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1 **The major soybean allergen Gly m Bd 28K induces hypersensitivity reactions**
2 **in mice sensitized to cow's milk proteins.**

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24

25 **Abstract**

26

27 Reactions to soy have been reported in a proportion of patients with IgE-mediated cow's milk
28 allergy (CMA). In this work, we analyzed if Gly m Bd 28K /P28, one of the major soybean
29 allergens, is a cross-reactive allergen with cow milk proteins (CMP). We showed that P28 was
30 recognized by IgE sera from CMA patients and activated human peripheral basophils
31 degranulation. Moreover, IgE sera of mice exclusively sensitized to CMP recognized P28.
32 Splenocytes from sensitized animals secreted IL-5 and IL-13 when incubated with CMP or soy
33 proteins, but only IL-13 when treated with P28. In addition, skin test was strongly positive for
34 CMP and weakly positive for P28. Remarkably, milk-sensitized mice showed hypersensitivity
35 symptoms following sublingual challenge with P28 or CMP. Using bioinformatics' tools seven
36 putative cross-reactive epitopes were identified. In conclusion, using in vitro and in vivo test we
37 demonstrated that P28 is a novel cross-reactive allergen with CMP.

38

39 **Keywords:** Food allergy; P28 soybean protein; Bovine caseins; Cross-reactivity.

40

41 Introduction

42 Food allergies are currently an important medical and social problem. Approximately 8% of
43 children and 1-2% of adults have some type of food allergy¹. Particularly, cow's milk allergy
44 (CMA) constitutes the main food allergy observed in the pediatric population in Argentina and in
45 many other countries²⁻⁴. Soy-based formulas are frequently used as a dairy substitute during the
46 avoidance treatment. Although both milk and soy belong to the group of eight main allergenic
47 foods, often referred as the Big-8, soy allergy (SA) is not as prevalent as CMA⁵. However,
48 several cases of clinical soy-intolerance, at the beginning of the avoidance treatment in milk
49 allergic patients, have been reported⁶⁻⁹. This situation introduces a further complexity in the
50 management of patients. The identification and characterization of the components of soybean
51 allergens responsible for allergy may be useful not only to understand the cross-reactivity
52 observed, but also to raise a immunotherapy^{10,11} using mutated and hypoallergenic soy
53 allergens¹². In this regard, we are focused on the investigation of cross-recognition between CMP
54 and soy components, and we have found that bovine caseins (BC)^{13,14} cross-react with three soy
55 proteins (SP): Gly m 6.0401 sub G4, Gly m 5.0101 sub α and, more recently, Gly m Bd 30K)¹⁵⁻
56 ¹⁹.

57 In this work, we focused in Gly m Bd 28K/P28, major soybean allergen²⁰⁻²⁵. P28 is a seed
58 glycoprotein that was isolated from 7S globulin fraction and belongs to the cupin protein
59 superfamily. Several major plant food allergens are included in this superfamily, such as the
60 major storage globulins of legumes and nuts (Ara h 1, Ara h 3 from peanut, MP27/MP32 from
61 pumpkin, Gea 8 from carrot, and Gly m 5 and Gly m 6 from soybean)²⁶⁻²⁸. The cupin
62 superfamily includes a diverse group of proteins that share a beta-barrel core domain with very
63 low level of sequence identity²⁹. These structural domains are thought to be highly stable,

64 resisting thermal denaturation and certain types of proteolysis. Thus, the stability factor
65 conferred by this fold may potentiate protein immunogenicity and allergenicity³⁰. Gly m Bd 28 K
66 cDNA encodes for a 476 amino acids residues protein a (52,9 kDa), which exhibits a high
67 homology with the MP27/MP32 proteins in pumpkin seeds and carrot globulin-like protein^{21,31}.
68 In developing seeds, Gly m Bd 28 K preproprotein is processed into the mature 28kDa N-
69 terminal and 23 kDa C-terminal polypeptides^{20,21}. Both resulting polypeptides are recognized by
70 IgE antibodies of patients sensitive to soybean^{20,22,23,25}, and based on these assays Gly m Bd 28 K
71 is considered as a major soybean allergen. However, the *in vivo* ability of P28 to trigger
72 hypersensitive symptoms has not been evaluated. Currently, animal models are a valuable
73 biological tool to study the correlation of the immunochemical cross-reactivity assessed with the
74 clinical outcome. In this study, we aimed to investigate the clinical relevance of cross-reactivity
75 detected between Gly m Bd 28K /P28 and milk proteins.

76

77 **Materials and Methods**

78 **Protein extracts and antibodies.** Soybean protein extract was obtained from *Glycine max* L.
79 Merr. seeds as described previously¹⁶. Cow's milk protein extract was prepared from commercial
80 non-fat dry milk (Svelty, Nestle). Proteins were extracted with phosphate saline buffer pH 7.4
81 and filtered. The presence of soy components in the CMP extract was previously discarded by
82 indirect ELISA using a (SP)-specific rabbit antiserum.

83 Sera of 10 pediatric patients (ranged 9 month to 7 years old, 5 male 5 female) and diagnosed as
84 milk allergic according to history, skin Prick test, and serum specific IgE were used. Milk allergy
85 was not diagnosed with the double-blind placebo-controlled food challenge because it is not
86 currently performed in Argentina; instead milk elimination during two weeks and open challenge
87 is done. Soy allergy was ruled out by history. In addition, sera from healthy individuals, with no
88 allergy history and normal level of serum IgE according to age, or from patients allergic to
89 aeroallergens, with no CMP-specific IgE antibodies and history of food allergy, were included as
90 controls. Three monoclonal antibodies (mAb) with differential specificity for α -casein (1D5), β -
91 casein (4C3) and κ -casein (3B5), previously obtained and characterized¹³, were used.

92

93 **Cloning, expression and purification of Gly m Bd 28K/P28.** The cDNA encoding sequence
94 for P28 (GenBank: accession. [AB046874.2](#)) was obtained by PCR amplification of cDNA
95 library³². Amplified PCR products were cloned directionally into pENTR/D TOPO (Life
96 Technology, S.A. Argentina), and then transferred to pDEST-maltose-binding protein (MBP)
97 destination vector for expression³³. *E. coli* BL21 Codon Plus containing the constructs: pDEST
98 His-MBP-P28 and pDEST His-MBP were induced and recombinant proteins were purified as
99 described previously¹⁶. Depletion of lipopolysaccharide was carried out with a Sepharose-

100 polymyxin B resin (Sigma-Aldrich, St. Louis, MO, USA). Endotoxin determination was
101 performed with Limulus amoebocyte chromogenic assay (LONZA, Buenos Aires, Argentina).
102 Protein concentration was determined by the bicinchoninic acid assay with bovine serum albumin
103 as a standard (Pierce, Rockford, IL, USA).

104

105 **Immunochemical assays.** *i. Western blot.* Three μg of purified P28 and MBP were separated by
106 SDS-PAGE, and transferred to nitrocellulose membranes. Blocked membranes (3% horse serum
107 in phosphate buffer pH 7.4) were incubated with patient serum (1:5, overnight at 4 °C), followed
108 by biotinylated anti-human IgE monoclonal antibody (1:3000, 4 hs at 37 °C, Vector Laboratories
109 Inc, CA, USA), and finally, horseradish peroxidase (HRP) -streptavidin conjugate (1:3000, 30
110 min at 37 °C, Sigma-Aldrich, MO, USA). Membranes were exposed to Luminol
111 chemiluminescent substrate, revealed with X-ray film (Amersham Hyperfilm ECL, GE
112 Healthcare Bio-Sciences Corp., USA) and scanned. Additionally, membranes were also revealed
113 using the casein-specific mAbs (1 $\mu\text{g}/\text{ml}$) as primary antibody, followed by rat monoclonal anti-
114 mouse immunoglobulin G antibodies conjugated with HRP (1:3000, 1 h at 37°C Bio-Rad
115 Laboratories, CA, USA).

116 *ii. Flow cytometry-based modified basophil activation test (mBAT).* Basophils were obtained
117 from whole heparinized blood collected from non-allergic human donors using Ammonium-
118 Chloride-Potassium (ACK) buffer pH 7.2 (0.15M NH_4Cl , 10 mM KHCO_3 , 0.1 mM Na_2EDTA)
119 for erythrocyte lysis. Cells were re-suspended in ice-cold lactic acid buffer pH 3.9 (13.4 mM
120 lactic acid, 140 mM NaCl and 5 mM KCl) during 5 min for stripping. Cells were washed and re-
121 suspended in 2 ml of saline buffer containing 20 % of serum of milk allergic patients, 4 mM
122 EDTA and 10 mg/ml heparin (90 min at 37°C). Cells were washed with 20 nM HEPES buffer

123 pH 7.4, and incubated with HEPES buffer containing 1 mM CaCl₂ (30 min at 37 °C)³⁴. Then,
124 cells were challenged as described previously¹⁹ with whole milk proteins, soy proteins or the
125 recombinant P28 soy allergen at different concentrations. Saline buffer was used to stop cell
126 activation. Finally, washed cells were incubated with 7-Aminoactinomycin (7-AAD) (BD
127 Pharmingen, USA), biotinylated anti-human IgE monoclonal antibody (Vector Laboratories Inc,
128 CA, USA) and streptavidin-Allophycocyanin (APC) (eBioscience, USA), or Phycoerythrin (PE)-
129 conjugated antibody specific to human CD63 (BD Pharmingen, USA), monoclonal antibodies
130 specific to human CD203c (Macs, Miltenyi Biotec, Germany) followed by goat anti-mouse IgG
131 fluorescein isothiocyanate (FITC) (Santa Cruz Biotechnology, USA). Fluorescence data were
132 acquired with a Becton Dickinson FACSCalibur flow cytometer (Franklin Lakes, NJ, USA), and
133 analyzed with the BD CellQuest Pro Software and FlowJo software (Tree Star Inc, Ashland, OR,
134 USA). Controls with a non-related protein (OVA) or with sera from non allergic subjects were
135 included.

136 *ii. Competitive ELISA.* Polystyrene microtitre plates were coated with 0,25 µg/well of α-casein or
137 β-casein (Sigma-Aldrich, MO, USA) and blocked with 5% horse serum. Purified 1D5 or 4C3
138 mAbs were incubated with different quantities of soluble inhibitors: α-casein, β-casein, P28 or
139 ovalbumin (OVA) as a non-related protein (2 hs at 37°C). Then, this pre-mixed dilution was
140 added to the coated wells and incubated for 30 min at 37°C. Finally, rat monoclonal anti-mouse
141 immunoglobulin G antibodies conjugated with HRP was added (1:3000, 1 h at 37°C) and color
142 was developed with o-phenylenediamine. Optical density (OD) was measured at 492 nm.

143

144 **Experimental mouse model of food allergy and immune response to antigens.** *i. Mice*
145 *sensitization and mucosal challenges.* Male 6- to 8-week old Balb/c mice were purchased from

146 the School of Animal Sciences, University of La Plata, and kept under pathogen-free conditions
147 with water and commercial diet provided *ad libitum*. Mice were grouped into sensitization and
148 control group (n=8 per group), and sensitized as described previously¹⁵. Briefly, mice received 6
149 weekly intragastric (*ig*) doses of CMP (20 mg/dose) plus cholera toxin (10 µg/dose) (Sigma
150 Aldrich, St Louis, MO, USA) in a final volume of 200 ml of bicarbonate buffer (sensitization
151 group), or CMP (20 mg/dose) without cholera toxin (control group). Mice were fasted for 2 h
152 before sensitization, and 3% sodium bicarbonate solution was given 30 min before the
153 immunization. Ten days after the final boost mice were *ig* challenged with 20 mg of CMP, 5 mg
154 of SP. Additionally, some animals were sublingual (*sl*) challenged with 5 µg of P28, 5 µg of β-
155 lactoglobulin (β-Lg) or 10 µg of OVA (Sigma-Aldrich, St Louis, MO, USA).

156 *ii. In vitro evaluation of the allergic reaction.* Serum specific IgE antibodies were assessed by
157 western blot. Three µg of CMP, SP, purified P28 and MBP were separated by SDS-PAGE, and
158 transferred to nitrocellulose membranes. Blocked membranes (3% horse serum) were incubated
159 with sensitized mouse serum (1:2, overnight at 4 °C), followed by biotinylated anti-mouse IgE
160 monoclonal antibody (1:500, 3 hs at 37 °C, BD Pharmingen, USA), and finally, with horseradish
161 peroxidase (HRP) -streptavidin conjugate (1:1000, 1 hs at 37 °C, Sigma-Aldrich, MO, USA).
162 Membranes were incubated with luminol chemiluminescent substrate and exposed to X-ray film
163 (Amersham Hyperfilm ECL, GE Healthcare Bio-Sciences Corp., USA). In addition, serum
164 specific IgG1 and IgG2a were measured (ELISA) using CMP, SP, P28 or MBP as described
165 previously¹⁵. Briefly, microtitre plates were coated with CMP, SP, P28 1 µg/100 µl, or MBP 0,5
166 µg/100 µl in carbonate/bicarbonate buffer, pH 9.6. The assay was developed as indicated¹⁵.

167 To study T cell activation, 24 h following the oral challenge mice were killed, spleens were
168 resected, and spleen cells were stimulated for 72 h with CMP (350 µg/ml), SP (200 µg/ml), P28

169 (15 $\mu\text{g/ml}$) or MBP (7,5 $\mu\text{g/ml}$). Concentration of IL-5, IFN- γ (Invitrogen Corporation, USA)
170 and IL-13 (R&D Systems, UK) was assayed in the supernatants by ELISA following the
171 manufacturer's instructions.

172 *iii. In vivo evaluation of the allergic reaction.* Clinical symptoms were observed 30-60 min
173 following the oral challenge in a blinded fashion by 2 independent investigators, and scored
174 according to Table 1. Mice also underwent cutaneous tests: mice were injected into the pad of
175 either rear foot with 20 μg of CMP, 10 μg of P28, or 10 μg of MBP in 20 μl of sterile saline, and
176 saline in the contra-lateral footpad, as a negative control. Mice were also injected intravenously
177 (tail vein) with 100 μl of 0.1% Evans blue dye (Anedra, 19 Buenos Aires, Argentina). The local
178 presence of blue color minutes after the injection of proteins was considered a positive skin test,
179 and footpad swelling was measured with a digital micrometer with a minimum increment of 0.01
180 mm.

181
182 **Bioinformatics analysis. i. Sequence Alignment.** The nucleotide sequences of Gly m Bd 28K
183 [Glycine max] (Sequence ID: gi: 410067729 Length: 476 aas), α s1-casein [Bos Taurus]
184 (Sequence ID: gi:162792 Length: 199 aas), β -casein [Bos Taurus] (Sequence ID: gi: 162931
185 Length: 210 aas) and κ -casein [Bos taurus] (Sequence ID: gi: 1228078 Length: 169 aas) were
186 compared using BLAST (Basic Local Alignment Search Tool,
187 <http://blast.ncbi.nlm.nih.gov/Blast.cgi>)³⁵ to identify putative cross- reactive epitopes.

188 *ii. Property Distance index.* to quantify the levels of similarity between P28 putative cross-
189 reactive epitopes and caseins the Property Distance (PD) value was calculated using Property-
190 Based Peptide Similarity Index PD for Two Sequences tool
191 (http://fermi.utmb.edu/SDAP/sdap_pdi.html)³⁶.

192 *iii. Peptide similarity of P28 with known allergen:* The P28 putative cross-reactive epitopes
193 were assessed for similarity with known allergens in the Structural Database of Allergenic
194 Proteins (SDAP, <http://fermi.utmb.edu/SDAP/>) using the Peptide similarity tool
195 (http://fermi.utmb.edu/SDAP/sdap_pps.html)³⁶.

196 *iv. Prediction and analysis of the secondary structure:* P28 secondary structure was predicted
197 with PSIPRED (<http://bioinf.cs.ucl.ac.uk/psipred/>)³⁷ server.

198

199 **Ethical Consideration.** All experimental protocols of this study were conducted in strict
200 agreement with international ethical standards for animal experimentation (Helsinki Declaration
201 and its amendments, Amsterdam Protocol of welfare and animal protection and National
202 Institutes of Health, USA NIH, guidelines: Guide for the Care and Use of Laboratory Animals)
203 and were approved by the local Institutional Animal Care and Use Committee at the School of
204 Animal Science (CICUAL, 700-003068 /14, University of La Plata). All efforts were made to
205 alleviate suffering during the whole experiment.

206 For the human sample analysis a written informed consent was obtained from parents of patients
207 and the project was approved by the Ethics Committee of the Argentinean Association of Allergy
208 and Clinical Immunology (#00589, February 2012).

209

210 **Statistical analysis.** The values are expressed as means \pm standard error of the mean (SEM). All
211 statistical analyses were carried out using GraphPad Prism 5 software. The significance of the
212 difference was determined using an independent-sample t-test or ANOVA test. Statistically
213 significant differences were defined as $p < 0.05$.

214

215 **Results**

216 **Production of recombinant P28 in *Escherichia coli*.** The 28kDa soybean protein is considered
217 a major soybean allergen based on its interactions with IgE sera of soy sensitive patients
218 ^{20,22,23,25,38}. To further analyze its involvement in hypersensitivity reactions, the cDNA encoding
219 for the 52,9 kDa proprotein was cloned into pENTR/D-TOPO (Invitrogen) and transferred to the
220 destination vector pDEST-HisMBP to fuse in frame with the sequences encoding a hexa-
221 Histidine tag (His) and Maltose Binding Protein (MBP) at the N-terminal (Figure 1a). The
222 obtained pDEST-HisMBP-P28 expressions vector (Figure 1b) was introduced into *E.coli* BL21
223 Codon Plus to produce the recombinant fusion protein MBP-P28 (Figure 1c). As a control, a
224 pDEST-His-MBP, without the toxic *ccdB* gene, was also obtained and used to produce MBP.
225 Both MBP-P28 and MBP fusion proteins were purified from the insoluble fraction under
226 denaturing conditions by nickel affinity chromatography and purified proteins were analyzed by
227 SDS-PAGE (Figure 1d). A 97kDa and 44kDa bands corresponding to the fusion MBP-P28 and
228 MBP alone, respectively, were detected. These purified proteins were used in further analysis.

229
230 ***In vitro* recognition of P28 by immunochemical assays with CMP-specific IgE. i.**
231 ***Immunoblotting with sera of CMA patients.*** Human sera containing CMP-specific IgE antibodies
232 were used to assess the recognition of the recombinant P28. Figure 2a shows that P28 was
233 recognized as a coated antigen by ten IgE sera of milk allergic patients. No binding was detected
234 for MBP with sera of allergic patients, and for P28 with sera of non allergic subjects and non-
235 milk allergic patients (#11 and 12).

236 ***ii. mBAT with sera of CMA patients.*** To confirm the cross-reactivity a basophile activation test
237 (BAT) was performed with CMP-specific human IgE sera. Stripped basophils from healthy

238 donors were passively sensitized with individual serum containing IgE and then challenged with
239 different concentrations of milk or soy proteins. A minimum of 100000 events per sample were
240 analyzed by flow cytometry and cells were gated based on physical properties (Figure 2biI), 7
241 AAD⁻ live cells (Figure 2biII), IgE⁺ cells (Figure 2biIII) and further analyzed for membrane
242 CD63 and CD203c (Figure 2biI). Isotype controls were used to set the fluorescence thresholds.
243 Percentages of double positive cells were compared between basophils incubated with the
244 allergen or PBS. Figure 2biII showed that CMP and SP rendered a higher percentage of IgE⁺
245 CD63⁺ CD203c⁺ cells compared with PBS (18.45±1.35% and 18.80±0.5% vs 9.16±1.26%,
246 respectively). When P28 was used for the IgE-dependent basophil activation we found
247 15.18±0.26% of double positive cells. The non-related protein OVA rendered 8.37±0.31% of
248 double-positive cells at different concentrations. In addition, there were no statistically
249 significant differences in cell activation using sera from non-allergic patients (data not shown).

250

251 *iii. Immunoblotting and competitive ELISA with the casein-specific monoclonal antibodies.* To
252 rule out co-sensitization in patients, immunoblotting was performed using three casein-specific
253 monoclonal antibodies, and we found that only for anti β -bovine casein (BC) mAbs revealed a
254 weak band corresponding to P28 (Figure 3a). MBP was not recognized. Then to rule out the
255 possibility of an artifactual reactivity due to new epitopes created during the coating of antigens,
256 P28 was used as soluble inhibitor in a competitive ELISA using the β -casein-specific 4C3 mAb
257 (Figure 3b) and α -casein-specific 1D5 mAb (Figure 3c). The sigmoid-shape of the dose-response
258 inhibition curve obtained with P28 as inhibitor demonstrated the specificity of the antigen-
259 antibody reaction. The β -casein-specific 4C3 mAb to the immobilized β -casein (0,25 μ g/well)
260 was almost 100 % inhibited with 0,2 mg/ml of soluble β -casein, while 50% of inhibition (IC50)

261 was achieved with 0.06 mg/ml of β -casein, and 0.024 mg/ml of P28. In a separate assay antibody
262 binding to the immobilized α -casein (0,25 μ g/well) was 100 % inhibited with 0,002 mg/ml of
263 soluble α -casein, while 50% of inhibition (IC50) was achieved with 0.0002 mg/ml of α -casein,
264 however no inhibition was observed with α -casein-specific 1D5 mAb a wide range of
265 concentration of P28. No inhibition was observed a wide range of concentration of OVA for both
266 mAbs.

267
268 ***In vivo* evaluation of P28 used a specific mouse model of CMP.** Mice exclusively allergic to
269 milk were used to investigate the clinical relevance of the immunochemical co-recognition. The
270 experimental protocol used to sensitized mice is depicted in Figure 4a. Intragastrically sensitized
271 mice developed hypersensitivity symptoms, scored according to Supporting Table S1,
272 immediately after the oral challenge with the different allergens. CMP-sensitized animals
273 showed higher scores upon the challenge with SP or CMP, compared to sham mice (Figure 4bi).
274 When animals were sublingually challenged with individual proteins (β -lactoglobulin, P28 or the
275 unrelated OVA) we observed high clinical scores with β -lactoglobulin, and intermediates scores
276 with P28, while OVA was rendered no symptom (Figure 4bii). Finally, we examined the
277 functionality of IgE-sensitized mast cells and basophils in the footpads of mice. The cutaneous
278 injection of CMP or P28 led to a local and distant inflammation in minutes (Figure 4c). Footpad
279 swelling was statistically lower when injected P28 compared CMP mice (0.27 ± 0.05 and
280 0.49 ± 0.06 , respectively; and saline buffer 0.05 ± 0.02 as control).

281
282 **Humoral and cellular immune response to P28 fusion protein by sensitized mice of CMP.**
283 Afterwards, we questioned if the soy proteins are recognized by the secreted T-helper 2 (Th2) -

284 associated antibodies. We found that CMP and SP were highly and significantly recognized by
285 mouse CMP-specific IgE and IgG1 antibodies. However, P28 was significantly recognized by
286 CMP-specific IgG1 but weakly recognized by IgE antibodies (Figure 5ai and ii). No IgG2a
287 specific antibodies were detected in sensitized or sham animals (Figure 5aiii). Also, to correlate
288 humoral immune response with the cellular immunity we measured cytokine production in
289 culture supernatants of splenocytes after being stimulated with different proteins. We found a
290 significant increase of the secreted Th2 cell-associated cytokines (IL-5 and IL-13) in sensitized
291 mice challenged with CMP or SP, while IFN- γ remained unchanged (Figure 5b). When the
292 detoxified P28 protein was used to stimulate spleen cells, only IL-13 were secreted, while the
293 addition of MBP induced no cell stimulation.

294 To further investigate immune response of milk sensitized mice a western blot was performed.
295 Mouse Ig bound to CMP and SP (Figure 5ai). Interestingly, several components of the SP extract
296 (i.e. α' - α - and β - β conglycinin, AB glicinin, 40kDa, P28) were specifically recognized by sera
297 of mice exclusively sensitized to milk. Recombinant P28 was also recognized by IgE-containing
298 serum antibodies.

299
300 ***In silico* analysis of cross-reactive epitopes.** In order to understand the observed *in vitro* and *in*
301 *vivo* cross-reactivity, P28 and α S1, β - and κ caseins sequence alignments were performed. Two
302 small regions with discrete homology were detected when α S1 protein sequence was aligned
303 with P28 (Table S1). The first homology region of α S1-casein with P28 had a length of 19
304 amino acids with 37% of identity and 57% of similarity, while the second homology region had
305 only of 6 amino acids length, without gaps, with 83% of identity and similarity. Alignment of β -
306 casein with P28 gave four homology regions of 8, 15, 17 and 23 amino acids length without

307 gaps, with identity in the range of 26% to 50% and similarity between 47-87% (Table 1). Only
308 one discrete homology region of 12 amino acids was found in the alignment of κ -casein and P28
309 with 42% of identity and 75% of similarity. These putative cross-reactive epitopes were
310 numbered according to their location along P28 sequence. The peptides 1 and 2 overlap in a
311 sequence of 11 amino acids residues. Predictive cross-reactive epitopes were located in beta
312 sheet and α helix structure as well disordered regions (Figure 6). To evaluate meaningful
313 physical-chemical similarities, the Property Distance (PD) index between the bovine caseins and
314 P28 peptides was calculated using Property-Based Peptide Similarity Index PD for Two
315 Sequences (http://fermi.utmb.edu/SDAP/sdap_pdi.html)³⁶. Peptide 5 and 6 has a PD value of
316 5.17 and 4.07, respectively while the other P28 peptides have PD values in the range of 9.06-
317 12.19 (Table 1). Peptides with PD the value lower than 10 share high similarity³⁶, therefore P28
318 have several peptides with the potential cross-reactivity with caseins.

319 The seven putative cross-reactive epitopes were assessed for similarity with known allergens in
320 the Structural Database of Allergenic Proteins (SDAP) using peptide similarity tool³⁶. Cross
321 reactivity with Bos d 8 allergen, which include α S1, β and κ caseins, was detected for all
322 putative P28 cross-reactive epitopes except for peptide 2 and 4 (Supporting Table S2). In
323 addition, P28 peptides showed significant similarity with allergens of peanut, sesame, soybean,
324 carrot, pear, pistachio, etc, since the Property Distance (PD) score were lower than 10
325 (Supporting Table S2). Remarkably other IgE cross-reactive soybean allergens with bovine
326 caseins described by our group^{15,16,17,18}, were retrieved by the SDAP peptide similarity tool. :
327 Peptides 5 and 3 had PD value of 4.08, and 7.15 with alpha subunit beta conglycinin (Gly m
328 5.0101)^{16,18}, respectively. Peptides 5 had a PD value of 4.83 with glycinin G4 A4A5B3 (Gly m
329 6.0401)^{15,17}(Supporting Table S2).

330 Discussion

331 Gly m Bd 28K (P28) is a minor component of the 7S globulins^{20,21,25} and it has been described,
332 together with P34 and α subunit of β -conglycinin, as the third major soybean allergen^{24,31,38,39}.

333 We have shown that P28 was recognized by serum IgE from milk-allergic patients, CMP-
334 specific IgE and IgG1 antibodies from milk-allergic mice and activated basophils passively
335 sensitized with milk-allergic sera. Additionally, we showed that P28 was a good competitor of β -
336 casein in an inhibition ELISA performed with a specific β -casein mAb in concordance with the
337 recognition of P28 by this mAb in western blot. Furthermore, an immediate cutaneous reaction
338 and hypersensitivity symptoms were observed in sensitized mice following oral exposure to P28,
339 consistent with the Th2-biased immune response. These *in vivo* results are in good correlation
340 with the *in vitro* cross-reactivity.

341 In previous studies, we demonstrated that bovine caseins and soy proteins: Gly m 6 G4 (glycinin
342 A₄A₅B₃)^{15,17}, Gly m 5.0101(α subunit of β -conglycinin)^{16,18} and Gly m Bd 30K (P34)¹⁹, were
343 recognized by IgE from milk allergic patients and they are able to trigger hypersensitive reaction
344 in a mouse model to cow's milk allergy. These *in vivo* results partially explaining the clinical
345 intolerance observed in a restricted population of IgE-mediated CMA patients, primarily not
346 sensitized to SP³. However, and to rule out the possibility of co-sensitization of patients, casein-
347 specific monoclonal antibodies were used to identify cross-reactive SP. Epitopes in P34 and α
348 subunit of β -conglycinin were bound by α -, β - and κ -casein specific monoclonal antibodies^{18,19},
349 while P28 only reacted when confronted the β -casein specific mAb (4C3). Although α -casein
350 1D5 and κ -casein 3B5 monoclonal antibodies did not recognized P28, we cannot discard that
351 P28 share epitopes with α - or κ -casein than were not recognized by these mAbs. This distinctive
352 immunochemically cross-reactivity detected for P28, was also observed *in vivo* through IgE

353 mediated response of milk sensitized mice. CMA mice after sublingual challenge with P28,
354 showed lower clinical hypersensitivity score compared with those confronted with P34 or α
355 subunit of β -conglycinin^{18,19}. Besides, P28 skin test was also weakly positive compared with
356 CMP, P34 or α subunit of β -conglycinin that were strongly positive^{18,19}. In addition, stimulation
357 of sensitized mice spleen cells with P28 only induced IL-13 release, while the other cross-
358 reactive soybean epitopes triggered both IL-5 and IL-13 discharge. Consequently, P28 T cell
359 epitopes associated with Th2 responses are different from the ones present in bovine caseins,
360 P34 and α subunit of β -conglycinin^{18,19}.

361 Based on our findings we cannot assure that cross-reactivity between P28 and caseins is based
362 exclusively on sequential epitopes⁴⁰. We know that the allergenicity of caseins is due to some
363 small protein fragments containing IgE-binding epitopes, which are spread along the whole
364 protein sequence, able to cross-link two IgE molecules, and thereby elicit an allergic reaction. In
365 general, caseins appear to be the predominant allergen in patients with CMA⁴¹. The majority of
366 milk allergenic patients showed a strong humoral and cellular response to caseins, with high
367 titers of IgE specific for these proteins. Multiple B and T epitopes in caseins, ranging from 8 to
368 20 AA, have been identified by different immunoassays⁴²⁻⁵⁰. In agreement with this, when we
369 aligned the sequences of α S1-, β - and κ - caseins with P28, seven peptides with discrete
370 homology were distinguished (Table 1 and Figures 6). These peptides showed a length between
371 6 to 23 amino acids, degrees of identity from 26% to 83% and PD values from 4.07 to 12.19.
372 Although, the threshold between cross-reactive and non-reactive peptides might vary among
373 epitopes, PD values lower than 10 point out significant physical-chemical similarities³⁶. Peptides
374 1 and 4-7 had PD values lower than 10, therefore are potentially cross-reactive epitopes with
375 caseins. Thus, the *in silico* sequence analysis reveals that P28 had at least five cross-reactive

376 epitopes with bovine caseins, and only two are required for binding of IgE molecules and trigger
377 hypersensitivity symptoms. These results contribute to validate the utility of the property
378 distance (PD) scale for prediction of cross-reactivity.

379 Our group has described several soybean cross reactive allergens with caseins: P28 and α subunit
380 of β -conglycinin (Gly m 5.01)^{16,18} that are 7S storage proteins and glycinin A₄A₅B₃ (Gly m 6
381 G4)^{15,17} is an 11S storage proteins, P34 (Gly m Bd 30 K) a inactive cysteine protease of the
382 papain-superfamily¹⁹. In this work, we showed that the putative cross-reactive P28 peptides with
383 caseins, also share significant similarity with other member of the 7S and 11S soybean storage
384 protein as well as cupins from legumes and legumes and tree nuts (Supporting Table S2).
385 Although, it was thought that proteins with less of 50% sequence identity were rarely cross-
386 reactive, recently it has been shown Ara h 1, Ara h 2, and Ara h 3 the three major allergen of
387 peanut are cross-reactive⁵¹. This report is in coincidence with our results, that support cross-
388 reactivity among non structural related proteins such as bovine caseins, soybean 7S and 11S
389 proteins. It is important to remark that at difference of other reports that only analyzed in vitro Ig
390 E recognition; we have demonstrated in vivo allergenicity among these proteins.

391 Experimental animal models are important to confirm the immunochemical data and to
392 investigate the cross-allergenicity. Immunochemical cross-reactivity not always correlates with
393 cross-allergenicity. In this work we probed that P28 triggered hypersensitivity symptoms in
394 exclusively milk-sensitized mice, which was not observed with the un-related OVA antigen.
395 Furthermore, the functionality of the IgE antibodies was also evidenced with the skin test,
396 demonstrating that P28 contains at least two surface cross-reactive epitopes. This *in vivo*
397 characterization that cannot be carried out in patients is critical to generate knowledge to be
398 incorporated into the allergen databases.

399 In conclusion, P28 was recognized by IgE sera from CMA patients, activated human peripheral
400 basophils degranulation, produced a weakly positive skin test and elicited immediate
401 hypersensitivity symptoms in milk-sensitized mice. These in vitro and in vivo data confirmed
402 that P28 contains B and T cross-reactive epitopes with bovine caseins that were proposed using
403 different bioinformatics tools. Consequently, we demonstrated that P28 is a new cross-reactive
404 soybean allergen with bovine caseins. In vivo studies of cross-reactivity of allergens are scarcely
405 and this work is an important contribution for understanding physical-chemical characteristics of
406 allergens and for developing hypoallergenic soy formulations or news therapeutic intervention
407 protocols.

408

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418

419

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570

571

572

573 **Figure legends**

574 **Figure 1. Construct and synthesis of recombinant *Gly m Bd 28K*/P28 protein.**

575 (a) pDEST-HisMBP expression vector. The vector contains the *cm* and toxic *ccd B* genes
576 between the recombination attR1 and attR2 sites.

577 (b) Schematic representation of MBP-P28 and MBP constructs. The hexahistidine (His) and
578 maltose binding protein (MBP) tags were fused to P28 proprotein (476 AA) in the N terminal to
579 produce MBP-P28. The vector pHisMBP is a modified version without the *cm* and toxic *ccd B*
580 genes, that was obtained to produce MBP protein.

581 (c) SDS-PAGE of total extracts: lane 1, *E. coli* wild type; lane 2, *E. coli* expressing MBP; lane
582 3: *E. coli* expressing MBP-P28.

583 (d) SDS-PAGE of the purified proteins: lane 1, MBP; lane 2: MBP-P28.

584 *MW*, Molecular weight ranges; *MBP*, maltose-binding protein; *His*, hexahistidine tag; *cm*,
585 chloramphenicol resistance gen; *ccd*, toxin gene that targets DNA gyrase; *P28*, *Gly m Bd 28K*;
586 *MW of MBP*: 44 kDa; *MW of MBP-P28*: 97 kDa.

587

588 **Figure 2. Immunochemical assays of soy allergen protein with sera of CMA patients.**

589 (a) ***P28 recognition by Ig E sera of CMA patients***: P28 recombinant protein was separated by
590 SDS-PAGE and membrane was revealed with individual sera of CMA patients (#1-10). In
591 addition, incubations with a representative serum of a non-CMA patient (#11) and of a non
592 allergic subject (#12) were used as negative controls. To discard recognition of IgE sera to the
593 His-MBP region of MBP-P28 fusion, the binding HisMBP protein was also analyzed. Ponceau S
594 staining of the blotted membrane was performed to check the efficiency of recombinant proteins
595 transfer.

596 (b) *Basophil activation test by flow cytometry*. Basophils from healthy donor subjects were
597 stripped and passively sensitized with milk-specific IgE-containing sera from CMA patients
598 (n=5). i. Cells were selected according to physical parameters I, viable cells; II, viable IgE-
599 sensitized cells; III, Activated basophils: CD63⁺ / CD203c⁺ double positive cells.
600 ii. Activated basophils after stimulation with CMP, SP, and P28 at different concentrations were
601 analyzed. OVA and PBS were used as negative controls. Results of representative experiments
602 are showed.
603 iii. Percentage of activated basophils obtained with five CMA's IgE-sera. Data represent mean ±
604 SEM % of double positive basophils. Statistical significant difference with t-test: ***p<0.0005,
605 **p<0.001 vs PBS group.

606

607 **Figure 3. Immunochemical assays with casein specific mAbs.**

608 (a) Western blot of P28 and MBP revealed with different casein specific monoclonal antibodies.
609 Ponceau S staining showed that proteins were efficiently transferred to the membrane.

610 (b) Inhibition ELISA: wells were coated with β-casein (0,25 μg/well), and β-casein-specific
611 monoclonal antibody 4C3 was pre-incubated with different concentrations of β-casein, P28 or
612 OVA as soluble inhibitors.

613 (c) Inhibition ELISA: wells were coated with α-casein (0,25 μg/well), and α-casein-specific
614 monoclonal antibody 1D5 was pre-incubated with different concentrations of α-casein, P28 or
615 OVA as soluble inhibitors.

616 *MBP, maltose-binding protein; P28, MBP-P28 recombinant fusion protein; 1D5, α-casein-*
617 *specific monoclonal antibody; 4C3, β-casein-specific monoclonal antibody; 3B5, κ-casein*
618 *specific monoclonal antibody. MW of MBP: 44 kDa; MW of MBP-P28: 97 kDa.*

619

620 Figure 4. Allergic sensitization of mice to CMP and clinical symptoms .

621 (a) Schematic drawing of the experimental protocol: BALB/c mice (n=8 per group) were
622 subjected to weekly intragastric sensitization with cholera toxin and CMP from day 0 through
623 day 35. Challenge was performed at day 45 by intragastric (CMP or SP) (n=7 per group) or
624 sublingual (β -Lg, P28 or OVA) (n=5 per group) administration of proteins. Control mice only
625 received CMP and then they were orally challenged;

626 (b) Clinical symptoms were observed 30 min following the challenges with CMP or SP (i.) or
627 with β -Lg, P28 or OVA (ii.) and then scored according to Table S1;

628 (c) Cutaneous test: sensitized control mice were injected into the pad of either rear foot with
629 CMP, P28, MBP and PBS (n=4 per group).

630 i. Blue colour in the skin within minutes after injection, was considered a positive cutaneous test.
631 The results are from a single experiment, which is representative of three independent
632 experiments.

633 ii. Footpad swelling: the difference between the thickness of the footpad injected with allergens
634 and negative control was calculated. Data are expressed as mean \pm SEM. Statistical significant
635 difference with ANOVA test: ***p<0.005, **p<0.01.

636 *CMP, cow's milk proteins; SP, soy proteins; CT, cholera toxin; β -Lg, beta lactoglobulin; MBP,*
637 *maltose-binding protein, P28, MBP-P28 recombinant fusion protein; OVA, ovalbumin.*

638

639 **Figure 5. Humoral and cellular immune response of milk sensitized mice.** (a) i. Specific IgE
640 (by WB): ii. Specific IgG1 (by ELISA) and iii. Specific IgG2a (by ELISA) in serum of milk-
641 sensitized mice (mean values \pm SEM);

642 (b) i. Levels of IL-5, ii. IL-13 and iii. IFN- γ (by ELISA) in supernatants of stimulated spleen
643 cells (mean values \pm SEM). Results correspond to a single experiment with at least three mice
644 per condition; three independent experiments gave similar results. Statistically significant
645 differences with Student's *t* test: *** $p < 0.005$, ** $p < 0.01$, * $p < 0.05$.

646 *SP*, soy proteins; *CS*, Coomassie stained protein gel; *MBP*, maltose-binding protein, *P28*, MBP-
647 *P28 recombinant fusion protein*; *CMP*, cow's milk proteins, *AB Gly*: glycinin subunits, α' -, α -
648 and β - β conglycinin subunits.

649

650 **Figure 6. *In silico* analysis of putative cross-reactive epitopes on P28.**

651 P28 protein sequence was alignment with bovine caseins. Regions of similarity to α S1-casein
652 (peptides 3 and 6) are shown with dashed lines, to β -casein (peptides 2, 4, 5, and 7) with solid
653 lines and to κ -casein (peptides 1) with dotted lines. Amino acid residue letter corresponds to its
654 degree of conservation: red for identical and purple for conserved. P28 secondary structure was
655 analyzed with PSIPRED prediction server (<http://bioinf.cs.ucl.ac.uk/psipred/>). The alpha-helix
656 regions are indicated with red cylinder, beta-sheets are presented in yellow arrows and
657 disordered regions with solid lines.

658

659 **Abbreviations:** **BAT**, basophil activation test; **β -Lg**, beta lactoglobulin; **CMA**, cow's milk
660 allergy; **CMP**, cow's milk proteins; **CR**, cross reactivity; **CT**, cholera toxin; **EAST**, Enzyme
661 Allergo Sorbent Test; **ELISA**, Enzyme-linked immunosorbent assay; **Gly m**, Glycine max (L.)
662 Merr. allergen; **HRP**, horseradish peroxidase; **IC50**, inhibitory concentration of 50%; **Ig**,
663 immunoglobulin; **IFN**, interferon; **IL**, interleukin; **IUIS**, International Union of Immunological
664 Societies; **i.v.**, intravenously; **i.g.**, intragástrica; **s.l.**, sublingual; **s.c.**, subcutaneous; **mAb**,
665 monoclonal antibody; **MBP**, maltose binding protein; **OD**, optical density; **OVA**, ovalbumin;
666 **PD**, property distance score; **P28**, 28 kDa soybean protein; **SDAP**, Structural Database of
667 Allergenic Proteins; **SDS-PAGE**, sodium dodecyl sulphate-polyacrylamide gel electrophoresis;
668 **SP**, soybean proteins; **Th**, T-helper cell; **WB**, Western Blot; **WHO**, World Health Organization.

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671

672 **Table 1: Identification of putative cross reactive epitopes by alignment of amino acid**
 673 **sequences**
 674 **of P28 and bovine milk caseins.**

Cross-Reactive Epitopes	Casein (Bos d 8)	Matches regions in the alignment P28 vs caseins	AA length	Identities	Positives	Gaps	PD Value*
Peptide 1	κ casein	P28 70 HMHIGFISMEPK 81 H H+ F+++ PK K-cas 121 HPHLSFMAIPPK 132	12	5/12 (42%)	9/12 (75%)	0/12 (0%)	9.06
Peptide 2	β casein	P28 71 MHIGFISMEPKSLFVPQ 87 MH + P +F PQ b-cas 159 MHQPHQPLPPTVMFPPQ 175	17	6/17 (35%)	8/17 (47%)	0/17 (0%)	12.01
Peptide 3	α S1 casein	P28 115 RRLKGTGDLYMIPSGSAFYL 133 R+ D Y PSG+ +Y+ a-cas 166 RQFYQLDAY--PSGAWYV 182	19	7/19 (37%)	11/19 (57%)	2/19 (10%)	12.19
Peptide 4	β casein	P28 225 KDDKEQQLKMMQDQ 239 + +++QQ + +QD+ b-cas 49 QSEEQQQTEDELQDK 63	15	4/15 (27%)	11/15 (73%)	0/15 (0%)	9.13
Peptide 5	β casein	P28 241 EDDEEKQT 248 + EE++QT b-cas 49 QSEEQQQT 56	8	4/8 (50%)	7/8 (87%)	0/8 (0%)	5.17
Peptide 6	α S1 casein	P28 309 EPDIGV 314 EP IGV a-cas 148 EPMIGV 153	6	5/6 (83%)	5/6 (83%)	0/6 (0%)	4.07
Peptide 7	β casein	P28 329 VNPISDEYTIIVLSGYGELHIGYP 351 V P ++ ++ L+ LH+ P b-cas 131 VEPFTESQSLTLTDVENLHLPLP 153	23	6/23 (26%)	12/23 (52%)	0/23 (0%)	9.79

675 The nucleotide sequences of Gly m Bd 28K [Glycine max] (Sequence ID: gi: 410067729 Length:
 676 476 aas), α s1-casein [Bos Taurus] (Sequence ID: gi:162792 Length: 199 aas), β -casein [Bos
 677 Taurus] (Sequence ID: gi: 162931 Length: 210 aas) and κ -casein [Bos taurus] (Sequence ID: gi:
 678 1228078 Length: 169 aas) were compared using BLAST (Basic Local Alignment Search Tool,
 679 <http://blast.ncbi.nlm.nih.gov/Blast.cgi>)³⁵ to identify putative cross-reactive epitopes. Seven
 680 regions with high similarity were detected and these peptides were considered putative cross-
 681 reactive epitopes. The alignment of these peptides with caseins is shown: the identical amino
 682 acid residues are in red color, and the conserved residues are consigned with plus purple symbol.
 683 Statistics for each peptide is shown on the right columns. *The Property distance (PD) index
 684 between the bovine casein peptides and P28 epitopes identified in the alignment was calculated
 685 using Property-Based Peptide Similarity Index PD for Two Sequences
 686 (http://fermi.utmb.edu/SDAP/sdap_pdi.html)³⁶. PD value detects meaningful physical-
 687 chemical similarities.
 688

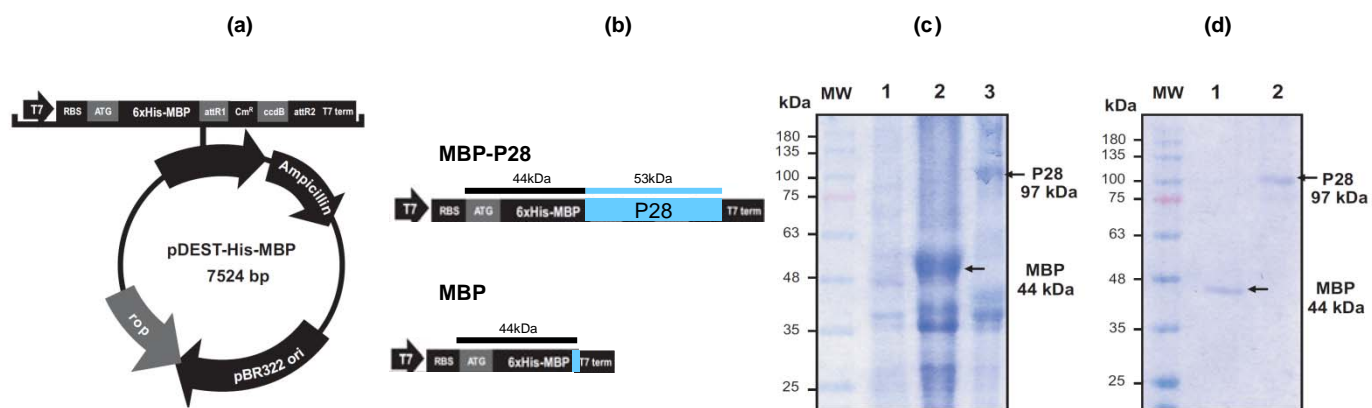
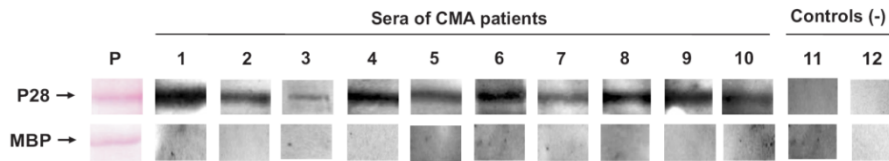


Figure 1

(a)



(b)

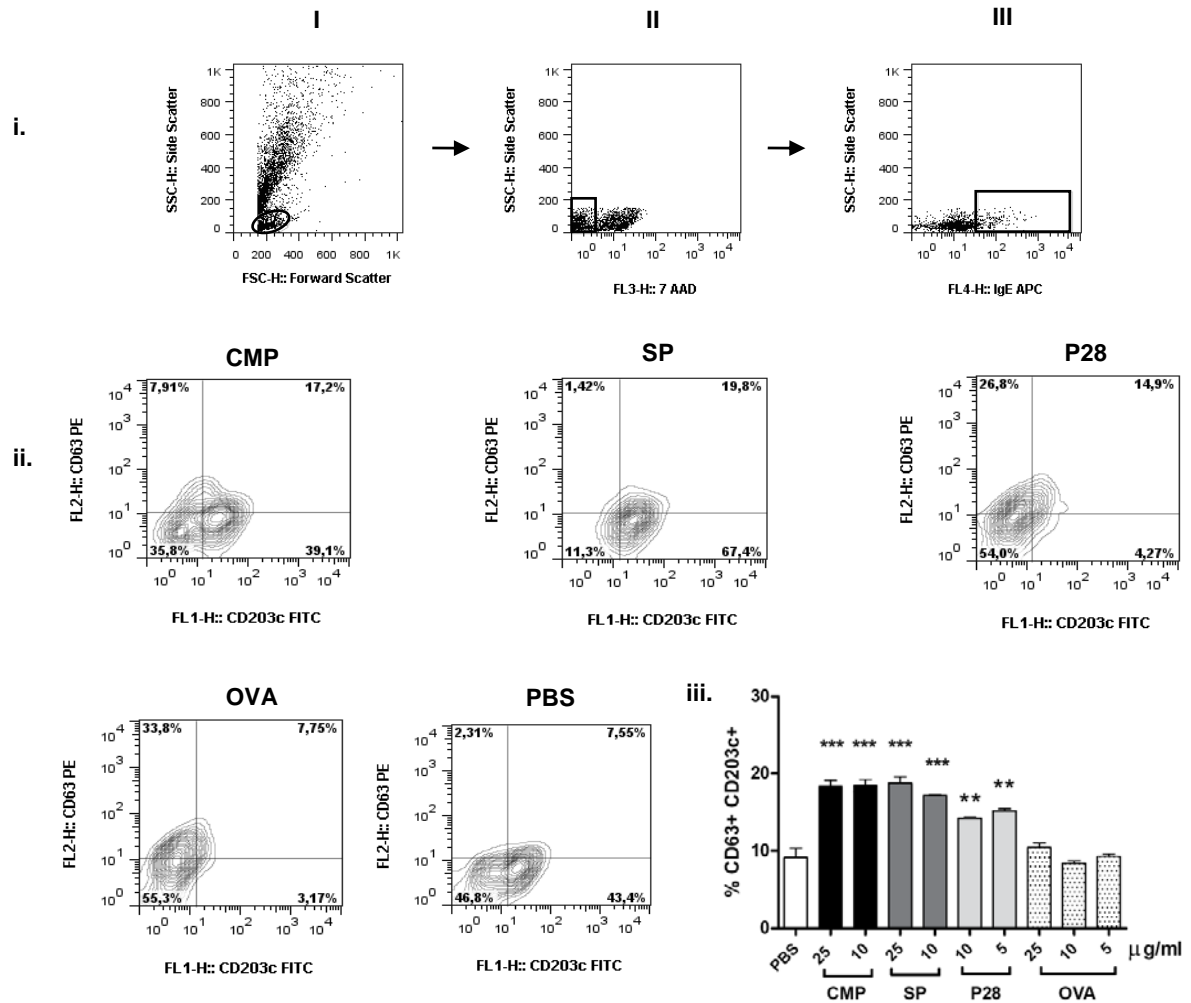
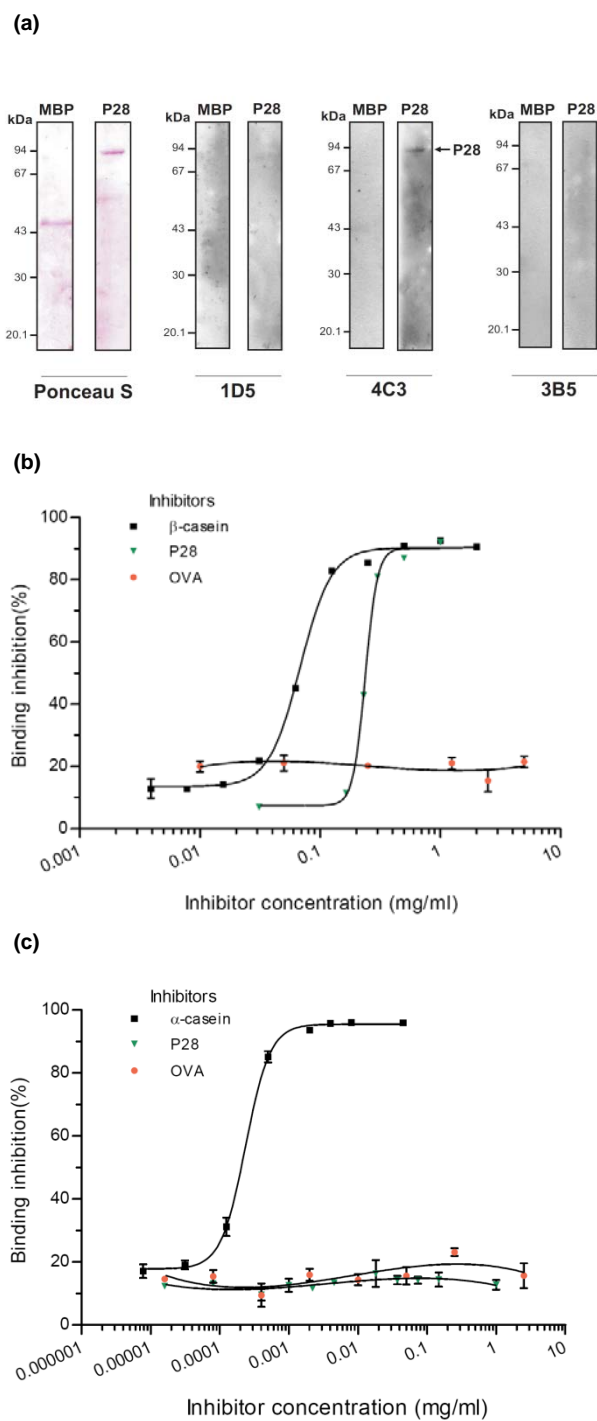
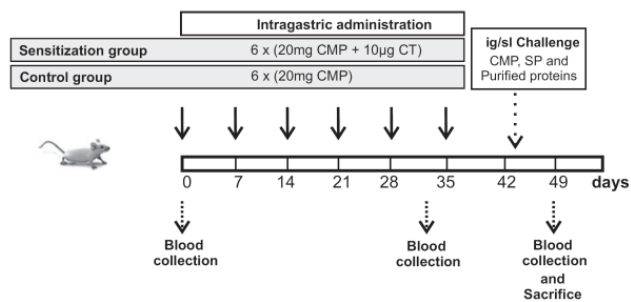


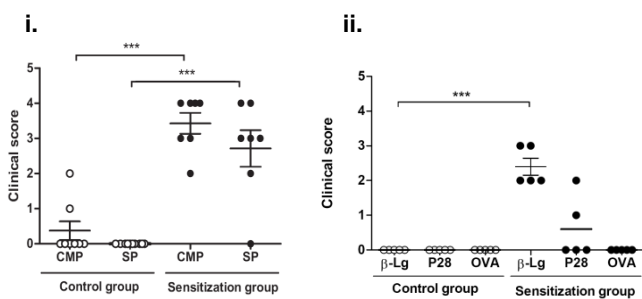
Figure 2

**Figure 3**

(a)



(b)



(c)

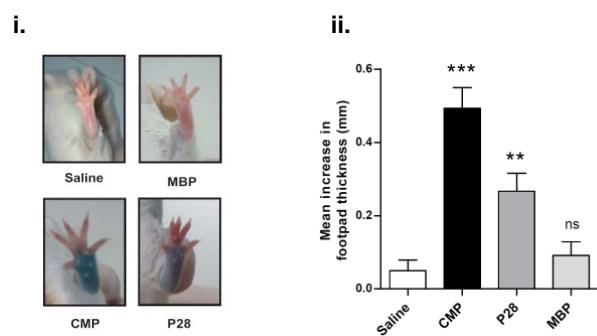


Figure 4

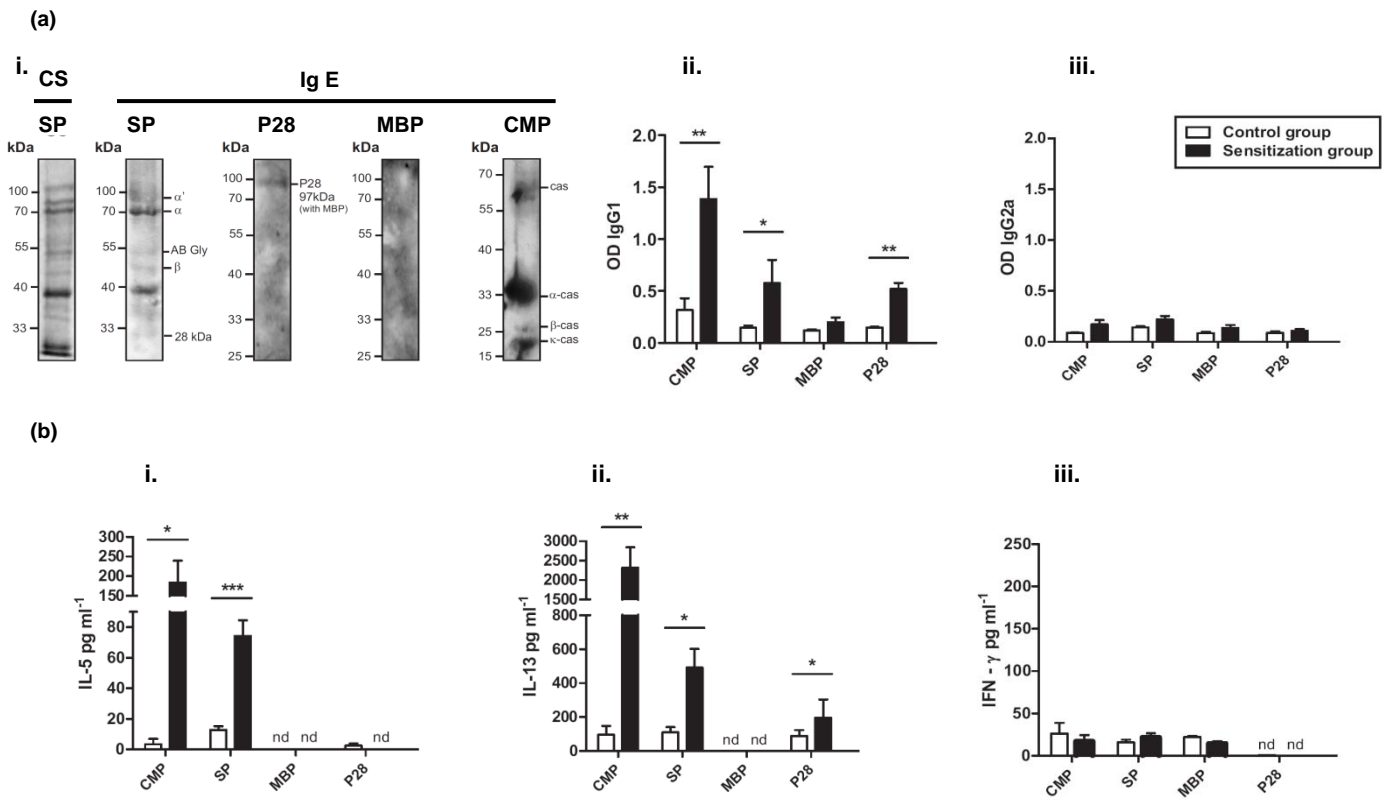
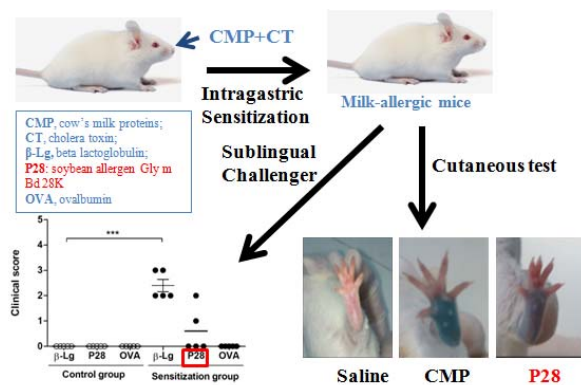


Figure 5



Figure 6



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