

Supplementary Information for

Deciphering a unique biotin scavenging pathway with redundant genes in the probiotic bacterium *Lactococcus lactis*

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Table S1 Comparative analyses of the genes involved in lipid metabolism of *L. lactis* and *S. suis* 2

Genes	<i>S. suis</i> 05ZYH33		<i>L. lactis</i> IL1403	
	Locus	GC %	Locus	GC %
Chromosome	/	41.1	/	35.3
<i>birA1</i>	SSU05_1625	44.0	L0191	34.2
<i>birA2</i>	/	/	L0192	34.4
<i>bioY1</i>	SSU05_0509	40.6	L24031	35.4
<i>bioY2</i>	/	/	L1011	37.5
<i>pycA</i>			L63652	38.1
<i>accA</i>	SSU05_1796	52.4	L0190	38.6
<i>accB</i>	SSU05_1801	53.1	L0187	38.7
<i>accD1</i>	SSU05_1797	51.3	L0180	37.8
<i>accD2</i>	SSU05_1798	42.9	/	/
<i>accC</i>	SSU05_1799	49.2	L0189	39.0
<i>fadA</i>	/	/	L25946	36.3
<i>fadB</i>	/	/	/	/
<i>fadL</i>	/	/	/	/
<i>fadD</i>	/	/	L54546	37.3
<i>fadE</i>	/	/	/	/
<i>fabF</i>	SSU05_1802	46.2	L0186	41.0
<i>fabG1</i>	SSU05_1803	43.4	L0185	36.9
<i>fabG2</i>	/	/	L27694	35.6
<i>fabG3</i>	/	/	L1530	41.7
<i>fabZ1</i>	SSU05_1800	45.5	L160425	34.9
<i>fabZ</i>	/	/	L0188	37.7
<i>fabI</i>	/	/	L161132	38.9
<i>fabH</i>	SSU05_1807	46.1	L0182	40.4
<i>fabD</i>	SSU05_1804	47.6	L0184	37.0
<i>fabK</i>	SSU05_1805	43.8	/	/
<i>acpP</i>	SSU05_1806	37.4	L0183	36.0
<i>acpD</i>			L115551	36.1
<i>acpH</i>	/	/	L186107	36.5
<i>fabT</i>	SSU05_1808	39.1	L180805	35.3
<i>fabM</i>	SSU05_1809	43.2	/	/
<i>acpS</i>	SSU05_1813	43.4	L61355	36.7
<i>lplA</i>	SSU05_1836	43.5	L64373	37.9