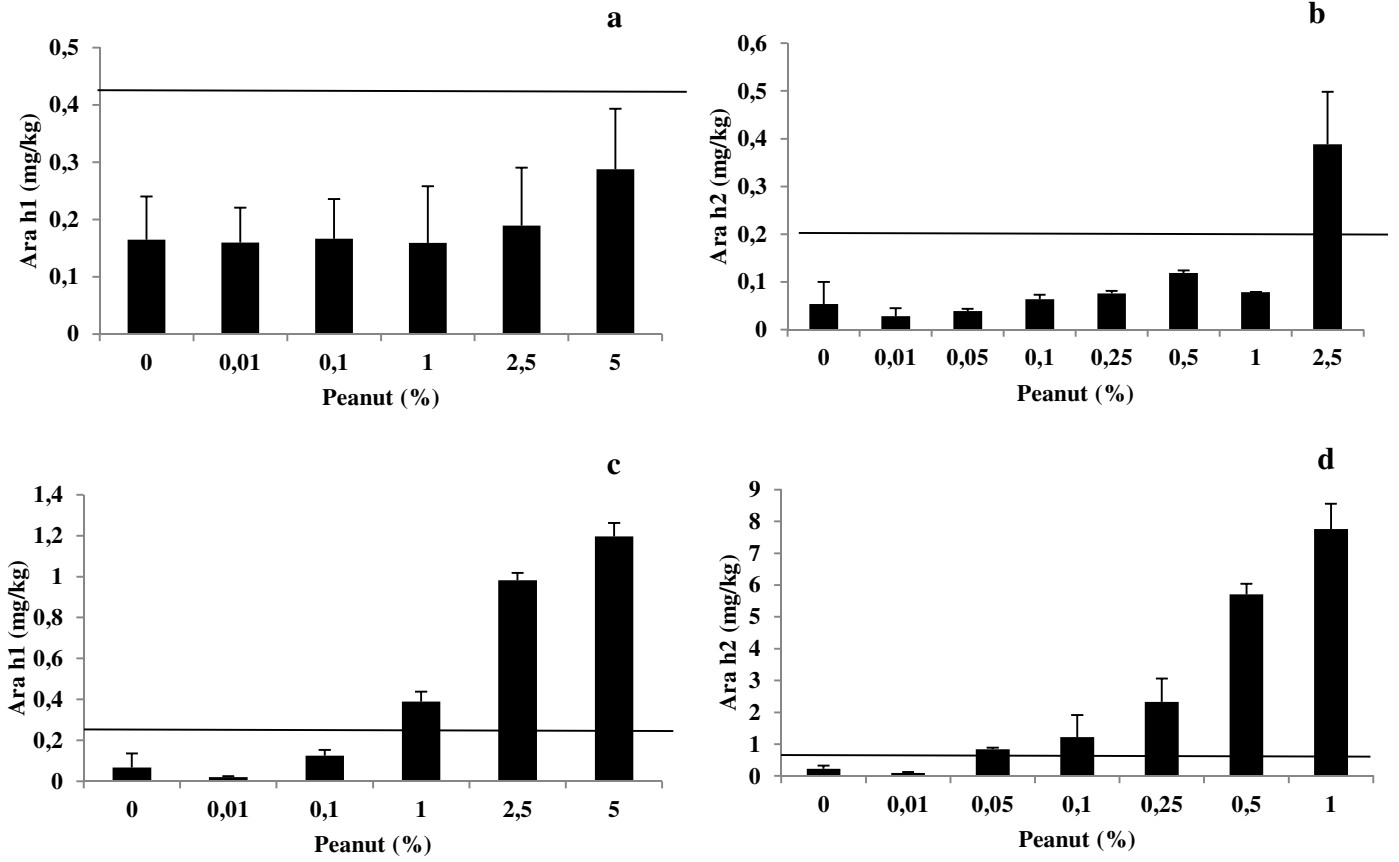


Table 1: Limit of detection (LOD) of the ELISA tests for Ara h1 and Ara h2 and cut-off establish for the ELISA tests to determine a biscuit sample as positive for peanut addition. Calibration points correspond to the protein concentration of standards used in each ELISA tests. Mean value + SD are given in brackets.

Test format	Target protein	LOD (mg/kg)	Cut-off (mg/kg)	Calibration points (mg/kg)
Sandwich	Ara h1	0.10 (0.04 ± 0.02)	0.42 (0.16 ± 0.08)	0-0.2-2.0-5.0-8.0
Sandwich	Ara h2	0.13 (0.11 ± 0.01)	0.20 (0.05 ± 0.05)	0-0.2-1.0-5.5-20.0
Competitive	Ara h1	0.19 (0.10 ± 0.03)	0.30 (0.07 ± 0.07)	0-0.2-2.0-8.0-20.0
Competitive	Ara h2	0.06 (0.02±0.011)	0.64 (0.24 ± 0.12)	0-0.2-1.0-5.5-20.0

Table 2: Results obtained by the four participating laboratories for the determination of Ara h2 (mg/kg) in model biscuits added with different percentages of peanut butter, using the direct competitive ELISA format.

Peanut Butter (%)	Assay 1				Assay 2			
	Lab 1	Lab 2	Lab 3	Lab 4	Lab 1	Lab 2	Lab 3	Lab 4
0	0.18 ± 0.16*	0.53 ± 0.11	0.60 ± 0.09*	0.38 ± 0.25*	0.30 ± 0.03*	0.61 ± 0.22*	0.19 ± 0.09*	0.60 ± 0.29*
0.01	0.12 ± 0.02*	0.81 ± 0.34	0.73 ± 0.21*	0.42 ± 0.36*	0.16 ± 0.13*	1.48 ± 0.31	0.48 ± 0.07*	0.40 ± 0.25*
0.05	0.95 ± 0.50	1.87 ± 0.11	1.38 ± 0.33	1.72 ± 0.46	1.20 ± 0.18	1.70 ± 0.20	0.98 ± 0.09	1.35 ± 0.28
0.10	1.03 ± 0.21	2.69 ± 0.59	2.31 ± 0.19	3.10 ± 0.11	1.76 ± 0.57	2.27 ± 0.42	1.04 ± 0.24	1.74 ± 0.21
0.25	1.82 ± 0.24	4.02 ± 0.52	3.06 ± 0.29	6.02 ± 1.13	2.76 ± 0.29	3.75 ± 0.47	2.60 ± 0.38	3.86 ± 1.27
0.50	6.10 ± 1.45	9.91 ± 0.93	5.69 ± 0.60	7.11 ± 0.65	5.53 ± 0.88	5.62 ± 1.18	4.65 ± 0.41	6.97 ± 0.74
1.00	7.93 ± 3.48	20.53 ± 2.11	14.33 ± 2.28	15.56 ± 1.03	8.33 ± 0.47	9.69 ± 0.37	6.58 ± 1.46	15.16 ± 2.00
2.50	62.75 ± 9.38	51.43 ± 20.5	21.87 ± 1.53	21.45 ± 7.69	49.32 ± 6.42	27.15 ± 5.09	44.55 ± 5.22	43.75 ± 2.21

*Food samples with concentration values below the cut-off established for the interlaboratory study.

Table 3: Results of the interlaboratory study. Performance criteria (repeatability and reproducibility data)

Performance characteristics	Abbreviation	Peanut Butter (%)							
		0.00	0.01	0.05	0.10	0.25	0.50	1.00	2.50
Total number of laboratories	P	4	4	4	4	4	4	4	4
Total number of replicates	n	8	8	8	8	8	8	8	8
Mean value	\bar{X}	0.42	0.57	1.39	1.99	3.49	6.45	12.26	40.28
Repeatability SD	S_r	0.169	0.253	0.221	0.721	0.856	1.572	4.714	14.924
Reproducibility SD	S_R	0.247	0.638	0.506	0.907	1.864	1.946	5.964	17.755
Repeatability RSD	RSD_r	39.91	44.07	15.83	39.19	24.56	24.38	38.44	37.05
Reproducibility RSD	RSD_R	58.32	111.13	36.39	45.55	53.47	30.18	48.62	44.07
Repeatability limit	r	0.473	0.708	0.618	2.018	2.397	4.401	13.199	41.788
Reproducibility limit	R	0.691	1.787	1.416	2.540	5.220	5.449	16.698	49.713

