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5 **PRODUCTION AND UTILISATION OF INTERLEUKIN-15 IN MALIGNANT**
6 **CATARRHAL FEVER**

7
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1 SUMMARY

2 Malignant catarrhal fever is an often fatal lymphoproliferative disease of ungulates caused by
3 either Alcelaphine herpesvirus-1 (AIHV-1) or Ovine herpesvirus-2 (OvHV-2). The pathogenesis
4 is poorly understood, but appears to involve an auto-destructive pathology where active cytotoxic
5 cells destroy areas of a variety of tissues. Interleukin-15 is a cytokine that develops and maintains
6 cytotoxic cells and may therefore have a role in the pathogenesis of MCF. We show herein that
7 the virus-infected large granular lymphocytes (LGL) obtained from MCF-affected rabbit tissues
8 exhibited a similar proliferative response to both IL-15 and to IL-2 in culture, but their activated
9 cytotoxic enzyme (BLT-esterase) content was maintained at higher levels in the presence of IL-
10 15 compared to IL-2. The LGLs did not produce IL-15 (either mRNA or protein). Abundant
11 expression of IL-15 mRNA and protein was detected in tissues rich in MCF lesions. Interestingly,
12 IL-15 production was associated with the MLN and appendix of OvHV-2-infected rabbits where
13 necrotic lesions were frequent, but not in the same tissues of AIHV-1-infected rabbits that lacked
14 necrotic lesions. The cellular source of the IL-15 was predominantly lymphocyte-like cells that
15 lacked B cell or monocyte- macrophage markers. Only a few IL-15+ cells (<10%) co-localised
16 with pan T cells or CD8+ T cells. The results show a marked abundance of IL-15 in MCF,
17 associated with tissue lesions that confers on the cytokine a potential role in MCF pathogenesis.

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1 INTRODUCTION

2 Malignant Catarrhal Fever (MCF) is an acute, often fatal lymphoproliferative disease of cattle,
3 deer and other susceptible ungulates (Reid, 2000; Coulter *et al.*, 2001). The known causative
4 agents are either of two related γ -herpesviruses – Alcelaphine herpesvirus-1 (AIHV-1) and Ovine
5 herpesvirus-2 (OvHV-2), the genome sequences of which have been obtained (Ensser *et al.*,
6 1997; Hart *et al.*, 2007). The natural reservoir host for AIHV-1 is the wildebeest (*Connochaetes*
7 *spp*), and for OvHV-2 is the sheep (*Ovis aries*). In these animals there is no apparent disease and
8 the virus persists as a benign infection. However, a proportion of cattle, deer or other susceptible
9 species that are in the vicinity of infected lambs or wildebeest calves can become infected and
10 develop MCF which is normally fatal. MCF is currently a serious problem in cattle in Indonesia
11 and Africa and bison in the USA. Rabbits develop MCF after experimental infection with AIHV-
12 1 or OvHV-2 that is characteristic of the disease in cattle (Buxton and Reid 1980; Buxton *et al.*,
13 1984).

14
15 MCF is characterised by hyperplasia of lymphoid organs and areas of necrosis in these and in
16 non-lymphoid tissues associated with the infiltration of large numbers of T lymphocytes and
17 lymphoblasts. Vasculitis is common (Liggitt and DeMartini 1980a,b; Anderson *et al.*, 2007). A
18 feature of MCF is that there is severe pathology in susceptible hosts in spite of the presence of
19 only a few virus-infected cells (as detected by virus DNA). The virus-infected cells in both
20 AIHV-1 and OvHV-2 MCF are lymphocytes at lesion sites in the tissues of MCF-affected
21 animals (Bridgen *et al.*, 1992; Simon *et al.*, 2003) leading to our current hypothesis that MCF is
22 caused by the auto-destruction of tissues by indiscriminately-cytotoxic lymphocytes, produced as
23 a consequence of MCF virus-infection (Swa *et al.*, 2001).

24
25 In contrast to the situation *in vivo*, virtually all large granular lymphocyte T cells (LGL) in culture
26 contain viral gene transcripts and antigen (Swa *et al.*, 2001). These are derived from the lymph

1 nodes, spleen, corneal and other tissues of MCF-affected animals (Reid *et al.*, 1983, 1989). The
2 LGL transmit MCF when injected into rabbits or other susceptible species. The phenotype of the
3 cultured LGL from OHV-2-infected cattle or AIHV-1 or OvHV-2-infected rabbits is generally
4 characteristic of T cells (Schock *et al.*, 1998; Swa *et al.*, 2001). Other characteristics of LGL in
5 culture are: constitutive, indiscriminate (i.e. non-MHC-associated) cytotoxicity; lack of mitogen-
6 stimulated proliferation in culture; and in the case of OvHV-2 cattle LGLs, lack of IL-2 mRNA
7 and protein expression, but expression of inflammatory cytokines (e.g. TNF- α and IL-1 β) and
8 IFN- γ (Schock *et al.*, 1998; Swa *et al.*, 2001). We have previously shown that the activated
9 phenotype of the LGLs, associated with MCF virus infection and in the absence of exogenous IL-
10 2, involves constitutively-activated Lck and Fyn src kinases (Swa *et al.*, 2001). The LGL
11 represent a population of virus-infected T cells found *in vivo* and are a valuable resource for
12 studying virus-cell interaction in the pathogenesis of MCF.

13
14 Interleukin-15 is a pleiotropic cytokine involved in both innate and adaptive immune responses to
15 infection (Ohteki 2002; Waldeman, 2006) that utilises a specific α receptor along with the
16 common β and γ receptor components of the four alpha helix bundle cytokine family that
17 includes IL-2. It is produced by a variety of tissue cells as part of an inflammatory response and
18 stimulates the survival and proliferation of cytotoxic T cells, NK cells and NKT cells. Its
19 production is normally highly-regulated. However, dysregulated expression of IL-15 is associated
20 with a spectrum of autoimmune diseases that involve cytotoxic T cells and/or NK cells (Ruchatz
21 *et al.*, 1998; Villasden *et al.*, 2003), and over-expression in transgenic mice leads to fatal T-NK
22 cell leukaemia (Fehninger and Caligiuri, 2001). This spectrum of activity could indicate a
23 possible contribution of IL-15 to the pathogenesis of MCF. We are testing the hypothesis that IL-
24 15, dysregulated as a consequence of infection, is involved in a pathway to recruit, activate and
25 maintain the cytotoxic activity of T cells in the tissues of MCF-affected animals. To test this

1 hypothesis, in the first instance we need to demonstrate that IL-15 is produced and utilised in
2 animals with MCF. Thus, the objectives of the present study were: first, to determine whether
3 LGLs from MCF-affected animals express IL-15 and/or respond to IL-15. Secondly, to determine
4 whether IL-15 was expressed in the lesion-containing tissues of MCF-affected animals.

5

6 **METHODS**

7 *Animals.* Nineteen New Zealand White rabbits, 2-3 months of age were used in this study. A
8 group of eight rabbits were inoculated intravenously (IV) with $1-2 \times 10^6$ AIHV-1-infected rabbit
9 lymphoid cells and another group of eight rabbits with $1-2 \times 10^6$ OvHV-2-infected rabbit
10 lymphoid cells (see below). A control group consisted of three uninfected rabbits. Rectal
11 temperatures were monitored and on the second day of a febrile response (defined as $\geq 40^\circ\text{C}$)
12 rabbits were euthanased. The animal experiments were carried out with the approval of the
13 Moredun Research Institute's experiments and ethics committee and complied fully with the
14 Home Office of Great Britain and Northern Ireland "Animals (Scientific Procedures) Act 1986".

15

16 *Viruses and the development of LGL cell lines.* AIHV-1 strain C500 (Handley *et al.*, 1995; Ensser
17 *et al.*, 1997; Swa *et al.*, 2001) was maintained by passage in rabbits using infected lymphoid
18 tissue as described above. OvHV-2 strain BJ1035 was originally from a cow with MCF in the UK
19 (Schock *et al.*, 1998; Swa *et al.*, 2001; Thonur *et al.*, 2006; Hart *et al.*, 2006) and was maintained
20 in rabbits by IV inoculation of infected LGL (described below) first, then subsequently with
21 rabbit lymphoid cells.

22 The AIHV-1-infected LGL cell lines (BJ1859, BJ1860, BJ1263) were generated by injecting
23 AIHV-1 C500-infected rabbit lymphoid cells (a pool of $\sim 10^6$ viable MLN and spleen cells from
24 frozen stock) intravenously into rabbits. Two days after a rise in rectal temperature $>40^\circ\text{C}$, the
25 rabbits were euthanased. MLN and spleen cells were harvested, single cell suspensions prepared

1 and cultured at $5-10 \times 10^6$ cells/ml Iscove's modified Dulbecco's medium containing 10% FCS
2 (IMDM) and 10U/ml interleukin-2 (IL-2, (Eurocetus, U.K.) in 25 cm² flasks.

3 The BJ1035, BJ1104, BJ1044 and BJ1196 LGL lines were derived from cattle naturally-infected
4 with OvHV-2 (Schock *et al.*, 1998; Swa *et al.*, 2001; and unpublished data). The BJ1857 LGL
5 line was derived from the lymphoid tissue of a rabbit with MCF induced by intravenous IV
6 injection of $\sim 10^6$ OvHV-2-infected rabbit lymphoid tissue cells (the OvHV-2 originally from a
7 clinical case of MCF and passaged in rabbits). The LGL were routinely passaged in IMDM
8 containing IL-2. AIHV-1 + and OvHV-2+ LGLs developed after 2-3 weeks of splitting and
9 refeeding the cells with medium containing IL-2 on an approximately weekly basis. After 8
10 weeks the phenotype of the cells had stabilised (Table 1). AIHV-1 genes and proteins were
11 detected in the LGL using PCR for AIHV-1 DNA and indirect immuno-fluorescence using serum
12 antibody to AIHV-1 proteins from an infected rabbit using techniques described previously (Swa
13 *et al.*, 2001). The presence of OvHV-2 DNA in the cells was detected by PCR as described
14 previously (Swa *et al.*, 2001). OvHV-2 virus proteins were detected within the cells using
15 antibodies from the serum of a convalescent cow and an indirect immunofluorescence technique
16 (Swa *et al.*, 2001). LGL phenotype analysis was by indirect immunofluorescence test using the
17 antibodies identified in Table 1.

18 Control rabbit or bovine cells were T cells derived from peripheral blood mononuclear cells
19 (PBMC) that were obtained by centrifugation of blood over lymphoprep (Nyegard, Oslo,
20 Norway). 2×10^6 /ml PBMC were stimulated with 5 μ g/ml Con-A for three days. 2×10^5
21 lymphoblasts/ml medium were expanded for 1-2 weeks in 10U/ml IL-2. The cells were split and
22 refed with the IL-2 and fresh medium every 3-5 days. The control cells were a mixture of CD8+
23 and CD4+ T cells (rabbit) and CD4+, CD8+ and $\gamma\delta$ TCR+ T cells (bovine) and were negative for
24 AIHV-1 or OvHV-2 viral DNA or antigens.

25

1 *BLT-esterase detection in LGLs.* Cytotoxic cell BLT-esterase activity was detected in the LGL
2 and control cells using the tryptase-specific substrate N- α -benzyloxy-carbonyl-L-lysine
3 thiobenzyl ester (BLT, Sigma, Poole, U.K.) as described previously (Haig *et al.*, 1996). Cell
4 lysates and cell-free supernates from culture (50 μ l) were added to 96-well flat-bottomed plates
5 and 200 μ l of a mixture of 0.3mM BLT and 0.33mM dithio-bis-(2-nitrobenzoic acid) in PBS pH
6 7.2 was added. Incubation was for 1 hour at 37°C. The reaction was terminated by placing the
7 plates on ice and adding 5mM Pefabloc[®] protease inhibitor (Boehringer-Mannheim)/ well. Plates
8 were read on an ELISA plate reader at an absorbance of 412nm. Enzyme activity was expressed
9 as Abs₄₁₂/hour at 37°C/ ml sample (Abs).

10

11 *Experimental procedure for cytokine stimulation of the LGLs*

12 LGL and control T cells were stimulated for 6 days with various doses of IL-2 or IL-15
13 (Peprotech Ltd). Viable cell numbers were counted, and the BLT-esterase assay performed on
14 culture supernates (not shown) and cell lysates.

15

16 *Tissue processing and sampling*

17 Rabbits were infected with either rabbit adapted (by *in vivo* passage) BJ1035 (the OvHV-2 virus
18 that has recently been sequenced (Hart *et al.*, 2007) or rabbit-adapted C500 (AIHV-1) as
19 described above. Uninfected rabbit tissues were collected as controls. At autopsy, various tissues
20 were examined for gross signs of MCF. Samples of appendix, kidney, liver, lung, mesenteric
21 lymph node (MLN) and spleen were collected and divided into aliquots for: (1) Formalin fixation
22 for histological analysis of H&E stained sections. (2) 4% paraformaldehyde fixation for IL-
23 15mRNA detection by the *in situ* hybridisation technique and IL-15 antigen by
24 immunohistochemistry (IHC)(see below); (3) Zn salts fixative for IHC using mabs to rabbit
25 alloantigens for cell phenotype analysis. In addition, aliquots of lymphoid cell samples (appendix,

1 spleen, MLN) were taken fresh to prepare single cell suspensions to obtain cell extracts for
2 western blot analysis of tissue IL-15. All fixation and tissue preparation procedures have been
3 described previously (Gonzales *et al.*, 2001; Anderson *et al.*, 2007). Standard 5µm-thick serial
4 sections were cut and mounted on Superfrost Plus glass slides (Menzel-Glaser, Braunschweig,
5 Germany) and dried overnight at 37°C. One section from each paraformaldehyde-fixed tissue was
6 stained with haematoxylin and eosin (H&E). Remaining sections were stored in slide boxes at
7 4°C until required.

8

9 *Histopathology*

10 In all infected rabbits (AIHV-1 or OvHV-2), MCF was confirmed by clinical and histological
11 criteria as described previously (Anderson *et al.*, 2007). The histopathology of MCF lesions in
12 the different tissues was defined in H&E sections as (a) areas of hyperplasia/ lymphoid cell
13 accumulation (including cells exhibiting mitotic indices) and (b) necrosis (focal tissue
14 destruction). In lymphoid tissues where hyperplasia and/or necrosis were present a score of + was
15 given. In tissues where no changes were present a score of - was assigned. In non-lymphoid
16 tissues, large interstitial lymphoid cell accumulations (affecting >25% of the area of tissue) were
17 scored as ++ and smaller less frequent interstitial accumulations (affecting less than 25% of the
18 area of the tissue) as +. Tissues with no interstitial lymphoid cell accumulations were given a
19 score of - .

20

21 *Immunohistochemistry*

22 Sections were dewaxed and hydrated using standard procedures. Immunohistochemical labelling
23 was performed manually using the Sequenza rack and coverplate technology (Shandon, Runcorn,
24 UK). The EnVision Plus HRP system (Dako, Ely, UK) was used to detect antibody binding as
25 described previously (Gonzales *et al.*, 2001; Anderson *et al.*, 2007). Labelled sections were
26 examined microscopically. Anti-rabbit alloantigen mabs used were: RTH1A (rabbit CD4-

1 specific), ISC29E (rabbit CD8-specific), RTH 26A (rabbit pan-T-specific), and MRB 29-A
2 (rabbit B cell-specific). These mabs have been described previously (Anderson *et al.*, 2007).
3 For IL-15 detection, mab 247 (R&D systems) was used on paraformaldehyde-fixed tissue treated
4 (after dewaxing) for antigen retrieval with sodium citrate buffer (10mM sodium citrate, 0.05%
5 Tween 20, pH 6.0) for 30 minutes at 95°C. The 247 mab has previously been shown to be
6 specific for rabbit IL-15 (Xiong *et al.*, 2005) as well as for human and murine IL-15.

7

8 *In situ hybridisation*

9 *In situ* hybridisation was used to detect and locate IL-15 mRNA in the tissues of MCF-affected
10 and uninfected control rabbits. Rabbit IL-15 cDNA was prepared from mRNA obtained from the
11 MLN of a MCF-affected rabbit and the DNA sequenced. The predicted amino acid sequence was
12 the same as that of rabbit IL-15 described previously (Xiong *et al.*, 2005). For riboprobe
13 preparation, restriction enzyme fragments (between 200bp and 500bp) of the IL-15 cDNA were
14 cloned into the transcription vector pSPT 18/19 (Roche diagnostics, Lewes, England). Both sense
15 and antisense RNA probes were prepared by *in vitro* transcription from the T7 and SP6 promoter
16 incorporating digoxigenin UTP (Roche). The concentration of each labelled probe was
17 determined by dot blot. *In situ* hybridisation was as described previously (Anderson *et al.*, 2001),
18 using an Omnislid thermal cycler and wash module (Hybaid, Ashford, England). Briefly,
19 sections were de-waxed and rehydrated and then treated with 7.5µg/ml proteinase-K in 200mM
20 HCL for 15 minutes at 37°C. After post-fixation with 4% paraformaldehyde, sections were
21 acetylated for 10 minutes with 100mM tri-ethanolamine pH 7.0, 0.25% acetic anhydride and then
22 pre-hybridised for 2h at 52°C in 6 x SSC, 45% de-ionised formamide, 5 x Denharts solution and
23 256µg per ml bovine liver RNA (Sigma). Hybridisation solution was 48% deionised formamide,
24 5 x Denharts solution, 1000U /ml heparin (Sigma), 0.1% Triton-X 100, 24mM EDTA, 24mM
25 Pipes (Sigma), 720 mM NaCl, 7.2% dextran sulphate (Sigma) and 512µg/ml bovine liver RNA
26 containing digoxigenin-labelled probes (sense or antisense) at concentrations between 0.1 to 0.4

1 ng/μl. Hybridisation was overnight at 52°C. Sections were washed and treated with RNase to
2 remove unbound probe. Bound probe was detected using anti-digoxigenin antibody Fab
3 fragments conjugated to alkaline phosphatase as described (Anderson *et al.*, 2001). Sections were
4 counterstained with haematoxylin.

5

6 *Western blots*

7 Both Mab 247 and the murine soluble IL-15 receptor protein (sIL15R, his-tagged; Ruchatz *et al.*,
8 1998; Smith *et al.*, 2000; Wei *et al.*, 2001) were used to detect rabbit IL-15 by western blot. A
9 single cell suspension of each tissue was prepared and the cells lysed in lysis buffer (0.8%
10 CHAPS in Tris-buffered saline pH8.3 with proteinase inhibitors ('Complete',Boehringer
11 Mannheim, UK). Lysates were adjusted to 1mg/ml and 10μl samples run on 15% SDS-PAGE gel
12 lanes under reducing conditions. After electro-transfer of proteins to nitrocellulose, the sheets
13 were treated with 1% casein in PBS for 1 hour. After several washes (PBS containing 0.5M NaCl
14 and 0.5% Tween 80), blots were incubated for 1 hour with either 2μg/ml sIL-15R or 1μg/ml
15 biotinylated mab 247 in wash buffer. After washing, bound receptor or antibody was detected by
16 either 2μg/ml anti-his-HRP conjugate (Roche) for the sIL-15R or 2μg/ml streptavidin-HRP
17 conjugate for mab 247 detection. Visualisation was by ECL chemiluminesence.

18

19 *Statistical analysis of data.*

20 Student's *t*-test was performed on data normalised by log₁₀ transformation.

21

22

1 RESULTS

2 *IL-15 supported large granular lymphocyte proliferation and BLT-esterase content.*

3 Table 1 shows the phenotype of the LGL cells used in this study. This was similar to virus-
4 infected LGL lines derived from MCF-affected animals and described previously (Schock et al.,
5 1998; Swa et al., 2001). The LGLs showed characteristics of T cells (CD2+ or CD5+, and with a
6 proportion of each line except BJ1035 expressing CD8). A constitutively-activated LGL
7 phenotype was indicated by the presence of BLT-esterase. LGLs were viral DNA positive by
8 PCR and >90% of the cells in each line were positive for virus antigen (either AIHV-1 or OvHV-
9 2 - not shown).

10

11 Figure 1 (rabbit LGLs and control T cells) and figure 2 (cattle BJ1035 LGL line and control T
12 cells) show that both the AIHV-1-infected LGL and OvHV-2-infected LGL lines exhibited IL-15
13 dose-dependent proliferation over 6 days in culture that was similar (in terms of growth curves)
14 both to the response of the cells to IL-2 and also to control non-infected T cell lines stimulated
15 with IL-15 or IL-2 (Fig. 1A, C, E, G; Fig. 2 A,C). The addition of a half maximal dose of IL-2
16 (1U/ml) with a range of IL-15 concentrations to the rabbit LGL or control T cell lines did not
17 reveal any synergistic effect of the combination over either cytokine alone, although an increased
18 proliferation compared to either cytokine alone was observed (Fig. 1A,C,E,G).

19

20 The content of BLT-esterase per 10^6 LGL cells in culture in general showed an inverse
21 relationship to cell proliferation in the presence of both IL-2 and IL-15 (Fig. 1D,F,H compared to
22 Fig. 1C, E,G and Fig. 2D compared to Fig. 2C) in that enzyme content per cell declined as cell
23 numbers increased. However, cells grown in IL-15 maintained higher levels of enzyme / 10^6 cells
24 over the dose range 10-100ng/ml than those in IL-2 (Fig. 1D,F,H and Fig. 2D, $P < 0.01$ for IL-15
25 at 10ng/ml, 50ng/ml or 100ng/ml compared to IL-2 at 1, 5 or 10U/ml). In contrast, in control
26 uninfected cells cultured in the higher doses of IL-15 (10ng/ml, 50ng/ml and 100ng/ml) there was

1 a dose-dependent increase in the content /10⁶ cells of BLT-esterase compared to cells in IL-2
2 (Fig. 1B, $P < 0.001$ for 50ng/ml or 100ng/ml IL-15 compared to 5U/ml and 10U/ml IL-2 and Fig
3 2B, $P < 0.007$ for 10ng/ml or 100ng/ml IL-15 compared to 5U/ml or 10U/ml IL-2). Thus, IL-15 at
4 higher doses maintained (LGLs) or induced BLT-esterase (control T cells) in the cells whereas
5 IL-2 did not. In the rabbit cells, the combination of 1U/ml IL-2 with different doses of IL-15
6 showed decreased cell content of BLT-esterase compared to cells stimulated with IL-15 alone.

7

8 *LGLs did not express IL-15*

9 The LGL lines (Table 1) were analysed for the expression of IL-15 mRNA by RT-PCR and for
10 IL-15 protein in cell lysates and supernates by western blot. In addition pelleted LGL cell pellets
11 (after centrifugation) were embedded in paraffin wax and taken through the *in situ* hybridisation
12 technique along with other tissues for the detection of IL-15mRNA. Neither IL-15 mRNA nor
13 IL-15 protein was detected in any of the LGL preparations. Figure 3A shows the absence of IL-
14 15 mRNA in the BJ 1859 AIHV-1+ LGL line by *in situ* hybridisation. Fig 3B and C show a
15 positive control and sense negative control respectively for the assay run at the same time.

16

17 *Tissue cells from MCF-affected animals expressed IL-15.*

18 IL-15 expression in cells or tissues was measured by *in situ* hybridisation (for IL-15 mRNA),
19 IHC and western blot (for IL-15 antigen). Table 2 shows a semi-quantitative score for the
20 frequency of IL-15 labelled cells measured by mab 247 in the various lymphoid and non
21 lymphoid tissues of AIHV-1-infected or OvHV-2-infected animals and uninfected controls and
22 comparing this with a lesion score for the tissues (Anderson *et al.*, 2007).

23 The results of the analysis revealed high frequencies of IL-15-expressing cells in both lymphoid
24 (appendix, MLN, spleen) and non-lymphoid tissues (particularly liver and lung) from rabbits
25 infected with OvHV-2 (Table 2; Fig 3B, Fig 4B,C,E). In general, IL-15 expression was
26 associated with MCF lesions (any or combinations of: areas of necrosis, lymphocyte

1 accumulations or lymphoid hyperplasia). In contrast to OvHV-2-infected rabbits, there was a
2 relative lack of IL-15-expressing cells in the MLN and appendix of AIHV-1-infected animals
3 (Table 2; Fig. 4B or C compared to Fig. 4A). There was an association between the presence of
4 necrotic lesions and IL-15-expressing cells in the MLN and appendix of OvHV-2+ MCF, and a
5 lack of necrotic lesions and a relatively low level of IL-15-expressing cells in the MLN and
6 appendix of AIHV-1+ MCF that was similar to that seen in uninfected control animals (Table 2).
7 In the spleen, necrotic lesions were not observed but hyperplasia was present in both AIHV-1
8 MCF and OvHV-2 MCF. There were less IL-15-expressing cells in the AIHV-1 group compared
9 to the OvHV-2 group, but this was not marked (Table 2). In non-lymphoid tissues, lymphoid cell
10 accumulations were more abundant in AIHV-1-infected compared to OvHV-2-infected animals,
11 but there was no obvious difference between the groups in the frequency of IL-15+ cells (Table
12 2; Fig. 3D,E,F for IL-15mRNA+ cells and controls in AIHV-1-infected rabbit liver).
13 The distribution of IL-15mRNA-expressing cells in the tissues was the same as for IL-15 antigen-
14 expressing cells. In all tissues in either infection, the IL-15-expressing cells were predominantly
15 mononuclear cells within areas of mainly lymphocyte accumulation and particularly located to
16 interfollicular (T cell) areas in lymphoid tissue (e.g. Fig. 4B,C,D). IL-15 mRNA and protein
17 expression was also associated with some epithelial and endothelial cells within or adjacent to
18 lesions (e.g. Fig 4E,F). However, in serial sections comparing IL-15-expressing cells and CD4+
19 cells or B cells (by digital overlay), there was no co-localisation. In spite of T cells labelled with
20 the pan T cell mab or CD8-specific mab and the majority of IL-15-expressing cells locating to
21 interstitial regions of lymphoid tissues and within mononuclear cell accumulations in non-
22 lymphoid tissues, only a minority of these (estimated at <10%) co-localised in the serial sections
23 (e.g. Fig. 4C compared to Fig. 4D).
24 The validity of the semi-quantitative scoring method was confirmed for lymphoid tissue
25 (appendix and MLN) from infected and control animals by the intensity of IL-15 protein bands
26 by western blot (Fig. 5A,B shows MLN samples). IL-15 was found in cell lysates but not in cell

1 culture concentrated supernates. Fig. 5A,B, shows in addition that the mab 247 and smIL-15R
2 respectively bind to rabbit IL-15, defined by its M.Wt (16kDa) and the fact that it binds to a sIL-
3 15 receptor.

4

5 **DISCUSSION**

6 In this study, we have shown that the virus-infected LGLs of cattle and a rabbit infected with
7 OvHV-2 or rabbits infected with AIHV-1 responded to IL-15 but did not produce it. In addition
8 we have shown that there was an abundance of IL-15 (mRNA+ cells, IL-15 antigen+ cells and
9 IL-15 western blot) expressed in lesion-rich areas of tissues of both AIHV-1 and OvHV-2 MCF-
10 affected rabbits. In disease-affected MLN and appendix there was a marked difference between
11 OvHV-2 MCF and AIHV-1 MCF as there was a high frequency of IL-15 expressing cells
12 associated with necrotic lesions in OvHV-2 MCF and a lack of IL-15 expressing cells and
13 necrotic lesions in these tissues in AIHV-1 MCF.

14

15 LGLs are virus-infected cells obtained from the tissues of MCF-affected animals and are thought
16 to represent the key cells involved in the pathogenesis of MCF (Reid *et al.*, 1983; Schock *et al.*,
17 1998; Swa *et al.*, 2001). They are indiscriminately cytotoxic for a variety of target cells in culture,
18 and exhibit a constitutively active *lck* and *fyn* kinase signalling pathway (Swa *et al.*, 2001). The
19 LGLs did not produce IL-15, which is therefore not involved in an autocrine loop maintaining the
20 phenotype of the LGLs. This is in contrast to HTLV-1-infected T cells where IL-15 and IL-15R
21 are induced, probably to expand the number of target T cells for HTLV-1 infection (Azimi *et al.*,
22 2000; Mariner *et al.*, 2001).

23

24 However, LGLs responded to IL-15 demonstrating that LGLs express a functional receptor for
25 IL-15 as well as for IL-2. This was shown for LGLs from rabbits infected with either OvHV-2 or
26 AIHV-1 and for a cattle LGL line infected with OvHV-2. IL-15 maintained the cell content of the

1 cytotoxicity-associated enzyme BLT-esterase in proliferating LGLs at higher levels than IL-2,
2 even though both cytokines stimulated similar rates of cell growth. IL-2 exhibited a dominant
3 negative effect on IL-15-mediated BLT-esterase maintenance per cell when both cytokines were
4 used together. Thus, IL-15 could be an important cytokine for maintaining active cytotoxic cells
5 in MCF. In control uninfected T cells, IL-15 but not IL-2 stimulated the accumulation of BLT-
6 esterase in a dose-dependent fashion, indicating its capacity to develop cytotoxic cells. The BLT-
7 esterase assay detects a tryptase that is part of the perforin-granzyme cytolytic pathway in
8 activated cytotoxic cells and a tryptase of mast cells (Garcia-Sanz *et al.*, 1987; Griffiths and
9 Mueller, 1991; Haig *et al.*, 1996). The results are consistent with the function of IL-15 in mouse
10 and man where it recruits, stimulates the development of and maintains the cytotoxic phenotype
11 of cytotoxic cells that include CD8 CTLs, CD4 CTLs, NK cells and NK T cells (Waldemann,
12 2006). IL-15 is also anti-apoptotic (Marks-Konczalik *et al.*, 2000) and dysregulation leading to
13 over-expression of IL-15 can lead to expansion of cytotoxic cells and autoimmunity (Waldeman,
14 2006). This may be occurring in MCF. Although IL-2 will stimulate proliferation and activation
15 of cytotoxic cells, it does not maintain active cytotoxic cell proliferation as it induces activation-
16 induced cell death (AICD) and the maintenance of regulatory T cells (T_{regs}) that limit T cell
17 responses (Marks-Konczalik *et al.*, 2000; Fontenot *et al.*, 2005).

18
19 We have shown that the IL-15 in MCF-affected animal tissues was produced mainly by
20 mononuclear cells and some epithelial and endothelial cells in areas containing MCF lesions
21 (lymphocyte accumulations in non-lymphoid tissue; hyperplasia and areas of necrosis in
22 lymphoid tissue). As mab 247 is the only antibody we have found that reacts with rabbit IL-15, a
23 quantitative assay (ELISA) of IL-15 production was not possible.

24
25 To determine the source of the IL-15, we looked for co-localisation (by IHC in serial sections) of
26 IL-15+ mononuclear cells with lymphocyte subsets. This was seen only with a few pan-T+ cells

1 and CD8⁺ T cells in interstitial areas of lymphoid tissue and in lymphocyte accumulations in
2 non-lymphoid tissue. There was no co-localisation with CD4⁺ T cells or B cells. CD14⁺
3 monocyte/macrophage cells are rare in the rabbit tissues studied (Anderson et al., 2007) and in
4 this study there was no co-localisation detected of IL-15⁺ cells with these (not shown). Thus, the
5 identity of the majority of IL-15⁺ cells in the mononuclear cell population is unknown. This
6 needs to be revisited using confocal techniques when more rabbit mononuclear cell-specific
7 antibodies and IL-15-specific antibodies become available. In mouse and man, the production of
8 IL-15 has been described mainly from monocytes, macrophages, dendritic cells, epithelial cells
9 and endothelial cells (Oppenheimer-Marks *et al.*, 1998; Dubois *et al.*, 2002; Ohteki *et al.*, 2001).
10 Although T cell expression of IL-15 has been recorded, there is little information on these cells.
11 Recently however, human T cells have been shown to constitutively produce low levels of IL-15
12 that acts through autocrine or juxtacrine loops to promote *ex vivo* homeostatic T cell proliferation
13 (that *in vivo* would maintain the T cell pool)(Miranda-Carus et al., 2005). Our study in rabbits is
14 the first we are aware of to show IL-15 production in abundance predominantly from
15 lymphocyte-like cells *in vivo* during a virus infection.

16
17 In other species, several isoforms of IL-15 have been described, the most common of which is a
18 membrane bound form (Dubois *et al.*, 2002; Sato *et al.*, 2007). This allows IL-15 -IL-15R
19 interactions to occur mainly by cell contact, involving low levels of cytokine maintained over a
20 long period of time. In this study, the nature of the IL-15 was not fully investigated, although
21 cytokine was found in cell lysates but not in concentrated supernates by western blot, suggesting
22 a predominance of cell-associated (non-secreted) IL-15.

23
24 In previous studies, we showed that there are differences in tissue tropism in the MCF caused by
25 AIHV-1 versus OvHV-2 (Buxton et al., 1984; Schock and Reid, 1996; Anderson et al., 2007). In
26 general, there was less tissue necrosis and more lymphoid cell accumulations in AIHV-1

1 compared to OvHV-2. In particular, there was marked tissue necrosis in the appendix and MLN
2 of OvHV-2-infected rabbits that was absent in AIHV-1-infected animals. In AIHV-1, MCF
3 lesions were more apparent in peripheral lymphoid tissues (popliteal lymph node, prescapular
4 lymph node and prefemoral lymph node) than in the central lymphoid tissues (appendix and
5 MLN)(Schock and Reid, 1996; Anderson *et al.*, 2007). This distribution was the converse for
6 OvHV-2 MCF. In this study, we confirm this observation with respect to MLN and appendix and
7 in addition demonstrate that IL-15 producing cells and the quantity of IL-15 were greatly
8 increased in the tissues from OvHV-2 –infected animals when compared to those of AIHV-1
9 infected rabbits. Importantly, it is known that lymphoid cell hyperplasia and lymphocyte
10 accumulation lesions *per se* are not associated with the onset of clinical MCF whereas necrotic
11 lesions are (Buxton *et al.*, 1984). The association of IL-15 with necrotic lesions is circumstantial
12 evidence of a role for IL-15 in the pathogenesis of MCF.

13

14 In other species, IL-15 is an inflammatory cytokine produced early in infections that can initiate
15 cytokine cascades (that include TNF- α and IFN- γ) and develop and expand activated cytotoxic
16 cells (McInnes *et al.*, 1997,1998). In this role, it can be involved in either host protective
17 responses to viruses or, if dysregulated, in autoimmune pathogenesis. In HSV infection for
18 example, IL-15 is required as part of a host inflammatory and immune response to recruit and
19 activate cytotoxic T cells and NK cells that are involved in virus clearance (Tsunobuchi *et al.*,
20 2000; Ashkar *et al.*, 2003). In addition to a protective role, IL-15 has also been implicated in
21 several autoimmune diseases. For example, in several rodent models of autoimmunity, blocking
22 IL-15 with antibodies or soluble receptors abrogated the severity of the disease (Ruchatz *et al.*,
23 1998; Smith *et al.*, 2000; Villadsen *et al.*, 2003). MCF-affected animals die in spite of a marked
24 IL-15 response in the tissues. This supports the contention that IL-15 is involved in MCF
25 pathogenesis and not virus clearance. The results of the present study are consistent with the view

1 that IL-15 is an active participant in MCF by supporting indiscriminate auto-destruction of host
2 tissues by cytotoxic T or NK cells induced as a consequence of infection.

3 We conclude that the abundance of IL-15 in MCF - affected tissues, associated with areas rich in
4 MCF lesions confers on the cytokine a possible role in the pathogenesis of MCF. This is further
5 supported by the fact that IL-15 stimulates the proliferation and maintenance of cytotoxic activity
6 in LGLs in culture. These cells, obtained from the tissues of MCF-affected animals are thought to
7 play a key role in MCF pathogenesis.

8

9 **ACKNOWLEDGEMENTS**

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12

13 **REFERENCES**

14 1. **Anderson, I., Reid, H.W., Nettleton, P.F., McInnes, C.J. & Haig, D.M. (2001).**

15 Detection of cellular cytokine mRNA expression during orf virus infection in sheep:
16 differential IFN- γ mRNA expression by cells in primary versus reinfection skin lesions.
17 *Vet Immunol Immunopathol* **83**, 161-176.

18 2. **Anderson, I., Buxton, D., Campbell, I., Russell, G., Davis, W.C., Hamilton, M.J. &**

19 **Haig, D.M. (2007).** Immunohistochemical study of experimental malignant catarrhal
20 fever in rabbits. *J Comp Path* **136**, 156 – 166.

21 3. **Ashkar, A.A. & Rosenthal, K.L. (2003).** Interleukin-15 and natural killer and NKT cells

22 play a critical role in innate protection against genital herpes simplex virus type 2
23 infection. *J Virol* **77**, 10168-10171.

24 4. **Azimi, N. Shiramizu, K.M. Tagaya, Y. Mariner, J. & Waldmann, T.A. (2000).** Viral

25 activation of interleukin-15 (IL-15): characterization of a virus-inducible element in the
26 IL-15 promoter region. *J Virol* **74**, 7338-7348.

- 1 5. **Bridgen, A., Munro, R. & Reid, H. W. (1992).** The detection of Alcelaphine
2 herpesvirus-1 DNA by *in situ* hybridization of tissues from rabbits affected with
3 malignant catarrhal fever. *J Comp Path* **106**, 351-9.
- 4 6. **Buxton, D. & Reid, H. W. (1980).** Transmission of malignant catarrhal fever to rabbits.
5 *Vet Record* **106**, 243-5.
- 6 7. **Buxton, D., Reid, H.W., Finlayson, & Pow, I. (1984).** Pathogenesis of 'sheep-associated'
7 malignant catarrhal fever in rabbits. *Res Vet Sci* **36**, 205-211.
- 8 8. **Coulter, L. J., Wright, H. & Reid, H. W. (2001).** Molecular genomic characterization of
9 the viruses of malignant catarrhal fever. *J Comp Path* **124**, 2-19.
- 10 9. **Dubois, S. Mariner, J. Waldmann, T.A. & Tagaya, Y. (2002).** IL-15Ralpha recycles
11 and presents IL-15 In trans to neighboring cells. *Immunity* **17**, 537-47.
- 12 10. **Ensser, A., Pflanz, R., & Fleckenstein, B. (1997).** Primary structure of the alcelaphine
13 herpesvirus 1 genome. *J Virology* **71**, 6517-6525.
- 14 11. **Fehninger, T.A. & Caligiuri, M.A. (2001).** IL-15: biology and relevance to human
15 disease. *Blood* **97**, 14-32.
- 16 12. **Fontenot, J.D., Rasmussen, J.P., Gavin, M.A. & Rudensky, A.Y. (2005).** A function
17 for interleukin-2 in FOXP3-expressing regulatory cells. *Nature Immunol* **6**, 1142-1151.
- 18 13. **Garcia-Sanz, J. A., Plaetinck, G., Velotti, F., Masson, D., Tschopp, J., MacDonald H.
19 R. & Nabholz, M. (1987).** Perforin is present only in normal activated Lyt2⁺ T
20 lymphocytes and not in L3T4⁺ cells, but the serine protease granzyme-A is made by both
21 subsets. *The EMBO Journal* **6**, 933-938.
- 22 14. **Gonzales, L., Anderson, I., Deane, D., Summers, C. & Buxton, D. (2001).** Detection of
23 immune system cells in paraffin wax-embedded ovine tissues. *J Comp Path* **125**, 41-47.
- 24 15. **Griffiths, G. M. & Mueller, C. (1991).** Expression of perforin and granzymes *in vivo*:
25 potential diagnostic markers for activated cytotoxic cells. *Immunology Today* **12**, 415-
26 419.

- 1 16. **Haig, D.M., Hutchison, G., Thomson, J., Yirrell, D. & Reid, H.W. (1996).** Cytolytic
2 activity and associated serine protease expression by skin and afferent lymph CD8⁺ T
3 cells during orf virus reinfection. *J Gen Virol* **77**, 953-961.
- 4 17. **Handley, J., Sargan, D.R., Herring, A.J. & Reid, H.W. (1995).** Identification of a
5 region of the alcelaphine herpesvirus-1 genome associated with virulence for rabbits. *Vet*
6 *Microbiol* **47**, 167-181.
- 7 18. **Hart, J., Ackermann, M., Jayawardane, G., Russell, G., Haig, D.M., Reid, H. &**
8 **Stewart, J.P. (2007).** Complete sequence and analysis of the ovine herpesvirus 2 genome.
9 *J Gen Virol* **88**, 28-39.
- 10 19. **Liggitt, H. D. & DeMartini, J. C. (1980a).** The pathomorphology of malignant catarrhal
11 fever. I. Generalized lymphoid vasculitis. *Vet Pathol* **17**, 58-72.
- 12 20. **Liggitt, H. D. & DeMartini, J. C. (1980b).** The pathomorphology of malignant catarrhal
13 fever. II. Multisystemic epithelial lesions. *Vet Pathol* **17**, 73-83.
- 14 21. **Mariner, J.M. Lantz, V. Waldmann, T.A. & Azimi, N. (2001).** Human T cell
15 lymphotropic virus type I Tax activates IL-15R alpha gene expression through an NF-
16 kappa B site. *J Immunol* **166**, 2602-2609.
- 17 22. **Marks-Konczalik, J. Dubois, S. Losi, J.M. Sabzevari, H. Yamada, N. Feigenbaum, L.**
18 **Waldmann, T.A. & Tagaya, Y. (2000).** IL-2-induced activation-induced cell death is
19 inhibited in IL-15 transgenic mice. *Proc Natl Acad Sci (USA)* **97**, 11445-11450.
- 20 23. **McInnes, I.B., Leung, B.P., Sturrock, R.D., Field, M. & Liew, F.Y. (1997).** IL-15
21 mediates T-cell-dependent regulation of TNF-alpha production in RA. *Nature Med* **3**,
22 189-195.
- 23 24. **McInnes, I.B. & Liew, F.Y. (1998).** IL-15: a proinflammatory role in RA synovitis.
24 *Immunol Today* **19**, 75-79.

- 1 25. **Miranda-Carus, M-E., Benito-Miguel, M., Llamas, M.A., Balsa, A. & Martin-Mola,**
2 **E. (2005).** Human T cells constitutively express IL-15 that promotes ex vivo T cell
3 homeostatic proliferation through autocrine /juxtacrine loops. *J Immunol* **175**, 3656-3662.
- 4 26. **Ohteki, T. Suzue, K. Maki, C. Ota, T. & Koyasu, S. (2001).** Critical role of IL-15-IL-
5 15R for antigen-presenting cell functions in the innate immune response. *Nature Immunol*
6 **2**, 1138-43.
- 7 27. **Ohteki, T. (2002).** Critical role for IL-15 in innate immunity. *Curr Mol Med* **2**, 371-80.
- 8 28. **Oppenheimer-Marks, N., Brezinschek, R.I., Mohamadzadeh, M., Vita, R. & Lipsky,**
9 **P.E. (1998).** IL-15 is produced by endothelial cells and increases the transendothelial
10 migration of T cells *in vitro* and in the SCID mouse-human rheumatoid arthritis model *in*
11 *vivo*. *J Clin Invest* **101**:1261-72.
- 12 29. **Reid, H. W. (2000).** Malignant catarrhal fever. *Infection Diseases Review* **2**, 20-22.
- 13 30. **Reid, H. W., Buxton, D., Pow, I., Finlayson, J. & Berrie, E. L. (1983).** A cytotoxic T-
14 lymphocyte line propagated from a rabbit infected with sheep associated malignant
15 catarrhal fever. *Res Vet Sci* **34**, 109-13.
- 16 31. **Reid, H. W., Buxton, D., Pow, I & Finlayson, J. (1989).** Isolation and characterisation
17 of lymphoblastoid cells from cattle and deer affected with 'sheep-associated' malignant
18 catarrhal fever. *Res Vet Sci* **47**:90-96.
- 19 32. **Ruchatz, H., Leung, B.P., Wei, X.Q., McInnes, I.B. and Liew, F.Y. (1998).** Soluble IL-
20 15 receptor α chain administration prevents murine collagen-induced arthritis: a role for
21 IL-15 in development of antigen-induced immunopathology. *J Immunol* **160**, 5654-5660.
- 22 33. **Sato, N., Patel, H.J., Waldemann, T.A. & Tagaya, Y. (2007).** The IL-15/IL-15R α
23 on cell surfaces enables sustained IL-15 activity and contributes to the long survival of
24 CD8 memory T cells. *Proc Natl Acad Sci (USA)* **104**, 588-593.
- 25 34. **Schock, A. & Reid, H. W. (1996).** Characterisation of the lymphoproliferation in rabbits
26 experimentally affected with malignant catarrhal fever. *Vet Microbiol* **53**, 111-119.

- 1 35. **Schock, A., Collins, R. A. & Reid, H. W. (1998).** Phenotype, growth regulation and
2 cytokine transcription in Ovine Herpesvirus-2 (OHV-2)-infected bovine T-cell lines. *Vet*
3 *Immunol Immunopathol* **66**, 67-81.
- 4 36. **Simon, S., Li, H., O'Toole, D., Crawford, T.B. & Oaks, J.L. (2003).** The vascular
5 lesions of a cow and bison with sheep-associated malignant catarrhal fever contain ovine
6 herpesvirus 2-infected CD8(+) T lymphocytes. *J Gen Virol* **84**, 2009-13.
- 7 37. **Smith, X.G., Bolton, E.M., Ruchatz, H., Wei, X., Liew, F.Y. & Bradley, J.A. (2000).**
8 Selective blockade of IL-15 by soluble IL-15 receptor alpha-chain enhances cardiac
9 allograft survival. *J Immunol* **165**, 3444-3450.
- 10 38. **Swa, S., Wright, H., Thomson, J., Reid, H. & Haig, D.M. (2001).** Constitutive
11 activation of Lck and Fyn tyrosine kinases in large granular lymphocytes infected with the
12 gamma-herpesvirus agents of malignant catarrhal fever. *Immunology* **102**, 44-52.
- 13 39. **Thonur, L., Russell, G.C., Stewart, J.P. and Haig, D.M. (2006).** Differential
14 transcription of ovine herpesvirus-2 genes in lymphocytes from reservoir and susceptible
15 species. *Virus Genes* **32**, 27-35.
- 16 40. **Tsunobuchi, H. Nishimura, H. Goshima, F. Daikoku, T. Suzuki, H. Nakashima, I.**
17 **Nishiyama, Y. & Yoshikai, Y. (2000).** A protective role of interleukin-15 in a mouse
18 model for systemic infection with herpes simplex virus. *Virology* **275**, 57-66.
- 19 41. **Villadsen, L.S., Schuurman, J., Beurskens, F., Dam, T.N., Dagnaes-Hansen, F., Skov,**
20 **L., Rygaard, J., Voorhorst-Ogink, M.M., Gerritsen, A.F., van Dijk, M.A., Parren,**
21 **P.W., Baadsgaard, O. & van de Winkel, J.G. (2003).** Resolution of psoriasis upon
22 blockade of IL-15 biological activity in a xenograft mouse model. *J Clin Invest* **112**,
23 1571-1580.
- 24 42. **Waldmann, T.A. (2006).** The biology of interleukin-2 and interleukin-15: implications
25 for cancer therapy and vaccine design. *Nature reviews immunology* **6**, 595-601.

- 1 43. **Xiong, C., Hixson, P.M., Mendoza, L.H. & Wayne-Smith, C. (2005).** Cloning and
2 expression of rabbit interleukin-15. *Vet Immunol Immunopathol* **107**, 131-141.
- 3 44. **Wei, X.Q., Orchardson, M., Gracie, J.A., Leung, B.P., Gao, B.M., Guan, H.,**
4 **Niedbala, W., Paterson, G.K., McInnes, I.B. & Liew, F.Y. (2001).** The Sushi domain
5 of soluble IL-15 receptor alpha is essential for binding IL-15 and inhibiting inflammatory
6 and allogeneic responses *in vitro* and *in vivo*. *J Immunol* **167**, 277-82.
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Table 1**Phenotype analysis of the LGLs and control T cells used in the study (% positive cells).**

Cells*	CD2	CD5	CD4	CD8	CD21	IgM	$\gamma\delta$ T	BLT- esterase
Control bovine T cells	50	99	43	18	<1	nd	32	48
Control rabbit T cells	nd	97	19	76	nd	<2	nd	34
BJ1035 OvHV-2+ bovLGL	98	98	<1	<1	<1	nd	<1	98
BJ1104 OvHV-2+ bovLGL	97	98	<2	93	<2	nd	5	94
BJ1044 OvHV-2+ bovLGL	96	96	32	56	<1	nd	11	nd
BJ1196 OvHV-2+ bovLGL	98	nd	<1	52	nd	nd	<1	nd
BJ1857 OvHV-2+rabLGL	nd	98	<1	76	nd	<1	nd	95
BJ1859 AIHV-1+ rabLGL	nd	50	<1	50	nd	<1	nd	83
BJ1860 AIHV-1+ rabLGL	nd	<1	<1	45	nd	<1	nd	88
BJ 1263 AIHV-1+ rabLGL	nd	99	2	50	nd	<1	nd	89

*Bov = bovine, rab = rabbit, nd = not done. All LGL exhibited the morphology of large granular lymphocytes and >91% of all LGL line cells were positive for either OvHV-2 or AIHV-1 antigen by immuno-fluorescence. The LGL lines were positive for OvHV-2 or AIHV-1 DNA by PCR, whereas the control uninfected cells did not contain virus DNA.

FACScan analysis of cells labelled by indirect immunofluorescence technique. Murine monoclonal antibodies used: clone CC42 anti-bovine CD2; clone CC8 anti-bovine CD4; clone CC17 anti-bovine CD5; clone CC63 anti-bovine CD8; clone CC15 anti-bovine $\gamma\delta$ T cell receptor ($\gamma\delta$ TCR); clone CC21 anti-bovine CD21; clone KEN-4 anti rabbit CD4; clone KEN-5 anti-rabbit CD5; clone 12C7 anti-rabbit CD8; clone NRBM anti-rabbit IgM. All from Serotec, U.K. or the Institute for Animal Health, Compton. BLT-esterase cytotoxic enzyme detection *in situ* by cytochemistry as described in Haig *et al.*, 1996.

1 **Table 2: Expression of IL-15 in AIHV-1 MCF compared to OvHV-2 MCF.**

Rab	virus	Lymphoid tissues									Non-lymphoid tissues					
		Appendix			MLN			Spleen			Kidney		Liver		Lung	
		IL15*	N†	H‡	IL15	N	H	IL15	N	H	IL15	L§	IL15	L	IL15	L
O1	OvHV2	3	+	+	3	+	+	3	-	+	nd	nd	3	+	nd	nd
O2	OvHV2	nd	nd	nd	3	+	+	2	-	+	1	-	1	-	1	-
O3	OvHV2	nd	nd	nd	3	+	+	3	-	+	2	+	1	-	1	-
O4	OvHV2	3	+	+	3	+	+	3	-	+	0	-	2	+	2	+
O5	OvHV2	3	+	+	3	+	+	3	-	+	1	+	3	+	1	-
O6	OvHV2	3	+	+	nd	nd	nd	3	-	+	0	+	2	+	2	+
O7	OvHV2	3	+	+	3	+	+	1	-	+	1	-	2	+	3	+
O8	OvHV2	3	+	+	3	+	+	2	-	+	0	-	1	+	3	+
A1	AIHV1	1	-	+	1	-	+	1	-	+	1	++	1	++	1	+
A2	AIHV1	1	-	+	1	-	+	1	-	+	1	++	-	+	1	++
A3	AIHV1	0	-	+	0	-	+	nd	nd	nd	0	++	2	++	2	++
A4	AIHV1	1	-	+	1	-	+	1	-	+	0	++	1	++	2	+
A5	AIHV1	0	-	+	0	-	+	3	-	+	1	+	3	++	3	++
A6	AIHV1	1	-	+	1	-	+	2	-	+	1	++	3	++	2	++
A7	AIHV1	1	-	+	1	-	+	3	-	+	0	+	3	++	3	++
A8	AIHV1	0	-	+	0	-	+	1	-	+	0	-	2	++	2	++
C1	control	0	-	-	1			1	-	-	0	-	-	-	-	-
C2	control	1	-	-	0			1	-	-	0	-	-	-	-	-
C3	control	1	-	-	1			2	-	-	0	-	-	-	-	-

2 *Total IL-15+ cells (>90% with lymphocyte morphology) scored as follows: 0 = <10/mm²; 1 = 10-
3 50/mm²; 2 = 50-200/mm²; 3 = >200/mm². †Areas of necrosis (N): - = none; + = areas of necrosis seen.
4 ‡Hyperplasia (H): - = no change compared to uninfected control tissue; + = increase with respect to
5 control uninfected tissue. §In non-lymphoid tissues, lymphoid cell accumulations (L) were scored as
6 follows: - = none; + = small infrequent accumulations; ++ = larger, more frequent accumulations. Rab =
7 rabbit, nd = not done.

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1 **Figure Legends**

2 **Figure 1.** Rabbit LGL and control T cell proliferation and BLT-esterase expression in culture
3 with interleukin-15 and interleukin-2. 5×10^5 LGL or control T cells/ml medium seeded on day 0
4 and cultured with the cytokines for 6 days. In the absence of IL-15 or IL-2, LGLs declined in
5 number over the 6 day period.

6
7 **Figure 2.** Bovine LGL and control T cell proliferation and BLT-esterase expression in culture
8 with interleukin-15 and interleukin-2. 2.5×10^5 LGL or control T cells/ml medium seeded on day
9 0 and cultured for 6 days. In the absence of IL-15 or IL-2, LGLs declined in number over the 6
10 day period.

11
12 **Figure 3.** IL-15 mRNA detected by *in situ* hybridisation in MCF-affected tissues, but not in an
13 LGL line. (A) AIHV-1+ LGL line 1859. (B) A positive control for (A) processed at the same
14 time, which shows IL-15mRNA+ mononuclear cells in the interstitial area of a MLN from an
15 OvHV-2-infected rabbit (O4 in Table 2). (C) Sense RNA probe, control for (B) at a lower
16 magnification. (D) H&E stained section of an AIHV-1-infected rabbit liver (A6 in Table 2)
17 showing a lymphocyte accumulation lesion. (E) serial section of (D), sense RNA probe negative
18 control for (F) which is a serial section of (D) showing antisense RNA probe detection of IL-
19 15mRNA+ mononuclear cells in the lymphocyte accumulation lesion.

20
21 **Figure 4.** Immuno-histochemistry detection of IL-15 antigen-expressing cells in MCF-affected
22 tissues using the 247mab. (A) Detail of an area of an AIHV-1-infected rabbit appendix (A2 in
23 Table 2), showing a cluster of three IL-15+ mononuclear cells (arrowhead). (B) IL-15+
24 mononuclear cells in the interstitial region of an appendix of an OvHV-2-infected rabbit (O4 in
25 Table 2). (C) Lower power magnification of (B) for a comparison with (D), a serial section of
26 (B/C) that shows T cells labelled with the pan-T mab. There is <10% overlap of +ve cells in this

1 comparison. (E) IL-15 antigen associated with appendix dome epithelium (arrowheads) from an
2 OvHV-2-infected rabbit (O8 in Table 2). (F) Control for (E) at a lower magnification using the
3 IHC labelling technique in the absence of primary 247 mab. N=area of necrosis.

4

5 **Figure 5.** Western blot for IL-15 detection in MLN lysates from uninfected control and virus-
6 infected rabbits. (A) 247 mab western blot showing IL-15 (M.Wt. 16kDa) in MLN cell lysates
7 from a control uninfected rabbit (C1 in Table 2), two AIHV-1-infected rabbits (A4 and A7 in
8 Table 2) and two OvHV-2-infected rabbits (O2 and O8 in Table 2). (B) The same samples as in
9 (G) probed with the murine sIL-15 receptor. The presence of a higher M.Wt. band shown here is
10 sometimes but not always seen in the IL-15 western blots with either sIL-15R or the 247 mab. All
11 lanes received 25ug protein.

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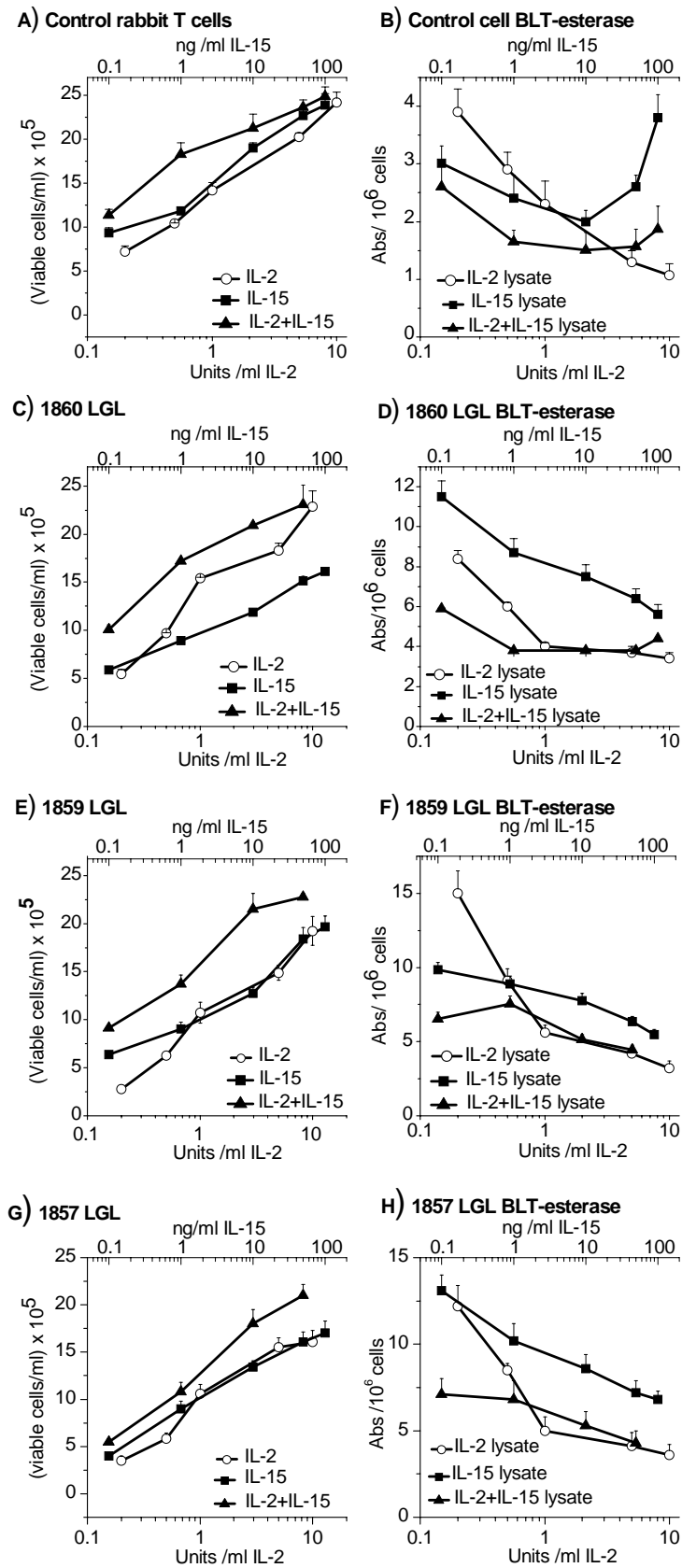
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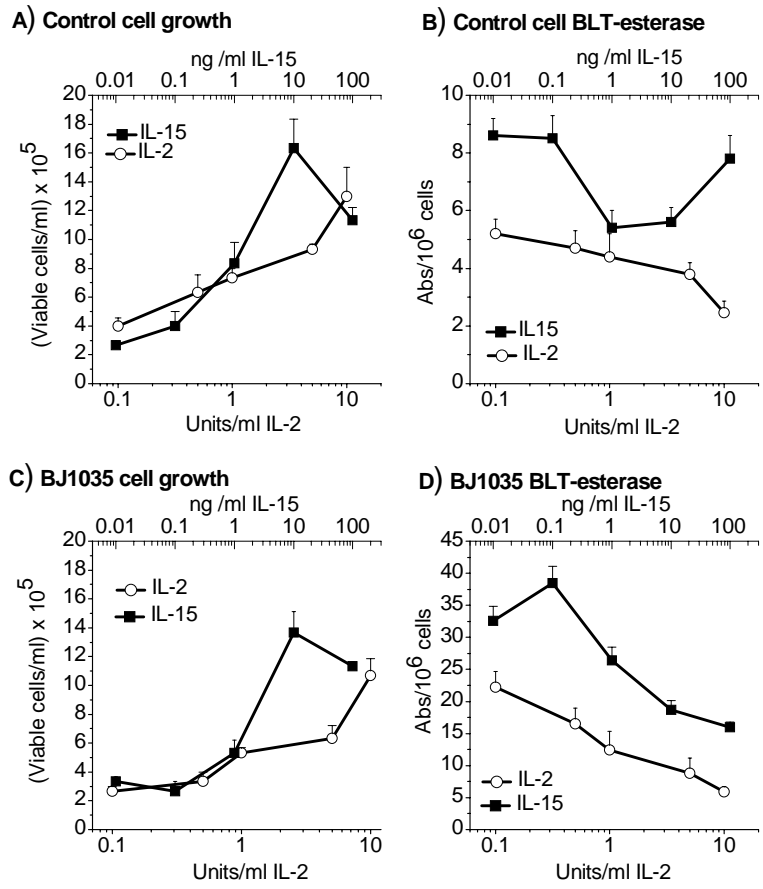
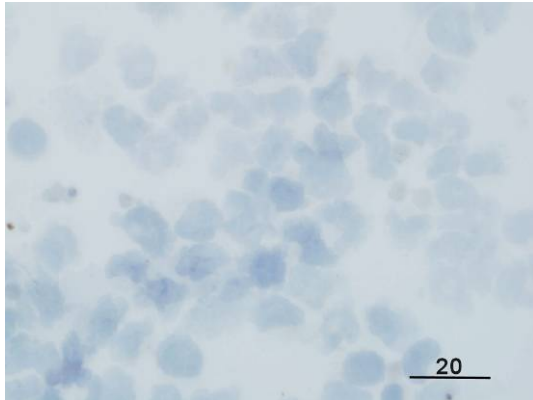


Figure 2.

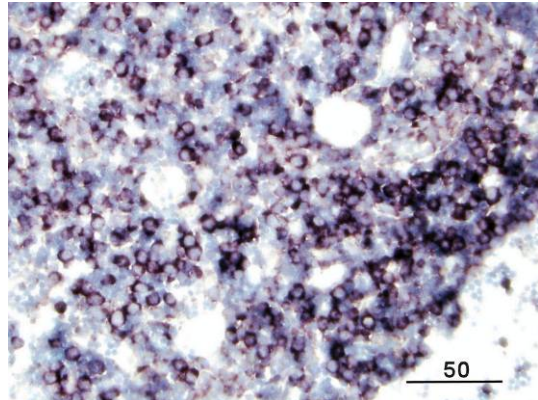
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Figure 3

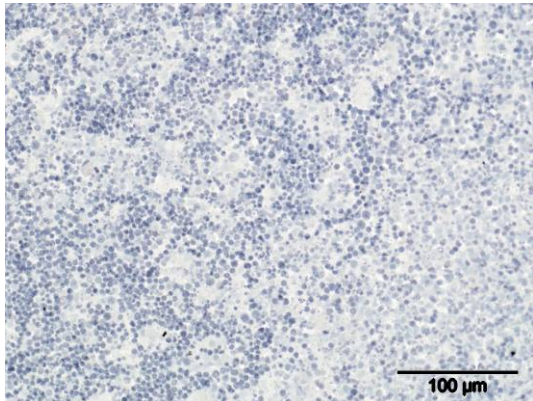
A)



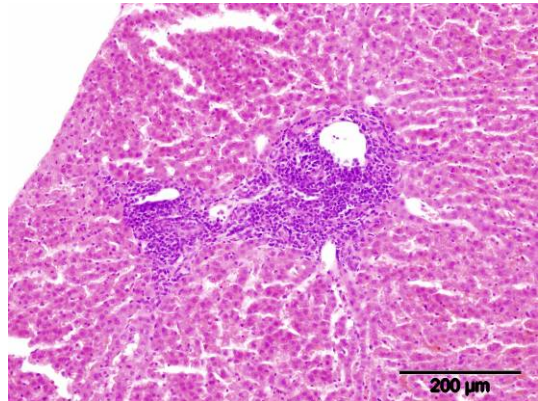
B)



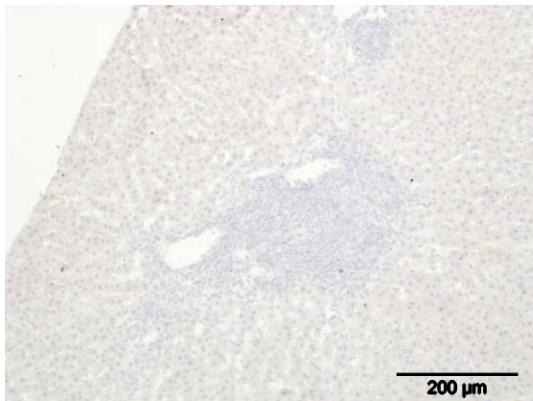
C)



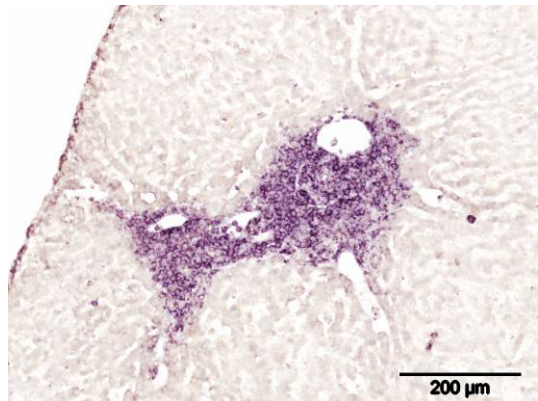
D)



E)



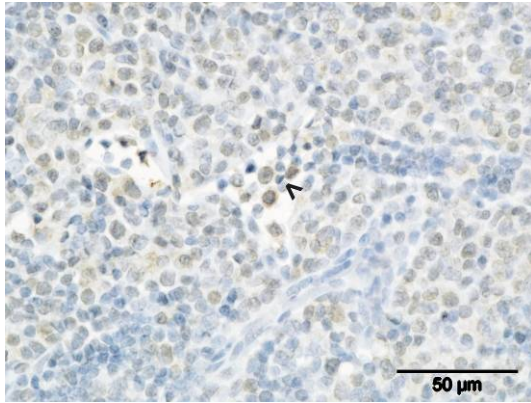
F)



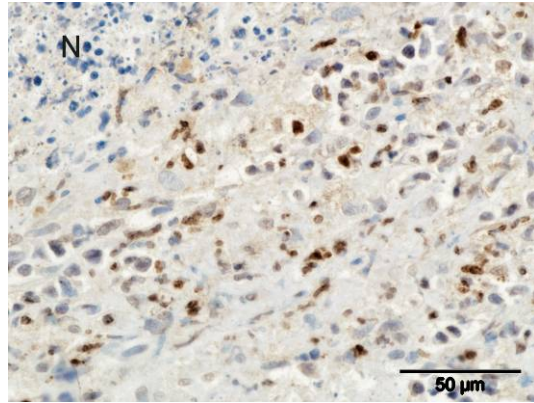
- 1
- 2
- 3
- 4
- 5

Figure 4

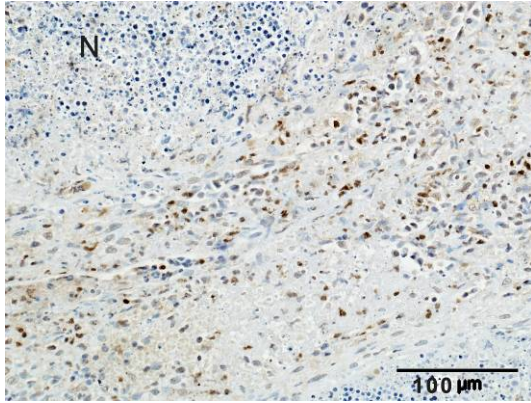
A)



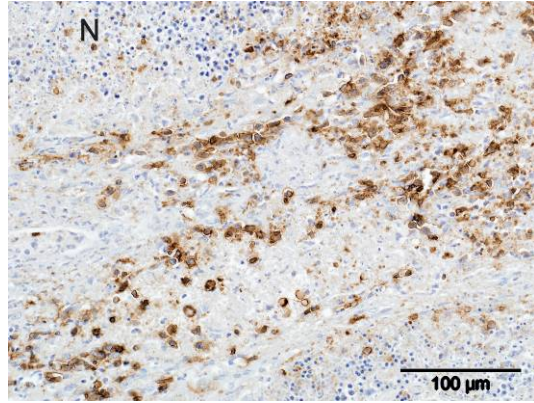
B)



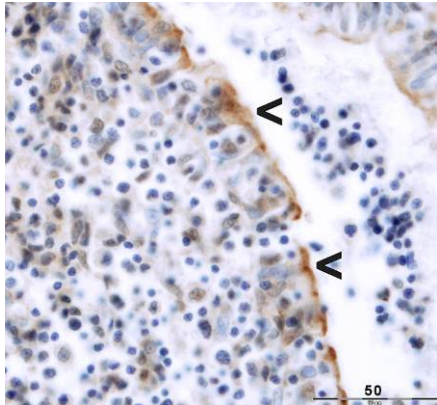
C)



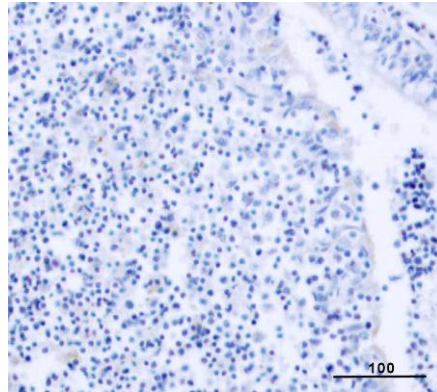
D)



E)



F)



1

2

3

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1 Figure 5

