

1 Supplementary Table 1. Summary of identified suggestive QTLs for fatty acid composition

SSC	Trait	Position	F-ratio <sup>1</sup>	Inheritance of MODE <sup>2</sup>	%Var <sup>3</sup>	SI marker interval <sup>4</sup>	Reference <sup>5</sup>
1	C10:0	167	8.6 <sup>†</sup>	A	0.9	SW1957-SW1307	Ramayo-Caldas et al. (2012)
	C16:1	123	9.6 <sup>†</sup>	A	1.0	SW962-SW1957	
	C17:1	167	7.5 <sup>†</sup>	AD	1.5	SW373-SW1301	
	C18:2	111	6.5 <sup>†</sup>	AD	1.3	SW1417-SW1301	
	C20:4	167	8.4 <sup>†</sup>	AD	1.7	SW974-SW1301	
	MUFA	0	12.8 <sup>†</sup>	A	1.3	SW1514-SW1515	
	PUFA	167	6.5 <sup>†</sup>	AD	1.3	SW2035-SW1301	
2	C17:1	10	12.1 <sup>†</sup>	A	1.2	SW2623-SW776	Ramayo-Caldas et al. (2012)
	C18:3	17	8.1 <sup>†</sup>	A	0.8	SW2623-S0370	
	C20:0	105	8.2 <sup>†</sup>	A	0.8	SW2623-SW1879	
	C20:4	23	6.9 <sup>†</sup>	A	0.7	SW2623-SW1879	
	SFA	25	11.7 <sup>†</sup>	A	1.2	SW2623-S0370	
	UFA	22	10.3 <sup>†</sup>	A	1.1	SW2623-SW1879	
3	C12:0	0	13.5 <sup>†</sup>	A	1.4	APR22-SW2021	Ramayo-Caldas et al. (2012)
	C18:3	69	8.1 <sup>†</sup>	A	0.8	APR22-SW1327	
4	C16:1	88	6.1 <sup>†</sup>	AD	1.3	SW489-MP77	Pérez-Enciso et al. (2000) and Ramayo-Caldas et al. (2012)
	C18:0	93	7.4 <sup>†</sup>	AD	1.5	SW1364-MP77	
	C18:1	44	7.7 <sup>†</sup>	A	0.8	SW489-MP77	
	C18:2	44	6.9 <sup>†</sup>	A	0.7	SW489-MP77	
5	MUFA	42	6.9 <sup>†</sup>	A	0.7	SW489-MP77	Pérez-Enciso et al. (2000), Nii et al. (2006), Guo et al. (2009), Uemoto et al. (2009), Ramayo-Caldas et al. (2012) and Uemoto et al. (2012)
	PUFA	79	12.2 <sup>†</sup>	A	1.3	APR8-SW122	
5	C12:0	101	8.4 <sup>†</sup>	A	0.9	SW2003-SW967	Uemoto et al. (2009)
	C18:2	123	7.1 <sup>†</sup>	A	0.7	SW2003-SW967	
	C20:0	122	8.8 <sup>†</sup>	A	0.9	SW963-SW967	
6	C12:0	14	7.9 <sup>†</sup>	AD	1.6	S0035-SW1057	Lee et al. (2010) and Ramayo-Caldas et al. (2012)
	C17:1	79	9.1 <sup>†</sup>	A	0.9	SW2406-S0059	
	C18:1	79	9.6 <sup>†</sup>	A	1.0	APR8-SW1055	
	C18:2	79	12.2 <sup>†</sup>	A	1.3	APR8-SW122	
	MUFA	79	8.3 <sup>†</sup>	A	0.9	APR8-SW1055	
6	PUFA	79	12.2 <sup>†</sup>	A	1.3	APR8-S0059	Lee et al. (2010)
	C14:0	134	8.8 <sup>†</sup>	A	0.9	SW147-SW764	Guo et al. (2009)
	C18:3	94	10.5 <sup>†</sup>	A	0.7	SW1369-SW764	
PUFA	62	12.8 <sup>†</sup>	A	1.3	SW1369-SW2108		
8	C12:0	116	9.9 <sup>†</sup>	A	1.0	SW1345-KS188	Uemoto et al. (2012)
	MUFA	4	7.4 <sup>†</sup>	A	0.8	SW2410-SW444	
9	C10:0	145	6.0 <sup>†</sup>	AD	1.2	SW2093-SW749	Uemoto et al. (2012)
	C16:1	145	7.7 <sup>†</sup>	AD	1.6	SW2093-SW749	
10	C16:0	54	9.1 <sup>†</sup>	A	1.0	SWR136-SW2000	Uemoto et al. (2012)
11	C12:0	47	5.3 <sup>†</sup>	AD	1.1	SW1460-SW1135	

	C18:1	61	6.5 <sup>†</sup>	A	0.7	SW1460-SW1135	
	MUFA	61	7.2 <sup>†</sup>	A	0.7	SW1460-SW1135	
12	C16:1	115	12.9 <sup>†</sup>	A	1.3	SO106-SWR1021	
13	C10:0	62	9.3 <sup>†</sup>	A	1.0	S0288-SW38	
	C16:1	74	12.5 <sup>†</sup>	A	1.3	SW1407-SW38	Guo et al. (2009)
	C18:2	29	9.4 <sup>†</sup>	A	1.0	SWR1941-SW38	
	C20:0	24	6.4 <sup>†</sup>	AD	1.3	SWR1941-S0215	
	C20:1	29	10.5 <sup>†</sup>	A	1.1	SW1378-S0283	
	PUFA	29	8.0 <sup>†</sup>	A	0.8	SWR1941-SW38	
14	C10:0	64	10.0 <sup>†</sup>	A	1.0	SW2038-SW2515	
	C17:0	58	8.7 <sup>†</sup>	A	0.9	SW2038-SW2515	
	C20:4	14	6.9 <sup>†</sup>	A	0.7	SW857-SW2515	
15	C20:0	78	9.1 <sup>†</sup>	A	0.9	SW1989-S0040	
	C20:1	130	6.7 <sup>†</sup>	AD	1.4	SW936-SWR2121	
	MUFA	141	6.0 <sup>†</sup>	AD	1.2	KS135-SWR2121	
16	C12:0	43	5.9 <sup>†</sup>	AD	1.2	SW419-SW2517	
	C17:0	59	6.5 <sup>†</sup>	AD	1.3	SW1809-S0105	
	C17:1	54	8.0 <sup>†</sup>	AD	1.6	SW419-SW2517	
	C18:1	55	7.3 <sup>†</sup>	AD	1.5	SW1035-SW1897	
	C18:2	53	5.0 <sup>†</sup>	AD	1.0	SW1035-SW1897	
	C20:4	41	7.1 <sup>†</sup>	AD	1.5	SW419-SW2517	
	MUFA	56	6.8 <sup>†</sup>	AD	1.4	SW1035-SW1897	
	PUFA	52	5.6 <sup>†</sup>	AD	1.2	SW1035-SW1897	
17	C10:0	11	11.1 <sup>†</sup>	A	1.1	SW1031-SW2142	
18	C10:0	22	6.7 <sup>†</sup>	AD	1.4	SY4-S0062	
	C16:1	24	10.2 <sup>†</sup>	A	1.1	SY4-SB58	

1 <sup>1</sup>Test statistic and level of significance: <sup>†</sup>suggestive 5% significance.

2 <sup>2</sup>A represents additive effect; AD represents additive and dominance effects.

3 <sup>3</sup>Flanking markers for 95% support intervals estimated by the 1.5-LOD drop method.

4 <sup>4</sup>Var % is the reduction in residual variance of the F<sub>2</sub> population obtained by inclusion of a  
5 QTL at the given position.

6 <sup>5</sup>Papers reporting QTL with similar effects at comparable positions in PigQTLdb.

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