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March 17, 2015

Dr. Michael J. Firko Deputy Administrator, Biotechnology Regulatory Services 4700 River Road, Unit 98 Riverdale, MD 20737

RE: Confirmation of regulatory status

Dear Dr. Firko:

I am writing to seek confirmation from Biotechnology Regulatory Services that a new maize crop line developed by Agrivida, Inc., does not meet the definition of a regulated article for purposes of 7 CFR Part 340. Agrivida, Inc. is a biotechnology company focusing on the development of new crop varieties targeted for the animal nutrition and agricultural processing industries.

Agrivida has introduced into maize a meganuclease enzyme that specifically recognizes and creates a deletion in exon for the endogenous maize gene. Since the gene responsible for expressing the meganuclease was introduced into the maize genome at a position that was not genetically linked to the target gene, it was then possible to segregate the introduced DNA from the variant allele following crosses with wild-type maize plants. In this manner, all introduced DNA was eliminated from those progeny plants containing the variant allele.

The process used to generate maize plants with the variant and allele involved the introduction of a meganuclease transgene via and the meganuclease transformation, utilizing the meganuclease expression cassette and the plant selectable marker included:

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from

Meganuclease expression cassette

- promoter
- A sequence derived from
- A meganuclease from
- terminator from

Selectable marker expression cassette

- promoter
- The gene encoding
 terminator from

Following crosses to wild-type maize plants to segregate away the meganuclease sequences, the absence of all introduced DNA sequences in the variant segregating lines was confirmed by both PCR and Southern blot analyses. We submit that the resulting null-segregant (NS) lines, which contain no introduced DNA sequences, are not, and would present no risk of creating, a plant pest, as defined under the Plant Protection Act.

In wild-type maize plants, **and the actions of several enzymes in a process that mediates the diurnal turnover of transient starch in maize. While heterozygous plants that contain one wild-type allele and one variant and** allele behave as wild-type plants, with no significant phenotypic variation, homozygous maize plants expressing the variant **allele** accumulate substantially higher levels of starch in their leaves and stalks than comparable wild-type plants. The increased starch phenotype observed in the variant **allele**, which causes gradual accumulation of starch in leaf and stalk tissue over the life span of these plants.

Because maize plants with the variant allele, derived from our targeted gene deletion strategy, contain no foreign DNA sequences and are modified only by specific removal of base pairs in exon of the endogenous gene, we submit that these modified materials do not comprise APHIS-regulated articles, for purposes of 7 CFR Part 340.¹

Please confirm that this understanding is correct.

Sincerely,

R. Michael Raab Chief Technology Officer Agrivida, Inc. 617-905-9500 Michael.Raab@Agrivida.com

¹ We note that APHIS has reached similar conclusions with respect to other analogous products. *See, e.g.,* Letter from Michael C. Gregoire (BRS) to Gary W. Rudgers (Dow AgroScience), March 8, 2012 (APHIS agrees that maize plants that have been engineered to contain a gene deletion by using zinc finger nuclease technology are <u>not</u> regulated articles for purposes of 7 CFR Part 340). *See also* Letter from Michael C. Gregoire (BRS) to Sally Mackenzie (University of Nebraska), June 6, 2012 (APHIS confirms that null segregant lines derived from plants that were genetically engineered utilizing *Agrobacterium*-mediated transformation, but that contain no transgenic material or plant pest sequences, are <u>not</u> regulated articles for purposes of 7 CFR Part 340).