

Duck hepatitis B virus, complete genome
NCBI Reference Sequence: NC_001344.1

FASTA Graphics

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LOCUS NC_001344 3027 bp DNA circular VRL 13-AUG-2018

DEFINITION Duck hepatitis B virus, complete genome.

ACCESSION NC_001344

VERSION NC_001344.1

DBLINK BioProject: PRJNA485481

KEYWORDS RefSeq; complete genome.

SOURCE Duck hepatitis B virus

ORGANISM Duck hepatitis B virus

Viruses; Riboviria; Pararnavirae; Artverviricota; Revtraviricetes;
Blubervirales; Hepadnaviridae; Avihepadnavirus.

REFERENCE 1 (bases 1 to 3027)

AUTHORS Tong,S., Mattes,F., Blum,H.E., Fernholz,D., Schneider,R. and Will,H.

TITLE Complete nucleotide sequence of a chinese Hepatitis B virus

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 3027)

CONSRTM NCBI Genome Project

TITLE Direct Submission

JOURNAL Submitted (01-AUG-2000) National Center for Biotechnology
Information, NIH, Bethesda, MD 20894, USA

REFERENCE 3 (bases 1 to 3027)

AUTHORS Schneider,R.

TITLE Direct Submission

JOURNAL Submitted (26-JUN-1991) R. Schneider, Max-Planck Inst fuer
Biochemie, Am Klopferspitz 18a, 8033 Martinsried, FRG

COMMENT PROVISIONAL REFSEQ: This record has not yet been subject to final
NCBI review. The reference sequence is identical to X60213.
COMPLETENESS: full length.

FEATURES Location/Qualifiers

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