



Spontaneously arising disease

Morphometry, cellular characterization and temporal evolution of granulomas induced by aluminium oxyhydroxide in sheep

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ABSTRACT

Persistent subcutaneous granulomas form at the injection site following administration of aluminium oxyhydroxide (AIOOH), a widely used vaccine adjuvant. Small ruminant lentiviruses (SRLVs) can infect macrophages within granulomas induced by commercial AIOOH-based vaccines in sheep. The entry of SRLVs into target cells involves the mannose receptor (MR), while catalytic polypeptide-like 3 protein containing Z1 domain (A3Z1) is considered a restriction factor for lentiviral replication. The objective of this study was to investigate the temporal evolution of AIOOH-induced post-vaccination granulomas in sheep experimentally infected with SRLVs. Twenty-four male lambs underwent two identical vaccination protocols and were challenged with SRLVs. Granulomas were detected in vaccinated groups only and progressively decreased in size. At post-mortem examination, 91.3% of the granulomas were recovered. Fistulas were present in granulomas following the second vaccination protocol. Central necrosis was present in 58.0% of granulomas and was associated with the vaccine used. Orthokeratotic keratin was seen within granulomas in 47.1% of the lambs. Considering all granulomas studied, significantly higher expression of MR was found compared with A3Z1. Differences in MR expression were related to the type of vaccine and the time since vaccination. A3Z1 expression was upregulated in granulomas from the infected groups. Macrophage polarization may influence SRLV infection of granulomas. While SRLV infection does not influence the architecture of post-vaccination granulomas, it may modulate their immune microenvironment. Further studies are needed to elucidate the complex interactions between AIOOH-induced granulomas and SRLV infection in sheep.

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1. Introduction

Adjuvants induce, enhance and maintain specific immune responses against inactivated antigens in vaccines [1]. Aluminium oxyhydroxide (AIOOH) is one of the most widely used adjuvants in licensed human and veterinary vaccines [2], yet its basic mechanism of action is poorly understood [3]. AIOOH is known to activate

innate and adaptive immune cells, leading to a Th2 biased response to vaccine antigens [3], and it can upregulate the expression of antigen-presenting and co-stimulatory molecules on cells of monocyte–macrophage lineage by an interleukin-4-dependent mechanism [4,5]. Internalized AIOOH can damage lysosomal membranes, triggering cathepsin-mediated cell death [6]. The cytotoxic effects of AIOOH activate the immune system through damage-associated molecular patterns (DAMPs), such as uric acid [7] and DNA [8]. Furthermore, this adjuvant promotes the recruitment of antigen-presenting cells (APCs) to the injection site and their systemic biodistribution [9–11].

Sheep health programmes include vaccination to prevent and control infectious diseases [12]. Consequently, multiple vaccines

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are often administered to a single animal each year depending on production targets, disease prevalence and geographical region [13]. The administration of AIOOH-based vaccines in sheep induces granulomas at the injection site, consisting of macrophages and fewer multinucleated giant cells (MGCs), lymphocytes and neutrophils [14]. AIOOH particles can persist at the injection site and be systemically distributed to distant tissues, including regional lymph nodes and the central nervous system [14,15].

Macrophages are plastic cells that display diverse phenotypes and functional responses under specific stimuli [16]. Two major pathways of macrophage activation are recognized: classically activated macrophages (M1) that promote inflammation and, alternatively activated macrophages (M2) that are involved in the resolution of inflammation [17,18]. Macrophage polarization is a key factor in the outcomes of infectious diseases caused by different pathogens, including porcine arteriviruses (eg, porcine reproductive and respiratory syndrome virus) [19], human lentiviruses (eg, human immunodeficiency virus) [20], small ruminant lentiviruses (SRLVs) [21] and ruminant mycobacteria [22–24].

In sheep, M1-like stimuli may enhance the expression of interferon-induced restriction factors, such as the apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 3 (APOBEC3), while M2-like stimuli are associated with the upregulation of mannose receptor (MR) and dendritic cell-specific intercellular adhesion molecule-3-grabbing non-integrin (DC-SIGN) [21]. APOBEC3 proteins containing Z1 domain (A3Z1) are restriction factors highly expressed in SRLV-resistant cells, such as monocytes and M1 macrophages [25]. In contrast, the MR expressed on M2 macrophages is proposed to be a key mediator for SRLV entry to target cells and tissue distribution [21,26]. Therefore, macrophage polarization in AIOOH-induced granulomas may influence SRLV infection and replication. Most knowledge about the polarization of SRLV-infected cells has come from in-vitro studies of blood monocyte-derived macrophages [21,25–27], with no information available on this process in in-vivo developed post-vaccination granulomas.

SRLVs cause progressive multisystemic inflammatory disease [28], directly affecting the welfare, production [29,30] and global trade of small ruminants [31]. The main target organs of SRLVs include the lungs, central nervous system, joints and mammary gland [32]. Currently, there are no treatments or vaccines to control SRLV infection, and attempts to develop an effective vaccine have failed [33]. However, SRLVs can infect granulomas induced by commercial AIOOH-based vaccines against various ovine pathogens, such as *Clostridium* spp, bluetongue virus and *Chlamydia abortus* [34].

Granuloma formation is a constant finding after AIOOH-based vaccine injections in sheep [35]. However, there is a notable knowledge gap regarding their evolution, cellular composition and potential interactions with infectious pathogens. The aim of this study was to investigate the temporal evolution of AIOOH-induced post-vaccination granulomas in lambs experimentally infected with SRLVs and to provide insights into factors that may influence their morphometry and composition, including SRLV infection.

2. Material and methods

2.1. Animals and experimental groups

The experimental design and all procedures were approved by the Ethical Committee of the University of Zaragoza (PI42/18) and conducted in accordance with Spanish Policy (Royal Decree 53/2013), which implements Directive 2010/63/EU of the European Parliament for the protection of animals used for scientific purposes. Twenty-four male Rasa Aragonesa lambs were sourced from a long-term SRLV-seronegative flock ('Masía El Chantre', Spain).

Newborn lambs and their ewes were kept in isolated pens and their seronegative status was confirmed with three commercial ELISA tests: (1) Eradikit SRLV Screening kit (In3Diagnostic, www.in3diagnostic.com); (2) Elitest MVV/CAEV (Hyphen Biomed, www.hyphen-biomed.com); and (3) INgezim Maedi screening (Eurofins Technologies Ingenasa, www.eurofins.com). Three-month-old post-weaned lambs were housed at the experimental facilities of the University of Zaragoza and divided into five treatment groups. At 6 months, the animals underwent an experimental protocol involving the subcutaneous administration of two commercial AIOOH-based vaccines or a saline solution, combined with a challenge with SRLVs or uninfected culture medium (Table 1).

2.2. Study design: vaccination schedule and small ruminant lentivirus challenge

All lambs in the vaccinated groups (1, 3 and 5) received two different commercial AIOOH-based vaccines: vaccine 1 against bluetongue virus and vaccine 2 against members of the Pasteurellaceae and Clostridiaceae families (Supplementary Table 1). The first vaccination protocol included a 2 ml dose of each vaccine at 0 days post treatment (dpt) and again at 30 dpt. This protocol was repeated at 135 and 165 dpt, resulting in each lamb receiving a total of eight vaccine doses (Supplementary Fig. S1). Lambs in each control group (2 and 4) received an identical number and volume of saline solution injections on the same dates. The anatomical locations of the injections are detailed in Supplementary Fig. 2.

The challenge with the previously used SRLVs strain ov496 [36] was carried out after completion of the first vaccination schedule, at 36 dpt. The same protocol was repeated at 93 dpt to ensure infection of the lambs (Supplementary Fig. 1). A total of 10⁶ tissue culture infectious dose (TCID₅₀) per animal was inoculated either intravenously (IV) or intratracheally (IT), depending on the group (Table 1). Lambs in group 5 were inoculated via the IT (n = 3) or IV (n = 3) route with non-infected culture medium (Gibco Dulbecco's Modified Eagle Medium [DMEM]; Thermo Fisher Scientific, www.thermofisher.com). Two animals, one from group 2 and one from group 5, died due to anaesthesia during the IT infection procedure. Therefore, a total of 22 lambs completed the study, which lasted about 8.5 months.

2.3. In-vivo studies

The evaluation of morphometric changes in AIOOH-induced post-vaccination granulomas was based on shape, as well as length and width measurements obtained with an automated calliper at 7, 16 and 35 days after injection. Due to the ellipse-like shape of the

Table 1
Experimental design

Groups	n	Vaccines/control	Viral challenge/control	Viral infection route
Group 1 AIOOH vaccines and SRLV IT	6	AIOOH vaccines	SRLV	IT
Group 2 SRLV IT only	2	Saline solution	SRLV	IT
Group 3 AIOOH vaccines and SRLV IV	6	AIOOH vaccines	SRLV	IV
Group 4 SRLV IV only	3	Saline solution	SRLV	IV
Group 5 AIOOH vaccines only	5	AIOOH vaccines	Non-infected culture medium	IT or IV

AIOOH, aluminium oxyhydroxide; IT, intratracheal; IV, intravenous; n, number of lambs per group at end of experiment; SRLV, ov496 strain of small ruminant lentivirus.

post-vaccination granulomas, Ramanujan's formula ($P \approx \pi [3(a+b) \sqrt{((3a+b)(a+3b))}]$) was used to estimate the perimeter [37].

2.4. Post-mortem studies

Lambs were humanely killed with an overdose of IV pentobarbital and a complete necropsy was performed. However, this report focuses exclusively on the post-vaccination granulomas. During the experiment, nine granulomas were biopsied (results to be described elsewhere) and only 127 out of 136 granulomas were expected to be detected at post-mortem examination. Animals were placed in lateral recumbency, the granulomas were located by palpation and the presence of fistulas was recorded. Skin and adipose tissue were removed to expose the subcutaneous tissue, and all post-vaccination granulomas found were sampled. Sections of each granuloma were trimmed and fixed in 10% neutral-buffered formalin, while a selection of granulomas ($n = 50$, Supplementary Table 2) were collected for molecular studies by immersion in RNAlater solution (Thermo Fisher Scientific) and stored at -20°C until processing. Formalin-fixed samples were routinely processed through graded alcohols, embedded in paraffin wax and $4 \mu\text{m}$ sections cut for haematoxylin and eosin (HE) staining and immunohistochemical procedures.

2.5. Pathological studies

Histological characterization of AIOOH-induced granulomas was based on a modified semiquantitative scoring system [14], as detailed in Supplementary Table 3. Immunohistochemical studies employed the avidin–biotin–peroxidase complex method (Vectastain Elite ABC Kit; Vector Laboratories, <https://vectorlabs.com>) with 3,3'-diaminobenzidine (Vector Laboratories) as chromogen. Details of the primary antibodies and technical aspects are summarized in Table 2. In brief, sections were subjected to specific pre-treatments for antigen retrieval. Non-specific binding sites were blocked by incubating sections at room temperature for 1 h with normal serum, obtained from the species corresponding to the secondary antibody, diluted in a solution of 1% polyvinylpyrrolidone, 1% bovine serum albumin and 1% tris-buffered saline (PVP-BSA-TBS). Ionized calcium-binding protein molecule 1 (Iba1) antibody, a pan-macrophage marker, was used for all granulomas ($n = 112$). For T lymphocytes (anti-CD3) and viral (anti-p28) detection, two granulomas from four different lambs in each vaccinated group (1, 3 and 5, $n = 24$) were investigated as described [38–40]. Ovine lymph nodes were used as positive controls for Iba1 and CD3 analyses, while SRLV-affected lungs were used for p28 detection. Granulomas from group 5 were included as SRLV-negative controls. The specificity of the technique was demonstrated by replacing the primary antibody with PVP-BSA-TBS solution and, in previous studies, by using isotype-matched antibodies against irrelevant antigens [38–40].

Table 2

Antibodies used for immunolabelling of aluminium oxyhydroxide-induced post-vaccination granulomas in experimental lambs

Antibody	Host	Type	Pre-treatment	Dilution	Incubation	Source
Anti-p28 (clone CAEP5A1)	Mouse	M	Microwave, citrate buffer, pH6	1:400	Overnight (room temperature)	VMRD (www.vmrld.com)
Anti-CD3	Rabbit	P	Pressure cooker, citrate buffer, pH6	1:100	Overnight (4°C)	Agilent Dako (www.agilent.com)
Anti-Iba1	Rabbit	P	Microwave, citrate buffer, pH6	1:300	Overnight (4°C)	FUJIFILM Wako Pure Chemical Corporation (www.fujifilm.com)

M, monoclonal; P, polyclonal.

2.6. Relative target gene expression in granulomas

Twenty mg of each granuloma were homogenized using steel beads in a vibration grinding mill (Mikro-Dismembrator U; Sartorius AG, www.sartorius.com), followed by RNA extraction and purification using a magnetic bead-based method (NucleoMag RNA-MagnetaPure32; MACHEREY-NAGEL, www.mn-net.com). The quantity and purity of the extracted nucleic acids were evaluated with a microvolume ultraviolet-visible spectrophotometer (Nano-Drop One/One; Thermo Fisher Scientific). A total of 1,000 ng of RNA was retrotranscribed into cDNA using the PrimeScript RT Reagent kit (Takara Bio, www.takarabio.com) with a single thermal cycle at 37°C for 15 min, followed by a cycle at 85°C for 15 s. Quantitative analyses of target genes (*A3Z1*, *DC-SIGN* and *MR*) were conducted in a total volume of 25 μl , using 5 μl of cDNA, DNA-binding dye (SYBR Green Master Mix; Bio-rad, www.biorad.com) and specific primers previously used in sheep samples [21]. β -actin was used as the housekeeping gene and no template controls were included in each run. Quantitative polymerase chain reactions (qPCRs) and data analyses were performed using the AriaMx Real-time PCR System (Agilent Technologies, www.agilent.com). Relative mRNA quantifications were determined by subtracting the β -actin Ct value from the target gene Ct value in each sample (ΔCt) and were expressed as $2^{-\Delta\text{Ct}}$ value [41].

2.7. Statistical analyses

Quantitative variables (eg, perimeter of granulomas and relative target gene expression) were tested for normality and equality of variance using the Shapiro–Wilk test and Levene's test, respectively. Unpaired comparisons between categorical variables with three or more categories (eg, treatment group) were analysed with the Kruskal–Wallis test, followed by Dunn's test as post-hoc test (non-normally distributed data). For variables with two categories (eg, type of vaccine), comparisons were performed using the Mann–Whitney U test (non-normally distributed data) or Student's t-test (normally distributed data). Paired comparisons within each categorical variable between sampling dates were carried out with the Wilcoxon test (non-normally distributed data) or Student's t-test (normally distributed data) [42]. Detection of fistulas and microscopic features were coded as binary variables (0 and 1, representing presence [P] and absence [A], respectively). The P/A of these variables was described using relative frequencies. Associations between qualitative and categorical variables were analysed using Pearson's Chi-square test; alternatively, Fisher's exact probability test was applied for 2×2 contingency tables and Likelihood Ratio test for the rest of the cases [43]. Data were analysed using SPSS Statistics 26.0 software (IBM Corporation, www.ibm.com), with a P value <0.05 considered statistically significant. Graphs were created by GraphPad Prism version 8.0 software (GraphPad, www.graphpad.com).

3. Results

3.1. Morphometric evolution of granulomas

Post-vaccination granulomas were detected only in the vaccinated groups (1, 3 and 5) as well-demarcated, soft and mobilizable subcutaneous nodules. Granulomas were absent in the groups (2 and 4) injected with saline solution. A significant ($P < 0.050$) and progressive decrease in granuloma size was seen during the first 35 days post vaccination, independent of the type of vaccine (Fig. 1). However, granulomas induced by vaccine 2 had a significantly greater mean perimeter than those induced by vaccine 1 at 7, 16 and 35 days post vaccination (Fig. 1). Only subtle differences in granuloma shape were seen; most granulomas that developed following vaccine 1 were plaque-like, while those from vaccine 2 tended to bulge from the skin (Fig. 2A).

3.2. Pathological characterization

In-vivo evaluation of injection sites revealed minimal macroscopic changes associated with ALOOH-induced granulomas; the skin over some granulomas was alopecic (Fig. 2A) and/or exhibited a communication with the outer skin (fistula) (Fig. 2A inset). A total of 116 out of 127 post-vaccination granulomas (91.3%) were recovered from the subcutaneous tissue at post-mortem examination (Fig. 2B). At this time, fistulas were observed in nine out of 116 (7.8%) granulomas. Presence of fistulas was associated ($P = 0.012$, Fisher exact test) with vaccine type, being predominantly found in granulomas induced by vaccine 2 (8/9; 88.9%) compared with vaccine 1 (1/9; 11.1%). All fistulas (9/9) occurred in granulomas induced by the second vaccination protocol (at 135 and 165 dpt), with none observed after the first vaccination protocol (at 0 and 30 dpt).

For histopathological analyses, only 112 out of the 116 granulomas were evaluated, as four were minimally sized. ALOOH-induced granulomas were a highly cellular inflammatory reaction, primarily composed of epithelioid macrophages mixed with MGCs, surrounded by varying amounts of fibrous connective tissue. Over half of the ALOOH-induced granulomas (65/112; 58.0%) contained a variably sized area of central necrosis (Fig. 2C). Necrosis was associated ($P < 0.001$, Pearson's chi-square test) with the type of vaccine, occurring more frequently in granulomas from vaccine 1 (43/59; 72.9%) than from vaccine 2 (22/53; 41.5%). Foci of mineralization were observed in 35 out of 112 (31.3%) scattered throughout necrotic areas (Fig. 2C). Necrosis was also linked ($P = 0.007$, Pearson's chi-square test) to the presence of lymphoid aggregates but was independent of the time post vaccination ($P = 0.615$, Pearson's

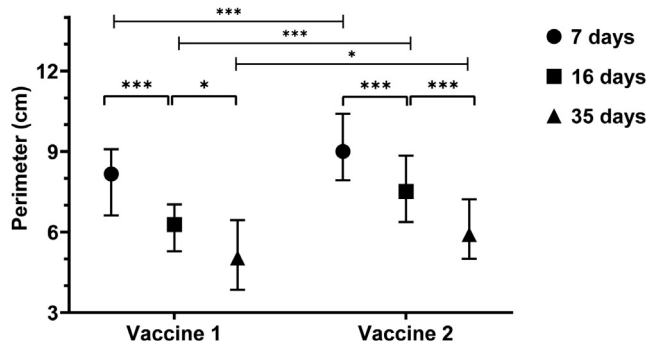


Fig. 1. Perimeter (cm) of post-vaccination granulomas following vaccines 1 and 2. Median values represented as circles (7 days post treatment, dpt), squares (16 dpt) and triangles (35 dpt). Whisker bars indicate interquartile range. *, statistical significance ($P < 0.05$); ***, statistical significance ($P < 0.001$).

chi-square test). A total of 55 out of 112 (49.1%) granulomas contained lymphoid cells arranged in various patterns (Supplementary Fig. 3A–C): (i) linear groups parallel to necrotic areas, (ii) follicular-like structures, and (iii) random interspersions with macrophages. Two of these patterns often co-occurred in the same granuloma. No statistical association was found between the presence of lymphoid aggregates and the type of vaccine or dpt ($P = 0.127$ and $P = 0.438$, Pearson's chi-square test, respectively).

Neutrophils were found in 25 out of 112 (22.3%) granulomas. Their presence was associated with dpt ($P = 0.046$, Pearson's chi-square test), being more common in granulomas formed at 135 and 165 dpt (18/61; 29.5%) compared with those at 0 and 30 dpt (7/51; 13.7%). The presence of neutrophils was also linked ($P = 0.008$, Pearson's chi-square test) to the type of vaccine, and these cells were found in a higher proportion of granulomas following vaccine 1 (19/59; 32.2%) compared with vaccine 2 (6/53; 11.3%). Scant to moderate layers of orthokeratotic keratin were seen between the granulomatous reaction and the necrotic content in nine out of 112 (8.0%) granulomas (Fig. 2D). Overall, eight out of 17 (47.1%) lambs had at least one granuloma with keratin. The presence of keratin was associated ($P = 0.004$, Fisher exact test) with dpt, as it was only found in 135 and 165 dpt granulomas. Keratin layers were occasionally lined by stratified squamous epithelium, with a small follicular-like cyst noted in one post-vaccination granuloma (Fig. 2D inset).

3.3. Immunohistochemical studies

Epithelioid macrophages and MGCs within granulomas had strong, membranous Iba1 labelling (Fig. 3A). CD3-positive lymphocytes exhibited cytoplasmic to membranous labelling and they were located at the periphery of the necrotic tissue (Fig. 3B) or formed lymphoid aggregates (Fig. 3C). SRLV protein was detected in the cytoplasm of macrophages and occasionally in MGCs of all examined granulomas from experimentally infected lambs, regardless of infection route (Fig. 3D). Granulomas from non-SRLV-infected lambs were consistently devoid of labelling.

3.4. Relative target gene expression in granulomas

MR expression was significantly higher ($P < 0.001$) than A3Z1 expression when granulomas from all vaccinated groups (1, 3 and 5) were considered. Differences in A3Z1 expression were found between groups (Fig. 4A): A3Z1 was significantly ($P = 0.020$) upregulated in granulomas from the infected group 3 compared with those from the non-infected group 5. Similarly, A3Z1 was upregulated, albeit not significantly ($P = 0.064$), in granulomas from the infected group 1 compared with those from group 5. There were no differences in A3Z1 expression between infected groups (1 and 3). Similarly, no significant differences in A3Z1 expression were observed in granulomas induced by the two vaccines or at different dpt (Supplementary Fig. S4). MR expression in granulomas was similar between all groups (Fig. 4A). However, MR expression was significantly ($P = 0.020$) higher in granulomas induced at 135 dpt by vaccine 1 compared with those induced by vaccine 2, with no significant differences in granulomas induced at 0 dpt (Fig. 4B). As DC-SIGN expression was minimal, further analyses using this gene were not performed.

4. Discussion

This study provides new insights into the histopathological features of ALOOH-induced post-vaccination granulomas in sheep. Advancing our understanding of the mechanisms underlying these chronic inflammatory reactions and their interactions with

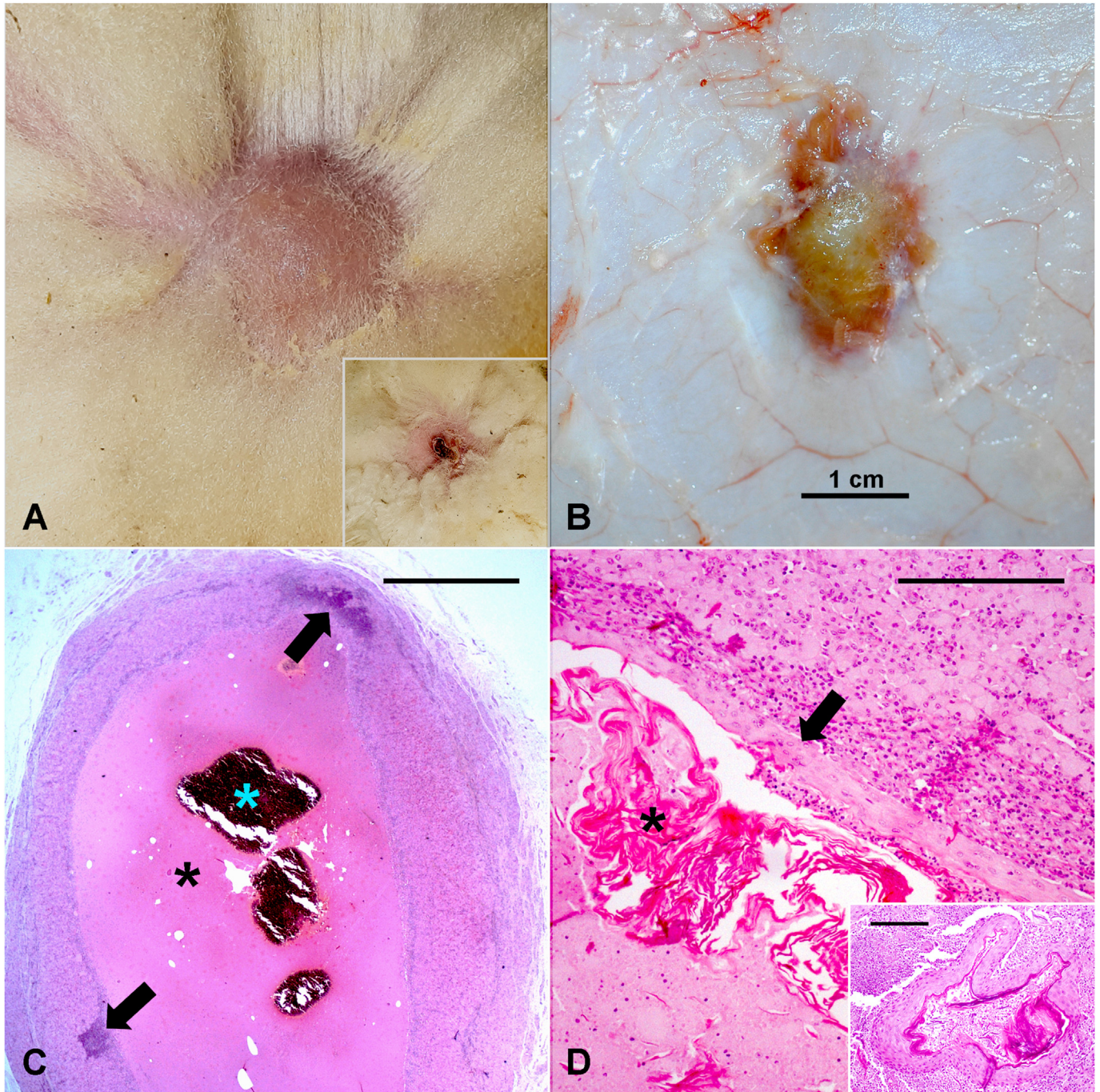


Fig. 2. Aluminium oxyhydroxide-induced post-vaccination granulomas, sheep. (A) In-vivo examination, vaccine 2. Well-demarcated, rounded and bulging granuloma (approximately 3–4 cm) covered by alopecic skin. Inset: fistula in a granuloma covered by a crust. (B) Post-mortem examination, vaccine 2. Granuloma in subcutaneous tissue following adipose tissue removal. (C) Granuloma architecture. Layers of macrophages with multifocal lymphoid aggregates (arrows) surrounding central necrotic area (black asterisk) with foci of mineralization (blue asterisk). HE. Bar, 300 µm. (D) Keratin layers (asterisk) lined by stratified epithelium (arrow) at periphery of necrotic area. HE. Bar, 300 µm. Inset: small follicular-like cyst in another granuloma. Bar, 500 µm.

lentiviruses is crucial, as granuloma formation is a constant local side effect following vaccination with AlOOH-based adjuvants.

In the present study, 91.3% of the expected granulomas were found at the end of the study, indicating a high persistence of these inflammatory reactions at the injection site. AlOOH-induced granulomas reached their maximum size at 7 days post vaccination, aligning with findings in mice in which inflammatory cell recruitment also peaked at this time [44]. The size reduction found in our study may be related to phagocytosis of AlOOH particles and

their migration to distant sites, as reported in sheep [14,15], rabbits [45] and mice [10,11]. In our study, granulomas induced by vaccine 2 were larger and more conspicuous than those from vaccine 1, probably due to the nearly double AlOOH content in vaccine 2 (Supplementary Table 1). However, Asín *et al* [14] noted that in sheep, larger granulomas were induced by commercial vaccines containing longer aluminium particles compared with those induced by the injection of an equal amount of AlOOH alone. This finding suggests the probable influence of other factors such as the

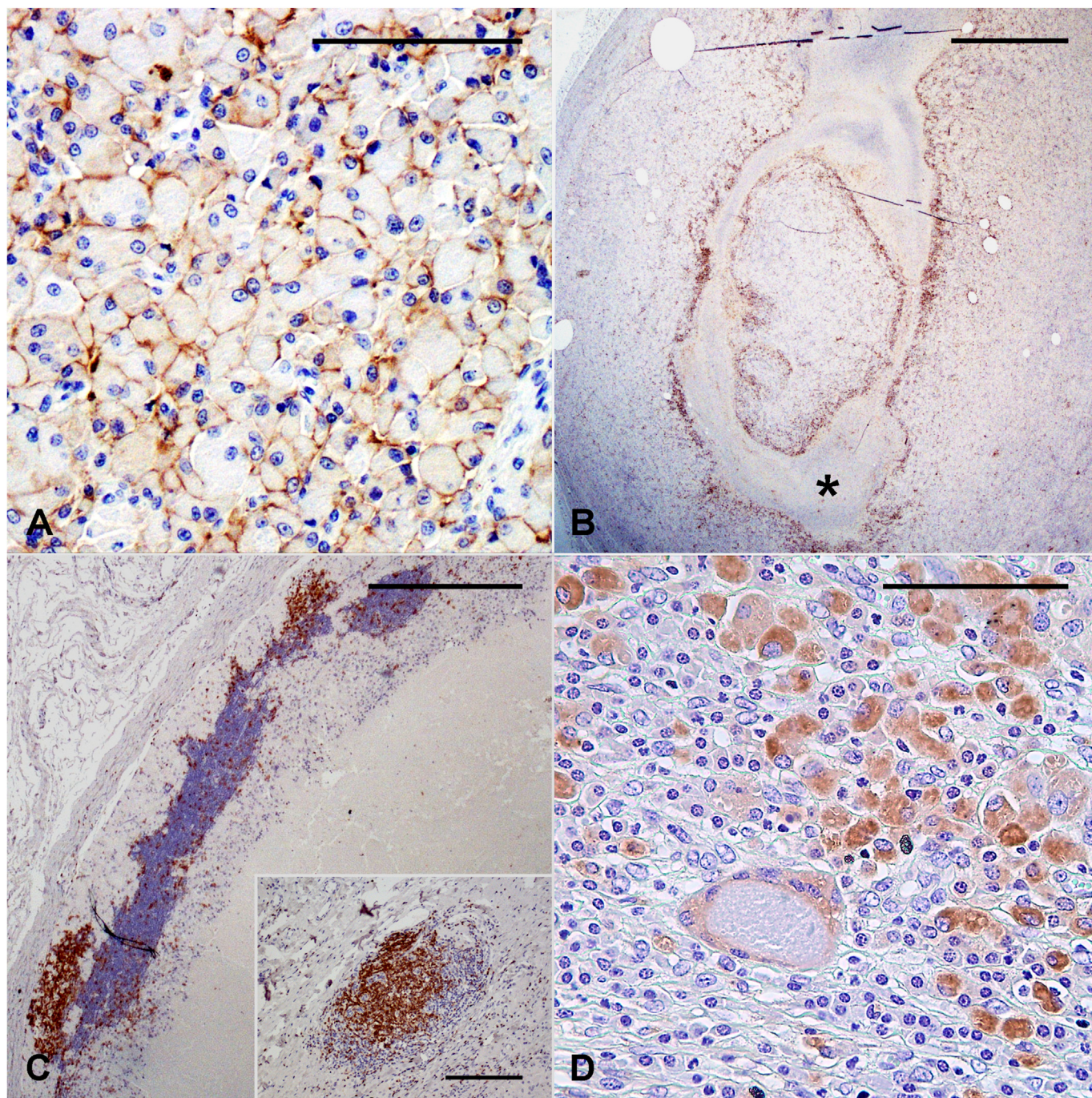


Fig. 3. Aluminium oxyhydroxide-induced post-vaccination granulomas, sheep, immunohistochemistry. (A) Iba1 immunolabelling of cell membranes of macrophages and multinucleated giant cell (MGCs). Bar, 50 μ m. (B) Dense band of CD3-positive lymphocytes border necrotic area (asterisk). Bar, 300 μ m. (C) CD3-positive lymphocyte aggregates at periphery of band of CD3-negative cells. Bar, 100 μ m. Inset: follicular lymphoid aggregate mainly composed of CD3-positive lymphocytes. Bar, 500 μ m (D) Granular to diffuse positive labelling for SRLV capsid protein p28 in cytoplasm of macrophages and MGCs. Bar, 50 μ m.

physicochemical properties of AlOOH–antigen complexes [46] or the effects of other components in vaccine formulations on the morphometry of granulomas.

Fistulas were primarily seen in vaccine 2 granulomas, probably due to their larger size and increased pressure in the subcutis, which may favour the discharge of necrotic content to the exterior. While fistulation of post-vaccination granulomas has been reported in sheep [14], our results indicate it is an uncommon event, mainly occurring in the first months after granuloma formation. The presence of necrotic areas may be also linked to the

physicochemical properties of the AlOOH–antigen complexes, rather than to higher AlOOH content, as they were more frequently found in granulomas induced by vaccine 1, which had a lower AlOOH concentration (Supplementary Table 1). Necrosis in granulomas was also associated with the presence of lymphoid aggregates. Indeed, AlOOH-induced cytotoxicity leads to the release of molecules recognized as DAMPs [7,8], which promote the activation of APCs and cytotoxic T lymphocytes, potentially driving so-called immunogenic cell death [47]. Neutrophils were more frequently seen in granulomas aged 3–4 months, declining thereafter, which

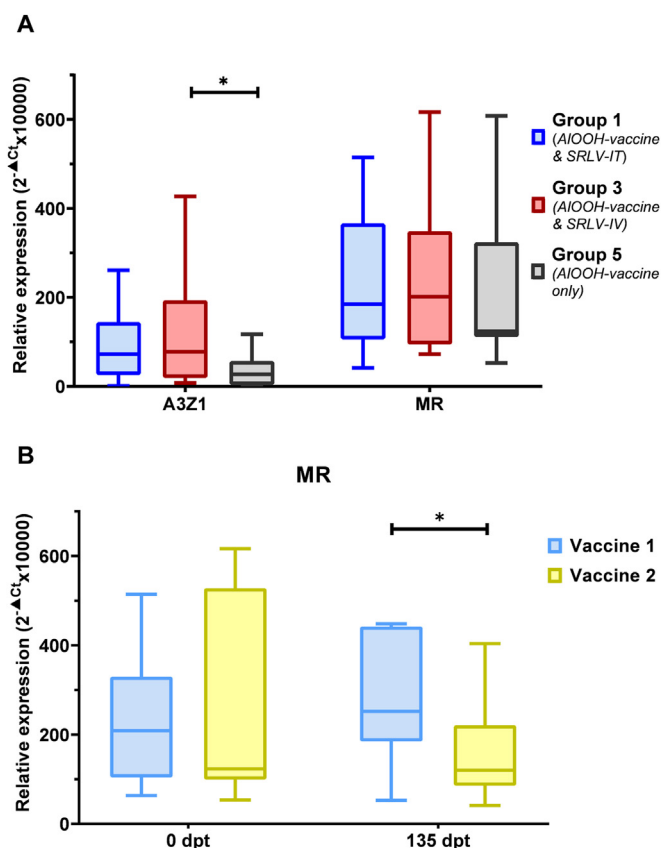


Fig. 4. Relative expression of target genes. (A) *APOBEC3Z1* (*A3Z1*) and mannose receptor (*MR*) expression in post-vaccination granulomas from vaccinated lambs. (B) *MR* expression in post-vaccination granulomas induced by vaccine 1 and vaccine 2 at 0 days post treatment (dpt) and 135 dpt. Data presented as $2^{-\Delta Ct} \times 10,000$ normalized to β -actin. Box plots represent median and quartiles, with whiskers indicating minimum and maximum values. *, statistical significance ($P < 0.050$). SRLV-IT, SRLV administered by intratracheal route; SRLV-IV, SRLV administered by intravenous route.

aligns to observations in mice [44]. The presence of keratin constituted a novel finding in sheep, observed exclusively in granulomas aged 3–4 months. Skin trauma and/or the expansion of the granulomatous reaction may cause hair entrapment, potentially resulting in the formation of follicular cysts, as suggested in dogs, cats [48] and sheep [49]. Follicular cyst formation has been reported in rabbits following intradermal injections [50,51] and in humans after vaccination [52]. Remarkably, human pilomatrixomas have also been linked to vaccine injections [53–56]. Further research is required to determine whether post-vaccination granulomas in sheep influence the development of certain skin conditions.

Our results indicate that SRLV infection does not influence the cellular architecture of post-vaccination granulomas but may modulate their immune microenvironment. The upregulation of *MR* found in ovine AIOOH-induced granulomas suggests a polarization of macrophages toward an M2-like phenotype, which may favour SRLV infection of these cells [26]. Interestingly, 4-month-old granulomas induced by vaccine 1 had significantly higher expression of *MR*, indicating that macrophage polarization may be influenced by the time that has elapsed since vaccine injection and the type of vaccine. Furthermore, vaccine 1 also produced more granulomas with necrosis, and *MR* expression is known to be induced following cellular death, as this receptor is involved in immune clearance [57]. In our study, *MR* expression was not altered by SRLV infection, consistent with previous in-vitro observations [25].

However, we detected an upregulation of *A3Z1* expression in granulomas from infected sheep, which contradicts in-vitro findings [25]. Our result may be linked to indirect activation mechanisms that stimulate *APOBEC3* expression via lentiviral-induced signalling molecules, as reported in the target organs of feline immunodeficiency virus in cats [58].

5. Conclusion

The present study advances understanding of the pathogenesis of AIOOH-induced post-vaccination granulomas in sheep. Morphometry and cellular composition of ovine AIOOH-induced granulomas are influenced by different factors such as vaccine type, time since vaccine administration and potential SRLV infection. Further studies are needed to investigate whether macrophage polarization can influence SRLV replication or if SRLVs modulate the granuloma microenvironment.

Statement of author contributions

Ana Rodríguez-Largo: Conceptualization; Data curation; Investigation; Writing – original draft, review & editing. **Álex Gómez:** Methodology; Visualization; Writing – review & editing. **Estela Pérez:** Methodology; Writing – review & editing. **Irti Moncayola:** Investigation. **Lucia Biagini:** Investigation; Visualization; Writing – review & editing. **Giacomo Rossi:** Investigation; Data curation; Visualization; Writing – review & editing. **Ignacio de Blas:** Data curation; Formal analysis. **Antonio Fernández:** Data curation; Investigation; Writing – review & editing. **Marta Pérez:** Funding acquisition; Methodology; Supervision; Writing – review & editing. **Idoia Glaria:** Conceptualization; Investigation; Methodology; Writing – review & editing. **Ramsés Reina:** Funding acquisition; Investigation; Methodology; Supervision; Writing – review & editing. **Lluís Luján:** Conceptualization; Funding acquisition; Supervision; Writing – review & editing.

Declaration on use of generative AI or AI-assisted technologies in the writing process

The authors declared that no generative artificial intelligence (AI) or AI-assisted technologies were used in the writing of this manuscript.

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Declaration of competing interests

The authors declared no conflicts of interest in relation to the research, authorship or publication of this article.

Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.jcpa.2024.11.002>.

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