

***Mycobacterium tuberculosis* and *Mycobacterium bovis* derived proteins induce caspase-independent apoptosis in bovine macrophages**

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Abstract

Species of the genus Mycobacterium are capable of inducing cell apoptosis. Infected cells with *M. tuberculosis* undergo apoptosis through a caspase-dependent pathway. We have previously shown that whole *Mycobacterium bovis* cells and derived crude protein extracts were able to trigger apoptosis through a caspase independent mechanism. However, the identity of the protein or protein fractions capable of inducing apoptosis has not been determined. In this study, bovine macrophages were incubated in the presence of *Mycobacterium tuberculosis* recombinant proteins and *Mycobacterium bovis* protein extracts to identify proteins capable of inducing apoptosis. A subgroup of proteins and protein extracts from strains of the *Mycobacterium tuberculosis* complex were shown to induce DNA fragmentation in bovine macrophages through a caspase independent pathway. Our findings suggest that bovine macrophages may elicit a different response to mycobacterium infection than that displayed by murine or human cells.

Keywords: Apoptosis, *M. bovis*, *M. tuberculosis*, Bovine macrophages, Caspase.

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Introduction

Tuberculosis is responsible for the deaths of millions of people each year and is the second leading cause of death due to an infectious disease after the human immunodeficiency virus (HIV). Human tuberculosis is mainly caused by *Mycobacterium tuberculosis* (*M. tuberculosis*), however, between 3 and 10% of cases are due to *Mycobacterium bovis* (*M. bovis*), the etiological agent of bovine tuberculosis.¹ In addition to the potential risks for human health, *M. bovis* generates enormous economic losses in the cattle industry globally.²

Pathogenesis of mycobacteria relates to their ability to survive inside host cells, especially in immune cells such as macrophages and monocytes.³ In fact, mycobacteria can disrupt the activation and phagocytic processes of immune cells.⁴ Species from the *M. tuberculosis* complex (MTC) can, for instance, survive in macrophages by inhibiting acidification of the phagosomal compartment thereby preventing the phagosome-lysosome fusion; thus not only adapting to the intracellular environment of the macrophage, but turning it into their replication niche.⁵

Macrophage response to MTC infection mediates the adaptive immune defense mechanisms, thus playing a central role in the outcome of the disease.⁶ Intracellular pathogens can induce activation of cell death mechanisms, such as apoptosis.⁷ During apoptosis, a cell undergoes volume reduction, rupture and condensation of chromatin, and formation of apoptotic bodies, which are vesicles created by cell membrane blebbing, that express elimination signals on their surface. Apoptotic bodies are avidly recognized and phagocytosed by professional phagocytes in a process called efferocytosis.⁸ Apoptosis is intracellularly directed through a biochemical cascade that activates a group of enzymes called caspases.⁹ These enzymes can be activated through an intrinsic pathway (*i.e.* mitochondrial) and an extrinsic pathway (*i.e.* death receptor), both of which converge to a common terminal or execution pathway, with caspase 3 as the main effector.⁸ However, apoptosis can also occur in the absence of caspase activation.¹⁰

Apoptosis of infected cells may benefit the host, as it removes the intracellular niche for pathogen replication. However, apoptosis can also benefit the pathogen by eliminating host defense cells, hence allowing penetration of epithelial barriers and favoring spread of infection by efferocytosis.¹¹ When attenuated strains of *M. bovis* or *M. tuberculosis* (such as strains BCG or H37Ra respectively), are studied, apoptosis seems to improve control of the infection by reducing pathogen survival and limiting its intracellular growth. In contrast, virulent strains of *M. tuberculosis* decrease or inhibit apoptosis in infected human derived macrophages.¹² Previous studies by our group have shown that apoptosis induction in bovine macrophages by *M. bovis* is time and dose dependent¹³ and does not entail activation of caspases.¹⁴ Moreover, we have established that apoptosis can also be induced by a crude protein extract from *M. bovis*.¹⁴ However, the identity of the protein or protein fraction capable of inducing apoptosis has not been determined. Therefore, the aim of this study was to identify *Mycobacterium* derived proteins capable of inducing apoptosis, through assessing the effect of *Mycobacterium tuberculosis* recombinant proteins and *Mycobacterium bovis* protein extracts on bovine macrophage cultures.

Material and methods

Bacteria

A *M. bovis* AN5 reference strain, donated by Dr. Rogelio Hernández-Pando of the National Institute of Medical Sciences and Nutrition, was used. Bacteria were incubated for 8 weeks in Sauton medium enriched with 0.5% glycerol (as sole carbon source) and 4 g/L of L-asparagine (as sole amino acid source), at 37 °C without shaking. These culture conditions provided the greatest amount of proteins produced with minimal breakdown.

M. bovis protein extracts

The bacterial biomass from the *M. bovis* culture was filtered eight weeks after incubation using 0.45 and 0.22 µm pore membranes (Merk, Millipore, MA). The culture filtrate extract (CFE) was then precipitated with 0.5 g/ml of ammonium sulfate at 70% saturation and stirred for 24 hours at 4 °C. Proteins were subsequently obtained by centrifugation at 10,300 × g for 30 minutes and then dialyzed for 4 days using a 3,500 MWCO membrane (Spectra/Por).

The *M. bovis* biomass was resuspended in 1X PBS and sonicated at 71.25W in ice for 2 hours, with 5-minute pulses, followed by a one-minute rest (Virtis™). The total extract was then centrifuged at 10,000 × g for 15 minutes and the supernatant (soluble extract, SE), recovered.

Total proteins from both extracts (SE and CFE), were quantified in a microplate using the bicinchoninic acid technique (Micro BCA Pierce™). Proteins from a single sample of each extract were also resolved by SDS-PAGE at 12% and stained with Coomassie brilliant blue R-250.

M. tuberculosis recombinant proteins

M. tuberculosis recombinant proteins employed in this study have 99-100% of identity with their orthologous genes in *M. bovis* (Table 1).

Bovine peripheral blood monocytes

Macrophages derived from bovine monocytes were obtained by jugular puncture of a healthy adult donor from a tuberculosis free herd (not vaccinated, exposed or challenged), according to a method previously described by our laboratory.^{10,13,14} Only one donor was used to avoid individual variations in the *in vitro* experiments. Briefly, blood was collected in 60 ml syringes, pre-loaded with 8 ml of ACD. The leukocyte pack was subsequently obtained by centrifugation at 500 × g for 30 minutes, diluted 1:1 with a phosphate plus citrate solution, and placed in a 2:1 Percoll gradient (GE Healthcare Bio-Sciences AB) with a specific density of 1.077. It was then centrifuged again at 1200 × g for 30 minutes, the leukocyte interface recovered and then washed on two occasions with a solution of 45 ml phosphates plus citrate with added autologous plasma (30 ml and 15 ml, respectively). Finally, the cell pellet was resuspended in RPMI-1640 medium (Gibco® Cell Culture) with 4% of autologous serum to facilitate monocyte adhesion to the Teflon bottom Erlenmeyer flasks. After being seeded into flasks, monocytes were incubated for 24 hours at 37 °C with a humid atmosphere and 5% CO₂. Unbound cells were then removed

Table 1. Recombinant proteins used for apoptosis induction in bovine macrophages

Gene	Synonymy	Access No. Mtb/Mb (Identity %)*	Weight (kDa)	Function	Apoptosis
dnaK	hsp70	Rv0350/Mb0358 (100%)	66.83	Intracellular cytoplasmic chaperone. Possible ATPase activity.	NA
glnA1	glnA	Rv2220/Mb2244 (100%)	53.53	Glutamin-synthetase	NA
PE_PGRS33	-	Rv1818c/Mb1849c (99%)	40.75	Unknown function. It seems to influence surface interactions.	Caspase-dependent apoptosis in murine cells ¹⁹
	dPGRS33	-	-	-	-
	dPE	-	-	-	-
mce2A	mce2	Rv0589/Mb0604 (99%)	43.45	Invasine.	NA
pstS1	phoS1, phoS, Ag-38kDa	Rv0934/Mb0959 (100%)	38.21	Lipoprotein of unknown function.	Caspase-dependent apoptosis in human MDMs ²⁰
fbpB	Ag85-B, mpt59 85B	Rv1886c/Mb1918c (100%)	34.58	Micolyl-transferase	NA
Apa	Mpt32, modD	Rv1860/Mb1891 (99%)	32.72	Fibronectin binding glycoprotein.	NA
lprG	P27	Rv1411c/Mb1446c (100%)	24.54	Lipoprotein of unknown function.	NA
sigD	-	Rv3414c/Mb3448c (100%)	22.91	Probable alternative sigma factor	NA
hbhA	-	Rv0475/Mb0485 (100%)	21.53	Methylated protein with heparin binding and sulfate dextran.	Caspase-dependent apoptosis in murine and human macrophages ²¹
hspX	Acr	Rv2031c/Mb2057c (100%)	16.22	Alpha-crystalline	NA
lpqH	P-19kDa	Rv3763/Mb3789 (100%)	15.11	Lipoprotein secreted and associated with the cell wall.	Caspase dependent apoptosis in murine and human macrophages ¹⁷
esxA	esat-6	Rv3875/Mb3905 (100%)	9.90	Antigenic secretory protein with unknown function.	Apoptosis by ER stress ¹⁸

NA= Not associated to apoptosis in monocytes or macrophages.

Mtb= *Mycobacterium tuberculosis*, Mb= *Mycobacterium bovis*

*Identity percentage according to the Protein BLAST Alignment.

and a RPMI-1640 medium was added with 12% autologous serum. Monocytes were kept under the described conditions for a further 12 days. Cell viability was assessed by Trypan blue staining following recovery and before each trial. Experiments were performed with 95% cell viability.

Apoptosis induction in macrophages

For apoptosis induction, bovine macrophages were plated in Teflon flasks, using a cell density of 4×10^5 cells. Twenty-four hours after seeding, cells were incubated with 100 $\mu\text{g}/\text{ml}$ of either SE or CFE proteins for 16 hours. Apoptosis induction was also assessed with several *M. tuberculosis* recombinant proteins (10 $\mu\text{g}/\text{ml}$, Table 1) under the same culture conditions. Camptothecin (25 $\mu\text{g}/\text{ml}$) was employed as a positive control (Sigma Aldrich). Basal apoptosis was measured in non-stimulated macrophages. Cells were fixed after the incubation period with 1% paraformaldehyde.

hyde at 4 °C for 15 minutes. They were then washed twice with 1 ml of PBS-1x and stored in a 70% ethanol solution at -20°C until the DNA fragmentation analysis.

DNA fragmentation

DNA fragmentation was determined by the TUNEL technique, using the APO-BrdU™ (Invitrogen) commercial kit following the manufacturer's specifications as previously described.^{10,14} Cells were stained in presence or absence of terminal deoxynucleotidyl transferase (TdT) as control for assay specificity. Fluorescein isothiocyanate (FITC) labeling was used to identify DNA fragmentation with a total of 10⁴ cells registered per assay with the flow cytometer (Facs Aria III, BD). Data were analyzed using the Infinicyt™ Flow Cytometry Software 1.7 version. Camptothecin-treated macrophages were used as a positive control for apoptotic features, whereas non-stimulated macrophages were used as negative controls. Values are represented as the mean ± SE of three independent assays.

Caspase 3 activity

To determine caspase 3 activity, a total of 10⁶ cells were incubated for 24 hours with 100 µg/ml of either SE or CFE proteins, or with 10 µg/ml of each of the recombinant proteins. Cells were then washed with 1 ml of 1X PBS and incubated at 4 °C for 15 minutes with lysis buffer (10 mM Tris-HCl, 10 mM NaH₂PO₄, 130 mM NaCl and Triton x100 at 1%). Subsequently, 100 µl of assay buffer were added (40 mM HEPES, 20% glycerol, 1M DDT, 10 mM AcDEVD-AM) and cells were incubated for a further 5 minutes at room temperature. Finally, emission fluorescence in the 380/430-460nm range was determined for 20 minutes using the Synergy HT Bio Tek microplate reader. Values are represented as the mean ± SE of four independent assays.

Statistical analysis

Effect of treatments on induction of DNA fragmentation and the caspase 3 activity was assessed by ANOVA and the Tukey's multiple comparison test. The statistical program GraphPad Prism 7.0 was used.

Results and discussion

The main goal of this study was to assess the effect of *Mycobacterium tuberculosis* recombinant proteins, and of *Mycobacterium bovis* derived protein extracts on apoptosis induction in bovine macrophages. A total of 6 L of *Mycobacterium bovis* were cultured in Sauton medium to obtain 174 mg of the soluble protein extract (SE) and 7.6 mg of the culture filtrate protein extract (CFE). **Figure 1** shows a 12% polyacrylamide gel representative of the total proteins from both extracts. In the SE, twenty-six protein bands with molecular weights from 100 to 6 kDa were observed, whereas 10 bands were found in the CFE, with molecular weights of proteins ranging from 80 to 6 kDa. These results are similar to previously reported data.¹⁵

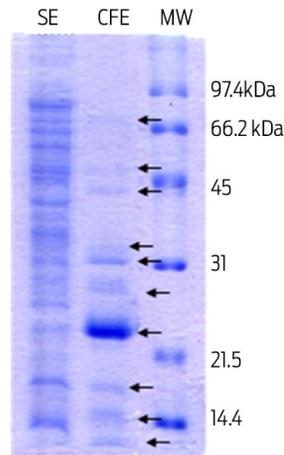


Figure 1. 12% SDS PAGE stained with Coomassie R-250 blue, showing bands from a soluble protein extract (SE) and culture filtrate protein extract (CFE) derived from a *M. bovis* AN5 culture. A total of 20 µg and 10 µg of protein were analyzed from the SE and the CFE respectively. MW: low-range Bio-Rad molecular weight.

Induction of apoptosis in bovine macrophages was evaluated with 100 µg of total proteins from extracts (SE and CFE) derived from *M. bovis*. Percentage of DNA fragmented cells after incubation with each protein extract is shown in [Figure 2A](#). CFE and SE prompted DNA fragmentation in $14.11 \pm 4.15\%$ and $26.05 \pm 4.49\%$ of cells respectively. DNA fragmentation was observed in $4.04 \pm 4.29\%$ of control cells. Induced apoptosis by both protein extracts was shown to be caspase 3 independent ([Figure 2B](#)).

The culture filtrate protein extract is usually enriched with secretion proteins, where the most abundant are the Ag85, MPT64 and ESAT-6 family proteins. For the soluble extract, it is possible to find both cytosolic and membrane proteins, depending on the separation method used.¹⁶ We intend to identify specific proteins from extracts in subsequent studies.

Few factors capable of triggering apoptosis have been identified in mycobacteria thus far, from which some examples are the LpqH,¹⁷ Esat-6,¹⁸ PE-PGR33,¹⁹ PstS-1²⁰ and HbhA²¹ proteins. We evaluated a panel of 13 recombinant *M. tuberculosis* proteins with known functions and molecular weights in apoptosis induction in bovine macrophages ([Table 1](#)). All proteins induced DNA fragmentation in cells, however the highest percentages of DNA fragmented bovine macrophages were found with the Hsp70 and HbhA proteins ($32.7 \pm 11.63\%$ and $29.69 \pm 17.41\%$, respectively; [Figure 3A](#)). Induced apoptosis was independent of the caspase 3 pathway ([Figure 3B](#)).

The Hsp70 protein has not been shown to elicit apoptosis in cells. It can however activate NF-κB through TLR-2 and TLR-4,²² to promote the synthesis of pro-inflammatory cytokines such as TNF-α,²³ which trigger apoptosis through the extrinsic pathway. In contrast, the HbhA protein can induce apoptosis in murine macrophages through a caspase-dependent mechanism.²¹ However, none of the recombinant proteins employed in this study activated caspase 3, suggesting that bovine macrophages can differentially respond to factors of mycobacterial origin. Other recombinant proteins, which are commonly described as apoptosis inducers, such as ESAT-6, PstS-1, LpqH, PE-PGR33, marginally increased the percentage of DNA fragmented cells, but this outcome was not statistically significant. It is

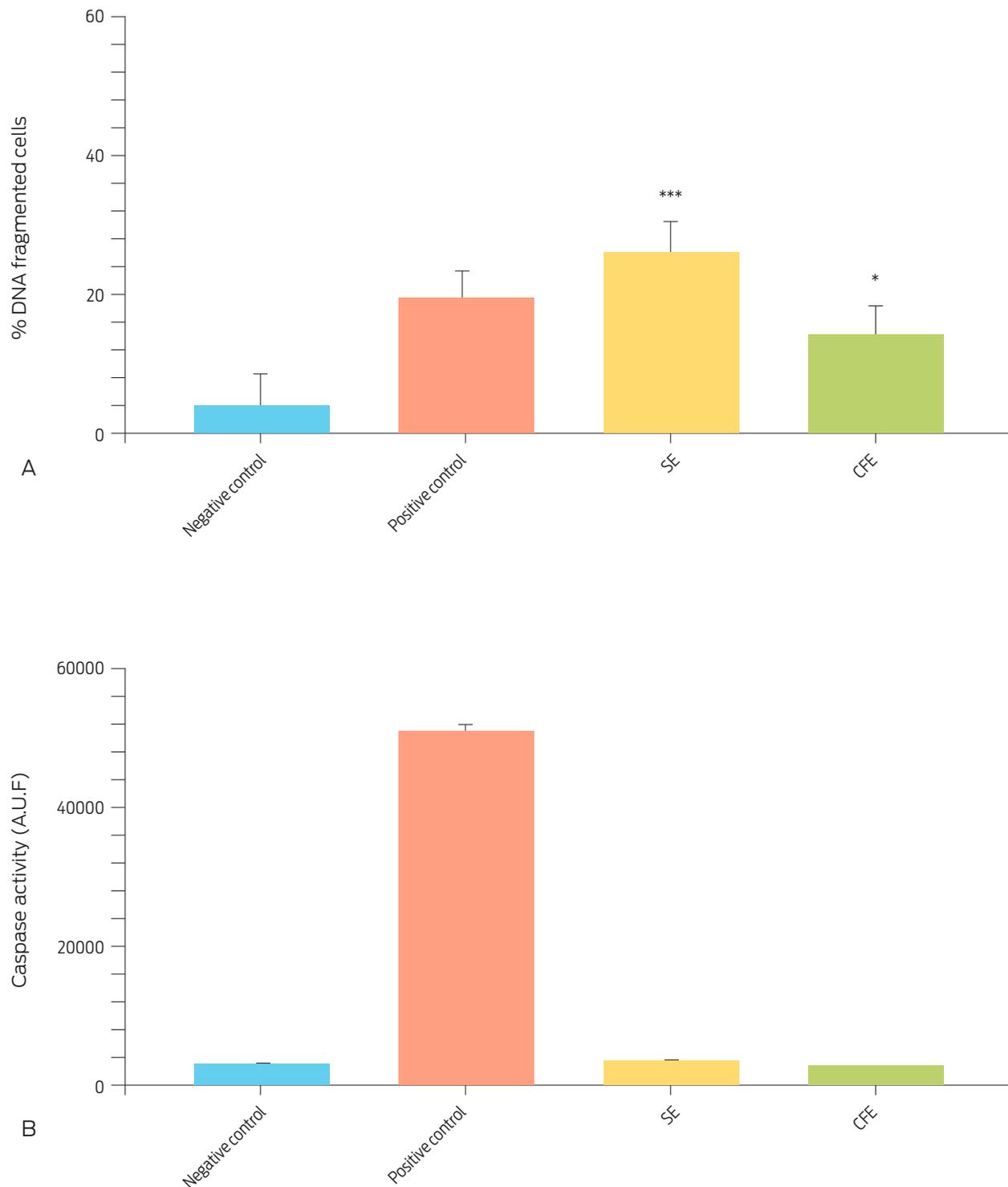


Figure 2. Percentage of DNA fragmented bovine macrophages and caspase 3 activity after incubation with 100 $\mu\text{g/ml}$ of *M. bovis* soluble protein extract (SE) or of culture filtrate protein extract (CFE) for 16 hours. Camptothecin (20 $\mu\text{g/ml}$) was used as a positive control and RPMI as a negative control. A) DNA fragmentation was assessed by TUNEL assay. Results are expressed as the mean \pm S.E. of three independent assays. (***) $P = 0.005$, * $P = 0.0438$). B) Caspase activity was measured through arbitrary units of fluorescence (A.U.F.) and expressed as means \pm SE of four independent assays ($P = 0.0001$).

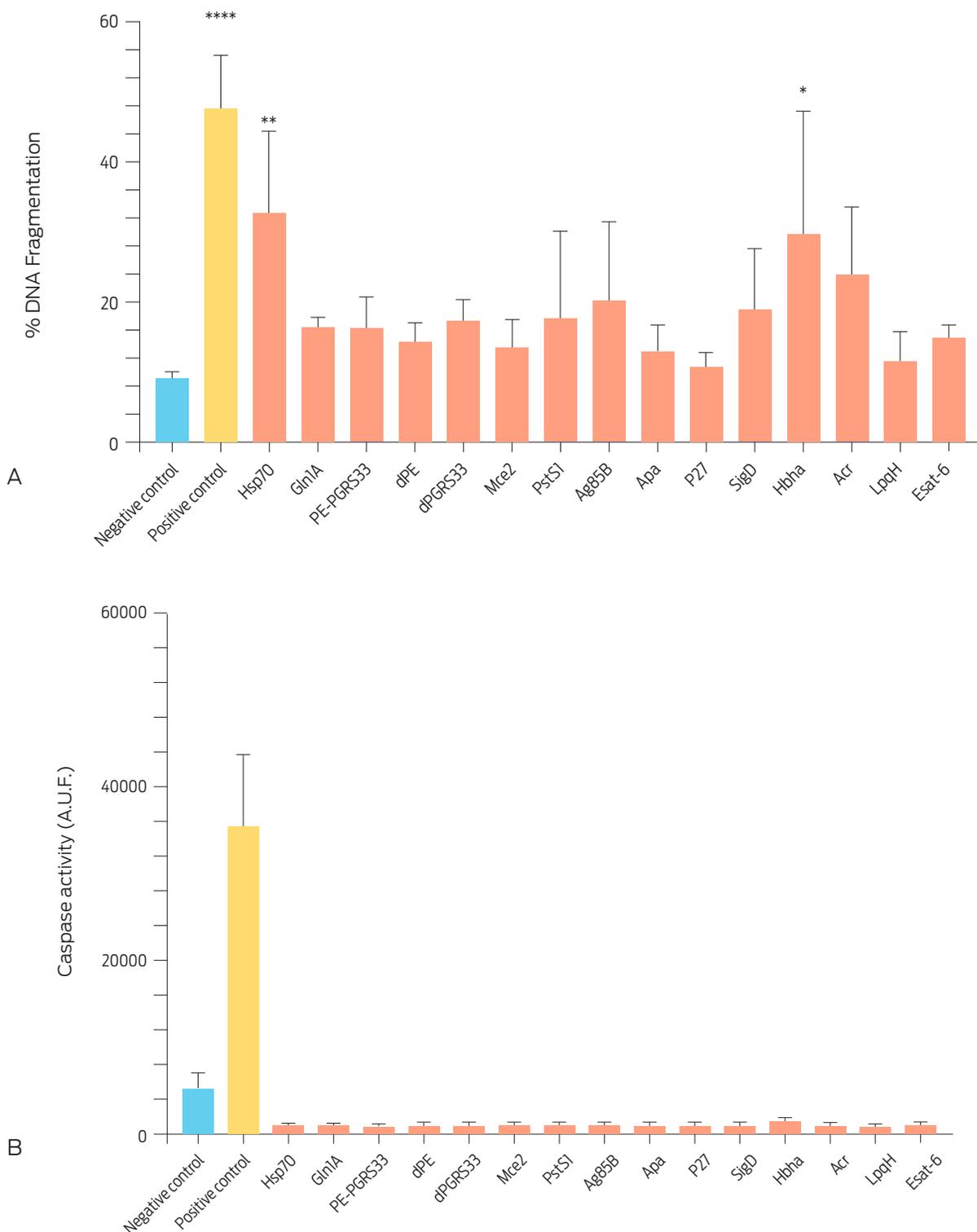


Figure 3. Percentage of DNA fragmented bovine macrophages and caspase 3 activity after culture with *Mycobacterium tuberculosis* complex recombinant proteins for 16 hours. A) Macrophages were incubated with 10 µg/ml of each protein and 20 µg/ml of camptothecin and RPMI as positive and negative controls, respectively. DNA fragmentation was assessed by TUNEL assay. Results are expressed as means ± S.E of three independent assays (P < 0.05). B) Caspase activity was measured using arbitrary units of fluorescence (A.U.F.), in 10⁶ bovine macrophages incubated for 24 hours with 10 µg/ml of each of the recombinant proteins Data are expressed as means ± S.E. of two independent assays with two replicas each. Recombinant proteins Ag85-B, Apa and Gln1A were not included when analyzing caspase activity.

not clear why apoptosis was not observed in a larger number of macrophages after incubation with these proteins as has been reported in other studies; however, specific characteristics of the used cell types could allow for different results.

Pathogenesis of mycobacteria relates to their ability to survive within macrophages and to modulate apoptosis induction of infected cells. In addition to multiplicity of infection and virulence of the mycobacterial strain, the course of an infection by the *M. tuberculosis* complex is determined by the type of triggered cell-death.^{6,7,11} Apoptosis in human or mouse macrophages after *Mycobacterium tuberculosis* infection is normally due to caspase activation. However, when the multiplicity of infection is 25:1, macrophages undergo apoptosis independently of the caspase pathway. Bovine macrophages infected by *M. bovis* also undergo apoptosis¹² independently of caspases.¹³ Hence, apoptotic pathway taken by the cell after infection could have important implications for disease outcome and host survival.⁶⁻⁸ Our main finding is that a subgroup of proteins and protein extracts from strains of the *Mycobacterium tuberculosis* complex were able to induce DNA fragmentation in bovine macrophages through a pathway in which caspases were not activated. This suggests that response of bovine macrophages to infection by tuberculosis bacilli differs from that seen in murine or human cells.¹⁶ We further speculate that the caspase-independent apoptosis mechanism aims to protect the organism from pathogens and is activated when the caspase pathway has failed. Mycobacteria may prevent caspase activation and thus apoptosis induction of its host cell to survive, whilst the host may trigger a caspase independent apoptosis pathway in an attempt to kill the pathogen. Our findings support the concept that apoptosis may work as a natural resistance mechanism of the host.^{14,24}

Conclusions

Apoptosis is a host defense mechanism against mycobacterial infection that can also benefit the pathogen by eliminating host defense cells. It is thus necessary to understand the mechanisms used by bacteria to modulate apoptosis. Our data show that bovine macrophages respond differently than murine or human cells to mycobacterial infection. In addition, this is the first report of cell-death induction in bovine macrophages by two *M. tuberculosis* recombinant proteins, not previously identified as apoptotic, through a caspase independent pathway. Moreover, soluble protein (SE) and culture filtrate protein (CFE) extracts from cultured *M. bovis* also induce apoptosis through a caspase independent mechanism. This pathway had been only previously observed with high multiplicities of infection with *M. tuberculosis*.

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Author contributions

A.B.G., E.A.S., J.A.G.P.: experimental design; A.M.R., I.J.V., O.C.B.: laboratory work; A.M.R., S.F.V., I.J.V., A.B.G., J.A.G.P.: data analysis; C.E.P., J.M., L.A.-P., J.A.G.P.: contributed reagents/materials/analysis tools; S.F.V., J.A.G.P.: document drafting.

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