

Supplemental Table S1. Spearman's rho values between fecal microbial phyla and fatty acids for each diet: CTRL-AL, HF-AL, and HF-TRF (n=14 for each group). Blue shading highlights statistically significant negative Spearman's Rho. Green shading indicates positive associations. Normal font indicates p < 0.05; bold font indicates p < 0.01.

SFA	Actinobacteria			Bacteroidetes			Deferrribacteres			Firmicutes			Verrucomicrobia		
	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF
12:0	-0.34	-0.60	-0.01	0.30	0.63	0.13	-0.11	-0.02	-0.43	-0.03	-0.30	-0.15	-0.27	-0.24	0.27
14:0	-0.05	-0.54	0.07	0.44	0.61	0.10	-0.48	0.04	-0.42	-0.43	-0.27	-0.16	-0.18	-0.11	0.38
15:0	0.11	-0.29	-0.09	0.31	0.34	0.43	-0.52	0.20	-0.67	-0.35	-0.21	-0.53	-0.22	0.25	0.39
16:0	-0.09	-0.57	0.03	0.54	0.58	0.08	-0.57	-0.09	-0.42	-0.44	-0.28	-0.13	-0.37	-0.13	0.34
17:0	-0.16	-0.59	-0.01	0.46	0.62	0.09	-0.54	-0.05	-0.48	-0.36	-0.32	-0.18	-0.18	-0.12	0.41
18:0	-0.10	-0.61	0.00	0.55	0.60	0.11	-0.59	-0.09	-0.46	-0.50	-0.32	-0.13	-0.29	-0.14	0.31
20:0	-0.25	-0.53	-0.03	0.60	0.58	0.01	-0.57	-0.17	-0.44	-0.69	-0.37	-0.06	0.11	-0.11	0.31
21:0	0.07	-0.11	-0.14	-0.07	-0.04	-0.15	0.02	-0.10	-0.55	-0.10	-0.15	0.07	0.42	0.06	0.17
23:0	-0.39	-0.41	-0.16	0.40	0.17	-0.08	-0.34	0.03	-0.53	-0.35	-0.05	0.07	-0.01	-0.08	0.13
24:0	-0.26	-0.47	-0.10	0.48	0.33	-0.09	-0.40	-0.04	-0.49	-0.52	-0.17	0.06	0.19	-0.13	0.22
Total SFA	-0.10	-0.57	0.02	0.58	0.59	0.08	-0.58	-0.09	-0.43	-0.49	-0.30	-0.12	-0.29	-0.11	0.32
BCFA															
13:0, 12-Me	-0.20	-0.09	-0.17	0.47	0.42	0.17	-0.51	0.00	-0.30	-0.56	-0.61	-0.63	0.56	0.72	0.88
14:0, 13-Me	-0.14	-0.38	-0.03	0.69	0.36	0.18	-0.60	0.17	-0.39	-0.69	-0.30	-0.52	-0.03	0.59	0.79
14:0, 12-Me	-0.01	-0.27	-0.11	0.49	0.59	0.21	-0.67	-0.10	-0.37	-0.60	-0.60	-0.58	0.05	0.63	0.81
15:0, 14-Me	0.02	-0.64	0.10	0.35	0.56	0.08	-0.34	0.17	-0.55	-0.35	-0.45	-0.34	-0.02	0.27	0.69
16:0, 15-Me	0.06	-0.37	0.07	0.33	0.29	0.37	-0.40	0.12	-0.80	-0.13	-0.28	-0.44	-0.53	0.51	0.36
16:0, 14-Me	-0.08	-0.48	0.20	0.44	0.79	0.47	-0.28	-0.18	-0.39	-0.36	-0.60	-0.58	-0.10	0.17	0.44
17:0, 16-Me	-0.02	-0.34	0.08	0.59	0.66	0.24	-0.52	-0.12	-0.67	-0.58	-0.59	-0.23	-0.18	0.47	0.16
21:0, 18-Me	-0.11	-0.65	0.13	0.58	0.57	-0.12	-0.58	0.06	-0.27	-0.68	-0.30	-0.01	0.26	-0.17	0.45
Total BCFA	-0.05	-0.42	0.13	0.55	0.64	0.26	-0.64	-0.04	-0.45	-0.62	-0.62	-0.49	0.08	0.61	0.69
MUFA															
16:1n-7, Z	0.09	-0.34	0.11	0.52	0.73	0.49	-0.58	-0.15	-0.64	-0.51	-0.63	-0.59	-0.12	0.34	0.43
18:1n-9,E	-0.26	-0.56	-0.27	0.65	0.65	0.08	-0.64	0.07	-0.74	-0.57	-0.43	-0.07	0.03	0.23	0.19
18:1n-9,Z	-0.13	-0.59	-0.01	0.42	0.65	0.16	-0.42	-0.03	-0.47	-0.30	-0.35	-0.15	-0.19	-0.12	0.26
18:1n-7, Z	0.12	-0.49	0.01	0.32	0.75	0.31	-0.50	-0.09	-0.62	-0.32	-0.56	-0.38	-0.33	0.23	0.41
18:1n-6, Z	0.39	-0.01	-0.07	-0.17	0.41	0.29	-0.22	0.02	-0.18	0.05	-0.43	-0.50	-0.22	0.37	0.51
20:1n-9	-0.08	-0.65	0.07	0.56	0.54	-0.06	-0.35	0.05	-0.34	-0.75	-0.32	-0.06	0.23	-0.10	0.43
22:1n-9	-0.50	0.06	0.25	0.41	-0.21	0.17	-0.27	0.31	-0.43	-0.25	0.21	-0.07	0.46	-0.08	-0.10
24:1n-9	-0.10	-0.41	-0.22	0.35	-0.08	-0.16	-0.39	0.35	-0.48	-0.46	0.11	0.13	0.14	-0.06	0.06
Total MUFA	-0.22	-0.63	-0.10	0.53	0.64	0.19	-0.60	0.04	-0.55	-0.46	-0.28	-0.21	-0.12	-0.10	0.32
n-6 PUFA															
18:2, 9Z,11E	0.12	-0.17	0.12	0.29	0.15	-0.30	-0.29	-0.21	-0.39	-0.45	-0.26	0.24	0.19	-0.06	0.02
18:2 Z/Z	0.16	0.06	-0.01	-0.42	0.18	0.08	0.04	0.14	-0.32	0.31	-0.32	-0.22	-0.29	0.10	0.22
18:2n-6, Z,Z	-0.02	-0.44	0.12	0.19	0.55	0.12	-0.25	-0.18	-0.32	-0.18	-0.26	-0.19	-0.23	-0.16	0.39
20:3n-6	-0.39	-0.49	-0.14	0.49	0.40	-0.21	-0.53	0.03	-0.42	-0.47	-0.28	0.16	0.19	-0.04	0.22
20:4n-6	0.17	-0.38	0.04	0.16	0.33	-0.45	-0.10	-0.05	-0.12	-0.42	-0.28	0.19	0.47	-0.04	0.46
22:4n-6	-0.11	-0.45	0.41	0.23	0.09	-0.02	-0.33	0.15	-0.04	-0.40	-0.05	0.15	0.27	0.21	-0.08
22:5n-6	0.03	0.04	-0.19	0.10	0.00	-0.04	-0.52	-0.36	-0.56	-0.31	-0.11	0.05	0.04	-0.05	0.12
Total n-6 PUFA	-0.11	-0.56	0.07	0.37	0.57	-0.05	-0.53	-0.08	-0.32	-0.45	-0.37	-0.04	0.17	-0.07	0.41
n-3 PUFA															
18:3n-3	-0.11	-0.49	0.04	0.23	0.55	0.13	-0.22	-0.18	-0.40	-0.21	-0.20	-0.16	-0.12	-0.29	0.31
20:5n-3	0.02	-0.10	0.14	0.16	0.00	-0.34	-0.37	0.06	-0.31	-0.25	-0.15	0.30	0.12	0.08	-0.05
22:5n-3	-0.17	0.16	0.26	0.18	-0.11	0.18	-0.35	-0.05	-0.31	-0.39	0.23	-0.27	0.24	0.22	0.51
22:6n-3	-0.31	0.31	0.11	0.22	-0.28	-0.18	-0.35	0.04	-0.47	-0.24	0.12	0.24	0.45	-0.01	-0.19

Total n-3 PUFA	-0.09	-0.49	0.13	0.18	0.52	-0.11	-0.39	-0.15	-0.29	-0.25	-0.25	0.03	0.23	-0.24	0.37
Total FA	-0.14	-0.58	0.02	0.64	0.59	0.08	-0.63	-0.07	-0.43	-0.59	-0.28	-0.12	-0.14	-0.12	0.32

AL, ad libitum; CTRL, control diet; HF, high fat diet; TRF, time-restricted feeding; BCFA, branched-chain fatty acids; FA, fatty acids; MUFA, monounsaturated fatty acids; PUFA, polyunsaturated fatty acids; SFA, saturated fatty acids.

Supplemental Table S2. Spearman's rho values between fecal microbial families and fatty acids. Blue shading highlights statistically significant negative Spearman's Rho. Green shading indicates positive associations. Normal font indicates $p < 0.05$; bold font indicates $p < 0.01$.

SFA	<i>Bifidobacteriaceae</i>	<i>Bacteroidaceae</i>	<i>Tannerellaceae</i>	<i>Deferrribacteraceae</i>	<i>Clostridiaceae</i>	<i>Lachnospiraceae</i>	<i>Desulfovibrionaceae</i>	<i>Akkermansiaceae</i>
	CTRL- HF-AL	CTRL- HF-TRF	CTRL- HF-AL	CTRL- HF-TRF	CTRL- HF-AL	CTRL- HF-TRF	CTRL- HF-AL	CTRL- HF-TRF
12:0	-0.22 -0.65 NA	0.30 0.64 0.29	0.06 0.28 0.02	-0.11 -0.02 -0.43	0.33 0.32 0.83	0.22 -0.06 0.01	-0.26 -0.56 -0.40	0.27 -0.24 0.27
14:0	0.02 -0.55 NA	0.57 0.67 0.29	0.36 0.31 0.00	-0.48 0.04 -0.42	0.50 0.35 0.83	-0.21 -0.05 0.01	-0.52 -0.57 -0.41	-0.18 -0.11 0.38
15:0	0.10 -0.30 NA	0.58 0.50 0.43	0.71 0.44	-0.52 0.20 -0.67	0.27 -0.06 0.03	-0.45 -0.46 -0.39	-0.34 -0.03 -0.71	-0.22 0.25 0.39
16:0	0.02 -0.62 NA	0.60 0.63 0.27	0.44 0.25 -0.01	-0.57 -0.09 -0.42	0.51 0.30 0.82	-0.22 0.02 0.04	-0.55 -0.56 -0.42	-0.37 -0.13 0.34
17:0	-0.21 -0.65 NA	0.59 0.71 0.31	0.33 0.41 0.01	-0.54 -0.05 -0.48	0.49 0.28 0.78	-0.35 -0.05 0.00	-0.49 -0.53 -0.48	-0.18 -0.12 0.41
18:0	-0.01 -0.64 NA	0.56 0.65 0.28	0.38 0.28 0.01	-0.59 -0.09 -0.46	0.56 0.33 0.82	-0.35 -0.01 0.03	-0.57 -0.56 -0.44	-0.29 -0.14 0.31
20:0	-0.18 -0.59 NA	0.61 0.65 0.24	0.56 0.34 -0.05	-0.57 -0.17 -0.44	0.27 0.26 0.79	-0.73 -0.05 0.11	-0.57 -0.51 -0.43	0.11 -0.11 0.31
21:0	-0.08 -0.05 NA	-0.01 0.17 0.04	0.08 0.20 0.00	0.02 -0.10 -0.55	-0.42 -0.01 0.31	-0.54 0.14 0.22	0.14 0.11 -0.63	0.42 0.06 0.17
23:0	-0.31 -0.47 NA	0.48 0.40 0.07	0.42 0.34 -0.03	-0.34 0.03 -0.53	0.26 -0.04 0.44	-0.66 0.16 0.23	-0.36 0.01 -0.63	-0.01 -0.08 0.13
24:0	-0.20 -0.57 NA	0.51 0.48 0.12	0.54 0.34 -0.09	-0.40 -0.04 -0.49	0.09 0.00 0.58	-0.73 0.09 0.24	-0.44 -0.13 -0.59	0.19 -0.13 0.22
Total	0.01 -0.62 NA	0.66 0.64 0.26	0.51 0.27 -0.01	-0.58 -0.09 -0.43	0.46 0.32 0.81	-0.31 0.00 0.05	-0.58 -0.58 -0.43	-0.29 -0.11 0.32
BCFA								
13:0, 12-Me	-0.29 0.22 NA	0.68 0.41 0.34	0.48 0.50 0.44	-0.51 0.00 -0.30	0.05 0.37 -0.03	-0.70 -0.61 -0.53	-0.46 -0.26 -0.39	0.56 0.72 0.88
14:0, 13-Me	-0.08 -0.12 NA	0.82 0.50 0.27	0.73 0.46 0.36	-0.60 0.17 -0.39	0.28 0.07 0.22	-0.56 -0.29 -0.37	-0.69 -0.17 -0.46	-0.03 0.59 0.79
14:0, 12-Me	0.09 -0.06 NA	0.78 0.73 0.35	0.73 0.69 0.43	-0.67 -0.10 -0.37	0.13 0.45 0.09	-0.50 0.57 -0.43	-0.49 -0.42 -0.54	0.05 0.63 0.81
15:0, 14-Me	0.00 -0.46 NA	0.40 0.67 0.24	0.33 0.68 0.15	0.34 0.17 -0.55	0.26 0.40 0.37	-0.58 -0.37 -0.18	-0.39 -0.30 -0.51	-0.02 0.27 0.69
16:0, 15-Me	0.12 -0.13 NA	0.28 0.38 0.37	0.34 0.46 0.45	-0.40 0.12 -0.80	0.43 0.02 0.37	-0.07 -0.09 -0.36	-0.40 -0.14 -0.67	-0.53 0.51 0.36
16:0, 14-Me	-0.04 -0.48 NA	0.46 0.77 0.61	0.36 0.73 0.64	-0.28 -0.18 -0.39	0.28 0.09 0.15	-0.30 -0.43 -0.59	-0.53 -0.51 -0.58	-0.10 0.17 0.44
17:0, 16-Me	-0.05 -0.30 NA	0.64 0.59 0.28	0.55 0.65 0.23	-0.52 -0.12 -0.67	0.33 0.29 0.33	-0.47 -0.51 -0.14	-0.57 -0.42 -0.64	-0.18 0.47 0.16
21:0, 18-Me	0.03 -0.63 NA	0.66 0.63 0.19	0.61 0.39 -0.13	-0.58 0.06 -0.27	0.24 0.25 0.78	-0.47 0.00 0.15	-0.62 -0.46 -0.35	0.26 -0.17 0.45
Total	-0.04 -0.20 NA	0.80 0.72 0.26	0.64 0.67 0.30	-0.64 -0.04 -0.45	0.23 0.29 0.24	-0.54 -0.53 -0.35	-0.57 0.39 -0.44	0.08 0.61 0.69
MUFA								
16:1n-7, Z	0.05 -0.36 NA	0.60 0.67 0.51	0.52 0.71 0.45	-0.58 -0.15 -0.64	0.32 0.24 0.20	-0.41 -0.68 -0.44	-0.54 -0.42 -0.71	-0.12 0.34 0.43
18:1n-9, E	-0.23 -0.44 NA	0.78 0.79 0.09	0.63 0.76 0.07	-0.64 0.07 -0.74	0.29 0.32 0.39	-0.48 -0.31 0.10	-0.60 -0.38 -0.62	0.03 0.23 0.19
18:1n-9, Z	-0.11 -0.63 NA	0.58 0.69 0.29	0.31 0.40 0.02	-0.42 -0.03 -0.47	0.31 0.26 0.85	-0.07 -0.12 0.00	-0.33 -0.53 -0.41	-0.19 -0.12 0.26
18:1n-7, Z	0.06 -0.35 NA	0.31 0.87 0.46	0.17 0.52 0.21	-0.50 -0.09 -0.62	0.54 0.14 0.71	-0.13 -0.37 -0.22	-0.39 -0.59 -0.58	-0.33 0.23 0.41
18:1n-6, Z	0.35 -0.04 NA	-0.16 0.29 0.45	-0.09 0.56 0.23	-0.22 0.02 -0.18	0.36 0.41 0.09	-0.06 -0.61 -0.35	-0.07 -0.33 -0.35	-0.22 0.37 0.51
20:1n-9	-0.02 -0.58 NA	0.70 0.66 0.22	0.60 0.43 -0.07	-0.35 0.05 -0.34	0.02 0.26 0.75	-0.55 0.01 0.10	-0.48 -0.38 -0.41	0.23 -0.10 0.43
22:1n-9	-0.52 0.10 NA	0.43 0.08 -0.04	0.31 0.27 0.37	-0.27 0.31 -0.43	-0.10 -0.12 -0.34	-0.33 -0.02 -0.17	-0.20 0.49 -0.46	0.46 -0.08 -0.10
24:1n-9	-0.17 -0.35 NA	0.53 0.14 -0.11	0.49 0.37 -0.04	-0.39 0.35 -0.48	0.00 -0.16 0.24	-0.80 0.12 0.28	-0.22 0.37 -0.57	0.14 -0.06 0.06
Total	-0.18 -0.65 NA	0.73 0.69 0.31	0.46 0.40 0.08	-0.60 0.04 -0.55	0.41 0.23 0.79	-0.32 -0.08 -0.06	-0.46 -0.54 -0.45	-0.12 -0.10 0.32
n-6 PUFA								
18:2, 9Z,11E	0.18 -0.24 NA	0.31 0.18 -0.06	0.52 0.24 -0.13	-0.29 -0.21 -0.39	0.12 0.09 0.33	0.59 0.06 0.35	-0.28 -0.08 -0.45	0.19 -0.06 0.02
18:2 Z/Z	0.35 0.36 NA	-0.40 0.22 0.06	-0.17 0.34 0.17	0.04 0.14 -0.32	0.22 0.19 -0.23	0.09 -0.43 -0.19	0.22 -0.10 -0.14	-0.29 0.10 0.22
18:2n-6, Z/Z	0.06 -0.60 NA	0.37 0.52 0.35	0.09 0.24 0.02	-0.25 -0.18 -0.32	0.21 0.27 0.90	0.29 -0.04 -0.05	-0.07 -0.54 -0.32	-0.23 -0.16 0.39
20:3n-6	-0.35 -0.55 NA	0.49 0.57 -0.04	0.44 0.58 -0.14	-0.53 0.03 -0.42	0.28 0.19 0.48	-0.71 -0.16 0.34	-0.46 -0.12 -0.53	0.19 -0.04 0.22
20:4n-6	0.15 -0.35 NA	0.25 0.44 -0.09	0.34 0.37 -0.27	-0.10 -0.05 -0.12	-0.30 0.04 0.46	-0.52 0.02 0.36	-0.09 -0.29 -0.23	0.47 -0.04 0.46
22:4n-6	-0.21 -0.40 NA	0.25 0.35 -0.03	0.14 0.52 -0.08	-0.33 0.15 -0.04	0.19 0.16 0.17	-0.73 0.05 0.14	-0.20 0.11 -0.18	0.27 0.21 -0.08
22:5n-6	0.02 -0.16 NA	0.20 0.01 0.07	0.14 -0.05 -0.02	0.52 -0.36 -0.56	0.33 0.01 0.48	0.61 0.28 0.21	0.19 -0.07 -0.62	0.04 -0.05 0.12
Total	-0.10 -0.59 NA	0.65 0.68 0.23	0.36 0.42 -0.10	-0.53 -0.08 -0.32	0.15 0.08 0.80	-0.34 -0.10 0.13	-0.29 -0.42 -0.38	0.17 -0.07 0.41
n-3 PUFA								
18:3n-3	-0.01 -0.70 NA	0.37 0.50 0.34	0.10 0.16 0.01	-0.22 -0.18 -0.40	0.15 0.31 0.89	0.28 0.05 -0.01	-0.13 -0.57 -0.37	-0.12 -0.29 0.31
20:5n-3	0.08 -0.12 NA	0.24 0.10 -0.16	0.25 0.45 -0.02	-0.37 0.06 -0.31	-0.01 0.02 0.12	-0.32 -0.15 0.28	-0.06 0.26 -0.32	0.12 0.08 -0.05
22:5n-3	-0.06 -0.02 NA	0.38 -0.16 0.24	0.13 0.01 0.12	-0.35 -0.05 -0.31	0.37 -0.14 0.30	-0.39 0.08 -0.10	-0.28 0.00 -0.33	0.24 0.22 0.51
22:6n-3	-0.24 0.30 NA	0.33 -0.12 -0.32	0.11 0.09 0.06	-0.35 0.04 -0.47	0.09 -0.22 -0.43	-0.13 0.03 0.15	-0.13 0.52 -0.39	0.45 -0.01 -0.19
Total	0.01 -0.53 NA	0.46 0.62 0.14	0.18 0.21 -0.12	-0.39 -0.15 -0.29	0.09 0.06 0.73	-0.04 0.08 0.19	-0.14 -0.42 -0.40	0.23 -0.24 0.37
Total FA	-0.04 -0.64 NA	0.76 0.63 0.26	0.60 0.27 -0.01	-0.63 -0.07 -0.43	0.40 0.32 0.81	-0.39 0.00 0.05	-0.63 -0.58 -0.43	-0.14 -0.12 0.32

AL, *ad libitum*; CTRL, control diet; HF, high fat diet; TRF, time-restricted feeding; FA, fatty acids; BCFA, branched chain fatty acids; MUFA, monounsaturated fatty acids; PUFA, polyunsaturated fatty acids; SFA, saturated fatty acids; NA, not applicable due to all zero relative abundance in HF-TRF.

Supplemental Table S3. Spearman's rho values for fecal microbial phyla by bile acids. Blue shading highlights statistically significant negative Spearman's Rho. Green shading indicates positive associations. Normal font indicates $p < 0.05$; bold font indicates $p < 0.01$.

Bile Acids	Actinobacteria			Bacteroidetes			Deferribacteres			Firmicutes			Patescibacteria			Verrucomicrobia			
	CTRL-	AI	HF-AL	CTRL-	AI	HF-AL	CTRL-	AI	HF-AL	CTRL-	AI	HF-TRF	CTRL-	AI	HF-AL	HF-TRF	CTRL-	AI	HF-AL
α -MCA	-0.20	0.30	0.39	0.24	-0.22	-0.34	-0.41	-0.30	0.75	-0.22	0.24	0.62	-0.59	-0.05	0.01	-0.24	0.00	-0.56	
β/ω - MCA	-0.10	-0.18	0.50	0.24	0.15	-0.25	-0.10	-0.31	0.47	-0.09	0.12	0.47	-0.54	-0.15	-0.03	-0.14	-0.13	-0.38	
DCA	0.02	0.43	-0.21	0.05	-0.17	-0.18	-0.55	-0.21	0.29	-0.05	0.10	0.45	0.17	0.34	0.47	-0.15	0.04	-0.66	
t-DCA	-0.11	0.48	-0.28	0.25	-0.09	-0.53	-0.24	-0.43	0.39	-0.29	-0.25	0.58	-0.31	-0.02	0.11	0.07	0.45	-0.05	
HDCA	0.00	0.29	-0.63	-0.15	0.27	0.12	-0.01	-0.12	-0.26	0.10	-0.47	-0.08	0.20	0.14	0.25	-0.14	0.26	-0.16	
t-HDCA	-0.07	0.41	-0.62	-0.16	-0.17	-0.31	0.22	0.09	0.10	0.10	-0.19	0.39	0.08	0.16	0.35	-0.07	0.19	-0.43	
t-CDCA	-0.18	0.40	-0.39	-0.02	-0.61	-0.47	0.04	0.31	0.37	-0.04	0.27	0.61	0.19	0.09	0.40	0.12	-0.01	-0.42	
LCA	0.15	0.27	-0.43	-0.06	-0.20	-0.15	-0.51	0.06	-0.09	-0.03	0.08	0.44	0.24	-0.19	0.51	-0.34	0.17	-0.36	
CA	-0.27	0.68	0.22	0.08	-0.02	-0.36	-0.23	-0.72	0.68	-0.04	-0.21	0.55	-0.21	-0.21	-0.02	0.02	0.45	-0.51	
α -t-CA	-0.23	0.62	-0.27	0.09	-0.59	-0.52	-0.02	0.05	0.27	-0.08	0.25	0.69	0.06	0.11	0.30	-0.02	0.10	-0.51	
t-CA	-0.22	0.61	-0.37	0.01	-0.72	-0.40	0.18	0.09	0.31	-0.04	0.42	0.52	0.08	0.15	0.43	0.14	-0.02	-0.42	
ω -t-CA	0.25	0.06	-0.11	-0.29	-0.48	-0.52	0.33	0.28	0.08	0.19	0.35	0.73	0.01	0.09	0.11	0.12	-0.02	-0.40	
UDCA	0.10	0.38	-0.41	0.03	-0.66	0.30	-0.52	-0.05	0.07	-0.01	0.50	-0.09	0.10	0.14	0.58	-0.18	0.16	-0.39	
Taurine conjugates	-0.07	0.46	-0.30	-0.10	-0.71	-0.54	0.21	0.20	0.18	0.06	0.38	0.69	0.03	0.10	0.26	0.08	0.07	-0.41	
Total bile acids	-0.13	0.21	0.00	0.22	-0.07	-0.24	-0.41	-0.36	0.49	0.16	0.15	0.60	0.27	0.17	0.41	-0.17	0.04	-0.74	

AL, *ad libitum*; CTRL, control diet; HF, high fat diet; TRF, time-restricted feeding; α -MCA, α -muricholic acid; β/ω muricholic acids; DCA, deoxycholic acid; t-DCA, taurine deoxycholic acid; HDCA, hyodeoxycholic acid; t-HDCA, taurine hyodeoxycholic acid; t-CDCA, taurine chenodeoxycholic acid; LCA, lithocholic acid; CA, cholic acid; α -t-cholic acid, alpha taurine cholic acid; t-CA, taurine cholic acid; ω -t-cholic acid, alpha taurine cholic acid; UDCA, ursodeoxycholic acid.

Supplemental Table S4. Spearman's rho for fecal microbial families by bile acids. Blue shading highlights statistically significant negative Spearman's Rho. Green shading indicates positive associations. Normal font indicates $p < 0.05$; bold font indicates $p < 0.01$.

Bile Acids	Clostridiales																							
	<i>Bifidobacteriaceae</i>			<i>Bacteroidaceae</i>			<i>Tannerellaceae</i>			<i>Deferrribacteraceae</i>			<i>Family XIII</i>			<i>Lachnospiraceae</i>			<i>Peptococcaceae</i>			<i>Akkermansiaceae</i>		
	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF	CTRL-AL	HF-AL	HF-TRF
α -MCA	-0.18	0.11	NA	0.24	-0.04	-0.33	0.14	-0.19	-0.51	-0.41	-0.30	0.75	-0.11	-0.03	0.29	-0.06	0.41	0.53	-0.12	-0.34	0.39	-0.24	0.00	-0.56
β/ω - MCA	-0.12	-0.43	NA	0.12	0.20	-0.24	0.12	-0.10	-0.34	-0.10	-0.31	0.47	-0.11	0.09	0.02	0.13	0.43	0.32	0.12	-0.20	0.47	-0.14	-0.13	-0.38
DCA	0.08	0.11	NA	0.09	-0.05	-0.20	0.16	0.28	-0.07	-0.55	-0.21	0.29	0.00	0.23	0.69	-0.34	-0.20	0.29	0.03	-0.66	-0.27	-0.15	0.04	-0.66
t-DCA	0.03	0.45	NA	0.31	-0.12	-0.37	0.34	0.02	-0.48	-0.24	-0.43	0.39	-0.01	0.33	0.28	-0.23	-0.06	0.57	0.33	-0.40	0.32	0.07	0.45	-0.05
HDCA	0.26	0.37	NA	-0.03	0.34	0.25	0.19	0.46	0.22	-0.01	-0.12	-0.26	-0.29	-0.01	0.38	0.03	-0.78	-0.02	0.00	-0.30	-0.69	-0.14	0.26	-0.16
t-HDCA	0.17	0.61	NA	-0.13	-0.16	-0.21	0.11	0.09	-0.10	0.22	0.09	0.10	-0.15	-0.05	0.69	0.09	-0.45	0.36	0.14	-0.07	-0.42	-0.07	0.19	-0.43
t-CDCA	-0.08	0.64	NA	0.05	-0.52	-0.55	0.22	-0.25	-0.66	0.04	0.31	0.37	-0.07	0.01	0.61	-0.32	0.09	0.66	0.18	-0.02	0.06	0.12	-0.01	-0.42
LCA	0.17	0.47	NA	0.03	0.03	-0.23	0.10	0.06	-0.35	-0.51	0.06	-0.09	-0.13	-0.13	0.29	-0.34	-0.10	0.49	-0.19	-0.14	-0.34	-0.34	0.17	-0.36
CA	-0.22	0.43	NA	-0.05	-0.08	-0.12	-0.04	-0.09	-0.50	-0.23	-0.72	0.68	0.23	0.51	0.22	-0.24	-0.02	0.48	0.10	-0.35	0.22	0.02	0.45	-0.51
α -t-CA	-0.18	0.67	NA	-0.04	-0.60	-0.63	0.10	-0.37	-0.61	-0.02	0.05	0.27	0.21	0.15	0.52	-0.35	0.04	0.66	0.22	-0.10	0.06	-0.02	0.10	-0.51
t-CA	-0.15	0.61	NA	-0.14	-0.74	-0.56	0.02	-0.55	-0.41	0.18	0.09	0.31	0.25	0.16	0.62	-0.23	0.19	0.48	0.49	-0.16	0.05	0.14	-0.02	-0.42
ω -t-CA	0.21	0.24	NA	-0.27	-0.22	-0.59	-0.14	-0.24	-0.70	0.33	0.28	0.08	0.02	-0.07	0.35	-0.07	0.31	0.75	0.70	0.10	0.21	0.12	-0.02	-0.40
UDCA	0.21	0.35	NA	0.07	-0.66	0.06	0.19	-0.70	0.44	-0.52	-0.05	0.07	-0.01	-0.02	0.45	-0.20	0.50	-0.28	-0.04	-0.34	-0.13	-0.18	0.16	-0.39
Taurine conjugated	-0.04	0.64	NA	-0.13	-0.59	-0.66	-0.02	-0.47	-0.63	0.21	0.20	0.18	0.04	0.04	0.51	-0.21	0.23	0.69	0.57	-0.04	0.13	0.08	0.07	-0.41
Total fecal BA	0.00	-0.08	NA	0.20	0.14	-0.19	0.32	0.02	-0.24	-0.41	-0.36	0.49	-0.10	0.08	0.55	-0.17	0.18	0.44	0.18	-0.66	-0.08	-0.17	0.04	-0.74

AL, *ad libitum*; CTRL, control diet; HF, high fat diet; TRF, time-restricted feeding; α -MCA, α -muricholic acid; β/ω muricholic acids; DCA, deoxycholic acid; t-DCA, taurine deoxycholic acid; HDCA, hyodeoxycholic acid; t-HDCA, taurine hyodeoxycholic acid; t-CDCA, taurine chenodeoxycholic acid; LCA, lithocholic acid; CA, cholic acid; α -t-cholic acid, alpha taurine cholic acid; t-CA, taurine cholic acid; ω -t-cholic acid, alpha taurine cholic acid; UDCA, ursodeoxycholic acid; NA, not applicable due to all zero relative abundance in HF-TRF.

Supplemental Table S5. Spearman's rho values for fecal microbial phyla by short chain fatty acids. SCFA, short chain fatty acids. Blue shading highlights statistically significant negative Spearman's Rho. Green shading indicates positive associations. Normal font indicates $p < 0.05$; bold font indicates $p < 0.01$.

SCFA	Phylum		<u>Actinobacteria</u>			<u>Bacteroidetes</u>			<u>Cyanobacteria</u>			<u>Deferribacteres</u>			<u>Firmicutes</u>			<u>Verrucomicrobia</u>		
	CTRL-	AL	HF-AL	HF-TRF	CTRL-	AL	HF-AL	HF-TRF	CTRL-	AL	HF-TRF	CTRL-	AL	HF-TRF	CTRL-	AL	HF-TRF	CTRL-	AL	HF-TRF
% Acetate	0.53	0.54	-0.21	-0.54	-0.43	-0.14	-0.34	-0.05	-0.37	0.10	-0.05	-0.32	0.45	0.26	0.23	-0.34	-0.01	-0.42		
% Propionate	-0.33	0.12	0.22	0.45	0.16	0.11	0.10	-0.14	0.60	0.09	-0.06	0.34	-0.42	-0.24	-0.39	0.38	0.22	0.62		
% Butyrate	-0.51	-0.64	0.15	0.62	0.36	0.16	0.21	0.13	0.27	-0.30	0.02	0.27	-0.58	-0.18	-0.12	0.36	0.01	0.20		
acetate	0.22	0.01	0.13	0.48	0.01	0.12	-0.13	0.07	0.18	-0.42	0.06	0.19	-0.58	-0.27	-0.07	0.01	0.31	0.20		
propionate	-0.23	0.12	0.19	0.63	0.05	0.23	0.10	-0.07	0.46	-0.15	-0.07	0.31	-0.63	-0.25	-0.27	0.26	0.32	0.37		
butyrate	-0.12	-0.41	0.19	0.64	0.28	0.22	0.13	0.17	0.25	-0.31	0.06	0.24	-0.65	-0.32	-0.16	0.19	0.13	0.19		
isobutyrate	-0.09	0.16	0.14	0.68	-0.09	0.30	0.15	-0.08	0.07	-0.32	0.02	0.07	-0.53	-0.12	-0.15	0.12	0.42	-0.10		
isovalerate	-0.24	0.25	0.11	0.63	-0.26	0.32	0.48	0.15	-0.11	-0.32	0.13	-0.16	-0.45	-0.03	-0.08	0.03	0.32	-0.20		
valerate	0.04	0.05	0.17	0.59	0.02	0.22	-0.03	0.10	0.00	-0.41	-0.08	0.14	-0.63	-0.29	-0.02	0.11	0.49	-0.15		
Total SCFA	0.06	-0.01	0.16	0.59	0.07	0.21	-0.09	0.09	0.21	-0.40	0.04	0.23	-0.66	-0.30	-0.15	0.08	0.29	0.19		

AL, *ad libitum*; CTRL, control; HF, high fat diet; TRF, time-restricted feeding; SCFA, short chain fatty acids.

Supplemental Table S6. Spearman's rho for fecal microbial families by short chain fatty acids. Blue shading highlights statistically significant negative Spearman's Rho. Green shading indicates positive associations. Normal font indicates $p < 0.05$; bold font indicates $p < 0.01$.

Family	<i>Clostridiales</i>												<i>Uncultured (Alpha)</i>																							
	<i>Eggerthellaceae</i>			<i>Bacteroidaceae</i>			<i>Flavobacteriaceae</i>			<i>Muribaculaceae</i>			<i>Clostridiaceae_1</i>			<i>Family XIII</i>			<i>Lactobacillaceae</i>			<i>Peptococcaceae</i>			<i>Desulfovibrionaceae</i>			<i>Enterobacteriaceae</i>			<i>Proteobacteria</i>			<i>Tannerellaceae</i>		
SCFA	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF	CTRL-	HF-AL	HF-TRF						
% Acetate	0.05	0.21	-0.18	0.49	-0.51	-0.24	-0.37	0.52	0.27	-0.20	-0.15	-0.39	0.03	-0.06	-0.60	-0.10	0.39	0.34	-0.37	0.60	-0.10	-0.40	0.35	-0.55	0.58	0.33	-0.06	0.38	0.03	0.23	-0.43	-0.15	-0.23	-0.50	-0.11	0.03
% Propionate	0.13	-0.29	0.28	0.53	0.46	0.16	0.59	-0.26	-0.09	0.23	-0.15	0.32	-0.21	-0.18	0.11	-0.01	-0.35	-0.43	0.48	-0.16	0.31	0.35	-0.20	0.56	-0.46	0.08	0.01	-0.56	-0.32	-0.19	0.23	0.22	0.35	0.48	0.32	0.13
% Butyrate	0.05	-0.14	0.10	0.59	0.30	0.17	0.10	-0.45	-0.17	0.19	0.15	0.30	0.01	0.04	0.71	0.12	-0.25	-0.24	0.31	-0.55	0.04	0.24	-0.22	0.44	-0.58	-0.35	0.12	-0.16	0.14	0.00	0.52	0.03	0.13	0.59	-0.10	-0.08
acetate	-0.08	-0.61	0.12	0.52	0.26	0.01	-0.12	-0.56	-0.40	0.25	-0.20	0.53	0.11	-0.18	0.53	0.23	-0.58	-0.15	0.13	-0.32	0.18	-0.33	0.10	0.54	-0.47	0.16	-0.07	-0.14	-0.43	-0.15	0.58	0.14	0.25	0.57	-0.04	-0.06
propionate	-0.02	-0.44	0.23	0.67	0.35	0.19	0.42	-0.45	-0.27	0.34	-0.23	0.52	-0.09	-0.23	0.46	0.09	-0.47	-0.25	0.48	-0.23	0.28	0.03	-0.17	0.55	-0.62	0.12	-0.03	-0.46	-0.38	-0.18	0.51	0.16	0.39	0.71	0.10	0.03
butyrate	-0.09	-0.37	0.16	0.60	0.39	0.10	0.25	-0.67	-0.27	0.34	0.01	0.50	0.06	-0.08	0.63	0.30	-0.51	-0.20	0.38	-0.57	0.14	0.00	-0.11	0.51	-0.68	-0.16	0.05	-0.41	-0.10	0.00	0.70	0.09	0.25	0.67	-0.03	-0.02
isobutyrate	-0.15	-0.31	0.05	0.60	0.15	0.08	0.12	-0.06	-0.48	0.12	-0.58	0.49	0.00	-0.60	0.18	0.16	-0.38	0.13	0.35	-0.03	0.19	-0.19	0.18	0.03	-0.69	0.29	-0.33	-0.43	-0.12	-0.10	0.72	-0.17	0.31	0.72	0.04	0.06
isovalerate	-0.47	-0.50	0.05	0.38	-0.04	-0.07	0.15	-0.44	-0.43	0.17	-0.29	0.57	0.17	-0.38	0.11	0.30	-0.64	0.13	0.29	-0.15	-0.01	-0.10	0.11	-0.06	-0.74	0.36	-0.27	-0.39	-0.52	-0.04	0.77	0.20	0.22	0.61	-0.14	-0.01
valerate	-0.08	-0.53	0.09	0.61	0.26	-0.01	0.07	-0.63	-0.42	0.25	-0.30	0.52	0.10	-0.26	0.36	0.22	-0.57	0.05	0.28	-0.30	0.14	-0.15	-0.11	0.22	-0.61	0.05	-0.10	-0.25	-0.42	-0.02	0.67	0.16	0.16	0.66	-0.06	-0.08
Total SCFA	-0.05	-0.54	0.14	0.64	0.31	0.10	0.07	-0.59	-0.38	0.31	-0.19	0.55	0.11	-0.20	0.52	0.20	-0.58	-0.13	0.30	-0.33	0.20	-0.26	0.02	0.51	-0.59	0.10	-0.04	-0.25	-0.38	-0.15	0.62	0.13	0.30	0.66	-0.01	-0.01

SCFA, short chain fatty acids; AL, *ad libitum*; CTRL, control; HF, high fat diet; TRF, time-restricted feeding; SCFA, short chain fatty acids.