

Figure S1. Aligned partial (64 bp) segment 10 sequences for 24 tilapia lake virus strains illustrating *in silico* specificity of the TaqMan RT-qPCR primers developed in this study.

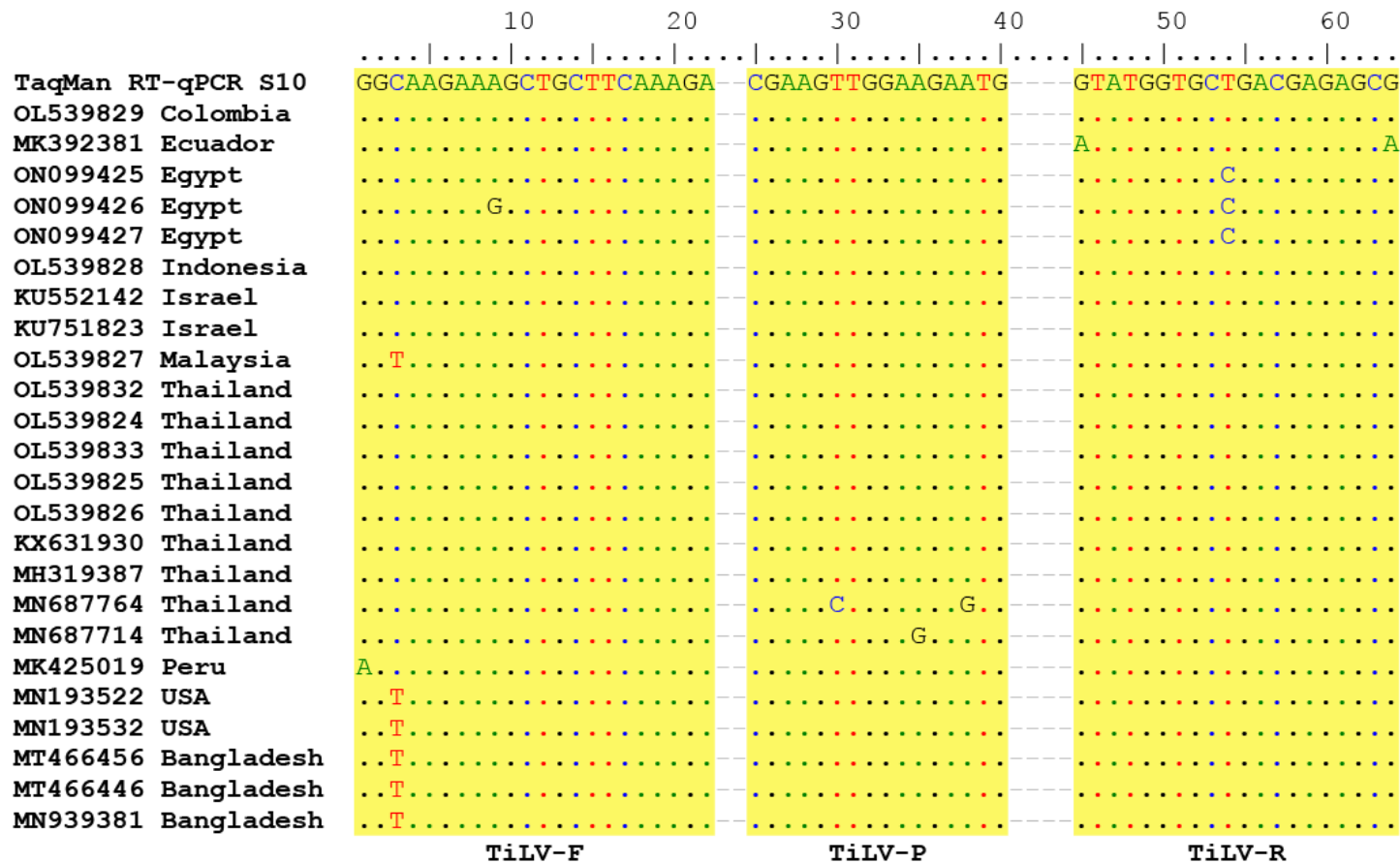


Figure S2. Aligned partial (491 bp) segment 3 sequences for 24 tilapia lake virus strains illustrating *in silico* specificity of the RT-PCR and nested RT-PCR primers developed by Eyngor et al. (2014) and Kembou Tsofack et al. (2017), respectively.

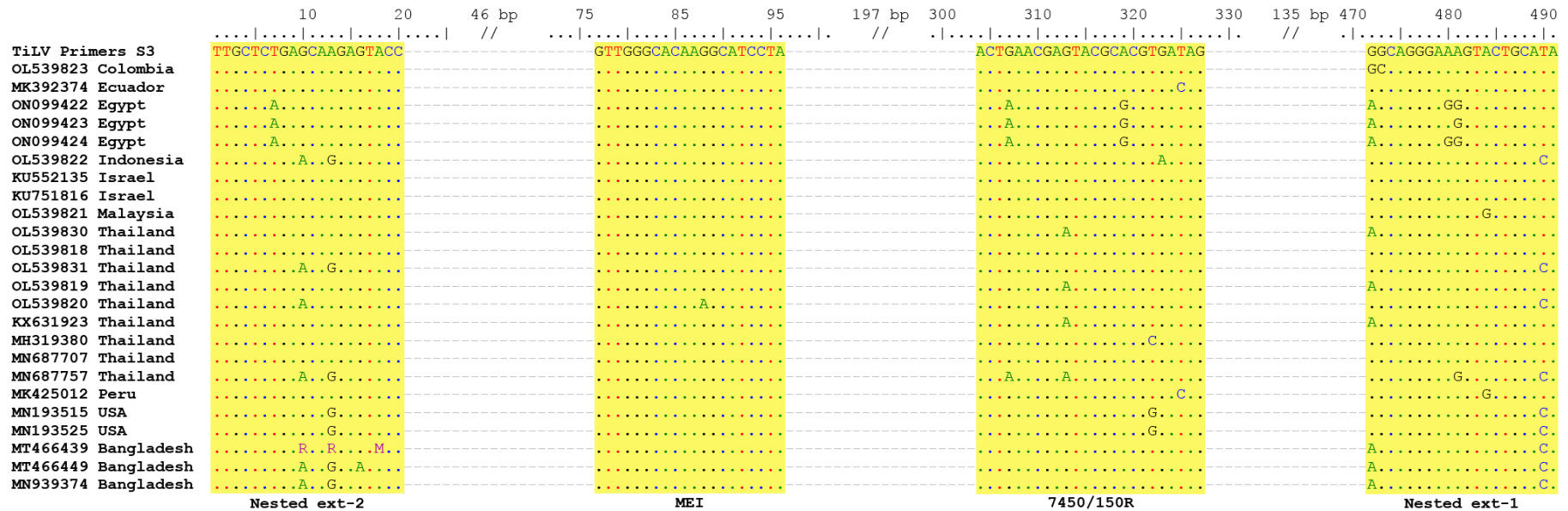
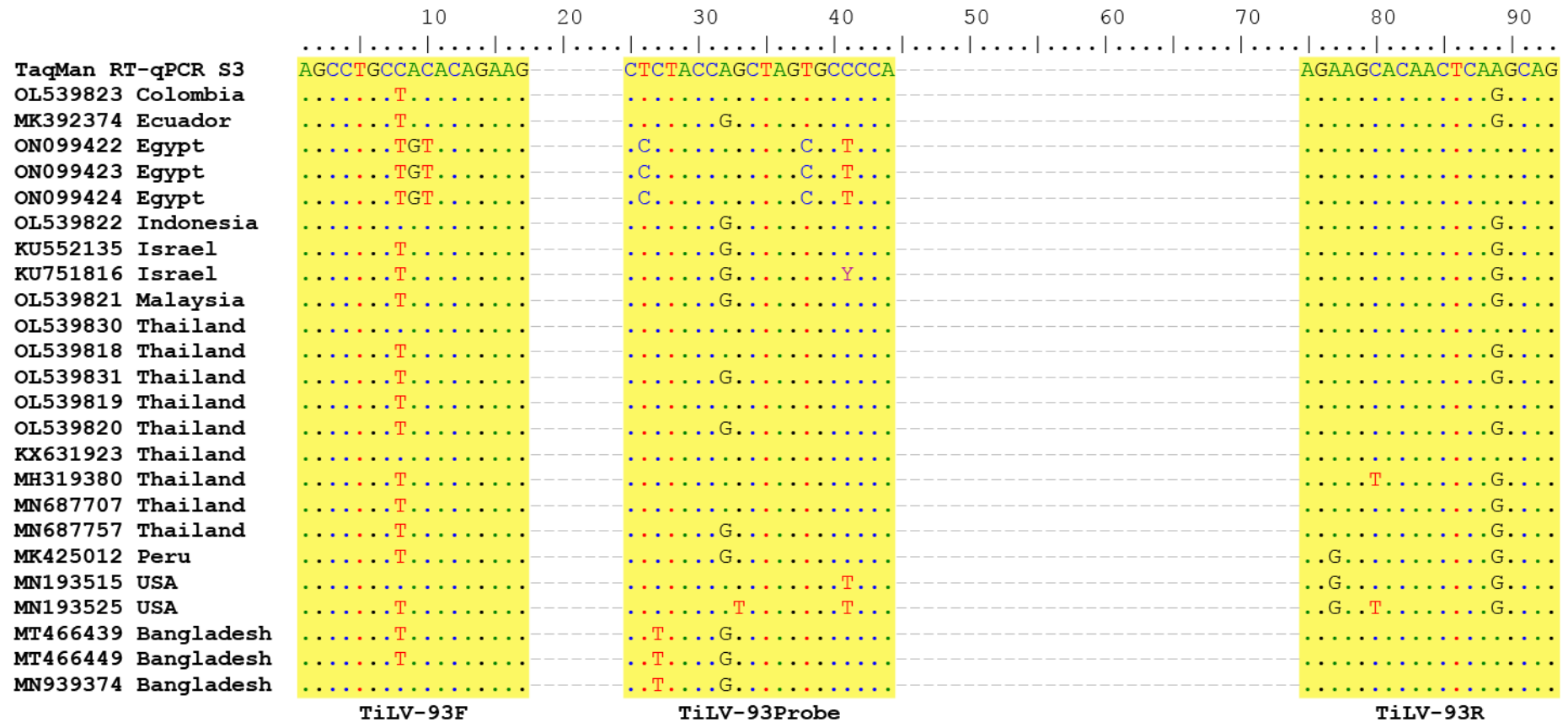


Figure S4. Aligned partial (93 bp) segment 3 sequences for 24 tilapia lake virus strains illustrating *in silico* specificity of the TaqMan RT-qPCR primers developed by Waiyamitra et al. (2018).



LITERATURE CITED

- Eyngor M, Zamostiano R, Kembou Tsofack JE, Berkowitz A and others (2014) Identification of a novel RNA virus lethal to tilapia. *J Clin Microbiol* 52: 4137–4146
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