

## Erratum

### Taura syndrome virus from *Penaeus vannamei* shrimp cultured in Korea

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- In Fig. 2 the Taura syndrome virus (TSV) isolate with the GenBank Accession No. AF277675 was referred to as a French isolate. However, this TSV isolate is not a French but a Hawaiian TSV isolate identical to HI94TSV (GenBank Accession No. AF510518). We have removed the TSV isolate represented by AF277675 from Fig. 2. The corrected figure is shown here.

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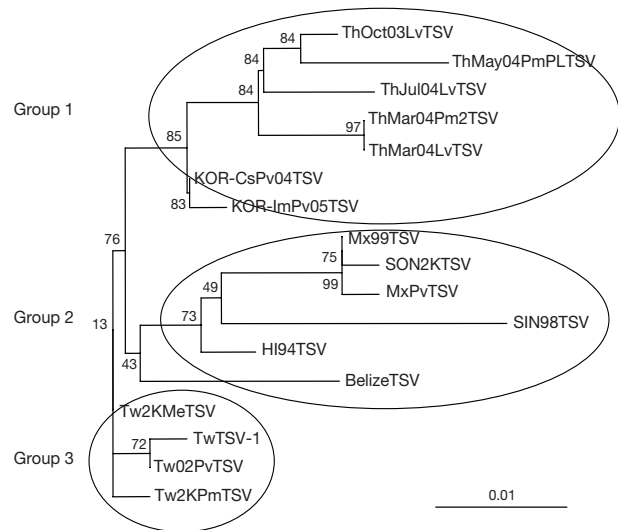


Fig. 2. Phylogenetic analysis of TSV VP1. Partial nucleotide sequences of the VP1 of 2 Korean isolates of TSV (KOR-CsPv04TSV, DQ099912; KOR-ImPv05TSV, DQ099913) were compared with those from 15 TSV strains. GenBank accession numbers for nucleotide sequences are as follows: Tw2KMeTSV (*Metapenaeus ensis*, Taiwan), AY355310; Tw02PvTSV (*Penaeus vannamei*, Taiwan), AY355311; Tw2KpMtsV (*P. monodon*, Taiwan), AY355309; ThMar04Pm2TSV (*P. monodon*, Thailand), AY912506; ThMar04LvTSV (*P. vannamei*, Thailand), AY912504; ThJul04LvTSV (*P. vannamei*, Thailand), AY912508; ThOct03LvTSV (*P. vannamei*, Thailand), AY912503; ThMay04PmPLTSV (*P. monodon*, Thailand), AY912507; HI94TSV (*P. vannamei*, Hawaii), AF510518; SIN98TSV (*P. vannamei*, Mexico), AF510515; Mx99TSV (*P. vannamei*, Mexico), AF510516; SON2KTSV (*P. vannamei*, Mexico), AF510517; TwTSV-1 (Taiwan), AF406789; BelizeTSV (Belize), AY590471; MxPvTSV (*P. vannamei*, Mexico), AF277378. The multiple alignments of the TSV VP1 nucleotide sequence were performed using ClustalW, and phylogenetic analysis was constructed using TreeView. The bootstrap values are indicated on the tree