



UNIVERSITI PUTRA MALAYSIA

***EXPRESSIONS OF TOLL-LIKE RECEPTORS AND CYTOKINE
MODULATION OF ANTIVIRAL INNATE IMMUNE RESPONSES IN
FELINE INFECTIOUS PERITONITIS VIRUS INFECTION***

MEGAT HAMZAH BIN MEGAT MAZHAR KHAIR

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By

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**Thesis Submitted to the School of Graduate Studies, Universiti Putra
Malaysia, in Fulfilment of the Requirements for the Degree of Master of
Science**

September 2020

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This thesis is dedicated to my beloved mother and father, who have been the support of my life.

And to my sisters Putri Raihanah and Putri Safiyah, and my brother Megat Hasan for their constant encouragement for the completion of this thesis.

Without them and the Almighty, I would not have come this far in life



Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfilment of the requirement for the degree of Master of Science

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MODULATION OF ANTIVIRAL INNATE IMMUNE RESPONSES IN FELINE
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September 2020

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Feline infectious peritonitis (FIP) is a fatal immune-mediated disease of domestic cats caused by feline infectious peritonitis virus (FIPV), a member of coronavirus family. Previous studies have shown that the lack or absence of adaptive cellular immunity and aberrant antibody production may lead to cats succumbing to FIP. FIPV viral protein has been shown to antagonize the production of type I interferon (IFN), the key cytokine crucial during early stage of viral infection and downregulate the production of anti-inflammatory cytokine such as IL-10. Natural killer (NK) cells, another important innate immune cell has been shown to be dysregulated in FIPV infection, suggesting the importance of innate immunity in FIP pathogenesis. Despite these findings, the information on pattern recognition receptors (PRRs) that play a role in the detection of common molecules on pathogens, or also known as pathogen-associated molecular patterns (PAMPs) is still lacking. Therefore, this study attempts to investigate the role of the Toll-like receptors (TLRs), one of the PRRs in FIPV infection *in vitro* and *ex vivo*. Stimulation of TLR pathway activates the NF- κ B and type I IFN-related pathways which in turn produce pro-inflammatory cytokines such as TNF- α , and type-I IFN (IFN- α , IFN- β). To achieve these objectives, Crandell-Rees Feline Kidney (CRFK) cells and feline CD14+ monocytes were infected with FIPV 79-1146 and harvested at 4, 12, and 24 hours post-infection (hpi). The infection of FIPV into these cells was confirmed by immunofluorescence (IF) assay. The mRNA expression of several TLRs (TLR3, TLR7, and TLR9) and some of downstream cytokines (TNF- α , IL-10, and IFN- β) were measured using real time PCR (qPCR). The results were then correlated with the viral load copy number. Results from the *in vitro* study revealed the involvement of TLR9 in TNF- α , and IFN- β cytokine modulation in CRFK cells. However, TLR3 was expressed at low level and remained stable throughout the *in vitro* infection while TLR7 was not detectable. In contrast, TLR7 expression was induced upon *ex vivo* infection of feline monocytes at earlier time point indicating the activation of TLR7 by binding to its ligand, single-stranded RNA. However, its expression was significantly reduced at later time points which could be due to the immune evasion strategy posed by FIPV. A similar trend was also observed

for TNF- α expression postulating the role of TLR7 in the regulation of TNF- α which has been implicated as the major pro-inflammatory cytokine seen in FIP cats. Furthermore, IFN- β gene was also expressed throughout the course of infection which could be mediated by TLR7 activation and independent of TLR3 and TLR9 signaling pathways as the level of expression of these two TLRs remained stable. In general, the viral replication kinetics in CRFK cells and feline monocytes were consistent with other studies where the virus increased from 4 hpi to 12 hpi and maintained or reduced from 12 hpi to 24 hpi. Interestingly, the expression of TLR7 in monocytes of one seronegative cat was sustained throughout the infection, and the expression of TNF- α , IL-10, and IFN- β were markedly upregulated suggesting a control of viral protein synthesis by the immune system which was confirmed by the absence of viral antigen by IF, although the viral RNA was present. Taken together, this study provides a new insight on the role of TLRs in modulating the immune responses in FIPV infection. Although different cells express a different set of TLRs as observed in this study, this report implicates the role of TLR7 and TLR9 in FIPV infection, therefore setting up an avenue for further investigation into their signaling pathway and possible formulation of new therapeutic strategies.

Keywords: Feline infectious peritonitis, Toll-like receptors, Cytokines, Gene expression, Monocytes

Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Master Sains

EKSPRESI RESEPTOR SEPERTI TOLL (TLR) DAN MODULASI SITOKIN DARI ANTIVIRUS IMUN SEMULAJADI DI DALAM JANGKITAN VIRUS PERITONITIS BERJANGKIT FELIN (FIPV)

Oleh

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Peritonitis berjangkit felin (FIP) adalah penyakit lantaran imun maut di dalam kucing domestik yang disebabkan oleh virus peritonitis berjangkit felin (FIPV), anggota keluarga coronavirus. Kajian terdahulu telah menunjukkan bahawa kekurangan atau ketiadaan keimunan sel penyesuaian dan penghasilan antibodi yang aberan boleh menyebabkan kucing tunduk kepada FIP. Protein yang dihasilkan oleh FIPV telah menunjukkan penentangan pengeluaran interferon jenis I, sitokin utama yang kritikal semasa peringkat awal jangkitan virus dan pengurangan kawal atur sitokin anti-radang seperti IL-10. Sel pembunuh semulajadi (NK) adalah sel keimunan semula jadi yang penting yang telah dinyahatur dalam jangkitan FIPV, mencadangkan kepentingan keimunan semula jadi dalam patogenesis FIP. Walaupun penemuan ini telah wujud, maklumat mengenai reseptor pengenalan pola (PRR) yang memainkan peranan dalam pengesanan molekul umum pada patogen, atau juga dikenali sebagai pola molekul yang berkaitan patogen (PAMP) masih kurang. Oleh itu, kajian ini cuba menyiasat peranan reseptor seperti toll (TLR), salah satu PRR dalam jangkitan FIPV *in vitro* dan *ex vivo*. Rangsangan laluan TLR mengaktifkan laluan yang berkaitan dengan NF- κ B dan IFN jenis I yang seterusnya menghasilkan sitokin pro-radang seperti TNF- α , dan IFN jenis I (IFN- α , IFN- β). Untuk mencapai objektif ini, sel Crandell-Rees Feline Kidney (CRFK) dan monosit CD14+ felin telah dijangkiti dengan FIPV 79-1146 dan dituai pada 4, 12 dan 24 jam selepas jangkitan (hpi). Jangkitan FIPV ke dalam sel ini telah disahkan oleh ujian imunopendarfluor (IF). Ekspresi mRNA untuk beberapa TLR (TLR3, TLR7, dan TLR9) dan beberapa sitokin hilirannya (TNF- α , IL-10, dan IFN- β) telah diukur dengan menggunakan tindak balas berantai polimerase masa nyata (qPCR). Keputusannya kemudian dikaitkan dengan jumlah salinan beban virus. Keputusan daripada kajian *in vitro* menunjukkan penglibatan TLR9 dalam pemodulatan sitokin TNF- α , dan IFN- β dalam sel CRFK. Namun, TLR3 telah diekspresi pada paras yang rendah dan tetap stabil sepanjang jangkitan *in vitro* manakala TLR7 tidak dikesan. Sebaliknya, ekspresi TLR7 dalam monosit felin telah terinduksi pada masa yang lebih awal setelah dijangkiti secara *ex vivo*, menunjukkan pengaktifan TLR7 melalui ikatan dengan ligandnya, RNA bebenang tunggal. Namun, ekspresinya telah berkurang dengan

signifikan pada masa kemudian yang mungkin disebabkan oleh strategi pengelakan imun yang dimiliki FIPV. Trend yang sama juga telah diperhatikan pada ekspresi TNF- α yang mempostulatkan peranan TLR7 dalam mengawal atur TNF- α dimana ia telah terlibat sebagai sitokin pro-radang utama dalam kucing FIP. Tambahan pula, gen IFN- β juga telah diekspresi sepanjang jangkitan berlaku yang mungkin diperantara oleh pengaktifan TLR7 dan bebas daripada laluan isyarat TLR3 dan TLR9 yang mana paras ekspresi kedua-dua TLR tersebut tetap stabil. Secara umum, kinetik replikasi virus dalam sel CRFK dan monosit felin konsisten dengan kajian-kajian terdahulu dimana virus telah meningkat daripada 4 hpi ke 12 hpi dan tetap atau berkurang dari 12 hpi ke 24 hpi. Menariknya, ekspresi TLR7 dalam monosit seekor kucing seronegatif kekal sepanjang jangkitan, dan ekspresi TNF- α , IL-10, dan IFN- β telah meningkat dengan ketara yang mencadangkan bahawa kawalan protein sintesis virus oleh sistem imun yang telah disahkan dengan ketiadaan antigen virus menggunakan IF, walaupun terdapat kewujudan RNA virus. Secara kebersamaan, kajian ini menyediakan pemahaman baru tentang peranan TLR dalam modulasi tindak-balas imun dalam jangkitan FIPV. Walaupun sel-sel berlainan mengekspresi set TLR yang berlainan seperti yang diperhatikan dalam kajian ini, laporan ini telah membabitkan peranan TLR7 dan TLR9 dalam jangkitan FIPV, maka membuka sebuah jalan untuk penyiasatan lanjut ke dalam laluan isyaratnya dan formulasi termungkin untuk strategi terapeutik yang baru

Kata Kunci: Peritonitis berjangkit felin, reseptor seperti toll (TLR), sitokin, ekspresi gen, monosit

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This thesis was submitted to the Senate of Universiti Putra Malaysia and has been accepted as fulfilment of the requirement for the degree of Master of Science. The members of the Supervisory Committee were as follows:

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TABLE OF CONTENTS

	Page
ABSTRACT	i
ABSTRAK	iii
ACKNOWLEDGEMENTS	v
APPROVAL	vi
DECLARATION	viii
LIST OF TABLES	xii
LIST OF FIGURES	xiii
LIST OF ABBREVIATIONS	xiv
CHAPTER	
1 INTRODUCTION	1
2 LITERATURE REVIEW	4
2.1 Feline Coronavirus (FCoV)	4
2.1.2 FCoV Classification	4
2.2 Feline Infectious Peritonitis (FIP)	5
2.2.1 Background	5
2.2.2 Causes	6
2.2.3 Pathogenesis	7
2.2.4 Infection and Immunity	7
2.2.5 Diagnosis of FIP	9
2.2.6 Treatment of FIP	10
2.3 The Toll-Like Receptors (TLRs)	10
2.3.1 History	10
2.3.2 Family	13
2.3.3 Protein Structure	14
2.3.4 TLR Expression and Ligands	16
2.3.5 TLR Signaling	20
2.3.6 TLR in Feline Species	22
3 MATERIALS AND METHODS / METHODOLOGY	25
3.1 Cell and Virus Culture	25
3.1.1 Cells, Virus and Reagents	25
3.1.2 CRFK Cell Culture	25
3.1.3 Virus Propagation	26
3.1.4 FIPV Confirmation	26
3.2 <i>In Vitro</i> FIPV Infection in CRFK Cells	27
3.2.1 FIPV Infection in CRFK Cells	27
3.2.2 RNA Extraction and cDNA Synthesis	27
3.3 <i>Ex Vivo</i> FIPV Infection in Feline Blood Monocyte (CD14+)	28
3.3.1 Institutional Animal Care and Use Committee (IACUC) and Cat's Criteria	28
3.3.2 Blood Collection and PBMC Isolation	28

3.3.3	FIPV Infection in Monocytes (CD14+)	28
3.3.4	Flow Cytometry	29
3.3.5	RNA Extraction and cDNA Synthesis	29
3.4	Taqman-Based Real Time PCR (qPCR)	30
3.4.1	Viral Quantification (Absolute Quantification)	30
3.4.2	Gene Expression (Relative Quantification)	31
3.4.3	Data Analysis	32
4	RESULTS	33
4.1	Titration (TCID ₅₀) and Confirmation of FIPV 79-1146	33
4.2	<i>In Vitro</i> FIPV Infection in CRFK Cells	36
4.3	<i>Ex Vivo</i> FIPV Infection in Feline Monocytes from Seropositive Cats	37
4.4	<i>Ex Vivo</i> FIPV Infection in Feline Monocyte from One Seronegative Cat	45
5	DISCUSSION AND FUTURE RECOMMENDATIONS	48
	REFERENCES	52
	APPENDICES	68
	BIODATA OF STUDENT	75

LIST OF TABLES

Table		Page
2.1	TLRs and their ligands	19
3.1	Heat cycle for the TaqMan detection of FCoV 7b gene	30
3.2	List primers and probes for TLRs and innate immune cytokines	31
4.1	Information of cat samples used for monocytes isolation	40
4.2	Comparison of gene expression and viral load between monocytes from seronegative cat (n = 1) and monocytes from seropositive cats (n = 8) after FIPV 79-1146 infection	47

LIST OF FIGURES

Figure		Page
2.1	FCoV structure and genome	5
2.2	Clinical features of FIP	6
2.3	The development of TLR field over time	13
2.4	Crystal structure of TLRs LRR and their ligands	16
2.5	Cellular localization of TLRs and their signaling pathways	22
4.1	Confirmation of FIPV using ICC-IF	34
4.2	RT-qPCR standard curve that detects FCoV 7b gene	35
4.3	FIPV 79-1146 infection induced the expression of TLR9, TNF- α , and IFN- β in CRFK after 24 h	37
4.4	Representative of gating strategy to determine monocytes purity for cat 16 (A) and cat 20 (B)	38
4.5	Representative of gating strategy to determine CD4 to CD8 ratio for cat 16	39
4.6	Representative of gating strategy to determine CD4 to CD8 ratio for cat 20	40
4.7	FIPV 79-1146 infection downregulated the expression of TLR7 and TNF- α in monocytes from seropositive cats after 24 and 12 hpi respectively	42
4.8	Type II FIPV infection in monocytes originating from seropositive cats	43
4.9	FIPV detection for cat 16 using immunofluorescence at 4 hours post-infection (hpi) (A), 12 hpi (B), and 24 hpi (C)	44
4.10	FIPV detection for cat 20 using immunofluorescence at 4 hours post-infection (hpi) (A), 12 hpi (B), and 24 hpi (C)	45

LIST OF ABBREVIATIONS

Ab	Antibody
ADE	Antibody-Dependent Enhancement
Ag	Antigen
AP-1	Activator Protein 1
APN	Aminopeptidase N
BMDC	Bone Marrow-Derived DC
BSL2	Biosafety Level 2
CCL8	Chemokine (C-C Motif) Ligand 8
CCV	Canine Coronavirus
CD	Cluster of Differentiation
CD14	Cluster of Differentiation 14
CD4	Cluster of Differentiation 4
CD8	Cluster of Differentiation 8
cDNA	Complementary DNA
CMI	Cell-Mediated Immunity
CPE	Cytopathic Effect
CpG	5'Cysteine-Phosphate-Guanine 3'
CRFK	Crandell Reese Feline Kidney
Ct	Threshold Cycle
CXCL10	C-X-C Motif Chemokine 10
DAPI	4',6-Diamidino-2-Phenylindole
DC	Dendritic Cell
DD	Death Domain

Dif	Dorsal-Related Immune Factor
DMSO	Dimethylsulfoxide
DSH	Domestic Short Hair
dsRNA	Double-Stranded RNA
E	Envelope
EDTA	Ethylenediaminetetraacetic Acid
EMCV	Encephalomyocarditis Virus
ER	Endoplasmic Reticulum
F	Forward
FBS	Fetal Bovine Serum
FCGS	Feline Chronic Gingivostomatitis
FCoV	Feline Coronavirus
fcwf-4	<i>Felis Catus</i> Whole Fetus
FECV	Feline Enteric Coronavirus
FeLV	Feline Leukemia Virus
FIP	Feline Infectious Peritonitis
FIPV	Feline Infectious Peritonitis Virus
FIV	Feline Immunodeficiency Virus
GAPDH	Glyceraldehyde 3-Phosphate Dehydrogenase
G-CSF	Granulocyte-Colony Stimulating Factor
GM-CSF	Granulocyte Macrophage-Colony Stimulating Factor
HBV	Hepatitis B Virus
HCV	Hepatitis C Virus
HIV-1	Human Immunodeficiency Virus 1
hpi	Hours Post Infection

HSV-1	Herpes Simplex Virus 1
HSV-2	Herpes Simplex Virus 2
hToll	Human Toll
IBV	Infectious Bronchitis Virus
ICAM-1	Intracellular Adhesion Molecule 1
ICC	Immunocytochemistry
IF	Immunofluorescence
IFAT	Immunofluorescent Antibody Test
IFN	Interferon
IFN- α	Interferon Alpha
IFN- β	Interferon Beta
IFN- γ	Interferon Gamma
IFN- ω	Interferon Omega
IgA	Immunoglobulin A
IgG	Immunoglobulin G
IHC	Immunohistochemistry
IKK	I κ B Kinase
IL-10	Interleukin 10
IL-12	Interleukin 12
IL-17	Interleukin 17
IL-1RI	IL-1 Receptor Type I
IL-1 β	Interleukin 1 Beta
IL-6	Interleukin 6
IL-8	Interleukin 8
ILC	Islet-Like Cluster

IRAK1	IL-1R-Associated Kinase 1
IRAK2	IL-1R-Associated Kinase 2
IRAK4	IL-1R-Associated Kinase 4
IRF	Interferon Regulatory Factor
I κ B	Inhibitor of Kappa B
Jak-STAT	Janus Kinases Signal Transducer and Activator of Transcription Proteins
LPL	Lamina Propria Lymphocytes
LPS	Lipopolysaccharide
LRR	Leucine-Rich Repeat
LTA	Lipoteichoic Acid
LTC	Liposome-TLR Complex
M	Membrane
MAPK	Mitogen-Activated Protein Kinase
MAVS	Mitochondrial Antiviral-Signaling Protein
MCMV	Mouse Cytomegalovirus
MD-2	Myeloid Differentiation 2
MEM	Minimal Essential Media
MERS-CoV	Middle East Respiratory Syndrome Coronavirus
MHV	Mouse Hepatitis Virus
miRNA	Micro RNA
MLN	Mesenteric Lymph Node
MMP-9	Matrix Metalloproteinase-9
MOI	Multiplicity of Infection
mRNA	Messenger RNA

Mx	Myxovirus Resistance
MX1	MX Dynamin-Like Gtpase 1
MyD88	Myeloid Differentiation Primary Response 88
N	Nucleocapsid
NCBI	National Center for Biotechnology Information
NEMO	NF-Kappa-B Essential Modulator
NF-κB	Nuclear Factor Kappa-Light-Chain-Enhancer of Activated B Cells
NK	Natural Killer
Nrdp1	Neuregulin Receptor Degradation Protein-1
NRT	No Reverse Transcriptase
NT	N Terminal
ODN-2216	Oligodeoxynucleotide-2216
ORF	Open Reading Frame
PAMP	Pathogen-Associated Molecular Pattern
PBMC	Peripheral Blood Mononuclear Cells
PBS	Phosphate Buffered Saline
PBST	Phosphate Buffered Saline 0.1% Tween
PC	Peritoneal Cells
pDC	Plasmacytoid Dendritic Cell
PEDV	Porcine Epidemic Diarrhea Virus
PFA	Paraformaldehyde
PI	Post Infection
PMN	Polymorphonuclear
Poly (I:C)	Polyinosinic:Polycytidylic Acid

PRR	Pathogen Recognition Receptors
qPCR	Real-Time PCR
R	Reverse
R-848	Resiquimod-848
RHD	Rel Homology Domain
RHIM	Receptor-Interacting Protein Homotypic Interaction Motif
RIG-1	Retinoic Acid-Inducible Gene 1
RIM	Rapid Immunochromatography
RIP1	Receptor-Interacting Protein 1
rpm	Rotation Per Minute
RT-PCR	Reverse Transcriptase Polymerase Chain Reaction
RT-qPCR	Real-Time Reverse Transcriptase Polymerase Chain Reaction
S	Spike
SARS	Severe Acute Respiratory Syndrome
SARS-CoV	Severe Acute Respiratory Syndrome Coronavirus
ssRNA	Single-Stranded RNA
STAT2	Signal Transducer and Activator of Transcription 2
TAB2	TGF-Beta-Activated Kinase 1 and MAP3K7-Binding Protein 2
TAB3	TGF-Beta-Activated Kinase 1 and MAP3K7-Binding Protein 3
TAK1	Transforming Growth Factor- β (TGF- β)-Activated Kinase 1
TANK	TRAF Family Member-Associated NF-Kappa-B Activator
TBEV	Tick-Borne Encephalitis Virus
TBK1	TANK-Binding Kinase 1
TCID ₅₀	Tissue Culture Infectious Dose 50
TGEV	Transmissible Gastroenteritis Virus

Th1	T Helper 1
Th17	T Helper 17
TICAM-2	TIR-Containing Adaptor Molecule 2
TIR	Toll-IL-1-Resistant
TIRAP	TIR Domain-Containing Adaptor Protein
TLR	Toll-Like Receptor
TNFR	Tumor Necrosis Factor Receptor
TNF- α	Tumor Necrosis Factor Alpha
TRADD	TNFR-Associated Death Domain
TRAF3	TNF Receptor-Associated Factor 3
TRAF6	TNF Receptor-Associated Factor 6
TRAM	TRIF-Related Adaptor Molecule
TRIF	TIR-Domain-Containing Adapter-Inducing Interferon- β
tRNA	Transfer RNA
Ubc13	Ubiquitin Conjugating Enzyme E2
Uev1A	Ubiquitin-Conjugating Enzyme E2 Variant 1A
Unc93b1	Unc-93 homolog B1
VCAM-1	Vascular Cell Adhesion Molecule 1
VEGF	Vascular Endothelial Growth Factor
VSV	Vesicular Stomatitis Virus
WNV	West Nile Virus
β 2M	Beta-2-Microglobulin

CHAPTER 1

INTRODUCTION

Feline infectious peritonitis (FIP) is a severe inflammatory disease affecting young domestic cats caused by FIP virus (FIPV), one of the two biotypes of feline coronavirus (FCoV). The other biotype is called feline enteric coronavirus (FECV) that caused mild illnesses such as self-limiting enteritis in kittens (Pedersen, 2009). The FCoV is categorized under the Coronaviridae family that has the largest RNA genome among other families of RNA viruses, with size of about 32 kB of positive-strand RNA (Dye and Siddell, 2007). Other members of Coronaviridae species that caused significant disease in animals are infectious bronchitis virus (IBV) affecting chickens and transmissible gastroenteritis virus (TGEV) affecting pigs that caused huge economic losses in poultry and farm industry.

The frequency of cats infected with FECV is extremely common especially in catteries and shelters, yet the chance of them to develop FIP is very rare which could be due to the avirulent nature of most field strain (Addie, 2000). However, several key events can trigger the development of FIP. During an intense viral replication in the gut, there is a higher chance of internal mutation events to occur inside FECV that can lead to a biotype shift towards FIPV. Yet, this internal mutation is not the only determinant of FIP as other factors like host's genetics, geographical region, and immune status of the cat also play critical role on the disease outcome (Myrrha et al., 2011).

FIP has two forms which are the effusive ('wet') FIP which contributes largely in clinical cases and has higher percent mortality; and the non-effusive ('dry') FIP. Symptoms of both forms include fever, weight loss, and loss of appetite; but the distinguishing feature between the two is that wet FIP cats show distended abdomen that is filled with protein-rich fluid while dry FIP cats usually is presented with granulomatous lesions and phlebitis in multiple organs (Kipar and Meli, 2014). The underlying reason for these forms to arise is largely due to the immune strength of the host, where the cell-mediated immune (CMI) responses of wet FIP cats are generally abrogated while dry FIP cats can mount some undetermined level of CMI.

There are several indirect approaches used to diagnose FIP which includes examining the clinical history such as cat's previous exposure to multi-household environment, stress level, presence of fluctuating fever, and loss of appetite. The diagnosis can be followed by analysis of serum albumin to globulin (A:G) ratio, FCoV-specific serology test, and reverse transcriptase polymerase chain reaction (RT-PCR) detecting viral RNA in tissues or effusion. The use of one test is usually not sufficient to suggest for FIP because each test has its limitation, therefore, the combination of different tests is very important to provide definitive diagnosis of FIP (Tasker, 2018).

Various efforts have been established to develop safe and effective therapy for FIP. Many of the earliest treatment in the 70's to late 90's involved the use of

immunosuppressive drugs in combination with antibiotics such as prednisolone and penicillin. This strategy was proven to be ineffective as many cats suspected to have FIP succumb to the disease (Hartmann and Ritz, 2008). The approaches used today are essentially similar which mainly involved the combination of several therapeutics agents such as anti-viral drugs such as ribavirin and interferon- α . Nevertheless, the number of proper clinical trials and controls to test the safety and efficacy of such regime is still lacking, therefore, definitive treatment for FIP is still unavailable.

In an effort to answer the perplexing questions surrounding FIP and its treatment, many interests have drifted towards investigating the role of innate immunity in positively modulating the immune system to counteract with FIP. Although it was discovered in the 90's from experiments involving the drosophila Toll protein, significant progress has been made that further expands our knowledge of this system. Amongst the earliest component of innate immunity in mammalian cells discovered was the toll-like receptors (TLRs) family which functions as unique sensors for numerous types of microbial. Recognition of microbial by TLRs activate the nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) and interferon regulatory factor (IRF) pathways which are the molecular switches that lead to the production of various cytokines and type I interferons (IFN) for the clearance of pathogens (Akira and Takeda, 2004).

There are currently ten human TLRs and twelve mice TLRs being discovered and characterized. Feline tissues are also shown to express at least nine TLRs (Ignacio et al., 2005). TLR3, TLR7, and TLR8 specifically detects virus-associated nucleic acid such as single-stranded (ss) RNA or double-stranded (ds) RNA (Jiménez-Dalmaroni et al., 2015). A number of viruses that are being recognized by these TLRs do elicit protective immune response that ultimately results in virus clearance. However, some viruses also exploit the TLRs signaling in order to provide favorable conditions for their replication. Therefore, characterization of TLR-recognition pathways in virus infection is pertinent in order to design immunotherapies that can either stimulate or suppress the TLRs pathways so that specific viral infection can be controlled (Patel et al., 2014).

To provide better understanding on the role of TLRs on the pathogenesis of FIP, whether they provide protection to the host or use by the virus to evade immune responses, the gene expression of nucleic acid-sensing TLRs - TLR3, TLR7, and TLR9 were measured upon infection with FIPV in vitro using probe-based quantitative real-time PCR (qPCR). The expression of these TLRs to the production of pro-inflammatory cytokines and type I IFN were then correlated to the FIPV load at different periods of infection. The significance of our study is then verified by using cell types that are susceptible to FIPV infection both ex vivo and in vitro which are the peripheral blood monocytes and Crandell-Rees Feline Kidney (CRFK) cells.

The objectives of the study are summarized as follows:

1. To measure the gene expression of TLR3, TLR7, and TLR9, and the expression of pro-inflammatory cytokines and type I IFN upon infection with FIPV at different time points in vitro.
2. To measure the gene expression of TLR3, TLR7, and TLR9, and the expression of pro-inflammatory cytokines and type I IFN upon infection with FIPV ex vivo.
3. To determine the correlation between viral load, the TLR3, TLR7 and TLR9 and cytokines expression upon infection with FIPV.

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APPENDICES

Appendix A: Titration (TCID₅₀) and confirmation of FIPV 79-1146

Dilution	Dead	Alive	Cumulative			Percent Mortality
			Dead	Alive	Ratio	
10 ⁻¹	10	0	64	0	64/64	100
10 ⁻²	10	0	54	0	54/54	100
10 ⁻³	10	0	44	0	44/44	100
10 ⁻⁴	10	0	34	0	34/34	100
10 ⁻⁵	10	0	24	0	24/24	100
10 ⁻⁶	10	0	14	0	14/14	100
10 ⁻⁷	3	7	4	7	4/11	36.3636364
10 ⁻⁸	1	9	1	16	1/17	5.88235294
10 ⁻⁹	0	10	0	26	0/26	0
10 ⁻¹⁰	0	10	0	36	0/36	0

$$\frac{\% \text{ Above } 50\% - 50\%}{\% \text{ Above } 50\% - \% \text{ below } 50\%} = \frac{100 - 50}{100 - 36.363}$$

$$= 0.785714286$$

$$\text{Dilution at which \% above } 50\% = 10^{-6} = 6$$

$$\text{TCID}_{50} = 6 + 0.785714286 = 6.786$$

$$\text{TCID}_{50}/\text{ml} = 10^{6.786}/0.1 \text{ ml} = \mathbf{6.115 \times 10^7}$$

A.1: Calculation of TCID₅₀ for FIPV 79-1146

Appendix B: *In vitro* type II FIPV infection in CRFK cells

Normality and Lognormality Tests						
Tabular results						
	A	B	C	D	E	F
	CRFK TLR3	CRFK TLR9	CRFK IL-10	CRFK IFN β	CRFK TNF α	CRFK Viral Load
1	Test for normal distribution					
2	Anderson-Darling test					
3	A2*	0.5847	0.4085	1.101	1.220	0.8900
4	P value	0.0901	0.2701	0.0035	0.0016	0.0131
5	Passed normality test (alpha=0.05)?	Yes	Yes	No	No	No
6	P value summary	ns	ns	**	**	*
7						
8	D'Agostino & Pearson test					
9	K2	2.445	1.379	3.444	2.421	2.665
10	P value	0.2945	0.5019	0.1787	0.2981	0.2638
11	Passed normality test (alpha=0.05)?	Yes	Yes	Yes	Yes	Yes
12	P value summary	ns	ns	ns	ns	ns
13						
14	Shapiro-Wilk test					
15	W	0.8691	0.8944	0.7375	0.7231	0.7871
16	P value	0.1203	0.2211	0.0039	0.0026	0.0145
17	Passed normality test (alpha=0.05)?	Yes	Yes	No	No	No
18	P value summary	ns	ns	**	**	*
19						
20	Kolmogorov-Smirnov test					
21	KS distance	0.2839	0.2130	0.3198	0.3826	0.3207
22	P value	0.0351	>0.1000	0.0085	0.0004	0.0082
23	Passed normality test (alpha=0.05)?	No	Yes	No	No	No
24	P value summary	*	ns	**	***	**
25						
26	Number of values	9	9	9	9	9

B.1: Normality test for viral load quantification and gene expression in CRFK cells.

Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	0.000	No	ns	>0.9999	A-B
7	4 vs. 24	4.000	No	ns	0.2209	A-C
8	12 vs. 24	4.000	No	ns	0.2209	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	6.333	6.333	0.000	3	3
12	4 vs. 24	6.333	2.333	4.000	3	3
13	12 vs. 24	6.333	2.333	4.000	3	3

B.2: Kruskal-Wallis test comparing TLR3 expression between different time points in CRFK cells.

ANOVA results		Multiple comparisons				
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-3.333	No	ns	0.4081	A-B
7	4 vs. 24	-5.667	Yes	*	0.0338	A-C
8	12 vs. 24	-2.333	No	ns	0.8902	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	2.000	5.333	-3.333	3	3
12	4 vs. 24	2.000	7.667	-5.667	3	3
13	12 vs. 24	5.333	7.667	-2.333	3	3
					Z	
						1.491
						2.534
						1.043

B.3: Kruskal-Wallis test comparing TLR9 expression between different time points in CRFK cells.

ANOVA results		Multiple comparisons				
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-0.3333	No	ns	>0.9999	A-B
7	4 vs. 24	-4.667	No	ns	0.1107	A-C
8	12 vs. 24	-4.333	No	ns	0.1579	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	3.333	3.667	-0.3333	3	3
12	4 vs. 24	3.333	8.000	-4.667	3	3
13	12 vs. 24	3.667	8.000	-4.333	3	3
					Z	
						0.1491
						2.087
						1.938

B.4: Kruskal-Wallis test comparing IL-10 expression between different time points in CRFK cells.

ANOVA results		Multiple comparisons				
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-3.000	No	ns	0.5391	A-B
7	4 vs. 24	-6.000	Yes	*	0.0219	A-C
8	12 vs. 24	-3.000	No	ns	0.5391	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	2.000	5.000	-3.000	3	3
12	4 vs. 24	2.000	8.000	-6.000	3	3
13	12 vs. 24	5.000	8.000	-3.000	3	3
					Z	
						1.342
						2.683
						1.342

B.5: Kruskal-Wallis test comparing IFN- β expression between different time points in CRFK cells.

ANOVA results		Multiple comparisons				
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-3.000	No	ns	0.5391	A-B
7	4 vs. 24	-6.000	Yes	*	0.0219	A-C
8	12 vs. 24	-3.000	No	ns	0.5391	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	2.000	5.000	-3.000	3	3
12	4 vs. 24	2.000	8.000	-6.000	3	3
13	12 vs. 24	5.000	8.000	-3.000	3	3
						Z
						1.342
						2.683
						1.342

B.6: Kruskal-Wallis test comparing TNF- α expression between different time points in CRFK cells.

ANOVA results		Multiple comparisons				
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-4.000	No	ns	0.2209	A-B
7	4 vs. 24	-5.000	No	ns	0.0760	A-C
8	12 vs. 24	-1.000	No	ns	>0.9999	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	2.000	6.000	-4.000	3	3
12	4 vs. 24	2.000	7.000	-5.000	3	3
13	12 vs. 24	6.000	7.000	-1.000	3	3
						Z
						1.789
						2.236
						0.4472

B.7: Kruskal-Wallis test comparing viral load between different time points in CRFK cells.

Appendix C: *Ex vivo* type II FIPV infection in feline monocytes

Normality and Lognormality Tests								
Tabular results								
	A	B	C	D	E	F	G	
	TLR7	TLR3	TLR9	IL-10	IFNb	TNFa	Viral Load	
1	Test for normal distribution							
2	Anderson-Darling test							
3	A2*	2.125	1.886	0.7767	2.723	4.143	0.3034	0.2882
4	P value	<0.0001	<0.0001	0.0375	<0.0001	<0.0001	0.5459	0.5873
5	Passed normality test (alpha=0.05)?	No	No	No	No	No	Yes	Yes
6	P value summary	****	****	*	****	****	ns	ns
7								
8	D'Agostino & Pearson test							
9	K2	34.24	34.88	4.418	45.08	21.25	4.993	2.331
10	P value	<0.0001	<0.0001	0.1098	<0.0001	<0.0001	0.0824	0.3118
11	Passed normality test (alpha=0.05)?	No	No	Yes	No	No	Yes	Yes
12	P value summary	****	****	ns	****	****	ns	ns
13								
14	Shapiro-Wilk test							
15	W	0.6946	0.7193	0.9239	0.5965	0.5939	0.9444	0.9660
16	P value	<0.0001	<0.0001	0.0712	<0.0001	<0.0001	0.2040	0.5695
17	Passed normality test (alpha=0.05)?	No	No	Yes	No	No	Yes	Yes
18	P value summary	****	****	ns	****	****	ns	ns
19								
20	Kolmogorov-Smirnov test							
21	KS distance	0.2412	0.2449	0.1731	0.2502	0.3247	0.09552	0.09852
22	P value	0.0009	0.0007	0.0610	0.0004	<0.0001	>0.1000	>0.1000
23	Passed normality test (alpha=0.05)?	No	No	Yes	No	No	Yes	Yes
24	P value summary	***	***	ns	***	****	ns	ns
25								
26	Number of values	24	24	24	24	24	24	24

C.1: Normality test for viral load quantification and gene expression in feline monocytes.

Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	7.875	No	ns	0.0778	A-B
7	4 vs. 24	12.75	Yes	***	0.0009	A-C
8	12 vs. 24	4.875	No	ns	0.5038	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	19.38	11.50	7.875	8	8
12	4 vs. 24	19.38	6.625	12.75	8	8
13	12 vs. 24	11.50	6.625	4.875	8	8

C.2: Kruskal-Wallis test comparing TLR7 expression between different time points in feline monocytes.

ANOVA results		Multiple comparisons				
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	2.125	No	ns	>0.9999	A-B
7	4 vs. 24	0.5000	No	ns	>0.9999	A-C
8	12 vs. 24	-1.625	No	ns	>0.9999	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	13.38	11.25	2.125	8	8
12	4 vs. 24	13.38	12.88	0.5000	8	8
13	12 vs. 24	11.25	12.88	-1.625	8	8

C.3: Kruskal-Wallis test comparing TLR3 expression between different time points in feline monocytes.

ANOVA results		Multiple comparisons				
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-2.000	No	ns	>0.9999	A-B
7	4 vs. 24	-2.125	No	ns	>0.9999	A-C
8	12 vs. 24	-0.1250	No	ns	>0.9999	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	11.13	13.13	-2.000	8	8
12	4 vs. 24	11.13	13.25	-2.125	8	8
13	12 vs. 24	13.13	13.25	-0.1250	8	8

C.4: Kruskal-Wallis test comparing TLR9 expression between different time points in feline monocytes.

ANOVA results		Multiple comparisons				
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-1.000	No	ns	>0.9999	A-B
7	4 vs. 24	-2.000	No	ns	>0.9999	A-C
8	12 vs. 24	-1.000	No	ns	>0.9999	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	11.50	12.50	-1.000	8	8
12	4 vs. 24	11.50	13.50	-2.000	8	8
13	12 vs. 24	12.50	13.50	-1.000	8	8

C.5: Kruskal-Wallis test comparing IL-10 expression between different time points in feline monocytes.

ANOVA results						
Multiple comparisons						
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-6.750	No	ns	0.1687	A-B
7	4 vs. 24	-3.375	No	ns	>0.9999	A-C
8	12 vs. 24	3.375	No	ns	>0.9999	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	9.125	15.88	-6.750	8	8
12	4 vs. 24	9.125	12.50	-3.375	8	8
13	12 vs. 24	15.88	12.50	3.375	8	8
						Z
						1.909
						0.9546
						0.9546

C.6: Kruskal-Wallis test comparing IFN- β expression between different time points in feline monocytes.

ANOVA results						
Multiple comparisons						
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	9.750	Yes	*	0.0175	A-B
7	4 vs. 24	5.625	No	ns	0.3348	A-C
8	12 vs. 24	-4.125	No	ns	0.7300	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	17.63	7.875	9.750	8	8
12	4 vs. 24	17.63	12.00	5.625	8	8
13	12 vs. 24	7.875	12.00	-4.125	8	8
						Z
						2.758
						1.591
						1.167

C.7: Kruskal-Wallis test comparing TNF- α expression between different time points in feline monocytes.

ANOVA results						
Multiple comparisons						
Kruskal-Wallis test						
Multiple comparisons						
1	Number of families	1				
2	Number of comparisons per family	3				
3	Alpha	0.05				
4						
5	Dunn's multiple comparisons test	Mean rank diff.	Significant?	Summary	Adjusted P Value	
6	4 vs. 12	-6.125	No	ns	0.2496	A-B
7	4 vs. 24	-1.375	No	ns	>0.9999	A-C
8	12 vs. 24	4.750	No	ns	0.5373	B-C
9						
10	Test details	Mean rank 1	Mean rank 2	Mean rank diff.	n1	n2
11	4 vs. 12	10.00	16.13	-6.125	8	8
12	4 vs. 24	10.00	11.38	-1.375	8	8
13	12 vs. 24	16.13	11.38	4.750	8	8
						Z
						1.732
						0.3889
						1.344

C.8: Kruskal-Wallis test comparing viral load between different time points in feline monocytes.

BIODATA OF STUDENT

Megat Hamzah Megat Mazhar Khair was born on the 12th of October 1993 in Gombak, Selangor. He finished his primary education in national elementary school in 2005 and completed his secondary high school education in 2010. He later pursued his Bachelor of Science degree in molecular biosciences and biotechnology overseas in Rochester, United States of America from 2012 to 2016. Upon his return to Malaysia, he filled in the position of research assistant (RA) under Dr. Farina Mustaffa Kamal for a project relating to TLR modulation in FIPV infection. He went on to continue his Master of Science degree under the same supervisor and project until now.





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