

Catalog No. AQ 076 TCK 267/2367

Intended Use

This TotalTrait™ Comb is designed to extract and detect the presence of certain proteins at the levels typically expressed in genetically modified soybeans. The TotalTrait Comb may contain the following QuickStix™:

Protein/Trade Name	Sensitivity
CP4 EPSPS / Roundup Ready®	1 soybean in 400 (0.25%)
DMO / Roundup Ready 2 Xtend	1 soybean in 400 (0.25%)
PAT/pat / LibertyLink®	1 soybean in 200 (0.50%)
2m EPSPS / Balance GT27™	1 soybean in 1000 (0.10%)

How the Test Works

In order to detect the proteins expressed by genetically modified bulk grain, the sample must first be extracted to solubilize the protein. All QuickStix included in the TotalTrait Comb are specially formulated to detect their analytes in a Common Extraction.

Each QuickStix Strip has an absorbent pad at each end. The protective tape with the arrow indicates the end of the strips to insert into the reaction cup. The sample will travel up the membrane strips and be absorbed into the larger pad at the top of the strip. The portion of the strip between the protective tape and the absorbent pad at the top of the strip is used to view the reactions as described under “Interpreting the Results.” Results may then be scanned and interpreted quantitatively with the EnviroLogix QuickScan System. Please avoid bending the comb.

Sample Preparation

Step 1: Determine Number and Size of Sub-samples

1. Collect a composite sample according to USDA/FGIS instructions found in the reference documents listed in Precautions and Notes.
2. The following is another helpful reference for use in designing a sampling plan: Remund, K.M., Dixon, D.A., Wright D.L., Holden, L.R. “Statistical considerations in seed purity testing for transgenic traits”, Seed Science Research, June 2001, Vol. 11 No.2, pp. 101-119.
3. To select the appropriate sample size, determine the purity standard and the degree of confidence required. Confidence level means the statistical probability that the % of GMO soybeans in the lot is below the selected purity standard. This calculation should be done for each trait tested, then choose the largest sub-sample volume result.

Step 2: Determine Sub-sample Weight, Jar Size, Grind Times and Water Volume Needed for Sample Preparation

1. Determine the **average weight** of the grain from the lot to be tested. Count and weigh 100 kernels/seeds, then divide by 100.
2. Calculate the sub-sample weight (g) needed for testing, (number of seeds X **average seed weight**).
3. Calculate water volume needed for sample preparation. The Soybean Common Extraction uses a water volume to sample weight ratio of **5 to 1**.

Example Calculation using a 100 seed sub-sample with an average seed weight of 0.15g.
 $0.15g \times 100 = 15g \times 5 \text{ mL} = 75 \text{ mL water for extraction}$

4. Choose an appropriate jar size and grind time based on the type of blender available for sub-sample preparation (see Table 2). Oster Sunbeam Blender with ice crusher blade is recommended over the Waring Blender for its bean grinding efficiency. (Note that bean grind time is longer and requires additional steps* when using a Waring Blender). The grind quality should be such that 40-50% passes through a 20-mesh sieve.

Table 2 - Soybeans

# of Beans (approximate)	Blender Type	Sub-sample weight (g)	Jar size (oz.)	Grind Time on High Speed
100-200	Oster Sunbeam	16-38	8	20 seconds
100	Waring	16-38	8	60 seconds (2 X 30 sec.*)
200-400	Waring	38-65	16	60 seconds (2 X 30 sec.*)

* For best results blend beans for ½ of total time, remove the jar and shake to redistribute larger particles, replace and resume grinding.

Contents of Kit:

- 100 or 40 TotalTrait Combs, packaged 5 combs per foil pouch
- Sample cups and disposable transfer pipettes

Items Not Provided:

- QuickScan System*
- Blender for sample prep:
 1. Oster® Sunbeam blender model #4094 or equivalent (with 4 oz. polystyrene blender jar, ice crusher blade, gasket, and blender base) ~or~
 2. Waring blender model 31BL91 or equivalent (with glass Mason jars and jar adapter [Eberbach #E8495] along with protective cover ~or~
 3. BUNN coffee grinder (industrial style grinder set on AutoDrip setting), plus extraction vessels for sample and water
- Graduated cylinder
- Tap water
- Protective cover for blender jar while grinding
- Digital scale for weighing samples
- Timer
- Scissors

Step 3: Prepare the Sample

1. Weigh sample into the appropriate vessel.
2. Put protective cover over glass jars.
3. Grind sample on high speed all whole beans are finely ground.
4. Add the volume of tap water calculated above (see formula above or at left).
5. Cap and shake jar vigorously for 30 seconds, ensuring that the entire sample is wet. Sample will begin to settle immediately and liquid can be drawn off at that time. Avoid pulling up particles with the transfer pipette.
6. Transfer 12 mL of the liquid portion from above the settled sample into the sample cup. The level should be above the bottom of the arrows but below the top of the lower colored portion of the TotalTrait Comb strips.
7. To prevent cross-contamination, thoroughly clean blender parts and jars to remove dust and residue prior to preparation of each sample and use a new sample cup for each. If pipetting, use a new tip or disposable pipette for each sample.

How to Run the TotalTrait Comb Test

1. Allow refrigerated foil pouch to come to room temperature before opening. Remove the combs to be used. Avoid bending the strips or handling the loose comb end.
2. Place the comb into the three-ounce cup containing 12 mL of the liquid soybean extract. The sample will travel up the strips.
3. Allow the comb to develop for 5 minutes before making final assay interpretations.
4. Remove the TotalTrait Comb from the cup. Cut off and discard the bottom section of each strip covered by the arrow tape, and insert into QuickScan for reading.

Note: Use extreme caution to prevent sample-to-sample cross-contamination with grain, fluids or disposables.

Use of the QuickScan System

Detailed instructions for use of the QuickScan System are supplied with each unit and can also be found at envirologix.com/support. In summary, the TotalTrait Comb is inserted into the reader and the strips are read by touching or clicking on the “Read Test” area of the screen. TotalTrait combs utilize the QuickScan Software feature “TotalTrait GMO Soy Sum,” a proprietary feature than when activated returns the most accurate GMO total when screening for today’s traitlet soybeans. Activation and use of the TotalTrait GMO Soy Sum is recommended over the legacy GMO Soy Sum. Results are then recorded in an electronic worksheet, allowing each user to report and track data easily.

Interpreting the Results

Development of the Control Lines within 5 minutes indicates that the strips have functioned properly. Any strip that does not develop a Control Line should be discarded and the sample re-tested using another comb or corresponding strip.

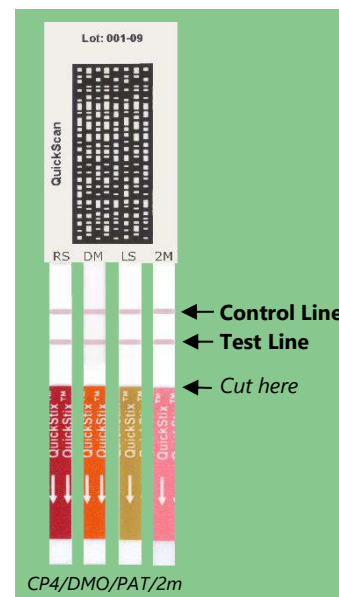
Results are scanned and interpreted quantitatively with the QuickScan System. QuickScan will return a result as “% GMO” or “<LOD” (less than the Limit of Detection). Please consult the QuickScan User Manual for details.

Kit Storage

This Kit may be stored at room temperature or refrigerated for longer shelf life. Please note the shelf life on the kit box for each storage temperature. The Kit may be used in field applications; however, prolonged exposure to high temperatures may adversely affect the test results. **Important: do not open the foil pouch until ready to use the combs. Allow container to come to room temperature before opening to prevent condensation.**

Precautions and Notes

- This kit is designed to be read quantitatively when used with the QuickScan System.
- As with all tests, it is recommended that results be confirmed by an alternate method when necessary.
- The assay has been optimized to be used with the protocol provided in the kit. Deviation from this protocol may invalidate the results of the test.
- The results generated through the proper use of this kit reflect the condition of the working sample directly tested. Extrapolation as to the condition of the originating lot, from which the working sample was derived, should be based on sound sampling procedures and statistical calculations which address random sampling effects, non-random seed lot sampling effects and assay system uncertainty. A negative result obtained when properly testing the working sample does not necessarily mean the originating lot is entirely negative for the analyte or protein in question.
- Protect all components from hot or cold extremes of temperature when not in use. Do not leave in direct sunlight or in a vehicle.
- CAUTION: Tightly closed containers of soy extract, if left sitting for several hours, may ferment and cause the lid or container to burst. Dispose of extract when testing is complete.



- USDA/AMS/FGIS (formerly GIPSA) References:
 - www.gipsa.usda.gov/fgis/handbook/BK1/Book1_2015-09-18.pdf - USDA Grain Inspection Handbook, Book 1, Grain Sampling.
 - www.gipsa.usda.gov/fgis/biotech/sample2.htm - Guidance document entitled Sampling for the Detection of Biotech Grains.
 - www.gipsa.usda.gov/fgis/biotech/sample1.htm - Practical Application of Sampling for the Detection of Biotech Grains.
 - www.gipsa.usda.gov/fgis/biotech/samplingplan1.xlsx - This website provides a simple to use Sample Planner (29K Excel Spreadsheet). The planner allows you to enter different assumptions in terms of sample size, number of samples, acceptable quality level and to determine the probability of accepting lots with given concentration levels. It also plots the probabilities in graph form for easy interpretation. Specific data can be saved for documentation and future analyses.

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For Technical Support
Contact Us At:

EnviroLogix
 500 Riverside Industrial Parkway
 Portland, ME 04103-1486 USA
Tel: (207) 797-0300
Toll Free: 866-408-4597
Fax: (207) 797-7533

e-mail: info@envirologix.com

website:
www.envirologix.com