

Aeromonas hydrophila, an Emerging Causative Agent of Freshwater-farmed Whiteleg shrimp *Litopenaeus vannamei*

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Table S1. Susceptibility of isolate WS05 to veterinary antibiotics.

Antibiotics	Content (µg/disc)	Inhibition zone diameter (mm)
Amoxicillin	10	20.3±0.5 ^S
Cotrimoxazole	23.75/1.25	23.3±0.9 ^S
Cefotaxime	30	46.7±0.5 ^S
Doxycycline*	30	23.7±0.5 ^S
Enrofloxacin*	5	34.3±0.5 ^S
Florfenicol*	30	35.0±0.8 ^S
Gentamicin	10	28.7±0.5 ^S
Kanamycin	30	25.3±0.5 ^S
Nalidixic acid	30	34.3±0.5 ^S
Neomycin*	30	23.7±0.5 ^S
Netilmicin	30	28.3±0.5 ^S
Novobiocin	30	9.0±0.8 ^R
Oxacillin	1	18.3±0.5 ^S
Polymyxin B	30	15.3±0.5 ^S
Rifampicin	5	11.0±0 ^R
Streptomycin	10	24.3±0.5 ^S
Tetracycline*	30	26.3±0.5 ^S
Tobramycin	10	27.3±0.5 ^S

Data are presented as the mean ± standard deviation of three independent experiments. ^SSensitive; ^RResistant.*Veterinary antibiotics used in aquaculture.

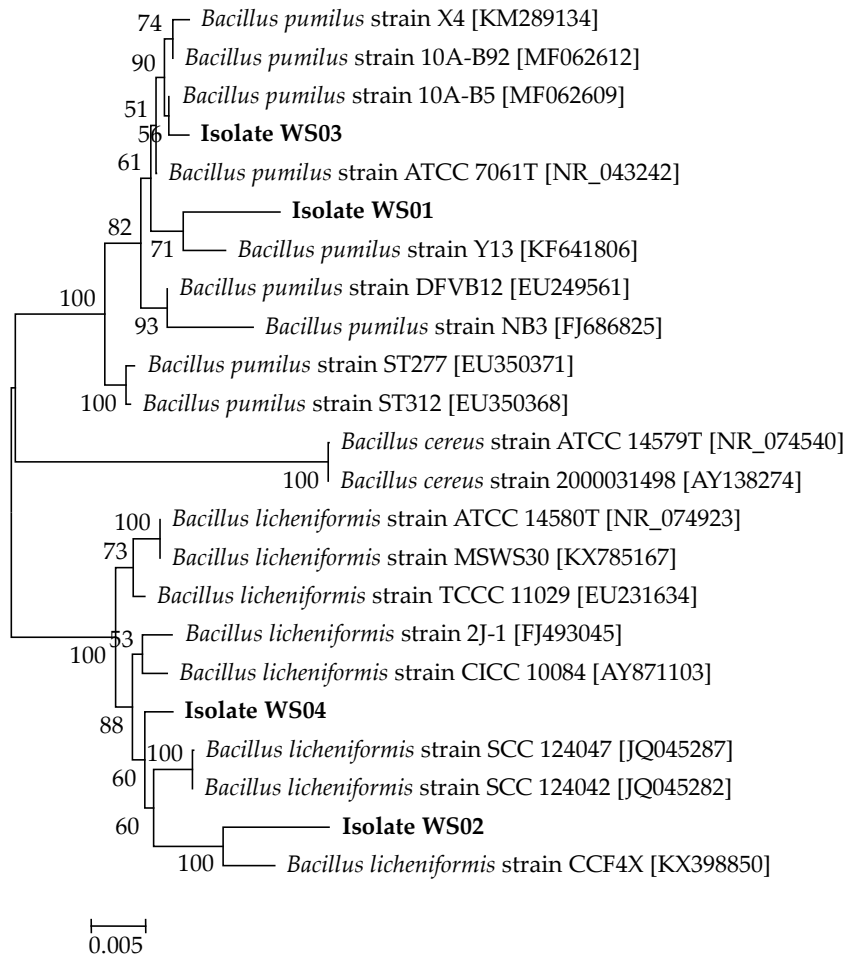


Figure S1. The 16S rRNA phylogenetic tree of 19 known bacteria and isolates WS01, WS02, WS03 and WS04 constructed using neighbor-joining method. The length of aligned sequences is 1207 bp. The bootstrap values (%) are shown besides the clades, accession numbers are indicated beside the name of strains, and scale bars represent distance values.