Variola virus strain Iran 1972 2602 Tabriz, complete genome

GenBank: DQ437587.1

FASTA Graphics

Go to:

LOCUS DQ437587 731 bp DNA linear VRL 19-AUG-2006

DEFINITION Variola virus strain Iran 1972 2602 Tabriz, complete genome.

ACCESSION DQ437587 REGION: 135107..135837

VERSION DQ437587.1

KEYWORDS .

SOURCE Variola virus

ORGANISM Variola virus

Viruses; Varidnaviria; Bamfordvirae; Nucleocytoviricota;

Pokkesviricetes; Chitovirales; Poxviridae; Chordopoxvirinae;

Orthopoxvirus; Orthopoxvirus variola.

REFERENCE 1 (bases 1 to 731)

AUTHORS Esposito,J.J., Sammons,S.A., Frace,A.M., Osborne,J.D.,

Olsen-Rasmussen,M., Zhang,M., Govil,D., Damon,I.K., Kline,R.,

Laker,M., Li,Y., Smith,G.L., Meyer,H., Leduc,J.W. and

Wohlhueter,R.M.

TITLE Genome sequence diversity and clues to the evolution of variola

(smallpox) virus

JOURNAL Science 313 (5788), 807-812 (2006)

PUBMED 16873609

REFERENCE 2 (bases 1 to 731)

AUTHORS Sammons,S.A., Frace,A.M., Olsen-Rasmussen,M.S., Osborne,J.D.,

Damon,I.K., Wohlhueter,R.M. and Esposito,J.J.

TITLE Direct Submission

JOURNAL Submitted (08-MAR-2006) National Center for Infectious Diseases, Centers for Disease Control and Prevention, 1600 Clifton Rd, NE, Atlanta, GA 30333, USA

FEATURES Location/Qualifiers

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CDS 1..507
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CDS 549..731
/locus_tag="VARV_IRN72_tbrz_147"

/note="similar to VACV-WR-158 and VACV-COP-A35R"

/codon_start=1

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