Published online 2021 March 28.

Research Article

Molecular and Immunologic Detection of Canine Distemper Virus (CDV) and Investigation of Co-infection with *Candida albicans* in Dogs

Fatemeh Ahmadi Sheikh Shabani 1, Azam Mokhtari 1, * and Somayeh Shahrokh Shahraki

¹Department of Pathobiology, Faculty of Veterinary Medicine, Shahrekord University, Shahrekord, Iran ²Research Institute of Zoonotic Diseases, Shahrekord University, Shahrekord, Iran

corresponding author: Department of Pathobiology, Faculty of Veterinary Medicine, Shahrekord University, Shahrekord, Iran. Tel: +98 -3832324401, Fax: +98-3832324427, Email: mokhtari_dvm@yahoo.com

Received 2021 February 01; Revised 2021 March 01; Accepted 2021 March 01.

Abstract

Background: Canine distemper (CDV) is a highly transmissible serious disease of carnivores. Distemper virus has immunosuppression effects which, in turn, causes opportunistic infections.

Objectives: The present study was performed to detect CDV by the genomic and immunological methods and investigate its coinfection with *Candida albicans*.

Methods: In this study, from spring of 2018 to winter of 2019, blood, eye, respiratory and digestive system samples were collected from 50 CDV suspected dogs and 50 seemingly healthy dogs. Rapid distemper immunochromatography kit was applied for the primary detection of CDV. RT-PCR and PCR tests were performed using special primers for molecular investigation.

Results: Using immunochromatography, twenty-nine cases and one case had positive results for CDV among dogs suspected of having the disease and seemingly healthy dogs, respectively. After RT-PCR and PCR assays, 37 samples were CDV-positive, and four were *C. albicans*-positive in the first group. While three CDV and one *C. albicans*-positive samples were found in the second group. In total, the frequency of co-infection was 4%.

Conclusions: In the present study, there was an association between distemper and *C. albicans* using statistical tests. Conducting such studies in an appropriate sample population gives more accurate results.

Keywords: Distemper, Candida albicans, Genomic Investigation, Immunological Detection, Dog

1. Background

Canine distemper virus (CDV) is a member of the family *Paramyxoviridae* (1). The virus is endemic in many areas such as India, Denmark, Finland, Brazil, and North America (2). There are some reports of CDV in dogs of Iran. A high prevalence (55.6%) of CDV has been reported among rural dogs in the north of Iran (3). In a study performed by Avizeh et al. (2007), the prevalence of CDV was 17.52% (4) Although the CD vaccine has significantly reduced the disease, CD is endemic in many parts of the world (5, 6).

The CDV inhibits immunity in vulnerable species, and its receptor is the signaling lymphocyte activation molecule 4 (SLAM 4) (5, 6). Recently, new reports have been published showing that CDV infection can facilitate the development of diseases caused by common or emerging viruses (7-11). Morbilliviruses are transmitted by aerosols and cause clinical signs such as respiratory and gastrointestinal symptoms, which are often complicated by the addition of other pathogens. In addition, one of the features

of morbillivirus infection is severe, temporary suppression of the host immune system (7).

During the first 24 hours after CDV infection, the virus accumulates in macrophages, B and T cells, and later, the virions go to the lymphoid tissues. Then, multiply in the reticuloendothelial organs (12, 13). The initial proliferation of the virus in lymphoid tissues leads to severe suppression of the immune system (14-16). After the viremia, CDV spreads to a number of epithelial tissues and the central nervous system (CNS) (13). Penetration of T lymphocytes and dendritic cells (DC) causes cutaneous manifestations of the disease (17). Lymphopenia occurs due to the reduction of the population of CD4+ T cells, CD8+ T cells, and CD21+ B cells, and CD4+ T cells are the main targets of CDV (18). CD4+ lymphocytes will be probably apoptotic and die (16), while CD8+ cells are more resistant (17).

The decrease in circulating immune system cells may be due to the reduction of cells in the lymphatic organs as well as the apoptosis of white blood cells. Programmed cell death can also happen in non-infected cells, indicat-

Copyright © 2021, Jentashapir Journal of Cellular and Molecular Biology. This is an open-access article distributed under the terms of the Creative Commons Attribution-NonCommercial 4.0 International License (http://creativecommons.org/licenses/by-nc/4.0/) which permits copy and redistribute the material just in noncommercial usages, provided the original work is properly cited.

ing an additional non-viral mechanism of apoptosis (19). *Candida albicans* (*C. albicans*) is one of the microflora of the body, which in some cases causes intense mucosal and skin infection, vaginal candidiasis, oral candidiasis, and other fungal diseases (20). People with immunosuppression are particularly vulnerable to infections like candidiasis (21). *Candida* species in growth medium are in the form of little yeast cells (3 - 6 microns), egg-shaped with thin shells, and Blastoconidium. In the growth time, buds do not separate from the mother cell and create a pseudohypha (18).

The attack of *C. albicans* on the skin happens through impact and creating laceration or burns, and when the skin moisture is high, the spots form. Onychomycosis is one of its other consequences (22). Thrush and oral candidiasis involve the oral cavity, tongue, gums, or oral palate, which may lead to bleeding (22). The attack of yeasts to vaginal mucus results in vulvovaginitis and causes itch, inflammation, and an increase in vaginal discharge (23).

2. Objectives

The aim of this study was the molecular and immunological investigation of CDV and its co-infection with *C. albicans.*

3. Methods

3.1. Sampling

In this cross-sectional study, the samples were taken from dogs in clinics of Isfahan (2018 - 2019). The age was determined based on the statements of the animal owner and the examination of the animal's teeth. For sampling, the animal was first physically restrained and examined for clinical signs. After the restrain of the animals [ketamine (Alfasan, Netherlands)-acepromazine (Hogestrati, Belgium)], respiratory and gastrointestinal tract swabs were taken from 50 symptomatic and 50 asymptomatic unvaccinated dogs for examination by a rapid immunochromatography detection kit of distemper and placed in sterile PBS. One of the limitations of the present study was the achievement of a large number of symptomatic CDV cases. For this reason, a statistically appropriate sample size was not obtained.

3.2. Rapid Distemper Immunochromatography

Rapid immunochromatography kit of distemper; VetALL SensPERT Canine Distemper Virus Test Kit, CDV Ag (Ngaio Diagnostics, New Zealand) was applied for detection of CDV in eye-respiratory swabs. The samples were firstly diluted in the kit buffer. Then 100 μ L of it was poured into the sample well, and from this moment, the time was taken. Then, the sample was dripped on a pad, and after the adsorption of dripped samples, two red bands showing the positive result appeared in the C (control) and T (test) positions.

3.3. Culture Test, Giemsa Staining

Culture tests were conducted on Sabouraud dextrose agar plate. After the smears were prepared on the slides, they were fixed with methanol and dried completely (30 seconds). Consequently, slides were immersed in a freshly prepared 5% Giemsa stain solution for 15 - 30 minutes. The slides were washed and dried with water and finally observed with a Bright Field microscope.

3.4. RNA/DNA Extraction and RT Reaction

QIAamp Viral RNA Mini Kit (Qiagen, Cat No: 52904) was used for RNA extraction from buffy coat. In order to determine the quality of the extracted RNA samples, the optical density (OD) of the samples was measured with a spectrophotometer at 260/280 nm. TaqMan RT Kit was used for cDNA synthesis (Invitrogen, USA) for applying in PCR reaction. The reaction was performed in a total volume of 25 μ L using recommended concentration of RT buffer, MgCl2, dNTPs, Random hexamers, RNase inhibitors, and reverse transcriptase (MMLV RT). the RNA template (400 - 500 ng/ μ L) and distilled sterile water up to 10 μ L. The thermal program included: 25°C for 10 min, 48°C for 30 min, and 95°C for 5 min. For investigation of *C. albicans*, DNA was extracted using the protocol suggested by Doyle et al. (1990) (cited in Davies et al.) (12).

3.5. Statistical Analysis

RT-PCR and PCR assays were performed with primers designed for detection of CDV by Beacon designer software. The sequences of specific primers of C. albicans species were extracted from Khan et al. (2001) (23). The sequences of primers were as follows: CDVF: AAGCCTCACACTGTTCAAG-3', CDVR: GATTAGGACTATAATGACATGC-3' and C.albF: ATGGGTGGTCAACATACTTC-3', C.albR: 5'-TACATCTATGTCTACCACC-3'. For detection of C. albicans, the PCR thermal cycle programs consisted of 94°C for 5 min, 30 cycles at 94°C for 45s, 60°C for 45 s and 72°C for 45 s, and 72°C for 5 min. The PCR cycling and temperature for investigation of CDV were the same as C. albicans. The only difference was that the annealing temperature was 58°C. The negative and positive controls were applied in each test. The PCR products electrophoresed in 1.5% agarose with 0.01% Green viewer (Parstous, Cat No.: B111151) along with a 50-base pair ladder (Fermentas, Cat No.: SM1133). Then two positive PCR products were sent for sequencing (Bioneer, Korea). Finally, the data were analyzed using Chi-square test by SPSS-16 software.

4. Results

There were 29 and one positive cases among symptomatic and asymptomatic dogs using a rapid distemper immunochromatography kit. The CDV and *C. albicans* positive samples and controls had the bands with the sizes of 100 bp and 496 bp, respectively (Figures 1 and 2).

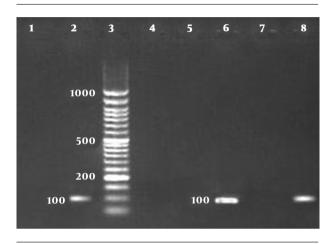


Figure 1. Electrophoresis gel of CDV RT-PCR products, 1) Negative control, 2and8) CDV-positive RT-PCR products (100 bp), 6) Positive control (100 bp), 3) Ladder (50 bp)

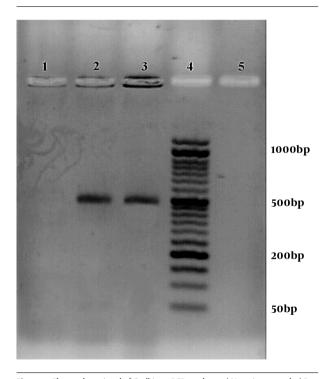


Figure 2. Electrophoresis gel of *C. albicans* PCR products; 1) Negative control, 2) Positive PCR product (496 bp), 3) *C. Albicans* (496 bp), 4) Ladder (50 bp)

The sequencing results of RT-PCR/PCR products showed that the degree of identity of the detected sequences with the sequences registered in the gene bank for distemper virus and *C. albicans* were high, and it was a good indicator of PCR efficiency (Figures 3 and 4).

On Sabouraud dextrose agar, the smooth creamy white colonies with a characteristic yeast odor were seen. In the Giemsa stained smears, budding yeast cells and pseudomycelium were typical (Figures 5 and 6).

In total, using RT-PCR/PCR test, among 50 symptomatic and 50 asymptomatic dogs, there were 37 and three CDVpositive results, respectively. The frequency of C. albicans samples in the first and second groups were four and one cases, respectively. The rate of CDV and C. albicans coinfection was 4%, only among dogs suspected of having distemper clinical signs. All of the 30 samples were positive using a rapid distemper immunochromatography test showed positive results by RT-PCR performed for detection of CDV. There was a statistically significant relationship between the results of these two tests (P < 0.05). Furthermore, there was a statistically significant association between CDV rapid test results and possess the symptoms of distemper. No statistically significant relationship was found between two diseases, C. albicans PCR results and gender; and CDV RT-PCR results and gender (P > 0.05). However, the relationship between C.albicans PCR tests and CDV RT-PCR results and CDV signs and age groups was statistically significant (P < 0.05) (Table 1).

5. Discussion

Almost all dogs with distemper experience immune suppression. As a result, other infections can be added to the disease and make these dogs sicker (3). Therefore, immune-suppressing mechanisms may affect the cells that are not directly involved with viral infection. Although the exact mechanism of immunosuppression due to distemper virus infection is not well understood, the weakening of the immune system due to this virus is accepted by all scientists. Therefore, and considering the possibility of further infections due to immunosuppression resulting from distemper virus, the current study was done to investigate co-infection of distemper and C. albicans as an opportunistic pathogen. However, the coinfection rate was 4% of cases, there was no statistically considerable relationship between distemper and C. albicans. In this study, the overall prevalence of C. albicans was 5%.

The limitation of the present study in accessing a higher number of symptomatic dogs and, consequently, low sample size may be one of the reasons for not achieving a considerable association between these two pathogens.

COVID-19 is an Emerging, Rapidly Evolving Situation. Get the Latest Public Health Information from CDC: <u>https://www.coronavinus.gov.</u> Get the Lates Research from NH: <u>https://www.nib.gov/cononavinus.</u> Find NCBI SARS-CoV-2 Literature, Sequence, and Clinical Content: <u>https://www.ncbi.nlm.nib.gov/sars-cov-2/</u>.

BLAST ® » Blastn Suite »Results for RID-JRMEK640016

Job Title	Cad F. ab1
RID	JRMEK640016 Search Expires on 08-07 17:48 pm
Program	BLASTN
Database	nt
Query ID	Icl Query_41277
Description	None
MoleculeType	dna
Query Length	100

Descriptions

Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
Canine morbillivirus isolate BJ16C9, complete genome	185	185	100%	2e-43	100.00%	MF926604.1
Canine morbillivirus isolate BJ16C8, complete genome	185	185	100%	2e-43	100.00%	MF926603.1
Canine morbillivirus isolate BJ16C7, complete genome	185	185	100%	2e-43	100.00%	MF926602.1
Canine morbillivirus isolate BJ16C0, complete genome	185	185	100%	2e-43	100.00%	MF926601.1
Canine morbillivirus isolate BJ16B35, complete genome	185	185	100%	2e-43	100.00%	MF926599.1
Canine distemper virus isolate Nobi, complete genome	185	185	100%	2e-43	100.00%	KY971530.1
Canine distemper virus isolate Bucharest, complete genome	185	185	100%	2e-43	100.00%	KY971529.1
Canine distemper virus strain CDV-L, complete genome	185	185	100%	2e-43	100.00%	KM926612.1
Canine distemper virus strain Snyder Hill culture-collection ATCC:VR-1587, complete genome	185	185	100%	2e-43	100.00%	<u>JN896987.1</u>
Canine distemper virus strain Shuskiy, complete genome	185	185	100%	2e-43	100.00%	HM063009.1
Canine distemper virus strain Phoca/Caspian/2007, complete genome	185	185	100%	2e-43	100.00%	HM046486.1
Canine distemper virus strain recombinant Snyder Hill, complete genome	185	185	100%	2e-43	100.00%	<u>GU138403.1</u>
Canine distemper virus isolate CDV3-C nucleocapsid protein (C) mRNA, partial cds	185	185	100%	2e-43	100.00%	FJ977579.1
Canine distemper virus strain CDV3, complete genome	185	185	100%	2e-43	100.00%	EU726268.1
Canine distemper virus strain Lederle nucleoprotein gene, complete cds	185	185	100%	2e-43	100.00%	EF418783.1
Canine distemper virus strain CDV3 nucleocapsid protein (NP) gene, complete cds	185	185	100%	2e-43	100.00%	EF375619.1

Sequence ID: MF926604.1 Length: 15690 Number of Matches: 1 Range 1: 125 to 224

Range I. 120					_			
Score	Expect	Identities	Gaps	Strand	Frame			
185 bits(100)	u u	100/100(100%)	0/100(0%)	Plus/Plus				
Query 1 Sbict 125	AAGCCTCACACTG	TCAAGAGGACTCGGG			3G 60 3G 184			
	AAGEETCACACTG		ACCAACCCCCTCTTGO		36 184			
Query 61 Sbjct 185			TAGTCCTAATC 100					
50jct 185	AOCAATAAOAOGA	ATAAAGCATGTCATTA	TAGICCIAATC 224					
Canine Morbi	llivirus Isolate BJ1	6C8, Complete Gen	ome					
Sequence ID Range 1: 125		igth: 15690 Numbe	r of Matches: 1					
-								
Score	Expect	Identities	Gaps	Strand	Frame			
185 bits(100)		100/100(100%)	0/100(0%)	Plus/Plus				
Query 1	AAGCCTCACACTG	TTCAAGAGGACTCGGG						
Sbjct 125	AAGCCTCACACTG	TTCAAGAGGACTCGGG			ŚĠ 184			
Query 61		ATAAAGCATGTCATTA	111111111111					
Sbjct 185	AUCAATAAUAUGA	ATAAAGCATGTCATTA	tágtéétááté 224					
Canine Morbi	llivirus Isolate BJ1	6C7, Complete Gen	ome					
		igth: 15690 Numbe	er of Matches: 1					
Range 1: 125	to 224							
Score	Expect	Identities	Gaps	Strand	Frame			
185 bits(100)	u u	100/100(100%)	0/100(0%)	Plus/Plus				
Query 1	AAGCCTCACACTG							
Sbjct 125	AAGCCTCACACTG			******	5G 184			
Query 61	AGCAATAAGAGGA		TAGTCCTAATC 100					
Sbjct 185	AGCAATAAGAGGA	ATAAAGCATGTCATTA	tágtéétááté 224					
Canine Morbi	llivirus Isolate BJ1	6C0,Complete Gen	ome					
Sequence ID: MF926601.1 Length: 15690 Number of Matches: 1								
Range 1: 125	10 224							
Score	Expect	Identities	Gaps	Strand	Frame			
185 bits(100)	2e-43()	100/100(100%)	0/100(0%)	Plus/Plus				
Query 1	AAGCCTCACACTG	TTCAAGAGGACTCGGG	ACCAACCCCCTCTTGC					
Sbjct 125	AAGCCTCACACTG	ttčAAGAGGACTCGGG	ACCAACCCCCCTCTTGC	ttttggttctggg	Ġ 184			
-								
Query 61	AGCAATAAGAGGA	ATAAAGCATGTCATTA	TAGTCCTAATC 100					
-	AGCAATAAGAGGA AGCAATAAGAGGA	ATAAAGCATGTCATTA	TAGTCCTAATC 100					
Query 61	AGCAATAAGAGGA AGCAATAAGAGGA	ATAAAGCATGTCATTA 1 4TAAAGCATGTCATTA	11111111111					

Canine Morbillivirus Isolate BJ16B35, Complete Genome Sequence ID: MF926599.1 Length: 15690 Number of Matches: 1 Range 1: 125 to 224

Figure 3. Alignment of CDV RT-PCR product sequence with the sequences registered in the gene bank.

Table 1. The Results of Chi-Square Tests for the Desired Variables ^a										
Variables	CDV Rapid Test	CDV Sign	Candida albicans PCR Gender		Age Group	CDV RT-PCR				
CDV rapid test	-	Significant	Significant	Not significant	Significant	Significant				
CDV Sign	Significant	-	Significant	Not significant	Not significant	Significant				
C. albicans PCR	Significant	Significant	-	Not significant	Significant	Not significant				
Gender	Not significant	Not significant	Not significant		-	Not significan				
Age group	Significant	Not significant	Significant		-	Significant				
CDV RT-PCR	Significant	Significant	Not significant	Not significant	Significant					

^aP-Values of less than 0.05 were considered significant

The use of a rapid distemper diagnostic kit is the best option for diagnosing the virus in veterinary clinics. Its specificity and sensitivity for conjunctival specimens are similar to nested-PCR (24). There is no interference between vaccination and this diagnostic method (25). So, this assay was used for the initial diagnosis. Previously, some research groups have worked on the issue of CDV co-infection with other pathogens, some of which are mentioned below:

We could not find any report of simultaneous infections with distemper in Iran; there are some reports of CDV

Edit Search	1	Cummanus						n -		
		h Summary 🗸		w to read this report	? 🗗 E	BLAST Help	Videos	DBack to	Tradition	nal Results Pa
b Title	CanR. ab1			r Results						
D		pires on 01-15 13:27 pm Downlo		anism only top 20	will appe	ar				exclud
ogram	BLASTN @ <u>Citation</u> ~			pe common name			r group n	ame		
lery ID	nt <u>See details</u> ~			Add organism	, 511011	iui, tuxiu o	Bioabi	iunite.		
escription	Icl Query_47579 None									
olecule type	dna		Per	cent Identity	EV	alue		Que	ry Cover	age
uery Length	498			to		to			to	
her reports Descriptions	Distance tree of results M	ISA viewer 😨 Alignments Taxono	omy						Filter	Reset
Sequences	producing significant a	lignments		Download	× (Select	column	s × S	how 1	100 🗸 😗
select all	100 sequences selected			GenBank	<u>Gra</u>	phics D	stance tre	ee of resu	ults Ne	MSA Viewe
		Description		Scientific Name		Total Quer		Per. Ident	Acc. Len	Accession
Candida alt	picans strain 256PhC Erg11 (Erg11	1) gene, complete cds		Candida albicans	Ŧ	891 99%	0.0	99.19%	1840	KY967260.1
		erol 14-alpha-demethylase (ERG11) g	pene, partial cds	Candida albicans	891	891 99%	0.0	99.19%	1548	MN537036.1
Synthetic co	onstruct isolate SCERG11R1S1C1	1 14aloha-lanosterol demethylase (EF	RG11) gene, complete co	is synthetic construct	891	891 99%	0.0	99.19%	1587	MK416241.1
		P450 lanosterol 14-alpha-demethylas				891 99%	0.0	99.19%	1587	KM609920.1
		450 L1A1 (ERG11) gene, complete co P450 L1A1 (ERG11) gene, complete o		Candida albicans Candida albicans		891 99% 891 99%	0.0	99.19% 99.19%	1810 1830	GU371859.1 GU371857.1
		P450 L1A1 (ERG11) gene, complete (Candida albicans		891 99%	0.0	99.19%	1832	GU371857.1
	and the second	450 L1A1 (ERG11) gene, complete c		Candida albicans		891 99%	0.0	99.19%	1840	GU371853.1
Candida alt	picans strain ATCC 90028 cytochro	ome P450 L1A1 (ERG11) gene. comp	plete cds	Candida albicans	891	891 99%	0.0	99.19%	1840	GU371851.1
	bicans strain 4242 lanosterol 14-al	lpha-demethylase (ERG11) gene, con	mplete cds	Candida albicans	891	891 99%	0.0	99.19%	1587	HM194180.1
		e P-450 lanosterol 14-alpha-demethy				891 99%	0.0	99.19%	1587	EU885938.1
		rome P-450 lanosterol 14-alpha-dem- ochrome P-450 lanosterol 14-alpha-de				891 99% 891 99%	0.0	99.19% 99.19%	1370 1370	FJ232384.1 FJ159492.1
		ochrome P-450 lanosterol 14-alpha-de				891 99%	0.0	99.19%	1370	FJ159492.1
		ochrome P-450 lanosterol 14-alpha-de				891 99%	0.0	99.19%	1370	FJ159484.1
Candida alt	picans strain 03-200663 7921 cyto	ochrome P-450 lanosterol 14-alpha-de	emethylase (ERG11) gen	Candida albicans	891	891 99%	0.0	99.19%	1370	FJ159483.1
	oicans strain CA56292 cytochrome	e P450 lanosterol 14-alpha-demethyla	ase (ERG11) gene. comp	ol Candida albicans	891	891 99%	0.0	99.19%	1587	<u>GQ202075.1</u>
		-450 lanosterol 14-alpha-demethylase				891 99%	0.0	99.19%	1587	FJ002303.1
		2-450 lanosterol 14-alpha-demethylasi 2-450 lanosterol 14-alpha-demethylasi				891 99% 891 99%	0.0	99.19% 99.19%	1587 1587	EU980038.1 EU848303.1
		-450 lanosterol 14-alpha-demethylas			891	891 99%	0.0	99.19%	1587	EU819550.1
		-450 lanosterol 14-alpha-demethylas			891	891 99%	0.0	99.19%	1587	EU819549.1
		cytochrome P450 L1A1	(ERG11) gene, c	omplete cds						
	ID: GU371859.1 Length: 1 1043 to 1536 GenBank Gran		▼ Next Ma	tch 🔺 Previous Match						
Score 891 bits(Expect Iden	ntities Gaps 0/494(99%) 0/494(09	Strand							
Query 1	ATGGGTGGTCAACATACTT	CTGCTTCTACTTCTGCTTGGTTCTTG	STTACATTTAGGTGAA	60						
Sbjct 1 Query 6		ĊŦĠĊŦŦĊŦĂĊŦŦĊŦĠĊŦŦĠĠŦŦĊŦŦĠ ŧŦŦAŦŦŦAŦĊAAGAAGŦŦĠŦŦĠAAŦŦA		1102 120						
Sbjct 1	103 AAACCTCATTTACAAGATG	TTATTTATCAAGAAGTTGTTGAATTA	ATTGAAAGAAAAAGGT	1162						
Query 1 Sbjct 1		CTTATGAAGATTTACAAAAATTACCA CTTATGAAGATTTACAAAAATTACCA		180 1222						
Query 1 Sbjct 1		TGCATATGCCATTACATTCTATTTT		240 1282						
Query 2	41 CCATTAAGAATCCCTGAAA	CCAATTATATTGTTCCAAAAGGTCAT	TATGTTTTAGTTTCT	300						
Sbjct 1		kécaattatattéttéédaaaágétéat Igtgaaagatattttgataaccetgaa		1342 360						
Query 3		GTGAAAGATATTTTGATAACCCTGAA	AGATTTTGATCCAACT	1402						
Sbjct 1		CCAAAGCTAATICTGTTTCATTTAAC	TOTICIGATGAAGTT	420						
Sbjct 1 Query 3 Sbjct 1	61 AGATGGGATACTGCTGCTG 403 AGATGGGATACTGCTGCTG	CCAAAGCTAATTCTGTTTCATTTAAC	CTCTTCTGATGAAGTT	1462						
Sbjct 1 Query 3	AGATGGGATACTGCTGCTG 403 AGATGGGATACTGCTGCTG 21 GATTATGGGTTTGGGAAAG		TCTTCTGATGAAGTT	1462 480						

Figure 4. Alignment of C. albicans PCR product sequence with the sequences registered in the gene bank



Figure 5. The culture of C. albicans on Sabouraud dextrose agar

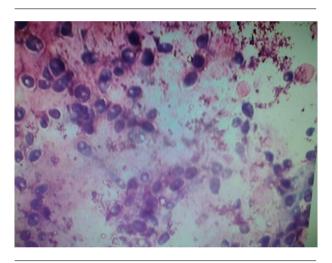


Figure 6. Giemsa staining of C. albicans

infection from some cities, such as a study in which serologic detection of CDV in unvaccinated dogs from Ahvaz were performed by Avizeh et al. in 2007. Furthermore, Namroudi et al. (2015) detected the Arctic and European cluster of CDV in the north and center of Iran (3, 4). In 2008, Gabriel et al. reported simultaneous infections of canine adenoviruses type-2 (CAV-2), CDV and different types of candidas in two 3-month-old dogs (5). In 2015, Headley et al. reported a new pneumonia case in a dog caused by *talaromyces marneffei* along with distemper infection (26). In another work, in 2015, Kubiski et al. reported co-infections of CDV and Sarcocystis species (27).

In 2006, Panasiti et al. reported a patient infected by simultaneous infections of Herpes simplex virus type-2 (HSV-2), human immunodeficiency virus (HIV), and *C. albicans* (28). Toplu et al. (2009) reported the co-infection of CDV, hepatozoonosis, and toxoplasmosis in 6 dogs with visceral leishmaniosis (29). Furthermore, Lemberger et al. (2005) reported the simultaneous infections of distemper virus and *Neospora caninum* in a dog (30).

Mochizuki et al. (2008) studied the pathogens of upper respiratory tract infections in domestic dogs. Canine parainfluenza virus (4.7%), canine coronavirus group 1 (4.4%), canine adenovirus type 2 (2.9%), canine coronavirus group 2 (1.5%), and canine distemper virus (1.5%), were detected and two cases of co-infections were reported in this study (11).

In the research performed by Damián et al. (2005), CDV was detected in 77%, CAV in 57%, and CpiV in 51% of cases, respectively. The most common co-infection was CDV-CpiV in 14% of cases (21). In the study of Headley et al. (2018), CDV-associated infections were reported as follows: *N. caninum* (100%), CPV-2 (100%), CAdV-1 (100%), and CAdV-2 (100%) (11). Aguiar et al. (2012) investigated the co-infection of CDV and *T. gondii* in dogs with neurological signs. Accordingly, 80.9% of dogs were positive for CDV using RT-PCR and 38.1% were positive for anti-*T. gondii* antibodies. Moreover, 41.1% were positive for both factors (6). In 2015, Headley et al. detected the co-infection of distemper virus with canine herpesvirus type 1, adenovirus and parvovirus in domestic dogs in southern Brazil (6).

The results of all mentioned researches indicate the high rate of co-infection of distemper with other pathogens confirms the hypothesis of increasing the susceptibility to other pathogens due to CDV immunosuppression. However, performing more similar studies on other hosts of the virus can give us complete information about this issue.

5.1. Conclusions

In total, the frequency of co-infection was four cases among the samples taken from symptomatic dogs and any statistically considerable association was not observed between distemper and *C.albicans*.

Footnotes

Authors' Contribution: Azam Mokhtari and Somayeh Shahrokh Shahraki conceived and designed the study and Seyedeh Fatemeh Ahmadi performed the experiments and analyzed the samples. All authors interpreted the data, critically revised the manuscript for important intellectual content, and approved the final version of the manuscript.

Ethical Approval: This study was performed in accordance with the Declaration of Helsinki guidelines (ethics code: 180/1345).

Funding/Support: This study was supported by a grant from Shahrekord University (grant number: 98GRD30M1801).

References

- Isfahani MS, Rostami A, Bahonar A, Barin A, Memarian I. Serologic survey for canine distemper virus in free-ranging wild canids in the northeast of Iran. *Revue de Médecine Vétérinaire*. 2017;**168**(10-12):247–51.
- 2. Namroudi S, Rostami A, Barin A, Ardebili KM. [Antibody monitoring to canine distemper virus in unvaccinated rural dogs in the southern coastal region of Caspian sea]. *J Vet Res.* 2013;**68**(3):209–15. Persian.
- Beineke A I, Puff C, Seehusen F, Baumgartner W. Pathogenesis and immunopathology of systemic and nervous canine distemper. *Vet Immunol Immunopathol*. 2009;**127**(1-2):1-18. doi: 10.1016/j.vetimm.2008.09.023. [PubMed: 19019458].
- Aguiar D, Amude A, Santos LGF, Ribeiro M, Ueno TEH, Megid J, et al. Canine distemper virus and Toxoplasma gondii co-infection in dogs with neurological signs. *Arquivo Brasileiro de Medicina Veterinária e Zootecnia*. 2012;64(1):221-4. doi: 10.1590/s0102-09352012000100032.
- Gabriel AL, Masuda EK, Ramos AT, Inkelmann MA, Azambuja E, Graça DL. Canine adenovirus type-2 and canine distemper virus pulmonary co-infection in two Chow-Chow puppies with Candida sp esophagitis. *Braz J Vet Pathol.* 2008;1:47-51.
- Headley SA, Alfieri AA, Fritzen JT, Garcia JL, Weissenbock H, da Silva AP, et al. Concomitant canine distemper, infectious canine hepatitis, canine parvoviral enteritis, canine infectious tracheobronchitis, and toxoplasmosis in a puppy. *J Vet Diagn Invest*. 2013;**25**(1):129–35. doi: 10.1177/1040638712471344. [PubMed: 23293164].
- Headley SA, Oliveira TES, Pereira AHT, Moreira JR, Michelazzo MMZ, Pires BG, et al. Canine morbillivirus (canine distemper virus) with concomitant canine adenovirus, canine parvovirus-2, and Neospora caninum in puppies: a retrospective immunohistochemical study. *Sci Rep.* 2018;8(1):13477. doi: 10.1038/s41598-018-31540-0. [PubMed: 30194440]. [PubMed Central: PMC6128882].
- Lavan R, Knesl O. Prevalence of canine infectious respiratory pathogens in asymptomatic dogs presented at US animal shelters. *J Small Anim Pract.* 2015;56(9):572-6. doi: 10.1111/jsap.12389. [PubMed: 26199194]. [PubMed Central: PMC7166506].
- Mochizuki M, Yachi A, Ohshima T, Ohuchi A, Ishida T. Etiologic study of upper respiratory infections of household dogs. *J Vet Med Sci.* 2008;**70**(6):563–9. doi: 10.1292/jvms.70.563. [PubMed: 18628596].
- Vandevelde M, Zurbriggen A. Demyelination in canine distemper virus infection: a review. *Acta Neuropathol.* 2005;**109**(1):56–68. doi: 10.1007/s00401-004-0958-4. [PubMed: 15645260].
- Siering O, Sawatsky B, Pfaller CK. C protein is essential for canine distemper virus virulence and pathogenicity in ferrets. *J Virol.* 2020. doi: 10.1128/JVI.01840-20. [PubMed: 33239455]. [PubMed Central: PMC7851556].
- Buragohain M, Goswani S, Kalita D. Clinicopathological findings of canine distemper virus infection in dogs. J Entomol Zool Stud. 2017;5(6):1817–9.
- Petrova VN, Sawatsky B, Han AX, Laksono BM, Walz L, Parker E, et al. Incomplete genetic reconstitution of B cell pools contributes to prolonged immunosuppression after measles. *Sci Immunol.* 2019;4(41). doi: 10.1126/sciimmunol.aay6125. [PubMed: 31672862].
- Damian M, Morales E, Salas G, Trigo FJ. Immunohistochemical detection of antigens of distemper, adenovirus and parainfluenza viruses in domestic dogs with pneumonia. *J Comp Pathol*. 2005;**133**(4):289–93. doi: 10.1016/j.jcpa.2005.05.005. [PubMed: 16202422].
- Feng H, Dong J, Chen Y, Shi L, Shen B, Sun X, et al. Nanoparticles of conformation-stabilized canine distemper virus hemagglutinin are

highly immunogenic and induce robust immunity. *Virol J.* 2020. doi: 10.21203/rs.3.rs-74893/v1.

- Moro L, de Sousa Martins A, de Moraes Alves C, de Araujo Santos FG, dos Santos Nunes JE, Carneiro RA, et al. Apoptosis in canine distemper. Arch Virol. 2003;148(1):153–64. doi: 10.1007/s00705-002-0903-6. [PubMed: 12536301].
- Moro L, Martins AS, Alves CM, Santos FG, Del Puerto HL, Vasconcelos AC. Apoptosis in the cerebellum of dogs with distemper. J Vet Med B Infect Dis Vet Public Health. 2003;50(5):221–5. doi: 10.1046/j.1439-0450.2003.00657.x. [PubMed: 12864896].
- Patil S, Majumdar B, Sarode SC, Sarode GS, Awan KH. Oropharyngeal Candidosis in HIV-infected patients-An update. *Front Microbiol.* 2018;9:980. doi: 10.3389/fmicb.2018.00980. [PubMed: 29867882]. [PubMed Central: PMC5962761].
- Schobesberger M, Summerfield A, Doherr MG, Zurbriggen A, Griot C. Canine distemper virus-induced depletion of uninfected lymphocytes is associated with apoptosis. *Vet Immunol Immunopathol.* 2005;**104**(1-2):33–44. doi: 10.1016/j.vetimm.2004.09.032. [PubMed: 15661329]. [PubMed Central: PMC7127010].
- Brown AJ. Morphogenetic signaling pathways in Candida albicans. In: Calderone RA, editor. *Candida and candidiasis*. ASM Press; 2002. p. 95–106.
- McCarthy MW, Walsh TJ. Candidemia in the cancer patient: diagnosis, treatment, and future directions. *Expert Rev Anti Infect Ther*. 2018;**16**(11):849–54. doi: 10.1080/14787210.2018.1536546. [PubMed: 30322269].
- Wang Y. Looking into Candida albicans infection, host response, and antifungal strategies. *Virulence*. 2015;6(4):307-8. doi: 10.1080/21505594.2014.1000752. [PubMed: 25590793]. [PubMed Central: PMC4601349].
- Khan Z, Mustafa A. Detection of Candida species by polymerase chain reaction (PCR) in blood samples of experimentally infected mice and patients with suspected candidemia. *Microbiol Res.* 2001;**156**(1):95– 102. doi: 10.1078/0944-5013-00072. [PubMed: 11372661].
- 24. Mosalanezhad B, Avizeh R, Seyfiabad Shapouri M R, Ramesh B. Antigenic detection of Canine Parainfluenza virus in urban dogs with respiratory disease in Ahvaz area, southwestern Iran. *Archives of Razi Institute,*. 2009;**2**:115–22.
- An DJ, Kim TY, Song DS, Kang BK, Park BK. An immunochromatography assay for rapid antemortem diagnosis of dogs suspected to have canine distemper. *J Virol Methods*. 2008;**147**(2):244–9. doi: 10.1016/j.jviromet.2007.09.006. [PubMed: 17950910]. [PubMed Central: PMC7119767].
- Headley SA, Pretto-Giordano LG, Lima SC, Suhett WG, Pereira AHT, Freitas LA, et al. Pneumonia due to Talaromyces marneffei in a dog from Southern Brazil with concomitant canine distemper virus infection. J Comp Pathol. 2017;157(1):61–6. doi: 10.1016/j.jcpa.2017.06.001. [PubMed: 28735673].
- Kubiski SV, Siso S, Church ME, Cartoceti AN, Barr B, Pesavento PA. Unusual necrotizing encephalitis in Raccoons and Skunks concurrently infected with canine distemper Virus and Sarcocystis sp. Vet Pathol. 2016;53(3):674–6. doi: 10.1177/0300985815603430. [PubMed: 26374278].
- Toplu N, Avci H, Metin N. Canine visceral leishmaniosis and concurrent infections (hepatozoonosis, toxoplasmosis and canine distemper). *Revue Méd. Vét.* 2009;160(7):341–8.
- Panasiti V, Devirgiliis V, Borroni RG, Spataro A, Melis L, Petrella MC, et al. Atypical cutaneous manifestation of HSV-2 with Candida albicans co-infection in a patient with HIV-1. *J Infect*. 2007;**54**(2):e55-7. doi: 10.1016/j.jinf.2006.03.036. [PubMed: 16716405].
- Lemberger KYL, Gondim FP, Pessier AP, McAllister MM, Kinsel MJ. Neospora caninum infection in a free-ranging raccoon (Procyon lotor) with concurrent canine distemper virus infection. *J Parasitol.* 2005;91(4):960–1. doi: 10.1645/GE-407R.1. [PubMed: 17089776].