

Alterations with age in peripheral blood lymphocyte subpopulations and cytokine synthesis in beagles

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Purpose: The immune system is considered to be affected by aging, which is linked to various immune pathogenesises. The purpose of this study was to determine age-associated changes in immune function of healthy dogs (beagles), specifically those of naive and memory T lymphocytes, based on cytokine synthesis.

Patients and methods: Blood samples were obtained from 44 healthy beagles that were divided into three age-groups: young (<4 years), middle-aged (4–8 years), and older dogs (>8 years). Subpopulations of T lymphocytes were determined by flow cytometry. Transcriptional (mRNA) levels of cytokines were determined for primary-cultured leukocytes using quantitative real-time polymerase chain reaction.

Results: There were negative correlations between dogs' ages and the number of peripheral blood mononuclear cells, T cells, and B cells. In particular, the number of naive CD4⁺ CD45RA⁺ T cells and CD8⁺ CD45RA⁺ T cells significantly decreased with age. The mRNA levels for interleukin (IL)-2, IL-2R α , and interferon-gamma were significantly higher in young or middle-aged dogs ($P < 0.05$), whereas IL-4 mRNA expression was not significantly different over the different age-groups. IL-2R γ mRNA expression tended to decrease with age.

Conclusion: Decreases of naive CD4⁺ and naive CD8⁺ T cells may be related to age-related immunosenescence in dogs. With regard to cytokine production, leukocyte IL-4 and IL-10 mRNA levels did not change with age, whereas IL-2, IL-2R α , and IL-2R γ mRNA levels decreased with age. These altered cytokine mRNA expression patterns may contribute to decreased naive T-cell function(s) with aging.

Keywords: aging, leukocyte subpopulation, cytokine, dog, naive T cell, IL-2R

Introduction

The occurrence of age-related disorders, such as tumors in dogs¹ and renal inflammation in cats,² increases with extended life span. The immune system is considered to be affected by aging, which is linked to various immune pathogenesises.^{3,4} The following age-related immunosenescent changes have been reported for dogs: reduced proliferative responses of peripheral blood mononuclear cells (PBMCs);^{5–7} reduced major histocompatibility complex (MHC) class II expression by peripheral blood lymphocytes;⁸ reduced numbers of peripheral blood T and B cells;^{9,10} reduced numbers of CD4⁺ T cells;^{1,11} and reduced CD4/8 T-cell ratios.^{8,12–14} Moreover, compared with young dogs, aged dogs have reduced numbers of circulating naive T cells and increased numbers of memory T cells.¹⁵ Naive T cells are important to mount an immune response to various newly encountered antigens; naive CD4⁺ and CD8⁺ T cells, in particular, are important for defenses against externally derived (bacteria, viruses) and internally derived (neoplastic cells) antigenic challenges, respectively.

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To assess the basic immune functions of T cells, it is useful to determine their cytokine synthesis capability. CD4⁺ T cells are divided into two distinct subsets based on their cytokine production patterns: type 1 (Th1) and type 2 (Th2). Th1 cytokines, such as interleukin (IL)-2 and interferon (IFN)- γ , are required for the generation of cytotoxic T cells and the activation of natural killer cells. IL-2 produced by activated T cells promotes the proliferation, differentiation, and survival of mature T and B cells.¹⁶

We investigated the mRNA expressions of Th1 and Th2 cytokines and the IL-2 receptor (IL-2R) in peripheral blood leukocytes of dogs (beagles) as a function of age. Our purpose was to determine if there were age-associated changes in immune functions in dogs, specifically those of naive and memory T lymphocytes, based on cytokine synthesis.

Materials and methods

Animals

Normal healthy beagles (n = 44) were used for this study. All dogs had been referred to the veterinary teaching hospitals of Kitasato University during 2007–2008. Dogs were divided into three age-groups, as in a previous report:¹⁷ young (1–3 years old; n = 18); middle-aged (4–7 years old; n = 13); and older dogs (8–12 years old; n = 13). A body-condition score (BCS) was assessed and expressed on a five-point scale: 1, thin; 2, lean; 3, optimal; 4, obese; 5, gross, as described by Laflamme.¹⁸ This study followed the Guidelines for Institutional Laboratory Animal Care and Use of the School of Veterinary Medicine of Kitasato University.

Biochemical and hormonal analysis of blood

Peripheral blood was collected from a cervical vein into sodium fluoride-coated tubes without an anticoagulant. To obtain serum, a sample was centrifuged at 1500 g for 10 minutes. Biochemical analyses of these samples used an AU400 Clinical Chemistry Analyzer (Olympus, Tokyo, Japan).

Classification of canine peripheral blood mononuclear cells

Peripheral blood from a cervical vein was collected into EDTA tubes. PBMCs were isolated by gradient centrifugation at 750 g for 60 minutes using Lymphoprep (Axis-Shield, Oslo, Norway; specific gravity = 1.077 \pm 0.001). Cells were washed twice with phosphate-buffered saline (PBS). Lymphocytes were labeled with CD molecules by incubating cells with anti-mouse immunoglobulin (Ig)G antibodies (CD4, CD8, CD21, MHC, CD45R, and CD14; Serotec, Kidlington, UK;

VMRD, Pullman, WA; Beckman Coulter, Fullerton, CA) for 60 minutes at 4°C. The cells were then washed and incubated with anti-mouse IgG antibodies labeled with phycoerythrin or fluorescein isothiocyanate for 30 minutes at 4°C. Cells were washed with PBS, and CD antigen expressions were analyzed using a FACScan flow cytometer (Becton Dickinson, Mountain View, CA). The CellQuest software package (BD, Franklin Lakes, NJ) was used for data acquisition and analysis. Absolute counts for lymphocyte subsets were determined by: (lymphocyte absolute counts \times percentage of fluorescent positive cells within lymphogate)/100.

Primary culture of PBMCs

Peripheral blood from a cervical vein was collected into heparin tubes, and PBMCs were prepared using Lymphoprep. These cells were cultured at a density of 5×10^6 cells/mL in Roswell Park Memorial Institute 1640 culture medium supplemented with 10% fetal calf serum at 37°C in 5% CO₂ for 12 hours with or without stimulation of PBMC. After culture, canine cells were harvested by centrifugation and total RNA was extracted using TRIzol (Invitrogen, Carlsbad, CA).

RNA purification

RNA purification, reverse transcription, and mRNA expression were examined according to the manufacturer's instructions. Briefly, after the addition of 0.2 mL of chloroform and thorough mixing, an RNA solution in TRIzol was incubated on ice for 5 minutes. Then, the sample was centrifuged at 7500 g for 15 minutes at 4°C. After the addition of 0.2 mL of isopropanol to the separated aqueous layer, the sample was incubated on ice for 15 minutes. After centrifugation at 7500 g for 15 minutes at 4°C, the supernatant was removed, 70% ethanol was added, and it was dried. Finally, total RNA was dissolved in 20 μ L of nuclease-free water. RNA concentration was determined with a Nano Drop ND-1000 Spectrophotometer (LMS, Tokyo, Japan). cDNA synthesis used 1 μ g of RNA, adjusted to 10 μ L with nuclease-free water, which was reverse-transcribed with oligo dT primers. After incubation at 68°C for 5 minutes, the sample was added to a mixed liquor containing 5 \times first-strand buffer (4 μ L), 10 mM dNTP mix (1 μ L), 25 mM MgCl₂ solution (2 μ L), 0.1 M DTT (2 μ L), and Superscript II RNaseH Reverse Transcriptase (0.7 μ L) (Invitrogen, Carlsbad, CA). This sample was first incubated at 42°C for 50 minutes and then at 68°C for 15 minutes.

Real-time PCR analysis

Quantification of mRNA used real-time polymerase chain reaction (PCR) analysis. The sequences for canine IL-2,

IFN- γ , IL-4, IL-10, IL-2 receptors (IL-2R α and IL-2R γ), and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNA were obtained from GenBank. Our designed primers were; IFN- γ : 5'-CCAGATGTATCGGAGCGGTGGTTATCGCCTTGCCTGGACC-3', IL-2: 5'-GCATCGCACTGACGCTTGTA TTGCTCCATCTGTTGCTCTGTT-3', IL-2R γ 5'-CAAGAGCATCTGCAAAACCA GCTTGCTTGAAGCTCTTCGT-3', IL-2R α : 5'-CCCCATGTTACACCCTAAAGCCTGAA AGGTTTCCGGGCCCTCACATTG-3', IL-4: 5'-ATGGGTCTCACCTCCAACCTG TCAATGCCTGTAGTATTTCTTC-3', IL-10: 5'-TACCTGGTTGCCAAGCCCT TTCACAGA-GAAGCTCAGTAAAT-3', GAPDH: 5'-GGGGCCATCACAGTCTTCT GCCAAAAGGGTCATCATCTC-3'. Mixed solutions of primers and SYBR Green PCR Master Mix (PE Applied Biosystems, Foster City, CA) used a 7700 Sequence Detector (PE Applied Biosystems) with denaturing at 50°C for 30 minutes and 95°C for 10 minutes, followed by 45 cycles at 95°C for 15 seconds and 60°C for 1 minute. The target gene expression was normalized against endogenous GAPDH giving a Δ CT value and calibrated with the Δ CT value from the sample with lowest expression giving a $\Delta\Delta$ CT. Relative expression values were transformed from logarithmic to linear form putting $\Delta\Delta$ CT into the equation $2^{-\Delta\Delta$ CT.

Statistical analysis

Results are given as mean \pm standard error. Relationships between dogs' age and peripheral blood leukocyte subpopulations were assessed using Spearman's rank-order correlation coefficients. Statistical analysis using the group comparisons was done by Student's *t*-test or one-way analysis of variance using GraphPad Prism software (GraphPad, San Diego, CA). *P*-values of <0.05 were considered significant.

Results

For this study, blood samples were obtained from healthy beagles of various ages. These were divided into three groups based on age: young (1–3 years old; $n = 18$); middle-aged (4–7 years old; $n = 13$); and older dogs (8–12 years old; $n = 13$). Their BCSs ranged from 2.5 to 4.0. There was no significant correlation between age and BCS (Table 1). There were no significant differences in white blood cell and red blood cell counts, except for mean corpuscular volume: middle-aged dogs had greater mean corpuscular volume values than dogs in the other age-groups ($P < 0.05$; Table 1).

Table 1 Clinical characteristics of dogs

Variable	Young (18)	Middle-aged (13)	Aged (13)
BCS	3.13 \pm 0.09	2.96 \pm 0.09	3.17 \pm 0.17
Age (years)	1.72 \pm 0.14 ^a	5.02 \pm 0.23 ^b	10.24 \pm 0.52 ^c
WBC ($\times 10^3$ /mL)	139.39 \pm 10.24	129.69 \pm 12.16	113.23 \pm 7.94
RBC ($\times 10^6$ /mL)	690.00 \pm 30.20	705.15 \pm 34.71	637.54 \pm 12.66
Hb (g/dL)	15.94 \pm 0.70	16.61 \pm 0.84	14.92 \pm 0.31
Ht (%)	45.17 \pm 1.97	47.82 \pm 2.35	42.05 \pm 0.76
MCV (fl)	65.56 \pm 0.63 ^a	67.84 \pm 0.62 ^b	66.02 \pm 0.58 ^a
MCH (pg)	23.12 \pm 0.25	23.55 \pm 0.29	23.42 \pm 0.30
MCHC (g/dL)	35.28 \pm 0.17	34.70 \pm 0.19	35.48 \pm 0.25
PLT ($\times 10^9$ /mL)	21.22 \pm 2.45	16.69 \pm 2.42	19.08 \pm 1.92

Notes: Values are represented as mean \pm standard deviation; different letters, a, b, c indicate statistically significant differences between groups ($P < 0.05$).

To determine if there were alterations with age in peripheral blood leukocyte subpopulations in dogs, lymphocyte subsets were evaluated using a flow cytometric immunophenotyping method. Representative subpopulations are shown in the scatter diagrams (Figure 1). There were significant negative correlations between dogs' ages and the number of PBMCs, CD3⁺ T cells, CD4⁺ T cells, CD8⁺ T cells, and CD21⁺ B cells. Notably, the number of CD4⁺ CD45R⁺ T cells (naive helper T cells) and CD8⁺ CD45R⁺ T cells (naive killer T cells) decreased significantly with age. The numbers of CD45R⁻ T cells (memory T cells) did not change with age for either CD4⁺ or CD8⁺ T cells.

The mRNA levels for various cytokines in cultured PBMCs were analyzed using quantitative real-time PCR (Figure 2). Leukocytes were cultured for 12 hours under optimized conditions. mRNA levels for IL-2 and IL-2R α were significantly increased only in the young and middle-aged groups, respectively. IL-2R γ mRNA expression tended to decline with age. By comparison, IL-4 and IL-10 mRNA expressions did not change with age.

Discussion

In the present study, no changes were found in the clinical conditions and the biochemical examinations of healthy beagles in different age-groups. However, for peripheral blood leukocyte subpopulations, there were significant negative correlations between the numbers of PBMCs, CD4⁺ T cells, CD8⁺ T cells, CD4⁺ CD45R⁺ T cells (possibly naive cells), CD8⁺ CD45R⁺ T cells, CD21⁺ B cells and dogs' ages. In contrast, there were no significant correlations between the numbers of CD4⁺ CD45R⁻ T cells (possibly memory cells) and CD8⁺ CD45R⁻ T cells and dogs' ages. Thus, the subpopulations that were reduced in older healthy dogs were characteristically naive T cells. These results are in agreement

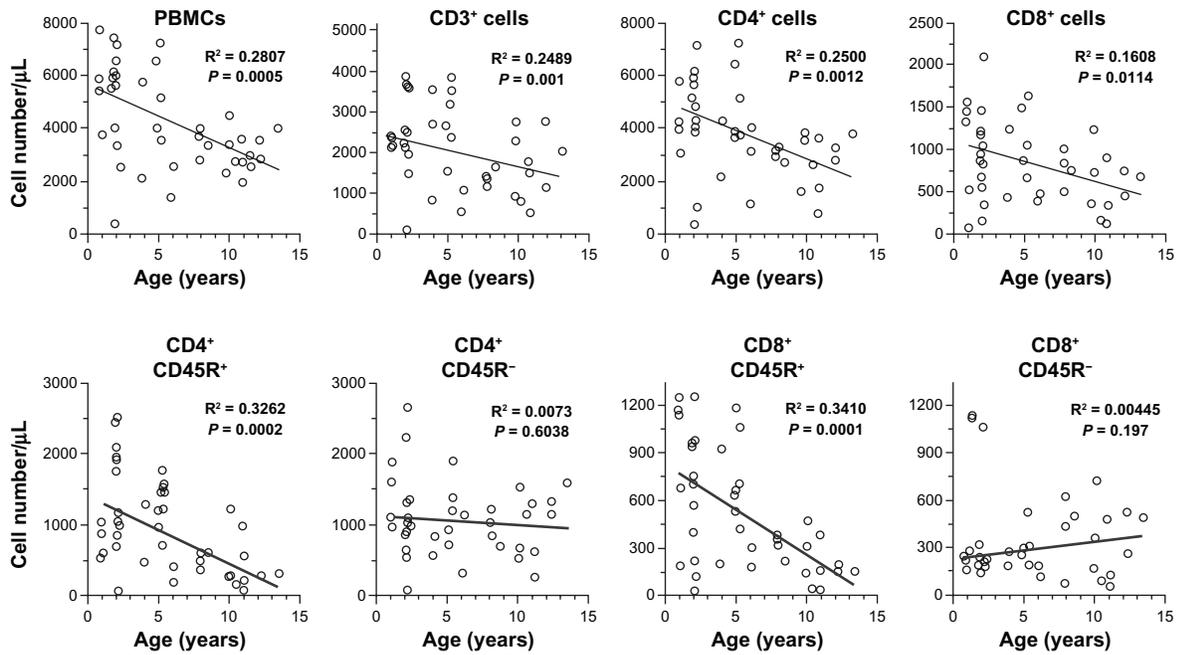


Figure 1 Scatter diagrams for age-related changes in peripheral blood leukocyte subpopulation in beagles.

Notes: Numbers of peripheral blood mononuclear cells (PBMCs), CD3⁺ T cells, CD4⁺ T cells, CD8⁺ T cells and CD45R^{+/−} cells, CD4⁺ or CD8⁺ T cells versus age in individual dogs are shown by open circles (n = 44). All of these variables, except for CD4⁺ CD45R[−] T cells and CD8⁺ CD45[−] T cells, were significantly negatively correlated with dogs' age ($P < 0.05$).

with human studies.^{19,20} Thus, the decrease in naive T cells may contribute to reduced T-cell responses to newly encountered antigens in aged dogs, as is observed in humans.²¹

In this study, the IL-2 and IFN- γ mRNA levels in Con A-stimulated PBMCs were significantly increased in cells from young and middle-aged dogs, respectively. In rats,

naive T cells are the best IL-2 producers.²² IFN- γ is also primarily produced by naive but not memory T cells.²³ Therefore, the age-related reductions of IL-2 and IFN- γ mRNA expressions may be due to the reduced numbers of naive T cells in older dogs. In contrast, IL-4 and IL-10 mRNA expressions appeared not to change with age. Memory T

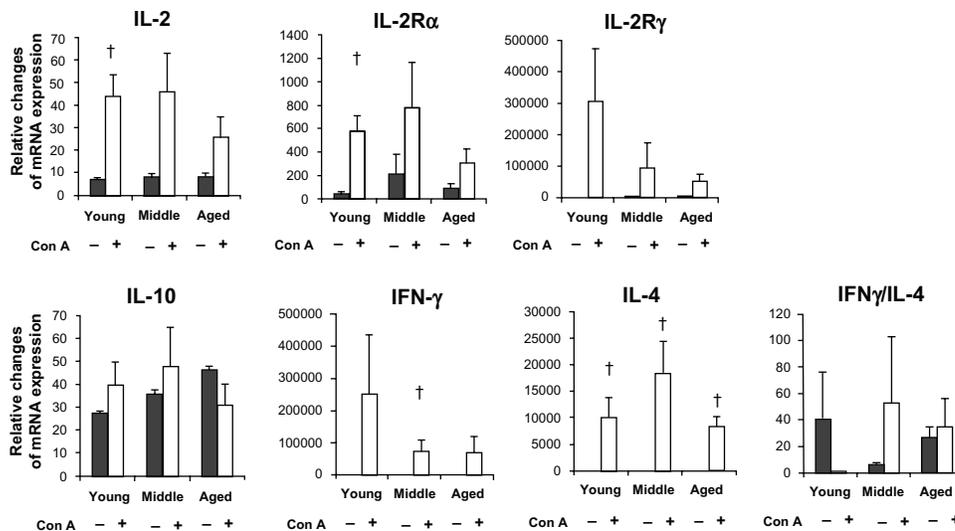


Figure 2 Age-related changes in the transcriptional (mRNA) levels for IL-2, IL-2R α , IL-2R γ , IL-10, IFN- γ , IL-4, and IFN γ /IL-4. mRNA levels were determined 12 hours after stimulation of PBMCs with Con A.

Notes: The control mRNA levels (gray bars) and mRNA levels induced by con-A stimulation (open bars) in PBMCs are shown in each age-group. IL-2, IL-2R α , and IFN γ mRNA levels were significantly increased only in young or middle-aged dogs ($P < 0.05$). IL-4 mRNA expression was significantly increased in all age-groups after Con-A stimulation ($P < 0.05$).

Abbreviations: IL, interleukin; PBMCs, peripheral blood mononuclear cells; IFN, interferon.

cells can produce significant amounts of IL-4.²³ Thus, it is possible that memory T cells, which did not change with age, may be primarily related to age-independent IL-4 mRNA expression.

Age-associated alterations in IL-2R mRNA expression will require further investigation. In this study, IL-2R γ mRNA expression tended to be downregulated with dogs' age. IL-2R γ -related cytokines appear to be essential for naive cell survival through the maintenance of antiapoptotic pathways.^{24,25} These changes in IL-2R γ mRNA patterns may also contribute to the decreases in naive T-cell functions observed with aging. In addition, IL-2R α mRNA expression decreased in an age-dependent manner. IL-2R α can effectively increase the formation of ternary complexes.²⁶ Because IL-2R α transmits much stronger proliferative and differentiation signals, age-related decreases in these IL-2R mRNA expressions may play an essential role in the decline of naive T cells.

The decline in immune function in older animals may be related to an increased incidence of certain diseases. However, a direct association between a decline in immune function and an increased susceptibility to diseases has not been reported. From the results of this study, the age-related decreases in naive T cells and IL-2R mRNA expression may contribute to the decline in immune function. Further studies will be necessary to determine whether these results are directly linked to clinical problems.

Conclusion

In summary, we analyzed the relationship between peripheral blood leukocyte subpopulations and their cytokine synthesis and age in healthy dogs. First, the numbers of naive T cells decreased with age, which may contribute to the reduced cytokine production by these cells. Second, the IL-2R α and IL-2R γ mRNA expression levels in PBMCs decreased with age. These reduced mRNA expressions may contribute to the decline in naive T cells. Reduced numbers of naive T cells and IL-2R mRNA expression could affect each other and be related to immunosenescence associated with aging.

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Disclosure

This work was supported by the Japan Pet Nutrition Society. The authors declare no conflicts of interest in this work.

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