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CoeEFV_consensus ATGTATGGTCAAATAAATAACAATTGACTAATGAGACAGTTTGAATCCAGATAGAGAGCCCTTGCATCAATAAATACTAATCTAAT
contig270160 ATGTATGGTCAAATAAATAACAATTGACTAATGAGACAGTTTGAATCCAGATAGAGAGCCCTTGCATCAATAAATACTAATCTAAT
contig184752 ATGTATGGTCAAATAAATAACAATTGACTAATGAGACAGTTTGAATCCAGATAGAGAGCCCTTGCATCAATAAATACTAATCTAAT
contig185880 ATGTATGGTCAAATAAATAACAATTGACTAATGAGACAGTTTGAATCCAGATAGAGAGCCCTTGCATCAATAAATACTAATCTAAT
contig245863 ATGTATGGTCAAATAAATAACAATTGACTAATGAGACAGTTTGAATCCAGATAGAGAGCCCTTGCATCAATAAATACTAATCTAAT
contig236769 ATGTATGGTCAAATAAATAACAATTGACTAATGAGACAGTTTGAATCCAGATAGAGAGCCCTTGCATCAATAAATACTAATCTAAT

CoeEFV_consensus ATTCAAACAGCCCTAATCACCAATAGAGATAACAGAGCATACCAAGTGTATTGATGACCAATCCAGACCATCCACACACAGTACACAGTACCATGTCTGATATA
contig243355 ATTCAAACAGCCCTAATCACCAATAGAGATAACAGAGCATACCAAGTGTATTGATGACCAATCCAGACCATCCACACACAGTACACAGTACCATGTCTGATATA
contig219087 ATTCAAACAGCCCTAATCACCAATAGAGATAACAGAGCATACCAAGTGTATTGATGACCAATCCAGACCATCCACACACAGTACACAGTACCATGTCTGATATA

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Figure S4. Alignment of the two sets of sequences used for dating CoEFV invasion. Flanking sequences are shown for each sequence set with consensus genomic sequence of CoeEFV.