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Author manuscript

*Vet Immunol Immunopathol*. Author manuscript; available in PMC 2016 April 15.

Published in final edited form as:

*Vet Immunol Immunopathol*. 2015 April 15; 164(0): 127–136. doi:10.1016/j.vetimm.2015.02.001.

## **CHEMOKINE RECEPTOR 7 (CCR7)-EXPRESSION AND IFN**γ **PRODUCTION DEFINE VACCINE-SPECIFIC CANINE T CELL SUBSETS**

**Ashley N. Hartley**1,2,3 and **Rick L. Tarleton**1,4

<sup>1</sup>Center for Tropical and Emerging Global Diseases, University of Georgia, Athens, GA, 30602, USA

<sup>2</sup>Department of Infectious Diseases, University of Georgia, Athens, GA, 30602, USA

<sup>3</sup>Department of Cellular Biology, University of Georgia, Athens, GA, 30602, USA

## **Abstract**

Canines suffer from and serve as strong translational animals models for many immunological disorders and infectious diseases. Routine vaccination has been a mainstay of protecting dogs through the stimulation of robust antibody responses and expansion of memory T cell populations. Commercially available reagents and described techniques are limited for identifying and characterizing canine T cell subsets and evaluating T cell-specific effector function. To define reagents for delineating naïve versus activated T cells and identify antigen-specific T cells, we tested anti-human and anti-bovine T-cell specific cell surface marker reagents for cross-reactivity with canine peripheral blood mononuclear cells (PBMCs. Both  $CD4<sup>+</sup>$  and  $CD8<sup>+</sup>$  T cells from healthy canine donors showed reactivity to CCL19-Ig, a CCR7 ligand, and coexpression with CD62L. An *in vitro* stimulation with concanavalin A validated downregulation of CCR7 and CD62L expression on stimulated healthy control PBMCs, consistent with an activated T cell phenotype. Anti-IFN $\gamma$  antibodies identified antigen-specific IFN $\gamma$ -producing CD4<sup>+</sup> and CD8<sup>+</sup> T cells upon *in vitro* vaccine antigen PBMC stimulation. PBMC isolation within 24 hours of sample collection allowed for efficient cell recovery and accurate T cell effector function characterization. These data provide a reagent and techniques platform via flow cytometry for identifying canine T cell subsets and characterizing circulating antigen-specific canine T cells for potential use in diagnostic and field settings.

#### **Conflict of Interest Statement**

The authors declare they have no competing interests.

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Address correspondence and reprint requests to Dr. Rick L. Tarleton, Center for Tropical and Emerging Global Diseases, University of Georgia, 500 D.W. Brooks Drive, N310 Coverdell Center, University of Georgia, Athens, GA 30602. Telephone: 706-542-3362

Fax: 706-542-3582 tarleton@uga.edu.<br><sup>3</sup>Current address: Small Animal and Clinical Science, College of Veterinary Medicine, University of Tennessee, Knoxville, TN, 37996, USA;

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## **Keywords**

dog; T cell; CCR7; CD62L; flow cytometry; vaccine

## **2. Introduction**

Domestication and tractability have allowed do gs to serve as research subjects for caninespecific diseases as well as models for human disorders. In particular, dogs serve as robust translational models in cardiovascular (Hohnloser et al., 2009), neoplastic (Khanna et al., 2006; Klopfleisch et al., 2010), immunological (Creevy et al., 2003; Marsella and Girolomoni, 2009), neurological (Awano et al., 2009; Selkoe et al., 1987), and genetic (Wilbe et al., 2010) research studies. Canines are also susceptible to and serve as models of zoonotic diseases such as leishmaniasis and American trypanosomiasis and hence used to evaluate anti-parasitic chemotherapeutic regimens (Guedes et al., 2002). Routine vaccination in canines allows an opportunity to assess the development of an appropriate immunological response to foreign antigens. Techniques and commercially available reagents are scarce for studying the canine immune system, especially as compared to those available for humans. As basic research pursues translational applications in animals more physiologically similar to humans, and veterinary medicine strives for more individualized patient therapies, an increasing need exists for identifying, characterizing, and monitoring the canine immune response.

The First International Canine Leukocyte Antigen Workshop (CLAW) was a significant step in identifying canine homologs of human CD antigens that delineated leukocyte populations by monoclonal antibodies (Cobbold and Metcalfe, 1994). Clusters of antibodies collected from several sources identified canine equivalents of CD4, CD8, and Thy1.1 antigens from peripheral blood. Additional antibodies reactive to canine leukocyte antigens including CD45R (Aguiar et al., 2005) CD45RA (Caniatti et al., 1996), CD11/CD18 (Danilenko et al., 1992a; Moore et al., 1990), and CD62L (Crockett-Torabi and Fantone, 1997) and to platelet and erythrocyte antigens (Schuberth et al., 2007) have been described separately from the CLAW workshop. Testing of monoclonal antibodies specific for cytokines in other species have also identified IL-4-, IL-8-, and IFN-γ-producing canine PBMCs and expanded the repertoire of canine specific reagents (Pedersen et al., 2002). However, despite these advances, delineating and characterizing naïve, activated, and memory T cell subsets in canines has remained limited.

The aim of this project was to identify and validate immunological reagents for characterizing canine T cells through phenotypic and effector function evaluation-based assays. Detection of the canine cross-reactive CCL19-hIg, a ligand for CCR7, identified naïve and antigen-experienced but not recently activated canine T cells. CCR7 cell surface expression was consistent with CD62L, an L-selectin expressed by naïve and central memory T cells during homing to secondary lymphoid organs. Decreases in CCR7 and CD62L expression following antigen stimulation or mitogen activation correlated with upregulation of the activation marker, CTL2.58, and delineated activated T cells. IFN $\gamma$ production following PBMC whole vaccine stimulation defined antigen-specific T cell

effector function. Extended time between blood collection and PBMC isolation of up to twenty-four hours revealed no significant loss in identifying vaccine-specific IFNγproducing T cells. These data provide a reagent platform for identifying and characterizing canine T cell populations and assessing antigen-specific effector function.

## **3. Materials and Methods**

#### **3.1. Animals and isolation of mononuclear cells**

Approximately 40–50mls of blood from four clinically healthy adult (>3 years of age) mixed breed dogs were drawn into heparinized tubes (Vacutainer, Becton-Dickinson, Franklin Lakes, NJ, USA) by venipuncture. Isolation of peripheral blood mononuclear cells (PBMCs) occurred immediately following collection or as otherwise indicated and as previously described for human subjects (Albareda et al., 2009). PBMCs were washed in Hank's buffered balance salt solution (Mediatech Inc., Manassas, VA, USA) and resuspended in RPMI-1640 (Mediatech Inc.) completed with 50uM 2-β-mercaptoethanol, 2mM Lglutamine, 25µg/mL gentamicin, 200U/mL penicillin (Mediatech Inc), 2µg/mL streptomycin (Mediatech Inc), 1mM sodium pyruvate, and 10% heat-inactivated (30min, 56°C) and aggregate-removed (800gx30min) fetal calf serum (HI-FCS) (HyClone Laboratories, ThermoScientific, Logan, UT, USA). Resuspended cells were frozen in media containing 10% dimethyl sulfoxide (Acros Organics, Fair Lawn, NJ, USA) in liquid nitrogen for longterm storage. Prior to use, PBMCs were recovered, thawed at 37°C, washed and resuspended in complete RPMI-1640 + 10% HI-FCS. These purification, storage, and recovery procedures consistently yielded >95% viability, as determined by microscopic examination of trypan blue dye exclusion. All animal use protocols were approved by the University of Georgia Institutional Animal Care and Use Committee.

#### **3.2. PBMC antibody reactivity**

For testing canine T cells antibody reactivity, a minimum of  $2\times10^5$  PBMCs, were stained at 1:50 antibody dilution in PBS containing 1% BSA and 0.05% sodium azide (PAB; both from Sigma-Aldrich, St. Louis, MO, USA) at 4°C for 45 min. Antibodies tested were those defined in Table 1. Cells were fixed in 2% formaldehyde prior to flow cytometric collection.

#### **3.3. T cell stimulation and proliferation assessment assays**

A total of  $4\times10^5$  PBMCs were stimulated in a 96-well flat-bottom tissue culture plate (Costar, Corning, NY, USA) at 37°C in the presence of media, 15µg/mL anti-canine CD3 (AbD Serotec, Raleigh, NC, USA), or 0.25µg/mL concanavalin A (Sigma-Aldrich) for the indicated days. For assessment of proliferation, PBMCs were washed twice with PBS, incubated with 5µM CFSE (Molecular Probes, Eugene, OR), quenched with FCS, and plated. Two days post-stimulation, centrifuged and washed cells were incubated with or without CCL19-hIg (ELC; (Hargreaves et al., 2001)) supernatant at 4°C for 45 min. Antibodies used were anti-CD8-Pacific Blue, anti-CD62L-PE (AbD Serotec), anti-human IgG-AF488 (Molecular Probes), and a cocktail containing anti-dog Pan T cell-APC, anti-B cell-PE, and anti-dog T cell Activation marker-FITC (Dog Activated T Lymphocyte Cocktail, BD Pharmingen, BD Biosciences, San Jose, CA). 7-amino-actinomycin D (7AAD; BD Pharmingen) was included for live/dead cell discrimination. Cells were stained in PAB for 45 min at 4°C, washed, and fixed in 2% formaldehyde.

#### **3.4 Intracellular cytokine staining**

For assaying IFN $\gamma$  levels,  $4\times10^5$  PBMCs were stimulated for 5hr in the presence of 2ng/mL phorbol 12-myristate 13-acetate (PMA),  $4\mu$ g/mL Ca<sup>2+</sup> ionomycin (Sigma-Aldrich), and brefeldin A (BD GolgiPlug; BD Biosciences) (Pedersen et al., 2002). For polyclonal activation, PBMCs were plated with 15µg/mL anti-CD3 (AbD Serotec) or diluted whole vaccine antigens, incubated overnight at 37°C and brefeldin A added 5hr prior to end of incubation. Canine vaccines were IMRAB 3TF (Rabies; Merial, Athens, GA, USA), Duramune 5 (Canine distemper-Adenovirus Type 2-Parainfluenza-Parvovirus (DAPP); Fort Dodge Animal Health, Fort Dodge, IA, USA), and Leptovax 4 (Leptospirosis bacterial extract; Fort Dodge Animal Health) vaccines. Cells were stained with anti-CD8-Pacific Blue and anti-CD4-FITC (AbD Serotec) followed by intracellular staining with anti-bovine anti-IFN-γ AF647 (AbD Serotec) according to the BD Cytofix/Cytoperm kit (BD Biosciences).

#### **3.5 Flow Cytometry and Analysis**

A minimum of 250,000 events were collected for each sample on a CyAn ADP using Summit, version 4.3 (Beckman Coulter, Fullerton, CA, USA). Where appropriate, fluorescence minus one controls were used to establish gating measures. FlowJo flow cytometry analysis software, version 9 (Tree Star, Ashland, OR, USA) was used for analyses.

## **4. Results**

As  $CD4^+$  and  $CD8^+$  T cells play critical roles in various infections, our primary efforts focused on identifying T cells and discriminatory surface markers for naïve, activated, and memory T cells subsets. A reagent panel used for testing of cross-reactivity with canine PBMCs included antibodies purchased commercially or kindly provided by other investigators (Table 1). A reagent was judged as positive if reactive with  $>2\%$  of PBMCs. Utilizing this strategy, fifteen of twenty-six reagents were cross-reactive with canine PBMCs as denoted in Table 1, with eleven of twelve (91.7%) previously referenced reactive antibodies confirmed in this study. If a reagent distinguished distinct cell populations individually or following incubation in a T cell stimulation protocol, the reagent was found to be discriminatory, as noted in Table 1.

For antibodies recognizing putative cell surface markers on T cells, we repeated staining to validate marker expression on  $CD4^+$  and  $CD8^+$  T cells. Canine T cell reactivity was found with CCL19-Ig (ELC), a fusion protein serving as a chemokine ligand of CCR7 (Hargreaves et al., 2001) and previously described to have reactivity with naïve and central memory ovine T cells (Debes et al., 2005). As circulating T cells traffic from the blood and home to lymph nodes, naïve and central memory T cells express CCR7 and other adhesion molecules, like L-selectin (CD62L) and LFA-1, to tether and arrest to endothelium for transmigration across high endothelial venules. To investigate if circulating canine CD4<sup>+</sup> and CD8+ T cells express both CCR7 and L-selectin (CD62L), PBMCs were stained with

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anti-CD4, anti-CD8, CCL19-Ig, and anti-CD62L. In clinically healthy canine donors, CD4<sup>+</sup> and CD8+ T cells expressed measureable levels of CCR7 (Fig 1A). Unlike CCR7 expression in human CD8+ T cells (Campbell et al., 2001), CCR7 expression by canine CD8+ T cells varied greatly between individual dogs. In comparison to humans and sheep where >85% of CD4+ T cells express CCR7 (Campbell et al., 2001; Debes et al., 2005), approximately 40– 50% of canine CD4+ T cells from healthy donors expressed CCR7. The observed difference is likely explained by the decreased ability of CCL19-hIg to delineate distinct peaks associated with CCR7+ and CCR7− canine CD4+ T cells populations (Fig 1A) versus a difference in species or health status of the hosts. Exploration of alternative secondary fluorophore combinations yielded similar CCR7+CD4+ T cell percentages (not shown). Both canine  $CD4^+$  and  $CD8^+$  T cells also showed reactivity to the lymph node homing marker CD62L (Fig 1B) and this expression agreed with that of CCR7 (Fig 1C), as the predominance of CD4<sup>+</sup>CD62L<sup>hi</sup> T cells expressed CCR7, results consistent with that observed in human peripheral blood (Campbell et al., 2001). Taken together, the cell surface markers of CCR7 and CD62L identified canine CD4<sup>+</sup> and CD8<sup>+</sup> T cell populations consistent with the phenotype of naïve or central memory T cells.

To evaluate if CCR7 a nd other cross-reactive antibodies identified in Table 3.1 were discriminatory for naïve versus activated T cells, a method and assay for T cell activation was used. For this purpose, CFSE-labeled canine PBMCs were incubated in media alone or with the T cell mitogen concanavalin A (ConA), and stimulation and proliferation was evaluated by changes in cell size and CFSE dilution. A comparison of forward versus side scatter revealed PBMC stimulated with ConA were on average volumetrically larger than unstimulated cells, consistent with cellular activation (Fig 2A). CFSE dilution indicated cell division of ConA-stimulated  $CD8<sup>+</sup> T$  cells as early as three days post stimulation (Fig 2B).

As we proposed the CCR7 and CD62L reagents identified naïve and central memory canine T cells, we hypothesized that following T cell activation canine T cells would decrease expression of CCR7 and CD62L. To address this hypothesis, canine PBMCs were cultured in the presence of anti-CD3, ConA, or media alone. Even before detectable proliferation by CFSE dilution two days post stimulation (Fig 3A), CD8+ T cells expressed less CCR7 (Fig 3B) and CD62L (Fig 3C) when activated. Expression levels of these two markers were consistent with the strength of stimulus, as ConAstimulated CD8+ T cells expressed less CCR7 and CD62L than cells stimulated with anti-CD3. Co-expression of both CCR7 and CD62L also decreased in stimulated cells, with ConA-stimulated cells showing greater decreases than cells incubated with anti-CD3 (Fig 3D). Down-regulation of CCR7 and CD62L cell surface expression further correlated with increased expression of the activation marker (CTL2.58<sup>+</sup>) on stimulated T cells (Fig 3E) and CD8<sup>+</sup> T cells (Fig 3F). These experiments identified markers for defining activated CTL2.58+CD8+ T cells and naïve or central memory-like CCR7+CD62LhiCD8+ T cells in dogs. Testing of other canine T cell reactive reagents listed in Table 1, including CD45RA, CD45RO, and CD11a, in media versus stimulatory conditions did not alter antibody T cell reactivity in our stimulation assay or were not detectable using secondary antibody reagents and therefore were not investigated further. These results are noted as "not discriminatory" in Table 1.

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Cognate antigen recognition through the TCR results in effector cytokine production, and thereby provides a method for evaluating and delineating antigen-specific T cells. To assay the production of the effector cytokine IFN $\gamma$  by canine T cells in response to previously seen antigens, PBMCs were isolated from three healthy dogs vaccinated three weeks prior with IMRAB 3TF (Rabies), Duramune 5 (Canine distemper-Adenovirus Type 2-Parainfluenza-Parvovirus (DAPP)), and Leptovax 4 (Leptospirosis bacterial extract) vaccines. Following incubation of cells with media or vaccine antigen overnight, intracellular  $IFN<sub>Y</sub>$  staining was performed to identify vaccine-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cells. CD4<sup>+</sup> T cells from three of three dogs produced IFN $\gamma$  in response to rabies vaccine, whereas only one of three had IFN $\gamma$ -producing cells when stimulated with DAPP vaccine antigens (Fig 4A). CD8<sup>+</sup> T cells from two of three dogs produced IFN $\gamma$  in response to rabies vaccine, but all appeared to not be significantly different from media controls when stimulated with the DAPP vaccine (Fig 4B). Despite differences in vaccine sources and stimulus concentrations, the low levels of detectable of IFNγ-producing T cells following canine distemper vaccine (DAPP) stimulation were consistent with previously published ELISAs of stimulated PBMC culture supernatants (Valli et al., 2010). Both  $CD4^+$  and  $CD8^+$  T cell subsets failed to produce robust IFNγ in response to the Leptovax 4 vaccine, suggesting this vaccine failed to elicit a strong T cell specific immune response three weeks post administration, consistent with the predominant humoral response elicited by the bacterial antigens of the leptospiral vaccine (Adler and Faine, 1977). These results provide a flow cytometric method for identifying antigen-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cells and evaluating IFN $\gamma$  production for individual canine T cells.

In fieldwork settings, circumstances may arise where PBMC isolation will occur hours following sample collection, as is the case in field sampling. We asked how this timing would affect sample quality and ability to identify antigen-specific and cytokine-producing T cells. To address this question, we determined recovery and effector function of T cells isolated at various times post-blood collection. Incubation of cells with media or  $PMA/Ca^{2+}$ ionomycin followed by intracellular cytokine staining revealed no significant loss in quantities or percentages of IFNγ-producing T cells isolated up to 24 hours post collection (Fig 5). In PBMCs purified 48 hours following venipuncture, a five-fold reduction in recovery of both  $CD4^+$  and  $CD8^+$  T cell as compared to the 24 hours post collection time point was observed. Additionally, unstimulated CD4+ T cells produced significantly higher levels of IFNγ. These data provide an optimal method for PBMC isolation for application to fieldwork settings and for accurate evaluation of T cell effector function.

## **5. Discussion and Conclusion**

T cells play a central role in the initiation and regulation of the immune response. T cells modulate cell surface expression of ligands and receptors to execute effector function and communicate with the environment. T-cell receptor complex signaling and CD28 costimulation results in naïve T cells downregulating lymph node homing receptors (CD62L and CCR7) while upregulating expression of cell adhesion molecules (CD44), T-cell receptor signaling (CD45RO), and early activation (CD69) antigens. Reliable expression of these cell surface molecules following cognate antigen encounter affords T cell monitoring during disease processes. Diagnosis and progression of human T cell lymphoproliferative

disorders have been examine through flow cytometic immunophenotyping. Similar individualized diagnostics are being pursued for veterinary patients, but the paucity of literature describing the detection and characterization of antigen-driven T cell responses limits reaching this goal.

To address the growing need of canine immune system characterization for individualized diagnoses and toward furthering dogs as suitable translational research models, we aimed to identify and validate canine-specific reagents for discriminating naïve versus activated T cells. Novel canine T cell reactivity of CCL19-Ig defined CCR7 expression in CD4+ and CD8+ T cells of healthy donors. CCR7 co-expression with CD62L and decreased expression following stimulation validated these markers for defining naïve or central memory CD8+ T cells. The activation marker CTL2.58 further delineated activated T cells. Combining these reagents into a single panel would allow for identification of naïve or central memory T cells as CCR7<sup>+</sup>CD62L<sup>hi</sup>CTL2.58<sup>−</sup> and activated T cells as CCR7<sup>−</sup>CD62L<sup>lo</sup>CTL2.58<sup>+</sup>. To our knowledge, these results are the first to describe and validate these markers for identifying naïve and activated canine T cell subsets.

With cognate antigen recognition stimulating cytokine-mediated T cell effector function, identification of IFNγ-producing T cells is critical for evaluating appropriate T cell responses. Monitoring antigen-specific T cell responses induction is a critical component for evaluating efficacy of T cell-mediated vaccines, especially in the era of canine oral malignant melanoma DNA vaccines (Grosenbaugh et al., 2011) where cytotoxic T cells would be proposed to mediate protection. Here, we described a method for defining vaccinespecific  $CD4^+$  and  $CD8^+$  T cells through intracellular cytokine staining following rabies, canine distemper-adenovirus-parainfluenza-parvovirus, and leptospirosis bacterial extract vaccination. Determining induction, kinetics, and effector function of these antigen-specific T cells are ideal for evaluating appropriate and protective vaccine responses and defining goals for future vaccines, where T cells are primary mediators of immune control for the targeted disease.

In instances where immediate PMBC isolation may be unavailable due to geographical location, evaluating appropriate storage conditions, subsequent cell isolation, and cytokine production following extended times post blood sampling is critical. For example, immunophenotyping of lymphoma of a dog living in a rural area or research studies involving characterization of T cell responses of dogs living in an international locale would likely require significant time between blood collection and PBMC isolation and storage. Data herein illustrates these canine subjects can be examined and immune responses characterized, expanding the translational application of this study beyond laboratory settings. These techniques and reagents are powerful in scenarios where human vaccination efforts would be unethical or inaccessible and dogs serve as models of natural infection and appropriate targets for control efforts, such as American trypanosomiasis (Padilla et al., 2010).

In conclusion, limited resources and techniques for phenotyping and characterizing canine T cells have restricted canine adaptive immune system studies. The tools defined here provide a platform for defining T cell subsets and identifying circulating antigen-specific T cells.

The impact of this work for improving canine health and facilitating the translational applications of dogs for human disease modeling is promising. As individualized T cell immunophenotyping becomes more accessible for human patients, application to veterinary medicine and animal patients is encouraging.

## **Acknowledgements**

We greatly acknowledge Lisa Reynolds of the University of Georgia College of Veterinary Medicine and Julie Nelson of the Center for Tropical and Emerging Global Diseases Flow Cytometry Facility at the University of Georgia, and members of the Tarleton Research Group for helpful discussion. ANH was supported in part by an NIH training grant T32 AI-060546 to the CTEGD. This work was supported by NIH Grant R01 AI089952 awarded to RLT.

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**Figure 1. Co-expression of CCR7 and CD62L on canine CD8+ and CD4+ T cells** Canine PBMCs from two clinically healthy dogs (#1 and #2) were incubated for 1 hour at 37°C and stained with or without CCL19-hIg (ELC), a CCR7 ligand. Cells were stained with anti-CD8, anti-CD4, anti-CD62L, anti-human secondary antibody, and 7AAD. Percentages of CD8+7AAD− and CD4+7AAD− T cells expressing CCR7 (A) and CD62L (B). Shaded lines are fluorescence-minus-one (FMO) controls for the indicated marker. C, Percentages of CD8+7AAD− or CD4+7AAD− T cells co-expressing CCR7 and CD62L. Gating strategies are determined by FMO controls, especially noting the highest CCR7 peak present in experimental and FMO control samples is due to nonspecific secondary antibody binding.

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#### **Figure 2. Measuring stimulation and proliferation using CFSE time-series data**

A, Forward scatter (fsc) versus side scatter plots (ssc) of CFSE-labeled PBMCs incubated with media or concavalin A (ConA) for 4 days. Gated cells are displayed in the fsc histogram, comparing ConA-stimulation (blue line) to media alone (grey). B, CFSE-labeled PBMCs were harvested on the indicated days and stained with 7AAD and anti-CD8 antibodies. Histograms represent percentages of CD8+7AAD− T cells with CFSE dilution following incubation with media (grey shaded) versus ConA (green line).

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**Figure 3. CCR7, CD62L, and CTL2.58 expression of canine PBMCs in stimulation conditions** Canine PBMCs incubated two days with media, anti-CD3, or ConA, harvested, and stained with various antibody combinations. A, Proliferation of CFSE-labeled CD8<sup>+</sup> T cells. B–D, PBMCs incubated with (blue line) and without (grey shaded) CCL19-hIg and stained with anti-human secondary antibody and anti-CD62L. Histograms of CCR7 (B) and CD62L (C) expression of CD8<sup>+</sup> T cells. D, Percentages of CD8<sup>+</sup> T cells co-expressing CCR7 and CD62L. E–F, PBMCs stained with the dog activated T lymphocyte antibody cocktail, 7AAD, and anti-CD8. Percentages of stimulated  $PanT^+(E)$  and  $CD8^+(F)$  T cells reactive to the CTL2.58 antibody, an activation marker.

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#### **Figure 4. IFN**γ **production by canine CD4+ and CD8+ T cells upon stimulation with vaccine antigens**

PBMCs collected from three dogs three weeks post vaccination with IMRAB 3TF (Rabies), Duramune 5 (DAPP), and Leptovax 4 (Lepto) were incubated with media,  $PMA/Ca^{2+}$ ionomycin, and vaccines. GolgiPlug was added 5hr prior to end of incubation. Cells were stained with anti-CD8, anti-CD4, and anti-bovine IFNγ. A–B, Numbers indicate percentages of  $CD4^+$  (A) and  $CD8^+$  (B) T cells producing cytokine in each condition.

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## **Figure 5. Viability and effector function of canine CD4+ and CD8+ T cells measured over a 48hr period**

Canine blood was collected, incubated on ice, and PBMCs isolated at the indicated time points. PBMCs were incubated at 37°C with PMA/ionomycin or media alone in the presence of GolgiPlug for 5 hours. Cells were stained with anti-CD8 and anti-CD4, permeabilized, and stained for IFNγ. Numbers indicate percentage (gated) and count (bottom right) of  $CD8<sup>+</sup>$  (A) and  $CD4<sup>+</sup>$  (B) T cells producing IFN<sub>Y</sub> at each isolation time point.

 Author Manuscript Author Manuscript **Table 1**

Reagent panel used to study canine peripheral mononuclear cells. Reagent panel used to study canine peripheral mononuclear cells.







Abbreviations: MHCI, major histocompatibility complex I; R', receptor; APC, antigen presenting cell.  $\frac{1}{6}$