

Supplementary Table S3. Summary of 16 predicted genus-specific genes of *Aeromonas*, 50 predicted horizontally transferred genes in core genome and 46 predicted virulence genes present in all 29 strains.

Gene family	Gene symbol	Genome type	Type	Category
Aeromonas_cluster634	<i>AHA_1282</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster20	<i>AHA_3819</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster956	<i>AHA_2198</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster717	<i>AHA_1552</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster160	<i>AHA_0150</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster693	<i>AHA_1510</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster354	<i>AHA_0627</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster311	<i>AHA_0510</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster281	<i>AHA_0437</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster534	<i>AHA_1006</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster690	<i>AHA_1499</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster1612	<i>AHA_4060</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster1324	<i>AHA_3339</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster356	<i>AHA_0638</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster1415	<i>AHA_3607</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster426	<i>AHA_0787</i>	Core	Aeromonas-specific	Specific for the genus <i>Aeromonas</i>
Aeromonas_cluster1058	<i>AHA_2472</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1072	<i>AHA_2580</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1084	<i>AHA_2616</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1091	<i>AHA_2629</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1096	<i>nth</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster111	<i>AHA_0046</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1215	<i>crr</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1231	<i>AHA_3118</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1234	<i>gshB</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1255	<i>rpe</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1310	<i>AHA_3322</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1313	<i>AHA_3325</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1322	<i>glyA</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1342	<i>mogA</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1419	<i>gap-2</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1457	<i>AHA_3698</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster147	<i>glyQ</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1507	<i>AHA_3830</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1551	<i>rpsI</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1552	<i>rplM</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1597	<i>rplA</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1622	<i>pntB</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1623	<i>pntA</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1642	<i>potI</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1654	<i>AHA_4189</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1656	<i>AHA_4196</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1677	<i>kbl</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1678	<i>tdh</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1693	<i>atpE</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster241	<i>rplD</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster246	<i>rpsC</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster249	<i>rpsQ</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster252	<i>rplE</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster314	<i>menB</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster32	<i>rfbB</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster341	<i>metJ</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster410	<i>xenB</i>	Core	HGT	Predicted horizontal transfer gene

Aeromonas_cluster411	<i>AHA_0738</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster412	<i>AHA_0739</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster501	<i>hfq</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster506	<i>rpmA</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster526	<i>crp</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster54	<i>zntA</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster664	<i>AHA_1404</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster732	<i>gmhA</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster794	<i>AHA_1732</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster813	<i>fdx</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster843	<i>AHA_1821</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster852	<i>lrp</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster890	<i>sucC</i>	Core	HGT	Predicted horizontal transfer gene
Aeromonas_cluster1135	<i>nueA</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1160	<i>fleR/flrC</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1161	<i>fleS/flrB</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1165	<i>flgK</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1166	<i>flgJ</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1167	<i>flgI</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1168	<i>flgH</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1169	<i>flgG</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1170	<i>flgF</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1171	<i>flgE</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1172	<i>flgD</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1173	<i>flgC</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1174	<i>flgB</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1175	<i>cheR-3</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1176	<i>cheV</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1177	<i>flgM</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1178	<i>flgN</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster1259	<i>tapP</i>	Core	Tap type IV pili	Predicted virulence gene
Aeromonas_cluster1260	<i>tapO</i>	Core	Tap type IV pili	Predicted virulence gene
Aeromonas_cluster1261	<i>tapM</i>	Core	Tap type IV pili	Predicted virulence gene
Aeromonas_cluster329	<i>exeC</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster330	<i>exeD</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster331	<i>exeE</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster332	<i>exeF</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster333	<i>exeG</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster334	<i>exeH</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster335	<i>exeI</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster336	<i>exeJ</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster337	<i>exeK</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster338	<i>exeM</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster339	<i>exeN</i>	Core	T2SS	Predicted virulence gene
Aeromonas_cluster367	<i>motX</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster519	<i>hutB</i>	Core	Iron uptake	Predicted virulence gene
Aeromonas_cluster520	<i>hutX</i>	Core	Iron uptake	Predicted virulence gene
Aeromonas_cluster521	<i>hutZ</i>	Core	Iron uptake	Predicted virulence gene
Aeromonas_cluster522	<i>ASA_3331</i>	Core	Iron uptake	Predicted virulence gene
Aeromonas_cluster523	<i>ASA_3330</i>	Core	Iron uptake	Predicted virulence gene
Aeromonas_cluster62	<i>mshJ</i>	Core	MSHA type IV pili	Predicted virulence gene
Aeromonas_cluster63	<i>mshI</i>	Core	MSHA type IV pili	Predicted virulence gene
Aeromonas_cluster653	<i>cheB-2</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster654	<i>pomB</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster655	<i>B565_1123</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster656	<i>cheW</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster782	<i>flaG</i>	Core	Polar flagella	Predicted virulence gene
Aeromonas_cluster7	<i>flaB</i>	Core	Polar flagella	Predicted virulence gene

Aeromonas_cluster818

tapF

Core

Tap type IV pili

Predicted virulence gene
