

Eukaryotic expression of recombinant Pso o 1, an allergen from *Psoroptes ovis*, and its localization in the mite

A. J. NISBET^{1*}, A. MACKELLAR¹, K. MCLEAN¹, G. P. BRENNAN² and J. F. HUNTLEY¹

¹ Moredun Research Institute, Pentlands Science Park, Bush Loan, Penicuik, EH26 0PZ, Scotland

² School of Biology and Biochemistry, Queen's University Belfast, Lisburn Road, Belfast BT9 7BL, Northern Ireland

(Received 16 June 2006; revised 17 July 2006; accepted 17 July 2006; first published online 18 September 2006)

SUMMARY

A cDNA encoding the immunogen Pso o 1 from *Psoroptes ovis* was obtained by polymerase chain reaction (PCR) amplification. The amplicon contained the entire coding sequence for the prepro-enzyme in an open reading frame (ORF) of 966 bp. This gene encoded a predicted protein of 322 amino acids (aa) with 64% aa identity (80% similarity) to the major house dust mite faecal allergen Der f 1. The pro-enzyme form of Pso o 1 was expressed as a recombinant protein in the *Pichia pastoris*-eukaryotic expression system. Maturation of the recombinant pro-enzyme by autocatalytic activation was not observed, and such maturation could not be achieved using a number of techniques known to activate recombinant Der p 1 and Der f 1 expressed in the same system. Serum raised against recombinant Pso o 1 cross-reacted with mature Der p 1 and allowed Pso o 1 to be immunolocalized to the gut of *P. ovis*.

Key words: *Psoroptes*, allergen, Pso o 1, cysteine proteinase.

INTRODUCTION

The exudative lesion, which is a clinical feature of sheep scab, appears rapidly after *Psoroptes ovis* mites are applied to the skin of the ovine host and is the result of pro-inflammatory factors derived from the parasite (van den Broek and Huntley, 2003). The continued presence of the sheep scab mites on the skin, and their feeding, excretion and secretion of immunogenic compounds, provokes both intra-dermal inflammatory responses and the generation of mite-specific IgE and IgG antibodies (van den Broek *et al.* 2000, 2003). *P. ovis* allergens have therefore been implicated both in the pathogenesis and immunity to infection (Huntley *et al.* 2004). As a result, these allergens may offer novel targets for diagnosis, therapies and vaccine development (Nisbet and Huntley, 2006).

Lee *et al.* (2002) described the isolation of a cDNA representing a truncated form of Pso o 1 from *P. ovis* that encoded a cysteine proteinase with 54% amino acid (aa) identity (69% similarity) to the major house dust mite allergen Der f 1 from *Dermatophagoides farinae*. The group I allergens, which include Der f 1, Der p 1 and Eur m 1 from *D. farinae*, *D. pteronyssinus* and *Euroglyphus maynei*, respectively (Chua *et al.* 1988; Dilworth *et al.* 1991; Kent *et al.* 1992) belong to the papain-superfamily of cysteine proteinases,

and approximately 90% of humans allergic to house dust mites possess IgE reactivity to these allergens (Arlan and Platts-Mills, 2001). Using a bacterially-expressed recombinant truncated Pso o 1, Lee *et al.* (2002) demonstrated that sheep developed a pronounced IgG response to this proteinase within the first 3 weeks of a primary *P. ovis* infestation and that some sheep also developed a weak, specific IgE response.

Der p 1 is synthesized as a prepro-enzyme of 320 amino acids (aa), composed of an 18-aa signal peptide and 80-aa N-terminal pro-sequence (Chua *et al.* 1988; Dilworth *et al.* 1991). Autocatalytic cleavage of the pro-region activates the mature enzyme, and studies using the specific and irreversible Der p 1 proteinase inhibitor PTL11028 have shown that the prevention of cleavage of the low-affinity receptor for IgE from human B cells by Der p 1 reduced significantly house dust mite allergen-induced hypersensitivity-related pathogenesis (John *et al.* 2000). This information indicates that the enzymatic activity of the cysteine proteinase is important or essential for the development of an allergic response. In addition, the correct tertiary structure of proteins is often critical for IgE binding. The previously characterized, truncated recombinant Pso o 1 protein lacked one of the C-terminal catalytic site residues characteristic for cysteine proteinases (Asn¹⁹¹ in the mature form of Der p 1) and was therefore unlikely to be enzymatically active (Lee *et al.* 2002). Recent reports (e.g. Takai *et al.* 2002) have demonstrated the success of producing mature recombinant Der p 1 with enzymatic activity and IgE binding-capacity in

* Corresponding author: Moredun Research Institute, Pentlands Science Park, Bush Loan, Penicuik, EH26 0PZ, Scotland. Tel: +44 (0)131 445 5111. Fax: +44 (0)131 445 6235. E-mail: Alasdair.Nisbet@moredun.ac.uk

the yeast *Pichia pastoris*, by expressing the protein either as a prepro-enzyme or a pro-enzyme and allowing self-activation. In addition, expressed sequence tag (EST) analysis of *P. ovis* has shown the presence of a number of transcripts encoding proteins with a significant homology to both Pso o 1 and Der p 1 (Kenyon *et al.* 2003). The aims of the current study were to use data from the EST project to determine the entire coding sequence of *Pso o 1*, to express the recombinant protein in a eukaryotic system in order to increase the likelihood of obtaining an enzymatically active recombinant protein and to determine the tissue distribution of the native enzyme to infer its potential biological role.

MATERIALS AND METHODS

Molecular cloning and expression of Pso o 1

A contiguous sequence was formed of the known *P. ovis* sequences and expressed sequence tags (ESTs) encoding Der p 1 homologues (NCBI Accession AF495854 and EMBL Accession BQ834765). Oligonucleotide primers were designed to the contig, to amplify by the polymerase chain reaction (PCR) the putative 966 bp ORF of the gene encoding prepro-Pso o 1. The sequences of these primers were: 5'-ATGAAATTTGTTTTGGCCATCGCCTC-3' (ppPsoo1Bf) and 5'-CTCAAAGCATTGCAGCCAATGGG-3' (ppPsoo1Br).

The PCR was performed using the Advantage[®]2 PCR system (BD Biosciences) according to the manufacturer's instructions, employing a *P. ovis* cDNA library lysate (diluted 1:100) as the template (Kenyon *et al.* 2003) and employing the following cycling parameters: 95 °C 1 min (1 cycle), 95 °C 15 sec, 68 °C 1 min (30 cycles). The resultant amplicon was column-purified (QIAquick[®] PCR purification kit, Qiagen), diluted 1:100 and used as a template to amplify cDNAs encoding prepro-Pso o 1 and pro-Pso o 1 (without a signal peptide) while simultaneously incorporating restriction sites for the enzymes *Xho*1 and *Not*1 at the 5' and 3' ends of the coding sequence, respectively. The forward primer for prepro-Pso o 1 was thus *Xho*1Psoo1B; 5'-GTA-TCTCTCGAGAAGAGAATGAAATTTGTTTT-TGG-3' and that for pro-Pso o 1 was *Xho*XPsoo1B; 5'-GTATCTCTCGAGAAGAGAGCTTACCCATCAG-3' (*Xho*1 cleavage sites are underlined). The latter (*Xho*XPsoo1B) was designed to amplify from the putative cleavage site of the signal peptide from the mature molecule (between Ala¹⁸ and Tyr¹⁹, see Fig. 1). The reverse primer in these reactions was *Not*1Psoo1B; 5'-AAAGCTGGCGGCCGCTCAA-AGCATT-3' (*Not*1 restriction site underlined); the cycling parameters were 95 °C 1 min (1 cycle), 95 °C 15 sec, 53 °C 30 sec, 68 °C 1 min (30 cycles), 70 °C 10 min (1 cycle), using the Advantage[®]2 PCR system. The resultant amplicons were digested with

the relevant restriction enzymes and ligated into the vector pPICZaC (Invitrogen) in frame with the secretion signal of the vector. The constructs were transformed into *Escherichia coli* JM109 (Promega) cultured in low-salt Luria Bertani medium (LB) prior to selection on low-salt LB agar containing Zeocin[™] (25 µg/ml). Colonies with plasmids containing the cDNA inserts were isolated and plasmids extracted, using Wizard[®] Plus SV Minipreps (Promega), after overnight liquid culture in low-salt LB containing Zeocin[™] (25 µg/ml). Automated sequencing using ET terminator chemistry on a MegaBACE DNA analyser confirmed the frame and sequence of each construct. Plasmids were linearised using *Pme*1 (New England BioLabs), column-purified (QIAquick[®] PCR purification kit, Qiagen), precipitated, and used to transform *Pichia pastoris* (Km71 strain) by electroporation, as described by the supplier (Invitrogen). Transformants, isolated by Zeocin[™] selection on yeast peptone dextrose (YPD) agar plates, were grown in 5 ml of buffered complex medium with 1% glycerol (BMGY) for 16 h at 30 °C before centrifugation (1500 g for 10 min) and resuspended in the same medium containing 0.5% methanol (BMMY). Protein expression was induced by adding methanol to a final concentration of 0.5% at 24 h intervals thereafter for 96 h. For large-scale cultures, transformants were grown for 30 h at 30 °C in 500 ml BMGY in 2 l capacity baffled flasks with vigorous shaking, centrifuged and resuspended in 50 ml of BMMY in 250 ml flasks. Protein expression was induced as above for 72 h. Culture fluid was collected at 24 h intervals and centrifuged (13 000 g for 10 min) to separate soluble proteins from the insoluble fraction. Pro-Pso o 1 was purified from the culture supernatant by gel filtration using a Sepharose S300 column, equilibrated and eluted with phosphate-buffered saline, pH 7.4 (PBS).

Verification of Pso o 1 identity by Western blot and matrix-assisted laser desorption/ionization time of flight (MALDI-ToF) mass spectrometric analysis

Culture supernatant was used for electrophoresis on a NuPAGE[®] Bis-Tris 4–12% gel (Invitrogen) under reducing conditions. The gel was stained with SimplyBlue[™], according to the manufacturer's instructions (Invitrogen), and the band detected (see Fig. 2) was excised, destained and reductively alkylated using dithiothreitol and iodoacetamide. The gel pieces were then digested overnight with trypsin or chymotrypsin at 37 °C. Digests were analysed using an Ultraflex II MALDI-ToF mass spectrometer (Bruker Daltonics), scanning the 600–5000 Dalton (Da) region in the reflectron mode producing monoisotopic resolution. The spectra generated were mass-calibrated using known standards and the peaks deisotoped. Databases were searched with the masses obtained using the MASCOT search engine

1 M K F V L A I A S L M V L S V V Y A Y P
 1 ATGAAATTTGTTTTGGCCATCGCCTCATTGATGGTGTGAGCGTTGTTTACGCTTACCCA
 21 S E I R T F E E F K K A S N K H Y V T P
 61 TCAGAAATCAGAACATTCGAAGAATTCAAAAAGCATCCAAACACATATGTAACACCT
 41 E A E Q E A R Q N F L A S L E H I E K A
 121 GAAGCAGAACAGAAGCTCGTCAAAATTTCTTAGCTTCATGGAAACACATGAAAAAGCT
 61 G K G R I N Q F S D M S L E E F K N Q Y
 181 GAAAAGGTGCGCATCAATCAATTCAGATATGTCATTGGAAGAATTCAAAAACCAATAT
 81 L M S D Q A S E A L K K E F D L D A G A
 241 TTGATGAGTGATCAAGCATAAGCTTTAAAAAGAATTTGATTTAGATGCTGGAGCT
 101 Q A C Q I G A V N I P N E I D L R A L G
 301 CAAGCTTGCCAAATCGGCGCGTAAACATTCCAAATGAAATTTGATTTACGTGCTTTGGGT
 121 Y V T K I K N Q V A C G S C W A F S G V
 361 TATGTAACAAAAATCAAGAATCAAGTTGCCTGTGGTTCATGCTGGGCTTTCTCTGGTGT
 141 A T V E S N Y L S Y D N V S L D L S E Q
 421 GCTACAGTCGAATCAAATTTATTCATACGATAATGTATCATTAGATCTTTCTGAACAA
 161 E L V D Q A S Q H G C G G D T V L N G L
 481 GAATTGGTTGACTGTGCTTACAACACGGTTGCGGTGGTGATACAGTTCTAAACGGTTTA
 181 R Y I Q K N G V V E E Q S Y P Y K A R E
 541 CGTTATATCAAAAAGAAATGGTGTGGTTGAAGAGCAAAGCTATCCATACAAAGCTCGGAA
 201 G R C Q I R D N A K R Y G I K D L C Q I Y
 601 GGACGATGCCAAAGACCAATGCTAAACGATACGGTATCAAAGATTTATGTCAAATTTAT
 221 P P N G D K I R T Y L A T K Q A A L S V
 661 CCACCAAACGGTGATAAAATCCGTACCTATCTTGCAACAAAAACAAGCAGCCCTTTTCAGTC
 241 I I G I R D L D S F R H Y D G R T I L Q
 721 ATCATTGGTATCCGAGATTTAGATTTCTTTCCGTATTATGATGGCCGAACAATTTTGCAA
 261 S D N G G K R D F H A I N I V G Y G S K
 781 AGTGATAATGGTGGCAACGAGATTTCCATGCAATCAATATTGTTGGTTACGGATCAAAA
 281 Q G V R Y W I I R N S W D T T W G D K G
 841 CAAGCGTTTCGATATTGGATTATACGAAACAGTTGGGACACAACCTTGGGGTGATAAAGGC
 301 Y G Y F V A D K N L M G I E K F P L A A
 901 TATGGCTATTTTGTGTGCTGATAAAAATTTGATGGGTATCGAAAAATTCACATGGCTGCA
 321 M L *
 961 ATGCTTTGA

Fig. 1. Nucleotide and deduced amino acid sequence of *prepro-Pso o 1*. The putative signal peptide is underlined and amino acid residues differing from the deduced amino acid sequence derived from a truncated version of *Pso o 1* (Lee *et al.* 2002) are shown as white text in black boxes. N-terminal and C-terminal residues not previously determined (Lee *et al.* 2002) are shown in italics. A potential N-glycosylation site is shown boxed. The consensus thiol proteinase cysteine active site (prosite motif PS00139) is grey shaded and that of the thiol proteinase asparagine active site (prosite motif PS00640) is double underlined.

(Swiss-Prot database) and a 50 ppm mass tolerance window. Significant matches from the Peptide Mass Fingerprint data were confirmed by MS/MS analysis using the search criteria described and an MS/MS-tolerance window of 0.5 Da.

For Western blots, culture supernatant, removed from a pro-*Pso o 1* culture 72 h after the induction of protein expression, or purified *Pso o 1* (before and after induction of proteinase activity), were subjected to electrophoresis in NuPAGE[®] (Invitrogen) Bis-Tris 4–12% gels under reducing conditions employing NuPAGE[®] MES SDS running buffer (Invitrogen). Proteins were transferred to a nitrocellulose membrane according to the manufacturer's instructions (Invitrogen). After transfer, the membranes were washed briefly in Tris-buffered saline, pH 7.4 (TBS; 25 mM Tris, 137 mM NaCl, 2.7 mM KCl) containing 0.1% (v/v) Tween 20 (TBST) and then incubated in TBST containing 5% (w/v) skimmed milk powder for 4 h at room temperature (RT; 22–24 °C) to block non-specific protein adsorption. To detect *Pso o 1*, the blotted proteins were incubated, for 1 h at RT, with serum produced in sheep against bacterially-expressed truncated *Pso o 1* (Lee *et al.* 2002). Control blots

were performed by omitting the test serum during this primary incubation step. Following extensive washing in TBST, membranes were incubated in affinity-purified donkey anti-sheep IgG horse-radish peroxidase (HRP) conjugate (Sigma) in TBST containing 5% (w/v) skimmed milk powder. Following incubation at RT for 1 h, and extensive washing, peroxidase activity was revealed using 3,3'-diaminobenzidine (DAB) as a substrate.

Measurement of enzymatic activity of recombinant *Pso o 1*

Activation of the recombinant pro-enzyme was attempted using each of the following methods. Dialysis against 100 mM acetate buffer (pH 4) for 72 h at 4 °C or at RT (Yasuhara *et al.* 2001; Takai *et al.* 2002); deglycosylation with endoglycosidase H (van Oort *et al.* 2002); exogenous activation with native Der p 1 (van Oort *et al.* 2002); heating to 60 °C in the presence of 20 mM cysteine (Jacquet *et al.* 2000; van Oort *et al.* 2002). In addition, the recombinant protein was incubated, for 10 min at 37 °C, with soluble extracts of *P. ovis* (0.02–0.4 µg protein, prepared as described previously (Nisbet

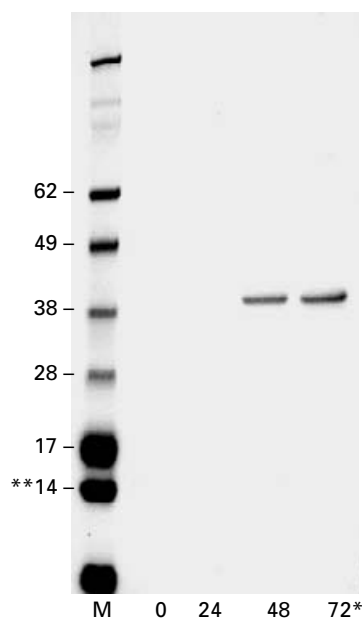


Fig. 2. Expression of Psoo1 in *Pichia pastoris*. Culture supernatant, removed from a pro-Pso o 1 culture 0–72 h after the induction of protein expression, was used for electrophoresis on a NuPAGE[®] Novex Bis-Tris 12–4% gel under reducing conditions and stained with SimplyBlue[™] according to the manufacturer's instructions (Invitrogen). * Hours after induction of protein synthesis, ** Molecular mass (kDa).

and Billingsley, 2000) in 100 mM phosphate buffer, pH 5.8, containing cysteine (10 mM) prior to the testing in the enzyme assay. The enzymatic activity was assessed using the substrates butyloxycarbonyl-Val-Leu-Lys-MCA (Sasaki *et al.* 1984) and Z-Phe-Arg-MCA (Bando *et al.* 1986). Positive control reactions were performed for both substrates using a soluble extract of *P. ovis* and/or purified native Der p 1 as an enzyme source. Native Der p 1 was a generous gift from Dr Colin R. A. Hewitt, University of Leicester, UK.

Immunolocalization of Pso o 1

Mites (adults, nymphs and larvae) were collected from infested donor sheep (× Suffolk sheep, aged 1–2 years), fixed in Carnoy's fluid and stored at 4 °C until processing. Fixative was removed and mites were pelleted in 1% molten agarose and left to solidify before processing into paraffin wax. Sections (5 µm) were dried at 40 °C, dewaxed in xylene and rehydrated. Sections were treated to remove endogenous peroxidase (EnVision Kit[™], DAKO) and blocked in 25% normal goat serum in TBS, pH 7.6, for 30 min before an overnight incubation at 4 °C in serum raised in rabbits against recombinant pro-Pso o 1 (1 in 2000 dilution in PBS). Following washing and incubation in goat anti-rabbit HRP conjugate (1:1000 dilution) (DakoCytomation) for 1 h at RT, the sections were washed, and DAB substrate

(Sigma) was used to detect the goat antibody conjugate. Negative controls included were a 'pre-bleed' serum from rabbits prior to injection with Pso o 1 (1 in 2000 dilution in PBS) and a 'conjugate only' control. Sections were washed in water, counterstained in haematoxylin for a few sec, washed in water, dehydrated, cleared and mounted in a resin-based mountant (Nisbet *et al.* 2006).

RESULTS

Molecular cloning and expression of Pso o 1

The PCR product obtained by amplification using primers ppPsoo1Bf and ppPsoo1Br was sequenced, and shown to contain an open reading frame (ORF) of 966 bp (nucleotide sequence data available in the EMBL, GenBank[™] and DDJB databases under Accession number AM269885). This gene encoded a 322 aa predicted protein with 64% identity (80% similarity) to the major house dust mite faecal allergen Der f 1 (prepro-enzyme, Accession number BAC53948). The predicted protein, Pso o 1, also shared 99% identity over 263 aa residues with the previously characterized, truncated form (Lee *et al.* 2002). However, the asparagine residue involved in the catalytic activity of cysteine proteinases (Asn²⁹⁰ in the prepro-enzyme sequence) was present in the C-terminus of the full-length Pso o 1 described herein (Fig. 1).

Pro-Pso o 1 was expressed efficiently in yeast within 48 h of induction (Fig. 2), but the expression of prepro-Pso o 1 was unsuccessful in this system (results not shown). The apparent molecular mass of the recombinant protein was 40 kDa, ~6 kDa larger than that based on the cDNA sequence. The recombinant protein reacted strongly with sheep anti-truncated Pso o 1 serum on Western blot (Fig. 3, lane 1); both the chymotrypsin and trypsin digests and MALDI-ToF analysis of the expressed protein confirmed that the recombinant protein was Pso o 1.

Measurement of enzymatic activity of recombinant Pso o 1

Large-scale expression of pro-Pso o 1 was performed, and attempts were made to activate the recombinant protein to a mature active enzyme. Enzymatic activity was assessed using the fluorogenic substrates butyloxycarbonyl-Val-Leu-Lys-MCA or Z-Phe-Arg-MCA. Activation of the recombinant enzyme was attempted employing each of the following treatments: dialysis against acidic buffer for 72 h; deglycosylation; activation with mite proteinases and heating to 60 °C. With the exception of the dialysis, each treatment reduced the apparent molecular mass of the major band by 2–4 kDa (Fig. 3), but none of the treatments resulted in a protein of the same molecular mass as native Pso o 1

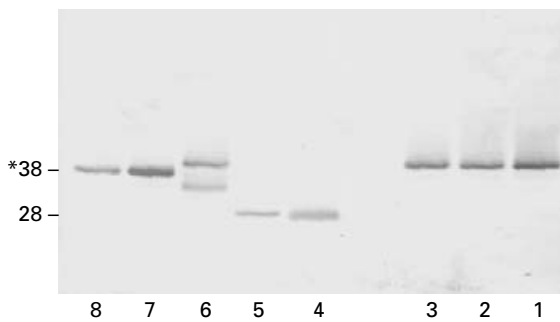


Fig. 3. Western blot of Pso o 1 and its derivatives. Lanes on blot are as follows: (1) pro-Pso o 1 culture supernatant (CS); (2) pro-Pso o 1 CS dialysed against acidic (pH 4) buffer for 72 h; (3) pro-Pso o 1 purified from CS by gel filtration; (4) purified native Der p 1; (5) soluble *P. ovis* extract containing native Pso o 1; (6) pro-Pso o 1 CS incubated at 60 °C for 10 min; (7) pro-Pso o 1 CS treated with 12.5 mu endoglycosidase H; (8) purified pro-Pso o 1 treated with 12.5 mu endoglycosidase H. To detect Pso o 1, Der p 1 and derivatives, the blotted proteins were incubated with serum raised in sheep against bacterially-expressed truncated Pso o 1 before being incubated in donkey anti-sheep IgG horse radish peroxidase (HRP) conjugate and peroxidase activity was revealed using DAB as a substrate. * Molecular mass (kDa).

or Der p 1 (Fig. 3 lanes 4 and 5). Soluble extract of *P. ovis* and native Der p 1 positive controls efficiently hydrolysed both substrates, but none of the treatments produced active enzyme from Pso o 1. In addition, treatment of the recombinant protein with a range of concentrations of a soluble extract of *P. ovis* failed to produce active enzyme.

Immunolocalization of Pso o 1

Pso o 1 was localized specifically to the gut in sections of adult *P. ovis* (Fig. 4A). The labelling of Pso o 1 was most intense near the luminal surface of the gut. Minor non-specific reactivity of the secondary antibody (goat anti-rabbit IgG) and/or the substrate was observed in the tissues underlying the cuticle in both the negative controls and the sections treated with rabbit anti-Pso o 1 (Fig. 4). Negative controls were identical when primary antibody was omitted (Fig. 4B) and when rabbit 'pre-bleed' serum was used in the primary incubation step (not shown).

DISCUSSION

The entire coding sequence of Pso o 1 was determined to augment the truncated sequence which had already been published (Lee *et al.* 2002). The newly determined sequence encoded a protein with a predicted signal peptide which is cleaved between Ala¹⁸ and Tyr¹⁹ [predicted using SignalP 3.0 (Dyrlov Bendtsen *et al.* 2004)] and possessed significant homology to the house dust mite group 1 allergens Der f 1, Der p 1 and Eur m 1, suggesting that Pso o 1

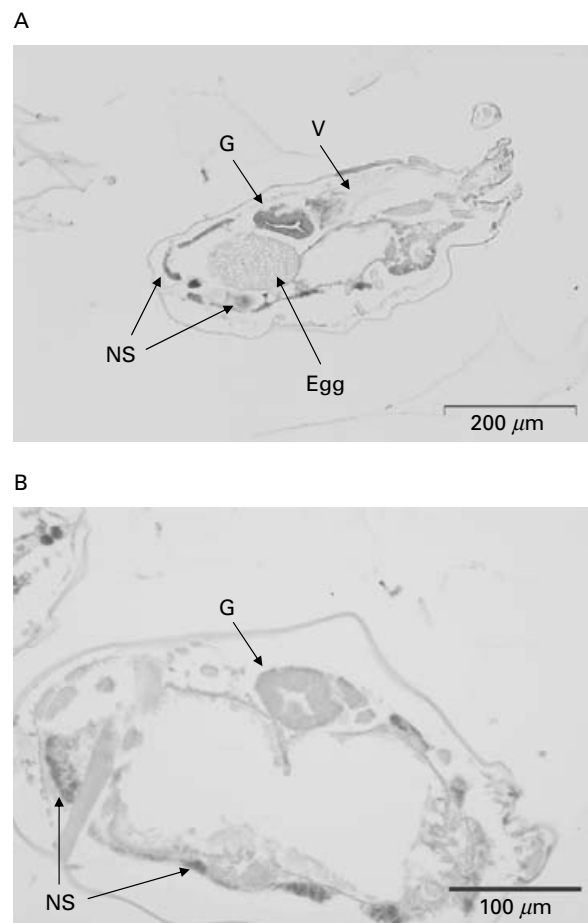


Fig. 4. Immunolocalization of Pso o 1 in sections of adult *Pichia ovis*. (A) Incubated with rabbit anti-Pso o 1 (1 in 2000 dilution) as primary antibody; (B) primary antibody omitted. Immunoreactivity was visualized using goat anti-rabbit HRP conjugate followed by incubation in DAB substrate. (B) Conjugate-only control. Areas annotated with 'G' show sections through gut; 'V', ventriculus; 'NS' denotes non-specific staining resulting from binding of the secondary antibody and/or reactivity of the substrate.

is secretory in nature and that it may also act as an allergenic component in the pathogenesis of sheep scab disease. Lee *et al.* (2002) noted that Pso o 1 possessed a high degree of amino acid homology within the residues surrounding the propeptide cleavage site involved in the maturation of active enzyme in house dust mite group 1 allergens and suggested that this area may also be involved in the maturation of Pso o 1 cysteine proteinase activity. These residues are present in the full-length Pso o 1 molecule described herein. Interestingly, only 50% of the 12 residues surrounding the proposed cleavage site (between Ala⁹⁸ and Gly⁹⁹ in Fig. 1) are identical between Der p 1 and Pso o 1, compared with >80% identity in the same region between the different house dust mite allergens. This information may explain the failure of autocatalytic processing of the recombinant Pso o 1, suggesting that alternative processing may be required, though endogenous

proteinase activity in soluble *P. ovis* extracts also failed to activate the recombinant enzyme. The ~6 kDa discrepancy between the observed and calculated masses for pro-Pso o 1 suggests hyperglycosylation of Pso o 1 in the yeast expression system, a phenomenon which has also been reported from a recent expression study of the group 1 house dust mite allergens (Takai *et al.* 2002).

Cysteine proteinase activity is abundant in the soluble extracts of a number of mite species, including *P. ovis* (see Nisbet and Billingsley, 2000). These enzymes are involved in fundamental processes in the pathogenesis of sheep scab, including the degradation of connective tissue and extracellular matrix molecules and the catalysis of immunoglobulins (Kenyon and Knox, 2002). By analogy to dust mite allergens, cysteine proteinases and Pso o 1, in particular, may also be involved in allergen-induced hypersensitivity-related effects (John *et al.* 2000). Studies using both storage mites (*Acarus farris*) and dust mites (*D. farinae*), fed on diets containing cysteine proteinase inhibitors, have also shown that the inhibition of cysteine proteinase activity in the guts of mites retarded their development and reduced survival, suggesting critical roles in the digestive process (Pernas *et al.* 1998; Sánchez-Ramos *et al.* 2004).

In addition, Der p 1 has been localized in the oral cavity, gut epithelium, gut contents and faecal pellets of *D. pteronyssinus* and is thought to be synthesized by cells lining the gastrointestinal tract (Rees *et al.* 1992). The present study demonstrated that Pso o 1 is localized within the gut of the *P. ovis*, making it potentially accessible to the immunoglobulin component of the exudates which have been shown to be imbibed by sheep scab mites feeding on the host (Pettit *et al.* 2000). Thus, an attractive possibility is that vaccination with gut cysteine proteinases, such as Pso o 1, will induce the production of 'blocking antibodies' which inhibit cysteine proteinase activity in the mite gut, resulting in a retardation of mite development and/or survival (Nisbet and Huntley, 2006). By further analogy with Der p 1, another possible beneficial effect of vaccination with Pso o 1 may be a down-regulation of the allergic inflammatory Th2 response involved in the pathogenesis of sheep scab disease (John *et al.* 2000).

Vaccination with cysteine proteinases has been shown to be effective against a number of other parasites. For example, the vaccination of dogs with Ac-CP-2, a recombinant cysteine proteinase derived from the hookworm *Ancylostoma caninum*, resulted in high antibody responses, the stunting of adult worms and a marked reduction in faecal egg output (Loukas *et al.* 2004). Also, IgG purified from the serum of vaccinated dogs inhibited the enzymatic activity of Ac-CP-2 by 73% (Loukas *et al.* 2004). Although enzymatic activity of recombinant proteins may be desirable for enhanced protection, it is

not always essential. For example, vaccination with inactive recombinant cysteine proteinases (hmcpl1, 4 and 6) from *Haemonchus contortus* resulted in a significant reduction in the intensity of infection in sheep (Redmond and Knox, 2004, 2006). Thus, although the recombinant Pso o 1 expressed in yeast herein was inactive as a cysteine proteinase, it may represent a vaccine candidate. Future studies will focus on this possibility.

A. J. N. and J. F. H. gratefully acknowledge funding by the Department for Environment Food and Rural Affairs, UK Research Contract OD0544 and J. F. H., A. McK and K. McL acknowledge funding by the Scottish Executive Environment and Rural Affairs Department. The authors are grateful to George Newlands, Jim Redmond and Karina Reinhardt, Moredun Research Institute, for their assistance in the expression, purification of recombinant enzymes and DNA analyses, respectively, and to Toshiro Takai, Juntendo University School of Medicine, Japan, for advice and useful discussions during this project. We are grateful to Dr Colin R. A. Hewitt, University of Leicester, for the gift of native Der p 1.

REFERENCES

- Arlian, L. G. and Platts-Mills, T. A. E.** (2001). The biology of dust mites and the remediation of mite allergens in allergic disease. *Journal of Allergy and Clinical Immunology* **107**, S406–S413.
- Bando, Y., Kominami, E. and Katunuma, N.** (1986). Purification and tissue distribution of rat cathepsin L. *Journal of Biochemistry* **100**, 35–42.
- Chua, K. Y., Stewart, G. A., Thomas, W. R., Simpson, R. J., Dilworth, R. J., Plozza, T. M. and Turner, K. J.** (1988). Sequence analysis of cDNA coding for a major house dust mite allergen, Der p 1. Homology with cysteine proteases. *Journal of Experimental Medicine* **307**, 175–182.
- Dilworth, R. J., Chua, K. Y. and Thomas, W. R.** (1991). Sequence analysis of cDNA clone coding for a major house dust mite allergen Der f I. *Clinical and Experimental Allergy* **21**, 25–32.
- Dyrløv Bendtsen, J., Nielsen, H., von Heijne, G. and Brunak, S.** (2004). Improved prediction of signal peptides: SignalP 3.0. *Journal of Molecular Biology* **340**, 783–795.
- Huntley, J. F., Machell, J., Nisbet, A. J., van den Broek, A., Chua, K. Y., Cheong, N., Hales, B. and Thomas, W. R.** (2004). Identification of tropomyosin, paramyosin and apolipoprotein/vitellogenin as three major allergens of the sheep scab mite, *Psoroptes ovis*. *Parasite Immunology* **26**, 335–342.
- Jacquet, A., Haumont, M., Massaer, M., Daminet, V., Garcia, L., Mazzu, P., Jacobs, P. and Bollen, A.** (2000). Biochemical and immunological characterisation of a recombinant precursor form of the house dust mite allergen Der p 1 produced by *Drosophila* cells. *Clinical and Experimental Allergy* **30**, 677–684.
- John, R. J., Rusznak, C., Ramjee, M., Lamont, A. G., Abrahamson, M. and Hewitt, E. L.** (2000). Functional effects of the inhibition of the cysteine protease activity of the major house dust mite allergen Der p 1 by a novel peptide-based inhibitor. *Clinical and Experimental Allergy* **30**, 784–793.

- Kent, N. A., Hill, M. R., Keen, J. N., Holland, P. W. H. and Hart, B. J.** (1992). Molecular characterization of group I allergen *Eurm 1* from house dust mite *Euroglyphus maynei*. *International Archives of Allergy and Immunology* **99**, 150–152.
- Kenyon, F. and Knox, D.** (2002). The proteinases of *Psoroptes ovis*, the sheep scab mite – their diversity and substrate specificity. *Veterinary Parasitology* **105**, 317–325.
- Kenyon, F., Welsh, M., Parkinson, J., Whitton, C., Blaxter, M. L. and Knox, D. P.** (2003). Expressed sequence tag survey of gene expression in the scab mite *Psoroptes ovis* – allergens, proteases and free-radical scavengers. *Parasitology* **126**, 451–460.
- Lee, A. J., Machell, J., van den Broek, A. H. M., Nisbet, A. J., Miller, H. R. P., Isaac, R. E. and Huntley, J. F.** (2002). Identification of an antigen from the sheep scab mite, *Psoroptes ovis*, homologous with house dust mite group I allergens. *Parasite Immunology* **24**, 413–422.
- Loukas, A., Bethony, J. M., Williamson, A. L., Goud, G. N., Mendez, S., Zhan, B., Hawdon, J. M., Bottazzi, M. E., Brindley, P. J. and Hotez, P. J.** (2004). Vaccination of dogs with a recombinant cysteine protease from the intestine of canine hookworms diminishes the fecundity and growth of worms. *Journal of Infectious Diseases* **189**, 1952–1961.
- Nisbet, A. J. and Billingsley, P. F.** (2000). A comparative survey of the hydrolytic enzymes of parasitic and free-living mites. *International Journal for Parasitology* **30**, 19–28.
- Nisbet, A. J. and Huntley, J. F.** (2006). Progress and opportunities in the development of vaccines against mites, fleas and myiasis-causing flies of veterinary importance. *Parasite Immunology* **28**, 165–172.
- Nisbet, A. J., Huntley, J. F., MacKellar, A., Sparks, N. and McDevitt, R.** (2006). A house dust mite allergen homologue from poultry red mite *Dermanyssus gallinae* (De Geer). *Parasite Immunology* **28**, 401–405. doi:10.1111/j.1365-3024.2006.00862.x
- Pernas, M., Sánchez-Monge, R., Gómez, L. and Salcedo, G.** (1998). A chestnut seed cystatin differentially effective against cysteine proteinases from closely related pests. *Plant Molecular Biology* **38**, 1235–1242.
- Pettit, D., Smith, W. D., Richardson, J. and Munn, E. A.** (2000). Localisation and characterisation of ovine immunoglobulin within the sheep scab mite, *Psoroptes ovis*. *Veterinary Parasitology* **89**, 231–239.
- Redmond, D. L. and Knox, D. P.** (2004). Protection studies in sheep using affinity-purified and recombinant cysteine proteinases of adult *Haemonchus contortus*. *Vaccine* **22**, 4252–4261.
- Redmond, D. L. and Knox, D. P.** (2006). Further protection studies using recombinant forms of *Haemonchus contortus* cysteine proteinases. *Parasite Immunology* **28**, 213–219.
- Rees, J. A., Carter, J., Sibley, P. and Merrett, T. G.** (1992). Localisation of the allergen Der p 1 in the gut of the house dust mite *Dermatophagoides pteronyssinus* by ImmuStain. *International Archives of Allergy and Applied Immunology* **94**, 365–367.
- Sánchez-Ramos, I., Hernández, C. A., Castañera, P. and Ortego, F.** (2004). Proteolytic activities in body and faecal extracts of the storage mite, *Acarus farris*. *Medical and Veterinary Entomology* **18**, 378–386.
- Sasaki, T., Kikuchi, T., Yumoto, N., Yoshimura, N. and Murachi, T.** (1984). Comparative specificity and kinetic studies on porcine calpain I and calpain II with naturally occurring peptides and synthetic fluorogenic substrates. *Journal of Biological Chemistry* **259**, 12 489–12 494.
- Takai, T., Mineki, R., Nakazawa, T., Takaoka, M., Yasueda, H., Murayama, K., Okumura, K. and Ogawa, H.** (2002). Maturation of the activities of recombinant mite allergens Der p 1 and Der f 1, and its implication in the blockade of proteolytic activity. *FEBS Letters* **531**, 265–272.
- van den Broek, A. H. and Huntley, J. F.** (2003). Sheep scab: the disease pathogenesis and control. *Journal of Comparative Pathology* **128**, 79–91.
- van den Broek, A. H., Huntley, J. F., Machell, J., Taylor, M., Bates, P., Groves, B. and Miller, H. R. P.** (2000). Cutaneous and systemic responses during primary and challenge infestations of sheep with the sheep scab mite, *Psoroptes ovis*. *Parasite Immunology* **22**, 407–414.
- van den Broek, A. H., Huntley, J. F., Machell, J., Taylor, M. and Miller, H. R. P.** (2003). Temporal pattern of isotype-specific antibody responses in primary and challenge infestations of sheep with *Psoroptes ovis* – the sheep scab mite. *Veterinary Parasitology* **111**, 217–230.
- van Oort, E., de Heer, P. G., van Leeuwen, W. A., Derksen, N. I., Muller, M., Huveneers, S., Aalberse, R. C. and van Ree, R.** (2002). Maturation of *Pichia pastoris*-derived recombinant pro-Der p 1 induced by deglycosylation and by the natural cysteine protease Der p 1 from house dust mite. *European Journal of Biochemistry* **269**, 671–679.
- Yasuhara, T., Takai, T., Yuuki, T., Okudaira, H. and Okumura, Y.** (2001). Biologically active recombinant forms of a major house dust mite group 1 allergen Der f 1 with full activities of both cysteine protease and IgE binding. *Clinical and Experimental Allergy* **31**, 116–124.