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Supplementary Material



An Investigation of Honey Bee Viruses Prevalence in Managed Honey Bees (*Apis mellifera* and *Apis cerana*) Undergone Colony Decline

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Abstract:

Objective:

In the absence of known clinical symptoms, viruses were considered to be the most probable key pathogens of honey bee. Therefore, the aim of this study was to investigate the prevalence and distribution of honey bee viruses in managed *Apis mellifera* and *Apis cerana* in China.

Methods:

We conducted a screening of 8 honey bee viruses on *A. mellifera* and *A. cerana* samples collected from 54 apiaries from 13 provinces in China using RT-PCR.

Results:

We found that the types and numbers of viral species significantly differed between *A. mellifera* and *A. cerana*. Black Queen Cell Virus (BQCV), Chronic Bee Paralysis Virus (CBPV), *Apis mellifera* filamentous virus (AmFV), and Kakugo virus (DWV-A/KV) were the primary viruses found in *A. mellifera* colonies, whereas Chinese Sacbrood Bee Virus (CSBV) and Sacbrood Bee Virus (SBV) were the primary viruses found in *A. cerana*. The percentage infection of BQCV and CSBV were 84.6% and 61.6% in all detected samples. We first detected the occurrences of *Varroa destructor* virus-1 (VDV-1 or DWV-B) and DWV-A/KV in China but not ABPV in both *A. mellifera* and *A. cerana*.

Conclusion:

This study showed that BQCV and CSBV are the major threat to investigated *A. mellifera* and *A. cerana* colonies.

Keywords: Honey bee viruses, BQCV, CBPV, AmFV, CSBV, *A. mellifera*, *A. cerana*.

Article History

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Table S1. Numbers of apiaries selected in different province.

Virus	<i>A. mellifera</i>	<i>A. cerana</i>
Zhejiang	1	1
Henan	11	3
Hubei	0	3
Anhui	11	
Guangdong	0	2
Liaoning	13	2
Hunan	0	1
Beijing	2	1
Neimenggu	7	0
Heilongjiang	2	0

(Table S1) contd.....

Virus	<i>A. mellifera</i>	<i>A. cerana</i>
Chongqing	0	2
Jiangsu	1	0
Yunnan	0	2

Table S2. Primers used for PCR detection in present study.

Virus	Forward Primer Reverse Primer	Reference
AmFV	CAGAGAATTCGGTTTTTGTGAGTG CATGGTGGCCAAGTCTTGCT	Hartmann <i>et al.</i> , 2015
IAPV	AGACACCAATCACGGACCTCAC AGATTTGTCTGTCTCCCAGTGCAC	Maori <i>et al.</i> , 2007
SBV	ATATACGGTGCGAGAAGTGC CTCGGTAATAACGCCACTGT	Hou <i>et al.</i> , 2014
ABPV	TTATGTGTCAGAGACTGTAT GTCCTATTGCTCGGTTTTTC	Blanchard <i>et al.</i> , 2007
BQCV	TGGTCAGCTCCCACTACCTAAAC GCAACAAGAAGAAACGTAAACCAC	Benjeddou <i>et al.</i> , 2001
CBPV	TCAGACACCGAATCTGATTATTG ACTACTAGAAAACCTCGCTTCG	Berényi <i>et al.</i> , 2006
VDV-1	CATAGCGAATTACGGTGCAA GAGGGTCCCTACTCTACCG	Hou <i>et al.</i> , 2014
DWV	CTTACTCTGCCGTCGCCCA CCGTTAGGAACCTATTATCGCG	Chen <i>et al.</i> , 2005
CSBV	CCTGGGAAGTTTGCTAGTATTTACG CCTATCACATCCATCTGGGTCAG	Ma <i>et al.</i> , 2013
KBV	TATGCTGAACAACGCAAAGA ACAACACGATGTCTGGGTTT	Stolz <i>et al.</i> , 1995
KV	GACTGAACCAAATCCGATGTC TCTCAAGTTCGGGACGCATTC	Fujiyuki <i>et al.</i> , 2009

Table S3. Results of chi-square test for all types of co-infection in *A. mellifera*.

Number of Virus	Type of Co-infection	Chi-square (df=1)	P Level
2	BQCV; KV	1.81	0.18
-	BQCV; CBPV	0.22	0.64
-	CBPV; DWV	0.51	0.48
-	BQCV; DWV	1.82	0.18
-	BQCV; AmFV	0.42	0.48
-	IAPV; DWV	0.05	0.18
3	BQCV; AmFV; KV	2.56	0.11
-	BQCV; CBPV; DWV	2.56	0.11
-	BQCV; CBPV; AmFV	1.01	0.32
-	BQCV; CSBV; CBPV	2.56	0.11
-	IAPV; SBV; CSBV	1.01	0.32
-	IAPV; SBV; CBPV	0.19	0.66
-	IAPV; DWV; VDV-1	1.01	0.32
4	IAPV; SBV; CSBV; DWV	0.0026	0.95
-	IAPV; BQCV; DWV; VDV-1	12.61	0.00038
5	IAPV; DWV; VDV-1; CBPV; AmFV	4.61	0.031
-	IAPV; SBV; BQCV; DWV; CSBV	4.61	0.031
6	IAPV; BQCV; CBPV; DWV; VDV-1; AmFV	18.88	<10-5

Table S4. Results of chi-square test for all types of co-infection *A. cerana*.

Number of Viruses	Type of Co-infection	Chi-square (df=1)	P Level
2	BQCV; AmFV	0.0009	0.02
	BQCV; CSBV	0.0009	0.02
	SBV; CSBV	0.061	0.19
	BQCV; DWV	<10-5	0.004
	BQCV; CBPV	0.0009	0.02
3	BQCV; CSBV; KV	0.04	0.15
	SBV; CSBV; AmFV	0.04	0.15
	IAPV; SBV; CSBV	0.04	0.15

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