NIST IR 8163

Dietary Supplement Laboratory Quality Assurance Program: Exercise G Final Report

Melissa M. Phillips Catherine A. Rimmer Laura J. Wood

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ABSTRACT

The NIST Dietary Supplement Laboratory Quality Assurance Program (DSQAP) was established in collaboration with the National Institutes of Health (NIH) Office of Dietary Supplements (ODS) in 2007 to enable members of the dietary supplements community to improve the accuracy of measurements made in compliance with various regulations including the dietary supplement current good manufacturing practices (cGMPs). Exercise G of this program offered the opportunity for laboratories to assess their in-house measurements of nutritional elements (Na), contaminants (Pb), water-soluble vitamins (folic acid), fat-soluble vitamins (ß-carotene), and anthocyanins in foods and/or botanical dietary supplement ingredients and finished products.

INTRODUCTION

The dietary supplement industry in the U.S. is booming, with two-thirds of adults considering themselves to be supplement users.¹ Consumption of dietary supplements, which includes vitamin and mineral supplements, represents an annual US expenditure of more than \$25 billion. These figures represent an increasing American trend, and as a result, it is critically important that both the quality and safety of these products are verified and maintained.

The Dietary Supplement Health and Education Act of 1994 (DSHEA) amended the Federal Food, Drug and Cosmetic Act to create the regulatory category called dietary supplements. The DSHEA also gave the FDA authority to write current Good Manufacturing Practices (cGMPs) that require manufacturers to evaluate the identity, purity, and composition of their ingredients and finished products. In addition, the DSHEA authorized the establishment of the Office of Dietary Supplements at the National Institutes of Health (NIH ODS). To enable members of the dietary supplements community to improve the accuracy of the measurements made in compliance with these and other regulations, NIST established the Dietary Supplement Laboratory Quality Assurance Program (DSQAP) in collaboration with the NIH ODS in 2007.

The program offers the opportunity for laboratories to assess their in-house measurements of active or marker compounds, nutritional elements, contaminants (toxic elements, pesticides, mycotoxins), and fat- and water-soluble vitamins in foods as well as botanical dietary supplement ingredients and finished products. Reports and certificates of participation are provided and can be used to demonstrate compliance with the cGMPs. In addition, NIST and the DSQAP assist the ODS Analytical Methods and Reference Materials program (AMRM) at the NIH in supporting the development and dissemination of analytical tools and reference materials. In the future, results from DSQAP exercises could be used by ODS to identify problematic matrices and analytes for which an AOAC INTERNATIONAL Official Method of Analysis would benefit the dietary supplement community

NIST has experience in the area of quality assurance programs, but the DSQAP takes a unique approach. In other NIST quality assurance programs, a set of analytes is measured repeatedly over time in the same or similar matrices to demonstrate laboratory performance. In contrast, the wide range of matrices and analytes under the "dietary supplement" umbrella means that not every

¹ Walsh, T. (2012) Supplement Usage, Consumer Confidence Remain Steady According to New Annual Survey from CRN. Council for Responsible Nutrition, Washington, DC.

laboratory is interested in every sample or analyte. The constantly changing dietary supplement market, and the enormous diversity of finished products, makes repeated determinations of a few target compounds in a single matrix of little use to participants. Instead, participating laboratories are interested in testing in-house methods on a wide variety of challenging, real-world matrices to demonstrate that their performance is comparable to that of the community. In an area where there are few standard methods, the DSQAP offers a unique tool for assessment of the quality of measurements, provides feedback about performance, and can assist participants in improving laboratory operations.

This report contains the results from the seventh exercise of the DSQAP, Exercise G. Seventyeight laboratories responded to the call for participants distributed in May 2011. Samples were shipped to participants in July 2011, and results were returned to NIST by October 2011. The information and data contained in this report was disseminated to the participants in June 2011.

OVERVIEW OF DATA TREATMENT AND REPRESENTATION

Individualized data tables and certificates are provided to the participants that have submitted data in each study, in addition to this report. Examples of the data tables using NIST data are also included in each section of this report. Community tables and graphs are provided using randomized laboratory codes, with identities known only to NIST and individual laboratories. The statistical approaches are outlined below for each type of data representation.

Statistics

Data tables and graphs throughout this report contain information about the performance of each laboratory relative to that of the other participants in this study and relative to a target around the expected result, if available. The consensus mean and standard deviation are calculated according to the robust algorithm outlined in ISO 13528:2005(E), Annex C.² The algorithm is summarized here in simplified form.

Initial values of the consensus mean, x^* , and consensus standard deviation, s^* , are estimated as

x^* = median of x_i	(i = 1, 2,, n)
$s^* = 1.483 \times \text{median of } x_i - x^* $	(i = 1, 2,, n).

These initial values for x^* and s^* are updated by first calculating the expanded standard deviation, δ , as

 $\delta = 1.5 \times s^*$.

Then each x_i is compared to the expanded range and adjusted to x_i^* as described below to reduce the effect of outliers.

If $x_i < x^* - \delta$, then $x_i^* = x^* - \delta$. If $x_i > x^* + \delta$, then $x_i^* = x^* + \delta$.

² ISO 13528:2005(E), Statistical methods for use in proficiency testing by interlaboratory comparisons, pp. 14-15.

Otherwise, $x_i^* = x_i$.

New values of x^* , s^* , and δ are calculated iteratively until the process converges. Convergence is taken as no change from one iteration to the next in the third significant figure of s^* and in the equivalent digit in x^* :

$$x^* = \frac{\sum_{i=1}^n x_i^*}{n}$$

$$s^* = 1.134 \times \sqrt{\frac{\sum_{i=1}^n (x_i^* - x^*)}{n-1}}.$$

Individualized Data Table

The data in this table is individualized to each participating laboratory and is provided to allow participants to directly compare their data to the summary statistics (consensus or community data as well as NIST certified, reference, or estimated values). The upper left of the data table includes the randomized laboratory code. Tables included in this report are generated using NIST data to protect the identity and performance of participants.

Section 1 of the data table contains the laboratory results as reported, including the mean and standard deviation when multiple values were reported. A blank indicates that NIST does not have data on file for that laboratory for a particular analyte or matrix. An empty box for standard deviation indicates that only a single value was reported and therefore that value was not included in the calculation of the consensus data.²

Also in Section 1 are two Z-scores. The first Z-score, Z_{comm} , is calculated with respect to the community consensus value, using x* and s*:

$$Z_{comm} = \frac{x_i - x_*}{s_*}.$$

The second Z-score, Z_{NIST} , is calculated with respect to the target value (NIST certified, reference, or estimated value), using x_{NIST} and U_{95} (the expanded uncertainty) or s_{NIST} (the standard deviation of NIST measurements):

$$Z_{NIST} = \frac{x_i - x_{NIST}}{U_{95}}$$

or

$$Z_{NIST} = \frac{x_i - x_{NIST}}{s_{NIST}}.$$

The significance of the Z-score is as follows:

- |Z| < 2 indicates that the laboratory result is considered to be within the community consensus range (for Z_{comm}) or NIST target range (for Z_{NIST}).
- 2 < |Z| < 3 indicates that the laboratory result is considered to be marginally different from the community consensus value (for Z_{comm}) or NIST target value (for Z_{NIST}).

• |Z| > 3 indicates that the laboratory result is considered to be significantly different from the community consensus value (for Z_{comm}) or NIST target value (for Z_{NIST}).

Section 2 of the data table contains the community results, including the number of laboratories reporting more than a single value for a given analyte¹, the mean value determined for each analyte, and a robust estimate of the standard deviation of the reported values.³ Consensus means and standard deviations are calculated using the laboratory means; if a laboratory reported a single value, the reported value is not included.³ Additional information on calculation of the consensus mean and standard deviation can be found in the previous section.

Section 3 of the data table contains the target values for each analyte. When possible, the target value is a certified or reference value determined at NIST. Certified values and the associated expanded uncertainty (U_{95}) have been determined with two independent analytical methods at NIST, by collaborating laboratories, or in some combination. Reference values are assigned using NIST values obtained from the average and standard deviation of measurements made using a single analytical method or by measurements obtained from collaborating laboratories. For both certified and reference values, at least six samples have been tested and duplicate preparations from the sample package have been included, allowing the uncertainty to encompass variability due to inhomogeneity within and between packages. For samples in which a NIST certified or reference value is not available, the analytes are measured at NIST using an appropriate method. The NIST-assessed value represents the mean of at least three replicates. For materials acquired from another proficiency testing program, the consensus value and uncertainty from the completed round is used as the target range.

Summary Data Table

This data table includes a summary of all reported data for a particular analyte in a particular study. Participants can compare the raw data for a single laboratory to data reported by the other participating laboratories or to the consensus data. A blank indicates that the laboratory signed up and received samples for that particular analyte and matrix, but NIST does not have data on file for that laboratory.

Graphs

Data Summary View (Method Comparison Data Summary View)

In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). Data points that are unfilled represent laboratories that reported a single value for that analyte and therefore were not included in the consensus mean. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. Where appropriate, two consensus means may be calculated for the same sample if bimodality is identified in the data. In this case, two consensus means and ranges will be displayed in the data summary view. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified, reference, or estimated value bounded by twice its uncertainty (U_{95}) or standard deviation. For the purpose of the DSQAP, a target range spanning twice the uncertainty in the NIST value is selected because participants are only asked to make a limited number of

³ ISO 13528:2005(E), Statistical methods for use in proficiency testing by interlaboratory comparisons, Annex C.

observations. The size of the y-axis on the data summary view graph represents the consensus mean bounded by 2δ . In this view, the relative locations of individual laboratory data and consensus zones with respect to the target zone can be compared easily. In most cases, the target zone and the consensus zone overlap, which is the expected result. The major program goals are to reduce the size of the consensus zone and center the consensus zone about the target value. Analysis of an appropriate reference material as part of a quality control scheme can help to identify sources of bias for laboratories reporting results that are significantly different from the target zone. In the case in which a method comparison is relevant, different colored data points may be used to indicate laboratories that used a specific approach to sample preparation, analysis, or quantitation.

Sample/Sample Comparison View

In this view, the individual laboratory results for one sample (NIST SRM with a certified or reference value) are compared to the results for another sample (another NIST SRM with a more challenging matrix, a commercial sample, etc.). The error bars represent the individual laboratory standard deviation. The solid red box represents the target zone for the first sample (x-axis) and the second sample (y-axis). The dotted blue box represents the consensus zone for the first sample (x-axis) and the second sample (y-axis). The axes of this graph are centered about the consensus mean values for each sample or control, to a limit of zero and twice the consensus mean. Depending on the variability in the data, the axes may be scaled proportionally to better display the individual data points for each laboratory. In some cases, when the consensus and target ranges have limited overlap, the solid red box may only appear partially on the graph. If the variability in the data is high (greater than 100 % relative standard deviation (RSD)), the dotted blue box may also only appear partially on the graph. This view emphasizes trends in the data that may indicate potential calibration issues or method biases. One program goal is to identify such calibration or method biases and assist participants in improving analytical measurement capabilities. In some cases, when two equally challenging materials are provided, the same view (sample/sample comparison) can be helpful in identifying commonalities or differences in the analysis of the two materials.

NUTRITIONAL ELEMENTS (Na) IN POWDERED MATRICES

Study Overview

In this study, participants were provided with two NIST SRMs, SRM 1573a Tomato Leaves and SRM 3281 Cranberry (Fruit). Participants were asked to use in-house analytical methods to determine the mass fraction of sodium in each matrix and report values on an as-received basis.

Sample Information

Tomato Leaves. Participants were provided with one bottle containing approximately 50 g of dried, powdered tomato leaves. The material was prepared from tomato leaves collected in Pennsylvania that had been picked from an untreated area, rinsed, then dried at 60 °C to 70 °C. The dried leaves were ground and sieved prior to packaging. Before use, participants were instructed to thoroughly mix the contents of the bottle and use a sample size of at least 0.5 g. Participants were asked to store the material at controlled room temperature, 20 °C to 25 °C, and to prepare three samples and report three values from the single bottle provided. Prior to the study, the approximate analyte level was given as 100 mg/kg to 200 mg/kg. The certified value for sodium in SRM 1573a was determined at NIST using instrumental neutron activation analysis (INAA) and flame atomic emission spectrometry (FAES). The certified values and uncertainties for Na are provided in the table below, both on a dry-mass basis and on an as-received basis accounting for moisture of the material (0.97 %).

	Certified Mass Fraction	in SRM 1573a (mg/kg)
	<u>(dry-mass basis)</u>	(as-received basis)
Sodium (Na)	136 ± 4	132 ± 4

Cranberries. Participants were provided with one packet containing approximately 6 g of freezedried, powdered cranberries. The cranberry powder was blended, aliquotted, and heat-sealed inside 4 mil polyethylene bags, which were then sealed inside nitrogen-flushed aluminized plastic bags along with two packets of silica gel. Before use, participants were instructed to thoroughly mix the contents of each packet and use a sample size of at least 0.5 g. Participants were asked to store the material at controlled room temperature, 20 °C to 25 °C, and to prepare three samples and report three values from the single packet provided. Prior to the study, the approximate analyte level was given as 200 mg/kg to 300 mg/kg. The reference value for sodium in SRM 3281 Cranberry (Fruit) was determined at NIST using inductively coupled plasma optical emission spectroscopy (ICP-OES). The reference values and uncertainties for Na are provided in the table below, both on a dry-mass basis and on an as-received basis accounting for moisture of the material (2.39 %).

	Reference Mass Fraction	on in SRM 3281 (mg/kg)			
	<u>(dry-mass basis)</u>	(as-received basis)			
Sodium (Na)	259 ± 3	253 ± 3			

Study Results

• Thirty-seven laboratories enrolled in this study and received samples. Twenty-four laboratories reported results for both the samples (65 % participation).

- For both samples, the consensus ranges were wide but encompassed the NIST target ranges.
 - The consensus mean for sodium in tomato leaves was on the upper edge of the target range.
 - The consensus mean for sodium in the cranberries was above the target range.
- The between-laboratory variability for was high in both materials (29 % and 28 % RSD for the tomato leaves and cranberries, respectively).
- A majority of the laboratories reported using either open-beaker digestion (48 %) or microwave digestion (32 % to 36 %) for sample preparation. The remaining laboratories reported using hot block digestion, dry ashing, or dilution. One laboratory did not report the type of sample preparation technique that was used.
- A majority of the laboratories reported using either ICP-OES (56 %) or ICP-MS (28 %) as their analytical method. Two laboratories reported using atomic absorption spectroscopy, and one laboratory reported using ion chromatography with conductivity detection. One laboratory did not report the type of analytical technique that was used.
- A majority of the laboratories reported using an external standard approach to calibration (88 %). One laboratory reported using a standard addition approach, and one laboratory reported using an internal standard approach. One laboratory did not report the type of calibration approach that was used.

Technical Recommendations

The following recommendations are based on results obtained from the participants in this study.

- A significant difference was apparent between results obtained using open beaker and those using microwave digestion. Too few results were reported by other methods to identify any additional trends.
 - As shown in **Figure 1** and **Figure 2**, laboratories that reported high sodium values also reported using open beaker digestion. This trend may indicate contamination of samples with sodium from the environment during digestion.
 - To minimize contamination, work areas should be cleaned prior to sample preparation, glassware should be cleaned by acid wash prior to use, and exposure of samples to the laboratory environment should be limited.
 - If a soap solution is used for cleaning, sodium levels in blank solutions must be checked, as some soaps will give high sodium blank levels.
 - Extra procedural reagent blanks should be prepared along with samples to know the extent of sodium contamination from the analysis.
- No difference was apparent in the sodium results based on analytical method used (ICP-OES or ICP-MS). Too few results were reported by other methods to identify any additional trends.
- Additional sources for potential error in the final results may be errors in calibration and dilution.
- Samples with very low sodium levels may be inadvertently diluted below the calibration range and possibly out of the quantification or detection range of the instrument. Care should be taken to evaluate a more concentrated sample if the signal appears to be lower than expected or is below the lowest calibration point.

- Many of the calibration curves reported by participants extended far above the reported working range of sodium in the solutions to be analyzed. While the calibration curve may appear linear over a wide concentration range, a measured value near an end of the calibration curve may have significant bias. For the most accurate results, use calibration points closely surrounding the expected solution concentration, and ensure that the calibration curve is linear along the region of your expected solution concentrations.
 - The sample/sample comparison graph does not reflect an obvious calibration error. Some laboratories reported values that were high for one sample but were within range for the second sample. This type of trend may indicate individualized problems with the digestion of one matrix compared to another.
 - Quality assurance samples should always be used. These can be commercially available reference materials (CRMs, SRMs, or RMs) or prepared in-house, but need to be of known concentration.
 - They are used to ensure that the method is performing as expected.
 - They are useful in finding where errors are occurring, including calculation errors.
 - After checking for calculation errors, make sure results are reported correctly.

Table 1. Individual data table (NIST) for sodium in foods and dietary supplements.

National Institute of Standards & Technology

	Lab Code:	NIST		1. Your	Results		2. Co	mmunity]	Results	3. Targe	et Value
Analyte	Sample	Units	Mean	s total	Z _{comm}	Z _{NIST}	Ν	Mean	Std Dev	Value	U_{95}
Na	Tomato Leaves	mg/kg	132	3.9	-0.1	0.1	24	134	40	132	3.9
Na	Cranberry (Fruit)	mg/kg	253	2.9	-0.2	0.1	24	265	73	253	2.9

Exercise G - July 2011 - Nutritional Elements

Mean Average of reported values

s_{total} Standard deviation of reported values

Z_{comm} Z-score: (Lab Mean - Consensus Mean)/ Consensus Standard Deviation

Z_{NIST} Z-score: (Lab Mean - NIST Value or Label Claim)/ NIST or Label Claim Standard Deviation N Number of quantitative values reported

Mean Robust mean of the reported values

Std Dev Robust standard deviation

Value NIST-assessed value

 $U_{95} \pm 95\%$ confidence interval about the assessed value

NR No data reported

						So	odium				
		SRN	A 1573a T	lomato Le	aves (mg	/kg)	SRM 3281 Cranberry (Fruit) (mg/kg)				
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST				132	4				253	3
	G701	80	79	77	79	1	210	215	215	213	3
	G703	113	106	111	110	4	139	146	136	140	5
	G704										
	G709										
	G710	112	113	116	114	2	233	234	240	236	4
	G711	103	106	103	104	1	214	214	148	192	38
	G714	151	140	133	142	9	231	223	243	232	10
	G720										
	G721	707	391	452	517	167	911	902	889	901	11
	G723	119	116	118	117	2	232	230	224	229	4
	G724	130	131	142	134	7	260	252	248	253	6
	G728										
	G729										
	G734	219	220	218	219	1	572	573	554	566	11
s	G735	157	162	166	162	5	418	410	432	420	11
Individual Results	G736	194	181	187	187	6	403	436	416	418	17
Re	G737	163	163	169	165	3	502	491	485	493	9
dua	G738	138	138	139	138	1					
divid	G739										
Ine	G741	107	109	108	108	1	240	238	239	239	1
	G742										
	G743	180	175	200	185	13	299	286	296	294	7
	G747	305	245	271	274	30	228	229	225	227	2
	G749	116	130		123	9	294	314		304	14
	G752	122	122	117	120	3	287	257	263	269	16
	G754										
	G755										
	G756						260	233	246	246	14
	G757	111	113	110	111	2	223	232	230	228	5
	G758 G759	122	122	118	121	2	263	262	265	263	1
	G759 G761	122	122	110	121	2	205	202	203	205	1
	G762	113	113	112	113	1	237	237	238	237	1
	G763	105	100	102	102	2	230	232	230	231	1
	G764										
	G770 G778	128	123	120	124	4	183	184	181	183	1
y	0//0	Consensu	s Mean		136		Consensu	s Mean		268	
Community Results				Deviation					l Deviation		
ommun Results		Maximum			517		Maximum			901	
Cor R		Minimum			79		Minimum			140	
		Ν			23		Ν			23	

Table 2. Data summary table for sodium in foods and dietary supplements.

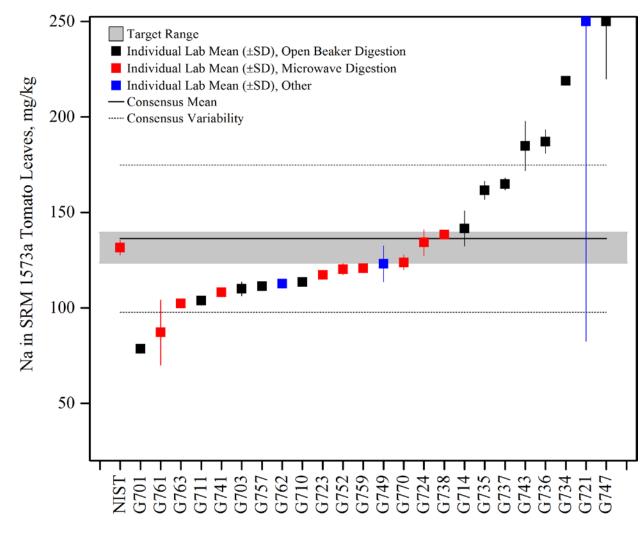


Figure 1. Sodium in SRM 1573a Tomato Leaves (data summary view – digestion method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The color of the data point represents the sample preparation (digestion) procedure employed. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST reference value bounded by twice its uncertainty (U_{95}).

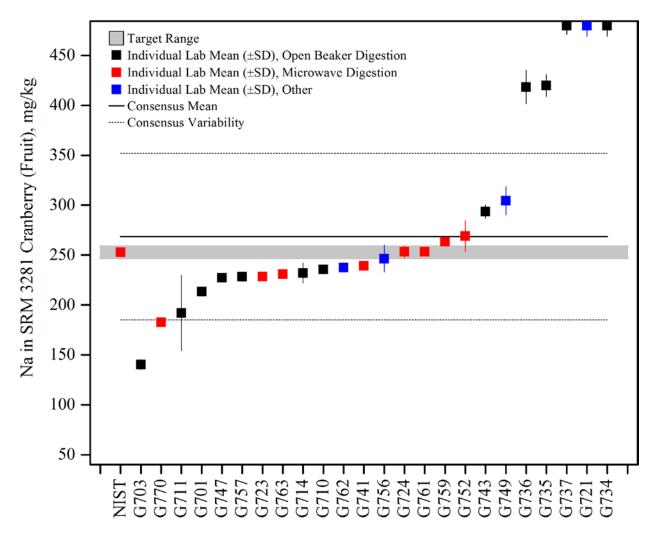


Figure 2. Sodium in SRM 3281 Cranberry (Fruit) (data summary view – digestion method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by digestion method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}).

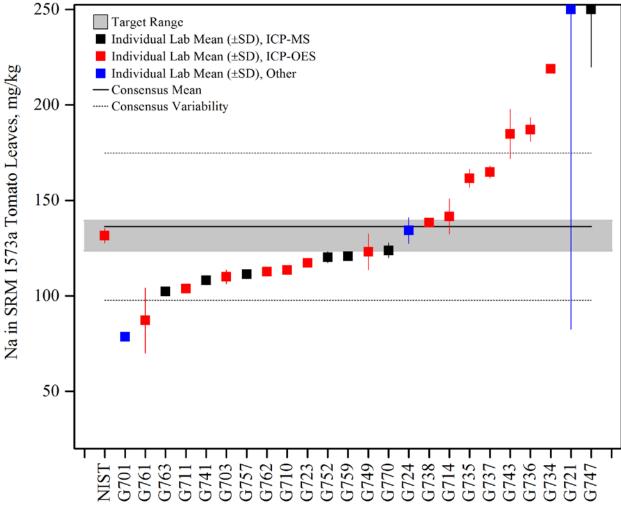


Figure 3. Sodium in SRM 1573a Tomato Leaves (data summary view – instrumental methods). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded twice by its uncertainty (U_{95}).

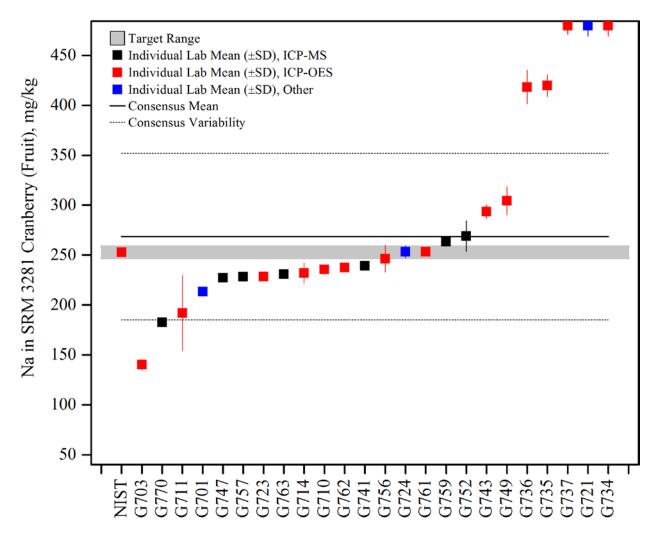


Figure 4. Sodium in SRM 3281 Cranberry (Fruit) (data summary view – instrumental methods). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}).

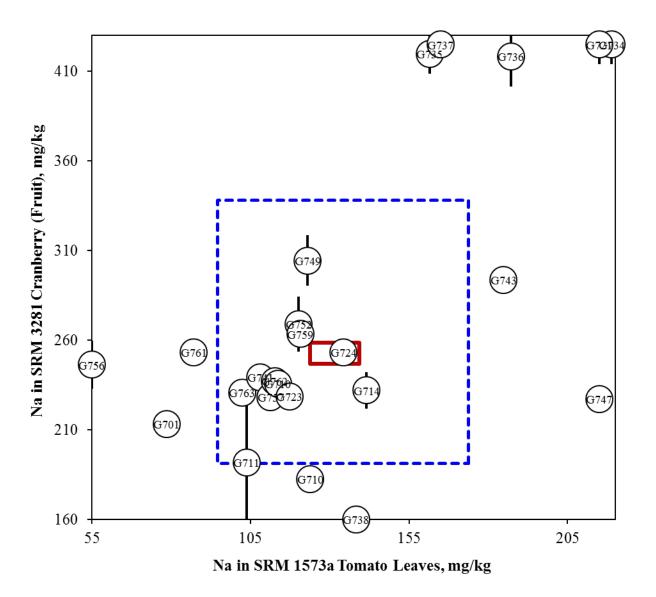


Figure 5. Sodium in SRM 1573a Tomato Leaves and SRM 3281 Cranberry (Fruit) (sample/sample comparison view). In this view, the individual laboratory results for one sample (tomato leaves) are compared to the results for a second sample (cranberry). The solid red box represents the target zone for the two samples, tomato leaves (x-axis) and cranberry (y-axis). The dotted blue box represents the consensus zone for tomato leaves (x-axis) and cranberry (y-axis).

TOXIC ELEMENTS (Pb) IN DIETARY SUPPLEMENTS

Study Overview

In this study, participants were provided with two NIST SRMs, SRM 3243 Ephedra-Containing Solid Oral Dosage Form and SRM 3280 Multivitamin/Multielement Tablets. Participants were asked to use in-house analytical methods to determine the mass fractions of lead (Pb) in each of the matrices and report values on an as-received basis.

Sample Information

Ephedra-Containing Tablets. Participants were provided with three bottles containing approximately 2.5 g of dried, powdered ephedra-containing tablets. The ephedra tablets were ground, homogenized, and packaged in amber high-density polyethylene bottles. Before use, participants were asked to thoroughly mix the contents of the bottles and use a sample size of at least 1.0 g. Participants were asked to store the material at controlled room temperature, 20 °C to 25 °C, and to prepare one sample and report one value from each bottle provided. Prior to the study, the approximate analyte level was given as 600 ng/g to 800 ng/g. The certified value for lead in SRM 3243 was determined at NIST using isotope dilution inductively coupled plasma mass spectrometry (ID-ICP-MS), in combination with data from two collaborating laboratories. The certified value and uncertainty for Pb are provided in the table below, both on a dry-mass basis and on an as-received basis accounting for moisture of the material (4.63 %).

	Certified Mass Fraction in SRM 3243 (ng/g)					
<u>Analyte</u>	<u>(dry-mass basis)</u>	(as-received basis)				
Lead (Pb)	692 ± 56	660 ± 53				

Multivitamin/Multielement Tablets. Participants were provided with three vials containing approximately 2.5 g of ground multivitamin/multielement tablets. The multivitamin/multielement tablets were ground, sieved, and packaged in amber glass vials. Before use, participants were asked to thoroughly mix the contents of the vial and to use a sample size of at least 0.25 g. Participants were asked to store the material at controlled room temperature, 20 °C to 25 °C, and to prepare one sample and report one value from each of the vials provided. Prior to the study, the approximate analyte level was given as 200 ng/g to 300 ng/g. The certified value for lead in SRM 3280 was determined at NIST using isotope dilution inductively coupled plasma mass spectrometry (ID-ICP-MS). The certified values and uncertainties are reported in the table below, both on a dry-mass basis and on an as-received basis accounting for moisture of the material (1.37 %).

	Certified Mass Fraction	on in SRM 3280 (ng/g)
<u>Analyte</u>	(dry-mass basis)	(as-received basis)
Lead (Pb)	273 ± 2	269 ± 2

Study Results

- Forty-eight laboratories enrolled in this exercise and received samples. Thirty-six laboratories reported results for lead in Ephedra tablets (75 % participation). Thirty-seven laboratories reported results for lead in multivitamin tablets (77 % participation).
- The consensus means for lead in both matrices were within the target ranges. The betweenlaboratory variability for lead in the Ephedra tablets was acceptable (13 % RSD), but variability for lead in the multivitamin tablets was high (23 % RSD).
- A majority of the laboratories reported using microwave digestion (58 %) or open beaker digestion (36 %) for sample preparation. Hot block digestion was also reported as a method of sample preparation by two laboratories (6 %). One laboratory did not report the sample preparation technique used.
- Most laboratories reported using ICP-MS as their analytical method for analysis (78 %). Laboratories also reported using ICP-OES (17 %) and AAS (6 %). One laboratory did not report the analytical method used.
- A majority of the laboratories reported using an external standard approach to calibration (89 %). One laboratory reported using a standard addition approach (6 %), and one laboratory reported using an internal standard approach (6 %). One laboratory did not report the type of calibration approach that was used.

Technical Recommendations

The following recommendations are based on results obtained from the participants in this study.

- Lead is generally easily digested, and digestion with HNO₃ is recommended. Digestion with HCl may form a PbCl₂ precipitate that is difficult to dissolve. While the sample solution may look clear, results may be biased low if solid PbCl₂ remains. The Cl can be removed by repeatedly drying the sample solution using HNO₃.
- Some laboratories reported high values for one sample, but not both, which may indicate more difficulty in digestion of one matrix over the other. Because each matrix is different, some may digest more readily, and the digestion quality must be evaluated for each sample prior to analysis. A quality assurance sample of a similar matrix prepared alongside unknown samples will help to determine if the digestion method is appropriate for the determination of lead in these matrices.
- ICP-OES has low sensitivity for lead, making it difficult to measure low-level samples. The concentration of lead in solutions prepared from the multivitamin samples may be near the method detection limit of many ICP-OES systems, leading to inaccurate results. Sufficient procedural reagent blanks should be prepared along with samples to determine the method detection limit for lead.
- Many of the calibration curves reported by participants extended far below the reported working range of lead in the solutions to be analyzed. While the calibration curve may appear linear over a wide concentration range, a measured value outside of the calibration curve may have significant bias. For the most accurate results, use calibration points closely surrounding the expected solution concentration, and ensure that the calibration curve is linear along the region of your expected solution concentrations.
 - The sample/sample comparison graph does not reflect an obvious calibration error. Some laboratories reported values that were high for one sample but were within range for the second sample. This type of trend may indicate individualized problems with the digestion of one matrix compared to another.

- Quality assurance samples should always be used. These can be commercially available reference materials (CRMs, SRMs, or RMs) or prepared in-house, but need to be of known concentration.
 - They are used to ensure that the method is performing as expected.
 - They are useful in finding where errors are occurring, including calculation errors.
 - After checking for calculation errors, make sure results are reported correctly.

Table 3. Individual data table (NIST) for lead in dietary supplements.

National Institute of Standards & Technology

	Lab Code:	NIST		1. Your I	Results		_	2. Co	mmunity]	3. Target Value		
Analyte	Sample	Units	Mean	s total	Z _{comm}	Z _{NIST}		Ν	Mean	Std Dev	Value	U_{95}
Pb	Ephedra Tablets	ng/g	660	53	0.4	0.0		36	626	84	660	53
Pb	Multivitamin Tablets	ng/g	269	2.4	-0.3	0.0	_	36	291	66	269	2

Exercise G - July 2011 - Toxic Elements

Mean Average of reported values

 s_{total} Standard deviation of reported values Z_{comm} Z-score: (Lab Mean - Consensus Mean)/ Consensus Standard Deviation Z_{NIST} Z-score: (Lab Mean - NIST Value or Label Claim)/ NIST or Label Claim Standard Deviation N Number of quantitative values reported

Mean Robust mean of the reported values

Std Dev Robust standard deviation

Value NIST-assessed value

 $U_{95} \pm 95\%$ confidence interval about the assessed value

NR No data reported

ata s	umma	ary tabl	e for l	lead in	dietar	y supp	olemen	ts.			
							ead				
	-	SRM 3243 Ephedra Tablets (ng/g) SRM 3280 Multivitamin						Tablets (ng/g)		
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST				660	53				269	2
	G703	661	678	621	653	29	279	288	275	281	7
	G706	690	717	728	712	20	228	232	240	233	6
	G709	636	641	627	634	7	265	253	273		10
	G710	665	644	584	631	42	282	277	277	279	3
	G711										
	G713	620	649	624	631	16	271	283	280		6
	G714						219	235	341		66
	G718	587	601	549	579	27	319	297	269		25
	G719	500	559	549	536	32	210	225	214	216	8
	G720										
	G723	704	687	664	685	20	226	229	226	227	2
	G724	711	681	674	689	20	258	257	252	256	3
	G725	682	615	614	637	39	279	288	287	285	5
	G726	711	672	693	692	20	324	328	292	315	20
	G728										
	G729										
	G734	472	466	505	481	21	616	627	658	634	22
	G735	538	569	522	543	24	618	554	554	576	37
	G736	659	489	553	567	86	769	818	811	799	27
Individual Results	G737	960	678	946	861	159	657	628	638	641	15
	G738	670	610	680	653	38	290	280	290	287	6
	G739										
idu	G742										
vibu	G743	531	546	524	534	11	233	230	225	230	4
П	G745	667	560	576	601	58	275	285	272	277	7
	G746	614	614	614	614	0	326	370	360	352	23
	G747	606	596	573	592	17	242	245	222	236	13
	G748	699	644	736	693	46	299	200	333	277	69
	G749	1610		3400	2505	1266	530			530	
	G750	757	658	634	683	65	211	208	213	211	3
	G751						267		267		2
	G752										89 12
	G754 G755	552	302	545	540	15	320	303	324	519	12
	G756	574	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	238	5						
	G757	647	601	614	621	24	249	257	249	252	5
	G758										
	G759										0
	G762 G763										6 33
	G764	000	027	045	020	19	505	321	299	330	55
	G766	549	538	567	551	15	239	246	238	241	4
	G769									437	42
	G770									300	2
	G771										5
	G773 G774	929	//8	//4	827	88	592	357	443	397	43
	G778										
Ŷ	2.70	Consensus	Mean		626		Consensu	is Mean		256 285 315 634 576 799 641 287 230 277 352 236 277 530 211 268 727 319 238 252 280 283 330 241 437 300 89 397 291	
Community Results				l Deviation	84				1 Deviation		
ommun Results		Maximum			2505		Maximun	1			
Cor R		Minimum			273		Minimum				
		Ν			36		Ν			36	

 Table 4. Data summary table for lead in dietary supplements.

		Lead										
		SRI	EphedraTa	hlets (no		Tablets (ησ/σ)					
1	Lab	A	B	С	Avg	SD	A	B	С			
	NIST		D	C	660	53		D	C	269	2	
	G703	661	678	621	653	29	279	288	275	281	7	
	G706	690	717	728	712	20	228	232	240	233	6	
	G709	636	641	627	634	7	265	252	273	264	10	
	G710	665	644	584	631	42	282	255	275	279	3	
	G711	005	044	504	031	42	202	211	211	21)	5	
	G713	620	649	624	631	16	271	283	280	278	6	
	G714	020	047	024	0.51	10	219	235	341	265	66	
	G718	587	601	549	579	27	319	297	269	205	25	
	G719	500	559	549	536	32	210	225	214	216	8	
	G720	500	557	547	550	52	210	225	214	210	0	
	G723	704	687	664	685	20	226	229	226	227	2	
	G723	704	681	674	689	20	258	229	220	256	3	
	G724	682	615	614	637	39	279	288	232	285	5	
	G725	711	672	693	692	20	324	328	287	315	20	
	G728	/11	072	093	092	20	324	328	292	515	20	
	G728 G729											
		472	466	505	481	21	616	627	650	624	22	
	G734 G735	472 538	400 569	505 522	543	21	616 618	627 554	658 554	634 576	37	
			489							799		
Individual Results	G736	659		553	567 861	86	769	818	811		27	
	G737	960	678	946	861	159	657	628	638	641	15 6	
	G738	670	610	680	653	38	290	280	290	287	0	
	G739 G742											
	G742 G743	521	546	524	524	11	222	230	225	220	4	
Inc	G745	531 667	560	524 576	534 601	11 58	233 275	285	225 272	230 277	7	
	G745 G746	614	614	614	614	0	326	370	360	352	23	
	G740	606	596	573	592	17	242	245	222	236	13	
	G748	699	644	736	693	46	299	245	333	230	69	
	G749	1610	044	3400	2505	1266	530	200	333	530	09	
	G750	757	658	634	683	65	211	208	213	211	3	
	G751	614	689	589	631	52	267	270	267	268	2	
	G752	1042	718	849	870	163	649	706	824	727	89	
	G754	532	562	543	546	15	328	305	324	319	12	
	G755 G756	574	534	521	543	28	233	239	243	238	5	
	G757	647	601	614	621	20	233	257	243	252	5	
	G758			-	-					-	-	
	G759	637	646	642	642	4	281	280	280	280	0	
	G762	640	630	650	640	10	280	280	290	283	6	
	G763 G764	606	627	645	626	19	365	327	299	330	33	
	G764 G766	549	538	567	551	15	239	246	238	241	4	
	G769	163	153	504	273	200	467	456	389	437	42	
	G770	667	689	689	682	13	298	302	300	300	2	
	G771	421	507	391	440	60	94	91	83	89	5	
	G773	929	778	774	827	88	392	357	443	397	43	
	G774 G778											
y	0,70	Consensus	Mean		626		Consensu	s Mean		291		
Community Results		Consensus		d Deviation	84		Consensu	s Standard	l Deviation			
ommun Results		Maximum			2505		Maximum			799		
C01 R		Minimum			273		Minimum			89 26		
		N			36		Ν			36		

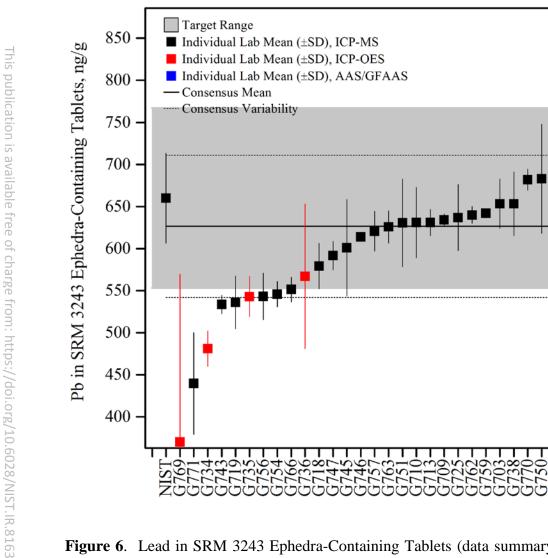


Figure 6. Lead in SRM 3243 Ephedra-Containing Tablets (data summary view – instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}) .

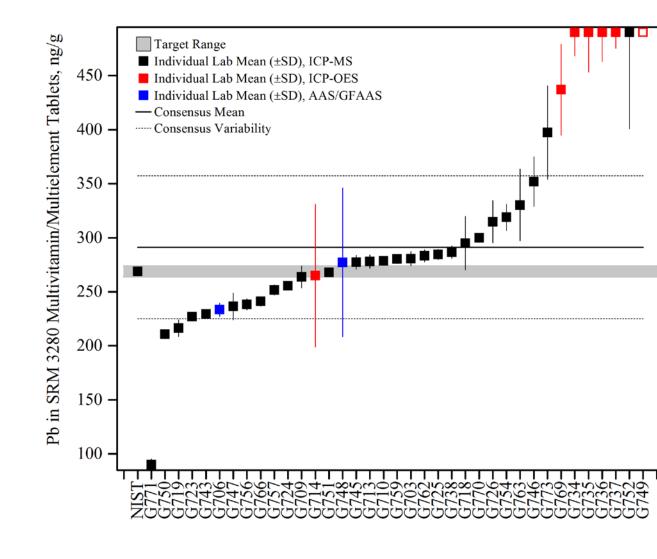


Figure 7. Lead in SRM 3280 Multivitamin/Multielement Tablets (data summary view – instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}).

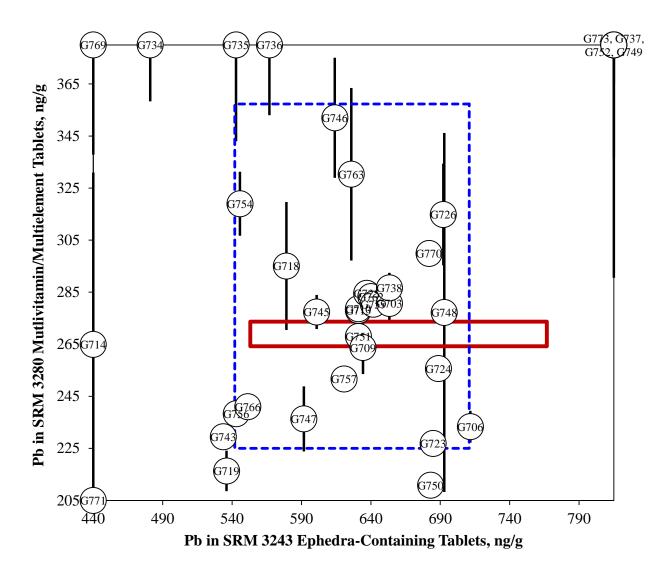


Figure 8. Lead in SRM 3243 Ephedra-Containing Tablets and SRM 3280 Multivitamin/Multielement Tablets (sample/control comparison view). In this view, the individual laboratory results for the one sample (SRM 3243 Ephedra-Containing Tablets) are compared to the results for a second sample (SRM 3280 Multivitamin/Multielement Tablets). The solid red box represents the target zone for the two samples, ephedra tablets (x-axis) and multivitamin (yaxis). The dotted blue box represents the consensus zone for the ephedra tablets (x-axis) and the multivitamin (y-axis).

FOLIC ACID IN FOOD MATRICES

Study Overview

In this study, participants were provided with one NIST SRM, SRM 3233 Fortified Breakfast Cereal, and a well-characterized fortified milk powder sample. Participants were asked to use inhouse analytical methods to determine the mass fraction of folic acid in each of the matrices and report values on an as-received basis.

Sample Information

Fortified Breakfast Cereal. Participants were provided with one bottle containing approximately 60 g of dried, powdered breakfast cereal. The cereal flakes were ground, homogenized, and packaged in amber glass bottles, and the bottles were capped, sealed with heat-shrink tape, and individually sealed in mylar bags. Before use, participants were instructed to thoroughly mix the contents of the bottle and use a sample size of at least 1.0 g. Participants were asked to store the material at controlled room temperature, 20 °C to 25 °C, and to prepare three samples and report three values from the single bottle provided. Prior to the study, the approximate analyte level was given as 15 mg/kg. The certified value for folic acid in SRM 3233 was determined at NIST by ID-LC-MS/MS following solvent extraction, in combination with data from numerous collaborating laboratories. The certified values and uncertainties are reported in the table below, both on a dry-mass basis and on an as-received basis accounting for moisture of the material (1.7 %).

	Certified Mass Fraction in S	RM 3233 (mg/kg)
<u>Analyte</u>	(dry-mass basis)	(as-received basis)
Folic Acid	15.1 ± 1.2	14.8 ± 1.2

Milk Powder. Participants were provided with three packets containing approximately 10 g of freeze-dried, powdered milk. The fortified milk powder was homogenized and packaged in nitrogen-flushed foil pouches with a paper over-wrap. Before use, participants were instructed to thoroughly mix the contents of the packet and use a sample size of at least 1.0 g. Participants were asked to store the material at -20 °C, and to prepare one sample and report one value from each packet of milk powder provided. Prior to the study, the approximate analyte level was given as 2.4 mg/kg. The target value for folic acid in the fortified milk powder was determined at NIST by ID-LC-MS/MS following solvent extraction and hydrolysis, in combination with data from two collaborating laboratories. The NIST-determined values and uncertainties are reported in the table below, both on a dry-mass basis and on an as-received basis accounting for moisture of the material (1.6 %).

	Mass Fraction in Mill	Milk Powder (mg/kg)				
<u>Analyte</u>	<u>(dry-mass basis)</u>	(as-received basis)				
Folic Acid	$2.35 ~\pm~ 0.06$	$2.31 \hspace{.1in} \pm \hspace{.1in} 0.06$				

Study Results

• Twenty-three laboratories enrolled in this exercise and received samples. Eleven laboratories reported results for folic acid in both the breakfast cereal and the milk powder

(48 % participation). One laboratory reported a single value for folic acid in the breakfast cereal and was therefore excluded from calculation of summary statistics.²

- The consensus ranges for folic acid in both matrices were wide but contained the target ranges. The consensus means for folic acid in both matrices were slightly above the target ranges.
 - For folic acid in the breakfast cereal, the between-laboratory variability was high (32 % RSD).
 - For folic acid in the milk powder, the between-laboratory variability was acceptable (15 % RSD).
- A majority of the laboratories reported using an enzymatic hydrolysis approach (50 %). A solvent extraction approach (25 %) and a shaking or sonication extraction approach (25 %) were also reported as methods of sample preparation.
- Most laboratories reported using LC with absorbance detection as the analytical method for analysis (67 %). Laboratories also reported using microbiological assay (17 %), LC-MS (8 %), and protein binding assay (8 %) as the instrumental approach.
- All participating laboratories reported the use of an external standard approach to quantitation.

Technical Recommendations

The following are recommendations based on results obtained from the participants in this study.

- No sample preparation approach or analytical method was identified as exceptionally good or problematic.
- Some high values were reported for either the sample or control (but not both). This may indicate chromatographic coelutions with matrix components. Because each matrix is different, the chromatographic method should be evaluated to confirm any potential biases from coeluting matrix components.
- NIST values in SRM 3233 and the fortified milk powder were determined using a gentle extraction procedure designed for determination of fortified folic acid in foods. As a result, the consensus means may be slightly higher than the NIST target ranges, reflecting the small fraction of endogenous folic acid that could be extracted by participating laboratories using more extensive extraction procedures such as enzymatic treatments. This contribution is expected to be relatively small, perhaps only 2 % of the total folic acid present in these highly fortified samples.

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								-				
	Lab Code:	NIST	1. Your Results				_	2. Co	mmunity l	3. Target Value		
Analyte	Sample	Units	Mean	s total	Z _{comm}	Z _{NIST}		Ν	Mean	Std Dev	Value	U_{95}
Folic Acid	Breakfast Cereal	mg/kg	14.8	1.2	-0.5	0.0		10	17.8	5.7	14.8	1.2
Folic Acid	Milk Powder	mg/kg	2.31	0.06	-0.4	0.0	_	11	2.44	0.37	2.31	0.06

Exercise G - July 2011 - Folic Acid

Mean Average of reported values

s_{total} Standard deviation of reported values Z_{comm} Z-score: (Lab Mean - Consensus Mean)/

Consensus Standard Deviation

Z_{NIST} Z-score: (Lab Mean - NIST Value or Label Claim)/ NIST or Label Claim Standard Deviation N Number of quantitative values reported

Mean Robust mean of the reported values

Std Dev Robust standard deviation

Value NIST-assessed value

 $U_{95} \pm 95\%$ confidence interval about the assessed value

NR No data reported

		SRM 32.	33 Fortifie	d Breakf	ast Cereal	l (mg/kg)	I	Fortified N	Ailk Powde	er (mg/kg			
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD		
	NIST				14.8	1.2				2.31	0.06		
	G703	13.1	13.1	13.9	13.4	0.5	2.33	2.22	2.32	2.29	0.06		
	G705												
	G706												
	G709	24.6	26.2	26.0	25.6	0.9							
	G711												
	G721												
	G727	14.0	13.9	13.6	13.8	0.2	2.20	2.14	2.15	2.16	0.03		
	G728												
lts	G729	15.8	15.5	14.8	15.4	0.5	2.61	1.90	2.72	2.41	0.44		
esu	G738	16.1			16.1		3.10	3.20	3.20	3.17	0.06		
al R	G739												
Individual Results	G740												
vibr	G744	14.9	15.6	14.1	14.9	0.8	2.97	2.70	2.59	2.75	0.20		
I	G746	24.4	22.5	22.3	23.1	1.2	5.00	4.90	4.80	4.90	0.10		
	G749	40.6	44.4	51.6	45.5	5.6	1.23	1.59	3.65	2.16	1.31		
	G752												
	G753	15.2	15.0	14.9	15.0	0.1	2.19	1.93	2.16	2.09	0.14		
	G754	14.8	15.6	15.2	15.2	0.4	2.27	2.28	2.29	2.28	0.01		
	G758												
	G759	15.5	15.7	15.4	15.5	0.2	2.02	2.33	2.28	2.21	0.16		
	G767												
	G768						2.74	2.72	2.24	2.57	0.28		
	G778												
y		Consensu	s Mean		17.8		Consensu	s Mean		2.44			
Community Results		Consensu	s Standard	Deviation	5.7		Consensu	s Standard	l Deviation	0.37			
ommuni Results		Maximum	1		45.5		Maximun	1		4.9			
Coi R		Minimum			13.4		Minimum			2.1			
		Ν			10		Ν			11			

 Table 6. Data summary table for folic acid in dietary supplements.

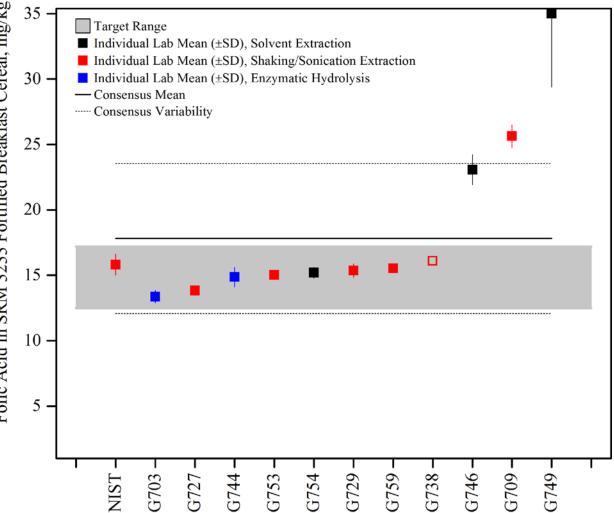


Figure 9. Folic acid in SRM 3233 Fortified Breakfast Cereal (data summary view – sample preparation method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by sample preparation method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}).

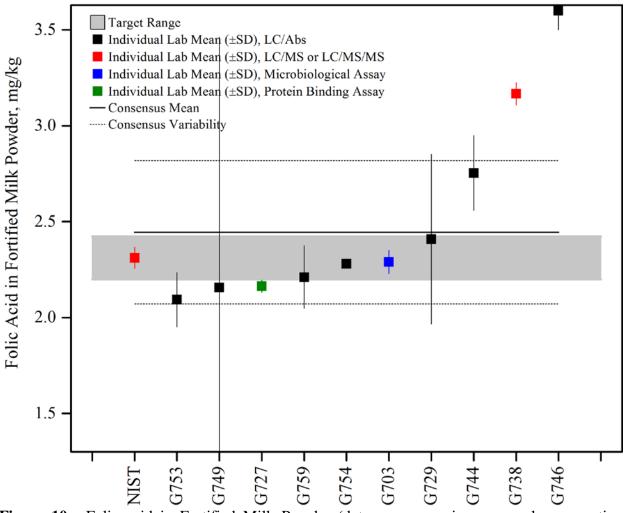


Figure 10. Folic acid in Fortified Milk Powder (data summary view – sample preparation method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by sample preparation method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST assigned value bounded by twice its standard deviation.

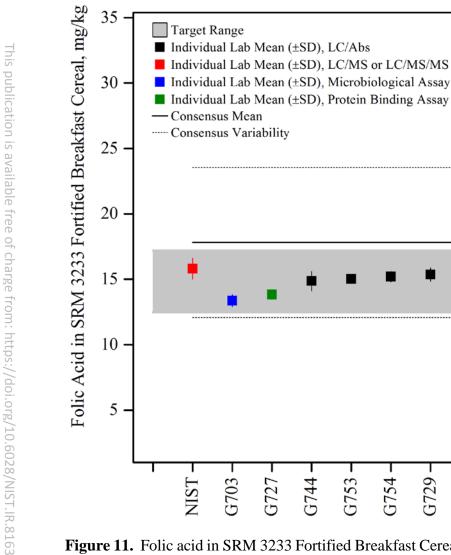


Figure 11. Folic acid in SRM 3233 Fortified Breakfast Cereal (data summary view - instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}) .

G754 -

G729.

G759 -

G738 -

G746 -

G709

G749

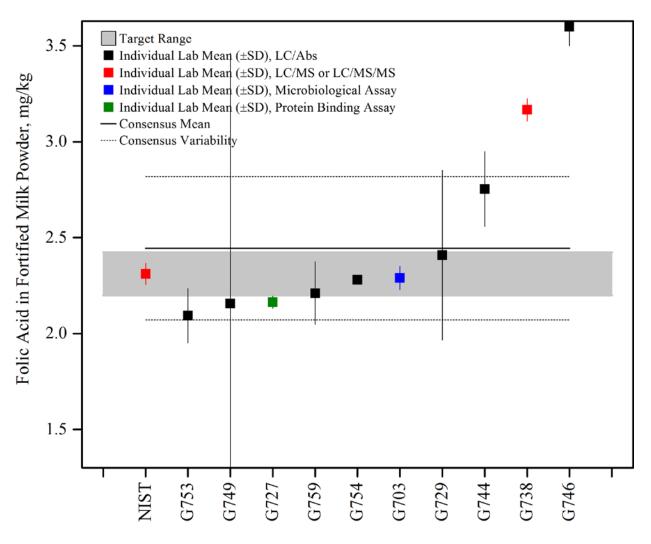


Figure 12. Folic acid in Fortified Milk Powder (data summary view – instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST assigned value bounded by twice its standard deviation.

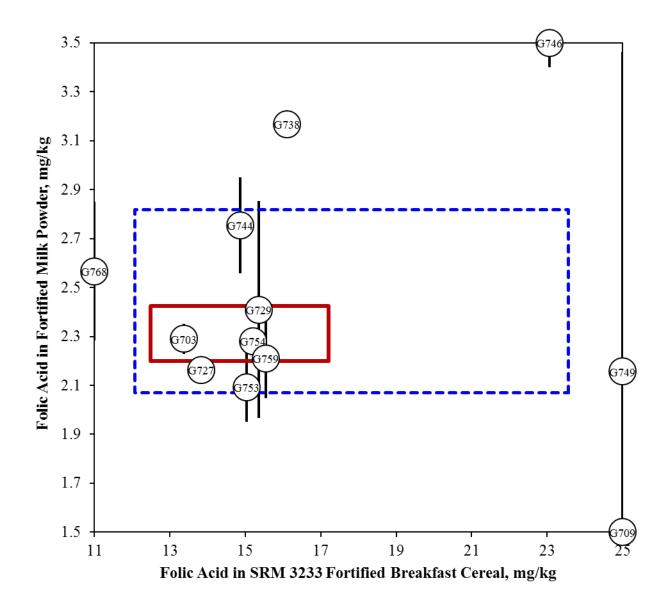


Figure 13. Folic acid in SRM 3233 Fortified Breakfast Cereal and Fortified Milk Powder (sample/sample comparison view). In this view, the individual laboratory results for one sample (SRM 3233 Fortified Breakfast Cereal) are compared to the results for a second sample (Fortified Milk Powder). The solid red box represents the target zone for the breakfast cereal (x-axis) and the milk powder (y-axis). The dotted blue box represents the consensus zone for the breakfast cereal (x-axis) and the milk powder (y-axis).

β-CAROTENE IN SOLUTIONS AND DIETARY SUPPLEMENTS

Study Overview

In this study, participants were provided with one NIST SRM, SRM 3251 *Serenoa repens* Extract, and two ethanolic solutions of β -carotene. Participants were asked to use in-house analytical methods to determine the mass fraction of total β -carotene and isomers in each of the matrices and report values on an as-received basis.

Sample Information

β-carotene Solution 1. Participants were provided with three vials containing approximately 2 mL of all-*trans*-β-carotene dissolved in ethanol containing 30 ppm BHT. Solution 1 did not contain significant amounts of *cis*-β-carotene isomers. Before use, participants were instructed to thoroughly mix the contents of the vials and use a sample size of at least 50 mg. Participants were asked to store the material in a freezer at -20 °C, and to prepare one sample and report one value per analyte from each vial provided. Prior to the study, the approximate level of total β-carotene was given as 0.9 mg/kg. The NIST value and uncertainty for total β-carotene in solution 1 were determined spectrophotometrically in ethanol with a molar absorptivity of 2620 dLg⁻¹cm⁻¹, and the β-carotene concentration decreased approximately 10 % from July 2011 until February 2012. The NIST-determined values and uncertainties are reported in the table below.

β-carotene Solution 2. Participants were provided with three vials containing approximately 2 mL of *Dunaliella* extract dissolved in ethanol containing 30 ppm BHT. In solution 2, approximately 30 % of the total β-carotene was in the form of the 9-*cis*-β-carotene isomer. Before use, participants were instructed to thoroughly mix the contents of the vials and use a sample size of at least 50 mg. Participants were asked to store the material in a freezer at -20 °C, and to prepare one sample and report one value per analyte from each vial provided. Prior to the study, the approximate level of total β-carotene was given as 4 mg/kg. The NIST value and uncertainty for total β-carotene in solution 2 were determined spectrophotometrically in ethanol with a molar absorptivity of 2620 dLg⁻¹cm⁻¹; the total β-carotene content of solution 2 was stabile throughout the time period of the study. The NIST-determined values and uncertainties are reported in the table below.

Saw Palmetto Extract. Participants were provided with three ampoules, each containing approximately 1 mL of Serenoa repens extract. The extract was packaged under nitrogen in amber glass ampoules. Before use, participants were instructed to thoroughly mix the contents of the ampoules and use a sample size of at least 50 mg. Participants were asked to store the material at controlled room temperature, 20 °C to 25 °C, and to prepare one sample and report one value per analyte from each ampoule provided. Prior to the study, the approximate level of total β -carotene was given as 50 mg/kg. The NIST certified values and uncertainties for *trans*- β -carotene, 9-*cis*- β -carotene, and total β -carotene in SRM 3251 were determined at NIST by LC-absorbance and LC-fluorescence (using different column chemistries), following gravimetric dilution and addition of an internal standard. The NIST-determined values and uncertainties are reported in the table below.

Analyte	Mass Fraction in Solution 1 (mg/kg)	Mass Fraction in Solution 2 (mg/kg)	Mass Frag in SRM 3 (mg/kg	3251
trans-β-carotene			36.4 ±	5.6
9- <i>cis</i> -β-carotene			10.40 \pm	1.20
Total β -carotene	0.900 ± 0.045	4.000 ± 0.200	46.8 ±	4.6

Study Results

- Forty-one laboratories enrolled in this exercise and received samples. Twenty-three laboratories reported results for total β -carotene in both of the β -carotene solutions (56 % participation). Twenty-four laboratories reported results for total β -carotene in the saw palmetto extract (59 % participation). One laboratory reported a single value for total β -carotene in the saw palmetto extract and was therefore excluded from calculation of summary statistics.²
- Few laboratories (6 or fewer) reported values for *cis* and *trans* isomers of β-carotene in any sample.
- The consensus ranges for total β -carotene in both solutions were wide (21 % to 23 % relative standard deviation, (RSD) respectively) but overlapped with the target ranges. The consensus means for total β -carotene in both solutions were significantly above the target ranges.
- The consensus range for total β -carotene in SRM 3251 was excellent (10 % RSD) and overlapped with the target range. The consensus mean for total β -carotene was within the target range.
- The consensus range for *trans*- β -carotene in SRM 3251 was excellent (7 % RSD) and overlapped with the target range. The consensus mean for *trans*- β -carotene was within the target range.
- The consensus range for 9-*cis*- β -carotene in SRM 3251 was reasonable (17 % RSD) and overlapped with the target range. The consensus mean for 9-*cis*- β -carotene was within the target range.
- A majority of the laboratories reported using simple dilution (48 %) or saponification with extraction (40 %). Other solvent extraction approaches (8 %) and hydrolysis extraction approaches (4 %) were also reported as methods of sample preparation.
- Most laboratories reported using LC with absorbance detection as the analytical method for analysis (92 %). Laboratories also reported using LC-MS or LC-MS/MS (4 %) and spectrophotometry (4 %) as the instrumental approach.
- All participating laboratories reported the use of an external standard approach to quantitation.

Technical Recommendations

The following are recommendations based on results obtained from the participants in this study.

- No sample preparation approach or analytical method was identified as exceptionally good or problematic.
- Although no sample pretreatment was necessary for the β -carotene solutions, many laboratories subjected the samples to saponification, enzymatic hydrolysis, and solvent extraction. Sample preparation can isomerize the β -carotene, which would not be evident

in the total β -carotene measurements, but would affect the ratio of the *cis/trans* isomers observed.

• As shown in **Figure 25**, **Figure 26**, and **Figure 27**, calibration errors may be possible. Laboratories tended to report consistently biased results for the samples and solutions. The most common calibration error in the measurement of carotenoids is caused by either neglecting to assign the concentration spectrophotometrically (carotenoid measurements are traceable to molar absorptivity) or using the wrong molar absorption coefficient (wrong solvent or temperature).

National Institute of Standards & Technology

	Lab Code:	NIST	Exercis		y 2011 - β- Results	carotene	2. C	o mmunity l	Results	3. Targe	et Value
Analyte	Sample	Units	Mean	s total	Z _{comm}	Z _{NIST}	N	Mean	Std Dev	Value	U_{95}
trans-β-carotene	Solution 1	mg/kg					6	1.05	0.227	NR	NR
trans-β-carotene	Solution 2	mg/kg					6	2.48	0.548	NR	NR
trans-β-carotene	Saw Palmetto Extract	mg/kg	36.4	5.6	0.6	0.0	7	34.9	2.35	36.4	5.6
9-cis-β-carotene	Solution 1	mg/kg					2	0.0113	0.00135	NR	NR
9-cis-β-carotene	Solution 2	mg/kg					6	2.21	0.74	NR	NR
9- <i>cis</i> -β-carotene	Saw Palmetto Extract	mg/kg	10.4	1.2	-0.7	0.0	6	11.8	2.04	10.4	1.2
13- <i>cis</i> -β-carotene	Solution 1	mg/kg					1	0.0364		NR	NR
13-cis-β-carotene	Solution 2	mg/kg					3	0.199	0.0427	NR	NR
13-cis-β-carotene	Saw Palmetto Extract	mg/kg					4	2.59	2.33	NR	NR
15-cis-β-carotene	Solution 1	mg/kg					3	0.0341	0.0347	NR	NR
15-cis-β-carotene	Solution 2	mg/kg					4	0.202	0.156	NR	NR
15-cis-β-carotene	Saw Palmetto Extract	mg/kg					4	2.75	2.37	NR	NR
Total β-carotene	Solution 1	mg/kg	0.9	0.045	-1.0	0.0	20	1.17	0.271	0.900	0.045
Total β-carotene	Solution 2	mg/kg	4	0.2	-1.4	0.0	21	5.61	1.19	4.00	0.20
Total β-carotene	Saw Palmetto Extract	mg/kg	46.8	4.6	-1.0	0.0	20	52.2	5.38	46.8	4.6

Mean Average of reported values

 s_{total} Standard deviation of reported values

Z_{comm} Z-score: (Lab Mean - Consensus Mean)/

Consensus Standard Deviation

Z_{NIST} Z-score: (Lab Mean - NIST Value or Label Claim)/ NIST or Label Claim Standard Deviation

N Number of quantitative values reported Mean Robust mean of the reported values Std Dev Robust standard deviation

Value NIST-assessed value

 $U_{95} \pm 95\%$ confidence interval about the assessed value

NR No data reported

								tra	ns-β-carot	ene						
			Solu	tion 1 (mg	/kg)				tion 2 (mg			SRM	3251 Saw	Palmetto	Extract (n	ng/kg)
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST														36.4	5.6
	G703	1.34	1.60	1.17	1.37	0.22	2.98	3.23	3.26	3.16	0.15	35.8	34.4	34.4	34.9	0.8
	G705															
	G706															
	G707															
	G709	1.13	1.00	0.95	1.03	0.10	3.48	2.51	2.26	2.75	0.64	33.0	33.8	34.0	33.6	0.5
	G710															
	G711 G712															
	G714 G716	0.75	0.92	1.00	0.89	0.13	1.33	2.16	1.78	1.76	0.42	34.8	34.7	41.8	37.1	4.1
	G717	0.75	0.92	1.00	0.89	0.15	1.55	2.10	1.78	1.70	0.42	54.8	34.7	41.8	37.1	4.1
	G717 G721															
	G722															
	G724															
	G727	18.61	22.90	17.56	19.69	2.83	53.80	47.72	50.17	50.56	3.06		37.2	37.5	37.4	0.2
	G728															
ts	G729															
Individual Results	G730															
al R	G731															
/idu	G732															
ndiv	G733															
	G738															
	G739															
	G740															
	G742															
	G744 G746															
	G749															
	G752															
	G753	1.06	1.11	1.09	1.09	0.03	2.48	2.54	2.67	2.56	0.10	35.0	35.5	37.0	35.8	1.0
	G755															
	G758 G760															
	G762															
	G764															
	G766	0.78	0.75	0.85	0.79	0.05	2.18	2.03	2.25	2.15	0.11	28.4	29.0	28.1	28.5	0.4
	G767 G770															
	G773															
	G774															
	G775	9					9			2.5.5		6			25.0	
Community Results		Consensu		Deviation	1.14 0.37		Consensu		l Deviation	2.76 0.96		Consensu		l Deviation	35.0 2.6	
sult		Maximum		- De viation	19.69		Maximum			50.56		Maximun			37.4	
Con		Minimum			0.79		Minimum			1.76		Minimum			28.5	
-		Ν			6		Ν			6		Ν			6	

Table 8. Data summary table for *trans*- β -carotene in solutions and dietary supplements.

								0 ~	is-β-caroto	no						
			Solu	tion 1 (mg	r/kg)				tion 2 (mg			SRM	3251 Saw	Palmetto	Extract (r	ng/kg)
	Lab	А	B	C	Avg	SD	А	B	C C	Avg	SD	A	B	C	Avg	SD
	NIST		D	U	ning.	55		D	U	1118	50		D	Ũ	10.4	1.2
	G703	0.0108	0.0146	0.0110	0.0121	0.0021	2.63	2.67	2.67	2.66	0.02	15.3	15.0	14.8	15.1	0.3
	G705															
	G706															
	G707															
	G709						1.81	2.46	2.44	2.23	0.37	11.5	12.2	12.2	12.0	0.4
	G710															
	G711															
	G712															
	G714															
	G716						0.28	0.41	0.34	0.34	0.07	10.3	10.1	11.7	10.7	0.8
	G717															
	G721															
	G722															
	G724															
	G727															
	G728															
s	G729															
Individual Results	G730															
l Re	G731															
qua	G732															
divi	G733															
In	G738															
	G739															
	G740															
	G742															
	G744															
	G746															
	G749															
	G752															
	G753						2.76	2.71	2.72	2.73	0.03	9.3	9.7	9.0	9.3	0.4
	G755															
	G758 G760															
	G762															
	G764															
	G766						1.80	1.77	1.82	1.80	0.03	12.9	16.3	10.3	13.2	3.0
	G767															
	G770 G773															
	G774															
	G775															
ity		Consensu			0.0121		Consensu			1.95		Consensu			12.0	
uni ılts			s Standard	Deviation					l Deviation				is Standard	Deviation	2.4	
Community Results		Maximum Minimum					Maximum Minimum			2.73 0.34		Maximun Minimum			15.1 9.3	
ŭ		Nimmum			1		Ninimum			0.34 5		Ninimum			9.3 5	
		14			1		11			5		13			5	

Table 9. Data summary table for 9-*cis*- β -carotene in solutions and dietary supplements.

								13 .	cis-β-carot	ene						
			Solut	tion 1 (m	a/ka)		1		tion 2 (mg			SDM	3251 Saw	Dalmatta	Extract (r	ng/kg)
	Lab	Α	B	C		SD	Α	B	C C		SD	A	3231 Saw B	C		SD
	NIST	A	D	t	Avg	50	A	D	C	Avg	50	A	Б	C	Avg	50
	G703															
	G705															
	G706															
	G707											0.00	6.40		6.07	0.00
	G709											8.00	6.40	6.51	6.97	0.89
	G710															
	G711															
	G712			_		_		_	_	_	_		_		_	
	G714															
	G716						0.160	0.220	0.190	0.190	0.030	0.73	0.66	0.70	0.70	0.04
	G717															
	G721															
	G722															
	G724															
	G727															
	G728															
ts	G729															
Individual Results	G730															
I R.	G731															
dua	G732															
divi	G733															
Ĩ	G738															
	G739															
	G740															
	G742															
	G744															
	G746															
	G749															
	G752															
	G753						0.360	0.340	0.340	0.347	0.012	3.20	2.90	2.60	2.90	0.30
	G755						0.000	010 10	0.010	01017	01012	0.20	2170	2.00	2100	0150
	G758															
	G760															
	G762															
	G764															
	G766 G767															
	G770															
	G773										_					
	G774															
	G775															
Community Results		Consensus					Consensu			0.268		Consensu			3.52	
nun ults		Consensus	Standard	Deviatior	1				l Deviation				is Standard	Deviation		
Rest		Maximum Minimum					Maximum Minimum			0.347 0.190		Maximun Minimum			7.0 0.7	
ŭΓ		N			0		N			0.190 2		N			3	
		19			U		1N			Z		IN			3	

Table 10. Data summary table for -*cis*- β -carotene in solutions and dietary supplements.

								15-0	cis-β-carot	ene						
			Solu	tion 1 (mg	/kg)				tion 2 (mg			SRM	3251 Saw	Palmetto	Extract (r	ng/kg)
1	Lab	Α	В	C	Avg	SD	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST															
	G703															
	G705															
	G706															
	G707															
	G709															
	G710															
	G711															
	G712															
	G714															
	G716	0.0300	0.0400	0.0400	0.0367	0.0058	0.150	0.190	0.170	0.170	0.020	1.93	1.90	2.23	2.02	0.18
	G717															
	G721															
	G722															
	G724															
	G727															
	G728															
lts	G729															
tesu	G730															
Individual Results	G731															
vidu	G732															
ndi	G733															
Ι	G738															
	G739															
	G740															
	G742															
	G744															
	G746															
	G749															
	G752 G753	0.0500	0.0600	0.0800	0.0633	0.0153	0.270	0.270	0.270	0.270	0.000	3.50	4.70	3.40	3.87	0.72
	G755	0.0500	0.0000	0.0800	0.0033	0.0155	0.270	0.270	0.270	0.270	0.000	3.30	4.70	3.40	5.87	0.72
	G758															
	G760															
	G762			_												
	G764 G766						0.350	0.310	0.370	0.343	0.031	5.63	4.88	4.26	4.92	0.69
	G767						0.550	0.510	0.370	0.545	0.051	5.05	+.00	7.20	7.72	0.09
	G770															
	G773															
	G774															
~	G775	Consensu	s Mean		0.0500		Consensu	s Mean		0.261		Consensu	is Mean		3.60	
ts ts				Deviation					l Deviation					Deviation	1.67	
nmu esult		Maximum			0.0633		Maximum	ı		0.343		Maximun			4.9	
Community Results		Minimum			0.0367		Minimum			0.170		Minimum			2.0	
Ē		Ν			2		Ν			3		Ν			3	

Table 11. Data summary table for -*cis*- β -carotene in solutions and dietary supplements.

								Tot	tal β-carot	ene						
			Solu	tion 1 (mg	g/kg)				tion 2 (mg			SRM	3251 Saw	Palmetto	Extract (r	ng/kg)
[Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST				0.90	0.05				4.00	0.20				46.8	4.6
	G703	1.46	1.73	1.27	1.48	0.24	6.47	6.76	6.79	6.67	0.18	57.7	55.4	54.5	55.9	1.7
	G705															
	G706	0.88	1.13	1.02	1.01	0.13	4.91	5.19	4.68	4.93	0.26	46.2	46.0	46.5	46.2	0.3
	G707															
	G709	1.13	1.00	0.95	1.03	0.10	5.29	4.97	4.69	4.98	0.30	52.5	52.4	52.7	52.6	0.2
	G710															
	G711	1.13	0.94	0.91	0.99	0.12	4.81	4.60	5.58	5.00	0.52	48.3	47.8	51.5	49.2	2.0
	G712															
	G714	0.94	0.77	1.18	0.96	0.21	4.33	6.44	6.74	5.84	1.31					
	G716	0.78	0.96	1.04	0.93	0.13	1.92	2.98	2.48	2.46	0.53	47.8	47.4	56.4	50.5	5.1
	G717															
	G721															
	G722	2.82	2.87	2.76	2.82	0.06	8.74	9.84	8.82	9.13	0.61	163.3	176.9	182.5	174.2	9.9
	G724	1.16	1.28	1.43	1.29	0.14	6.28	6.68	6.16	6.38	0.27	50.8	49.3	49.9	50.0	0.8
	G727															
	G728															
ults	G729	1.16	0.79	0.69	0.88	0.25	5.13	5.04	3.66	4.61	0.82	62.3	38.8	57.6	52.9	12.4
Individual Results	G730	1.50	1.30	1.30	1.37	0.12	6.30	6.10	6.20	6.20	0.10	57.5	56.9	58.4	57.6	0.8
I la I	G731	1.19	1.63	1.20	1.34	0.25	5.56	5.57	5.44	5.52	0.07	54.5	53.5	53.2	53.7	0.6
ıbivi	G732	1.18	1.25	1.15	1.19	0.05	5.67	5.88	5.65	5.73	0.13	49.8	50.7	51.8	50.7	1.0
Indi	G733	1.31	1.45	1.35	1.37	0.07	5.95	6.39	6.23	6.19	0.22	52.8	55.5	51.5	53.3	2.0
	G738	2.40	1.30	1.10	1.60	0.70	6.20	7.30	7.00	6.83	0.57	89.0	119.0	150.0	119.3	30.5
	G739															
	G740															
	G742	1.04	0.06	0.05	0.02	0.11	4.50	1.20	4.00	4.20	0.17		42.0	45.0	47.0	7.0
	G744 G746	1.04	0.86	0.85	0.92	0.11	4.50 3.40	4.20 3.30	4.20	4.30 3.87	0.17	55.5 69.6	42.0	45.8	47.8 57.4	7.0
	G740 G749	2.22	1.90	1.20	2.22	0.40	8.33	5.83	4.90 6.27	6.81	1.33	113.6	45.8 106.4	56.7 124.3	114.8	11.9 9.0
	G749	2.22			2.22		6.55	5.85	0.27	0.81	1.55	115.0	100.4	124.5	114.0	9.0
	G752	1.11	1.17	1.17	1.15	0.03	6.35	6.37	6.50	6.41	0.08	51.0	52.8	52.0	51.9	0.9
	G755															
	G758	1.05	1.00	1.15	1.1.6	0.10	4.00	5.00	1.07	1.00	0.10	<i></i>			c 4 7	
	G760 G762	1.05 1.32	1.30 1.19	1.15 1.08	1.16 1.20	0.12	4.89 6.37	5.09 6.29	4.97 6.25	4.98 6.30	0.10	64.5 56.2	54.5	55.6	64.5 55.4	0.9
	G764	1.52	1.19	1.00	1.20	0.12	0.37	0.29	0.25	0.50	0.00	50.2	54.5	55.0	55.4	0.9
	G766											47.0	50.1	42.7	46.6	3.7
	G767															
	G770	0.10	0.11	0.00	0.10	0.00	2.00	2.02	2.60	2.00	0.11	49.4	43.6	43.3	45.4	3.4
	G773 G774	0.10	0.11	0.08	0.10	0.02	3.90	3.83	3.68	3.80	0.11	0.1	0.1	0.1	0.1	0.0
	G775	0.90	0.95		0.93	0.04	6.29	5.91	6.60	6.27	0.35	54.2	52.0	51.4	52.5	1.5
ity		Consensu	s Mean		1.17		Consensu	ıs Mean		5.58		Consensu	s Mean		52.5	
nuni ılts				Deviation					l Deviation					1 Deviation		
Community Results		Maximum	I		2.82 0.88		Maximun			9.13 2.46		Maximun			174.2 46.2	
ŭ		Minimum N			0.88 19		Minimum N			2.46 20		Minimum N			46.2 19	
		1 N			19		IN			20		IN			19	

Table 12. Data summary table for total β -carotene in solutions and dietary supplements.

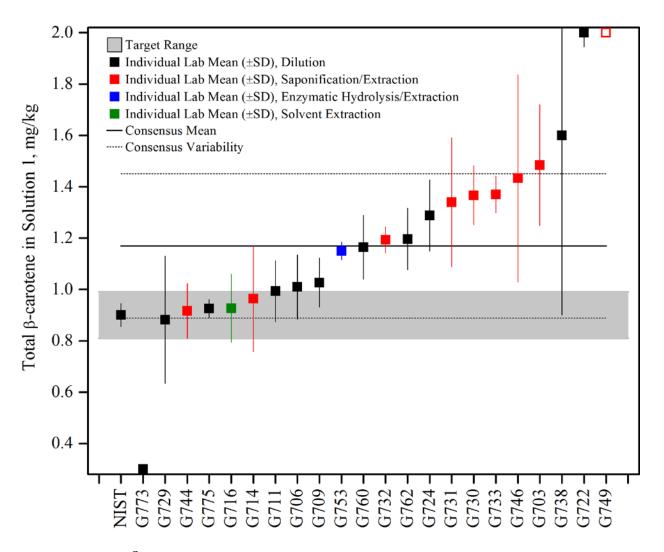


Figure 14. Total β -carotene in Solution 1 (data summary view – sample preparation method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by sample preparation method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST assigned value bounded by twice its standard deviation.

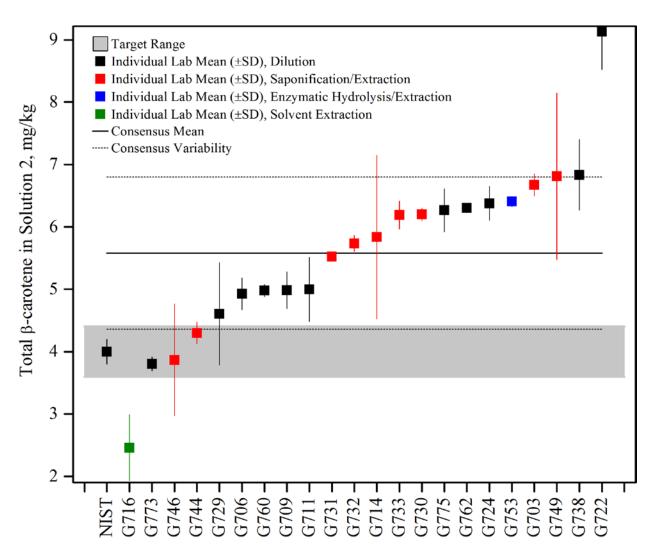


Figure 15. Total β -carotene in Solution 2 (data summary view – sample preparation method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by sample preparation method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST assigned value bounded by twice its standard deviation.

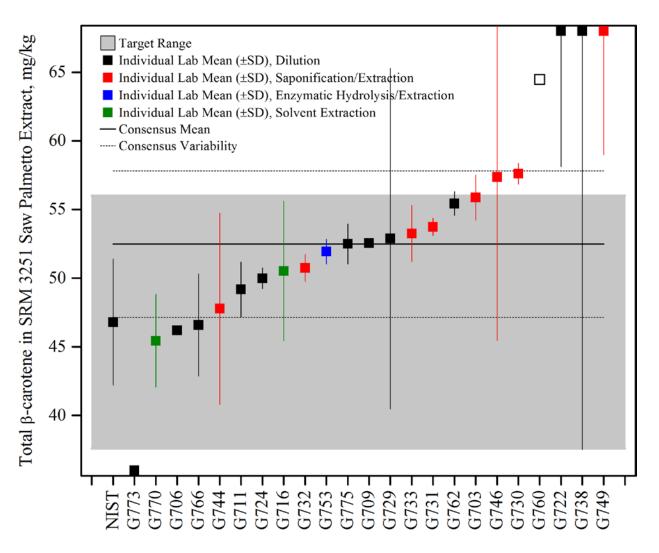


Figure 16. Total β -carotene in SRM 3251 *Serenoa repens* Extract (data summary view – sample preparation method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by sample preparation method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}).

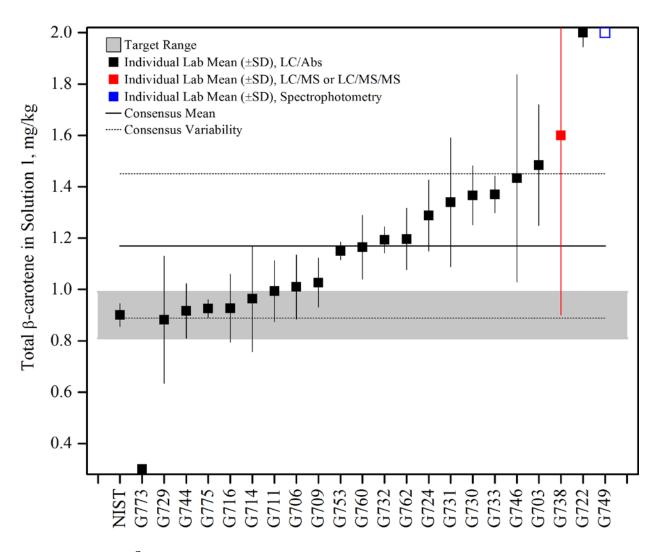


Figure 17. Total β -carotene in Solution 1 (data summary view – instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST assigned value bounded by twice its standard deviation.

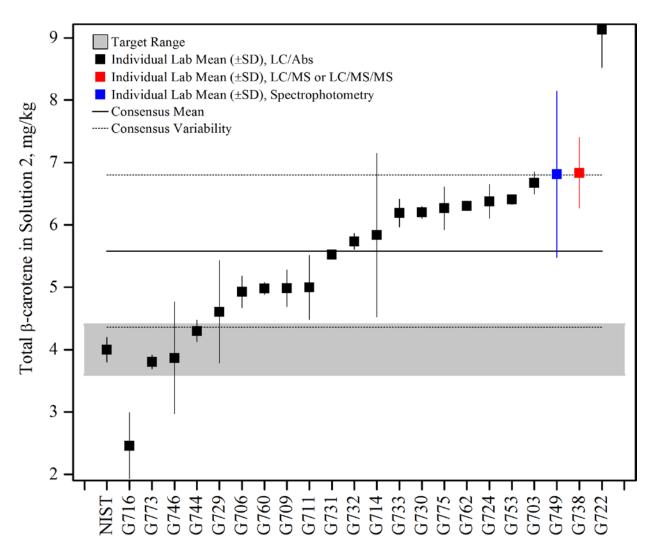


Figure 18. Total β -carotene in Solution 2 (data summary view – instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST assigned value bounded by twice its standard deviation.

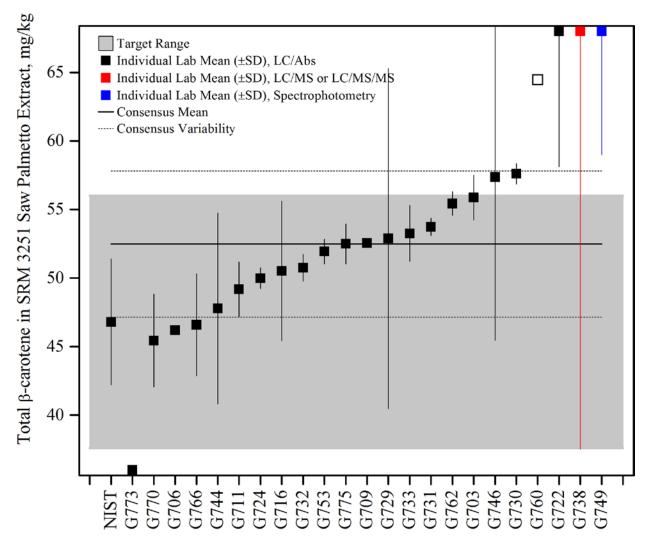


Figure 19. Total β -carotene in SRM 3251 *Serenoa repens* Extract (data summary view – instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}).

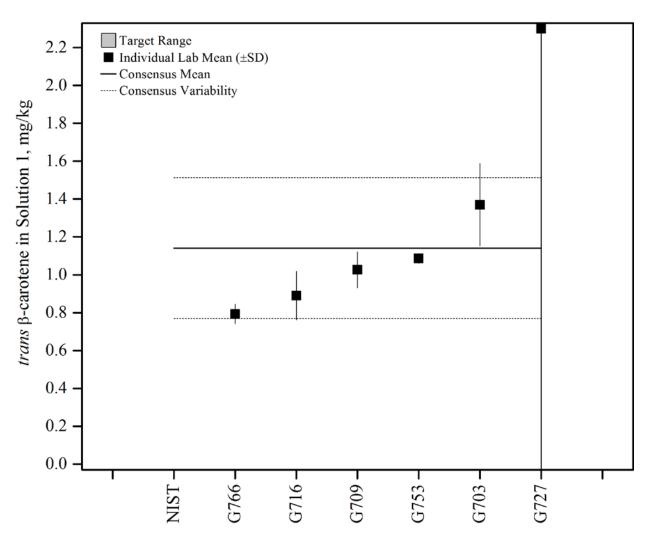


Figure 20. *trans*- β -carotene in Solution 1 (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

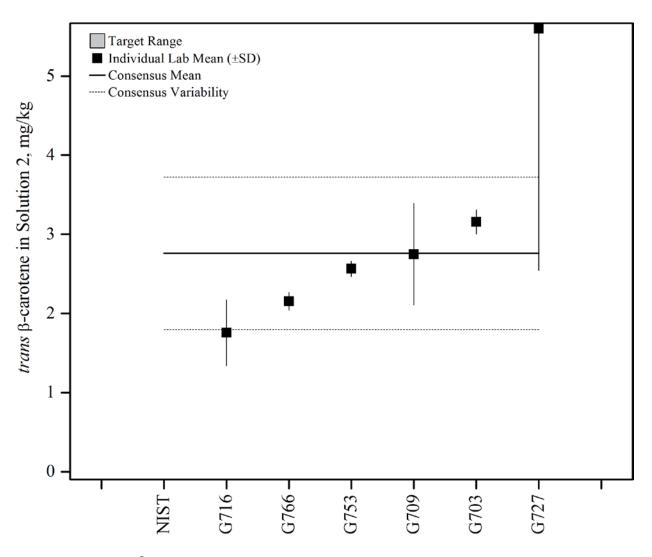


Figure 21. *trans*- β -carotene in Solution 2 (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

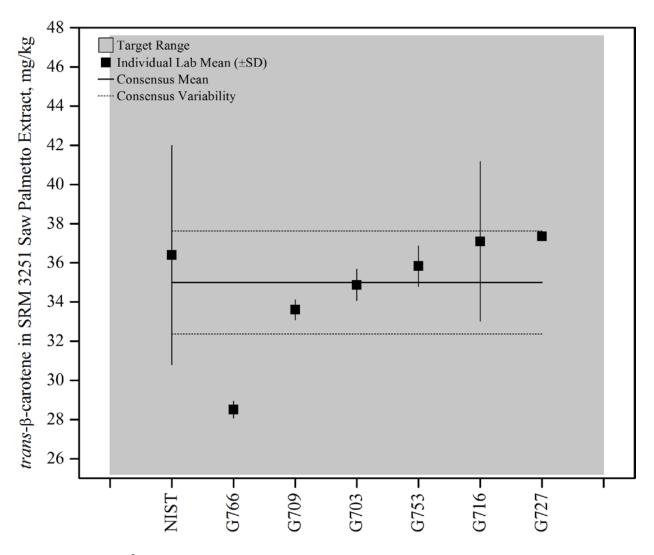


Figure 22. *trans*- β -carotene in SRM 3251 *Serenoa repens* Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}).

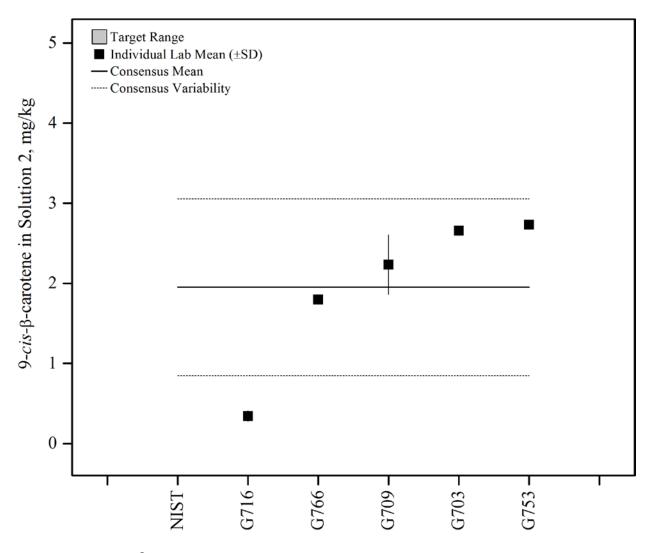


Figure 23. 9-*cis*- β -carotene in Solution 2 (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

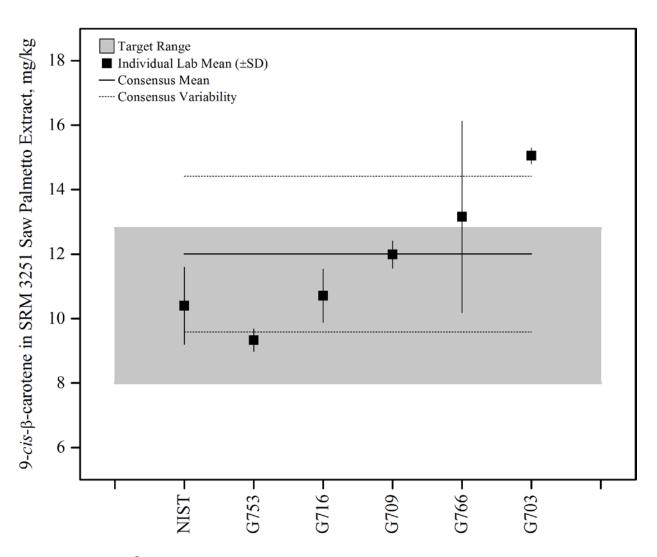


Figure 24. 9-*cis*-- β -carotene in SRM 3251 *Serenoa repens* Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean. The gray shaded region represents the target zone for "acceptable" performance, which encompasses the NIST certified value bounded by twice its uncertainty (U_{95}).

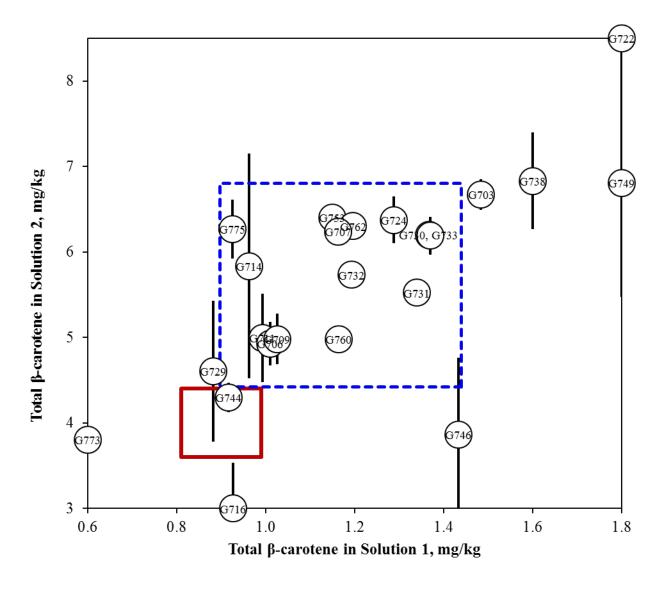


Figure 25. Total β -carotene in Solution 1 and Solution 2 (sample/sample comparison view). In this view, the individual laboratory results for one sample (β -carotene Solution 1) are compared to the results for a second sample (β -carotene Solution 2). The solid red box represents the target zone for solution 1 (x-axis) and solution 2 (y-axis). The dotted blue box represents the consensus zone for solution 1 (x-axis) and solution 2 (y-axis).

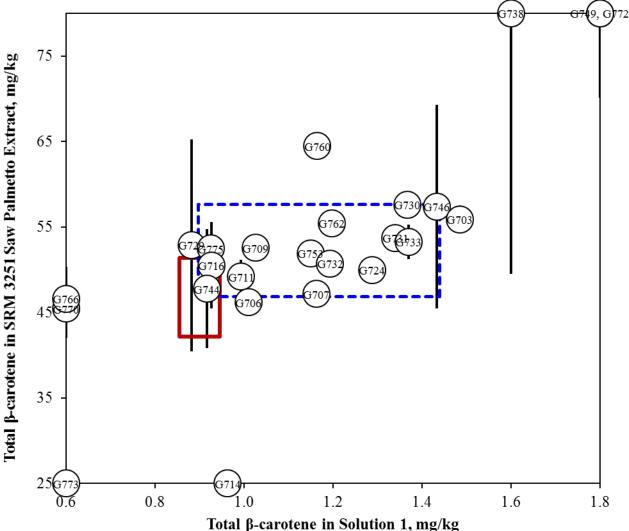


Figure 26. Total β -carotene in Solution 1 and SRM 3251 *Serenoa repens* Extract (sample/sample comparison view). In this view, the individual laboratory results for one sample (β -carotene Solution 1) are compared to the results for a second sample (saw palmetto extract). The solid red box represents the target zone for solution 1 (x-axis) and saw palmetto extract (y-axis). The dotted blue box represents the consensus zone for solution 1 (x-axis) and saw palmetto extract (y-axis).

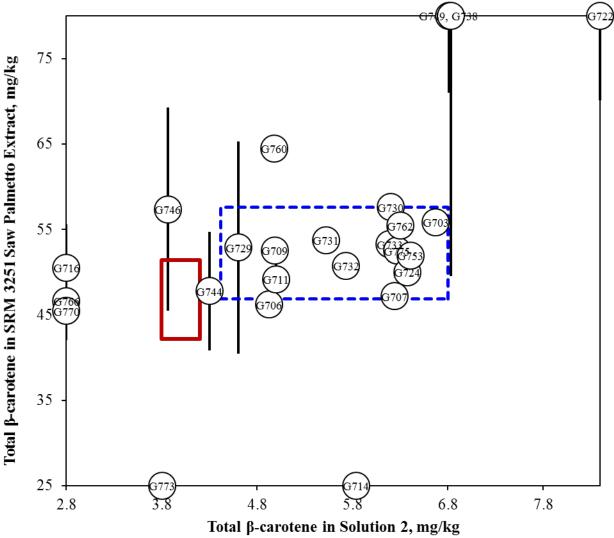


Figure 27. Total β -carotene in Solution 2 and SRM 3251 *Serenoa repens* Extract (sample/sample comparison view). In this view, the individual laboratory results for one sample (β -carotene Solution 2) are compared to the results for a second sample (saw palmetto extract). The solid red box represents the target zone for solution 2 (x-axis) and saw palmetto extract (y-axis). The dotted blue box represents the consensus zone for solution 2 (x-axis) and saw palmetto extract (y-axis).

ANTHOCYANINS & ANTHOCYANIDINS IN DIETARY SUPPLEMENTS

Study Overview

In this study, participants were provided with two NIST SRMs, SRM 3283 Cranberry Extract and SRM 3291 Bilberry Extract. Participants were asked to use in-house analytical methods to determine the mass fractions of anthocyanins and/or anthocyanidins in each of the matrices and report values on an as-received basis.

Sample Information

Cranberry Extract. Participants were provided with one packet containing approximately 2.5 g of cranberry extract. The cranberry extract was heat-sealed inside nitrogen-flushed 4 mil polyethylene bags, which were then sealed inside aluminized plastic bags with 2 packets of silica gel. Before use, participants were instructed to thoroughly mix the contents of the packet and use a sample size of at least 0.1 g. Participants were asked to store the material at controlled room temperature, 20 °C to 25 °C, and to prepare three samples and report three values from the single packet provided. Prior to the study, the approximate analyte levels were given as 1.8 mg/g cyanidin, 0.1 mg/g delphinidin, and 0.7 mg/g peonidin. NIST assigned values for anthocyanins and anthocyanidins were not available at the time of this report.

Bilberry Extract. Participants were provided with one packet containing approximately 1 g of bilberry extract. The bilberry extract was heat-sealed inside nitrogen-flushed 4 mil polyethylene bags, which were then sealed inside aluminized plastic bags with 2 packets of silica gel. Before use, participants were instructed to thoroughly mix the contents of the packet and use a sample size of at least 0.1 g. Participants were asked to store the material at controlled room temperature, 20 °C to 25 °C, and to prepare three samples and report three values from the single packet provided. Prior to the study, the approximate analyte levels were given as 40 mg/g cyanidin, 45 mg/g delphinidin, 25 mg/g malvidin, 10 mg/g peonidin, and 15 mg/g petunidin. NIST assigned values for anthocyanins and anthocyanidins were not available at the time of this report.

Study Results

- Thirty-four laboratories enrolled in this exercise and received samples.
- Five to ten laboratories reported results for anthocyanidins in both the cranberry extract and the bilberry extract (21 % to 38 % participation). One laboratory reported a single value for anthocyanidins in both samples and was therefore excluded from calculation of summary statistics.² Further discussion will only include studies in which at least eight laboratories reported data.
 - The consensus ranges for anthocyanidins in both matrices were wide (46 % to over 100 % RSD).
 - The consensus means were very low compared to the expected values reported prior to the study.
- Nine to fourteen laboratories reported results for anthocyanins in both the cranberry extract and the bilberry extract (27 % to 41 % participation).
 - The consensus ranges for anthocyanins in the cranberry extract ranged from acceptable (21 % RSD) to wide (29 % to over 100 % RSD).
 - The consensus ranges for anthocyanins in the bilberry extract ranged from acceptable (13 % to 23 % RSD) to wide (38 % to 56 % RSD).

- Nineteen laboratories reported results for total anthocyanins in both the cranberry extract and the bilberry extract (56 % participation). One laboratory reported a single value for total anthocyanins in both samples and was therefore excluded from calculation of summary statistics.²
 - The consensus range for total anthocyanins in the cranberry extract was wide (38 % RSD).
 - The consensus range for total anthocyanins in the bilberry extract was acceptable (22 % RSD).
- A majority of the laboratories reported using a solvent extraction approach (50 %) or a shaking or sonication extraction approach (39 %). Hydrolysis was also reported as a method of sample preparation by two laboratories (11 %).
- Most laboratories reported using LC with absorbance detection as the analytical method for analysis (78 %). Laboratories also reported using spectrophotometry (17 %) and a microbiological assay (6 %) as the instrumental approach.
- Most laboratories reported the use of an external standard approach to quantitation (89 %), while two laboratories reported calibration against a molar absorptivity.

Technical Recommendations

The following are recommendations based on results obtained from the participants in this study.

- The wide consensus ranges for the anthocyanidins may be related to the extraction approach utilized.
 - Laboratories using a hydrolysis approach were not included in the consensus calculations.
 - The variability in the data could be attributed to the extraction approach. The glycosides can be broken down into the aglycones, and differences in the extraction time, temperature, or solvent used could result in increased variability in the amount of glycosides converted to aglycones.
 - Many laboratories are not calibrating using reference standards for each of these compounds, and are instead using a single compound (e.g. cyanidin) for calibration of all five aglycones. If this were a major factor, the variability in the cyanidin consensus should be lower than that for the other aglycones, which was not observed.
- The wide consensus ranges for the anthocyanins may also be related to the extraction approach utilized.
 - Laboratories using a hydrolysis approach were not included in the consensus calculations.
 - As described above, the variability in the data could be attributed to the extraction approach. The glycosides can be broken down into the aglycones, and differences in the extraction time, temperature, or solvent used could result in increased variability in the amount of glycosides converted to aglycones.
 - Many laboratories are not calibrating using reference standards for each of these compounds, and are instead using a single compound (e.g. cyanidin glucoside) or a small subset of the glycosides for calibration of all fifteen glycosides. If this were a major factor, the variability in the cyanidin consensus should be lower than that for the other aglycones, which was not observed.

- After molar mass conversion to totals for each aglycone, the data for both hydrolysis and extraction approaches could be compared. In many cases, the consensus variability for a total aglycone was decreased compared to that for any of the other forms individually.
 - The value reported for each of the three glycosides was converted to the aglycone equivalent and summed, along with the value reported for the aglycone, to give a value for total aglycone.
 - The decrease in variability using this approach is further evidence in support of the theory provided above, of the differences in extraction procedure leading to increased variability. By reducing all forms to the base aglycone yields a sum that has a tighter consensus.
 - **Figure 73** does not reflect an obvious calibration error. Some laboratories reported values that were high for the one sample but were within range for the other sample. This type of trend may indicate individualized problems with the extraction or separation from one matrix compared to another.
- The consistency of the data is improved when all values are converted to a value for "equivalents" of each compound, and even further improved when converted to a value for total anthocyanidins.
 - The degree of sample pretreatment may affect the individual compounds measured, but would affect the total less. Harsher extraction conditions (elevated temperature, high acid content, etc.) could cause increased, unintended hydrolysis. Extraction conditions should be thoroughly optimized to ensure that conversion is not occurring unintentionally during the extraction process.
 - Lack of authentic standards for the large number of compounds may cause difficulties in chromatographic peak identification and quantitation. The best solution is to acquire as many standards as possible and use retention time (and m/z) to confirm peak identifications. Relying on literature or official methods has limitations, as variations in column chemistry, mobile phase composition, and temperature can all affect the chromatographic selectivity and therefore the retention times of all compounds.

Individual data table (NIST) for anthocyanins and anthocyanidins in dietary Table 13. supplements.

National Institute of Standards & Technology

Exercise	G - July	2011	- Anthocyanir	is and A	Anthocyanidin

		Exerci	se G - July 2	2011 - An	thocyanins	and Anthoc	yanidins				
	Lab Code:	NIST		1. Your	Results		2. Co	ommunity l	Results	3. Targe	et Value
Analyte	Sample	Units	Mean	s total	Z _{comm}	Z _{NIST}	N	Mean	Std Dev	Value	U_{95}
Cyanidin	Cranberry	mg/g					8	0.0859	0.0504	NR	NR
Cyanidin	Bilberry	mg/g					12	3.08	2.68	NR	NR
Delphinidin	Cranberry	mg/g					6	0.116	0.18	NR	NR
Delphinidin	Bilberry	mg/g					12	4.23	4.64	NR	NR
Malvidin	Cranberry	mg/g					2	0.0493	0.0269	NR	NR
Malvidin	Bilberry	mg/g					10	1.11	0.823	NR	NR
Peonidin	Cranberry	mg/g					6	0.0505	0.0045	NR	NR
Peonidin	Bilberry	mg/g					10	0.669	0.608	NR	NR
Petunidin	Cranberry	mg/g					2	0.133	0.0589	NR	NR
Petunidin	Bilberry	mg/g					11	0.757	0.348	NR	NR
Cyanidin Ara	Cranberry	mg/g					11	0.198	0.0574	NR	NR
Cyanidin Ara	Bilberry	mg/g					14	29.9	11.3	NR	NR
Cyanidin Gal	Cranberry	mg/g					13	0.132	0.0394	NR	NR
Cyanidin Gal	Bilberry	mg/g					14	35.5	7.32	NR	NR
Cyanidin Glu	Cranberry	mg/g					8	0.0586	0.0778	NR	NR
Cyanidin Glu	Bilberry	mg/g					13	43.2	13.3	NR	NR
Delphinidin Ara	Cranberry	mg/g					2	0.0615	0.0815	NR	NR
Delphinidin Ara	Bilberry	mg/g mg/g					13	37.3	7.64	NR	NR
Delphinidin Gal	Cranberry	00					3	0.0187	0.029	NR	NR
Delphinidin Gal	Bilberry	mg/g					13	41.9	6.45	NR	NR
		mg/g					4	0.0568	0.0621	NR	NR
Delphinidin Glu	Cranberry	mg/g									
Delphinidin Glu	Bilberry	mg/g					14	45.8	7.8	NR	NR
Malvidin Ara	Cranberry	mg/g						0.0514	0.0776	NR	NR
Malvidin Ara	Bilberry	mg/g						8.19	3.22	NR	NR
Malvidin Gal	Cranberry	mg/g					3	0.0595	0.0865	NR	NR
Malvidin Gal	Bilberry	mg/g					13	18.4	8.76	NR	NR
Malvidin Glu	Cranberry	mg/g					4	0.0881	0.0908	NR	NR
Malvidin Glu	Bilberry	mg/g					12	29.7	13.4	NR	NR
Peonidin Ara	Cranberry	mg/g					11	0.148	0.0479	NR	NR
Peonidin Ara	Bilberry	mg/g					12	2.44	0.921	NR	NR
Peonidin Gal	Cranberry	mg/g					12	0.201	0.0428	NR	NR
Peonidin Gal	Bilberry	mg/g					14	5.29	2.96	NR	NR
Peonidin Glu	Cranberry	mg/g					9	0.0467	0.0454	NR	NR
Peonidin Glu	Bilberry	mg/g					13	17.5	2.96	NR	NR
Petunidin Ara	Cranberry	mg/g					0			NR	NR
Petunidin Ara	Bilberry	mg/g					12	10.3	1.31	NR	NR
Petunidin Gal	Cranberry	mg/g					2	0.0993	0.109	NR	NR
Petunidin Gal	Bilberry	mg/g					11	16.8	3.84	NR	NR
Petunidin Glu	Cranberry	mg/g					4	0.0618	0.0998	NR	NR
Petunidin Glu	Bilberry	mg/g					14	32.7	5.96	NR	NR
Total Anthocyanins	Cranberry	mg/g					18	0.934	0.358	NR	NR
Total Anthocyanins	Bilberry	mg/g					18	354	78.2	NR	NR
Cyanidin	Cranberry	mg/g					15	0.259	0.103	NR	NR
Cyanidin	Bilberry	mg/g					15	76.5	23.7	NR	NR
Delphinidin	Cranberry	mg/g					9	0.134	0.166	NR	NR
Delphinidin	Bilberry	mg/g					15	84.7	14.4	NR	NR
Malvidin	Cranberry	mg/g					8	0.0624	0.0576	NR	NR
Malvidin	Bilberry	mg/g					15	34.2	12.7	NR	NR
Peonidin	Cranberry	mg/g					15	0.252	0.118	NR	NR
Peonidin	Bilberry	mg/g					15	17.6	4.37	NR	NR
Petunidin	Cranberry	mg/g					6	0.0942	0.0979	NR	NR
Petunidin	Bilberry	mg/g					15	38.4	7.06	NR	NR
i cruindin	Differry	mg/g					15	50.4	7.00	INIX	1111

Mean Average of reported values

- s_{total} Standard deviation of reported values
- Z_{comm} Z-score: (Lab Mean Consensus Mean)/
- Consensus Standard Deviation
- $Z_{NIST}\;$ Z-score: (Lab Mean NIST Value or Label Claim)/ NIST or Label Claim Standard Deviation

N Number of quantitative values reported Mean Robust mean of the reported values Std Dev Robust standard deviation

Value NIST-assessed value

 $U_{95} \pm 95\%$ confidence interval about the assessed value

NR No data reported

	. Dutt	. Summu	<u>, 1010</u>	101 0 7 0			nidin				
		SRM	4 3283 Ci	ranberry I	Extract (m	-	r	M 3291 F	Bilberry E	xtract (mg	/g)
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST										
	G701										
	G702										
	G703	0.068	0.067	0.069	0.068	0.001	3.33	3.31	3.36	3.33	0.03
	G704										
	G705	0.039	0.043	0.038	0.040	0.003	25.60	25.40	25.50	25.50	0.10
	G707										
	G708										
	G715	0.095	0.098		0.097	0.002	1.81	1.75		1.78	0.05
	G716										
	G717	0.053	0.051	0.048	0.050	0.002	0.25	0.25	0.25	0.25	0.00
	G718	0.060	0.059	0.057	0.059	0.002	0.74	0.64	0.67	0.68	0.05
	G722										
	G724										
s	G728						2.10	1.83	1.82	1.92	0.16
sult	G729										
l Re	G738		_		_	_		_			
dua	G739										
Individual Results	G740										_
In	G749										
	G752										
	G753 G757										
	G760										
	G762	0.126	0.128	0.126	0.127	0.001	3.88	3.60	3.41	3.63	0.24
	G765	0.120	0.120	0.120	0.127	0.001	1.26	1.20	1.30	1.25	0.24
	G766						1.20	1.20	1.50	1.25	0.05
	G767										
	G771										
	G772										
	G773						4.73	4.86	4.47	4.69	0.20
	G774										
	G775						2.22	2.23	2.28	2.25	0.03
	G776										
	G777	0.173	0.169	0.172	0.171	0.002	48.45	48.35	48.58	48.46	0.11
y		Consensu	s Mean		0.0859		Consensu	s Mean		3.08	
Community Results		Consensu	s Standard	Deviation	0.0504		Consensu	s Standard	l Deviation	2.68	
ommuni Results		Maximum	l		0.1713		Maximum	l		48.46	
Cor		Minimum			0.0402		Minimum			0.25	
		Ν			8		Ν			12	

 Table 14. Data summary table for cyanidin in dietary supplements.

					-		Arabinosid		5 11		
		SRM	4 3283 Ci	ranberry E	xtract (m	g/g)	SRI	M 3291 B	ilberry E	xtract (mg/	'g)
	Lab	А	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST										
	G701	0.150	0.160	0.160	0.157	0.006	53.2	53.1	48.4	51.6	2.7
	G702										
	G703	0.134	0.132	0.133	0.133	0.001	20.0	20.3	20.3	20.2	0.2
	G704										
	G705	0.157	0.175	0.154	0.162	0.011	42.8	42.5	42.5	42.6	0.2
	G707										
	G708	0.217	0.211	0.217	0.215	0.003	26.7	26.5	26.6	26.6	0.1
	G715						20.9	19.5		20.2	0.9
	G716										
	G717	0.210	0.211	0.215	0.212	0.003	21.5	20.9	19.3	20.6	1.2
	G718	0.235	0.236	0.235	0.235	0.001	25.2	24.2	25.1	24.8	0.6
	G722										
	G724	0.262	0.002	0.260	0.220	0.040	24.5	20.0	22.7	25.4	2.2
ts	G728	0.362	0.283	0.369	0.338	0.048	34.5	39.0	32.7	35.4	3.2
Individual Results	G729										
al R	G738 G739										
idu	G739 G740										
ndiv	G749										
I	G752										
	G753										
	G757										
	G760										
	G762	0.162	0.175	0.171	0.169	0.007	28.5	28.4	26.0	27.6	1.4
	G765	0.231	0.237	0.253	0.240	0.011	43.7	41.9	42.5	42.7	0.9
	G766	0.242	0.240	0.228	0.237	0.008	27.5	26.6	26.7	26.9	0.5
	G767										
	G771										
	G772										
	G773		0.130	0.130	0.130	0.000	19.7	19.7	17.5	19.0	1.3
	G774										
	G775						23.0	23.0	23.0	23.0	0.0
	G776										
	G777						42.3	42.4	42.3	42.3	0.1
ty		Consensu			0.198		Consensus			29.9	
Community Results				Deviation			Consensus	Standard	Deviation		
ommun Results		Maximum	L		0.338		Maximum			51.6	
C0 F		Minimum			0.130		Minimum			19.0	
		Ν			11		Ν			14	

 Table 15. Data summary table for cyanidin-3-arabinoside in dietary supplements.

				j		-	-	Galactosid		<i>J ~~</i> FF		
NIST 0.090 0.087 0.090 0.089 0.002 58.9 50.5 48.1 52.5 5.7 G702			SRM	A 3283 Ci	ranberry H	Extract (m	g/g)	SR	M 3291 E	Bilberry E	xtract (mg/	/g)
Figure G701 0.090 0.087 0.099 0.099 0.099 0.099 0.099 0.099 0.091 0.098 0.001 27.5 27.9 27.6 27.7 0.2 G702 G703 0.099 0.099 0.092 0.002 36.8 36.4 36.3 36.5 0.3 G706 G707 G708 0.149 0.150 0.155 0.151 0.000 26.3 27.1 27.0 26.8 0.44 G715 0.140 0.148 C0.155 0.157 0.004 34.3 34.0 33.7 34.0 0.3 G716 0.149 0.144 0.145 0.147 0.022 43.8 34.0 33.7 34.0 0.3 G718 0.152 0.159 0.157 0.004 34.3 33.2 34.5 34.0 0.7 G728 0.207 0.164 0.189 0.187 0.022 43.8 42.9 44.6 43.7 0.9 <		Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
Image: Note of the section o		NIST										
Image: Note of the section		G701	0.090	0.087	0.090	0.089	0.002	58.9	50.5	48.1	52.5	5.7
Nome Grand Grand <th< td=""><td></td><td>G702</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></th<>		G702										
Note Calce Out2 Out1 <		G703	0.099	0.099	0.097	0.098	0.001	27.5	27.9	27.6	27.7	0.2
Image: Note of the second se		G704										
Figure 1G7080.1490.1500.1510.00326.327.127.026.80.44G7150.1400.148 <td< td=""><td></td><td>G705</td><td>0.022</td><td>0.024</td><td>0.020</td><td>0.022</td><td>0.002</td><td>36.8</td><td>36.4</td><td>36.3</td><td>36.5</td><td>0.3</td></td<>		G705	0.022	0.024	0.020	0.022	0.002	36.8	36.4	36.3	36.5	0.3
Note:G7150.1400.1480.1440.00635.035.035.035.00.00G7160.1490.1440.1450.1460.00234.334.033.734.00.3G7180.1520.1590.1590.00434.333.234.534.00.7G7220.2070.1640.1890.1870.02243.842.944.643.70.9G7280.2070.1640.1890.1870.02243.842.944.643.70.9G7280.2070.1640.1890.1870.02243.842.944.643.70.9G7290.2070.1640.1890.1870.02243.842.944.643.70.9G7380.210.150.1870.02243.842.944.643.70.9G7390.210.1640.1870.0243.842.944.643.70.9G7300.210.1640.125.85.85.85.85.85.85.85.85.85.85.71.4G7500.1780.1780.1640.01258.555.856.757.01.41.4G7660.1590.1780.1640.01258.555.856.757.01.4G7670.1590.1690.1613.0.13.1.131.131.131.10.1G7760.159 <t< td=""><td></td><td>G707</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></t<>		G707										
Image: Note of the second se		G708	0.149	0.150	0.155	0.151	0.003	26.3	27.1	27.0	26.8	0.4
Normal G717 0.149 0.144 0.145 0.146 0.002 34.3 34.0 33.7 34.0 0.7 G718 0.152 0.159 0.159 0.157 0.004 34.3 33.2 34.5 34.0 0.7 G722			0.140	0.148		0.144	0.006	35.0	35.0		35.0	0.0
G7180.1520.1590.1590.1570.00434.333.234.534.00.7G722G7240.2070.1640.1890.1870.02243.842.944.643.70.9G729G7380.11G7390.11G7400.11G7520.11												
Nome Gamma Gamma <thg< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></thg<>												
G724G724G.0270.1640.1890.1870.02243.842.944.643.70.9G729G.297G.2070.1640.1890.1870.02243.842.944.643.70.9G738G.29G.207 <td< td=""><td></td><td></td><td>0.152</td><td>0.159</td><td>0.159</td><td>0.157</td><td>0.004</td><td>34.3</td><td>33.2</td><td>34.5</td><td>34.0</td><td>0.7</td></td<>			0.152	0.159	0.159	0.157	0.004	34.3	33.2	34.5	34.0	0.7
G7280.2070.1640.1890.1870.02243.842.944.643.70.9G729												
G729G729G729G729G730G7												
G752 G753 G754 G754 G753 G753 G754 G753 G753 G753 G753 G753 G713 G.113 O.111 O.002 37.4 37.2 34.4 36.3 1.7 G766 O.155 O.159 O.178 O.164 O.012 58.5 55.8 56.7 57.0 1.4 G766 O.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G767 G.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G771 G.130 O.140 O.160 O.143 O.015 27.3 28.3 25.4 27.0 1.5 G775 <td>s</td> <td></td> <td>0.207</td> <td>0.164</td> <td>0.189</td> <td>0.187</td> <td>0.022</td> <td>43.8</td> <td>42.9</td> <td>44.6</td> <td>43.7</td> <td>0.9</td>	s		0.207	0.164	0.189	0.187	0.022	43.8	42.9	44.6	43.7	0.9
G752 G753 G754 G754 G753 G753 G754 G753 G753 G753 G753 G753 G713 G.113 O.111 O.002 37.4 37.2 34.4 36.3 1.7 G766 O.155 O.159 O.178 O.164 O.012 58.5 55.8 56.7 57.0 1.4 G766 O.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G767 G.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G771 G.130 O.140 O.160 O.143 O.015 27.3 28.3 25.4 27.0 1.5 G775 <td>sult</td> <td></td>	sult											
G752 G753 G754 G754 G753 G753 G754 G753 G753 G753 G753 G753 G713 G.113 O.111 O.002 37.4 37.2 34.4 36.3 1.7 G766 O.155 O.159 O.178 O.164 O.012 58.5 55.8 56.7 57.0 1.4 G766 O.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G767 G.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G771 G.130 O.140 O.160 O.143 O.015 27.3 28.3 25.4 27.0 1.5 G775 <td>lRε</td> <td></td> <td></td> <td>_</td> <td></td> <td>_</td> <td>_</td> <td></td> <td>_</td> <td></td> <td>_</td> <td>_</td>	lRε			_		_	_		_		_	_
G752 G753 G754 G754 G753 G753 G754 G753 G753 G753 G753 G753 G713 G.113 O.111 O.002 37.4 37.2 34.4 36.3 1.7 G766 O.155 O.159 O.178 O.164 O.012 58.5 55.8 56.7 57.0 1.4 G766 O.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G767 G.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G771 G.130 O.140 O.160 O.143 O.015 27.3 28.3 25.4 27.0 1.5 G775 <td>dua</td> <td></td>	dua											
G752 G753 G754 G754 G753 G753 G754 G753 G753 G753 G753 G753 G713 G.113 O.111 O.002 37.4 37.2 34.4 36.3 1.7 G766 O.155 O.159 O.178 O.164 O.012 58.5 55.8 56.7 57.0 1.4 G766 O.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G767 G.173 O.152 O.156 O.160 O.011 38.1 37.7 37.8 37.9 O.2 G771 G.130 O.140 O.160 O.143 O.015 27.3 28.3 25.4 27.0 1.5 G775 <td>divi</td> <td></td>	divi											
G753 G757 Image: Second	In											
G757 G757 G760 Image: Constraint of the constrant of the constraint of the constraint of t												
G760G7760G7760G7760G.1100.1130.1110.0110.00237.437.234.436.31.7G7650.1550.1590.1780.1640.01258.555.856.757.01.4G7660.1730.1520.1560.1600.01138.137.737.837.90.2G767II												
G7620.1100.1130.1110.00237.437.234.436.31.7G7650.1550.1590.1780.1640.01258.555.856.757.01.4G7660.1730.1520.1560.1600.01138.137.737.837.90.2G767												
G7650.1550.1590.1780.1640.01258.555.856.757.01.4G7660.1730.1520.1600.01138.137.737.837.90.2G767			0.110	0.112	0.111	0.111	0.002	27.4	27.2	24.4	26.2	17
G7660.1730.1520.1560.1600.01138.137.737.837.90.2G767IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII												
G767 G767 Image: Second S												
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $			0.175	0.152	0.150	0.100	0.011	56.1	51.1	57.8	51.5	0.2
G772 G773 0.130 0.140 0.160 0.143 0.015 27.3 28.3 25.4 27.0 1.5 G774 427.0 1.50 6774 427.0 1.50 6774 6784 6784 6784 6784 6784 6784 6784 6784 6784 6784 6784 6784 6784												
G773 0.130 0.140 0.160 0.143 0.015 27.3 28.3 25.4 27.0 1.5 G774 I I IIII IIIII 31.1 31.1 31.1 31.1 0.0 G775 I I IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII												
G774 G775 G775 31.1 31.1 31.1 31.1 31.1 0.0 G776 G777 0.094 0.092 0.094 0.002 34.7 34.9 34.8 34.8 0.1 Maximum 0.132 Consensus Mean 0.132 Consensus Mean 35.5 7.3 Maximum 0.187 Maximum 57.0 57.0 Minimum 0.022 Minimum 26.8 57.0			0.130	0.140	0.160	0.143	0.015	27.3	28.3	25.4	27.0	1.5
G775 G775 31.1 31.1 31.1 31.1 31.1 0.0 G776 G777 0.094 0.092 0.096 0.094 0.002 34.7 34.9 34.8 34.8 0.1 G777 0.094 0.092 0.096 0.094 0.002 34.7 34.9 34.8 34.8 0.1 Maximum 0.132 Consensus Mean 0.132 Consensus Mean 35.5 7.3 Maximum 0.187 Maximum 57.0 57.0 57.0 Minimum 0.022 Minimum 26.8 57.0			01120	01110	01100	01110	0.010	2110	2010	2011	2110	110
G776 G777 0.094 0.092 0.096 0.094 0.002 34.7 34.9 34.8 34.8 0.1 No Consensus Mean 0.132 Consensus Mean 0.132 Consensus Standard Deviation 7.3 Maximum 0.187 Maximum 0.187 Maximum 57.0 Minimum 0.022 Minimum 26.8 Consensus								31.1	31.1	31.1	31.1	0.0
G777 0.094 0.092 0.096 0.094 0.002 34.7 34.9 34.8 34.8 0.1 Maximum 0.132 Consensus Mean 0.132 Consensus Mean 35.5 7.3 Maximum 0.187 Maximum 0.187 Maximum 57.0 Minimum 0.022 Minimum 26.8 26.8												
Normalized bitConsensus Mean0.132Consensus Mean35.5Consensus Standard Deviation0.039Consensus Standard Deviation7.3Maximum0.187Maximum57.0Minimum0.022Minimum26.8			0.094	0.092	0.096	0.094	0.002	34.7	34.9	34.8	34.8	0.1
Consensus Standard Deviation0.039Consensus Standard Deviation7.3Maximum0.187Maximum57.0Minimum0.022Minimum26.8			Consensu	s Mean	I			Consensus	Mean			
	unity ts				Deviation					l Deviation	7.3	
	umu sult											
	Con Rí		Minimum			0.022		Minimum			26.8	
	-		Ν			13		Ν			14	

 Table 16.
 Data summary table for cyanidin-3-galactoside in dietary supplements.

- 401		Cyanidin-3-Glucoside									
		SRM 3283 Cranberry Extract (mg/g)					SRM 3291 Bilberry Extract (mg/g)				
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST										
	G701	0.0086	0.0062	0.0068	0.0072	0.0012	93.4	83.5	88.0	88.3	4.9
	G702										
	G703	0.0136			0.0136		31.1	31.5	31.5	31.4	0.2
	G704										
	G705										
	G707										
	G708	0.0102	0.0100	0.0100	0.0101	0.0001	38.6	38.4	38.5	38.5	0.1
	G715						38.0	37.8		37.9	0.1
	G716										
	G717	0.0111	0.0108	0.0116	0.0112	0.0004	38.1	37.6	37.3	37.7	0.4
	G718	0.1110	0.1110	0.0930	0.1050	0.0104	93.1	90.0	93.3	92.1	1.9
sults	G722										
	G724										
	G728						47.2	46.0	48.3	47.2	1.1
	G729										
l Re	G738										
Individual Results	G739										
	G740										
	G749										
	G752										
	G753										
	G757			_							
	G760	0.0064	0.00.00	0.0000	0.0051	0.0000	10.0	40.0	25.5	20.0	1.0
	G762	0.0064	0.0068	0.0080	0.0071	0.0008	40.9	40.8	37.7	39.8	1.8
	G765	0.0180	0.0180	0.0160	0.0173	0.0012	62.8	60.1	60.7	61.2	1.4
	G766						42.0	41.0	41.1	41.4	0.5
	G767										
	G771 G772										
	G773						30.3	30.2	27.2	29.2	1.8
	G774						50.5	30.2	21.2	29.2	1.0
	G775	0.1711	0.1740	0.1643	0.1698	0.0050	33.9	34.0	34.0	34.0	0.1
	G776	0.1711	0.1740	0.1045	0.1098	0.0050	55.7	54.0	54.0	54.0	0.1
	G777	0.1400	0.1390	0.1440	0.1410	0.0026	38.1	38.2	38.2	38.2	0.0
Community Results	0,,,,	Consensu		011110	0.0586	0.0020	Consensus Mean			43.2	010
			s Standard	l Deviation			Consensus Standard Deviation				
		Maximum			0.1698		Maximum			92.1	
		Minimum			0.0071		Minimum			29.2	
 		N			8		N			13	
L					~						

 Table 17. Data summary table for cyanidin-3-glucoside in dietary supplements.

	Cyanidin Equivalents							*				
		SRM 3283 Cranberry Extract (mg/g)					SRM 3291 Bilberry Extract (mg/g)					
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD	
	NIST											
	G701	0.166	0.169	0.171	0.169	0.003	133.7	122.0	120.1	125.3	7.4	
	G702											
	G703	0.232	0.221	0.222	0.225	0.006	54.5	55.2	55.0	54.9	0.4	
	G704											
	G705	0.161	0.179	0.156	0.165	0.012	78.4	77.8	77.8	78.0	0.4	
	G707											
	G708	0.250	0.247	0.254	0.250	0.004	59.8	60.0	60.1	60.0	0.2	
	G715	0.184	0.193		0.189	0.006	62.7	61.7		62.2	0.8	
	G716											
ssults	G717	0.299	0.294	0.295	0.296	0.002	61.3	60.3	58.8	60.1	1.2	
	G718	0.389	0.393	0.379	0.387	0.007	99.4	95.9	99.6	98.3	2.1	
	G722								_			
	G724											
	G728	0.380	0.299	0.374	0.351	0.045	83.9	85.3	83.6	84.3	0.9	
	G729	0.007			0.007		24.5			24.5		
ıl Ro	G738	0.297			0.297		34.5			34.5		
Individual Results	G739 G740											
	G740 G749											
I	G752											
	G753											
	G757											
	G760											
	G762	0.311	0.325	0.319	0.318	0.007	73.5	72.9	67.3	71.2	3.4	
	G765	0.269	0.275	0.297	0.281	0.015	108.7	104.0	105.4	106.1	2.4	
	G766	0.276	0.262	0.256	0.265	0.011	70.0	68.5	68.7	69.1	0.8	
	G767											
	G771											
	G772	1.174	1.172	1.133	1.160	0.023	64.2	67.5	67.7	66.5	2.0	
	G773	0.083	0.179	0.191	0.151	0.059	55.0	55.7	50.1	53.6	3.1	
	G774											
	G775	0.109	0.111	0.105	0.109	0.003	59.5	59.6	59.7	59.6	0.1	
	G776											
	G777	0.323	0.317	0.325	0.322	0.004	124.0	124.1	124.2	124.1	0.1	
Community Results		Consensu			0.259		Consensus Mean 76.5					
		Consensu	s Standard	Deviation			Consensus	s Standard				
		Maximum			1.160		Maximum			125.3		
C0 F		Minimum			0.109		Minimum			34.5		
		Ν			15		Ν			15		

 Table 18. Data summary table for cyanidin equivalents in dietary supplements.

Lan	- 1/• L	Julu Sulli	mary ta				hinidin	ppiente			
		SRM	[3283 Cr	anberry E	xtract (m	-		M 3291 R	ilberrv Ex	ktract (mg/	/g)
	Lab	A	В	C	Avg	SD	A	B	C	Avg	SD
	NIST		2	U		52		2	0		02
	G701										
	G702										
	G703	0.011	0.015		0.013	0.003	4.50	4.39	4.48	4.46	0.06
	G704										
	G705						4.08	3.36	3.29	3.58	0.44
	G707										
	G708										
	G715						0.63	1.33		0.98	0.49
	G716										
	G717	0.007	0.007	0.006	0.007	0.000	0.91	0.83	0.92	0.89	0.05
	G718						0.97	0.92	0.96	0.95	0.03
	G722										
	G724										
	G728						1.17	1.21	1.23	1.20	0.03
ults	G729										
Individual Results	G738										
lual	G739										
livid	G740										
Ind	G749										
	G752										
	G753										
	G757										
	G760										
	G762	0.038	0.039	0.037	0.038	0.001	3.38	3.54	3.12	3.35	0.21
	G765						0.47	0.43	0.45	0.45	0.02
	G766										
	G767										
	G771										
	G772										
	G773	0.150	0.130		0.140	0.014	10.30	10.60	7.98	9.63	1.43
	G774										
	G775						2.92	3.00	3.01	2.97	0.05
	G776										
	G777	0.557	0.547	0.573	0.559	0.013	17.37	17.36	17.20	17.31	0.10
ity		Consensus		Derivi	0.116		Consensus		Derivi	4.23	
Community Results		Consensus		Deviation			Consensus		Deviation		
omr Res		Maximum Minimum			0.559		Maximum Minimum			17.31	
Ü		Minimum N			0.007		Minimum			0.45	
		Ν			6		Ν			12	

 Table 19. Data summary table for delphinidin in dietary supplements.

			j		-		Equivalents		<u> </u>	prement	
		SRN	A 3283 Ci	ranberry I	Extract (m	g/g)	SR	M 3291 B	Silberry E	xtract (mg/	g)
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST										
	G701	0.166	0.169	0.171	0.169	0.003	133.7	122.0	120.1	125.3	7.4
	G702										
	G703	0.232	0.221	0.222	0.225	0.006	54.5	55.2	55.0	54.9	0.4
	G704										
	G705	0.161	0.179	0.156	0.165	0.012	78.4	77.8	77.8	78.0	0.4
	G707										
	G708	0.250	0.247	0.254	0.250	0.004	59.8	60.0	60.1	60.0	0.2
	G715	0.184	0.193		0.189	0.006	62.7	61.7		62.2	0.8
	G716	0.000	0.004	0.005	0.006	0.000	<i>61.0</i>	60 0	5 0.0	60.4	1.0
	G717	0.299	0.294	0.295	0.296	0.002	61.3	60.3	58.8	60.1	1.2
	G718	0.389	0.393	0.379	0.387	0.007	99.4	95.9	99.6	98.3	2.1
	G722										
	G724 G728	0.380	0.299	0.374	0.351	0.045	83.9	85.3	83.6	84.3	0.9
lts	G729	0.380	0.299	0.374	0.331	0.045	03.9	85.5	83.0	64.5	0.9
esu	G738	0.297			0.297		34.5			34.5	
Individual Results	G739	0.277			0.277		54.5			54.5	
	G740										
ndi	G749										
-	G752										
	G753										
	G757										
	G760										
	G762	0.311	0.325	0.319	0.318	0.007	73.5	72.9	67.3	71.2	3.4
	G765	0.269	0.275	0.297	0.281	0.015	108.7	104.0	105.4	106.1	2.4
	G766	0.276	0.262	0.256	0.265	0.011	70.0	68.5	68.7	69.1	0.8
	G767										
	G771										
	G772	1.174	1.172	1.133	1.160	0.023	64.2	67.5	67.7	66.5	2.0
	G773	0.083	0.179	0.191	0.151	0.059	55.0	55.7	50.1	53.6	3.1
	G774										
	G775	0.109	0.111	0.105	0.109	0.003	59.5	59.6	59.7	59.6	0.1
	G776										
	G777	0.323	0.317	0.325	0.322	0.004	124.0	124.1	124.2	124.1	0.1
ity		Consensu			0.259		Consensus		D : .	76.5	
Community Results				l Deviation			Consensus		Deviation		
ommun Results		Maximum			1.160		Maximum			125.3	
Ŭ		Minimum			0.109		Minimum			34.5	
		Ν			15		Ν			15	

 Table 20.
 Data summary table for delphinidin-3-arabinoside in dietary supplements.

			Delphinidin-3-galactoside in circlary supprements. Delphinidin-3-Galactoside SRM 3283 Cranberry Extract (mg/g) SRM 3291 Bilberry Extract (mg/g)								
		SRM	• •								
	Lab										SD
	NIST									0	
	G701										
	G702										
	G703						35.2	35.7	35.6	35.5	0.3
	G704										
	G705						47.5	47.0	47.2	47.2	0.3
	G707										
	G708						43.6	43.6	43.7	43.6	0.1
	G715						42.2	42.5		42.4	0.2
	G716										
	G717						44.4	43.9	43.6	44.0	0.4
	G718						42.7	41.4	43.0	42.3	0.9
	G722										
	G724										
	G728						48.7	47.7	48.3	48.2	0.5
Individual Results	G729										
Re	G738										
lual	G739										
livid	G740										
Ind	G749										
	G752										
	G753										
	G757										
	G760										
	G762	0.0020	0.0069	0.0030	0.0040	0.0026	45.1	45.3	41.6	44.0	2.1
	G765						61.1	58.4	59.2	59.6	1.4
	G766						31.0	30.9	31.0	30.9	0.1
	G767										
	G771					_		_		_	
	G772										
	G773					_	34.7	34.8	31.4	33.6	1.9
	G774										
	G775	0.0488	0.0495	0.0464	0.0482	0.0016	38.0	38.3	38.3	38.2	0.2
	G776										
	G777	0.0040	0.0040	0.0040	0.0040	0.0000	42.1	42.1	42.1	42.1	0.0
ity		Consensu		D	0.0187		Consensus			41.9	
Community Results			s Standard	Deviation			Consensus Standard Deviation				
ommun Results		Maximum			0.0482		Maximum		59.6		
ວົ		Minimum			0.0040 Minimum			30.9			
		Ν			3		Ν			13	

 Table 21. Data summary table for delphinidin-3-galactoside in dietary supplements.

- 401			innary ta	.010 101 (±	0	-3-Glucosia		i y supp	tements.	
		SRM	A 3283 Ci	anberry I	Extract (m	-			Bilberry E	xtract (mg	/g)
	Lab	Α	В	C	Avg	SD	Α	В	C	Avg	SD
	NIST			-	8					8	~_
	G701						64.7	52.3	53.2	56.7	6.9
	G702										
	G703						38.1	38.6	38.6	38.4	0.3
	G704										
	G705						50.9	50.5	50.6	50.7	0.2
	G707										
	G708						46.5	46.2	46.5	46.4	0.2
	G715						45.2	45.2		45.2	0.0
	G716										
	G717						47.0	46.4	46.2	46.5	0.4
	G718	0.0530	0.0510	0.0570	0.0537	0.0031	45.7	44.2	45.9	45.2	0.9
	G722										
	G724										
	G728						52.2	51.1	51.3	51.5	0.6
Individual Results	G729										
l Re	G738										
dual	G739										
divio	G740										
In	G749										
	G752										
	G753										
	G757										
	G760	0.0242	0.0202	0.0100	0.0214	0.0024	40.2	40.4	44.5	47 1	2.2
	G762 G765	0.0242	0.0202	0.0198	0.0214	0.0024	48.3 64.7	48.4	44.5	47.1 63.1	2.2 1.5
	G766						33.1	61.8 32.8	62.7 33.0	33.0	0.1
	G767						55.1	32.8	55.0	55.0	0.1
	G771										
	G772										
	G773						36.7	36.9	33.4	35.7	2.0
	G774						2017	2017	5511	0011	210
	G775	0.1363	0.1370	0.1324	0.1352	0.0025	40.3	40.7	40.7	40.6	0.2
	G776										
	G777	0.0170	0.0160	0.0180	0.0170	0.0010	44.9	45.1	45.1	45.1	0.1
		Consensu			0.0568		Consensu			45.8	
Community Results		Consensu	s Standard	Deviation			Consensu	s Standard	l Deviation	7.8	
ommun Results		Maximum			0.1352		Maximum			63.1	
Con R(Minimum			0.0170		Minimum			33.0	
		Ν			4		Ν			14	

 Table 22. Data summary table for delphinidin-3-glucoside in dietary supplements.

		and built	Delphinidin Equivalents in dietary supplements. Delphinidin Equivalents SRM 3283 Cranberry Extract (mg/g) SRM 3291 Bilberry Extract (mg/g)								
		SRM 3283 Cranberry Extract (mg/g) SRM 3291 Bilberry Extract (mg/g)							/g)		
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST				0					0	
	G701						42.2	34.0	34.7	37.0	4.5
	G702										
	G703	0.011	0.015		0.013	0.003	72.8	73.6	73.8	73.4	0.5
	G704										
	G705	0.076	0.085	0.074	0.078	0.006	96.8	95.2	95.5	95.9	0.8
	G707										
	G708						92.3	91.3	91.9	91.8	0.5
	G715						83.5	84.5		84.0	0.7
	G716										
	G717	0.007	0.007	0.006	0.007	0.000	87.3	86.0	85.7	86.3	0.8
	G718	0.035	0.033	0.037	0.035	0.002	83.8	81.1	84.2	83.0	1.7
	G722										
	G724										
	G728						96.9	94.6	95.3	95.6	1.2
sults	G729										
Individual Results	G738	0.087			0.087		41.7			41.7	
lual	G739										
livid	G740										
Inc	G749										
	G752										
	G753										
	G757		_		_	_		_			
	G760										
	G762	0.055	0.056	0.052	0.054	0.002	92.2	92.4	84.9	89.8	4.3
	G765						120.4	114.9	116.6	117.3	2.8
	G766						60.0	59.5	59.8	59.7	0.2
	G767										
	G771	0.204	0.202	0.260	0.270	0.000	02.2	06.0	96.0	057	2.0
	G772	0.384	0.383	0.369	0.379	0.008	83.3	86.8	86.9	85.7	2.0
	G773	0.150	0.130		0.140	0.014	77.5	77.7	68.5	74.6	5.2
	G774	0.121	0.122	0.117	0.120	0.002	76.9	77 5	77.6	2 77	0.4
	G775	0.121	0.122	0.117	0.120	0.003	76.8	77.5	77.6	77.3	0.4
	G776 G777	0.580	0.566	0.594	0.580	0.014	100.4	100.6	100.3	100.4	0.1
	0///	Consensus		0.374	0.380	0.014	Consensu		100.3	84.7	0.1
nity ;				l Deviation					l Deviation		
Community Results		Maximum		. De viation	0.166		Maximum			117.3	
lom Re		Minimum			0.007		Minimum			37.0	
C		N			9		N			15	
		14)		14			15	

 Table 23. Data summary table for delphinidin equivalents in dietary supplements.

1 aux	- 47. 1	Jala Sull	iiiiai y ta				ary supp. vidin		5.		
		SRM	A 3283 Ci	ranberry H	Extract (m			M 3291 H	Bilberry Ex	tract (mg	/g)
	Lab	Α	В	C	Avg	SD	A	B	C	Avg	SD
	NIST			-	8				-	8	~-
	G701										
	G702										
	G703						1.19	1.19	1.21	1.20	0.01
	G704										
	G705						0.99	0.90	0.93	0.94	0.04
	G707										
	G708										
	G715	0.0300	0.0350		0.0325	0.0035	0.75	0.71		0.73	0.03
	G716										
	G717						0.34	0.34	0.32	0.33	0.01
	G718	0.0680	0.0640	0.0660	0.0660	0.0020	0.26	0.25	0.25	0.25	0.00
	G722										
	G724										
	G728						0.76	0.64	0.62	0.67	0.08
ults	G729										
Individual Results	G738										
ual	G739										
livid	G740										
Ind	G749										
	G752										
	G753										
	G757										
	G760										
	G762										
	G765										
	G766										
	G767										
	G771										
	G772										
	G773						2.06	1.60	2.51	2.06	0.46
	G774										
	G775						1.57	1.57	1.24	1.46	0.19
	G776										
	G777	G					4.72	4.68	4.81	4.74	0.07
ity		Consensu			0.0493		Consensus			1.11	
Community Results			is Standard	Deviation				Standard	1 Deviation	0.82	
ommun Results		Maximum			0.0660		Maximum			4.74	
Ŭ C		Minimum			0.0325		Minimum			0.25	
		Ν			2		Ν			10	

 Table 24. Data summary table for malvidin in dietary supplements.

Tubh			innary to				Arabinosi		ry suppr	ements.	
		SRN	A 3283 Ci	ranberry I					Bilberry E	xtract (mg	/g)
	Lab	Α	В	C	Avg	SD	Α	В	C	Avg	SD
	NIST			-	8				-	8	~-
	G701										
	G702										
	G703						6.89	7.12	7.08	7.03	0.12
	G704										
	G705						9.25	9.14	9.19	9.19	0.06
	G707										
	G708	0.0959	0.0966	0.1070	0.0998	0.0062	11.20	11.00	11.10	11.10	0.10
	G715						8.87	9.00		8.94	0.09
	G716										
	G717						8.43	8.23	8.40	8.35	0.11
	G718						9.21	8.48	8.71	8.80	0.37
	G722										
	G724										
	G728						12.34	11.36	12.04	11.91	0.50
ults	G729										
Individual Results	G738										
ual	G739										
ivid	G740										
Ind	G749										
	G752										
	G753										
	G757										
	G760										
	G762						10.50	10.20	9.47	10.06	0.53
	G765						11.52	11.11	11.79	11.47	0.34
	G766						5.03	4.84	4.91	4.93	0.10
	G767										
	G771										
	G772										
	G773						1.24	1.04	0.96	1.08	0.14
	G774										
	G775						7.87	7.85	7.82	7.85	0.03
	G776										
	G777	0.0030	0.0030	0.0030	0.0030	0.0000	1.50	1.49	1.52	1.50	0.01
ty		Consensu			0.0514		Consensu			8.19	
Community Results				l Deviation					l Deviation		
ommun Results		Maximum			0.0998		Maximum			11.91	
C0 F		Minimum			0.0030		Minimum			1.08	
		Ν			2		Ν			13	

 Table 25. Data summary table for malvidin-3-arabinoside in dietary supplements.

Malvidin-3-Galactoside SRM 3283 Cranberry Extract (mg/g) SRM 3291 Bilberry Extract (mg/g)											
		SRN	A 3283 Ci	ranberry l	Extract (m	g/g)	SR	M 3291 H	Bilberry E	xtract (mg	/g)
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST										
	G701	0.1400	0.1400	0.1600	0.1467	0.0115	26.1	23.1	21.0	23.4	2.5
	G702										
	G703	0.0125			0.0125		12.1	12.3	12.3	12.2	0.1
	G704										
	G705						16.3	16.1	16.2	16.2	0.1
	G707										
	G708										
	G715						12.5	12.6		12.5	0.1
	G716										
	G717						38.1	37.1	38.6	37.9	0.8
	G718	0.0520			0.0520		14.9	14.2	14.7	14.6	0.3
	G722										
	G724										
s	G728						19.2	18.1	19.4	18.9	0.7
Individual Results	G729										
l Re	G738										
dua	G739										
divi	G740										
In	G749										
	G752		_			_		_			
	G753										
	G757										
	G760	0.0252	0.0050	0.0050	0.0044	0.0010	14.0	110	12.0	12.5	0.6
	G762	0.0252	0.0270	0.0270	0.0264	0.0010	14.0	14.0	13.0	13.7	0.6
	G765						18.3	17.4	17.8	17.8	0.4
	G766						6.7	6.6	6.8	6.7	0.1
	G767 G771										
	G772										
	G773						27.9	27.9	25.3	27.0	1.5
	G774						21.9	21.9	25.5	27.0	1.5
	G775						33.3	33.3	33.2	33.3	0.1
	G776						55.5	55.5	55.2	55.5	0.1
	G777	0.0060	0.0050	0.0050	0.0053	0.0006	13.3	13.3	13.3	13.3	0.0
		Consensu			0.0595		Consensu			18.4	
nity s			s Standard	Deviation					l Deviation		
ommun Results		Maximum			0.1467		Maximum			37.9	
Community Results		Minimum			0.0053		Minimum			6.7	
		N			3		N			13	
L										-	

 Table 26. Data summary table for malvidin-3-galactoside in dietary supplements.

- 401	• 1	- ata Ball						lin-3-Glucoside				
		SRN	SRM 3283 Cranberry Extract (mg/g) SRM 3291 Bilberry A B C Avg SD A B C						Bilberry Ex	ktract (mg	/g)	
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD	
	NIST											
	G701											
	G702											
	G703						30.2	30.6	30.6	30.5	0.2	
	G704											
	G705											
	G707											
	G708	0.1720	0.1780	0.1800	0.1767	0.0042	37.3	37.0	37.2	37.2	0.2	
	G715	0.0270	0.0320		0.0295	0.0035	34.7	34.8		34.8	0.0	
	G716											
	G717	0.0128	0.0110	0.0117	0.0118	0.0009	14.5	14.3	14.2	14.3	0.2	
	G718						36.5	35.4	36.6	36.2	0.7	
	G722											
	G724											
	G728						43.8	42.5	43.3	43.2	0.6	
Individual Results	G729											
Re	G738											
lual	G739											
livid	G740											
Inc	G749											
	G752											
	G753											
	G757											
	G760											
	G762						35.8	35.6	32.9	34.8	1.6	
	G765						45.9	43.9	44.4	44.7	1.0	
	G766	0.1323	0.1340	0.1370	0.1344	0.0024	21.6	21.2	21.2	21.3	0.2	
	G767											
	G771		_					_				
	G772											
	G773						10.2	11.4	9.7	10.4	0.9	
	G774											
	G775		_				13.4	13.5	13.5	13.4	0.0	
	G776											
	G777						35.7	35.8	35.8	35.8	0.1	
ity		Consensu		·	0.0881		Consensus			29.7		
Community Results				l Deviation					l Deviation			
ommun Results		Maximum			0.1767		Maximum			44.7		
C		Minimum			0.0118		Minimum			10.4		
		Ν			4		Ν			12		

 Table 27. Data summary table for malvidin-3-glucoside in dietary supplements.

- 4.01	0, _		<u>iiiidi y</u> ta			-	Equivalents	~	suppren	10111051	
		SRM	M 3283 Ci	ranberry l	Extract (m	g/g)	SR	M 3291 I	Bilberry E	xtract (mg	/g)
	Lab	Α	В	C	Avg	SD	Α	В	C	Avg	SD
	NIST				0					0	
	G701	0.0940	0.0940	0.1074	0.0985	0.0078	17.5	15.5	14.1	15.7	1.7
	G702										
	G703	0.0084			0.0084		34.5	35.1	35.1	34.9	0.3
	G704										
	G705						18.5	18.2	18.4	18.4	0.1
	G707										
	G708	0.1840	0.1886	0.1974	0.1900	0.0068	33.1	32.7	32.9	32.9	0.2
	G715	0.0481	0.0565		0.0523	0.0059	38.8	38.9		38.9	0.1
	G716										
	G717	0.0086	0.0074	0.0079	0.0079	0.0006	41.7	40.7	41.8	41.4	0.6
	G718	0.1029	0.0640	0.0660	0.0776	0.0219	41.4	39.7	40.9	40.7	0.9
	G722										
	G724										
	G728						51.8	49.4	51.3	50.9	1.3
sult	G729										
Individual Results	G738						6.1			6.1	
dual	G739										
divi	G740										
In	G749										
	G752										
	G753										
	G757			_							
	G760	0.04.40					10.0	10.5		.	1.0
	G762	0.0169	0.0181	0.0181	0.0177	0.0007	40.9	40.6	37.6	39.7	1.8
	G765	0.0000	0.0000	0.0020	0.0002	0.001.6	51.3	49.1	50.2	50.2	1.1
	G766	0.0888	0.0900	0.0920	0.0903	0.0016	22.6	22.1	22.3	22.3	0.2
	G767										
	G771 G772						38.3	41.2	42.8	40.8	2.3
	G773						28.5	28.7	42.8 26.7	40.8 28.0	1.1
	G774						28.3	20.7	20.7	28.0	1.1
	G775						38.6	38.5	38.1	38.4	0.2
	G776						50.0	50.5	50.1	50.4	0.2
	G777	0.0062	0.0055	0.0055	0.0057	0.0004	38.7	38.8	38.9	38.8	0.1
	2,77	Consensu		0.0000	0.0624	0.0001	Consensu		20.7	34.2	5.1
nity s			is Standard	Deviation					l Deviation		
ommun Results		Maximum			0.1900		Maximum			50.9	
Community Results		Minimum			0.0057		Minimum			6.1	
		N			8		N			15	
		*1			0		- 1			15	

 Table 28. Data summary table for malvidin equivalents in dietary supplements.

		vata san	iiiiai y ta		<u>coman</u>	Peor	ary supp nidin				
		SRM	A 3283 Ci	ranberry H	Extract (m			M 3291 E	Bilberry Ex	xtract (mg	/g)
[Lab	Α	В	C	Avg	SD	Α	В	C	Avg	SD
	NIST				8					8	
	G701										
	G702										
	G703	0.0507	0.0526	0.0549	0.0527	0.0021	0.468	0.479	0.491	0.479	0.012
	G704										
	G705	0.0372	0.0411	0.0355	0.0379	0.0029	0.427	0.491	0.418	0.445	0.040
	G707										
	G708										
	G715	0.0510	0.0460		0.0485	0.0035	0.234	0.234		0.234	0.000
	G716										
	G717	0.0507	0.0507	0.0498	0.0504	0.0005	0.200	0.206	0.223	0.210	0.012
	G718		0.0510		0.0510		0.242	0.229	0.229	0.233	0.008
	G722										
	G724										
5	G728						0.299	0.316	0.323	0.313	0.012
sult	G729										
Individual Results	G738										
dual	G739										
divi	G740										
In	G749										
	G752		_		_			_		_	
	G753										
	G757										
	G760										
	G762										
	G765										
	G766										
	G767 G771										
	G772										
	G773						1.270	1.490	1.220	1.327	0.144
	G774						1.270	1.470	1.220	1.527	0.144
	G775						1.307	1.082	1.452	1.280	0.186
	G776						1.507	1.002	1.452	1.200	0.100
	G777	0.1260	0.1220	0.1320	0.1267	0.0050	1.491	1.476	1.534	1.500	0.030
		Consensu		I	0.0505		Consensu		I	0.669	
Community Results				l Deviation					Deviation		
ommun Results		Maximum			0.1267		Maximum			1.500	
Con Ré		Minimum			0.0379		Minimum			0.210	
		Ν			6		Ν			10	

 Table 29. Data summary table for peonidin in dietary supplements.

			u				Arabinosio				
		SRM	A 3283 Ci	ranberry E	Extract (m	g/g)	SR	M 3291 E	Bilberry Ex	ktract (mg	/g)
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST										
	G701	0.168	0.163	0.174	0.168	0.006	23.97	23.36	20.93	22.75	1.61
	G702										
	G703	0.103	0.101	0.107	0.104	0.003	1.63	1.67	1.65	1.65	0.02
	G704										
	G705	0.129	0.138	0.123	0.130	0.008	2.54	2.46	2.50	2.50	0.04
	G707										
	G708										
	G715	0.134	0.148		0.141	0.010	2.12	2.15		2.14	0.02
	G716										
	G717	0.165	0.165	0.171	0.167	0.003	1.77	1.69	1.58	1.68	0.10
	G718	0.192	0.195	0.192	0.193	0.002	2.12	2.02	2.09	2.08	0.05
	G722										
	G724	0.245	0.102	0.225	0.219	0.022	2.90	2 59	2.90	2.79	0.19
lts	G728 G729	0.245	0.183	0.225	0.218	0.032	3.89	3.58	3.89	3.78	0.18
esul	G729 G738										
Individual Results	G739										
	G740										
	G749										
Ι	G752										
	G753										
	G757										
	G760										
	G762	0.140	0.144	0.140	0.141	0.002	2.61	2.58	2.41	2.53	0.11
	G765	0.176	0.179	0.182	0.179	0.003	3.37	3.40	3.51	3.43	0.07
	G766										
	G767										
	G771										
	G772										
	G773			0.150	0.150		1.61	1.54	1.35	1.50	0.13
	G774										
	G775	0.045	0.045	0.050	0.046	0.003	2.01	1.99	1.93	1.98	0.04
	G776										
	G777	0.108	0.106	0.112	0.109	0.003	2.21	2.23	2.24	2.23	0.01
ity		Consensu			0.148		Consensu			2.44	
Community Results				Deviation					Deviation		
ommun Results		Maximum			0.218		Maximum			22.75	
ວົ		Minimum			0.046		Minimum			1.50	
		Ν			11		Ν			12	

 Table 30. Data summary table for peonidin-3-arabinoside in dietary supplements.

			j	<u>r</u>		Peonidin-3-Galactoside							
		SRM	/I 3283 Ci	ranberry E	xtract (m	g/g)	SRM	A 3291 B	ilberry E	xtract (mg/	/g)		
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD		
	NIST												
	G701	0.177	0.173	0.177	0.176	0.002	1.67	0.76	0.82	1.08	0.51		
	G702												
	G703	0.156	0.157	0.160	0.158	0.002	3.73	3.81	3.80	3.78	0.04		
	G704												
	G705	0.178	0.199	0.176	0.184	0.013	5.72	5.70	5.68	5.70	0.02		
	G707												
	G708	0.211	0.211	0.218	0.213	0.004	4.56	4.49	4.54	4.53	0.04		
	G715	0.177	0.204		0.191	0.019	8.55	8.19		8.37	0.25		
	G716												
	G717	0.221	0.223	0.224	0.223	0.002	3.76	3.78	3.97	3.84	0.12		
	G718	0.241	0.260	0.251	0.251	0.010	4.65	4.49	4.64	4.59	0.09		
	G722												
	G724												
s	G728	0.456	0.211	0.404	0.357	0.129	9.67	9.08	10.32	9.69	0.62		
sult	G729												
Individual Results	G738												
	G739												
	G740												
In	G749												
	G752		_		_	_		_		_			
	G753												
	G757												
	G760	0.172	0.176	0.172	0.174	0.000	5.15	c 17	1.60	5.00	0.00		
	G762	0.172	0.176	0.173	0.174	0.002	5.15	5.17	4.68	5.00	0.28		
	G765	0.235	0.240	0.243	0.239	0.004	7.47	7.06	7.23	7.25	0.21		
	G766	0.197	0.218	0.199	0.205	0.012	4.40	3.98	4.00	4.13	0.24		
	G767												
	G771												
	G772 G773	0.077	0.078	0.100	0.085	0.013	2.28	2.18	1.84	2.10	0.23		
		0.077	0.078	0.100	0.085	0.013	2.20	2.10	1.04	2.10	0.23		
	G774 G775						4.25	4.25	4.20	4.23	0.03		
	G776						4.23	4.23	4.20	4.23	0.05		
	G777						13.02	13.02	13.09	13.04	0.04		
	0,77	Consensu	s Mean		0.201		Consensus		15.07	5.29	0.07		
nity s				Deviation					Deviation				
ommun Results		Maximum			0.043Consensus Standard Deviation0.357Maximum				_ • • • • • • • • • • • • • • • • • • •	13.04			
Community Results		Minimum								1.08			
C0 F		N			12		N			14			
					12		· '			17			

 Table 31. Data summary table for peonidin-3-galactoside in dietary supplements.

Peonidin-3-Glucoside											
		SRN	A 3283 Ci	ranberry H					Bilberry Ex	ktract (mg/	/g)
	Lab	Α	В	C	Avg	SD	Α	В	C	Avg	SD
	NIST				0					0	
	G701										
	G702										
	G703	0.0286	0.0267	0.0235	0.0263	0.0026	14.2	14.4	14.4	14.3	0.1
	G704										
	G705	0.0245	0.0252	0.0224	0.0240	0.0015	18.9	18.7	18.7	18.8	0.1
	G707										
	G708	0.0367	0.0385	0.0710	0.0487	0.0193	31.7	31.6	31.6	31.6	0.1
	G715						16.2	16.3		16.2	0.1
	G716										
	G717	0.0298	0.0313	0.0296	0.0302	0.0009	16.9	16.8	16.6	16.8	0.1
	G718	0.1290	0.1350	0.1260	0.1300	0.0046	17.3	16.6	17.1	17.0	0.3
	G722										
	G724										
	G728						23.9	23.6	24.1	23.9	0.3
sults	G729										
Re	G738										
Individual Results	G739										
	G740										
Inc	G749										
	G752										
	G753										
	G757										
	G760										
	G762	0.0084	0.0089	0.0080	0.0084	0.0005	18.1	18.1	16.6	17.6	0.9
	G765	0.0320	0.0340	0.0450	0.0370	0.0070	28.0	26.6	27.7	27.4	0.7
	G766						16.8	16.3	16.1	16.4	0.4
	G767										
	G771										
	G772						15.5		10.6		1.0
	G773						15.5	14.6	13.6	14.6	1.0
	G774	0.1000	0.1104	0.11.40	0 11 4 4	0.0050	15.0	150	15.7	15.0	0.0
	G775	0.1090	0.1194	0.1148	0.1144	0.0052	15.8	15.8	15.7	15.8	0.0
	G776	0.0190	0.01/0	0.0170	0.0170	0.0010	17.0	17 1	12.0	157	2.2
	G777	0.0180 Consensu	0.0160	0.0170	0.0170	0.0010	17.0 Consensus	17.1 Mean	13.0	15.7 17.5	2.3
uity :				Deviation	0.0467				l Deviation		
Community Results		Maximum			0.0454		Maximum			31.6	
om		Minimum			0.1300		Minimum			14.3	
C		N			0.0084 9		N			14.5	
		IN			フ		1N			13	

 Table 32. Data summary table for peonidin-3-glucoside in dietary supplements.

		Peonidin Equivalents Peonidin Equivalents SRM 3283 Cranberry Extract (mg/g)											
		SRM	A 3283 Ci	ranberry I	Extract (m	g/g)	SR	M 3291 F	Bilberry E	xtract (mg	/g)		
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD		
	NIST												
	G701	0.232	0.226	0.236	0.231	0.005	17.7	16.7	15.1	16.5	1.3		
	G702												
	G703	0.242	0.242	0.249	0.244	0.004	13.3	13.5	13.5	13.4	0.1		
	G704												
	G705	0.259	0.283	0.250	0.264	0.017	18.2	18.1	18.0	18.1	0.1		
	G707												
	G708	0.161	0.162	0.188	0.170	0.015	23.6	23.5	23.5	23.5	0.1		
	G715	0.259	0.281		0.270	0.016	17.8	17.6		17.7	0.1		
	G716												
	G717	0.328	0.331	0.333	0.331	0.003	14.9	14.7	14.7	14.8	0.1		
	G718	0.374	0.443	0.379	0.399	0.039	16.0	15.4	15.8	15.7	0.3		
	G722												
so.	G724												
	G728	0.467	0.264	0.419	0.383	0.106	24.8	24.0	25.4	24.8	0.7		
Individual Results	G729												
l Re	G738	0.092			0.092		2.5			2.5			
lua	G739												
divi	G740												
Inc	G749												
	G752												
	G753												
	G757												
	G760												
	G762	0.215	0.220	0.215	0.217	0.003	16.9	16.9	15.5	16.5	0.8		
	G765	0.296	0.303	0.314	0.304	0.009	25.4	24.2	25.1	24.9	0.6		
	G766	0.128	0.142	0.129	0.133	0.008	13.8	13.2	13.0	13.3	0.4		
	G767												
	G771		_			_		_			_		
	G772	0.642	0.641	0.600	0.628	0.024	15.5	16.6	17.3	16.5	0.9		
	G773	0.050	0.051	0.169	0.090	0.069	13.9	13.5	12.2	13.2	0.9		
	G774												
	G775	0.102	0.109	0.109	0.107	0.004	15.7	15.5	15.7	15.6	0.1		
	G776								• • •				
	G777	0.213	0.206	0.221	0.213	0.007	22.6	22.6	20.1	21.7	1.4		
ity		Consensu			0.252		Consensus Mean17.6Consensus Standard Deviation4.4						
nun ults				Deviation									
Community Results		Maximum			0.628		Maximum			24.9			
		Minimum			0.090								
		Ν			15		Ν			15			

Table 33. Data summary table for peonidin equivalents in dietary supplements.

1 aux		Jala Sulli	mary ta				tary supj midin	Jement			
		SRM	I 3283 Ci	anberry E	xtract (m			M 3291 F	Bilberry E	xtract (mg	/g)
	Lab	A	В	C	Avg	SD	A	В	C	Avg	SD
	NIST				8	~-			-	8	~-
	G701										
	G702										
	G703						1.230	1.210	1.230	1.223	0.012
	G704										
	G705						0.750	0.742	0.765	0.752	0.012
	G707										
	G708										
	G715	0.086	0.107		0.097	0.015	0.766	0.724		0.745	0.030
	G716										
	G717						0.576	0.556	0.586	0.573	0.015
	G718						0.350	0.324	0.343	0.339	0.013
	G722										
	G724										
	G728						0.724	0.562	0.531	0.606	0.104
sults	G729										
Re	G738										
lual	G739										
Individual Results	G740										
Inc	G749										
	G752										
	G753										
	G757										
	G760										
	G762										
	G765						0.460	0.540	0.490	0.497	0.040
	G766										
	G767										
	G771										
	G772	0.150	0.190	0.190	0.170	0.017	0.540	0.670	0.610	0.007	0.065
	G773	0.150	0.180	0.180	0.170	0.017	0.540	0.670	0.610	0.607	0.065
	G774						1 497	1 479	1 471	1 470	0.009
	G775 G776						1.487	1.478	1.471	1.479	0.008
	G777						0.961	0.959	0.941	0.954	0.011
	0///	Consensus	Mean		0.122		Consensu		0.941	0.934	0.011
nity ;				Deviation	0.133 0.059				l Deviation		
Community Results		Maximum		Deviation	0.039		Maximum		. Deviation	1.479	
Com Re		Minimum			0.097		Minimum			0.339	
0		N			2		N			11	
		11			4		11			11	

Table 34. Data summary table for petunidin in dietary supplements.

Lann	I		Petunidin-3-Arabinoside										
		SRM	1 3283 C i	ranberry l	Extract (mg		1		Bilberry Ex	tract (mg	/g)		
	Lab	Α	В	C	Avg	SD	Α	В	C	Avg	SD		
	NIST									8			
	G701												
	G702												
	G703						8.3	8.3	8.4	8.3	0.1		
	G704												
	G705						11.1	11.0	11.1	11.1	0.1		
	G707												
	G708						10.9	10.8	10.8	10.8	0.1		
	G715						10.3	10.3		10.3	0.0		
	G716												
	G717						9.9	9.7	9.7	9.8	0.1		
	G718						10.2	9.9	10.2	10.1	0.2		
	G722												
	G724												
	G728						11.7	12.0	12.2	11.9	0.3		
Individual Results	G729												
Re	G738												
lual	G739												
livid	G740												
Inc	G749												
	G752						_						
	G753												
	G757												
	G760												
	G762		_				10.9	10.9	10.0	10.6	0.5		
	G765						14.3	13.7	13.9	14.0	0.3		
	G766						11.1	10.6	10.6	10.7	0.3		
	G767												
	G771												
	G772						0.1	0.2	7.0	07	0.7		
	G773						9.1	9.2	7.9	8.7	0.7		
	G774						0.2	0.4	0.2	0.2	0.0		
	G775 G776						9.3	9.4	9.3	9.3	0.0		
	G777												
	0///	Consensus	s Mean				Consensu	s Mean		10.3			
nity s				l Deviation				Deviation	1.3				
ommuni Results		Maximum					Consensus Standard Deviation Maximum						
Community Results		Minimum					Minimum			14.0 8.3			
		N			0		N			12			
L		- •			3		1 * '						

 Table 35. Data summary table for petunidin-3-arabinoside in dietary supplements.

		Petunidin-3-Galactoside										
		SRM	/I 3283 Ci	anberry F	xtract (m	g/g)	SR	M 3291 E	Bilberry Ex	tract (mg/	/g)	
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD	
	NIST											
	G701											
	G702											
	G703						13.1	13.3	13.3	13.2	0.1	
	G704											
	G705											
	G707											
	G708						16.7	16.5	16.7	16.6	0.1	
	G715	0.1700	0.1640		0.1670	0.0042	20.5	21.9		21.2	1.1	
	G716											
	G717						21.2	21.3	22.7	21.7	0.9	
	G718						16.2	15.6	16.2	16.0	0.3	
	G722											
	G724											
	G728						18.6	12.8	20.6	17.3	4.1	
Individual Results	G729											
Re	G738											
lual	G739											
livid	G740											
Ind	G749											
	G752											
	G753											
	G757											
	G760											
	G762	0.0297	0.0330	0.0319	0.0315	0.0017	16.8	16.8	15.5	16.4	0.8	
	G765						22.1	21.1	21.5	21.6	0.5	
	G766						13.2	12.4	12.5	12.7	0.5	
	G767											
	G771											
	G772											
	G773						13.1	13.5	12.0	12.9	0.8	
	G774											
	G775						15.2	15.2	15.2	15.2	0.0	
	G776											
	G777											
ty		Consensu			0.0993		Consensus			16.8		
Community Results				Deviation			Consensus		3.8			
ommun Results		Maximum			0.1670		Maximum			21.7		
C ₀		Minimum			0.0315		Minimum			12.7		
		Ν			2		Ν			11		

 Table 36. Data summary table for petunidin-3-galactoside in dietary supplements.

Petunidin-3-Glucoside											
		SRM	A 3283 Ci	ranberry I	Extract (m				Bilberry E	xtract (mg	/g)
	Lab	Α	В	C	Avg	SD	Α	В	C	Avg	SD
	NIST				0					0	
	G701						57.8	49.6	53.0	53.5	4.1
	G702										
	G703						27.3	27.7	27.6	27.5	0.2
	G704										
	G705						33.5	34.5	34.6	34.2	0.6
	G707										
	G708						34.0	34.0	34.2	34.1	0.1
	G715						29.4	30.6		30.0	0.8
	G716										
	G717						32.6	33.7	33.6	33.3	0.6
	G718	0.0440		0.0440	0.0440	0.0000	33.2	32.0	33.2	32.8	0.7
	G722										
	G724										
s	G728						39.9	37.5	39.6	39.0	1.3
sult	G729										
l Re	G738										
Individual Results	G739										
	G740		_			_		_			
	G749										
	G752										
	G753										
	G757										
	G760 G762	0.0056	0.0057	0.0060	0.0058	0.0002	34.0	33.9	31.0	33.0	1.7
	G765	0.0050	0.0037	0.0000	0.0058	0.0002	44.3	42.3	43.0	43.2	1.7
	G766						26.2	25.2	25.3	25.6	0.6
	G767						20.2	23.2	23.3	23.0	0.0
	G771										
	G772										
	G773		0.0180		0.0180		23.8	23.0	21.5	22.8	1.2
	G774										
	G775	0.1866	0.1918	0.1945	0.1910	0.0040	30.1	30.2	30.2	30.2	0.1
	G776										
	G777	0.0070	0.0060	0.0060	0.0063	0.0006	31.9	32.0	32.0	31.9	0.0
y		Consensu	s Mean		0.0618		Consensu	s Mean		32.7	
unit; ts		Consensu	s Standard	Deviation	0.0998		Consensus	s Standard	l Deviation	6.0	
ommun Results		Maximum	ı		0.1910		Maximum			53.5	
Community Results		Minimum			0.0058					22.8	
		Ν			4		Ν			14	

 Table 37. Data summary table for petunidin-3-glucoside in dietary supplements.

			<u>iiiidi y</u> ta			-	Equivalent		supple	ine inc.	
		SRN	A 3283 Ci	ranberry I	Extract (m	g/g)	SR	M 3291 H	Bilberry E	xtract (mg	/g)
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST										
	G701						38.3	32.8	35.1	35.4	2.7
	G702										
	G703						33.8	34.2	34.2	34.1	0.2
	G704										
	G705						30.8	31.3	31.5	31.2	0.4
	G707										
	G708						41.2	41.0	41.3	41.2	0.1
	G715	0.1985	0.2155		0.2070	0.0120	41.0	42.8		41.9	1.2
	G716										
	G717						43.1	43.8	44.7	43.9	0.8
	G718	0.0291		0.0291	0.0291	0.0000	40.3	38.9	40.2	39.8	0.8
	G722										
	G724										
s	G728						47.7	42.3	49.0	46.3	3.6
sult	G729										
l Re	G738						12.1			12.1	
Individual Results	G739										
	G740										
	G749										
	G752										
	G753										
	G757										
	G760 G762	0.0233	0.0256	0.0251	0.0247	0.0012	41.3	41.2	37.8	40.1	2.0
	G765	0.0233	0.0230	0.0231	0.0247	0.0012	54.5	52.2	53.0	53.2	1.2
	G766						34.0	32.4	32.4	32.9	0.9
	G767						54.0	52.4	52.4	52.7	0.9
	G771										
	G772						41.9	44.4	45.3	43.9	1.8
	G773	0.1500	0.1919	0.1800	0.1740	0.0216	31.4	31.3	28.4	30.4	1.7
	G774										
	G775	0.1235	0.1269	0.1287	0.1264	0.0027	38.0	38.2	38.1	38.1	0.1
	G776										
	G777	0.0046	0.0040	0.0040	0.0042	0.0004	22.1	22.1	22.1	22.1	0.0
y		Consensu	s Mean		0.0942		Consensu	s Mean		38.4	
unit. Its		Consensu	s Standard	Deviation	0.0979		Consensus Standard Deviat			7.1	
ommun Results		Maximum	1		0.2070		Maximum			53.2	
Community Results		Minimum			0.0042		Minimum			12.1	
		Ν			6		Ν			15	

 Table 38. Data summary table for petunidin equivalents in dietary supplements.

Total Anthocyanins											
		SRN	A 3283 Ci	anberry l	Extract (m	g/g)	SR	M 3291 H	Bilberry E	xtract (mg	/g)
	Lab	Α	В	С	Avg	SD	Α	В	С	Avg	SD
	NIST										
	G701	0.734	0.729	0.768	0.744	0.021	380	336	334	350	26
	G702										
	G703	0.676	0.651	0.644	0.657	0.017	310	314	314	313	2
	G704										
	G705	0.696	0.768	0.675	0.713	0.049	390	387	387	388	2
	G707										
	G708	0.891	0.895	0.958	0.915	0.038	376	374	376	375	1
	G715	0.910	0.982		0.946	0.051	366	368		367	2
	G716	1.700	1.800	1.900	1.800	0.100	290	280	270	280	10
	G717	0.798	0.796	0.808	0.801	0.006	371	367	367	368	2
	G718	1.337	1.321	1.280	1.313	0.029	424	409	424	419	9
	G722										
	G724	0.730	0.770	0.770	0.757	0.023	282	281	282	282	1
s	G728	1.270	0.841	1.187	1.099	0.228	457	443	457	453	8
Individual Results	G729										
l Re	G738	0.476			0.476		97			97	
lual	G739										
divid	G740										
Inc	G749										
	G752										
	G753										
	G757	10.100	11.100	10.200	10.467	0.551	433	444	419	432	13
	G760										
	G762	0.685	0.710	0.699	0.698	0.013	388	387	357	377	18
	G765	0.847	0.867	0.917	0.877	0.036	543	519	528	530	12
	G766	0.745	0.745	0.720	0.737	0.014	303	296	297	298	4
	G767										
	G771			_							_
	G772	2.200	2.196	2.102	2.166	0.055	243	257	260	253	9
	G773	0.507	0.676	0.720	0.634	0.112	302	303	272	292	17
	G774										
	G775	0.696	0.717	0.702	0.705	0.010	340	341	341	340	1
	G776							1.0.0			
	G777	1.267	1.233	1.292	1.264	0.030	428	428	424	427	2
ity		Consensu		D · · ·	0.934		Consensus			354	
Community Results			s Standard	Deviation			Consensus				
ommun Results		Maximum			10.467 Maximum 530						
Co F		Minimum			0.476 Minimum 97						
		Ν			18		Ν			18	

 Table 39. Data summary table for total anthocyanins in dietary supplements.

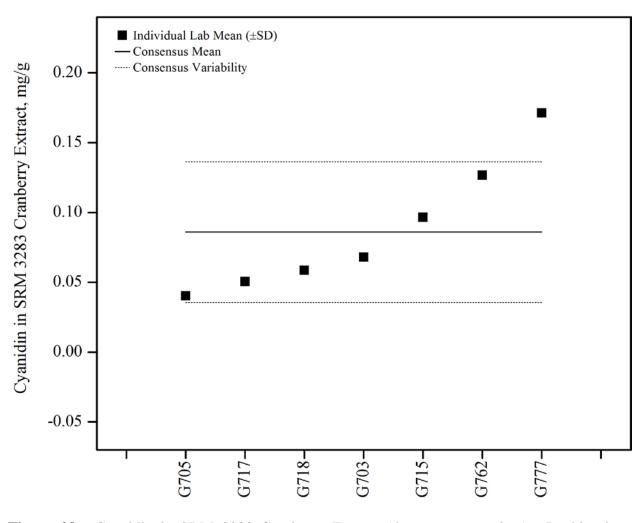


Figure 28. Cyanidin in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

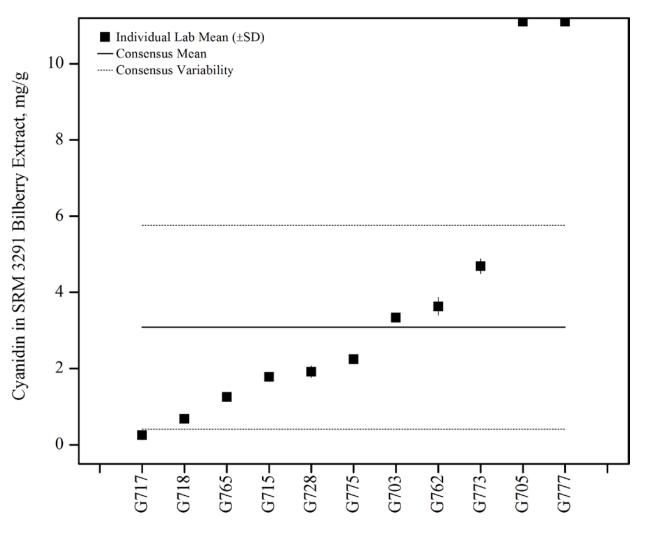


Figure 29. Cyanidin in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

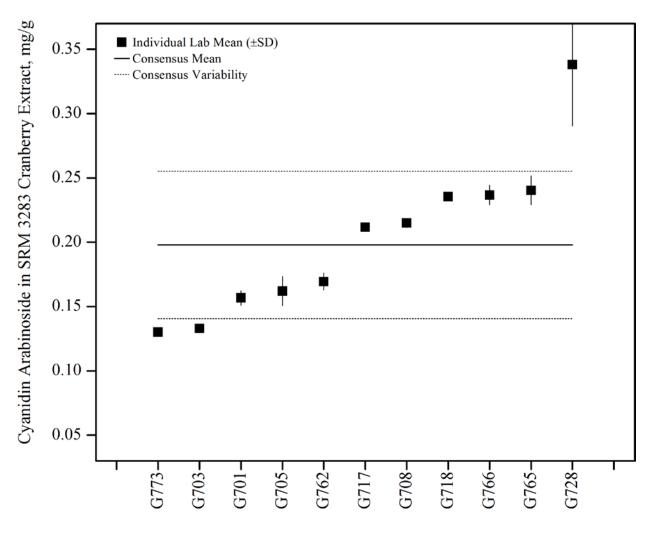


Figure 30. Cyanidin-3-arabinoside in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

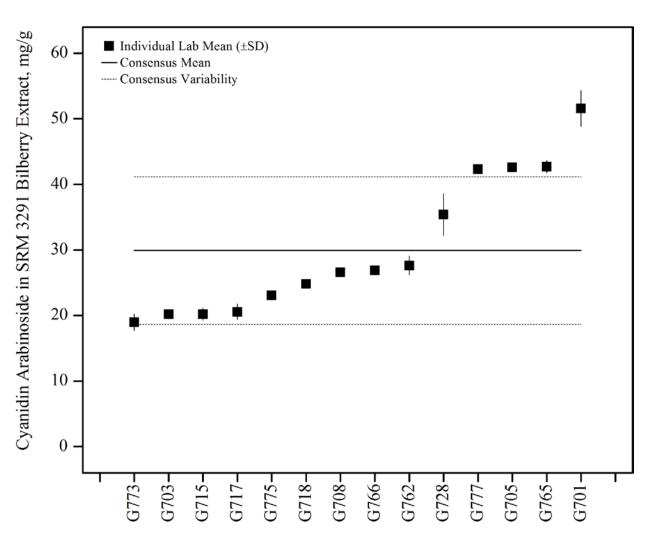


Figure 31. Cyanidin-3-arabinoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

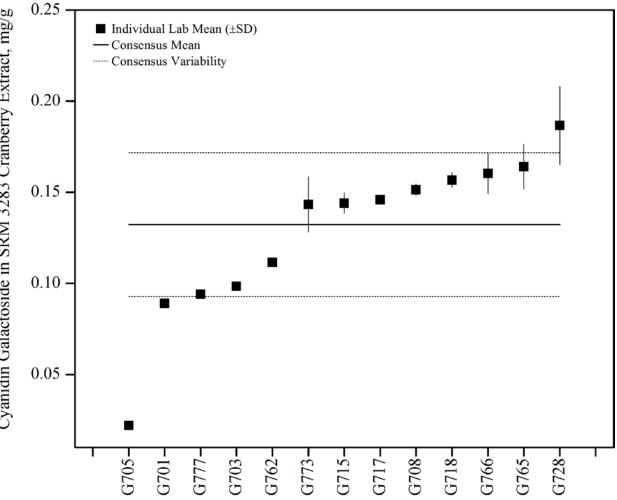


Figure 32. Cyanidin-3-galactoside in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

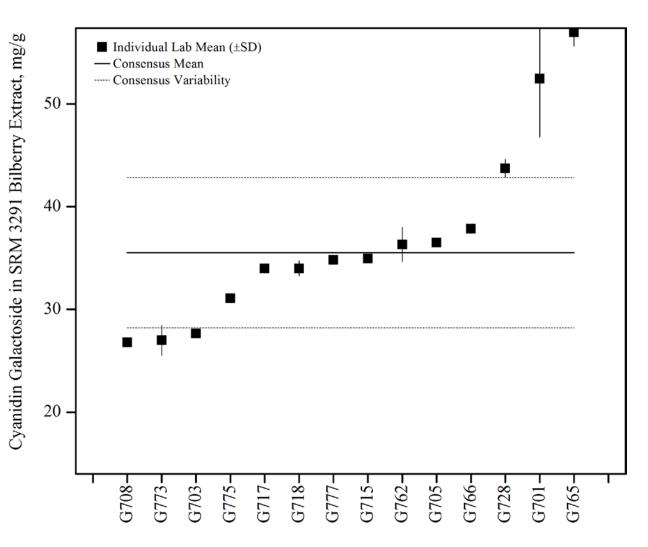


Figure 33. Cyanidin-3-galactoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

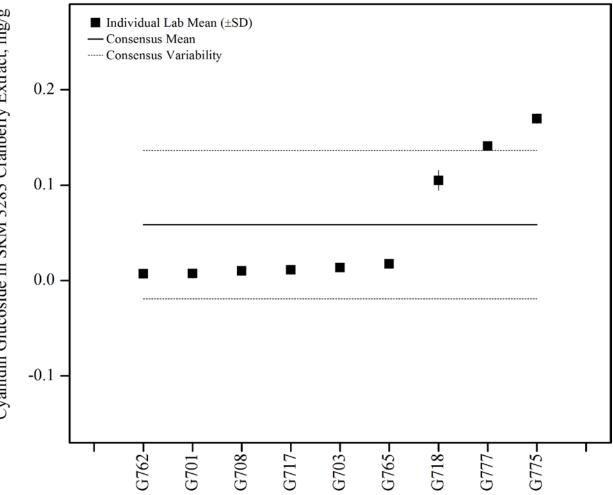


Figure 34. Cyanidin-3-glucoside in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

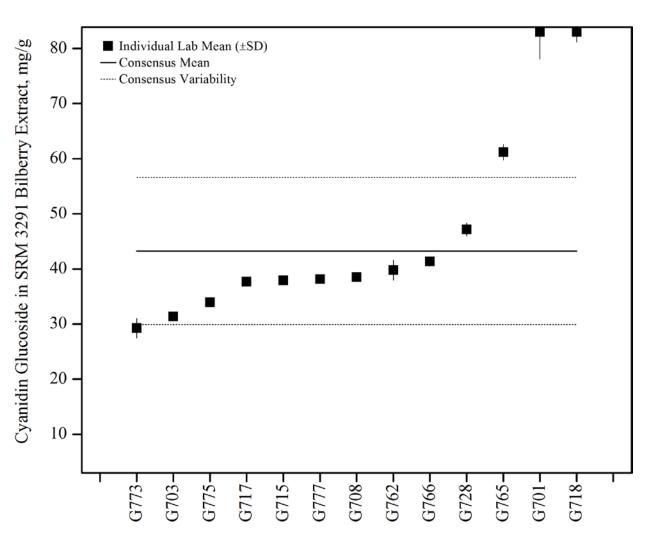


Figure 35. Cyanidin-3-glucoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

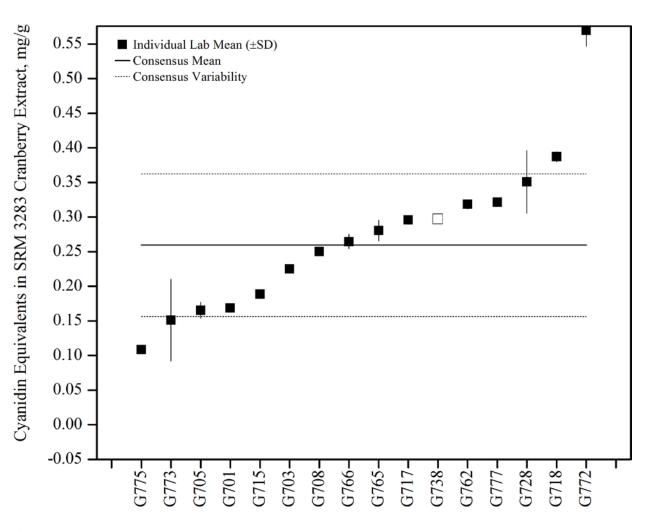
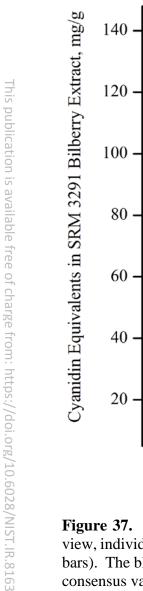


Figure 36. Cyanidin equivalents in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.



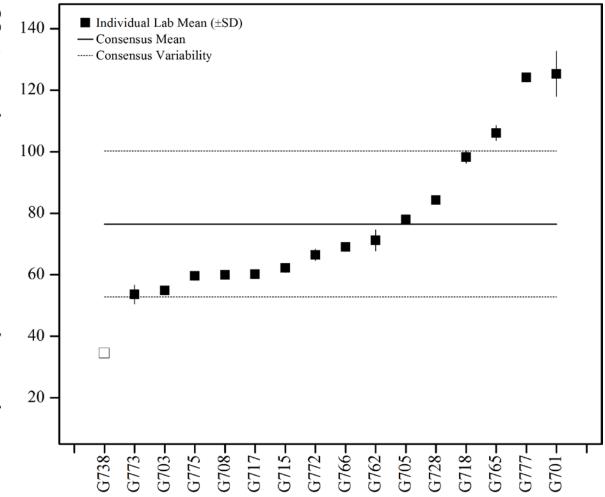


Figure 37. Cyanidin equivalents in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

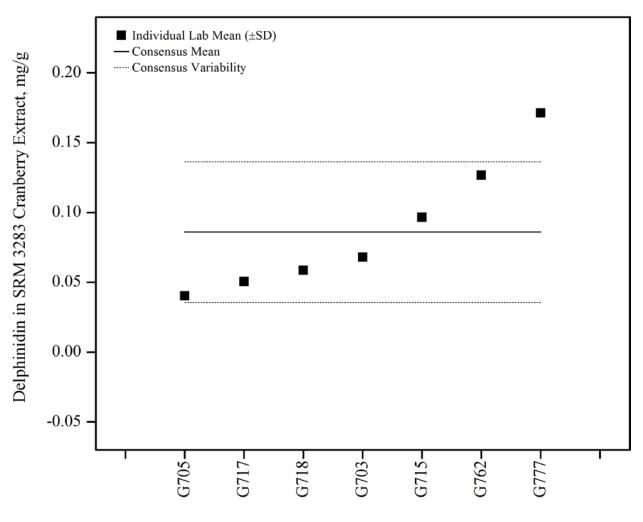
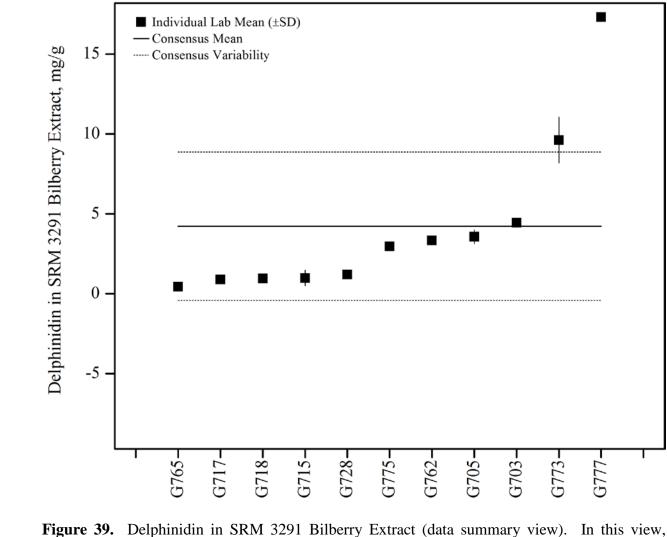


Figure 38. Delphinidin in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.



individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

G773-

G777.

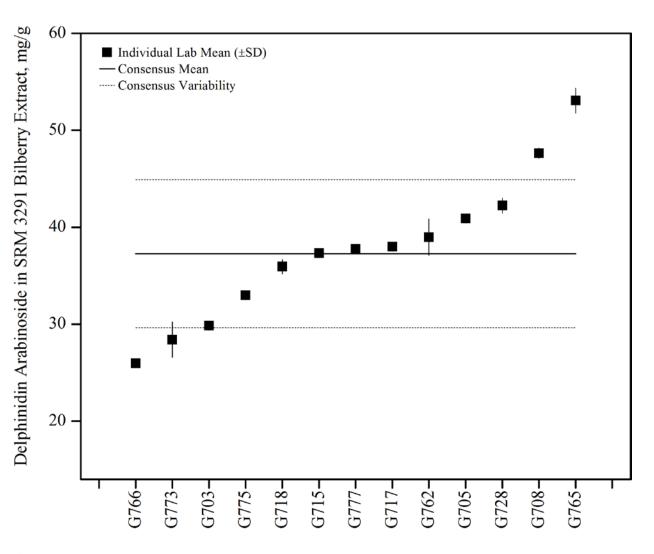


Figure 40. Delphinidin-3-arabinoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

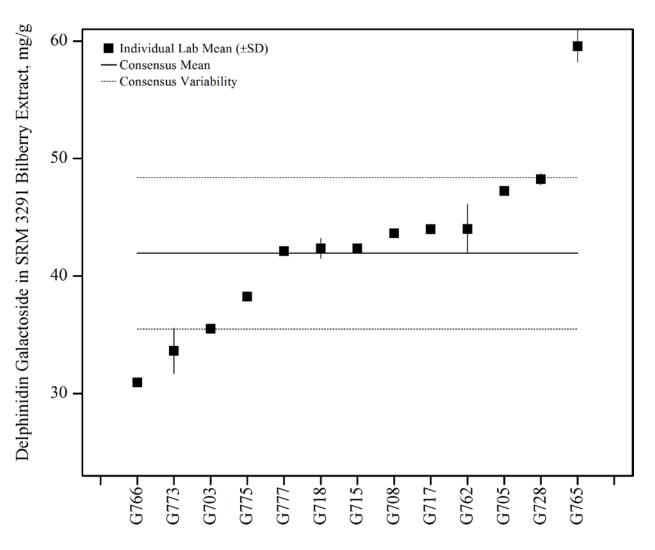


Figure 41. Delphinidin-3-galactoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

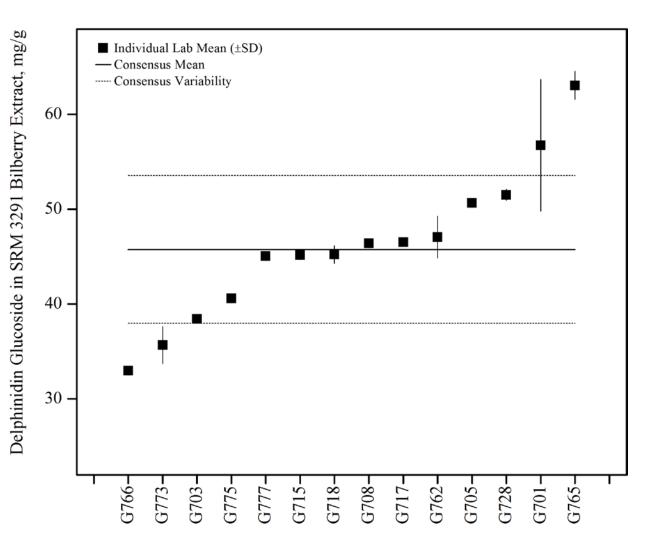


Figure 42. Delphinidin-3-glucoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

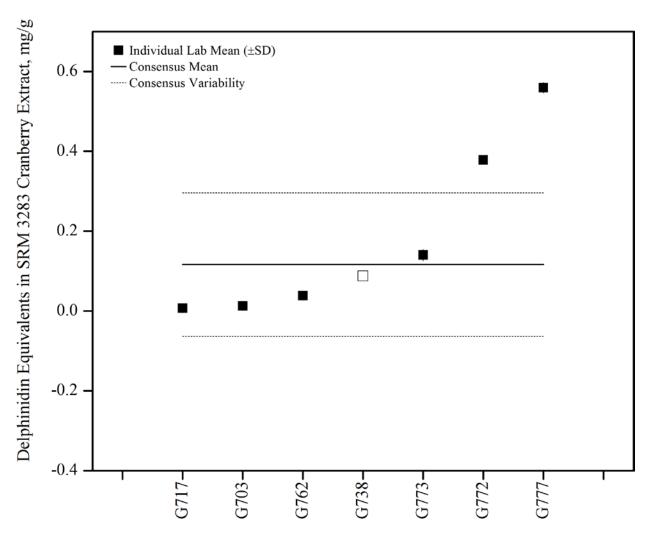


Figure 43. Delphinidin equivalents in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

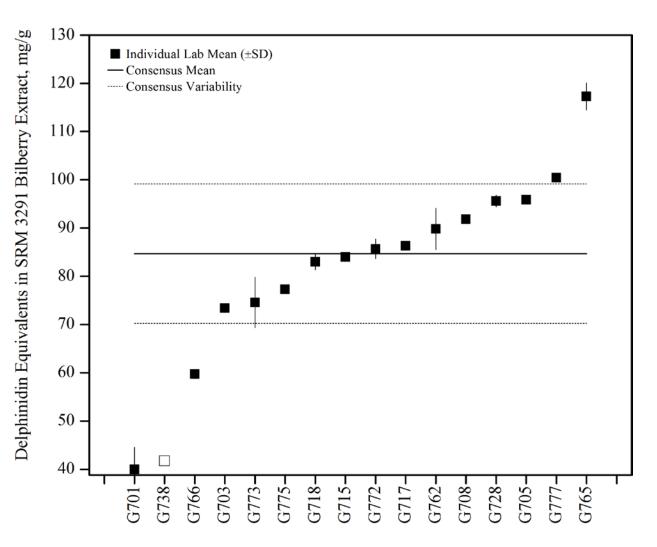


Figure 44. Delphinidin equivalents in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

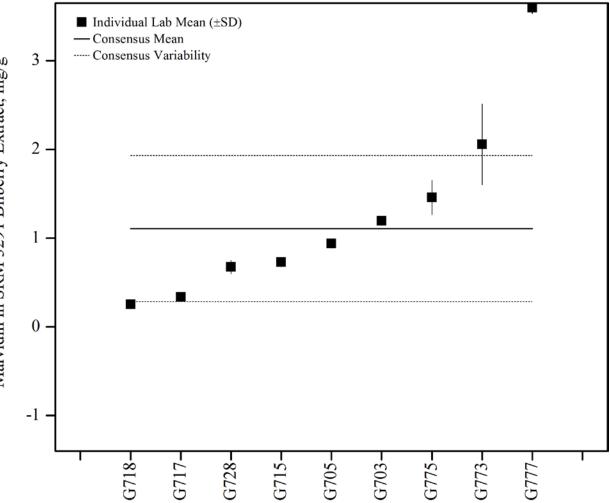


Figure 45. Malvidin in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

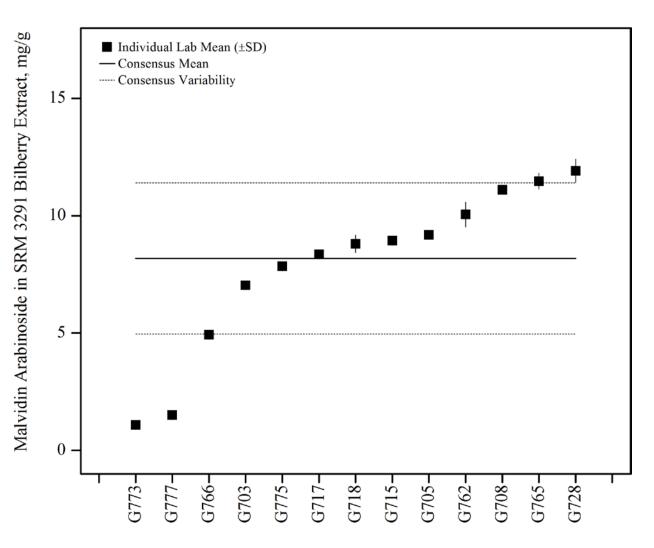


Figure 46. Malvidin-3-arabinoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

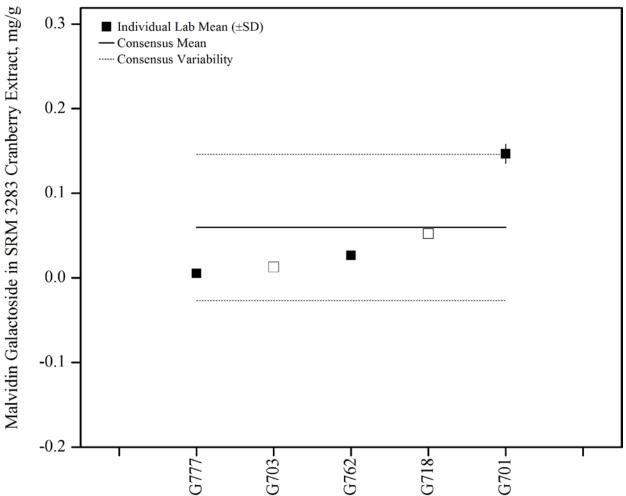


Figure 47. Malvidin-3-galactoside in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

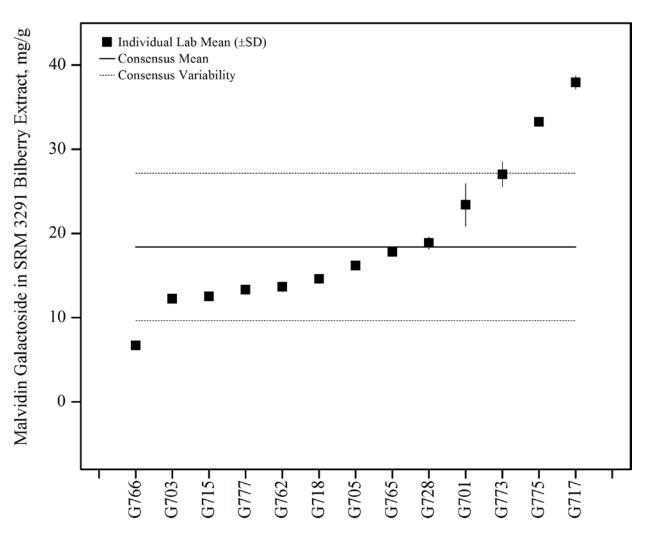


Figure 48. Malvidin-3-galactoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

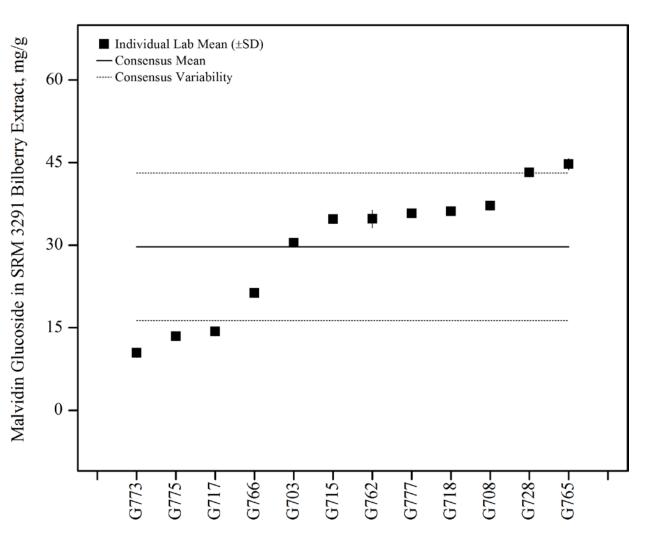


Figure 49. Malvidin-3-glucoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

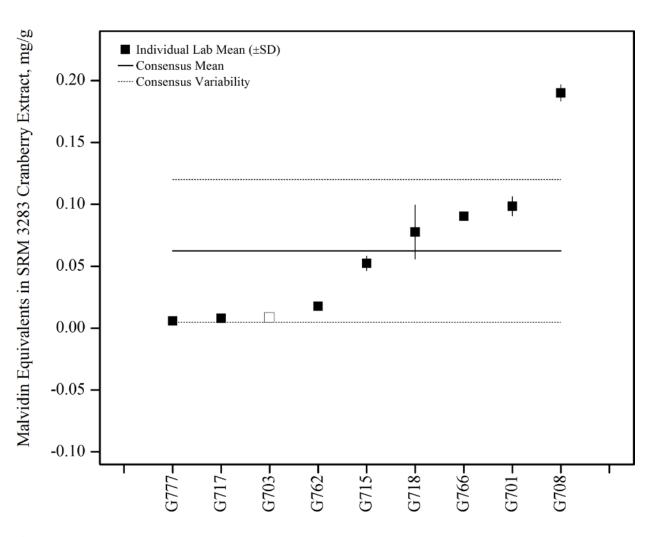


Figure 50. Malvidin equivalents in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

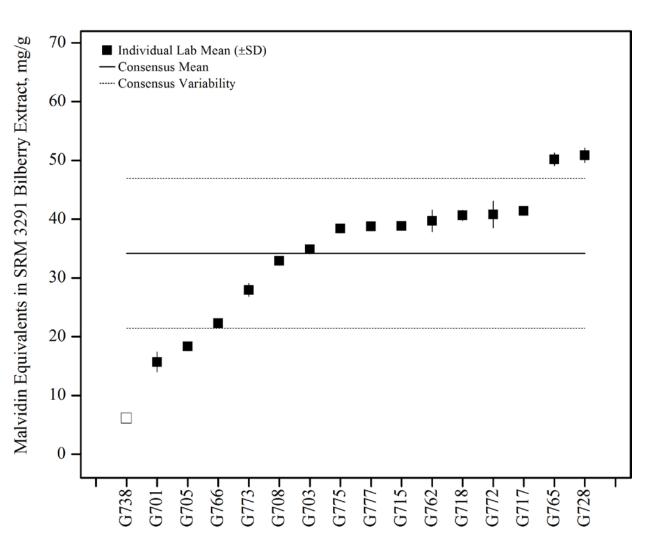


Figure 51. Malvidin equivalents in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

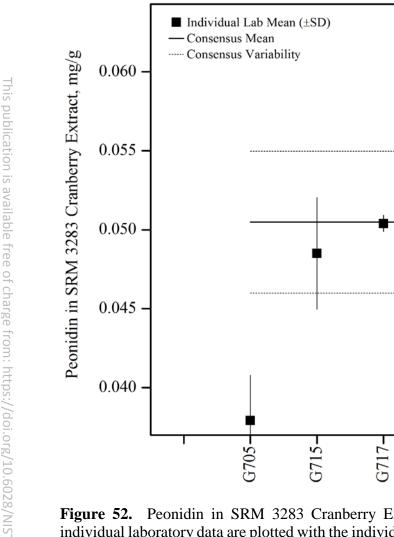


Figure 52. Peonidin in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

G718-

G703-

G777-

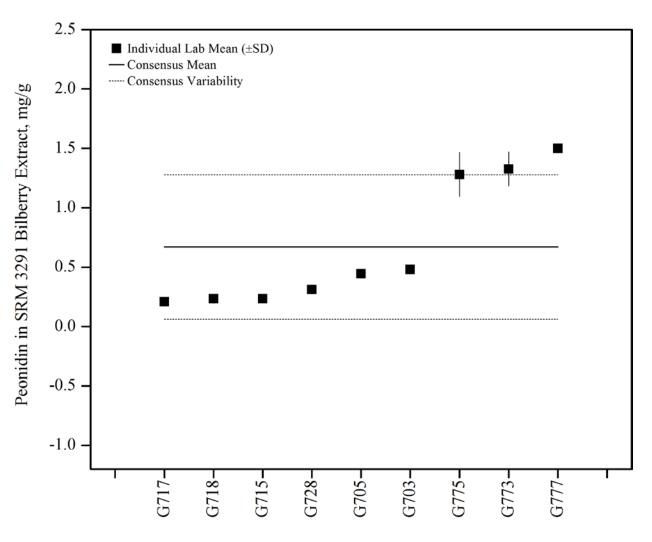


Figure 53. Peonidin in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

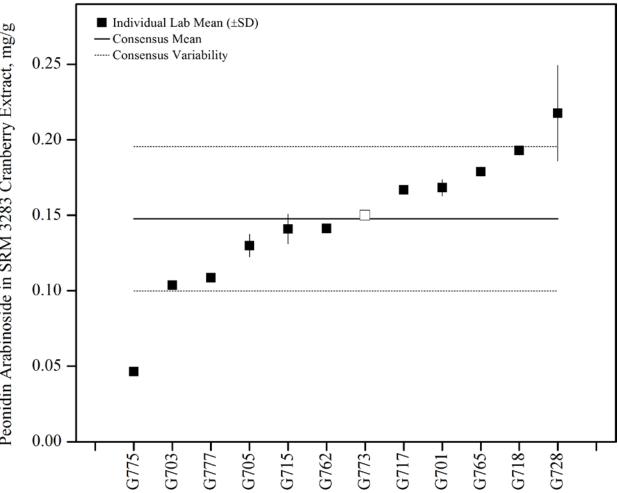


Figure 54. Peonidin-3-arabinoside in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

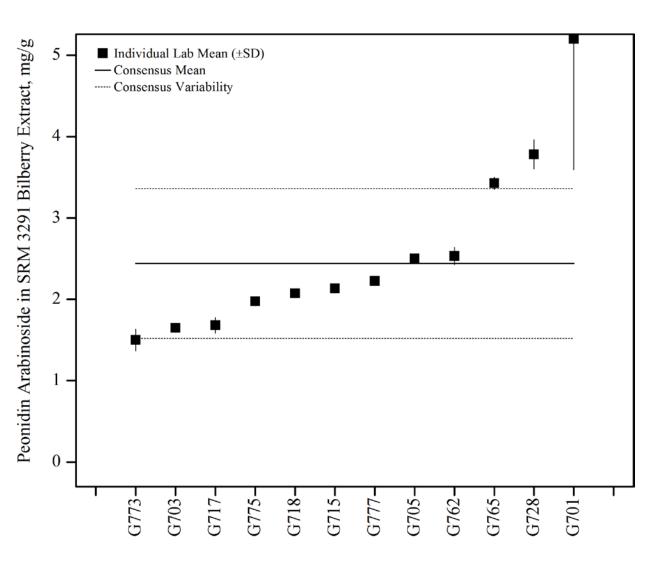
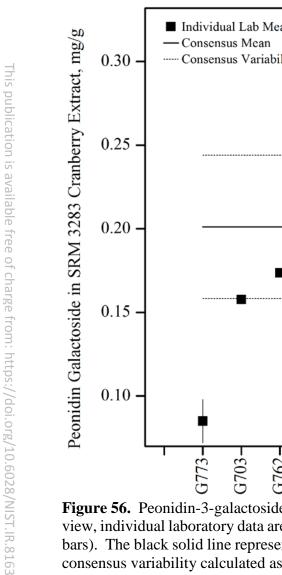
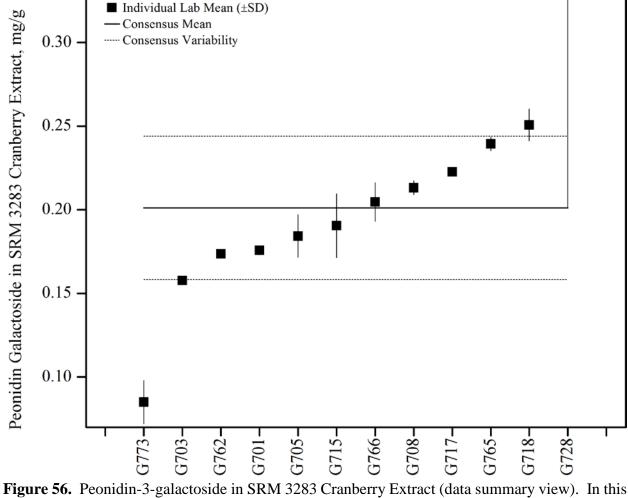


Figure 55. Peonidin-3-arabinoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.





view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

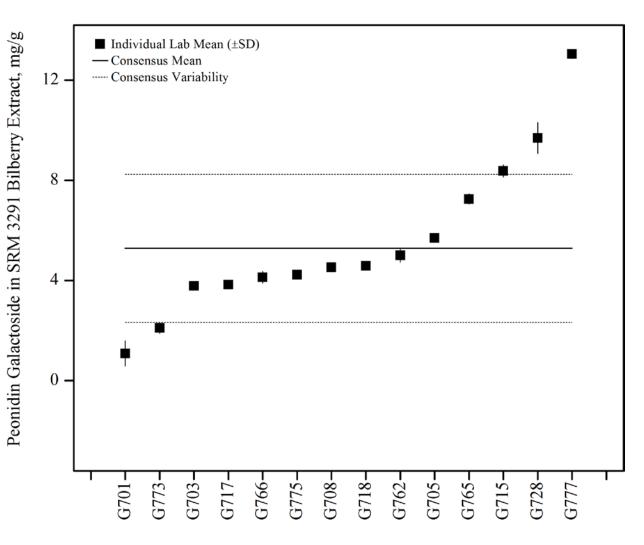


Figure 57. Peonidin-3-galactoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

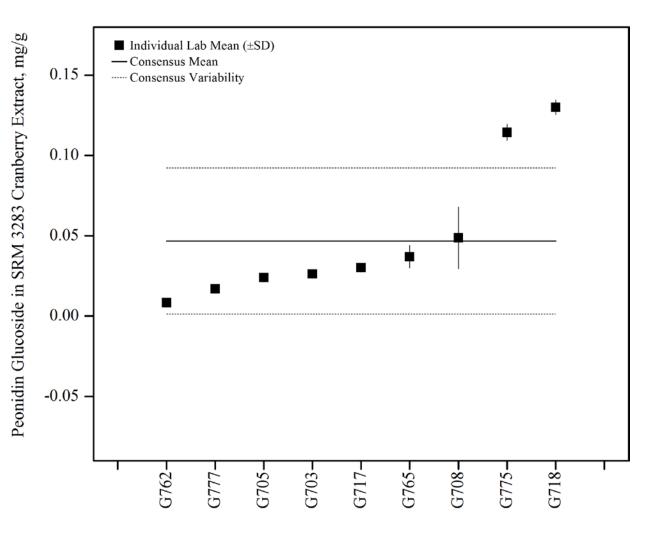


Figure 58. Peonidin-3-glucoside in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

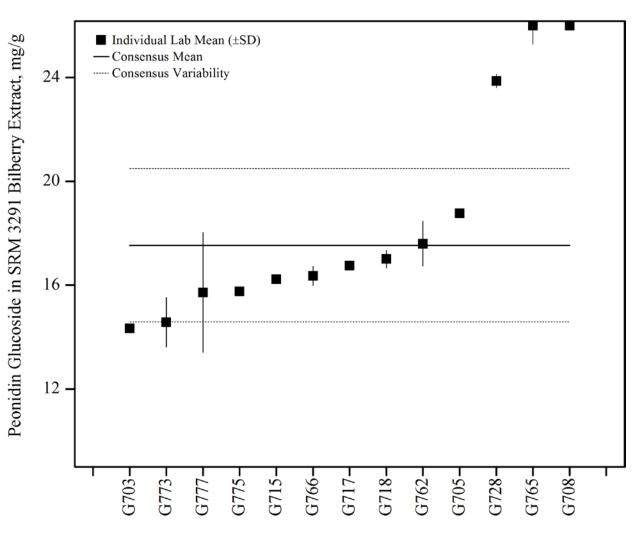


Figure 59. Peonidin-3-glucoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

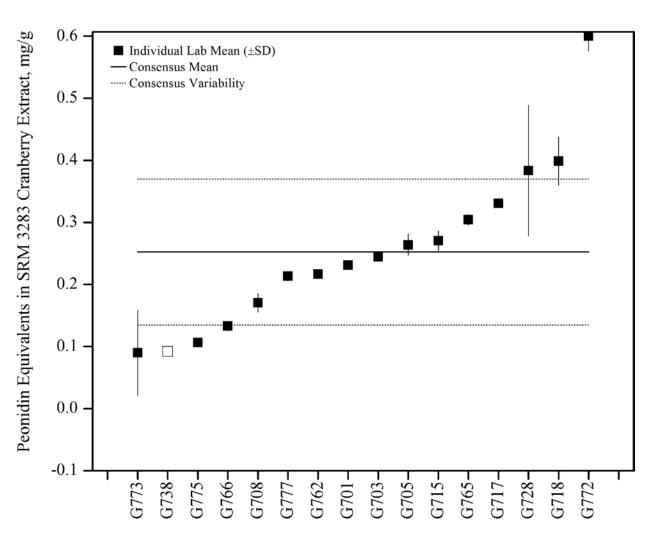


Figure 60. Peonidin equivalents in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

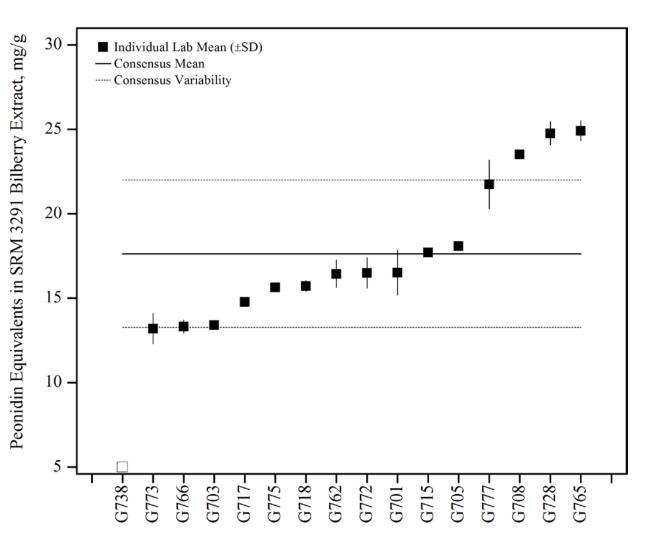


Figure 61. Peonidin equivalents in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

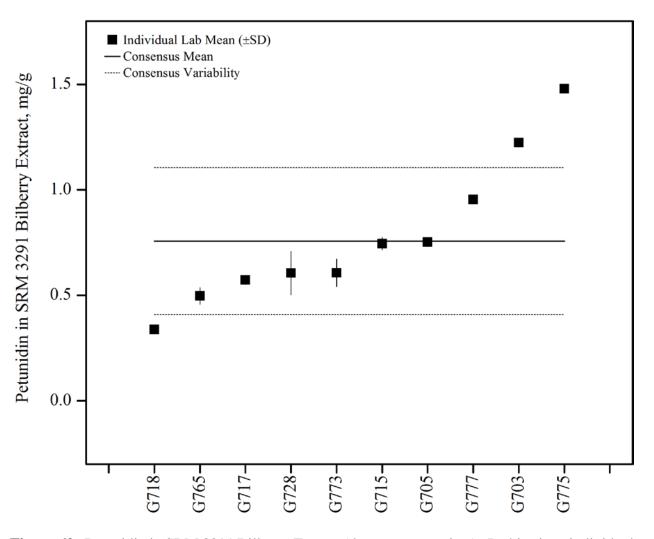


Figure 62. Petunidin in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

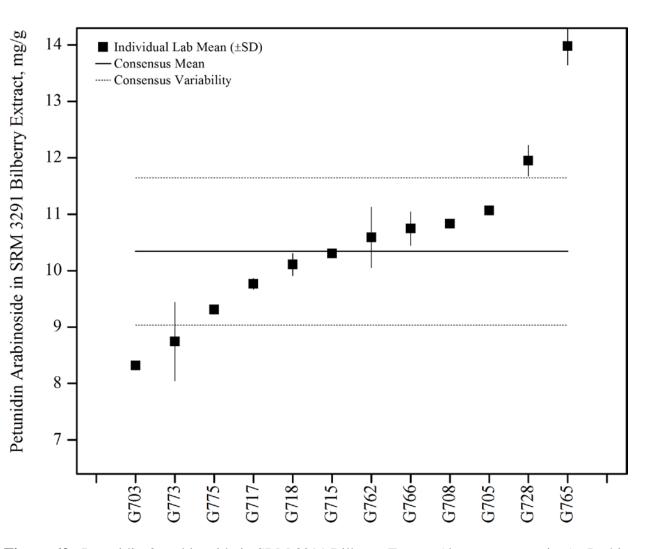


Figure 63. Petunidin-3-arabinoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

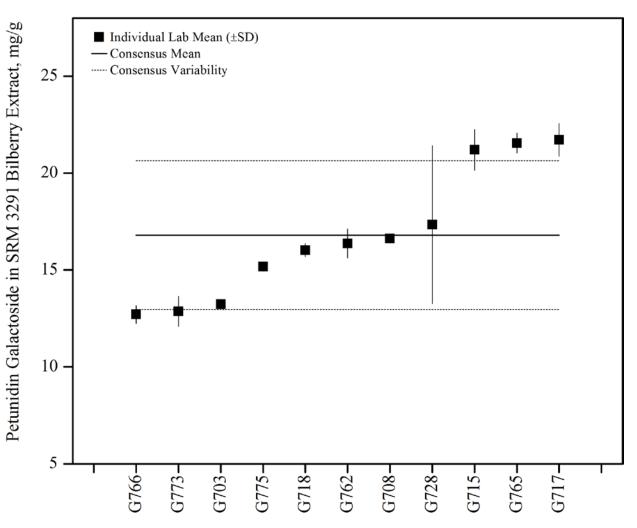


Figure 64. Petunidin-3-galactoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

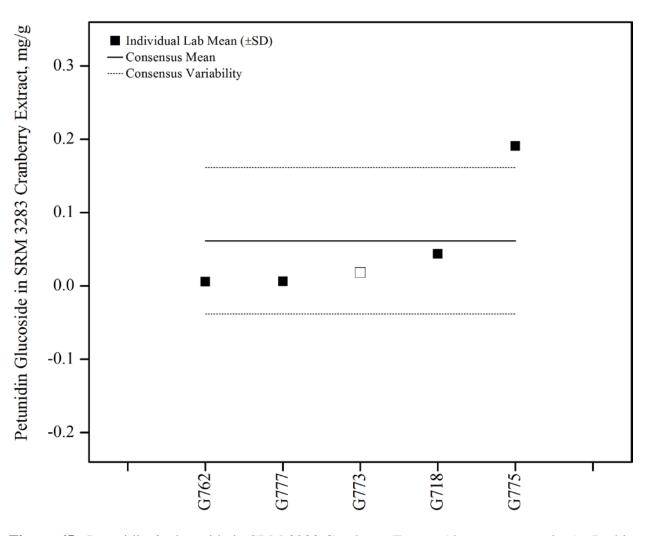


Figure 65. Petunidin-3-glucoside in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

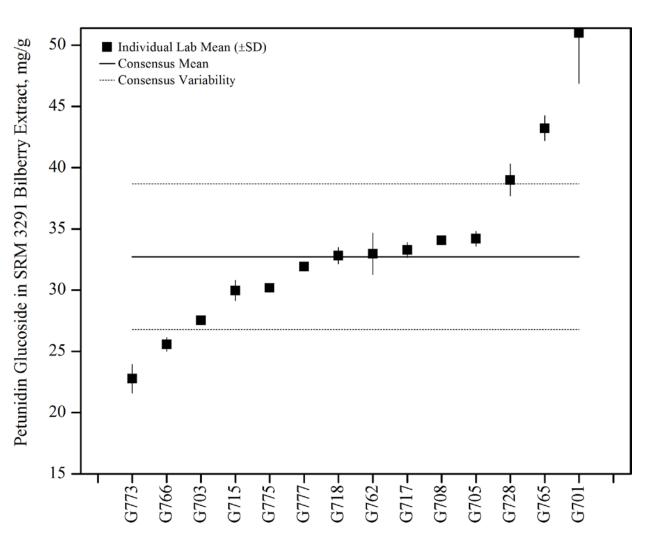


Figure 66. Petunidin-3-glucoside in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

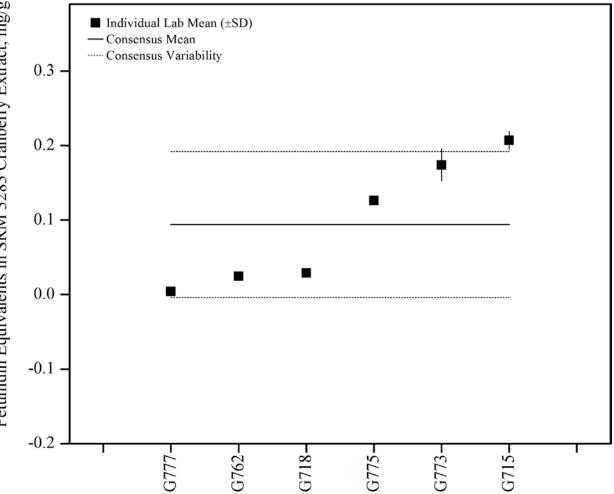


Figure 67. Petunidin equivalents in SRM 3283 Cranberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

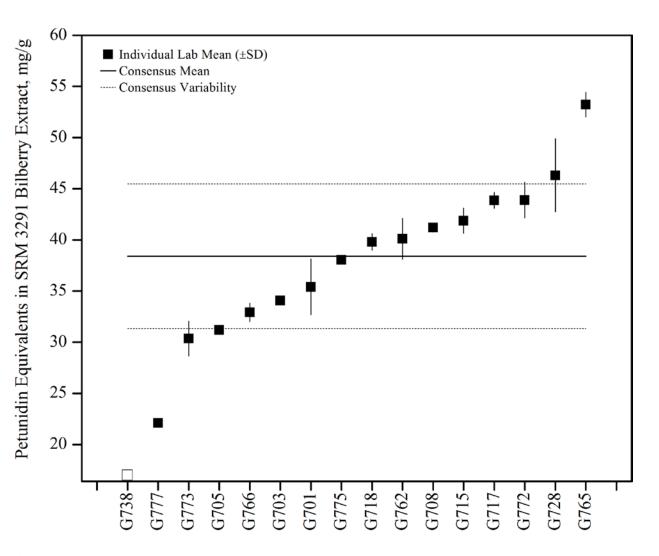


Figure 68. Petunidin equivalents in SRM 3291 Bilberry Extract (data summary view). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

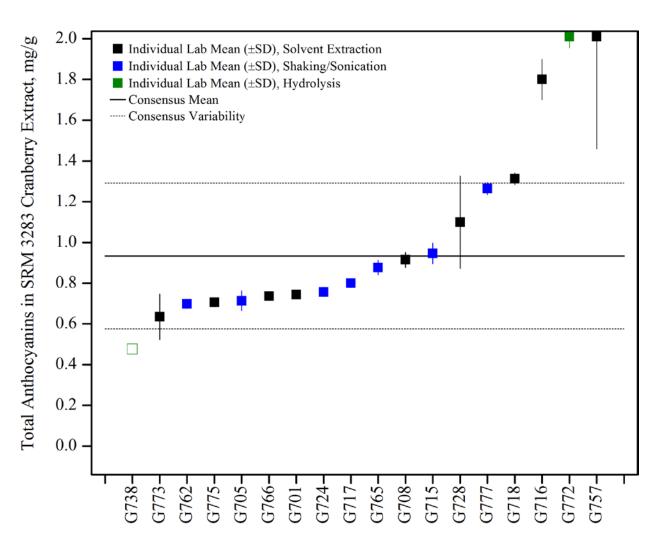


Figure 69. Total anthocyanins in SRM 3283 Cranberry Extract (data summary view – sample preparation method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by sample preparation method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

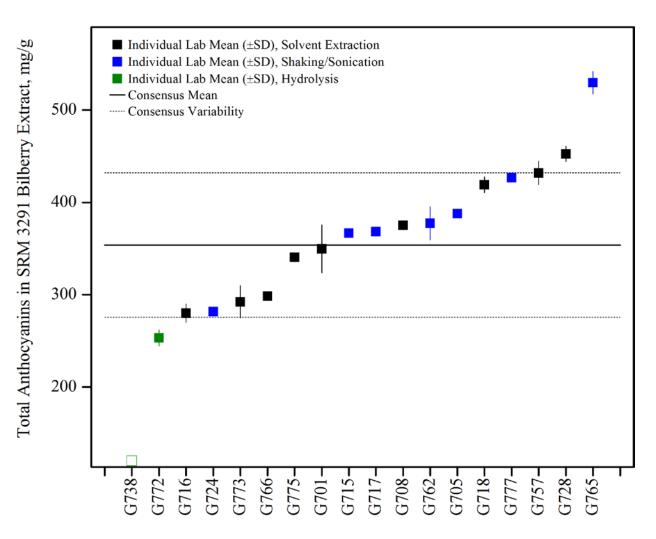


Figure 70. Total anthocyanins in SRM 3291 Bilberry Extract (data summary view – sample preparation method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by sample preparation method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

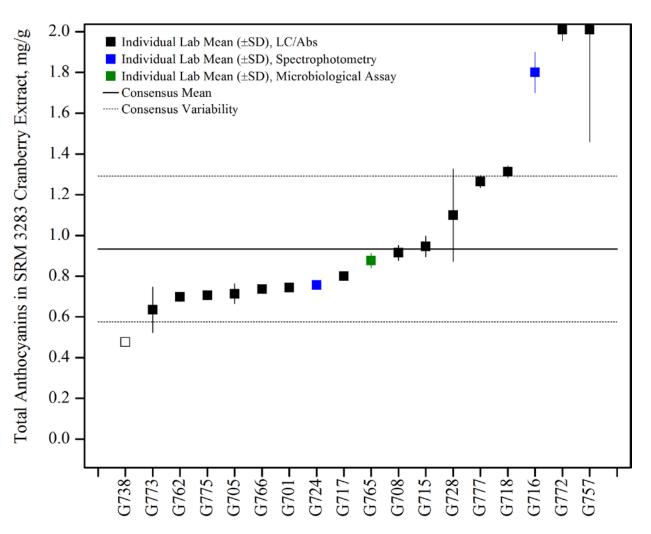


Figure 71. Total anthocyanins in SRM 3283 Cranberry Extract (data summary view – instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

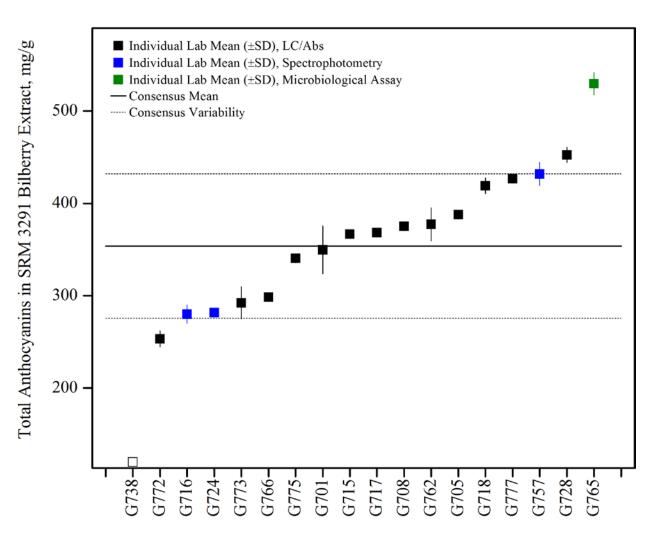


Figure 72. Total anthocyanins in SRM 3291 Bilberry Extract (data summary view – instrumental method). In this view, individual laboratory data are plotted with the individual laboratory standard deviation (error bars). The data are identified by instrumental method in this graph. The black solid line represents the consensus mean, and the black dotted lines represent the consensus variability calculated as one standard deviation about the consensus mean.

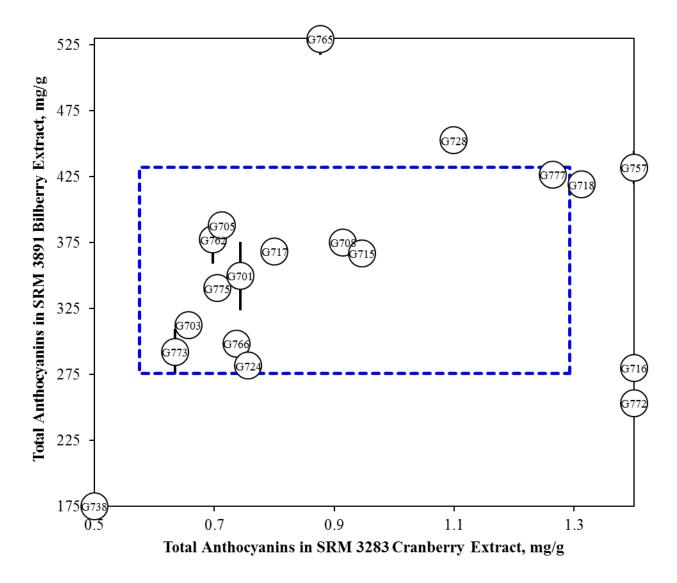


Figure 73. Total anthocyanins in SRM 3283 Cranberry Extract and SRM 3291 Bilberry Extract (sample/sample comparison view). In this view, the individual laboratory results for one sample (SRM 3283 Cranberry Extract) are compared to the results for a second sample (SRM 3291 Bilberry Extract). The dotted blue box represents the consensus zone for the cranberry extract (x-axis) and the bilberry extract (y-axis).