

This form should be used for all taxonomic proposals. Please complete all those modules that are applicable (and then delete the unwanted sections).

Code(s) assigned:	2008.005- 009V	(to be completed by ICTV officers)
Family Hepeviridae	the genus <i>Zetavirus</i> ;	eccies in the genus Hepevirus; formally recognize the s; re-classification of the family <i>Zetaviridae</i> etc.) $2 \boxtimes 3 \square 4 \square 5 \square$

#### Author(s) with e-mail address(es) of the proposer:

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## ICTV-EC or Study Group comments and response of the proposer:

The ICTV Hepeviridae Study Group has reviewed and approved the proposed changes.

# MODULE 2: **NEW FAMILY**

(if more than one family is to be created, please complete additional copies of this module)

Code	2008.005V	(assigned by ICTV officers)
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If there i	s no order, put "unassigned" h	nere.
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Code	2008.006V	(assigned by ICTV officers)
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Code	2009 0071/	
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#### Argument to justify the creation of a new family:

In the ICTV 8<sup>th</sup> Report, The genus "*Hepevirus*" was created which consisted of one type species "Hepatitis E virus" and one tentative species "Avian hepatitis E virus". However, the genus *Hepevirus* was not formally assigned to any family in the ICTV 8<sup>th</sup> Report even though many in the scientific community have already begun to use the family name "*Hepeviridae*" in publications.

With the identification of several strains of avian hepatitis E virus (from chickens) with only approximately 50% nucleotide sequence identity to the hepatitis E virus, members of the *Hepeviridae* Study Group have agreed that it is time to formally recognize the family *Hepeviridae* for this group of viruses. By formally recognizing a new family "*Hepeviridae*", we can now more appropriately place avian hepatitis E virus as a new species in the family (it was designated as a "tentative species in the genus" in the ICTV 8<sup>th</sup> Report).

### **Origin of the new family name:**

The family name is derived from the sigla of <u>*He*</u>patitis <u>*E*</u> <u>V</u>irus.

#### **References:**

Huang FF, Sun ZF, Emerson SU, Purcell RH, Shivaprasad HL, Pierson FW, Toth TE, Meng XJ. Determination and analysis of the complete genomic sequence of avian hepatitis E virus (avian HEV) and attempts to infect rhesus monkeys with avian HEV. J Gen Virol. 2004;85(Pt 6):1609-18.

Billam P, Sun ZF, Meng XJ. Analysis of the complete genomic sequence of an apparently avirulent strain of avian hepatitis E virus (avian HEV) identified major genetic differences compared with the prototype pathogenic strain of avian HEV. J Gen Virol. 2007;88(Pt 5):1538-44.

Guo H, Zhou EM, Sun ZF, Meng XJ, Halbur PG. Identification of B-cell epitopes in the capsid protein of avian hepatitis E virus (avian HEV) that are common to human and swine HEVs or unique to avian HEV. J Gen Virol. 2006;87(Pt 1):217-23.

#### **Annexes:**

Include as much information as necessary to support the proposal. The use of Figures and Tables is strongly recommended.