

Molecular Epidemiology Report Form

IAH-P-EP-MEG-FOR-005-1

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Serotype: O WRL Ref No: IRN/8/2006 Sender Ref: Iran-4 Date collected: 29/04/2006 Date received by WRLFMD: 22/05/2006 Date received for sequencing: 26/05/2006 Species: Cattle Material used: BTy1 Region sequenced: VP1 RT-PCR primers: O-1C272F/EUR-2B52R; O-1C244F/EUR-2B52R No. of Nt determined: 639 No. of ambiguities: 0 Gene length: 639	Report date: 07/06/2006 Reported by: N.J. Knowles Checked by: D.P. King Topotype: ME-SA Genotype/strain: PanAsia Sequence filename: IRN06-08.SEQ Date sequence last updated: 06/06/2006 Total no. of comparisons: 1847 Min. no. of nt for comparison: 300 Total turn-around time: 16 days Sequencing time: 12 days																																																																																								
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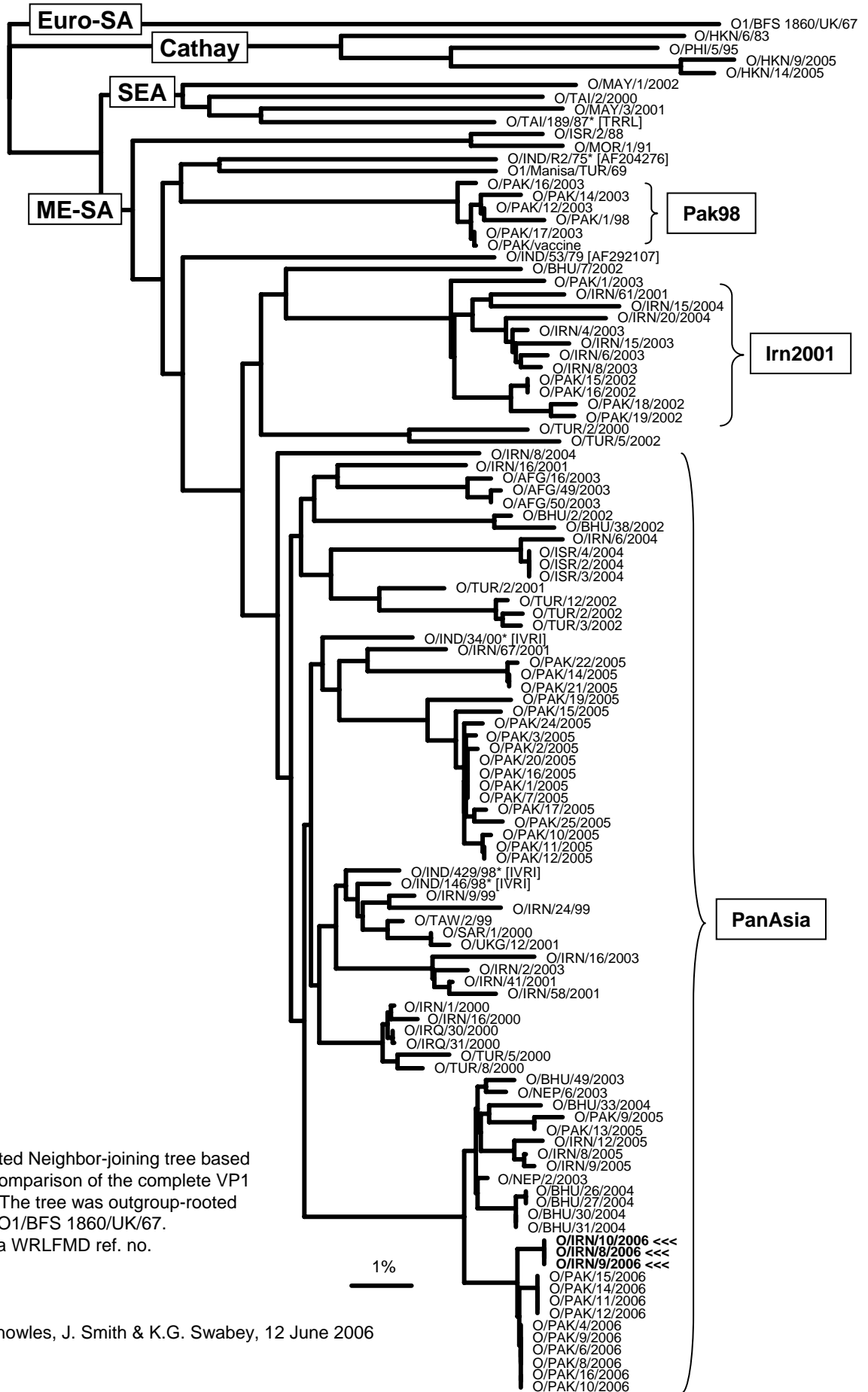
Molecular Epidemiology Report Form

IAH-P-EP-MEG-FOR-005-1

Page 1 of 1

Serotype: O WRL Ref No: IRN/10/2006 Sender Ref: Iran-1 Date collected: 09/05/2006 Date received by WRLFMD: 22/05/2006 Date received for sequencing: 26/05/2006 Species: Cattle Material used: BTy1 Region sequenced: VP1 RT-PCR primers: O-1C272F/EUR-2B52R; O-1C244F/EUR-2B52R No. of Nt determined: 639 No. of ambiguities: 0 Gene length: 639	Report date: 07/06/2006 Reported by: N.J. Knowles Checked by: D.P. King Topotype: ME-SA Genotype/strain: PanAsia Sequence filename: IRN06-10.SEQ Date sequence last updated: 07/06/2006 Total no. of comparisons: 1847 Min. no. of nt for comparison: 300 Total turn-around time: 16 days Sequencing time: 12 days																																																																																								
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Ten Most Closely Related Viruses																																																																																									
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Report on: O Iran 2006



Unrooted Neighbor-joining tree based on a comparison of the complete VP1 gene. The tree was outgroup-rooted using O1/BFS 1860/UK/67.

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1%