September 29, 2007

#### Dear Colleague:

Enclosed is the summary report of the results for the pilot exercise in the Dietary Supplement Quality Assurance (QA) Program. Included in this report are: a summary of data for all laboratories; the measurement comparability summary for evaluating laboratory performance; a summary of individual laboratory performance and interlaboratory accuracy and precision; and a summary of the NIST assigned value (NAV) vs. your laboratory value for the analytes you measured. This summary information is also displayed on a certificate. In this exercise, the NIST assigned values are the certified or reference values for the study materials and the controls, except for Standard Reference Material (SRM) 3278, where values for individual fatty acids will not be assigned in the SRM; in this case, NIST performed a single set of analyses for comparison with the results reported by study participants.

As program organizers, we have learned quite a bit from this study. We are hoping to make arrangements for an informal workshop at which we can discuss what worked and didn't work in this pilot exercise, exchange information on methodology, etc., but have not been able to do so yet. In the meantime, the QA program will be discussed by Kate Rimmer during an NIH ODS session entitled "Complying with GMPs: Tools for a QA Unit" being held at SupplySide West on the afternoon of November 6.

The Dietary Supplement Quality Assurance Program will begin in earnest with the next set of samples, which we expect to distribute in February with results due by the end of April. Prior to distribution of samples, you will be provided with a list of samples and analytes from which to choose, as you were in the pilot exercise. Either Kate Rimmer or Laura Wood will be contacting you in this regard, and Katrice Lippa will hereafter be responsible for data analysis. Dave Duewer and I will continue to be involved with the program, but Kate, Laura, and Katrice will be filling the roles that we filled during the pilot exercise.

We appreciate your participation in this pilot exercise. If you have any questions regarding this report, please contact David Duewer at 301-975-3935 or <a href="mailto:david.duewer@nist.gov">david.duewer@nist.gov</a>, or me at 301-975-3121 or <a href="mailto:katherine.sharpless@nist.gov">katherine.sharpless@nist.gov</a>.

Sincerely,

Katherine E. Sharpless, Ph.D. Research Chemist Analytical Chemistry Division Chemical Science and Technology Laboratory

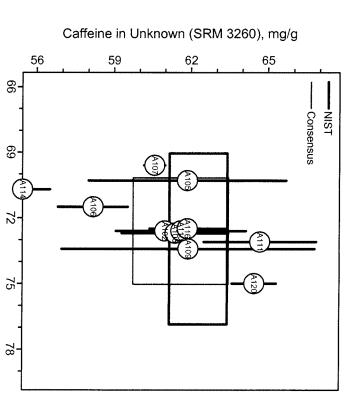
Enclosures

<ul> <li>A<sub>2</sub> Second measurement of first unit of unknown</li> <li>B<sub>1</sub> First measurement of second unit of unknown</li> </ul>	Col/Row Headers  Lab Control Unknown A <sub>1</sub> A <sub>2</sub> B <sub>1</sub> B <sub>2</sub> Mean
it of unknown	Lab Particpant code  Control Name of control material  Unknown Name of unknown material  A <sub>1</sub> First measurement of first unit of unknown  A <sub>2</sub> Second measurement of first unit of unknown
	$B_2$
	Mean
	S btwn
	Swthn
B <sub>2</sub> Second measurement of second unit of unknown  Mean Mean of each participant's replicate measurements  s <sub>bwn</sub> Participant's between-unit (sample preparation) standard deviation  s <sub>wthn</sub> Participant's within-unit (instrumental) standard deviation	S total
	z
	Average
	Median
	MADe
	%RSD
	NIST NIST value

 $\pm U_{95}$  Approximate 95% confindence interval on the NIST value

Caffeine, mg/g

±0 <sub>95</sub>	NIST	%RSD	MADe	Median	Mean	z	A120	A118	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101	Lab	
3.9	73.0	1.7	1.22	72.60	72.18	11	75.			72.5		70.7		73.1	72.6	73.42	72.7	69.6	71.5	70.3			72.6		SRM 3243	Control
							64.9			61.6		53.5		63.4	61.1	61.76	62.2	60.3	58.3	63.1			60.5		Ą	[
							64.2			61.1		51.		65.5	60.4	61.92	60.5	60.7	57.5	63.3			61.8		$A_2$	
							64.1			62.4		53.6		65.2	62.	64.38	62.	60.7	57.9	60.8			61.3		B,	Unk
							64.6			62.3		54.7		64.6	62.9	59.46	60.8	60.6	59.	60.2			60.3		<b>B</b> <sub>2</sub>	nown:
<u></u>	62.3	1.5	0.93	61.60	60.96	11	64.45			61.85		53.20		64.68	61.60	61.88	61.38	60.58	58.18	61.85			60.98		Mean	Unknown: SRM 3260
					0.75		0.			0.68		0.93		0.	1.13	0	0.	0.	0.	1.9			0		Sbtwn	60
					1.05		0.43			0.25		1.37		1.09	0.57	2.46	1.04	0.21	0.68	0.32			0.82		Swthn	
					1.29		0.43			0.73		1.65		1.09	1.27	2.46	1.04	0.21	0.68	1.92			0.82		S total	



Plot box (black) encloses ±10 % around consensus medians.

Consensus box (blue) encloses ±2×MADe around consensus medians.

Precision bars (thick black) span  $\pm 2 \times s_{\text{total}}$  about Lab mean values. NIST box (red) encloses  $\pm U_{95}$  region around NIST values.

Caffeine in Control (SRM 3243), mg/g

Lead, ng/g

NIST ±U <sub>95</sub>	%RSD	MADe	Median	Average	z	A118 A120	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101	Lab	_
660.0 53.4		32.32		656.15	2 2				657.6	614.						727.					626.		SRM 3243	Control
								***********	744.3	615.						813.					781.		Αı	
						**************************************			746.8	635.						733.					818.		$A_2$	
									729.7	645.						795.					799.		Ťα	U <sub>n</sub>
									744.3 746.8 729.7 766.8	631.						710.					824.		$B_2$	known:
738.2 8.5	5.8	43.4	754.8	736.7	4				746.9	631.5	*****					762.8					805.5		Mean	Unknown: SRM 3248
				1.6					0.							0.					0.		Mean S <sub>btwn</sub>	248
				33.2					18.6							58.4					22.3		Swthn	
				33.2					18.6	12.6						58.4					22.3		S total	
Consensus   Plot box (bla	Precision ba						65		ad	in     70		kno		n (S 50	SRI		324 300	8),		1/g 850	)		900	)
NIS I box (red) encloses ±0 <sub>95</sub> region around NIS I values.  Consensus box (blue) encloses ±2×MADe around consensus medians.  Plot box (black) encloses ±20 % around consensus medians.	Precision bars (thick black) span ±2×s <sub>total</sub> about Lab mean values.			Lead in Control (SRM 3243), ng/g	550 600 650 700 750	A114	) <del>-</del>				-				<u> </u>		(A102)	)—				Consensus	NIST	

Folic Acid, µg/g

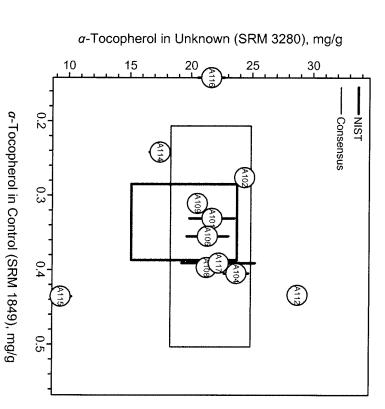
±U <sub>95</sub>	NIST	%RSD	MADe	Median		Average	z	A120	A118	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101	Lab	
0.14	2.11	12.2	0.24	2.01		2.18	10			1.99	ω	2.03	1.9		2.19	2.21	na	1.99		1.86	ļ	2.88		1.73		SRM 1849	Control
										440.	353.	626.	710.		377.	331.	419.	398.		363.	:	347		363.7		Ą	
										440.	353.	622.	540.		384.	345.	430.	417.		339.	į	372		365.8		Ą	
										440.	360.	593.	900.		373.	343.	423.	410.		353.	!	362		362.8		φ	Unl
										450.	350.	564.	820.		382.	345.	435.	425.		356.	:	374		363.7 365.8 362.8 375.2 366.9		$B_2$	(nown:
14.8	401.4	13.1	49.7	379.0		434.8	<u></u>			442.5	354.0	601.3	742.5		379.0	341.0	426.8	412.5		352.8	0	363.8		366.9		Mean s <sub>btwn</sub>	Unknown: SRM 3280
						46.8				0.	0.	30.5	152.3		0.	0	0.	0.		0	9	>		0		- 1	280
						29.8				5.0	5.0	14.6	93.9		5.7	7.1	8.1	12.1		12.1	;	130	,	ნ ა		Swthn	
						55,5				5.0	5.0	33.8	179.0		5.7	7.1	8. <u>1</u>	12.1		12.1		130		ნ ა		S total	
Plot box (black) encloses ±55 % around consensus medians	NIST box (red) encloses $\pm U_{95}$ region around NIST values. Consensus box (blue) encloses $\pm 2 \times MADe$ around consensus medians.	Precision bars (thick black) span ±2×s <sub>total</sub> about Lab mean values.			Folic Acid in Cont		1 1.5	2	00	Fo	olic	Ac	30 30		Jnk	A100		1 (S		)-	280	50 <u>'</u>		1	Consensus	NIST PAIN	
onsensus medians.	nd NIST values. around consensus medians.	about Lab mean values.			Folic Acid in Control (SRM 1849), µa/a		2 25 3							•		) (	)—			A113)							3

Retinol, mg/g

±U <sub>95</sub>	NIST	%RSD	MADe	Median		Average	z	A120 [	A118	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101	Lab	-
0.0011	0.0145	27.2	0.0043	0.0157		0.0240	<u>-</u>			0.0199	0.0064	0.0157	0.114	0.0177			0.0121	0.0128		0.0156		0.0188		0.0145	0.0165	SRM 1849	Control
										0.5	0.549	0.4	1.2	0.57			0.659	0.641		0.55		0.599		0.815	0.394	A <sub>1</sub>	
										0.54	0.547	0.479	1.18	0.534			0.653	0.625		0.52		0.658		0.825	0.379 0.344	$A_2$	
										0.59	0.573	0.443	1.22	0.565			0.652	ΝÞ		0.51 0.46		0.64		0.815 0.825 0.841 0.865 0.836 0.021 0.013 0.025	0.344	B <sub>1</sub>	Unk
										0.58	0.572	0.365	1.32	0.562			0.67	0.623				0.664 0.64		0.865	0.365	$B_2$	Unknown: SRM 3280
0.169	0.828	21.2	0.119	0.560		0.633	1			0.553	0.56	0.422	1.23	0.558			0.6590	0.628		0.51		0.64		0.836	0.37	Mean	SRM 32
						0.023				0.553 0.044 0.021	0.017	0.	0.044 0.051	0.				0.		0.029		0.		0.021	0.021	Sbtwn	280
						0.633 0.023 0.028 0.037					0.001			0.018				0.011		0.029 0.029 0.041		0.032 0.032		0.013	0.013	Swthn	
						0.037				0.048	0.017	0.056	0.067	0.018			0.009	0.011		0.041		0.032		0.025	0.025	S total	
Plot box (bl	NIST box (I	Precision b							0.		Reti	inol		Ur .4	ıkn	ow		SR ),6	M	328		, m 0.8		9		1.0	0
Plot box (black) encloses ±80 % around consensus medians.	NIST box (red) encloses $\pm U_{95}$ region around NIST values. Consensus box (blue) encloses $\pm 2 \times MADe$ around consensus medians.	Precision bars (thick black) span ±2×s <sub>total</sub> about Lab mean values.		,	Retinol in Control (SRM 1849), mg/g		0.005 0.01 0.015 0.02 0.025						A101)	A119	<b>—</b>	(A109) ~ (	(A119) (A117)	·	A109	)			(A10)		Consensus	TSIN	!

a-Tocopherol, mg/g

NIST ±U <sub>95</sub>	Median MADe %RSD	N Average	A118 A120	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101	Lab	
0.336 0.051	0.355 0.074 20.9	11 0.335		0.391	0.104	0.436	0.242	0.434			0.311	0.397		0.355		0.405		0.276	0.331	SRM 1849	Control
				20.	22.	9.97	17.	28.58			20.32	21.		22.		23.4		24.1	22.51	A <sub>1</sub>	
				23.	22.	<u> </u>	17.4	28.83			20.32 20.44 20.75 20.56 20.52	21.1		20.9		24.2		24.7	22.2	$A_2$	
				23.	21.4	9	18.	28.99			20.75	21.4		21.9		23.9		24.1	20.86 21.3	B <sub>1</sub>	Unk
				23.	21.1	9.05	17.3	28.5			20.56	21.6		20.6		23.3		24.6	21.3	<b>B</b> <sub>2</sub>	nown:
19.5 4.3	21.63 1.64 7.6	11 21.12		22.25	21.63	9.32	17.43	28.73			20.52	21.28		21.35		23.70		24.38	21.72	Mean	Unknown: SRM 3280
		0.33		0	0.52	0.2	0.14	0.			0.18	0.31		0.		0			0.88	Sbtwn	80
		0.33 0.60 0.68		1.50	0.15	0.41	0.40	0.28			0.11	0.11		0.85		0.50		0.39	0.27	Swthn	
		0.68		1.50	0.54	0.45	0.43	0.28			0.21	0.33		0.85		0.50	•••	0.39	0.92	S total	



Precision bars (thick black) span  $\pm 2 \times s_{total}$  about Lab mean values. NIST box (red) encloses  $\pm U_{95}$  region around NIST values. Consensus box (blue) encloses  $\pm 2 \times MADe$  around consensus medians. Plot box (black) encloses  $\pm 60$  % around consensus medians.

Calcium, mg/g

NIST ±U <sub>95</sub>	%RSD	MADe	Median	Average	? } } Z	A120 L	A118	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101	Lab	1
0.280 0.008	6.5	0.014	0.216	0.224	2 4					0.204		· · · · · · ·						0.223	• •••		0.209	0.258		SRM 2711	Control
										103.5								109.			112.3	124.5		Ą	
										103.1								114.			112.3 113.	132.4		$A_2$	
										103.2								107.			111.7	130.4		B <sub>1</sub>	Unk
										103.7								112.			111.7	127.3		В2	(nown:
109.2 5.2	5.9	6.53	111.34	113.67 0.31 2.77 2.79	4					103.5 103.1 103.2 103.7 103.37								110.50			111.7 111.7 112.18	128.65		Mean	Unknown: SRM 3280
				0.31	2					0.								0.			0.62	0.	1	Sbtwn	80
				2.//	1					0.30								3.54			0.35	4.24	- 1	Swthn	
				2.79	1					0.30								3.54			0.72	4.24		S total	
Cons Plot b	Preci							С	ald	ciun	n ir	ı Uı	nkr	nov	vn :	(SF	RM	32	80	), r	ng/	g			
box (re ensus t oox (bla	sion ba						80	)	1	1		10	0				120	)				140	)		
d) encloses ±U <sub>95</sub> re box (blue) encloses : ck) encloses ±35 %	rs (thick black) span		Calciulli	Calcium	0.15																		Consensus	TSIN—	
NIST box (red) encloses $\pm U_{95}$ region around NIST values. Consensus box (blue) encloses $\pm 2 \times$ MADe around consensus medians. Plot box (black) encloses $\pm 35$ % around consensus medians.	Precision bars (thick black) span ±2×s <sub>lotal</sub> about Lab mean values.		Calcium in Como (SKM Z/ 11), mg/g	in Control (SDM 37	0.2								(A118)	) -(	A103 A109	)	•							The state of the s	
es. ensus median: lians.	าก values.		711), 1119/9	144) mala	0.25							<b>L</b>					_		(A102)	)					
ò											-						···								

Iron, mg/g

Control         Uhrkmown: SRM 3280           RM 2711         A, A₂ B₁ B₂ B₁ B₂ Mean S <sub>sham</sub>	Median MADe %RSD NIST	Average	A118 A120	A117	A116	A114 A115	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101		1
Unknown: SRM 3280    B <sub>1</sub>   B <sub>2</sub>   Mean   S <sub>blvm</sub>   S <sub>withn</sub>   S <sub>total</sub>     13.4   12.9   13.03   0.   0.47   0.47     13.1   13.22   13.12   0.03   0.06   0.07     11.4   12.     11.78   0.   0.39   0.39     9.55   11.8     10.49   0.   2.22   2.22       12.10     0.01   1.15   1.15     12.40   1.00   8.1     12.2   0.9	0.256 0.030 11.9 0.281 0.006	0.263				0.274							0.237						SRM 2711	Control
Unknown: SRM 3280    B <sub>1</sub>   B <sub>2</sub>   Mean   S <sub>blvm</sub>   S <sub>withn</sub>   S <sub>total</sub>     13.4   12.9   13.03   0.   0.47   0.47     13.1   13.22   13.12   0.03   0.06   0.07     11.4   12.     11.78   0.   0.39   0.39     9.55   11.8     10.49   0.   2.22   2.22       12.10     0.01   1.15   1.15     12.40   1.00   8.1     12.2   0.9						8 39							11.6			13.07	12.5		A <sub>1</sub>	
Nknown: SRM 3280       B2     Mean     S blwn     S withn     S total       12.9     13.03     0.     0.47     0.47       13.22     13.12     0.03     0.06     0.07       12.     11.78     0.     0.39     0.39       5     11.8     10.49     0.     2.22     2.22       5     11.8     12.10     0.01     1.15     1.15       12.40     1.00       8.1       12.2     0.9					į	12 22							12.1			13.1	13.3		$A_2$	
Mean S bhwn S withn S total  13.03 0. 0.47 0.47 2 13.12 0.03 0.06 0.07  11.78 0. 0.39 0.39  10.49 0. 2.22 2.22  12.40 1.00 8.1  12.2 0.9					0	9 55							11.4			13.1	13.4		B <sub>1</sub>	둦
btwn Swthn Stotal  0. 0.47 0.47  0.03 0.06 0.07  0. 0.39 0.39  0. 2.22 2.22  0. 1.15 1.15						2 2							12.			13.22	12.9		<b>B</b> <sub>2</sub>	nown:
btwn Swthn Stotal  0. 0.47 0.47  0.03 0.06 0.07  0. 0.39 0.39  0. 2.22 2.22  0. 1.15 1.15	12.40 1.00 8.1 12.2 0.9	12 10				10 49							11.78			13.12	13.03		Mean	SRM 32
S total 0.47 0.07 0.39 1.15	6	0.01			?	)										0.03	0			280
S total 0.47 0.07 0.39 1.15	:	1 15			1	<b>&gt; &gt;&gt;</b>							0.39			0.06	0.47	- 1		
mg/g  SRM 3280), 4  Onsensus  Iron in Control (SRM 2711), mg/g  Precision bars (thick black) span ±2×s <sub>lotal</sub> about Lab mean values.  NIST box (red) encloses ±0 gs region around NIST values.  Consensus box (blue) encloses ±35 % around consensus medians.	:	1 15	·		1	) )) 							0.39			0.07	0.47		S total	
	Iron in Control (SRM 2711), mg/g  Precision bars (thick black) span ±2×S <sub>total</sub> about Lab mean values.  NIST box (red) encloses ±U 95 region around NIST values.  Consensus box (blue) encloses ±2×MADe around consensus medians.  Plot box (black) encloses ±35 % around consensus medians.	0.25	•			0	Jnk	nov		(SF					mg/	/g	<u></u>		TSIN	

Magnesium, mg/g

NIST ±U <sub>95</sub>	%RSD	MADe	Median	Average	z	A120	A118	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101	Lab	•
0.103 0.003	1.5	0.001	0.074	0.080	4					0.074								0.074			0.073	0.098		SRM 2711	Control
									•	53.05								62.3			68.38	77.8		<u> </u>	
										53.05 52.91								64.9			68.38 67.38	82.7		$A_2$	
										52.75								62.4			68.36	80.8		φ	Uni
										52.75 53.05								66.			68.36 69.03	78.2		$B_2$	known:
66.9 3.9	17.2	11.38	66.09	66.25 0.19	4					52.94								63.90			68.29			Mean	Unknown: SRM 3280
				0.19						0								0				0		S btwn	80
				1.80						0.17								2.22			0.60	2.77		Swthn	
				1.81						0.17								2.22		:	0.72	2.77		Stotal	
Consensus box (blue) encloses ±2×MADe around consensus medians.  Plot box (black) encloses ±45 % around consensus medians.	Precision bars (thick black) span ±2×s <sub>total</sub> about Lab mean values		Magnesium in Control (SRM 2711), mg/g		0.05	4(		1ag		50	ım	in I	Unl 60	410g	) -	n (\$		M 3	80		m,	. 9	90	TSIN	
sus medians. ns.	values.		711), mg/g		0 09												•		(A102)	)—	_				

Zinc, mg/g

±U <sub>95</sub>	NIST	%RSD	MADe	Median	Average	z	A120	A118	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101		7
0.005	0.341	1.3	0.004	0.332	0.328	4					0.335								0.335			0.315	0.329		SRM 2711	Control
											10.16								9.87 10.3			10.43	10.8		A <sub>1</sub>	
											10.02								10.3			10.25	<u>-1</u>		$A_2$	
											10.16 10.02 10.35 10.27								9.8			10.43 10.25 10.53 10.27	11.1		β	Ε̈́
											10.27								10.3			10.27	11.3		<b>B</b> <sub>2</sub>	(nown:
0.8	10.0	2.2	0.22	10.29	10.42	4					10.20								10.07			10.37	11.05		Mean	Unknown: SRM 3280
					0.12						10.20 0.14								0			0			S btwn	80
					0.12 0.20						0.08								0.33			0.16			Swthn	
					0.23						0.17								0.33			0.16			S total	
Plot box (black) encloses ±15 % around consensus medians.	NIST box (red) encloses $\pm U_{95}$ region around NIST values.  Consensus box (blue) encloses $\pm 2 \times MADe$ around consensus medians.	Precision bars (thick black) span ±2×s <sub>totel</sub> about Lab mean values.		E. Coling (Olaw 2711), 119/9	Zinc in Control (SRM 2711) o	0.3 0.33 0.36	•	9				n L	Jnk	10 10		(S	RM _			O), 1	mg.		.1		TSIN TSIN	

Fatty Acids in Unknown (SRM-3280), %

Sum Reported Fatty Acids	Behenic Acid [C22:0]	Gondoic Acid [C20:1]	Arachidic Acid [C20:0]	Linolenic Acid [C18:3]	Linoleic Acid [C18:2]	Vaccenic Acid [C18:1]	Oleic Acid [C18:1]	Stearic Acid [C18:0]	Margaric Acid [C17:0]	Palmitoleic Acid [C16:1]	Palmitic Acid [C16:0]	Analyte	
101.7	<0.01	Coelutes	0.430	1.789	31.57	1.304	56.34	3.395	$\sim$	0.1753	4.84	Mean	NIST
0.4		ıtes	0.014	0.068	0.66	0.046	0.83	0.061	0.0030	0.1753 0.0045	0.12	S total	] ]
94.2		0.439	0.395	1.735	27.55	2.273	51.53	3.208	0.0409	0.1283	4.86	Mean	A103
0.4		0.008	0.013	0.029	0.68	0.119	1.01	0.059	0.0020	0.0048	0.09	S total	03
96.0	0.700	0.300	0.325	1.725	28.63		55.48	3.300		<u>^0.1</u>	5.03	Mean	A1
0.1	0.000	0.000	0.050	0.050	0.13		0.27	0.000			0.05	S <sub>total</sub>	A106
92.4	0.692	0.350	0.438	1.763	26.43	1.030	51.45	3.263		0.2108	4.96	Mean	A1
0.4	0.018	0.012	0.014	0.046	0.53	0.022	0.99	0.067		0.2108 0.0119	0.11	S total	A116
5.7		0.254	0.187	2.105					0.0190	0.0776 0.010	2.84	Mean	A118
0.1		0.032	0.021	0.021					0.0025	0.0101	0.18	S total	18
	2	4	4	4	ω	2	ω	ω	2	ω	4	z	
	0.696	0.325	0.360	1.749	27.6	1.65	51.53	3.263	0.030	0.128	4.91	Median MADe	
	0.006	0.072	0.084	0.028	1.6	0.92	0.11	0.056	0.016	0.075	0.12		
	0.8	22.1	23.2	1.6	5.8	55.8	0.2	1.7	54.3	58.6	2.5	%RSD	

Palmitic Acid [C16:0], %

±U <sub>95</sub>	NIST	%RSD	MADe	Median	Ċ	Average	A120 L	A118	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101		7
0.10	1.36	1.7	0.022	1.345		1.334		1.34		1.37									1.35			1.277			SRM 3276	Control
								2.71		4.87									5.1			4.81			Ą	
								2.96		5.02									<u>ښ</u>			4.79			$A_2$	
								na		4.89									<u>.</u>			4.9			<u>~</u>	L L E
								na	***	5.05									<u>5</u>			4.94			<b>B</b> <sub>2</sub>	nown:
0.46	4.84	2.5	0.12	4.91	i	4 42		2.84		4.96									5.03			4.86			Mean	Unknown: SRM 3280
										0.									0.			0.09			S btwn	80
					-	0 05 0 11 0 12		0.18		0.11									0.05			0.02			Swthn	
					?	0 13		0.18		0.11					-,				0.05			0.09			S total	
Plot box (black) encloses ±20 % around consensus medians.	NIST box (red) encloses $\pm U_{95}$ region around NIST values. Consensus box (blue) encloses $\pm 2 \times$ MADe around consensus medians.	Precision bars (thick black) span $\pm 2 \times s_{\text{total}}$ about Lab mean values.			Palmitic Acid [C16:0] in Control (SRM 3276), %	1.1 1.2 1.3 1.4 1.5 1.	4.	Pali	miti	ic A		d [C		5:0]			1kn 5.0	ow -	n (	SR		328		), Consensus	TSIN ——	

Palmitoleic Acid [C16:1], %

NIST ±U <sub>95</sub>	MADe %RSD	Median	N Average	A118 A120	A117	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A103	• :	A101 A102	Lab	-
0.01 0.00	0.040 88.0	0.046	4 0.051	0.016	0.090	0 008								0.07		0.022	,		SRM 3276	Control
				0.085 0.07	0.203 0.210 0.2	0 205 0								<u>^</u> 0.1		0.126 0.125 0.135 0.127			Ą	
				).07	0.7.6	310								<0.1		).125	!		<del>2</del> 2	
				na	Ċ	ა ა								<u>^</u>		0.135			В	S S
				na	0.22	3 3								<u>6</u> .1		0.127			B <sub>2</sub>	known:
0.175 0.018	0.075 58.6	0.128	3 0.139	0.078	0.21	2								<u>^</u> 0.1		0.128			Mean	Unknown: SRM 3280
			0.002		Ċ	>										0.			S btwn	280
			3 0.139 0.002 0.009 0.009	0.010	210.0											0.004			Swthn	
			0.009	0.010 0.010	0.012 0.012	2										0.004 0.005			S total	
NIST box (red) encloses $\pm U_{95}$ region around NIST values. Consensus box (blue) encloses $\pm$ MADe around consensus medians. Plot box (black) encloses $\pm 90$ % around consensus medians.	Precision bars (thick black) span ±2×s <sub>total</sub> about Lab mean values.	Fairnitoleic Acid [C16:1] in Control (SRM 32/6), %	0.02 0.04 0.06	Palm	0.05		id		6: <sup>-</sup>	1] i	m L		15 [	wn		M 3		280), —— Consensus	%NIST	
κ		2/6), %	0.08																	

Stearic Acid [C18:0], %

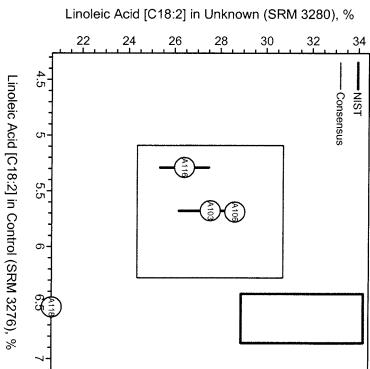
NIST ±U <sub>95</sub>	%RSD	MADe	Median	Avelage	2 2 2 2	A120 [	A118	A117	A116	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101	Lab	•
1.140 0.040	3.8	0.043	1.140	  	, , , ,		1.19		1.15									1.13			1.092			SRM 3276	Control
							na		3.21									ω ω			3.149			Ą	
							na		ယ ယ									ω ω			3.149 3.196 3.21			$A_2$	
							na		3.22									သ			3.21			₽,	_ Çr
							na		3.32									ა ა			3.279			$B_2$	hknown
3.40 0.24	1.7	0.056	3.263	3.23/	γ 2 3 3				3.263									3.300			3.208			Mean	Unknown: SRM 3280
				0.024	3				0									0						Sbtwn	280
				0.046					0.067									0.000			0.042			Swthn	
				3.237 0.024 0.046 0.052		W. Carlotte			0.067 0.067									0.000 0.000			0.04 0.042 0.059			S total	
Consensus Plot box (bl	Precision b					;	Ste		c A		-	:18 3.0	:0]	in 3.			owi	-	SR 3,6	м:	328	•		.0	
Consensus box (blue) encloses ±2×MADe around consensus medians. Plot box (black) encloses ±25 % around consensus medians.	Precision bars (thick black) span ±2×s <sub>lotal</sub> about Lab mean values. NIST box (red) encloses ± <i>U</i> or region around NIST values.			Stearic Acid [C18:0] in Control (SRM 3276), %	0.9 1 1.1 \(\frac{1.2}{2}\) 1.3 1	A118							•	(10)	(Airis		•						Consensus	TSIN ———	

Oleic Acid [C18:1], %

NIST box (red) encloses ±U <sub>95</sub> region around NIST values.  Consensus box (blue) encloses ±2×MADe around consensus medians.  Plot box (black) encloses ±25 % around consensus medians.
Precision bars (thick black) span +2×s about Lab mean values
0.83
OI 40
o g Ac
id [C 45
18:
 n L 50
Jnk
no
wn
0.27 &
RM
280
S total

Linoleic Acid [C18:2], %

NIST ±U <sub>95</sub>	Median MADe %RSD	N Average	A118 A120	A116 A117	A115	A114	A112	A111	A110	A109	A108	A107	A106	A105	A104	A103	A102	A101		1
6.640 0.220	5.684 0.297 5.2	4 5.799	6.54	5.29				****					5.69			5.677			SRM 3276	Control
			na	25.9									28.8			27.			Ą	
			na	26.6									28.6			27.15			$\frac{A}{2}$	
			na	26.2					-,,				28.5			27.15 27.89 28.17			Φ	_ G
			na	27.									28.6			28.17			В2	known:
31.6 2.6	27.55 1.59 5.8	3 27.53		26.43 0.									28.63			27.55			Mean	Unknown: SRM 3280
		0.39		0.									0.07			0.66			S btwn	.80
		0.33		0.53									0.11			0.16			Swthn	
		0.51		0.53	- <b>-</b>								0.13			0.68			S total	
Const Plot b	Preci		Line	oleic /	<b>Aci</b> c	) (	218	:2]	in	Un	ıkn	ow	n (	SR	M	328	80)	, %	)	



Precision bars (thick black) span  $\pm 2\times s_{local}$  about Lab mean values. NIST box (red) encloses  $\pm U_{95}$  region around NIST values. Consensus box (blue) encloses  $\pm 2\times MADe$  around consensus medians. Plot box (black) encloses  $\pm 25\%$  around consensus medians.