Quantification assay for the major allergen of *Cupressus sempervirens* pollen, Cup s 1, by sandwich ELISA

M.C. Arilla, I. Ibarrola, A. Martínez and J.A. Asturias


ABSTRACT

**Background:** The Cupressaceae are an important cause of pollinosis, particularly in Mediterranean countries. Cypress pollen allergenic extracts are difficult to produce since they have a low protein and a high carbohydrate content and consequently accurate standardization of these extracts is essential for diagnosis and immunotherapy.

**Method:** Natural Cup s 1 was purified by a combination of hydrophobic interaction, gel filtration and ion exchange chromatographies and its enzymatic activity was analyzed. The allergen was used as reference material in the ELISA standard curve. The assay was based on a specific monoclonal antibody (3D2) immobilized on ELISA plates and used to capture Cup s 1. Bound proteins were detected by a combination of biotinylated specific antiserum and peroxidase-conjugated streptavidin.

**Results:** Purified Cup s 1 is a functional pectate lyase enzyme with a specific activity of 750 U/mg protein. The developed ELISA measured Cup s 1 concentrations ranging from 31.25 to 250 ng/ml in the linear portion of the standard curve. The intra-assay and inter-assay variation coefficients in the working range were less than 8.1 % and 16 %, respectively. The assay was highly sensitive, with a detection limit of 3.8 ng/ml. The dose-response curves obtained with *C. sempervirens* pollen extracts and extracts belonging to other species from the Cupressaceae family showed a good parallelism compared with those obtained using the purified allergen, indicating that the same protein was measured.

**Conclusions:** The assay described is sensitive, specific and reproducible for the quantification of Cup s 1 in *C. sempervirens* pollen extracts for clinical use. This ELISA could also be useful for other Cupressaceae-related pollen extracts.

**Key Words:** Cupressus sempervirens. Cup s 1. Major allergen. Quantification. Monoclonal antibody. ELISA. Cupressaceae. Standardization. Pectate lyase.

RESUMEN

**Antecedentes:** Las cupresáceas son una importante causa de polinosis, particularmente en los países Mediterráneos. Los extractos alérgicos de polen de ciprés son difíciles de producir ya que tienen bajo contenido de proteínas y alto de carbohidratos, por lo que es esencial una precisa estandarización de estos extractos para su uso en diagnóstico e inmunoterapia.

**Método:** El alergeno natural Cup s 1 fue purificado mediante interacción hidrófoba, tamizado molecular, e intercambio iónico, y se analizó su actividad enzimática. El alergeno fue utilizado como referencia en la realización de la curva de calibrado del ensayo. El ensayo se basó en un anticuerpo monoclonal (3D2) inmovilizado en la placa y usado para capturar a Cup s 1. Posteriormente un antisuero es-
This moment for the standardization of allergenic extracts are difficult to carry out. Moreover, Cupressaceae family pollen extracts are characterized to have a high content in carbohydrates and a low protein content (3 %), what has hindered the possibility of producing good standardized extracts for diagnosis and immunotherapy.

Allergen content can be quantified in mass units by using monoclonal (mAb) and/or polyclonal antibodies specific for the allergen. The Cupressus sempervirens major allergen is a 43 kDa protein, named Cup s 1 that has been recently purified\(^1\) and cloned (EBI accession number AF7257491), sharing a high homology with Cup a 1. Recently, a mAb-based ELISA has been developed in our lab for quantification of Cup a 1\(^1\), but no accurate determination can be obtained for C. sempervirens and other Cupressaceae species. In this work we describe a two-site sandwich ELISA for the quantification of Cup s 1 in C. sempervirens pollen extracts, using a monoclonal antibody as capture and a monospecific polyclonal antibody as detector.

**MATERIALS AND METHODS**

Allergenic extracts and Cup s 1 purification

In the preparation of the allergenic extracts, pollen of *C. sempervirens*, *C. arizonica*, *Thuja plicata*, and *Juniperus communis* (Iber-Polen, Málaga, Spain) were defatted and extracted during 4 h at 4 °C in 50 mM phosphate buffer pH 8.0 containing 65 mM NaCl. The obtained extracts were clarified by centrifugation to 5000 xg for 30 min, filtered through AP glass fiber prefiltre (Millipore Corp., Bedford, Madison, USA) and dialyzed by ultrafiltration with a 5000 Daltons exclusion size in a Pellicon System (Millipore). In order to improve Cup s 1 purification, *C. sempervirens* pollen was washed for 10 min in 10 mM KH\(_2\)PO\(_4\) pH 4.3, followed by an extraction with 40 mM NH\(_4\)HCO\(_3\) at 4 °C.\(^2\) Water (NH\(_4\))\(_2\)SO\(_4\) was added to the clarified supernatant and was applied to a High Flow Phenyl-Sepharose 16/20 column equilibrated with 20 mM phosphate buffer pH 7.0, containing 2 M (NH\(_4\))\(_2\)SO\(_4\) in the AKTA-prime System\(^3\) (Amersham Biosciences, Buckinghamshire, UK). The bound fraction was eluted with 20 mM phosphate buffer pH 7.0, concentrated and applied to a Superdex S-200 16/60 column equilibrated with phosphate buffer saline (PPBS). The fraction containing the 43 kDa protein was collected, and after a dialysis with 50 mM sodium acetate pH 5.0, it was injected in a Mono S 1.6/5 ion exchange column into a Smart System\(^4\) (Amersham Biosciences). The purified al-
Cup s 1 were detected with Coomassie brilliant blue after being separated by SDS-PAGE at 12.5 % acrylamide under reducing conditions according to Laemmli method. For the immunodetection assay, the proteins were transferred to Hybond-P membranes (Amersham Biosciences) and blocked for 1 h at room temperature with PBS containing 0.1 % Tween-20. Next, the membranes were incubated under similar conditions with mAb 3D2 (0.5 μg/ml) or with anti-Cup s 1 rabbit antisera (diluted 1:500,000). After washing, blots were incubated with the corresponding peroxidase-labeled secondary antibody and developed by the ECL western blotting detection system (Amersham Biosciences).

Quantification of Cup s 1 by a two-site sandwich ELISA

Microwell plates (Greiner, Frickenhausen, Germany) were coated overnight at room temperature with 100 μg/well of the mAb 3D2 at 5 μg/ml in PBS. Next, plates were blocked by adding 200 μg/well of PBS-1 % BSA-0.05 % Tween 20 (PBS-B-T) and incubated 1 h at 37 °C. Afterwards, wells were incubated with 100 μg/well of purified Cup s 1 (2000-2 ng/ml), or with the C. sempervirens pollen extracts (10000-25 ng/ml) in PBS-B-T, followed by a second incubation with biotin-labeled anti-Cup s 1 rabbit polyclonal antibody (0.31 μg/ml) and finally with streptavidin-peroxidase (0.25 μg/ml) (Sigma). All the incubations were carried out at 37 °C during 1 h, and with 3 washes of 200 μg/well of PBS-T between incubations. Color development was performed at room temperature and in darkness with a solution of o-phenylenediamine (Sigma-Fast o-phenylenediamine dihydrochloride Tablet Sets, Sigma). The reaction was stopped at 30 min with 50 μg/ml of 3 M H₂SO₄, and the optical density was measured at 492 nm.

RESULTS

In the Cup s 1 purification, the pollen was subjected to a quick wash with low ionic content and acidic pH buffer, followed by an extraction at basic pH with the purpose of eliminating most of the non-protein components. The obtained supernatant was supplemented with 2 M (NH₄)₂SO₄ with the purpose of promoting the protein binding to phenyl-sepharose resin. After washing, the bound fraction was eluted with phosphate buffer, and contained a high concentration of proteins with low amounts of carbohydrates (fig. 1A). Further purification was obtained by gel filtration chromatography where Cup s 1 was contained in the 40-50 kDa peak (fig. 1B). This frac-
The purification of Cup s 1 from C. sempervirens extracts was subjected to a last purification step in a Smart System using a cation exchange column. Cup s 1 was eluted with 100 mM NaCl as a 43 kDa band while the rest of proteins come out in the non-bound fraction at pH 5.0 (fig. 1C). The purification yield was 1.5 % of the total extract proteins.

The identification of the purified protein as Cup s 1 was proven by immunoblotting with specific antibodies, as well as for its pectate lyase activity. Cup s 1 possesses the capacity to degrade pectin with a specific activity of 725 U/mg protein, and this activity is also completely dependent of calcium (fig. 1D). This is the first time it has been demonstrated that Cup s 1 is a functional active pectate lyase enzyme.

For the quantification of Cup s 1 in the two-site sandwich ELISA, the mAb 3D2 was used as capture
The extract dose-response curves showed a good parallelism when compared with the obtained with the purified allergen, indicating that the same protein was measured (fig. 3). The Cup s 1 standard curve was applied to the quantification of Cup s 1-like proteins in T. plicata and J. communis pollen extracts, with an allergen content of 75 and 50 g/mg of protein, respectively. Nevertheless, for accurate quantification, the corresponding purified group 1 allergen for each extract should be used as reference in the dose-response curve. The specificity of the assay was tested using extracts of non-plant origin (Alternaria alternata and Der- matophagoides pteronyssinus), and pollen extracts from Olea europaea, Platanus acerifolia, Betula verrucosa, Pinus radiata, Cedrus atlantica, Dactylis glomerata, Helianthus annuus, Parietaria judaica, Artemisia vulgaris and Plantago lanceolata. No reactivity was detected between 0.25 and 25 g/ml of protein in any case.

This assay was used to quantify the Cup s 1 content in different batches of C. sempervirens pollen extracts, ranging from 73 to 95 µg/mg of protein.

The extract dose-response curves showed a good parallelism when compared with the obtained with the purified allergen, indicating that the same protein was measured (fig. 3). The Cup s 1 standard curve was applied to the quantification of Cup s 1-like proteins in T. plicata and J. communis pollen extracts, with an allergen content of 75 and 50 µg/mg of protein, respectively. Nevertheless, for accurate quantification, the corresponding purified group 1 allergen for each extract should be used as reference in the dose-response curve. The specificity of the assay was tested using extracts of non-plant origin (Alternaria alternata and Dermatophagoides pteronyssinus), and pollen extracts from Olea europaea, Platanus acerifolia, Betula verrucosa, Pinus radiata, Cedrus atlantica, Dactylis glomerata, Helianthus annuus, Parietaria judaica, Artemisia vulgaris and Plantago lanceolata. No reactivity was detected between 0.25 and 25 µg/ml of protein in any case.

<table>
<thead>
<tr>
<th>Capture</th>
<th>Detection</th>
<th>Linear portion of the standard curve</th>
<th>Detection limit</th>
<th>Intrassay CV (31.25-250 ng/ml)</th>
<th>Interassay CV (31.25-250 ng/ml)</th>
</tr>
</thead>
<tbody>
<tr>
<td>mAb 3D2 (5 µg/ml)</td>
<td>Biotinylated polyclonal antibody (0.31 µg/ml)</td>
<td>31.25-250 ng/ml $y = 1.382 + 0.447x$ $r = 0.982$ ($n = 10$)</td>
<td>3.8 ng/ml</td>
<td>&lt; 8.1% ($n = 20$)</td>
<td>&lt; 16% ($n = 10$)</td>
</tr>
</tbody>
</table>
The allergic extract standardization is usually carried out by means of in vitro techniques using allergic patient sera, which are difficult to homogenize. In the case of Cupressaceae allergy, the finding of good sera becomes more difficult since these patients have low levels of specific IgE to cypress extracts. Therefore, the development of an immunosassay for the quantification of Cup s 1 is of great interest as a complementary tool for standardization. The success in the development of this type of procedures is closely linked to the existence of specific antibodies and the availability of a high grade purity standard. The mAb 3D2, used as capture mAb, reacted with Cup s 1, Cup a 1 and a corresponding protein of T. plicata and J. communis pollen extracts. Therefore, it seems to detect a conserved epitope presented in Cupressaceae proteins and becomes a useful tool in the immunosassays for the quantification of major allergen within the Cupressaceae family. Nevertheless, the corresponding purified allergen for each extract should be used for an accurate quantification of the allergen content expressed in mass units. The use of polyclonal antibodies as tracer did not diminish the assay specificity, since this second antibody did not show reaction to any other proteins, even from Pinus or Cedrus extracts, plants from the Pinaceae family included, as the Cupressaceae family, inside the Coniferal order. Using the described assay, allergen concentration of 7.3, 7.5 and 5 % for pollen extracts of C. sempervirens, T. plicata and J. communis pollen seems to be more compatible with data of protein yield after purification and immunoblotting than allergen concentration of 0.5, 1.2, and 1.6 % obtained previously using a sandwich ELISA based on two mAbs.

The ELISA could be applicable to the routine analyses of cypress extracts used to specific immunotherapy and diagnostic, given the high sensitivity and specificity requirements, was also fulfilled in the two-site sandwich ELISA, with acceptable intraassay and interassay variation coefficients. In conclusion, this paper describes a two-site sandwich ELISA procedure for measurement of Cup s 1 in C. sempervirens pollen extracts. This assay results useful due to its specificity, sensitivity and reproducibility, and it can be applicable to the quantification of the other Cupressaceae major allergens.

ACKNOWLEDGMENTS

This work was supported in part by Bial-Aristegui and by Grants FIT-090000-2003-61 from the Plan Nacional de I + D (Programa PRQFIT, Ministerio de...
Ciencia y Tecnología, Spain), and TEI-0163-2002 from the Programa INTEK (Departamento de Industria, Agricultura y Pesca, Gobierno Vasco).

REFERENCES

10. Arilla MC, Barris K, García R, de la Hoz B, Martínez A, Asturias JA. Quantification of the major allergen from cypress (Cupressus arizonica) pollen. Cup s 1, by monoclonal antibody-based ELISA. Int Arch Allergy Immunol 2004;134:10-6.

Allergol et Immunopathol 2004;32(6):319-25