

Cooperative harvesting of aquatic resources triggered the beginning of pottery production in north-eastern North America

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What benefits were derived from the invention of pottery? And why did ceramics remain marginal for so long? Increasing use of pottery has been seen as a response to large-scale harvesting in a model that favours economic advantage through increased efficiency. This paper challenges that view; combining carbon and nitrogen isotope and lipid analysis the authors argue that pottery was used selectively for storing or processing valued exchange commodities such as fish oil. Its use can be seen as part of broader developments in hunter-gatherer society, featuring seasonal gatherings, collective feasting and a new articulation of social relations.

Keywords: North America, hunter-gatherers, pottery production, aquatic resources, carbon isotopes, nitrogen isotopes, lipid analysis, Vinette 1 pottery, social gatherings

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Table S1. Charred deposits and ceramic samples selected for bulk isotope analysis, lipid analysis by GCMS and GC-c-IRMS. FA (Cx:y)=fatty acids with carbon length x and number of unsaturations y, br=branched chain acids, DCx=α,ω-dicarboxylic acids with carbon length x, ALx=alkanes with carbon length x, pri=pristanic acid, phy=phytanic acid, TMTD=4,8,12-trimethyltridecanoic acid, APFA (Cn)=ω-(o-alkylphenyl) alkanolic acids with carbon length n, ODA= dehydroabiatic acid, cholest=cholesterol and cholesterol by-products, lano=lanosterol, stig=stigmasterol, ergo=ergosterol, ket=long chain ketones. F=interior foodcrust sample, S=exterior carbonised deposit, I=interior ceramic sample, E=exterior ceramic sample. tr=trace, NA=not applicable. Lipid concentrations are expressed in ug mg⁻¹ for carbonised deposits, and in ug g⁻¹ for interior and exterior ceramic samples. In the column P/A of aquatic biomarkers, two ticks indicate complete sets of biomarkers and one tick indicates partial sets of biomarkers. Each line groups potsherds from the same vessel.

Sites	Sample ID	Lipid analysis		Bulk isotope analysis				GC-C-IRMS		
		Lipid conc.	Compound detected	Aquatic Biomarkers (P/A)	C16/C18	δ ¹⁵ N	δ ¹³ C	C:N	C _{16:0} δ ¹³ C (‰)	C _{18:0} δ ¹³ C (‰)
Bruce Boyd	AdHc4-1-F	NA	NA		NA	8.62	-24.94	14.75	NA	NA
	AdHc4-1-I	69.99	FA(C _{9:0-28:0} C _{16:1-18:1} C _{15-17br}), AL(C _{16-18,26}), DC(C ₅₋₁₂), APFA (C _{18(tr)}), phy	P(v)	1.68					
	BB01-I	191.48	FA(C _{5:0-22:0} C _{18:1} C _{17br}), AL(C ₁₄₋₂₈) DC(C ₉₋₁₃), APFA (C ₁₈), phy, ket	P(v)	0.45				-27.45	-27.93
	BB05-I	10.37	FA(C _{9:0-28:0(tr)} C _{15-17br}), AL(C ₁₆₋₂₅), DC(C ₉)	A	3.13					
	BB07-I	19.32	FA(C _{12:0-26:0} C _{14(tr),15br}), AL(C ₁₆₋₂₇), DC(C _{9(tr)})	A	1.84					
	BB08-I	5.18	FA(C _{12-26(tr)} C _{18:1} C _{15,17(br)}), AL(C ₁₇₋₂₈), TMTD(tr)	P(v)	1.28					
	BB09-I	27.14	FA(C _{12:0-26:0} C _{14(tr),15br}), AL(C ₁₆₋₂₉), DC(C _{9(tr)})	A	2.99					
	BB09-E	53.15	FA(C _{12:0-28:0} C _{16:1-18:1} C _{14,15,17br}), AL(C ₁₆₋₃₁), DC(C ₉₋₁₁), APFA(C ₁₆), TMTD	P(v)	1.90					

	BB10-I	13.82	FA(C _{12:0-26:0} C _{18:1} C _{14(tr),15-17br}), AL(C _{16-27,29}), DC(C _{9(tr)})	A	2.67			
	AfGr9-1-F	<0.5				6.72	-28.25	8.91
	AfGr9-1-I	122.53	FA(C _{10:0-26:0} C _{16:1-20:1} C _{16,17br}), AL(C _{16,17}), DC(C _{8,9,12,16}), APFA (C _{16-22(tr)}), pri, phy, ket (C _{16-16,16-18,18-18})	P(vv)	0.73			-27.85 -29.22
	PB01-I	206.26	FA(C _{8:0-26:0} C _{16:1,18:1,20:1} C _{15-17,19br}), AL(C ₂₂₋₃₀), APFA(C _{18,20}), phy, cholest	P(vv)	1.56			
	PB02-I	<5						
Peace Bridge	PB03-I	127.79	FA(C _{10:0-26:0} C _{18:1} C _{17br}), APFA (C _{18-22(tr)}), pri, phy, ket	P(vv)	0.42			-29.31 -30.88
	PB04-I	<5						
	PB06-I	<5						
	PB08-I	<5						
	PB09-I	6.25	FA(C _{12:0(tr)-28:0(tr)} C _{16:1,18:1} C _{15-17br}), AL(C ₁₆₋₂₀)	A	1.45			
	Cda17.3-1-F	<0.5				5.59	-25.80	11.63
	Cda17.3-1-I	50.70	FA(C _{10:0-28:0}), AL(C ₁₅₋₂₈)	A	1.02			
	Cda17.3-2-F	<0.5				6.39	-25.91	8.29
	Cda17.3-2-I	34.90	FA(C _{9:0-28:0} C _{15br}), AL(C ₁₅₋₂₈), DC(C ₆₋₉)	A	0.57			
	Cda17.3-3-I	6.09	FA(C _{12:0-26:0} C _{18:1,22:1} C _{14-17br}), AL (C ₁₆₋₂₇)	A	1.55			
Scaccia	Cda17.3-4-I	23.82	FA(C _{9:0-28:0} C _{15,17br}), AL(C ₁₆₋₂₇), DC(C ₆₋₉)	A	2.70			
	Cda17.3-5-I	<5						
	Cda17.3-6-I	42.62	FA(C _{9:0-28:0} C _{18:1,22:1} C _{17,18,22,24,25br}), AL(C ₁₅₋₂₄), DC(C ₉)	A	1.24			
	Cda17.3-6-E	17.16	FA(C _{9:0-10:0,12:0,14:0-18:0,20:0,22:0,24:0,26:0(tr)} C _{18:1,22:1} C _{15br}), AL(C ₁₇₋₂₉), DC(C ₆)	A	1.04			
	Cda17.3-7-I	14.01	FA(_{12:0-30:0} C _{18:1} C _{15,17,22,24br}), AL(C ₁₆₋₂₇), DC(C _{6,9})	A	1.52			
	Cda17.3-8-I	<5						

	Cda17.3-9-I	45.63	FA(C _{10:0-26:0}), AL(C ₁₄₋₂₅)	A	2.05				
	Cda17.3-10-I	440.74	FA(C _{9:0-30:0} C _{18:1} C _{15-17,26,27br}), AL(C ₁₄₋₂₈), DC(C ₅₋₁₄), pri, phy	P(v)	0.69				-31.11 -31.86
Vine Valley	VineValley1-F	<0.5				6.53	-27.95	11.51	
	VineValley1-I	305.56	FA(C _{8:0-28:0} C _{16:1-18:1} C _{16,17br}), AL(C ₁₄₋₂₇), DC(C _{6,8-11}), TMTD(tr), pri(tr), phy, ODA(tr)	P(v)	0.44				-27.34 -28.29
	VineValley2-F	<0.5				6.51	-27.53	10.29	
	VineValley3-F	<0.5				7.19	-25.96	7.01	
	VineValley3-I	201.28	FA(C _{8:0-28:0} C _{18:1} C _{15,17br}), AL(C ₁₅₋₂₆), DC(C _{6,8}), APFA(C ₁₈), phy, ODA(tr)	P(v)	0.69				
	VineValley4-F	<0.5				6.73	-26.02	6.62	
Dawson Creek	BaGn16-1-I	36.11	FA(C _{9:0-10:0,12:0-18:0,20:0,22:0-28:0} C _{18:1} C _{15br}), AL(C ₁₆₋₂₅), DC(C ₉)	A	4.70				
	BaGn16-2-I	21.60	FA(C _{9:0-10:0,12:0-18:0,20:0,22:0-28:0} C _{18:1} C _{15br}), AL(C ₁₆₋₂₅), DC(C ₉)	A	2.20				
	BaGn16-3-I	27.10	FA(C _{9:0-10:0,12:0-18:0,20:0,22:0-28:0} C _{18:1}), AL(C ₁₆₋₂₅), DC(C ₉)	A	2.36				
	BaGn16-3-E	32.63	FA(C _{9:0-26:0(tr)} C _{16:1-18:1} C _{14-17br}), AL(C ₁₆₋₂₇), DC(C _{6-11,13}), TMTD	P(v)	1.37				
	BaGn16-4-I	13.32	FA(C _{9:0-10:0,12:0-18:0,20:0,22:0-28:0} C _{18:1}), AL(C ₁₆₋₂₅), DC(C ₉)	A	2.53				
	BaGn16-5-I	<5							
BaGn16-6-I	<5								
Pointe-du-Buisson	BhFl1-1-F	0.8	FA(C _{9:0-26:0} C _{16:1-22:1} C _{15-17br}), AL(C ₂₂₋₃₁), APFA(C ₁₆₋₂₂), pri, phy	P(vv)	0.95	6.92	-27.47	16.96	
	BhFl1-1-I	227.70	FA(C _{10:0-26:0} C _{18:1,22:1} C _{15-17br}), AL(C ₁₅₋₁₉), DC(C ₇₋₁₂), APFA(C _{16-22(tr)}), phy, stig(tr)	P(vv)	1.01				-27.09 -28.13
	BhFl1-24-F	<0.5				8.78	-25.93	12.80	
	BhFl1-2-F	0.72	FA(C _{9:0-26:0} C _{16:1-22:1} C _{15-19br}), DC(C ₆₋₁₄), APFA(C ₁₆₋₂₀), TMTD, pri, phy	P(vv)	0.62	9.54	-26.78	15.93	-26.32 -27.65
	BhFl1-2-I	776.14	FA(C _{9:0-26:0} C _{16:1-20:1} C _{13-19br}), AL(C ₁₄₋₁₈), DC(C ₇₋₁₈), APFA(C ₁₆₋₂₂), TMTD, pri, phy	P(vv)	0.70				-27.05 -28.08
	BhFl1-3-F	1.25	FA(C _{12:0-24:0} C _{16:1-22:1} C _{15-19br}), AL(C ₂₄₋₂₈), DC(C ₉₋₁₁), APFA(C ₁₆₋₂₀), TMTD, pri, phy, cholest	P(vv)	1.15	10.26	-24.99	10.70	-26.03 -26.33

BhFI1-3-I	217.28	FA(C _{9:0-26:0} C _{16:1-20:1} C _{15-19br}), AL(C ₁₅₋₁₉), DC(C ₆₋₁₈), APFA(C ₁₆₋₂₂), TMTD, phy, cholest	P(vv)	1.34					-25.65	-26.37
BhFI1-19-I	121.98	FA(C _{9:0-30:0} C _{18:1-20:1} C _{15-19br}), AL(C ₁₇₋₁₈), DC(C _{6-12,14,16}), APFA(C _{16-22(tr)}), phy	P(vv)	1.52					-24.95	-25.20
BhFI1-4-F	Ind	Not completely derivatized			7.94	-26.61	14.30			
BhFI1-4-I	792.34	FA(C _{9:0-28:0(tr)} C _{16:1-22:1} C _{13,15-19br}), AL(C ₁₅₋₁₈), DC(C ₆₋₁₆), APFA(C ₁₄₋₂₄), TMTD, pri, phy, cholest, ergo(tr)	P(vv)	2.05					-26.82	-26.63
so	1.63	FA(C _{14:0-26:0} C _{16:1-22:1} C _{15-19br}), AL(C ₂₄₋₂₈), DC(C ₆₋₁₁), APFA(C ₁₆₋₂₀), TMTD, pri(tr), phy	P(vv)	1.12	10.86	-26.63	41.00			
BhFI1-23-F	<0.5				8.45	-25.31	8.80			
BhFI1-23-I	298.85	FA(C _{9:0-26:0} C _{16:1-20:1} C _{13-19br}), AL(C ₁₅₋₁₈), DC(C ₆₋₁₆), APFA(C ₁₆₋₂₂), TMTD, pri(tr), phy	P(vv)	1.68					-25.19	-25.41
BhFI1-6-F1	1.3	FA(C _{12:0-26:0(tr)} C _{16:1-22:1} C _{15-19br}), DC(C ₇₋₁₃), APFA(C ₁₆₋₂₂), TMTD, pri, phy	P(vv)	2.10	10.45	-24.12	9.90		-26.60	-26.15
BhFI1-6-F2	NA	NA			11.37	-24.50	9.40			
BhFI1-6-S	<0.5				10.59	-24.94	26.50			
BhFI1-6-I	57.72	FA(C _{9:0-26:0} C _{18:1} C _{15-19br}), AL(C ₁₆₋₁₉), DC(C ₇₋₁₂), APFA(C ₁₆₋₂₂), TMTD, phy	P(vv)	1.80					-24.63	-25.69
BHFI1-6-E	17.03	FA(C _{10:0-28:0} C _{16:1-18:1} C _{14-19br}), AL(C _{17,18,22-29})DC(C ₈₋₁₃), TMTD	P(v)	2.07						
BhFI1-7-F	0.54	FA(C _{12:0-28:0} C _{16:1-22:1} C _{15-19br}), AL(C ₂₄₋₃₀), DC(C ₇₋₁₁), APFA(C _{16(tr)-22}), TMTD, pri, phy	P(vv)	0.88	7.79	-26.19	10.20			
BhFI1-8-F	<0.5				8.47	-27.41	12.10			
BhFI1-21-F	<0.5				8.92	-25.02	9.10			
BhFI1-11-F	<0.5				7.89	-28.20	9.00			
BhFI1-11-I	64.92	FA(C _{9:0-26:0} C _{18:1} C _{15-18br}), AL(C ₁₆₋₁₉), DC(C ₆₋₁₂), APFA(C _{16-20(tr)}), phy	P(vv)	1.01					-26.55	-27.14
BhFI1-12-I	24.22	FA(C _{9:0-20:0,24:0,26:0} C _{18:1} C _{15,17,19br}), AL(C ₁₆₋₁₉), DC(C ₈₋₉)		1.10						
BhFI1-13-I	624.75	FA(C _{9:0-30:0} C _{16:1-20:1} C _{15-17,19,25br}), AL(C ₁₄₋₁₈), DC(C ₅₋₁₈), APFA(C _{16(tr),18,20(tr)}), TMTD, phy	P(vv)	0.71					-29.61	-31.93
BhFI1-14-F	1.68	FA(C _{14:0-28:0} C _{18:1} C _{15-19br}), AL(C ₂₃₋₃₀), APFA(C _{18,20}), pri, phy, cholest	P(vv)	0.35	6.80	-27.37	13.10		-28.01	-30.24
BhFI1-14-I	53.87	FA(C _{9:0-28:0(tr)} C _{18:1} C _{15,17br}), AL(C ₁₅₋₂₄), DC(C ₆₋₁₀), ODA(tr), APFA(C _{18,20}), phy	P(vv)	0.73					-27.27	-28.53

	BhFI1-15-I	145.84	FA(C _{9:0-30:0} C _{16:1-22:1} C _{15-18br,21br}), AL(C ₁₆₋₁₈), DC(C ₅₋₁₄), APFA(C _{18,20(tr)}), TMTD, pri(tr), phy, cholest(tr)	P(vv)	0.94				-25.17	-25.54
	BhFI1-17-I	1056.4	FA(C _{8:0-26:0} C _{16:1-20:1} C _{12-17br,19br}), AL(C ₁₄₋₁₈), DC(C _{6-13,16}), APFA(C _{16-22(tr)}), TMTD, phy	P(vv)	0.82				-27.78	-29.36
	BhFI1-18-I	187.37	FA(C _{9:0-26:0} C _{18:1-20:1} C _{13-18br}), AL(C _{15-18,20}), DC(C ₆₋₁₄), APFA(C _{16-22(tr)}), TMTD, phy	P(vv)	1.78				-23.98	-24.51
	BhFI1-20-I	82.28	FA(C _{9:0-28:0} C _{16:1-20:1} C _{15-17br,18-27br}), AL(C ₁₆₋₁₉), DC(C ₈₋₁₆), APFA(C ₁₈), TMTD, phy	P(v)	0.23				-24.95	-25.20
	BhFI1-22-F	<0.5				10.01	-24.09	9.30		
	BhFI1-25-F	<0.5				8.63	-26.57	9.70		
	BhFI1-26-I	139.23	FA(C _{9:0-30:0} C _{18:1} C _{15-19br}), AL(C ₁₅₋₂₇), DC(C ₆₋₁₂), APFA(C ₁₆₋₂₂), TMTD, pri(tr), phy	P(vv)	0.99				-24.40	-24.91
	CcFb1-1-F	<0.5				6.96	-26.86	11.70		
	CcFb1-1-I	402.84	FA(C _{9:0-26:0(tr)} C _{18:1} C _{15,17,19br}), AL(C _{15-18,20}) APFA(C _{18-22(tr)}), TMTD, phy, cholest	P(vv)	1.70				-27.71	-27.92
	CcFb1-2-I	65.48	FA(C _{14:0-28:0} C _{16:1-18:1} C _{15-17br}), AL(C ₁₇₋₂₃), DC(C _{9,11,14})	A	1.18				-28.56	-29.97
	CcFb1-3-I	215.76	FA(C _{10:0-26:0} C _{16:1-18:1} C _{15-17,25,26br}), AL(C ₁₇₋₂₉), DC(C ₉₋₁₃), stig, cholest	A	1.87				-26.90	-28.84
Batiscan	CcFb1-3-E	167.04	FA(C _{9:0-28:0} C _{16:1-18:1} C _{13-17,25br}), AL(C _{15-18,20,22-27}), DC(C ₇₋₁₄), TMTD, pri, phy	P(v)	1.81					
	CcFb1-4-I	229.94	FA(C _{12:0-28:0(tr)} C _{18:1} C _{15-18br}), AL(C ₁₇₋₂₃), DC(C _{11,16}), APFA(C ₁₈₋₂₂), phy, cholest	P(vv)	1.35				-27.09	-30.52
	CcFb1-5-I	141.56	FA(C _{14:0-26:0} C _{18:1} C _{15-17br}), AL(C ₁₆₋₂₄), APFA(C ₁₈), phy	P(v)	1.26				-29.69	-30.54
	CcFb1-6-I	471.38	FA(C _{12:0-26:0} C _{18:1} C _{15-19br}), AL(C _{16-20,24-27}), DC(C ₉₋₁₄), APFA(C _{16(tr),18-22(tr)}), phy, cholest	P(vv)	0.81				-28.67	-29.93
	CcFb1-7-I	57.00	FA(C _{14:0-26:0(tr)}), AL(C ₁₆₋₂₅)	A	1.61				-29.12	-29.92
	CcFb4-1-F	<0.5				9.50	-24.58	8.40		
Parc	CcFb4-1-I	302.70	FA(C _{14:0-26:0} C _{18:1,20:1} C _{15-19br}), DC(C _{8-11,16,18}), APFA(C _{18,20}), phy, cholest	P(vv)	1.35				-26.98	-27.24
des	CcFb4-2-F	<0.5				10.39	-24.36	8.40		
Pins	CcFb4-2-I	84.77	FA(C _{14:0-24:0} C _{16:1,18:1} C _{15-19br}), DC(C ₉), APFA(C _{18,20(tr)}), phy, cholest	P(vv)	1.78				-29.67	-30.07
	CcFb4-3-F	<0.5				9.12	-25.71	8.10		

	CcFb4-3-I	12.77	FA(C _{16:0-26:0(tr)} C _{18:1} C _{17br})	A	0.71			
	CcFb4-4-I	87.47	FA(C _{14:0-24:0} C _{18:1-20:1} C _{15-17br}), DC(C ₉₋₁₆), cholest	A	1.55		-28.82	-29.75
	CcFb4-5-I	23.85	FA(C _{9:0-26:0} C _{16:1-20:1} C _{15-19br}), DC(C _{8-11,18,22})	A	1.01			
	CcFb4-6-I	9.88	FA(C _{14:0-24:0} C _{18:1} C _{15-17br})	A	0.86			
	CcFb4-7-I	27.36	FA(C _{15:0-28:0} C _{16:1,18:1,18:2} C _{15-17,19br}), DC(C _{16,18,20,22})	A	0.92			
	CcFb4-8-I	31.45	FA(C _{12:0-26:0} C _{16:1,18:1,18:2} C _{15-18br})	A	1.25			
	CcFb4-9-I	15.46	FA(C _{15:0-26:0(tr)} C _{16:1,18:1} C _{16,17br})	A	1.11		-28.98	-29.40
	CeEu12-1-F	<0.5				7.82	-26.37	15.40
	CeEu12-1-I	111.27	FA(C _{9:0-26:0} C _{16:1(tr)-20:1} C _{15-19br}), AL(C ₁₅₋₂₀), DC(C ₆₋₁₁), APFA(C ₁₆₋₂₀), TMTD, pri, phy	P(vv)	2.26			
	CeEu12-2-F	<0.5				7.22	-26.45	16.04
	CeEu12-3-F	0.59	FA (C _{16:0,18:0})	A	1.34	6.58	-26.06	10.38
	CeEu12-3-S	1.46	FA (C _{16:0,18:0})	A	1.47	12.47	-24.31	27.41
	CeEu12-4-F	<0.5				9.04	-25.21	11.44
Lambert	CeEu12-4-S	<0.5						
	CeEu12-4-I	133.12	FA(C _{9:0-28:0} C _{16:1-22:1} C _{13,15-19br}), AL(C ₁₆₋₁₈), DC(C _{8,9,11,12}), APFA(C ₁₆₋₂₂), TMTD, pri, phy	P(vv)	1.88		-24.74	-26.25
	CeEu12-4-E	39.41	FA(C _{12:0-28:0} C _{16:1-22:1} C _{14-19br}), AL(C _{16-18,20}), DC(C ₆₋₁₄), APFA(C _{18,20}), TMTD, pri, phy	P(vv)	1.39			
	CeEu12-5-F	N/A	N/A			9.87	-23.89	10.50
	CeEu12-5-I	35.54	FA(C _{9:0-28:0(tr)} C _{18:1} C _{13,17,19br}), AL(C ₁₆₋₂₆) APFA(C _{18(tr),20(tr)})	P(vv)	0.46		-25.82	-26.41
	CeEu12-6-I	196.68	FA(C _{9:0-30:0} C _{16:1-20:1} C _{15-19br}), AL(C ₁₅₋₁₈), DC(C ₇₋₁₂), APFA(C ₁₆₋₂₂), TMTD, phy	P(vv)	0.87		-26.84	-27.12
Gasser	BgFg2-1-I	25.54	FA(C _{9:0-18:0,24:0} C _{16:1,18:1,22:1} C _{15br}), AL(C ₁₇₋₂₀), DC(C ₉), APFA(C _{18,20}), ODA	P(v)	1.62			
Carson	VT1-F	<0.5						

Farm	VT1-I	93.32	FA(C _{9:0-30:0} C _{16:1-20:1} C _{13-17,19-23,25-27br}), AL(C ₁₅₋₂₉), DC(C _{6-9,11-13}), APFA(C _{16(tr),18-22(tr)}), pri, phy	P(vv)	0.67			
	VT2-F	<0.5						
	VT2-I	62.52	FA(C _{9:0-28:0(tr)} C _{18:1} C _{15,17br}), AL(C ₁₄₋₂₉), DC(C ₆₋₉)	A	2.03			
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Fort Hunter	36Da159-1-I	<5						
	36Da11-1-F	<0.5				10.28	-23.72	10.85
Mc-Cormick Is.	36Da11-2-F	<0.5				11.06	-22.03	9.38
	36Da11-2-I	20.03	FA(C _{9:0-26:0} C _{18:1-24:1} C _{15-18br}), AL(C ₁₆₋₁₈), DC(C ₇₋₁₂), APFA(C ₁₈₋₂₂), TMTD, pri, phy	P(vv)	1.62			-23.47 -25.18
	36Da11-2-E	6.78	FA(C _{12:0-18:0,20:0,24:0} C _{18:1,22:1} C _{14-17br}), AL(C _{17-20,22,23}), DC(C ₉), APFA(C ₁₆), TMTD	P(v)	1.42			
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Sheldon	Sheldon-1-I	48.50	FA(C _{12:0-28:0} C _{16:1-22:1(tr)} C _{15-19br}), AL(C ₁₆₋₁₉), phy	P(v)	0.36			-28.89 -29.40
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	36Br58-1-I	<5						
Wilson	36Br58-2-I	53.78	FA(C _{10:0-28:0} C _{18:1}), AL(C ₁₅₋₂₈), DC(C ₇₋₉), ODA(tr)	A	0.39			
	36Br58-3-I	44.34	FA(C _{9:0-26:0} C _{18:1}), AL(C ₁₅₋₂₉), DC(C ₄₋₉), APFA(C ₁₈), phy, ODA(tr)	P(v)	0.78			
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Barton complex	18AG3-1-I	28.47	FA(C _{10:0-28:0} C _{16:1-18:1,22:1} C _{15,17,23,25br}), AL(C _{15-17,20}), cholest., ODA(tr)	A	1.84			
	18AG3-1-E	34.20	FA(C _{12:0-30:0} C _{16:1-18:1,22:1} C _{14-17,19-23,25br}), AL(C _{15-18,20,22-27}), DC(C ₉), cholest.	A	2.02			
	18AG8-1-I	28.94	FA(C _{12:0-28:0} C _{18:1,22:1} C _{15,25br}), AL(C _{15-20,22-25})	A	2.31			
	18AG8-2-I	142.04	FA(C _{10:0-30:0} C _{16:1-22:1} C _{15,17br}), AL(C _{15-18,22,24-28}), DC(C ₇₋₁₀), APFA(C ₁₈), phy	P(v)	0.65			-27.77 -29.25
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Zimmerman	36Pi14-1-F	<0.5				5.68	-24.31	14.31
	36Pi14-2-F	<0.5				4.95	-26.03	10.72
	36Pi14-2-I	<5						
	36Pi14-3-F	<0.5				7.77	-25.45	12.56
	36Pi14-3-I	107.51	FA(C _{8:0-26:0} C _{16:1-22:1} C _{16-18br}), AL(C ₁₇₋₁₈), DC(C ₈₋₁₄), APFA(C ₁₆₋₂₂), TMTD, pri, phy, stig(tr)	P(vv)	1.54			-25.35 -25.66

	36Pi14-3-E	20.25	FA(C _{12:0-26:0} C _{16:1-18:1,22:1} C _{15,17br}), AL(C _{17-18,22-24}), DC(C ₉₋₁₁), APFA (C ₁₆₋₁₈), TMTD, pri, phy	P(v)	1.63			
	36Pi14-4-F	<0.5				5.26	-24.92	11.22
	36Pi14-4-I	<5						
	36Pi14-5-F	<0.5				10.18	-24.69	12.10
	36Pi14-5-I	24.04	FA(C _{12:0-26:0} C _{16:1-18:1} C _{15-17br}), AL(C ₂₀₋₂₅), DC(C ₈₋₁₂), APFA (C _{16(tr),18,20(tr)}), pri, phy	P(vv)	1.34			
	28SX28-1-I	229.61	FA(C _{9:0-30:0} C _{16:1-22:1} C _{15-17br,19br,21br,23br}), AL(C ₁₄₋₁₉), DC(C ₈₋₁₈), APFA (C ₁₆₋₂₂), TMTD, pri, phy, stig	P(vv)	0.53			-26.67 -29.16
Minisink	28SX28-2-I	22.54	FA(C _{11:0-28:0} C _{18:1} C _{17br}), AL(C ₁₅₋₂₉), DC(C ₈₋₉)	A	5.41			
Island	28SX28-3-I	16.29	FA(C _{11:0-28:0} C _{18:1} C _{16-17br,19br}), AL(C ₁₆₋₂₇), DC(C ₆₋₉), APFA(C ₁₈₋₂₂), phy	P(vv)	0.81			-25.89 -26.51
	28SX28-4-I	69.21	FA(C _{11:0-30:0} C _{18:1-22:1} C _{17-19br,23br,25br}), AL(C ₁₅₋₁₉), DC(C _{8-10,14}), phy	P(v)	0.17			-27.03 -28.37
	27BK16-1-I	44.02	FA(C _{10:0-28:0} C _{16:1-24:1} C _{15,17br}), AL(C ₁₄₋₂₄), DC(C ₆₋₁₄), APFA(C ₁₆₋₂₂), pri, phy	P(vv)	1.68			
Drake	27BK16-2-I	106.81	FA(C _{10:0-28:0(tr)} C _{16:1-22:1} C _{15,17br}), AL(C ₁₅₋₁₈), DC(C _{8,11,13,14,16,18}), APFA(C ₁₆₋₂₂), pri(tr), phy, stig	P(vv)	0.96			
Hormell	27CA15-1	32.94	FA(C _{12:0-28:0} C _{18:1} C _{15-17br}), AL(C _{15-20,25,26,28}), DC(C ₆₋₁₁), stig, cholest	A	1.88			
	36La51-1-I	169.20	FA(C _{9:0-30:0} C _{18:1-24:1} C _{15-17br,23br}), AL(C ₁₄₋₂₇), DC(C ₅₋₁₁), APFA (C ₁₈₋₂₂), TMTD, pri, phy(tr)	P(vv)	2.13			
Roberts	36La51-2-I	8.96	FA(C _{10:0-18:0,24:0,26:0}), AL(C ₁₄₋₂₈), DC(C ₉)	A	2.73			
	36La51-3-I	91.97	FA(C _{9:0-30:0(tr)} C _{18:1-24:1} C _{15-18br}), AL(C ₁₅), DC(C ₅₋₁₈), APFA (C ₁₆₋₂₂), TMTD, pri, phy	P(vv)	2.27			
Kirby Brooks	6LF2-1-I	38.30	FA(C _{12:0-30:0(tr)} C _{15,16,24-26br}), AL(C ₁₆₋₂₅), TMTD	P(v)	2.75			
	6LF2-2-I	183.59	FA(C _{10:0-30:0} C _{18:1} C _{17,25,26br}), AL(C ₁₅₋₂₃), DC(C ₈₋₉)	A	0.49			
	6LF70-1-I	167.78	FA(C _{10:0-30:0} C _{18:1-22:1} C _{18br}), AL(C ₁₄₋₂₂), DC(C ₇₋₁₂), APFA (C ₁₈₋₂₂), TMTD, pri, phy, ergo	P(vv)	0.98			
Lovers Leap	6LF70-2-I	<5						
	6LF70-3-I	101.87	FA(C _{12:0-30:0} C _{18:1-20:1} C _{16,17,22-26br}), AL(C ₁₅₋₃₁), DC(C _{8,9,18}), APFA(C _{18-22(tr)}), phy	P(vv)	0.46			
	6LF70-4-I	10.83	FA(C _{12:0-16:0,18:0,20:0,23:0-26:0} C _{16:1-18:1}) AL(C ₁₆₋₂₇), ODA(tr)	A	0.91			

	6LF70-5-I	23.06	FA(C _{11:0-14:0,16:0,18:0,23:0-26:0}) AL(C ₁₅₋₂₇), DC(C ₉)	A	0.97			
	6LF70-6-I	138.92	FA(C _{10:0-30:0} C _{16:1-26:1,18:3} C _{17,18br}), AL(C ₁₆₋₂₄), DC(C ₄₋₁₄), APFA (C ₁₆₋₂₂), TMTD, pri, phy	P(vv)	3.26			
	6LF70-7-I	72.14	FA(C _{9:0-30:0} C _{18:1-22:1}), AL(C ₁₄₋₂₇), DC(C ₅₋₁₁), APFA(C _{18-22(tr)}), pri, phy	P(vv)	1.41			
	6LF70-8-I	264.90	FA(C _{9:0-30:0} C _{18:1-22:1} C _{15-17br}), AL(C ₁₄₋₂₀), DC(C _{6-12,14,16}), APFA (C ₁₈₋₂₂), TMTD, pri, phy, stig, ergo, lano	P(vv)	1.87			
Hopkins	6LF1-1-I	14.03	FA(C _{12:0-30:0} C _{16:1-18:1,18:2} C _{15,17br}), AL(C ₁₆₋₂₆), DC(C ₆₋₁₁), APFA (C _{18-22(tr)}), phy	P(vv)	2.16			
	NH38.6-1-F	<0.5				5.16	-25.00	20.46
	NH38.6-1-I	212.73	FA(C _{8:0-26:0} C _{14:1-18:1,18:2} C _{15-17br}), AL(C _{22,24-27}), DC(C ₅₋₁₃), TMTD, pri(tr), phy	P(v)	3.61			-24.96 -24.22
	NH38.6-2-I	<5						
	NH38.6-3-I	76.75	FA(C _{8:0-30:0} C _{16:1-24:1} C _{15-18br}), AL(C ₁₆₋₂₀), DC(C ₄₋₁₆), APFA(C ₁₆₋₂₂), TMTD, pri, phy	P(vv)	1.80			-23.94 -23.53
Eddy	NH38.6-3-E	32.52	FA(C _{11:0-30:0} C _{16:1-24:1} C _{15-18br}), AL(C ₁₆₋₂₀), DC(C ₇₋₁₆), APFA(C _{16(tr),18(tr),20(tr),22(tr)}), TMTD(tr), pri, phy	P(vv)	2.08			
	NH38.6-4-I	7.86	FA(C _{12:0-28:0} C _{16:1-18:1} C _{15br,17br}), AL(C _{16-19,22-26})	A	1.41			
	NH38.6-5-I	101.22	FA(C _{12:0-30:0} C _{18:1-24:1} C _{15-18br}), AL(C ₁₆₋₂₇), DC(C ₅₋₁₄), APFA(C ₁₈₋₂₂), TMTD, pri, phy	P(vv)	2.62			-25.67 -26.07
	NH38.6-6-I	75.05	FA(C _{9:0-30:0(tr)} C _{18:1-24:1} C _{17br,18br}), AL(C ₁₅₋₂₆), DC(C ₅₋₁₀), APFA(C _{16(tr),18,20(tr),22}), pri, phy, ODA, stig	P(vv)	1.13			
	NH38.6-7-I	88.78	FA(C _{11:0-30:0(tr)} C _{16:1-24:1} C _{17br}), AL(C ₁₅₋₂₂), DC(C ₈₋₁₄), APFA(C _{16(tr),18(tr),20,22}), pri, phy, ODA, stig, cholest	P(vv)	1.68			
Hoffman	M12.43-1-I	65.30	FA(C _{9:0-28:0} C _{15-17br}), AL(C ₁₅₋₂₉), DC(C ₉), APFA(C _{18,20(tr)}), TMTD	P(vv)	2.51			
	Mpt-1-I	21.31	FA(C _{14:0-28:0} C _{16:1-18:1} C _{15-18br}), AL(C ₂₂₋₂₄), DC(C ₈₋₁₁)	A	2.16			
Morrill Point	Mpt-2-I	204.42	FA(C _{11:0-30:0} C _{16:1-20:1} C _{15-17br}), AL(C _{17,28}), DC(C ₇₋₁₆), APFA(C ₁₆₋₂₄), TMTD, pri, phy	P(vv)	1.40			
	Mpt-3-I	7.61	FA(C _{14:0-28:0} C _{16:1-18:1} C _{14-16br}), AL(C _{17,18}), DC(C ₈)	A	2.51			
	M49.2-1-F	<0.5				11.51	-21.22	13.30
Hornblower	M49.2-2-F	<0.5				10.57	-22.49	10.60
	M49.2-3-F	<0.5				8.97	-23.47	9.20

	M49.2-4-F	<0.5					8.73	-23.23	9.90		
	M49.2-5-F	<0.5					8.07	-22.99	7.70		
	M49.2-5-I	99.66	FA(C _{9:0-28:0} C _{16:1-20:1}), AL(C ₁₄₋₂₆), DC(C ₆₋₁₀), APFA(C _{16(tr),18,20}), TMTD, phy		P(vv)	1.15					
	M49.2-5-E	64.92	FA(C _{9:0-24:0} C _{16:1} C _{15,17br}), AL(C ₁₇₋₂₇), DC(C ₆₋₉), ODA		A	1.11					
	M49.2-6-F	<0.5					9.04	-23.16	9.30		
	M49.2-6-I	79.81	FA(C _{8:0-28:0} C _{16:1-20:1} C _{13,15-17br}), AL(C ₁₅₋₂₂), DC(C ₈₋₁₀), APFA(C _{16(tr),18}), TMTD(tr), phy		P(v)	0.87					
	M49.2-10-F	<0.5					9.75	-22.75	10.50		
	M49.2-10-I	45.08	FA(C _{8:0-28:0} C _{16:1-18:1} C _{15br}), AL(C ₁₅₋₂₈), DC(C ₅₋₁₀)		A	1.36					
	M49.2-11-F	<0.5					8.93	-22.74	8.70		
	M49.2-12-F	<0.5					11.17	-22.42	9.30		
	M49.2-12-I	318.62	FA(C _{9:0-30:0} C _{18:1-20:1} C _{15-18br}), AL(C _{14-20,22-25}), APFA(C ₁₈), phy		P(v)	0.60					
	M49.2-13-F	<0.5					14.31	-21.89	9.70		
	M49.2-14-F	<0.5					14.27	-21.83	8.80		
	M49.2-14-I	160.76	FA(C _{8:0-30:0} C _{18:1-22:1} C _{15,17br}), AL(C ₁₄₋₂₇), DC(C ₆₋₁₂), APFA(C ₁₆₋₂₄), TMTD, pri, phy, cholest		P(vv)	0.83				-25.66	-25.78
	M38.7-1/2-F	0.54	FA(C _{12:0-18:0,20:0,22:0,24:0} C _{16:1-18:1} C _{15,17br}), AL(C ₁₆₋₂₂), DC(C ₈₋₁₁), TMTD(tr), phy(tr)		P(v)	2.03	14.29	-21.73	7.90		
	M38.7-2-I	25.72	FA(C _{8:0-26:0} C _{18:1} C _{15-17br}), AL(C ₁₆₋₂₆), DC(C ₈₋₁₁), TMTD(tr), ODA		P(v)	1.72					
Small swamp	M38.7-3-F	<0.5					13.29	-22.40	12.90		
	M38.7-4-F	<0.5					11.52	-21.43	11.60		
	M38.7-4-I	20.38	FA(C _{8:0-30:0} C _{18:1-24:1} C _{15-18br}), AL(C ₁₅₋₂₄), DC(C ₈₋₁₂), APFA(C ₁₆₋₂₄), TMTD, pri, phy, ODA		P(vv)	1.04				-23.74	-23.62
Rose	M38.11-1-F	<0.5					12.89	-21.26	7.50		
	M38.11-1-I	20.89	FA(C _{10:0-26:0} C _{16:1-18:1} C _{15,17br}), AL(C ₁₆₋₂₂), DC(C ₈₋₁₂), TMTD		P(v)	1.76					

RI 1428	RI1428-1-F	<0.5					13.82	-20.67	8.88			
	RI1428-4-F	<0.5					12.73	-22.72	13.61			
	RI1428-4-I	54.11	FA(C _{7:0-26} C _{16:1-24:1} C _{13-19br}), AL(C ₁₅₋₁₈), DC(C ₈₋₁₂), APFA(C ₁₆₋₂₄), TMTD, pri, phy, cholest, wax esters	P(vv)	3.80						-25.96	-24.18
	RI1428-2-F	<0.5					12.53	-18.60	7.14			
	RI1428-2-I	19.87	FA(C _{9:0-28:0} C _{16:1-22:1} C _{15-19br}), AL(C ₁₅₋₂₂), APFA(C ₁₆₋₂₄), pri, phy	P(vv)	1.46						-22.50	-21.99
	RI1428-2-E	<5										
	RI1428-3-F	<0.5					13.84	-19.30	10.12			
	RI1428-5-I	20.28	FA(C _{9:0-24:0} C _{16:1-18:1} C _{14-18br}), AL(C ₁₆₋₃₀), DC(C ₆₋₁₂), ODA(tr)	A	3.27							
	RI1428-6-I	<5										
	RI1428-7-I	38.97	FA(C _{12:0-26:0} C _{16:1-20:1} C _{15-17br}), AL(C _{17,18}), DC(C ₆₋₁₄), APFA (C ₁₆), TMTD, pri, phy	P(v)	2.54						-21.69	-22.85
Knox	Knox-1-I	100.90	FA(C _{14:0-28:0} C _{16:1-18:1} C _{15-17br}), AL(C ₁₆₋₂₃), DC(C ₁₆), cholest	A	1.16							
	Knox-2-I	22.45	FA(C _{12:0-30:0(tr)} C _{18:1} C _{15,17br}), AL(C ₁₆₋₂₅), DC(C _{8,9})	A	2.15							
Great Diamond Island	GDA-1-F	N/A	N/A				10.53	-25.33	10.28			
	GDA-1-I	22.80	FA(C _{9:0-24:0} C _{18:1} C _{15,17br}), AL(C ₁₆₋₂₅), DC(C _{6,8,9}), APFA(C _{16(tr),18}), phy	P(v)	1.66							
	GDA-2-F	<0.5					11.32	-25.47	13.58			
	GDA-2-I	96.74	FA(C _{11:0-28:0} C _{16:1-22:1} C _{15-17br}), AL(C ₁₆₋₁₈), DC(C ₇₋₁₆), APFA(C ₁₆₋₂₂), TMTD, pri(tr), phy	P(vv)	0.92						-26.42	-28.87
	GDA-3-F	<0.5					11.53	-24.42	13.59			
	GDA-4-F	<0.5					11.72	-24.58	11.51			
	GDA-4-I	14.36	FA(C _{9:0-24:0} C _{15br}), AL(C ₁₆₋₂₅)	A	1.37							
	GDA-5-F	<0.5					13.41	-23.01	9.56			
	GDA-5-I	14.43	FA(C _{9:0-24:0} C _{18:1} C _{15-17br}), AL(C ₁₇₋₂₀), DC(C _{6,8,9}), APFA(C ₁₈), phy	P(v)	1.10							

GDA-6-F	<0.5					13.93	-19.03	7.90		
GDA-6a-I	184.54	FA(C _{10:0-30:0} C _{16:1-22:1} C _{14-17br}), AL(C ₁₅₋₂₈), DC(C ₇₋₁₄), APFA(C ₁₆₋₂₂), TMTD, pri, phy, cholest	P(vv)	2.86					-24.15	-24.01
GDA-7-I	130.13	FA(C _{9:0-30:0} C _{16:1-22:1} C _{15-17br}), AL(C ₁₆₋₂₇), DC(C ₅₋₁₄), APFA(C ₁₆₋₂₂), TMTD, pri, phy	P(vv)	3.07					-23.81	-23.81
GDA-8-I	101.02	FA(C _{10:0-30:0} C _{18:1} C _{15-17br}), AL(C ₁₆₋₂₆), DC(C _{8,9}), ODA	A	1.55						
GDA-9-I	72.99	FA(C _{11:0-30:0} C _{18:1} C _{15br}), AL(C ₁₅₋₂₈), DC(C ₈₋₁₀), ODA	A	1.54					-26.57	-26.40
GDA-9-E	43.86	FA(C _{12:0-28:0} C _{18:1} C _{15,17br}), AL(C ₁₆₋₂₉), DC(C ₈₋₉), ODA	A	2.04						
GDA-10-I	237.75	FA(C _{10:0-30:0} C _{16:1-24:1} C _{15-17br}), AL(C ₁₅₋₂₂), DC(C ₆₋₁₄), APFA(C ₁₆₋₂₄), TMTD, pri, phy	P(vv)	1.39					-25.23	-27.93
GDA-11-I	46.11	FA(C _{11:0-28:0} C _{14-17,25br}), AL(C ₁₅₋₂₉), DC(C _{8,9}), ODA(tr)	A	3.46						
GDA-12-I	79.42	FA(C _{11:0-26:0} C _{16:1} C _{13-15br}), AL(C ₁₆₋₂₆), DC(C ₉), ODA	A	2.64						
GDA-13-I	144.21	FA(C _{10:0-28:0} C _{18:1} C _{14,15,17br}), AL(C ₁₅₋₂₆), DC(C ₇₋₁₁), APFA(C ₁₈), phy	P(v)	2.70					-26.69	-26.31
GDA-14-I	47.49	FA(C _{10:0-26:0} C _{16:1-18:1} C _{13-17,25br}), AL(C ₁₆₋₂₇), DC(C ₈₋₁₂), ODA	A	2.35					-23.97	-24.35

Table S2. Stable carbon isotopic measurements of *n*-hexadecanoic (C_{16:0}) and *n*-octadecanoic (C_{18:0}) acids of reference fats from modern authentic animal products. To facilitate comparison with archaeological data, all modern samples, including marine samples, were adjusted for the addition of the effects of post-industrial carbon (Friedli *et al.* 1986).

Category	Species	Sample type	Provenience	Mean $\delta^{13}\text{C}$ values (‰)		Reference
				C _{16:0}	C _{18:0}	
Wild non-ruminant	<i>Ursus americanus</i> (Black bear)	bone	Canada	-26.73	-27.00	This study
	<i>Castor canadensis</i> (Beaver)	soft tissue	Canada	-31.04	-31.20	This study
	<i>Lepus americanus</i> (Hare)	soft tissue	Canada	-32.15	-32.54	This study
	<i>Procyon lotor</i> (Raccoon)	soft tissue	Canada	-29.15	-28.65	This study
	<i>Ondatra zibethicus</i> (Muskrat)	soft tissue	Canada	-33.58	-33.02	This study
	<i>Lontra Canadensis</i> (Otter)	soft tissue	Canada	-31.76	-33.34	This study
Wild ruminant	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-28.13	-31.89	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-27.77	-31.54	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-28.78	-32.98	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-30.41	-34.06	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-29.49	-33.08	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-29.16	-33.43	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-30.75	-33.44	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-29.88	-33.47	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-29.31	-32.69	Craig <i>et al.</i> 2012
	<i>Cervus elaphus</i> (Red deer)	bone	Poland	-29.82	-33.41	Craig <i>et al.</i> 2012
	<i>Alces alces</i> (Moose)	bone	Canada	-29.12	-32.62	This study
	<i>Alces alces</i> (Moose)	bone	Canada	-30.27	-32.80	This study
	<i>Alces alces</i> (Moose)	soft tissue	Canada	-29.30	-32.21	This study
	<i>Alces alces</i> (Moose)	soft tissue	Canada	-29.43	-31.37	This study
	<i>Odocoileus virginianus</i> (White-tailed deer)	bone	Canada	-29.32	-31.01	This study
	<i>Odocoileus virginianus</i> (White-tailed deer)	soft tissue	Canada	-29.65	-31.51	This study
	<i>Odocoileus virginianus</i> (White-tailed deer)	soft tissue	Canada	-29.83	-30.28	This study
	<i>Rangifer tarandus</i> (Caribou)	bone	Canada	-25.40	-28.32	This study
	<i>Rangifer tarandus</i> (Caribou)	soft tissue	Canada	-24.87	-26.14	This study
	Freshwater	<i>Salmonidae</i> sp. (Salmon)	charred deposit	Japan	-28.01	-28.72
<i>Salmonidae</i> sp. (Salmon)		charred deposit	Japan	-25.22	-26.79	Craig <i>et al.</i> 2013
<i>Salmonidae</i> sp. (Salmon)		charred deposit	Japan	-26.01	-27.33	Craig <i>et al.</i> 2013
<i>Salmonidae</i> sp. (Trout)		charred deposit	Japan	-26.74	-26.72	Craig <i>et al.</i> 2013
<i>Salmonidae</i> sp. (Trout)		charred deposit	Japan	-27.64	-27.88	Craig <i>et al.</i> 2013
<i>Salmonidae</i> sp. (Trout)		charred deposit	Japan	-25.83	-26.24	Craig <i>et al.</i> 2013
<i>Rhynchocypris lagowskii</i> (Amur minnow)		charred deposit	Japan	-27.43	-28.13	Craig <i>et al.</i> 2013
<i>Pseudorasbora parva</i> (Topmouth gudgeon)		charred deposit	Japan	-26.95	-26.64	Craig <i>et al.</i> 2013
<i>Anguilla anguilla</i> (Eel)		soft tissue	Denmark	-28.96	-29.22	Craig <i>et al.</i> 2011
<i>Esox lucius</i> (Pike)		soft tissue	Denmark	-35.59	-35.84	Craig <i>et al.</i> 2011
<i>Tinca tinca</i> (Tench)		soft tissue	Denmark	-28.53	-29.60	Craig <i>et al.</i> 2011
<i>Tinca tinca</i> (Tench)		soft tissue	Denmark	-25.04	-27.14	Craig <i>et al.</i> 2011
<i>Tinca tinca</i> (Tench)		soft tissue	Denmark	-37.95	-37.27	Craig <i>et al.</i> 2011
<i>Ictalurus punctatus</i> (Channel catfish)		soft tissue	Canada	-27.41	-27.43	This study
<i>Ictalurus punctatus</i> (Channel catfish)		soft tissue	Canada	-26.07	-25.39	This study
<i>Ictalurus punctatus</i> (Channel catfish)		soft tissue	Canada	-26.71	-27.16	This study
<i>Ictalurus punctatus</i> (Channel catfish)		soft tissue	Canada	-26.15	-26.21	This study
<i>Ictalurus punctatus</i> (Channel catfish)		soft tissue	Canada	-24.75	-24.81	This study
<i>Ictalurus punctatus</i> (Channel catfish)		soft tissue	Canada	-26.04	-26.38	This study
<i>Microgadus tomcod</i> (Tomcod)		soft tissue	Canada	-34.38	-34.63	This study

	<i>Microgadus tomcod</i> (Tomcod)	soft tissue	Canada	-34.19	-33.95	This study
	<i>Microgadus tomcod</i> (Tomcod)	soft tissue	Canada	-33.86	-33.41	This study
	<i>Microgadus tomcod</i> (Tomcod)	soft tissue	Canada	-33.35	-33.28	This study
	<i>Microgadus tomcod</i> (Tomcod)	soft tissue	Canada	-32.58	-33.10	This study
Marine organisms	<i>Gymnocranius euanus</i> (Sea bream)	flesh	Japan	-22.10	-21.77	Craig <i>et al.</i> 2013
	<i>Gymnocranius euanus</i> (Sea bream)	flesh	Japan	-22.36	-22.19	Craig <i>et al.</i> 2013
	<i>Sebastes</i> sp. (Roackfish)	flesh	Japan	-23.38	-22.76	Craig <i>et al.</i> 2013
	<i>Mugil cephalus</i> (Flathead mullet)	flesh	Japan	-21.60	-20.97	Craig <i>et al.</i> 2013
	<i>Genyonemus lineatus</i> (Croaker)	flesh	Japan	-21.45	-21.09	Craig <i>et al.</i> 2013
	<i>Myoxocephalus scorpius</i> (Bull trout)	flesh	Denmark	-16.89	-17.89	Craig <i>et al.</i> 2011
	<i>Gadus morhua</i> (Atlantic cod)	flesh	Denmark	-22.73	-22.19	Craig <i>et al.</i> 2011
	<i>Gadus morhua</i> (Atlantic cod)	flesh	Denmark	-22.74	-24.12	Craig <i>et al.</i> 2011
	<i>Gadus morhua</i> (Atlantic cod)	flesh	Denmark	-22.04	-24.45	Craig <i>et al.</i> 2011
	<i>Zoarces viviparus</i> (Eelpout)	flesh	Denmark	-19.43	-21.01	Craig <i>et al.</i> 2011
	<i>Zoarces viviparus</i> (Eelpout)	flesh	Denmark	-21.15	-21.34	Craig <i>et al.</i> 2011
	<i>Platichthys flesus</i> (European flounder)	flesh	Denmark	-18.51	-19.82	Craig <i>et al.</i> 2011
	<i>Pleuronectes platessa</i> (Plaice)	flesh	Denmark	-19.81	-21.54	Craig <i>et al.</i> 2011
	<i>Pleuronectes platessa</i> (Plaice)	flesh	Denmark	-18.85	-20.07	Craig <i>et al.</i> 2011
	<i>Phoca largha</i> (Spotted seal)	blubber	Denmark	-20.00	-19.95	Craig <i>et al.</i> 2011
	<i>Phoca largha</i> (Spotted seal)	blubber	Denmark	-12.80	-14.26	Craig <i>et al.</i> 2011
	<i>Phoca vitulina</i> (Harbour seal)	blubber	Germany	-18.56	-20.21	Craig <i>et al.</i> 2011
	<i>Gadus morhua</i> (Atlantic cod)	soft tissue	Germany	-21.3	-21.70	Craig <i>et al.</i> 2011?
	<i>Clupea harengus</i> (Atlantic herring)	soft tissue	Germany	-23.2	-20.80	Craig <i>et al.</i> 2011?
	<i>Pinnipedia</i> sp. (Seal)	bone	Canada	-22.98	-24.15	This study
	<i>Pinnipedia</i> sp. (Seal)	bone	Canada	-22.16	-23.78	This study
	<i>Pinnipedia</i> sp. (Seal)	bone	Canada	-24.59	-24.28	This study

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