

Bovine adenovirus 3 strain HLJ0955, complete genome

GenBank: JN381195.1

[FASTA Graphics](#)

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LOCUS JN381195 34132 bp DNA linear VRL 28-DEC-2011

DEFINITION Bovine adenovirus 3 strain HLJ0955, complete genome.

ACCESSION JN381195

VERSION JN381195.1

KEYWORDS .

SOURCE Bovine adenovirus 3

ORGANISM [Bovine adenovirus 3](#)

Viruses; Varidnaviria; Bamfordvirae; Preplasmiviricota;

Tectiliviricetes; Rowavirales; Adenoviridae; Mastadenovirus;

Bovine

mastadenovirus B.

REFERENCE 1 (bases 1 to 34132)

AUTHORS Zhu,Y.M., Yu,Z., Cai,H., Gao,Y.R., Dong,X.M., Li,Z.L., Shi,H.F.,
Meng,Q.F., Lu,C. and Xue,F.

TITLE Isolation, identification, and complete genome sequence of a
bovine

adenovirus type 3 from cattle in China

JOURNAL Virol. J. 8 (1), 557 (2011)

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REFERENCE 2 (bases 1 to 34132)

AUTHORS Zhu,Y.M., Yu,Z., Cai,H., Gao,Y.R., Dong,X.M., Li,Z.L., Shi,H.F.,
Meng,Q.F., Lv,C. and Xue,F.

TITLE Direct Submission

JOURNAL Submitted (20-JUL-2011) Division of Livestock Infectious Diseases
State Key Laboratory of Veterinary Biotechnology, Harbin

Veterinary

Research Institute of Chinese Academy of Agricultural Sciences,

427

Maduan, Harbin, Heilongjiang 150001, P.R. China

FEATURES Location/Qualifiers

source 1..34132

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/strain="HLJ0955"

/serotype="3"

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