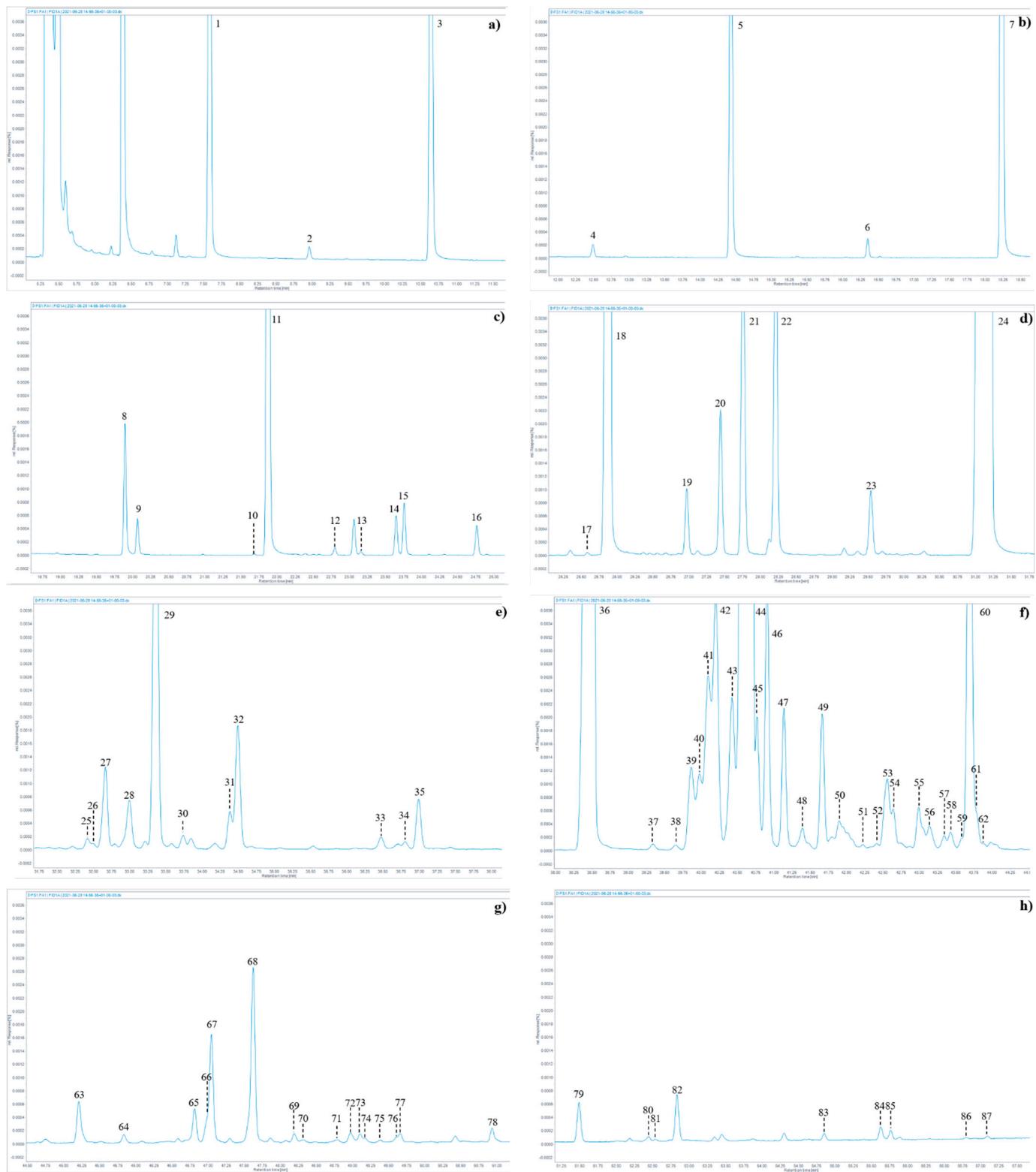
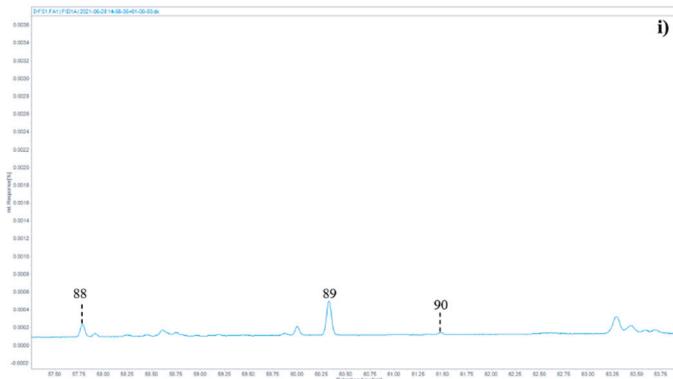


## Supplementary Data





**Figure S1:** Gas chromatogram illustrating the separation of fatty acid methyl esters in milk from the bulk tanks of 67 farms across Southern England over 12 months. The left y-axis represents arbitrary response and the x-axis represents the retention time from 5 to 70 min. Chromatographic conditions are presented in the Materials and Methods. The following peaks were identified: 1=C4:0, 2=C5:0, 3=C6:0, 4=C7:0, 5=C8:0, 6=C9:0, 7=C10:0, 8=C10:1 c9, 9=C11:0, 10=C11:1 c10 11=C12:0, 12= C13:0 iso, 13=C13:0 anteiso, 14=C12:1 c9, 15=C13:0, 16=C14:0 iso, 17=C13:1, 18=C14:0, 19=C14:1 t9, 20=C15:0, 21=C15:1 c9, 22=C15:0 anteiso, 23=C16:0 iso, 24=C16:0, 25=C16:1 t6-8, 26=C16:1 t9, 27=C17:0 iso, 28=C16:1 t11/13, 29=C16:1 + C17:0 anteiso, 30=C16:1 c11, 31=C16:1 c13, 32=C17:0, 33=C17:1 t10, 34=C18:0 iso, 35=C17: c9, 36=C18:0, 37=C18:1 t4, 38=C18:1 t5, 39=C18:1 t6-8, 40=C18:1 t9, 41=C18:1 t10, 42=C18:1 t11, 43=C18:1 c6/t12, 44=C18:1 c9, 45=C18:1 t15, 46=C18:1 c11, 47=C18:1 c12, 48=C18:1 c13, 49=C18:1 t6 + C18:1 c14, 50=C19:0 + C18:1 c15, 51=C18:2 t11t15, 52=C18:2 t9t12, 53=C18:2 c8c13, 54=C18:2 c10t14, 55=C18:2 c9c14, 56=C18:2 c9t12, 57=C18:1 c16, 58=C18:1 t11c15, 59=C18:2 t9c12, 60=C18:2 c9c12, 61=C18:2 t12 c15, 62=C19:1, 63=C20:0, 64=C18:3 c6 c9 c12, 65=C20:1 c8, 66=C20:1 c11, 67=C18:3 c9 c12 c15, 68=CLA c9 t11+t7c9 +t8c10+t6c8, 69-76=other CLA of unknown isomerism, 77=C20:2 c11c14, 78=C22:0, 79=C20:3 c8c11c14, 80=C22:1 c13, 81=C20:3 c11c14c17, 82=C20:4 c5c8c11c14, 83=C22:2 c13c16, 84=C20:5 c5c8c11c14c17 (EPA), 85=C24:0, 86=C24:1 c15, 87=C22:3 c13c16c19, 88=C22:4 c7c10c13c16, 89=C22:5 c7c10c13c16c19 (DPA), 90=C22:6 c4c7c10c13c16c19 (DHA).

**Table S1.** Mean, minimum and maximum values for the breed and diet composition of herds (%) from 67 farms in Southern England.

	Mean n=67	Min n=67	Max n=67	SE
Milking herd size (number of cows)	247	75	670	4.0
Milking cows (% herd)	86	30	100	0.4
Estimated Liveweight (kg) <sup>1</sup>	649	468	680	1.7
Holstein	74.8	0	100	1.36
British Friesian	3.28	0	100	0.493
NZ Friesian	0.56	0	100	0.257
Jersey	0.53	0	16.3	0.079
Scandinavian Red	0.42	0	41.0	0.070
Shorthorn	0.30	0	6.67	0.036
Ayrshire	4.76	0	100	0.723
Montbeliarde	0.06	0	6.45	0.017
Brown Swiss	0.95	0	51.4	0.192
Guernsey	0.03	0	1.57	0.007
Other breed or crossbreed	14.3	0	100	1.19
Offered feed (kg DM / day) <sup>2</sup>	20.2	6.80	31.7	0.09
Total Forage	66.2	39.4	95.6	0.40
Total concentrate	33.9	4.37	60.6	0.40
Predicted grazing intake	14.8	0	91.4	0.84
Total Silage intake	49.3	0	114	0.69
Grass silage	26.9	0	81.9	0.62
Grass:clover silage <sup>4</sup>	1.77	0	67.3	0.246
Maize silage	15.6	0	54.2	0.54
Cereal silage	0.15	0	35.9	0.066
Lucerne silage	12.5	0.25	0	0.05
Other mixed silage	53.7	1.12	0	0.17
Wholecrop	49.8	3.49	0	0.28
Hay and Straw	30.4	2.26	0	0.12
Moist by-products	25.6	3.03	0	0.19
Dry straights <sup>3</sup>	44.4	6.77	0	0.40
Cereals	21.0	3.06	0	0.18
Blends	59.1	20.5	0	0.45
Oil	4.03	0.51	0	0.029
Minerals (g/cow per day)	750	110	0	3.9
Vitamins (g/cow per day)	320	6.59	0	1.4

<sup>1</sup>Average herd live weights were estimated based on average breed live weights and the proportionate number of cows from each breed or crossbreed in the total herd, as described by Stergiadis, et al. [1].

<sup>2</sup>When cows had no access to pasture, this value reflects the summary of individual feeds DM as recorded in farmers' questionnaires. During the grazing season, this is predicted using equations published by [2].

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<sup>3</sup>Dry straights: single feeding stuffs from which compound feeds and protein concentrates are prepared (wheat, barley, flaked maize, field beans, groundnut cake and meal, soya bean meal).

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Cereal silage	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	2.00	1.76	0.56	0.271	0.086
Lucerne silage	0.33 <sup>abcd</sup>	0.27 <sup>abcd</sup>	0.24 <sup>cd</sup>	0.25 <sup>abcd</sup>	0.20 <sup>a</sup>	0.17 <sup>d</sup>	0.17 <sup>d</sup>	0.17 <sup>d</sup>	0.17 <sup>d</sup>	0.19 <sup>d</sup>	0.26 <sup>bcd</sup>	0.60 <sup>ac</sup>	0.161	0.027
Other mixed silage	0.89	1.26	1.14	1.12	0.89	1.14	0.77	0.47	1.10	0.59	1.48	2.66	0.542	0.200
Wholecrop	5.28 <sup>dcba</sup>	4.53 <sup>cd</sup>	4.39 <sup>cd</sup>	2.10 <sup>fg</sup>	1.12 <sup>gh</sup>	1.14 <sup>gh</sup>	0.43 <sup>h</sup>	2.56 <sup>fg</sup>	3.40 <sup>df</sup>	6.28 <sup>a</sup>	5.85 <sup>ac</sup>	4.84 <sup>bcd</sup>	0.895	<0.001
Hay and Straw	3.37 <sup>a</sup>	3.39 <sup>a</sup>	2.21 <sup>b</sup>	2.01 <sup>b</sup>	2.04 <sup>b</sup>	1.99 <sup>b</sup>	2.20 <sup>b</sup>	1.94 <sup>b</sup>	1.93 <sup>b</sup>	2.38 <sup>b</sup>	1.75 <sup>b</sup>	1.95 <sup>b</sup>	0.387	0.035
Moist by-products	3.02	3.13	3.32	2.91	2.92	3.16	3.25	2.98	3.18	2.8	3.09	2.62	0.654	0.359
Dry straights <sup>6</sup>	8.31	8.03	7.37	6.77	6.31	6.60	6.04	5.81	6.52	6.92	6.50	6.13	1.384	0.608
Cereals	3.67 <sup>abcd</sup>	3.36 <sup>abcd</sup>	3.56 <sup>ac</sup>	2.68 <sup>bcd</sup>	2.83 <sup>abcd</sup>	2.30 <sup>d</sup>	2.80 <sup>abcd</sup>	3.17 <sup>abcd</sup>	3.06 <sup>abcd</sup>	3.11 <sup>abcd</sup>	2.33 <sup>cd</sup>	3.90 <sup>ab</sup>	0.609	0.029
Blends	19.9	20.5	20.7	20.8	20.7	20.5	19.5	19.3	20.6	20.6	21.8	21.0	1.55	0.456
Oil	0.49	0.50	0.48	0.53	0.48	0.50	0.53	0.53	0.54	0.50	0.50	0.56	0.102	0.830
Minerals (g/cow per day)	112	116	111	113	100	95.4	98.6	101	106	120	118	123	13.3	0.236
Vitamins (g/cow per day)	8.80 <sup>abc</sup>	8.60 <sup>abc</sup>	8.60 <sup>abc</sup>	7.56 <sup>abc</sup>	7.39 <sup>abc</sup>	7.43 <sup>abc</sup>	6.57 <sup>abc</sup>	6.57 <sup>abc</sup>	6.57 <sup>ba</sup>	0.00 <sup>c</sup>	0.00 <sup>cb</sup>	11.2 <sup>a</sup>	4.40	0.001

<sup>1</sup>n is the number of records used to calculate means ± SE and P-values. Records with missing values were not included in the analysis.

<sup>2</sup>Significances were declared at P-value <0.05.

<sup>3</sup>Average herd live weights were estimated based on average breed live weights and the proportionate number of cows from each breed or crossbreed in the total herd, as described by Stergiadis, et al. [1].

<sup>4</sup>When cows had no access to pasture, this value reflects the summary of individual feeds DM as recorded in farmers' questionnaires. During the grazing season, this is predicted using equations published by Butler et al. (2008)

<sup>5</sup>In the conventional production system, this was predominantly perennial ryegrass silage, while in organic systems silage typically has variant grass:clover ratios.

<sup>6</sup>Dry straights: single feeding stuffs which from which compound feeds and protein concentrates are prepared (wheat, barley, flaked maize, field beans, groundnut cake and meal, soya bean meal).



	170 <sup>bc</sup>	168 <sup>bc</sup>	160 <sup>c</sup>	187 <sup>ab</sup>	186 <sup>b</sup>	185 <sup>b</sup>	179 <sup>bc</sup>	212 <sup>a</sup>	174.1 <sup>bc</sup>	176 <sup>bc</sup>	175 <sup>bc</sup>	174 <sup>bc</sup>	10.4	0.027
Fat concentrate efficiency (g fat/kg concentrate DM offered)	170 <sup>bc</sup>	168 <sup>bc</sup>	160 <sup>c</sup>	187 <sup>ab</sup>	186 <sup>b</sup>	185 <sup>b</sup>	179 <sup>bc</sup>	212 <sup>a</sup>	174.1 <sup>bc</sup>	176 <sup>bc</sup>	175 <sup>bc</sup>	174 <sup>bc</sup>	10.4	0.027
Protein efficiency (g protein/kg DM offered)	43.6 <sup>fgh</sup>	43.4 <sup>h</sup>	42.0 <sup>i</sup>	44.7 <sup>cdefg</sup>	45.1 <sup>dfcba</sup>	43.7 <sup>efgh</sup>	42.9 <sup>hi</sup>	43.2 <sup>ghi</sup>	43.5 <sup>fgh</sup>	43.8 <sup>defgh</sup>	46.8 <sup>ca</sup>	44.6 <sup>bcd</sup> <sup>efgh</sup>	0.94	<0.001
Protein non-grazing efficiency (g protein/kg non-grazing DM offered)	43.6 <sup>e</sup>	43.4 <sup>e</sup>	45.8 <sup>e</sup>	79.4 <sup>b</sup>	89.6 <sup>a</sup>	90.5 <sup>a</sup>	81.1 <sup>b</sup>	67.4 <sup>c</sup>	58.3 <sup>d</sup>	50.6 <sup>de</sup>	46.8 <sup>e</sup>	44.6 <sup>e</sup>	4.16	<0.001
Protein concentrate efficiency (g protein/kg concentrate DM offered)	140 <sup>bc</sup>	143 <sup>bc</sup>	133 <sup>c</sup>	160 <sup>ab</sup>	162 <sup>ab</sup>	158 <sup>b</sup>	158 <sup>b</sup>	180 <sup>a</sup>	145 <sup>bc</sup>	147 <sup>bc</sup>	146 <sup>bc</sup>	145 <sup>bc</sup>	8.9	0.014
Health parameters and indicators														
Mastitis (% of herd)	2.99 <sup>ab</sup>	2.17 <sup>cde</sup>	2.53 <sup>abc</sup>	1.92 <sup>de</sup>	1.81 <sup>e</sup>	2.30 <sup>cd</sup>	2.23 <sup>cde</sup>	2.89 <sup>ab</sup>	2.55 <sup>bcd</sup>	2.69 <sup>abc</sup>	3.07 <sup>a</sup>	2.24 <sup>cde</sup>	0.216	<0.001
Lameness (% of herd)	2.58	2.23	2.02	2.52	2.19	2.11	2.57	2.36	2.38	2.46	2.05	1.96	0.296	0.803
Other disease (% of herd)	0.61 <sup>bcd</sup>	0.56 <sup>cd</sup>	0.57 <sup>cd</sup>	0.55 <sup>cd</sup>	0.63 <sup>bcd</sup>	0.63 <sup>bcd</sup>	0.83 <sup>bcd</sup>	1.16 <sup>d</sup>	0.80 <sup>bcd</sup>	0.94 <sup>ab</sup>	0.81 <sup>abc</sup>	0.38 <sup>d</sup>	0.132	0.032
Fat:protein	1.21 <sup>a</sup>	1.18 <sup>bc</sup>	1.21 <sup>a</sup>	1.17 <sup>de</sup>	1.15 <sup>ef</sup>	1.17 <sup>dc</sup>	1.14 <sup>f</sup>	1.19 <sup>ba</sup>	1.20 <sup>ab</sup>	1.20 <sup>ab</sup>	1.19 <sup>ab</sup>	1.19 <sup>ab</sup>	0.009	<0.001
Milk SCC (x1000/ml milk)	149 <sup>cd</sup>	129 <sup>ef</sup>	156 <sup>abcd</sup>	151 <sup>cd</sup>	141 <sup>de</sup>	165 <sup>ac</sup>	149 <sup>bcd</sup>	160 <sup>abc</sup>	114 <sup>f</sup>	143 <sup>de</sup>	142 <sup>de</sup>	138 <sup>de</sup>	6.7	<0.001

<sup>1</sup>n is the number of records used to calculate means ± SE and P-values. Records with missing values were not included in the analysis.

<sup>2</sup>Significances were declared at P-value <0.05.

**Table S4.** Means  $\pm$  SE and P-values for the effect of production system (conventional, 41; Organic, 26) on the fatty acid profile of milk collected from 67 Farms across southern England

	Production System			
			Organic	P-value <sup>2</sup>
	Conventional	n <sup>1</sup> =485		
C4:0		26.2	26.8	0.08 <0.001
C5:0		0.26	0.24	0.003 <0.001
C6:0		18.5	19.3	0.06 <0.001
C7:0		0.25	0.23	0.004 <0.001
C8:0		11.6	12.0	0.05 0.029
C9:0		0.32	0.29	0.005 <0.001
C10:0		27.1	27.9	0.17 0.292
C10:1 <i>c</i> 9		2.64	2.85	0.020 <0.001
C11:0		0.59	0.54	0.010 0.004
C11:1 <i>c</i> 10		0.02	0.02	0.0004 0.056
C12:0		34.0	34.5	0.24 0.968
C13:0 iso		0.22	0.33	0.004 <0.001
C13:0 anteiso		0.09	0.11	0.001 <0.001
C12:1 <i>c</i> 9		0.86	0.87	0.008 0.841
C13:0		0.95	0.97	0.011 0.661
c14:0 iso		0.75	1.08	0.009 <0.001
C13:1		0.05	0.05	0.001 0.942
C14:0		106	113	0.4 <0.001
C14:1 <i>t</i> 9		1.97	2.65	0.020 <0.001
C15:0		4.15	4.87	0.0350 <0.001
C15:1 <i>c</i> 9		9.45	9.57	0.063 0.434
C15:0 anteiso		10.2	11.9	0.08 <0.001
C16:0 iso		1.88	2.30	0.0170 <0.001
C16:0		314	309	1.5 0.905
C16:1 <i>t</i> 6-8		0.31	0.27	0.002 <0.001
C16:1 <i>t</i> 9		0.09	0.06	0.002 <0.001
C17:0 iso		3.47	4.05	0.036 <0.001
C16:1 <i>t</i> 11/13		1.99	1.92	0.017 0.002
C16:1 + C17:0 anteiso		18.7	17.3	0.09 <0.001
C16:1 <i>c</i> 11		0.93	1.17	0.018 <0.001
C16:1 <i>c</i> 13		1.17	1.23	0.014 0.199
C17:0		4.90	5.96	0.028 <0.001
C17:1 <i>f</i> 10		0.52	0.54	0.004 0.018
C18:0 iso		0.41	0.52	0.008 <0.001
C17:0 <i>c</i> 9		1.97	2.28	0.014 <0.001
C18:0		99.2	102	0.59 0.080
C18:1 <i>t</i> 4		0.20	0.13	0.003 <0.001
C18:1 <i>t</i> 5		0.17	0.11	0.002 <0.001

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C18:1 <i>t</i> 6-8	3.37	2.37	0.033	<0.001
C18:1 <i>t</i> 9	2.56	1.80	0.025	<0.001
C18:1 <i>t</i> 10	5.79	3.93	0.105	<0.001
C18:1 <i>t</i> 11	11.9	16.8	0.30	<0.001
C18:1 <i>c</i> 6/ <i>t</i> 12	4.90	3.38	0.043	<0.001
C18:1 <i>c</i> 9	198	185	0.9	<0.001
C18:1 <i>t</i> 15	3.58	3.10	0.039	<0.001
C18:1 <i>c</i> 11	6.87	4.39	0.057	<0.001
C18:1 <i>c</i> 12	2.66	1.77	0.040	<0.001
C18:1 <i>c</i> 13	0.73	0.57	0.009	<0.001
C18:1 <i>t</i> 6 + C18:1 <i>c</i> 14	3.60	3.23	0.026	<0.001
C19:0 + C18:1 <i>c</i> 15	1.56	1.49	0.014	0.364
C18:2 <i>t</i> 11 <i>t</i> 15	0.19	0.38	0.008	<0.001
C18:2 <i>t</i> 9 <i>t</i> 12	0.05	0.03	0.001	<0.001
C18:2 <i>c</i> 8 <i>c</i> 13	2.39	1.86	0.024	<0.001
C18:2 <i>c</i> 10 <i>t</i> 14	1.11	1.12	0.014	0.727
C18:2 <i>c</i> 9 <i>c</i> 14	1.40	1.07	0.010	<0.001
C18:2 <i>c</i> 9 <i>t</i> 12	0.68	0.54	0.005	<0.001
C18:1 <i>c</i> 16	0.31	0.30	0.006	0.408
C18:1 <i>t</i> 11 <i>c</i> 15	1.32	2.16	0.042	<0.001
C18:2 <i>t</i> 9 <i>c</i> 12	0.13	0.10	0.004	0.211
C18:2 <i>c</i> 9 <i>c</i> 12	19.1	16.0	0.21	<0.001
C18:2 <i>t</i> 12 <i>c</i> 15	0.88	0.97	0.014	<0.001
C19:1	0.17	0.22	0.003	<0.001
C20:0	1.49	1.87	0.012	<0.001
C18:3 <i>c</i> 6 <i>c</i> 9 <i>c</i> 12	0.21	0.19	0.002	0.031
C20:1 <i>c</i> 8	1.05	1.25	0.009	<0.001
C20:1 <i>c</i> 11	0.54	0.56	0.013	0.076
C18:3 <i>c</i> 9 <i>c</i> 12 <i>c</i> 15	4.53	6.94	0.063	<0.001
CLA <i>c</i> 9 <i>t</i> 11 + <i>t</i> 7 <i>c</i> 9 + <i>t</i> 8 <i>c</i> 10 + <i>t</i> 6 <i>t</i> 8	5.94	8.00	0.123	<0.001
CLA other a	0.22	0.30	0.002	<0.001
CLA other b	0.17	0.40	0.008	<0.001
CLA other c	0.14	0.17	0.002	<0.001
CLA other d	0.41	0.38	0.006	0.031
CLA other e	0.22	0.20	0.001	<0.001
CLA other f	0.05	0.04	0.001	<0.001
CLA other g	0.12	0.21	0.004	<0.001
CLA other h	0.11	0.16	0.002	<0.001
C20:2 <i>c</i> 11 <i>c</i> 14	0.20	0.20	0.002	0.311
C22:0	0.51	0.85	0.007	<0.001
C20:3 <i>c</i> 8 <i>c</i> 11 <i>c</i> 14	0.87	0.71	0.007	<0.001
C22:1 <i>c</i> 13	0.16	0.55	0.023	<0.001
C20:3 <i>c</i> 11 <i>c</i> 14 <i>c</i> 17	0.07	0.13	0.002	<0.001

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C20:4 <i>c5c8c11c14</i>	1.11	0.96	0.009	<0.001
C22:2 <i>c13c16</i>	0.34	0.44	0.006	<0.001
C20:5 <i>c5c8c11c14c17</i> (EPA)	0.44	0.66	0.006	<0.001
C24:0	0.35	0.52	0.004	<0.001
C24:1 <i>c15</i>	0.06	0.10	0.002	<0.001
C22:3 <i>c13c16c19</i>	0.08	0.12	0.003	<0.001
C22:4 <i>c7c10c13c16</i>	0.05	0.07	0.001	<0.001
C22:5 <i>c7c10c13c16c19</i> (DPA)	0.74	1.05	0.007	<0.001
C22:6 <i>c4 c7c10c13c16c19</i> (DHA)	0.05	0.07	0.001	<0.001

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<sup>1</sup>n is the number of records used to calculate means. Records with missing data were not included in the means ± SE and P-values calculations.

<sup>2</sup>Significances were declared at P<0.05

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<sup>t6</sup>, C18:1 c15 (co-elutes with C19:0), C18:1 c16, C19:1, C20:1 c8, C20:1 c11, C22:1 c13, C24:1 c15.

<sup>5</sup>PUFA: C18:2 t11t15, C18:2 t9t12, C18:2 c9t13, C18:2 c10t14, C18:2 c9t14, C18:2 t11c15, C18:2 t9c12, C18:2 c9c12 (LA), C18:2 t12c15, C18:3 c6c9c12, C18:3 c9c12c15 (ALNA), 18:2 c9t11 conjugated (RA) (co-elutes with C18:2 t7c9 + t8c10 + t6c8), other conjugated FA of unknown isomerisms of C18:2, C20:2 c11c14, C20:3 c11c14c17, C20:4 c5c8c11c14, C22:2 c13c16, C20:5 c5c8c11c14c17 (EPA), C22:3 c13c16c19, C22:4 c7c10c16c19, C22:5 c7c10c13c16c19 (DPA), C22:6 c4c7c10c13c16c19 (DHA).

<sup>6</sup>cisMUFA: C10:1 c9, C11:1 c10, C12:1 c9, C13:1 c9, C14:1 c9, C16:2 c9 (co-elutes with C17:0 anteiso), C16:1 c11, C16:1 c13, C17:1 c9, C18:1 c6 (co-elutes with C18:1 t16), C18:1 c9 (OA), C18:1 c11, C18:1 c12, C18:1 c13, C18:1 c14 (co-elutes with C18:1 t6), C18:1 c15 (co-elutes with C19:0), C18:1 c16, c19:1 c9, C20:1 c8, C20:1 c11, C22:1 c13, C24:1 c15.

<sup>7</sup>transMUFA: C14:1 t9, C16:1 t6+t7+t8, C16:1 t9, C16:1 t11+t12+t13, C17:1 t10, C18:1 t4, C18:1 t5, C18:1 t6+t7+t8, C18:1 t9, C18:1 t10, C18:1 t11 (VA), C18:1 t12 (co-elutes with C18:1 c6), C18:1 t15, C18:1 t16 (co-elutes with C18:1 c14).

<sup>8</sup>cisPUFA: C18:2 c9c12 (LA), C18:3 c6c9c12, C18:3 c9c12c15 (ALNA), 20:2 c11c14, C20:3 c8c11c14, C20:3 c11c14c17, C20:4 c5c8c11c14, C22:3 c13c16, C20:5 c5c8c11c14c17 (EPA), C22:3 c13c16c19, C22:4 c7c10c13c16, C22:5 c7c10c13c16c19 (DPA), C22:6 c4c7c10c13c16c19 (DHA).

<sup>9</sup>transPUFA: C18:2 t11t15, C18:2 t9t12.

<sup>10</sup>cis,trans+trans,cis PUFA: C18:2 c9t13, C18:2 c10t14, C18:2 c9t14, C18:2 c9t12, C18:2 t11c15, C18:2 t9c12, C18:2 t12c15, C18:2 c9t11 (RA) (co-elutes with C18:2 t7c9 + t8c10 + t6c8), other conjugated FA of unknown isomerism (CLA other a-h).

<sup>11</sup>omega-3 PUFA (n-3): C18:2 t11t15, C18:2 t11c15, C18:2 t12c15, C18:3 c9c12c15, C20:3 c11c14c17, C20:5 c5c8c11c14c17 (EPA), C22:3 c13c16c19, C22:5 c7c10c13c16c19 (DPA), C22:6 c4c7c10c13c16c19.

<sup>12</sup>omega-6 PUFA(n-6): C18:2 t9t12, C18:2 c9t12, C18:2 t9c12, C18:2 c9c12 (LA), C18:3 c6c9c12, C20:2 c11c14, C20:3 c8c11c14, C20:4 c5c8c11c14, C22:2 c13c16, C22:4 c7c10c13c16.

<sup>13</sup>Cis n-3 PUFA: C18:3 c9c12c15, C20:3 c11 c14 C17, C20:5 c5c8c11c14c17, C22:3 c13C16C19, C22:5 c7c10c13C16C19, C22:6 c4c7c10c13C16C19.

<sup>14</sup>Cisn-6 PUFA: C18:2 c9c12, C18:3 c6c9c12, C20:2 c11c14, C20:4 c5c8c11c14, C22:2 c13c16.

<sup>15</sup>transFA: C14:1 t9, C16:1 t6+t7+t8, C16:1 t9, C16:1 t11+t12+t13, C17:1 t10, C18:1 t4, C18:1 t5, C18:1 t6+t7+t8, C18:1 t9, C18:1 t10, C18:1 t11 (VA), C18:1 t12 (co-elutes with C18:1 c6), C18:1 t15, C18:1 t16 (co-elutes with C18:1 c14), C18:2 t11t15, C18:2 t9t12.

<sup>16</sup>Atherogenicity index = (C12:0 + (4 × C14:0) + C16:0) / (MUFA + PUFA), as described in Šrednicka-Tober, et al. [3].

<sup>17</sup>Thrombogenicity index= (C14:0 + C16:0 + C18:0) / (0.5 × MUFA) + (0.5 × n-6) + (3×n-3) + (n-3:n-6) as described in Šrednicka-Tober, et al. [3].

<sup>18</sup>Hypocholesterolemic to hypercholesterolemic ratio = (C18:1 c9 + total PUFA) / (C12:0 + C14:0 + C16:0) as described in Mierlita [4].

<sup>19</sup>Δ9-desaturase activity index= (c9 C14:1 + c9 C16:1 + OA + RA) / (c9 C14:1 + c9 C16:1 + OA + RA + C14:0 + C16:0 + C18:0 + VA) as described in Kay, et al. [5].

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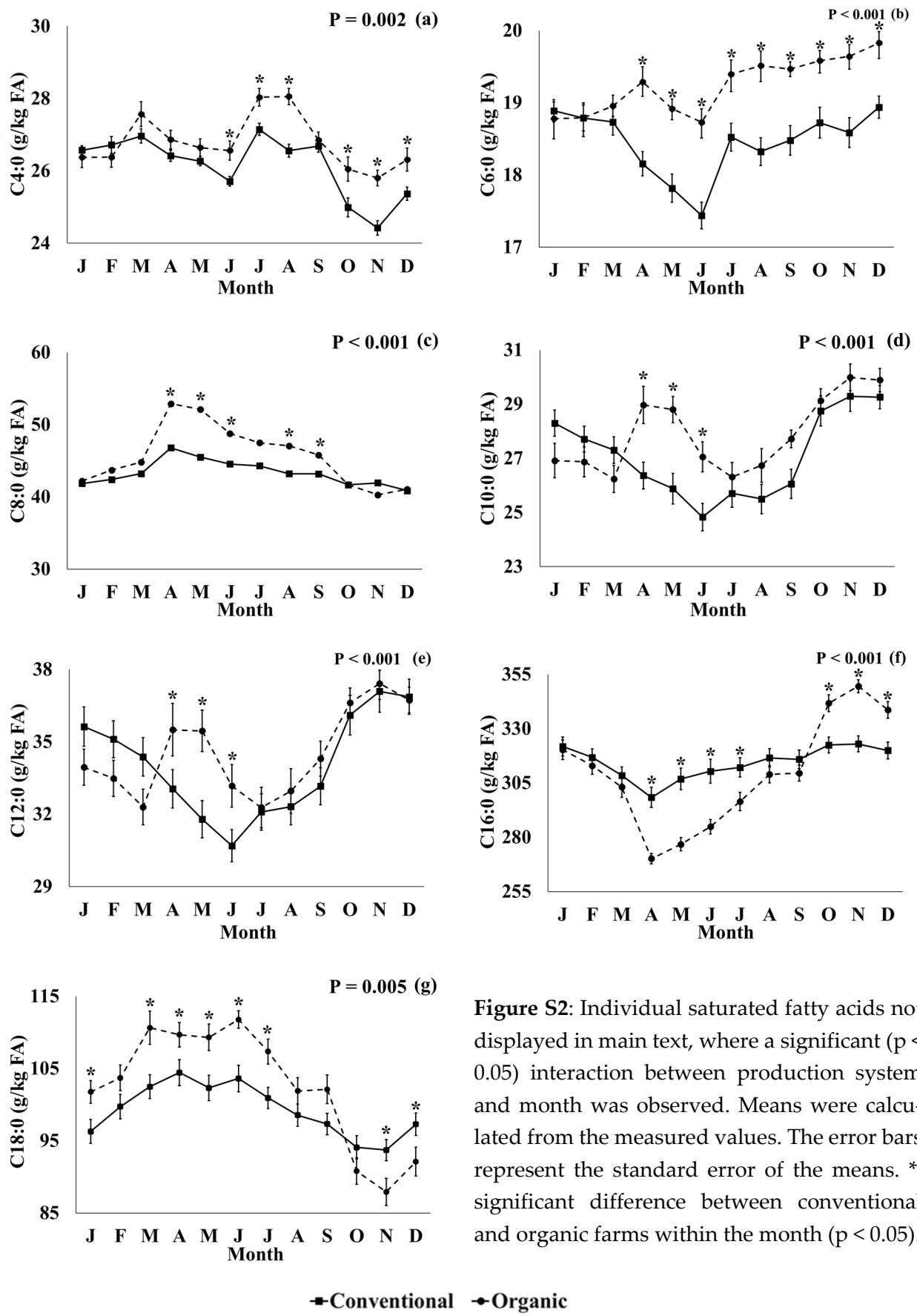


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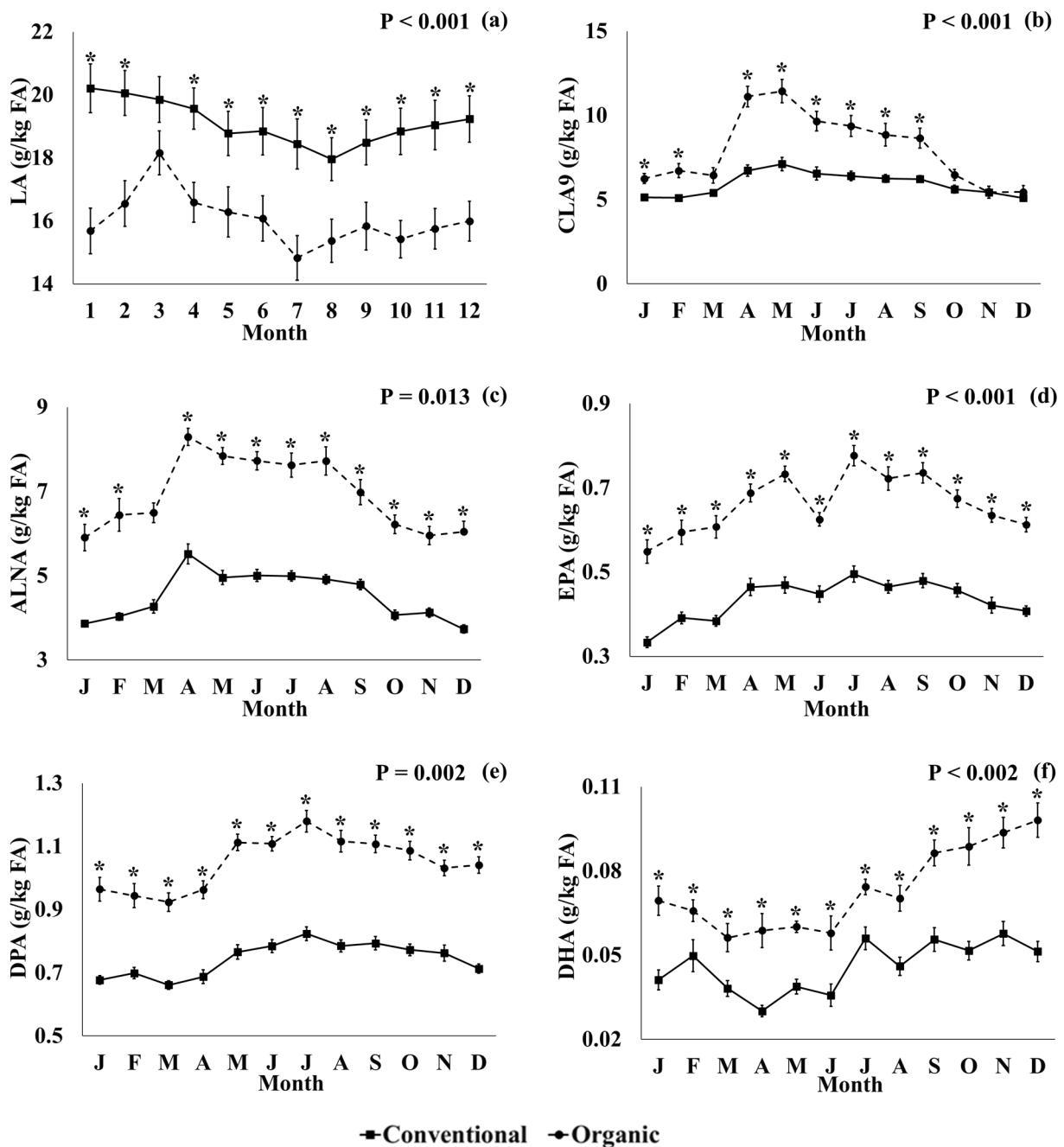
<sup>1</sup>n is the number of records used to calculate means ± SE and P-values. Records with missing values were not included in the analysis

<sup>2</sup>Significances were declared at P<0.05. Means for season within a row with different lower-case letters are significantly different according to Fisher's Least Significant Difference test (P<0.05).

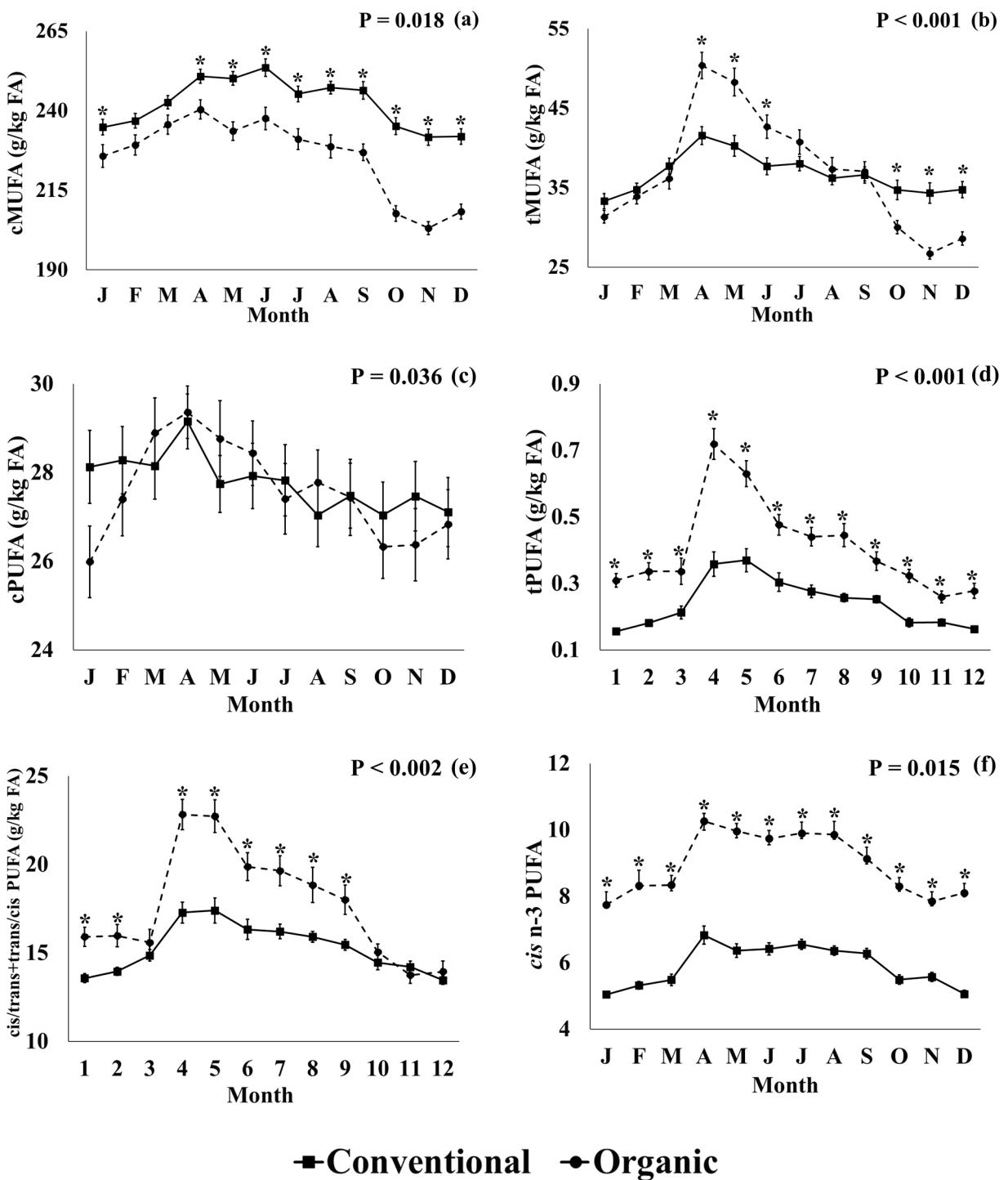
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**Figure S2:** Individual saturated fatty acids not displayed in main text, where a significant ( $p < 0.05$ ) interaction between production system and month was observed. Means were calculated from the measured values. The error bars represent the standard error of the means. \*: significant difference between conventional and organic farms within the month ( $p < 0.05$ ).



**Figure S3:** Individual polyunsaturated fatty acids not displayed in main text, where a significant ( $p < 0.05$ ) interaction between production system and month was observed. Means were calculated from the measured values. The error bars represent the standard error of the means. \*: significant difference between conventional and organic farms within the month ( $p < 0.05$ ).



**Figure S4:** FA groups not displayed in main text, where a significant ( $p < 0.05$ ) interaction between production system and month was observed. Means were calculated from the measured values. The error bars represent the standard error of the means. \*: significant difference between conventional and organic farms within the month ( $p < 0.05$ ).

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## References (supplementary material)

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