

Peripheral Lymphocyte Subsets as a Prognostic Indicator of Mortality and Morbidity in Healthy Dogs

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ABSTRACT. To evaluate the relationship among immune status and increased morbidity and mortality, peripheral blood lymphocytes (CD3⁺, CD4⁺, CD8⁺ and CD21⁺ cells) from 32 healthy dogs over 8 years of age were analyzed. Twenty-five of the 32 dogs were followed-up for 3 years after the analysis; and 14 dogs were found to be diseased, and nine dogs died. There was no notable difference between the ages of the dogs that died compared with the ones that survived. The relative percentage of CD4⁺ and the CD4⁺:CD8⁺ ratio decreased notably in dogs falling ill compared with healthy dogs. The relative percentage of CD3⁺ lymphocytes showed a notable decrease in dogs that died within 3 years in comparison with dogs that survived. In a discriminant analysis of morbidity and mortality, most patients were correctly classified as diseased or not and surviving or dead, respectively. These results indicate that the immunophenotypes of peripheral blood lymphocytes in older dogs offer promise as parameters for evaluating mortality and morbidity.

KEY WORDS: aging, canine, lymphocyte subset, morbidity, mortality.

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Changes in immune status are considered major contributing factors towards morbidity and mortality in aging humans. Many studies performed in human subjects and in experimental animals have suggested a correlation between immune function and age-related risk of morbidity and mortality. Immunosenescence has been associated with a numerical change in T-cell number [5, 22]. Previous studies suggested that a lower total lymphocyte count was associated with an increased risk of mortality [2, 7, 12]. Similarly, an association of high numbers of CD8⁺ and low numbers of CD4⁺ cells or low CD4⁺:CD8⁺ cell ratios with increased mortality has also been demonstrated [1, 3-5, 13, 19, 22]. Additionally, other evidence has been reported on the association of higher mortality with decreased lymphocyte proliferation to mitogens [14, 21].

Such immunological changes observed in aging humans have also been studied in dogs [9-11, 17, 18]. Our previous study demonstrated age-related changes in lymphocyte subsets (CD3, CD4, CD8, and CD21) in healthy dogs and a notable decrease in the absolute number and relative percentages of lymphocyte subsets in dogs with tumors compared with healthy controls. Furthermore, the relative percentages of particular lymphocyte subsets were significantly decreased in dogs with distant metastatic tumor in

comparison with dogs without metastases [20].

Despite extensive research in humans, few studies evaluating the relationship among age-related change in immune status and higher morbidity and mortality have been performed in dogs. In this study, we analyzed whether the immune status in clinically healthy dogs over 8 years of age that tumor incidence becomes higher [20] could predict future morbidity and mortality.

Peripheral blood was obtained from 32 (11 males and 21 females) dogs over 8 years of age that were evaluated as healthy by physical examination and blood tests at private veterinary hospitals. The blood drawing was conducted between May and June in 2007. None of the animals had chronic diseases or had received any immunosuppressive treatment, including corticosteroids. All blood samples were drawn into ethylenediaminetetraacetic acid (EDTA) tubes and complete blood count (CBC) tests were performed immediately. Retrospective analysis of mortality and morbidity was performed in June 2010, 3 years after the blood sample collection.

Two-color flow cytometry was used to calculate relative percentages of CD3⁺ (T cells), CD21⁺-like (B cells), CD4⁺ (T-helper cells) and CD8⁺ (T-cytotoxic cells) cells using canine-specific monoclonal antibodies (mAbs). Details for all mAbs used for flow cytometry assays are listed in Table 1. Each blood sample was prepared with the protocol that was described in our previous study [20]. Samples were analyzed using a Coulter Epics XL flow cytometer (Beckman-Coulter, Marseille, France) and the Expo32 ADC software (ver. 1.1c; Beckman-Coulter, Marseille, France). Absolute values for each subset were calculated using counts ob-

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Table 1. (A) Monoclonal antibodies used for flow cytometry and (B) staining protocol used

(A) Phenotype	Specificity	Clone	Conjugate	Manufacturer
CD3	Pan T cells	CA17.2A12	FITC	Serotec, Oxford, UK
CD4	Helper T cells	YKIX302.9	FITC	Serotec, Oxford, UK
CD8a	Cytotoxic T cells	YCATE55.9	RPE	Serotec, Oxford, UK
CD21-like	Pan B cells	CA2.1D6	RPE	Serotec, Oxford, UK

(B) Tube number	FL-1	FL-2
1	CD4 FITC	CD8a RPE
2	CD3 FITC	CD21 RPE

tained from white blood cell (WBC) analysis in combination with flow cytometry data. WBC count was analyzed using a veterinary automatic blood cell calculator (MICROS abc LC-152, Horiba, Tokyo, Japan).

The Kruskal-Wallis rank test with a Dunn's post test and discriminant analysis were used to identify lymphocyte subset patterns that differentiated healthy dogs from dogs that died or became diseased during the follow-up period of 3 years. Analyses were carried out using StatMate III for Windows (Atomusu, Tokyo, Japan). Differences were considered significant where $P < 0.05$. All values are presented as means \pm SD, medians, or ranges.

Sixteen dogs were alive at the 3-year follow-up; nine of the 25 dogs had died during the 3-year period. Seven were lost to follow-up. Among the dogs that died during the follow-up period, one dog died in the 1st year after blood sample collection, 2 dogs died in the second year, and 6 dogs died in the third year. The causes of death included cancer ($n=5$), infectious diseases ($n=1$), cardiovascular diseases ($n=1$), and unknown causes ($n=2$). There was no notable difference among the ages of the dogs that died and those that did not. Thirteen of the 25 dogs developed 1 or some diseases during the period after examination. These diseases included cancer ($n=5$), immune diseases ($n=3$), neuropathic diseases ($n=1$), cardiovascular diseases ($n=2$), disc hernia ($n=1$), diabetes mellitus ($n=1$), hepatic disease ($n=1$) and pyometora ($n=1$).

To investigate whether lymphocyte subsets correlated with future mortality, dogs were categorized into dogs that survived and those that died (Table 2). Of all the lymphocyte subsets analyzed in the present study, only the relative percentage of CD3⁺ lymphocytes showed any notable decrease in dogs that died in comparison with dogs that survived ($P < 0.05$) (Table 3). The relative percentages of CD4⁺, CD8⁺ and CD21⁺ lymphocytes and the absolute cell counts of CD21⁺ lymphocytes were decreased in dead dogs compared with the surviving dogs, but there were no statistical differences.

To investigate whether lymphocyte subsets correlated with future morbidity, dogs were categorized into dogs that remained healthy and those that were diseased (Table 2), and the lymphocyte subsets were compared (Table 4). The relative percentage of CD4⁺ lymphocytes demonstrated a significant decrease in dogs with onset of a disease compared with dogs without disease ($P < 0.01$). The relative percentages of

CD3⁺ and CD21⁺ lymphocytes were decreased in diseased dogs compared with healthy dogs, but there were no statistical differences. On the other hand, the relative percentage of CD8⁺ lymphocytes was increased in diseased dogs, but there was no statistical difference. The CD4⁺:CD8⁺ ratio notably decreased in dogs with onset of a disease ($P < 0.05$).

White blood cell count, absolute cell counts and relative percentages of all lymphocyte subsets (total lymphocyte, CD3⁺, CD4⁺, CD8⁺ and CD21⁺), CD4⁺:CD8⁺ ratio and CD3⁺:CD21⁺ ratio were included in a discriminant analysis to identify the relationship between lymphocyte subsets and increased mortality and morbidity. In a discriminant analysis of mortality, 92.0% of all patients were correctly classified as dead or surviving, as predicted by their immunophenotype. The linear discriminant function was $y = -100.6 + 2.4 \times [\text{lymphocytes (\%)}] + 0.35 \times [\text{CD3}^+ \text{ T cells (\%)}] + 0.37 \times [\text{CD4}^+ \text{ T cells (\%)}] + 0.74 \times [\text{CD8}^+ \text{ T cells (\%)}] - 3.0 \times [\text{CD21}^+ \text{ B cells (\%)}] + 0.0050 \times [\text{WBC (cells/\mu l)}] + 0.019 \times [\text{lymphocytes (cells/\mu l)}] - 0.040 \times [\text{CD3}^+ \text{ T cells (cells/\mu l)}] - 0.060 \times [\text{CD4}^+ \text{ T cells (cells/\mu l)}] - 0.039 \times [\text{CD8}^+ \text{ T cells (cells/\mu l)}] + 0.18 \times [\text{CD21}^+ \text{ B cells (cells/\mu l)}] + 11.1 \times [\text{CD4}^+:\text{CD8}^+] + 0.51 \times [\text{CD3}^+:\text{CD21}^+]$. This linear discriminate function estimated dogs to be surviving if $y < 0$ (dependent variable: dead=16; surviving=9). In a discriminant analysis of morbidity, 100% of the patients were correctly classified as diseased or healthy. Linear discriminant function was $y = -16.7 + 0.42 \times [\text{lymphocytes (\%)}] - 0.75 \times [\text{CD3}^+ \text{ T cells (\%)}] + 2.0 \times [\text{CD4}^+ \text{ T cells (\%)}] + 0.37 \times [\text{CD8}^+ \text{ T cells (\%)}] - 2.9 \times [\text{CD21}^+ \text{ B cells (\%)}] + 0.0027 \times [\text{WBC (cells/\mu l)}] + 0.018 \times [\text{lymphocytes (cells/\mu l)}] + 0.045 \times [\text{CD3}^+ \text{ T cells (cells/\mu l)}] - 0.14 \times [\text{CD4}^+ \text{ T cells (cells/\mu l)}] - 0.061 \times [\text{CD8}^+ \text{ T cells (cells/\mu l)}] + 0.15 \times [\text{CD21}^+ \text{ B cells (cells/\mu l)}] - 2.2 \times [\text{CD4}^+:\text{CD8}^+] - 0.15 \times [\text{CD3}^+:\text{CD21}^+]$. This linear discriminated function estimated dogs to be healthy if $y < 0$ (dependent variable: diseased=10; healthy=13). The largest coefficient of linear discriminant function to classify dogs as surviving or dead and healthy or diseased was for the CD4⁺:CD8⁺ ratio.

The data presented in the current study suggested that the immunophenotype of peripheral blood lymphocytes in older dogs can be a useful predictor for evaluating immunosenescence, mortality and morbidity in dogs. Age-related immunological changes are mainly represented by alteration in T-cell phenotype and function [15]. Immunosenescence is characterized by alterations in T cell subsets characterized by

Table 3. Differences in immunophenotypes of peripheral blood lymphocytes from dogs that died compared with age-matched controls

(%)	Lymphocytes		Phenotype			
			CD3 ⁺	CD4 ⁺	CD8 ⁺	CD21 ⁺
Survived (n=16)						
Median		20.3	70.1	34.3	30.0	12.0
Range		6.4–27.6	49.8–84.7	14.5–53.7	11.6–51.6	7.3–31.1
Dead (n=9)						
Median		24.4	59.0 ^{a)}	27.5	25.1	8.1
Range		14.1–30.7	32.4–83.5	12.6–44.2	11.7–41.4	1.7–36.2
(cells/ μ l)	WBCs	Lymphocytes	Phenotype			
			CD3 ⁺	CD4 ⁺	CD8 ⁺	CD21 ⁺
Survived (n=16)						
Median	8500	1621	1115	512	510	199
Range	3200–14700	511–2426	382–2038	181–1457	120–1189	45–468
Dead (n=9)						
Median	8400	1772	1069	537	453	154
Range	6100–9200	1182–2428	575–1476	172–863	209–971	21–692
			CD4 ⁺ :CD8 ⁺		CD3 ⁺ :CD21 ⁺	
Survived (n=16)						
Median			1.1		6.2	
Range			0.4–3.5		2.3–8.8	
Dead (n=9)						
Median			1.1		6.9	
Range			0.3–3.8		1.0–41.2	

a) Significant difference vs. survival group.

Table 4. Differences in immunophenotypes of peripheral blood lymphocytes from dogs with onset of a disease compared with age-matched controls

(%)	Lymphocytes		Phenotype			
			CD3 ⁺	CD4 ⁺	CD8 ⁺	CD21 ⁺
Healthy (n=10)						
Median		20.6	70.5	35.8	27.2	14.4
Range		6.4–27.6	62.3–84.7	25.7–53.7	11.6–39.2	7.3–31.1
Diseased (n=13)						
Median		22.7	61.2	25.2 ^{a)}	29.2	8.2
Range		11.9–25.5	32.4–83.5	12.6–48.4	13.3–51.6	1.7–36.2
(cells/ μ l)	WBCs	Lymphocytes	Phenotype			
			CD3 ⁺	CD4 ⁺	CD8 ⁺	CD21 ⁺
Healthy (n=10)						
Median	8100	1616	1082	523	517	259
Range	3200–14700	511–2426	382–2038	181–1457	120–668	45–468
Diseased (n=13)						
Median	8500	1655	1069	452	500	154
Range	6100–12700	1182–2251	575–1487	172–758	209–1189	21–692
			CD4 ⁺ :CD8 ⁺		CD3 ⁺ :CD21 ⁺	
Healthy (n=10)						
Median			1.2		5.1	
Range			0.7–3.5		2.3–8.8	
Diseased (n=13)						
Median			1.0 ^{a)}		7.0	
Range			0.3–2.1		1.0–41.2	

a) Significant difference vs. dogs that had no diseases.

Table 2. Immunophenotypes of peripheral blood mononuclear cells and outcomes in >8-year-old healthy dogs

No.	Age (years)		Sex	Breed	Survival	Cause of death	Disease	WBCs	Lymphocytes	Phenotype					
	Collection	Death								CD3 ⁺	CD4 ⁺	CD8 ⁺	CD21 ⁺	CD4 ⁺ :CD8 ⁺	CD3 ⁺ :CD21 ⁺
1	8	-	F	Great pyrenees	Surviving	-	No	(%) (cells/ μ l)	13.3 1048	68 698	41.1 425	11.6 120	25.7 264	3.5	2.6
2	8	-	M	Mongrel	Surviving	-	No	(%) (cells/ μ l)	6.4 936	77.7 708	53.7 545	21.5 218	31.1 283	2.5	2.5
3	8	-	M	Shih tzu	Surviving	-	No	(%) (cells/ μ l)	19.0 1574	66 1057	25.7 403	38.8 609	29.2 468	0.7	2.3
4	9	-	F	Shih tzu	Surviving	-	No	(%) (cells/ μ l)	27.6 2426	77.7 2038	53.4 1457	17.5 477	17.2 451	3.1	4.5
5	10	-	F	Pug	Surviving	-	No	(%) (cells/ μ l)	6.9 680	62.3 382	27.8 223	25.6 205	7.3 45	1.1	8.5
6	10	-	F	Shih tzu	Surviving	-	No	(%) (cells/ μ l)	27.3 1695	68.9 1123	33.3 624	35.7 668	7.8 127	0.9	8.8
7	10	-	F	Shih tzu	Surviving	-	No	(%) (cells/ μ l)	24.9 2238	72.1 1596	43.9 1011	24.9 574	13.5 299	1.8	5.3
8	11	-	F	Chihuahua	Surviving	-	No	(%) (cells/ μ l)	26.7 1657	74.1 1236	30.1 502	39.2 654	15.2 254	0.8	4.9
9	13	-	M	Shetland sheepdog	Surviving	-	No	(%) (cells/ μ l)	22.2 1665	66.5 1107	36.3 705	28.7 557	9.3 155	1.3	7.2
10	14	-	F	Maltese	Surviving	-	No	(%) (cells/ μ l)	16.0 511	84.7 447	35.3 181	31.3 160	9.9 52	1.1	8.6
11	9	-	M	Maltese	Surviving	-	Cardiac disease	(%) (cells/ μ l)	17.2 1548	77.7 1196	48.4 758	24 376	12.6 194	2.0	6.2
12	11	-	M	Beagle	Surviving	-	Disc herniation	(%) (cells/ μ l)	25.4 1750	79.6 1423	35.7 618	34.1 591	11.4 204	1.0	7.0
13	8	-	M	Yorkshire terrier	Surviving	-	Immune disease	(%) (cells/ μ l)	21.7 2213	67.5 1487	19.6 452	51.6 1189	8.8 194	0.4	7.7

Table 2. Continuu

No.	Age (years)		Sex	Breed	Survival	Cause of death	Disease	WBCs	Lymphocytes	Phenotype						
	Collection	Death								CD3 ⁺	CD4 ⁺	CD8 ⁺	CD21 ⁺	CD4 ⁺ :CD8 ⁺	CD3 ⁺ :CD21 ⁺	
14	8	-	F	Shih tzu	Surviving	-	Immune disease (cells/ μ l)	12700	11.9	61.2	25.2	31.5	24.7			
15	14	-	F	Mongrel	Surviving	-	Immune disease (cells/ μ l)	8700	1516	894	400	500	361	0.8	2.5	
16	13	-	M	Beagle	Surviving	-	Nervous disease (cells/ μ l)	6500	1586	1158	522	521	133	1.0	8.7	
17	8	10	F	Pomeranian	Dead	unknown	No	8700	20.4	70	44.2	11.7	13.4	0.5	6.1	
18	11	14	F	Welsh corgi	Dead	unknown	No	7900	1772	1285	800	212	246	3.8	5.2	
19	11	14	F	Shih tzu	Dead	Heart failure	Diabetes		30.7	55.6	36.8	41.4	8.8	0.9	6.3	
20	8	11	F	Great pyrenees	Dead	Infection	Pyometra		22.7	59	22.3	35.6	36.2	0.6	1.6	
21	9	11	F	Golden retriever	Dead	Tumor	Tumor		1932	1128	432	690	692	2.1	13.7	
22	11	13	M	Mongrel	Dead	Tumor	Tumor		24.5	46.7	28.1	13.7	3.4	1.1	1.0	
23	11	11	M	Irish setter	Dead	Tumor	Tumor		2251	1061	664	324	77	0.3	7.3	
24	11	14	F	Miniature schnauzer	Dead	Tumor	Tumor		24.9	32.4	30.9	29.2	30.9	1.0	41.2	
25	12	13	F	Golden retriever	Dead	Tumor	Tumor		1716	575	539	510	548	1.1	10.3	
									15.1	59.1	12.6	40.3	8.1	0.3	7.3	
									1347	773	172	549	106			
									14.1	70.1	24.1	25.1	1.7			
									1182	871	281	293	21			
									24.7	83.5	14.5	13.3	8.1			
									1509	1177	228	209	114			
									24.4	53.5	27.5	23.2	7.7			
									1852	1069	537	453	154			

an increase in CD8⁺ lymphocytes and CD8⁺CD28⁻ lymphocytes [5, 8] and a decrease in CD4⁺:CD8⁺ lymphocyte ratio in humans [5]. The increase in CD8⁺ lymphocyte population and an associated decrease in the CD4⁺:CD8⁺ ratio appear to be characteristic of an ageing immune system. This typical effect of aging has been observed in our previous study and other canine studies [6, 10, 11, 20]. In addition, a lower total count of lymphocytes was associated with an increased risk of mortality in older humans [2, 12]. The association of high CD8⁺ and low CD4⁺ T cells, or a low CD4⁺:CD8⁺ lymphocyte ratio, with increased mortality was reported in some studies [1–5, 7, 12, 13, 19, 22]. Other studies have reported that the absolute number of CD8⁺ memory T cells and decreased lymphocyte proliferation to mitogens correlated with increased mortality [14, 16, 21]. Our study demonstrated that the CD4⁺:CD8⁺ ratio significantly decreased in older dogs falling ill compared with dogs that remained healthy within the 3-year follow-up period. These data suggest a relationship between a low CD4⁺:CD8⁺ lymphocyte ratio and higher morbidity in older dogs.

In contrast, an association between T-cell subsets and higher mortality was not shown; however, this study may have the following major potential limitations. 1) The number of enrolled dogs was too small to clarify the statistical differences. 2) The cause of mortality was not selected. T-cell subset alterations and immunosenescence should be more relevant for certain conditions, such as mortality from sepsis, meningitis, other infectious diseases or tumors. 3) A follow-up period of 3 years was used to investigate the morbidity and mortality of older dogs, but there are no data as to whether a 3-year follow-up period is suitable to predict mortality and mobility from the lymphocyte subsets of dogs. 4) The immune system comprises various functions and consists of many types of cells, and it is therefore difficult to estimate immune status by one immunological parameter suitable for the assessment of immune functions in healthy dogs and patients suffering from various diseases.

The last limitation, however, may be resolved by the discriminant analysis performed in the current study. The discriminant analysis based upon the WBC count, absolute cell counts, relative percentages of all lymphocyte subsets (total lymphocytes, CD3⁺, CD4⁺, CD8⁺, and CD21⁺), CD4⁺:CD8⁺ ratio and CD3⁺:CD21⁺ ratio identified a relationship between lymphocyte subsets and mortality or morbidity in older dogs. In a discriminant analysis of mortality, 92.0% of the patients were correctly classified by their immunophenotype as survived or dead. Similarly, in a discriminant analysis of morbidity, 100% of the patients were correctly classified as diseased or healthy. However, due to the small population of dogs used in this study, the linear discriminant functions developed here are a temporary. More cases would likely increase the reliability of these discriminant functions, and whether the prognoses of dogs can be estimated should be examined in a prospective study.

The current study suggests that evaluating the peripheral immunophenotype in older dogs may provide a prognostic value of survival perhaps by detecting early indicators of disease not detected by a clinician or owner.

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