

SCST Proficiency Testing Task Force Report

Background

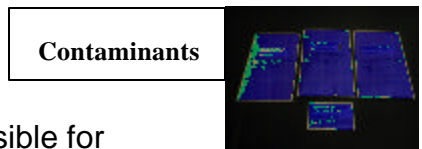
During the 2002 annual business meeting SCST members voted to send every Registered Seed Technologist (RST) three proficiency testing samples as a pilot proficiency-testing program. The results of these tests would be used to “take the pulse” of the Society and to help establish a permanent proficiency testing program. Members volunteered to assist with the process and a task force was formed. The task force met in September and chose the species and test to be performed. A purity and germination on *Triticum aestivum* (wheat), *Cucumis melo* (cantaloupe) germination and *Glycine max* (soybean) germination were selected. Volunteers were chosen to provide seed for the samples, prepare the samples, develop forms for recording results and to prepare education materials. A time line was developed for completion of these tasks.

Selection of Seed Lots and Preparation of samples



**Dividing down
with a riffle divider**

Seed lots exhibiting mid-range germination and low levels of inert matter were selected and sent to the sample preparation volunteers. The sample prep volunteers mixed the samples according to the ISTA Referee Sample Preparation Protocol. Samples were randomly selected for heterogeneity testing and sent to experienced RSTs for analysis (see Appendix A). The results were statistically analyzed to guarantee that the samples were uniform. In addition, the contaminants for the wheat samples were sent to a third party for species verification before they were added to the proficiency samples.



**Labeled and filled
envelopes**

The SCST Executive Director was responsible for providing labeled sample envelopes. Each RST was assigned a unique sample number. The same number was used for all three species. Envelopes were sent to the sample prep volunteers and the filled samples were returned to the Executive Director. Samples and result recording forms were sent to each RST. To ensure confidentiality only the Executive Director has access to the result forms and sample assignments.

Results were due February 1st for the wheat and cantaloupe and March 1st for the soybean. Three months were given to complete the tests and return the results.

Analysis

All the results received before March 15th were included in the analysis. The mean, median, and standard deviation were calculated for normal seedlings, abnormal seedlings, dead seed and the purity components. Graphs are used to illustrate results and where they fall in relation to one and two standard deviations.

The proficiency testing task force donated considerable time and resources to make this pilot proficiency testing a positive and educational experience for SCST members (see Appendix B). The task force will hold an open meeting at the annual meeting in Seattle, WA Monday June 9th. Everyone is invited to attend.

Respectfully submitted,

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Analysis of Results

Cucumis melo (cantaloupe) Germination Analysis

Normal seedling evaluation (118 results returned)

- 86% fell within one standard deviation of the mean
- 96% fell within two standard deviations of the mean
- Outlier sample numbers: 39, 68, 144, 148

Abnormal seedlings evaluation (116 results returned)

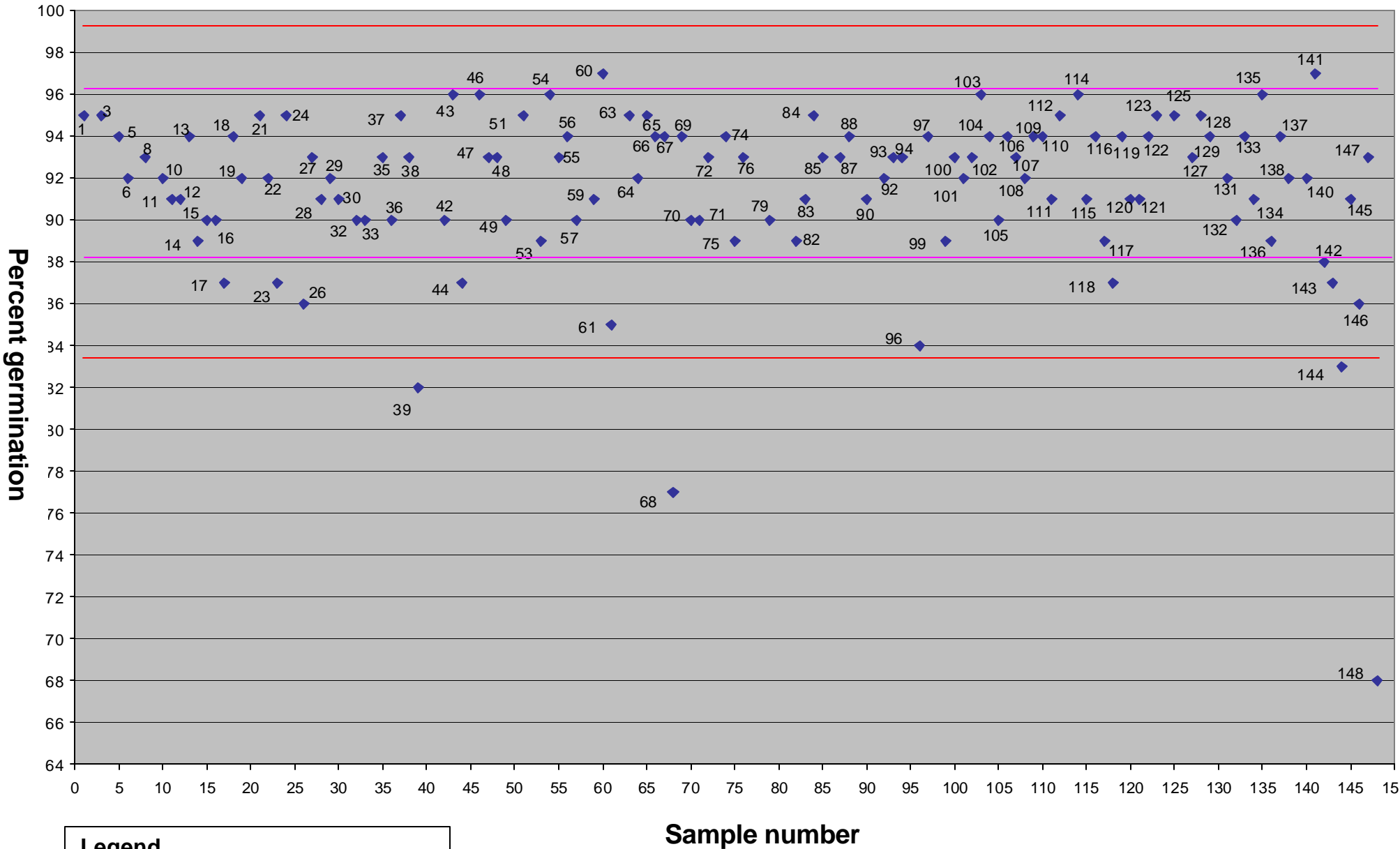
- 89% fell within one standard deviation of the mean
- 97% fell within two standard deviations of the mean
- Outlier sample numbers: 61, 68, 96, 144

Dead seed evaluation (116 results returned)

- 94% fell within one standard deviation of the mean
- 99% fell within two standard deviations of the mean
- Outlier sample number: 148

<i>Cucumis melo</i> germination	Normal seedlings	Abnormal seedlings	Dead seed
Mean	91.64	3.36	4.93
Median	92.5	3	5
Standard deviation	3.90	2.58	2.71
Mean + 1 standard deviation	95.54	5.94	7.64
Mean - 1 standard deviation	87.73	0.78	2.22
Mean + 2 standard deviation	99.44	8.52	10.34
Mean - 2 standard deviation	83.83	-1.80	-0.481

Cucumis melo Germination Graph



Glycine max (soybean) Germination Analysis

Normal seedling evaluation (111 results)

- 81% fell within one standard deviation of the mean
- 95% fell within two standard deviations of the mean
- Outlier sample numbers: 15, 32, 48, 51, 92, 136

Abnormal seedlings evaluation (111 returned)

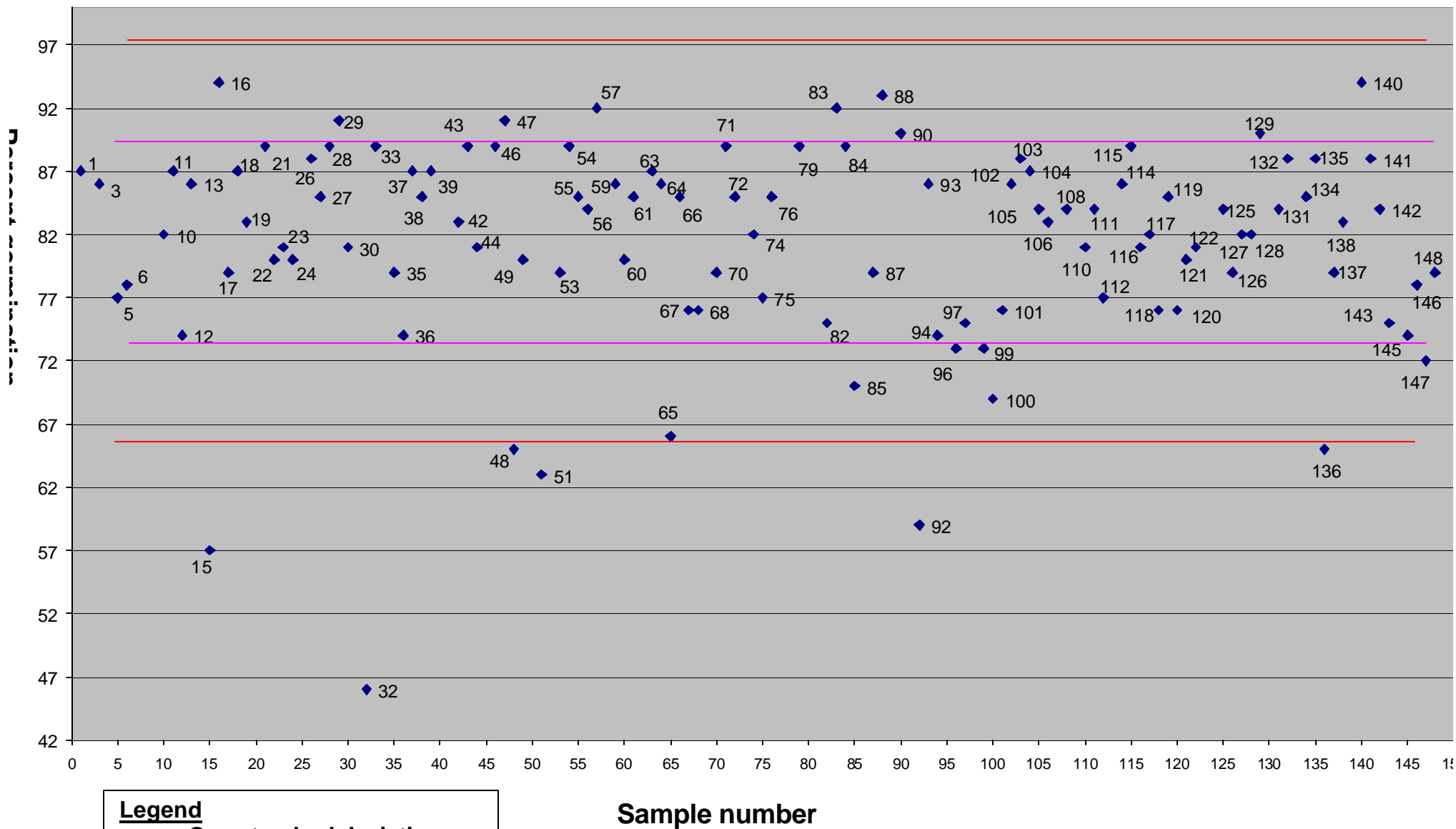
- 74% fell within one standard deviation of the mean
- 92% fell within two standard deviations of the mean
- Outlier sample numbers: 15, 16, 32, 48, 51, 57, 59, 92, 100

Dead seed evaluation (111 results returned)

- 73% fell within one standard deviation of the mean
- 95% fell within two standard deviations of the mean
- Outlier sample number: 32, 65, 136, 92, 94

Glycine max germination	Normal seedlings	Abnormal seedlings	Dead seed	Hard seed
Mean	81.50	13.49	5.05	0.05
Median	83	13	4	0
Standard deviation	7.95	5.83	4.11	0.33
Mean + 1 standard deviation	89.45	19.32	9.16	0.38
Mean - 1 standard deviation	73.56	7.65	0.93	-0.27
Mean + 2 standard deviation	97.40	25.15	13.27	0.71
Mean - 2 standard deviation	65.61	1.82	-3.18	-0.60

Glycine max Germination Graph



Legend

- One standard deviation
- Two standard deviations
- data marker w/sample number

Triticum aestivum (wheat) Germination Analysis

Normal seedling evaluation (116 results returned)

- 65% fell within one standard deviation of the mean
- 97% fell within two standard deviations of the mean
- Outlier sample numbers: 18, 46, 47, 72

Abnormal seedlings evaluation (116 results returned)

- 69% fell within one standard deviation of the mean
- 97% fell within two standard deviations of the mean
- Outlier sample numbers: 18, 44, 47, 120

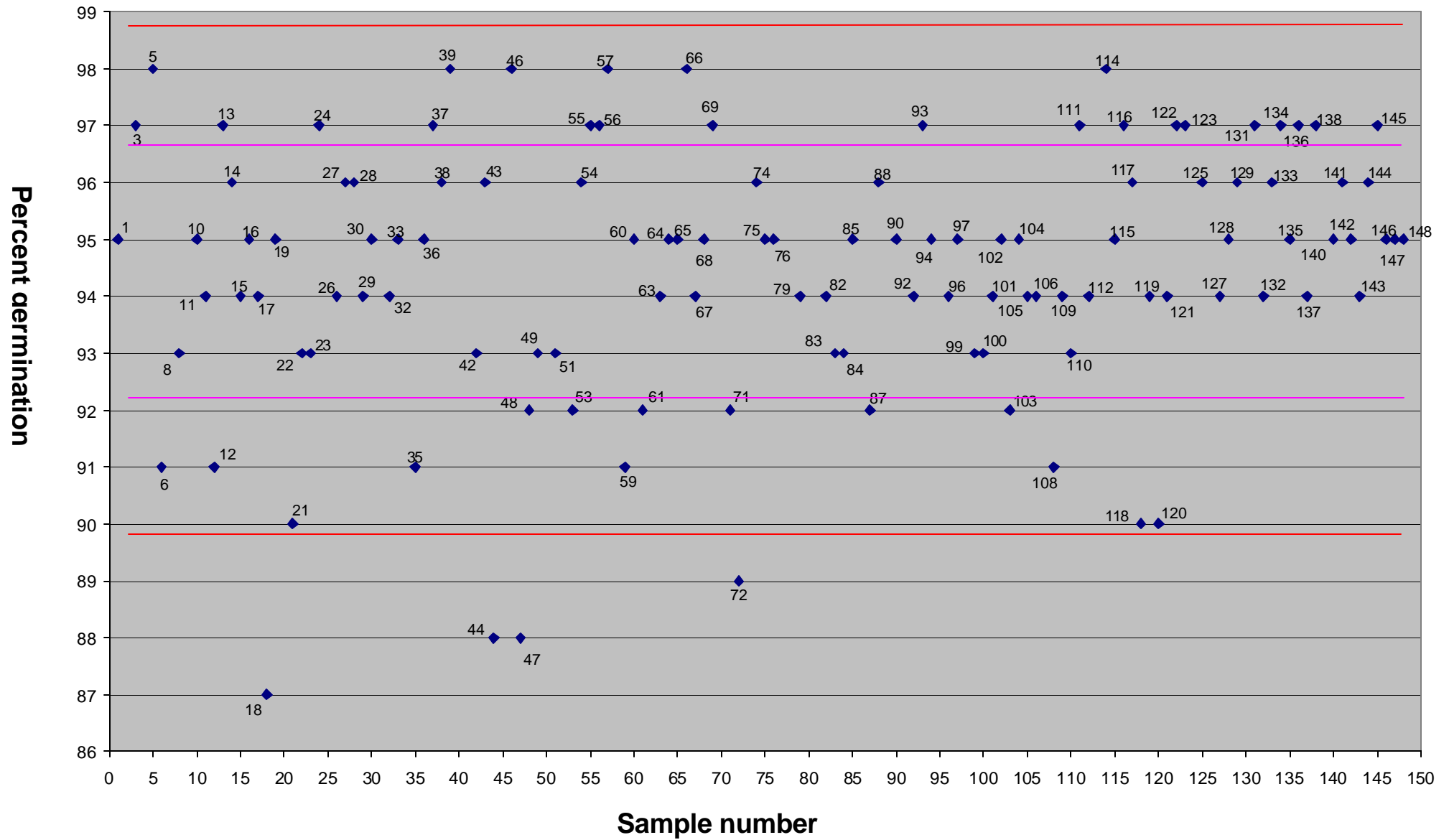
Dead seed evaluation (116 results returned)

- 66% fell within one standard deviation of the mean
- 99% fell within two standard deviations of the mean

Outlier sample number: 72

<i>Triticum aestivum</i> germination	Normal seedlings	Abnormal seedlings	Dead seed
Mean	94.48	3.42	2.09
Median	95	95	3
Standard deviation	2.24	2.24	1.97
Mean + 1 standard deviation	96.73	5.39	3.11
Mean - 1 standard deviation	92.24	1.46	1.08
Mean + 2 standard deviation	98.97	7.63	5.07
Mean - 2 standard deviation	89.99	-0.79	-0.88

Triticum aestivum Germination Graph



Legend

- One standard deviation
- Two standard deviations
- data marker w/sample number

Triticum aestivum (wheat) Purity Analysis

Pure seed evaluation (116 results returned)

- 73% fell within one standard deviation
- 94% fell within two standard deviations
- Outlier sample numbers: 8, 26, 44, 65, 67, 143, 145

Inert matter evaluation (116 results returned)

- 70% fell within one standard deviation
- 93% fell within two standard deviations
- Outlier sample numbers: 8, 61, 65, 67, 79, 128, 132

Weed seed evaluation (116 results returned)

- 86% fell within one standard deviation
- 95% fell within two standard deviations
- Outlier sample number: 15, 67, 96, 109, 143

Crop seed evaluation (116 results returned)

- 91% fell within one standard deviation
- 97% fell within two standard deviations
- Outlier sample number: 44, 48, 147

<i>Triticum aestivum</i> purity	% Pure seed	% Inert matter	% Weed seed	% Crop seed
Mean	99.10	0.60	0.16	0.14
Median	99.12	0.62	0.17	0.13
Standard deviation	0.22	0.20	0.04	0.09
Mean + 1 standard deviation	99.33	0.81	0.21	0.23
Mean - 1 standard deviation	98.88	0.40	0.12	0.05
Mean + 2 standard deviation	99.55	1.01	0.25	0.33
Mean - 2 standard deviation	98.66	0.20	0.08	-0.0422

Contaminants

The samples had an admixture of two *Convolvulus arvensis* (field bindweed), three *Aegilops cylindrica* (jointed goatgrass), 1 *Avena fatua* (wild oat), and 6 *Secale cereale* (rye). The lot itself contained *Hordeum vulgare* and *Triticum aestivum* ssp. (white wheat) these were reported in a few samples.

	C. arvensis	A. fatua	A. cyclidrica	S. cereale	Other
Mean	1.86	0.99	3	4.96	0.41
Median	2	1	3	6	0
St. deviation	0.39	0.09	0.81	1.58	1.5

Convolvulus arvensis:

- 88% reported two seeds of *Convolvulus arvensis*
- 11% reported one seed of *Convolvulus arvensis* and one seed of the following species:
 - 1 *Ipomea* sp.
 - 3 *Ipomea purpurea*
 - 2 *Convolvulus equitans*
 - 6 *Calystegia sepium*
 - 1 *Fallopia convolvulus*
- 1% did not report *Convolvulus arvensis*

Aegilops cylindrica

- 99% reported at least three seeds of *Aegilops cylindrica*
- 1% did not report *Aegilops cylindrical*

Avena fatua

- 99% reported one *Avena fatua*
- 1% did not report *Avena fatua*

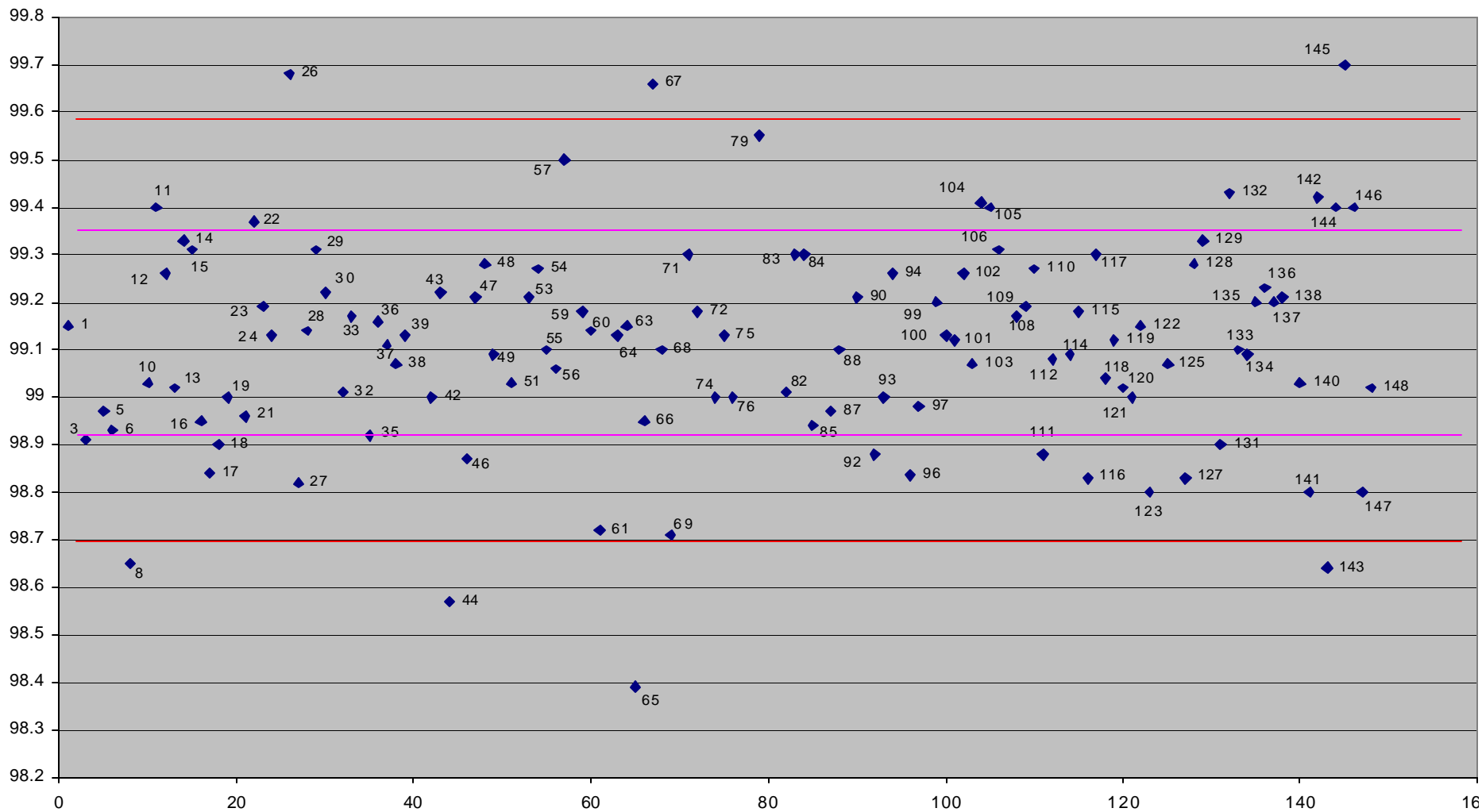
Secale cereale

- 2% did not report *Secale cereale*
- 3% reported one *Secale cereale*
- 5% reported two *Secale cereale*
- 4% reported three *Secale cereale*
- 14% reported four *Secale cereale*
- 21% reported five *Secale cereale*
- 48% reported six *Secale cereale*
- 1% reported seven *Secale cereale*
- 1% reported eight *Secale cereale*
- 1% reported nine *Secale cereale*

100% of identified seeds were correctly categorized as weed seed or crop seed.

Triticum aestivum Pure Seed Graph

Percent pure seed

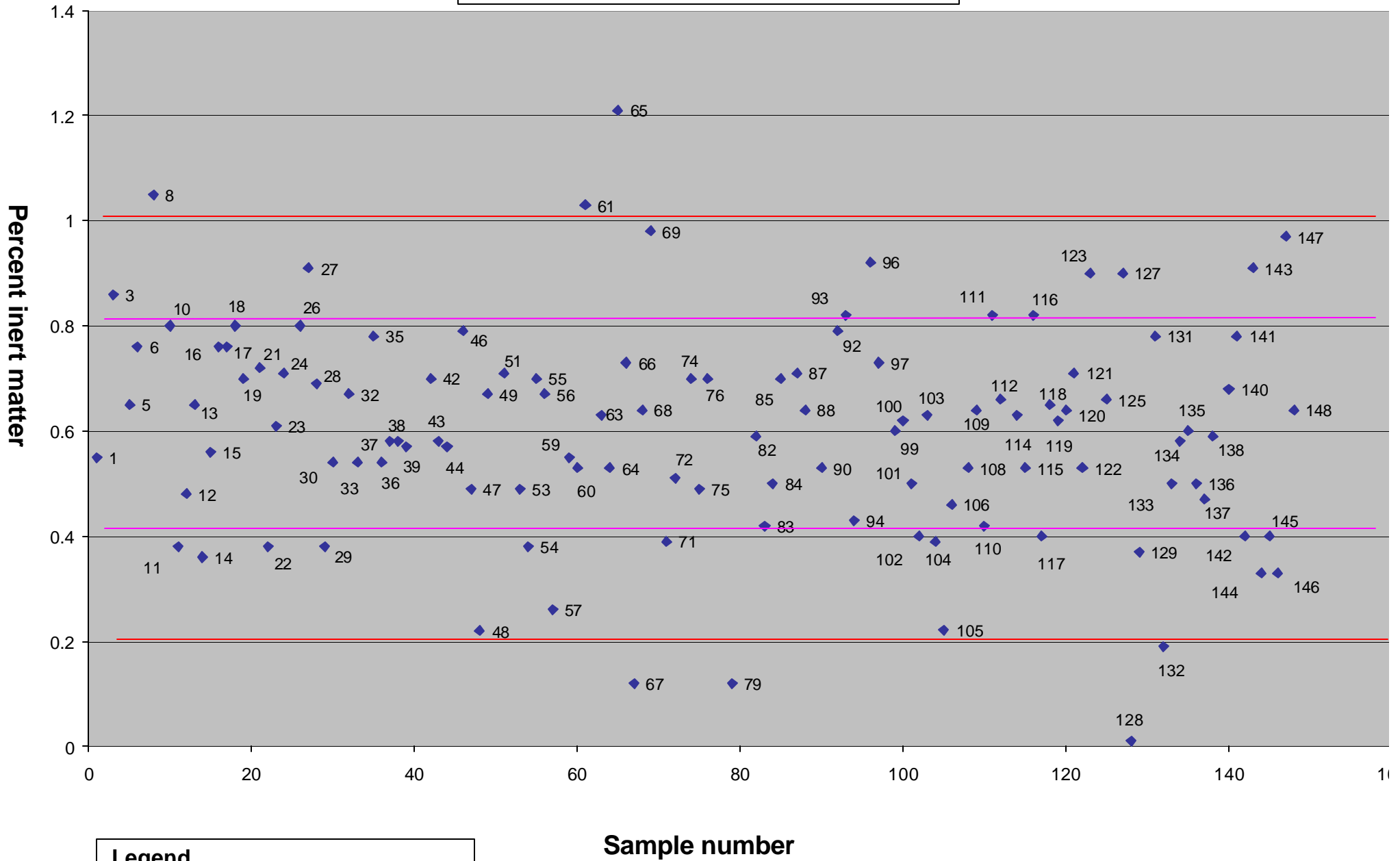


Sample number

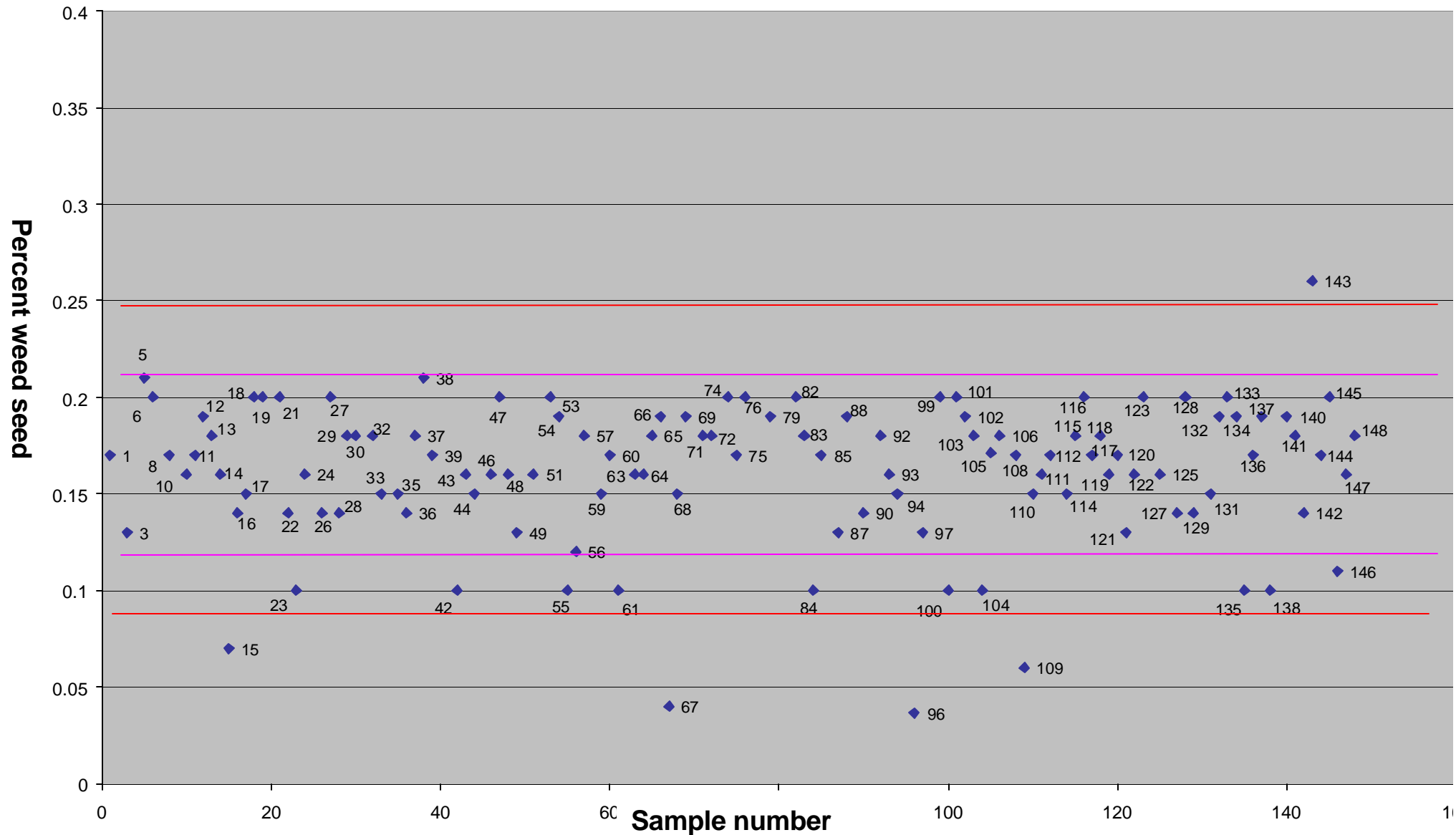
Legend

- One standard deviation
- Two standard deviations
- data marker w/sample number

Triticum aestivum Inert Matter Graph



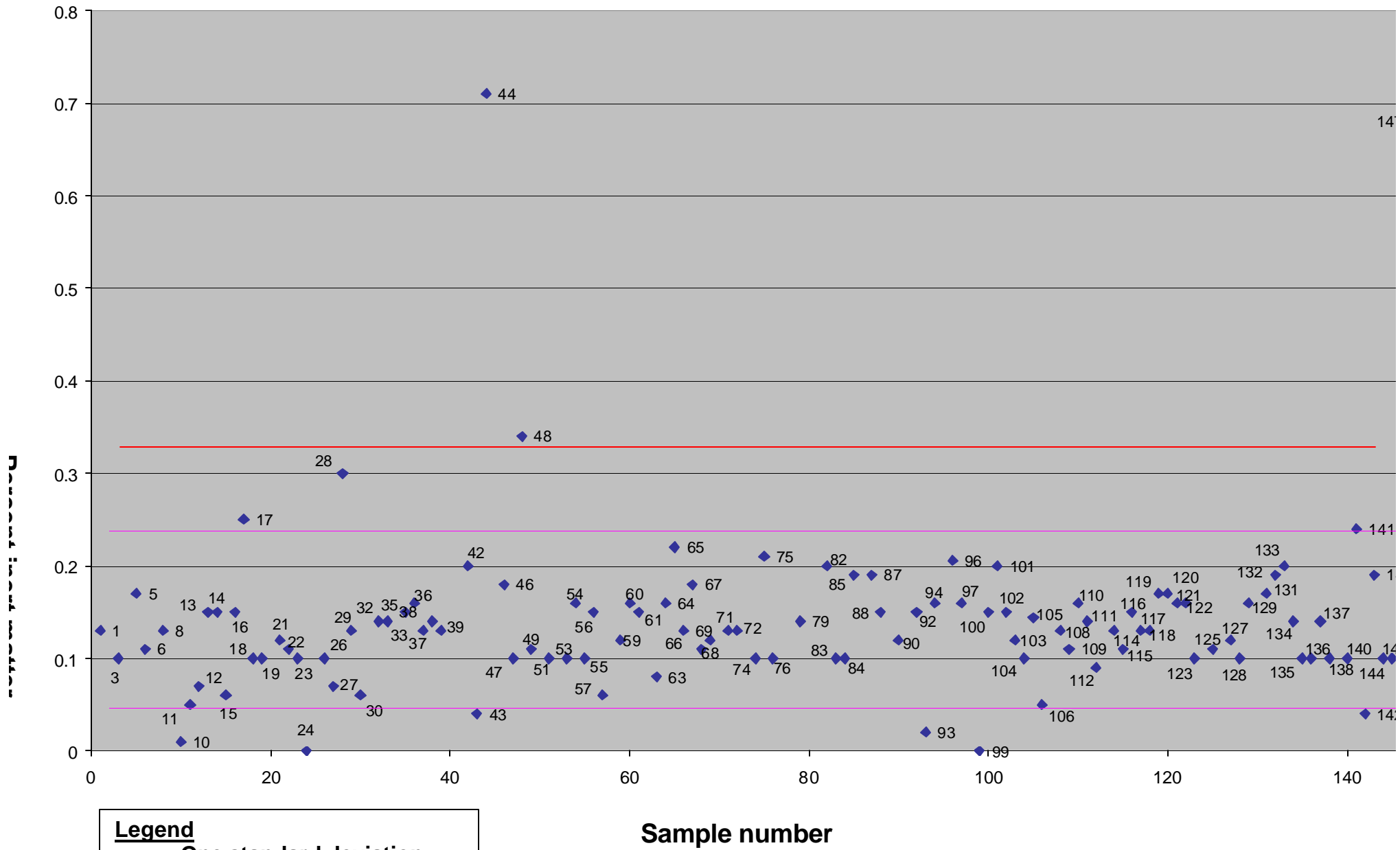
Triticum aestivum Weed Seed Graph



Legend

- One standard deviation
- Two standard deviations
- data marker w/sample number

Triticum aestivum Crop Seed Graph



Appendix A

Heterogeneity Testing

Cantaloupe germination- 13 samples tested in rolled towels @ 20-30 °C.

Sample #	Normal seedlings	Abnormal seedlings	Dead seed	
2	94	1	5	
4	92	1	7	
25	90	1	9	
45	91	2	7	
50	88	1	11	
73	89	2	9	
77	93	1	6	
81	92	1	7	
86	95	1	4	
95	93	2	5	
113	94	0	6	
130	92	1	7	
158	94	1	5	
Mean	92.08	1.15	6.77	
Median	92	1	7	

Soybean germination- 13 samples tested in rolled towels @ 25°C.

Sample #	Normal seedlings	Abnormal seedlings	Dead seed	Hard seed	
2	83	10	7	0	
4	83	10	7	0	
25	84	8	8	0	
45	83	7	10	0	
50	80	10	10	0	
73	81	10	9	0	
77	82	11	7	0	
81	89	7	4	0	
86	82	12	6	0	
95	84	12	4	0	
113	85	11	4	0	
130	84	11	5	0	
158	82	10	8	0	
Mean	83.23	9.92	6.85	0	
Median	83	10	7	0	

Wheat germination- 13 samples tested in rolled towels @ 15°C.

Sample #	Normal seedlings	Abnormal seedlings	Dead seed	
2	93	5	2	
4	95	3	2	
25	96	2	2	
45	93	5	2	
50	96	2	2	
73	96	2	2	
77	93	4	3	
81	97	1	2	
86	96	2	2	
95	96	2	2	
113	97	1	2	
130	95	3	2	
158	97	2	1	
Mean	95.38	2.62	2	
Median	96	2	2	

Wheat purity

Sample Number:	% Pure Seed	% Inert Matter	% Weed seed	% Crop seed
2	99.55	0.15	0.15	0.15
4	99.46	0.25	0.17	0.12
25	99.53	0.16	0.15	0.16
45	99.54	0.21	0.15	0.1
50	99.23	0.44	0.18	0.15
73	99.47	0.26	0.16	0.11
77	99.46	0.22	0.18	0.14
81	99.39	0.33	0.15	0.13
86	99.62	0.11	0.17	0.1
95	99.49	0.18	0.16	0.17
113	99.5	0.19	0.17	0.14
130	99.59	0.15	0.16	0.1
158	99.24	0.43	0.14	0.19
Mean	99.47	0.24	0.16	0.14
Median	99.49	0.21	0.16	0.14

Contaminants

Sample Number:	<i>Aegilops cylindrica</i>	<i>Convolvulus arvensis</i>	<i>Avena sativa</i>	<i>Secale cereale</i>
2	3	2	1	6
4	3	2	1	5
25	3	2	1	6
45	3	2	1	4
50	3	2	1	5
73	3	2	1	5
77	3	2	1	6
81	3	2	1	5
86	3	2	1	4
95	3	2	1	6
113	3	2	1	3
130	3	2	1	5
158	3	2	1	6
Mean	3	2	1	5.08
Median	3	2	1	5

Incidentals:

Hordeum vulgare and *Triticum aestivum* ssp. (white wheat) were reported in two of the samples.

Appendix B

Wheat sample preparation	Time in hours
Selecting lot for testing	6
Prep work & contaminant collection	3
Dividing & admixture preparation - 2 people	(combined time) 16
Packaging & shipping samples, misc.	2
Heterogeneity testing	9
Total	36

Soybean sample preparation	Time in hours
Selecting lot for testing	8
Dividing sample- 2 people	(combined time) 12
Packaging & shipping samples	1
Heterogeneity testing	6
Total	27

Canteloupe sample preparation	Time in hours
Selecting lot for testing	6
Dividing sample- 2 people	(combined time) 12
Packaging & shipping samples	1
Heterogeneity testing	6
Total	25

Statistical analysis of results	3
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Executive Director	Time in hours
Conference Calls (two)	3
Labeling samples	2
Packaging samples	7
Shipping samples	5
Recording results	12
Requesting results	2
Reporting & analysis of results	15
Total	46