

## **SUPPLEMENTARY INFORMATION**

### **Phylogeny of Flavobacteria Group Isolated from Freshwater Using Multilocus Sequencing Analysis**

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**Supplementary Table 1.** Information of nucleotide sequence accession numbers

	<i>tuf</i>	<i>murG</i>	<i>gyrB</i>	<i>glyA</i>	<i>dnaK</i>	<i>atpA</i>
<i>Flavobacterium aquatile</i> JCM 20475	JX256829	JX256836	JX256843	JX256850	JX256857	JX256864
<i>F. koreense</i> ARSA-15	JX256830	JX256837	JX256844	JX256851	JX256858	JX256865
<i>F. koreense</i> ARSA-19	JX256831	JX256838	JX256845	JX256852	JX256859	JX256866
<i>F. koreense</i> ARSA-42	JX256832	JX256839	JX256846	JX256853	JX256860	JX256867
<i>F. chungnamense</i> ARSA-103	JX256833	JX256840	JX256847	JX256854	JX256861	JX256868
<i>F. chungnamense</i> ARSA-108	JX256834	JX256841	JX256848	JX256855	JX256862	JX256869
<i>F. frigidimaris</i> JCM 12218	JX256835	JX256842	JX256849	JX256856	JX256863	JX256870