

October 16, 2023

Bernadette Juarez APHIS Deputy Administrator Biotechnology Regulatory Services

# Administratively revised version of 22-174-01rsr Contains Confidential Business Information

## Re: Request for Regulatory Status Review for Glufosinate Herbicide Tolerant Camelina sativa

Yield10 Bioscience is requesting a regulatory status review for its glufosinate herbicide tolerant *Camelina sativa* lines based on the provisions in 7 CFR § 340.4.

#### 1. Requestors

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# 2. Confidential Business Information (CBI) Statement

This RSR request contains CBI.

CBI Justification:

The Freedom of Information Act (FOIA) exempts federal agencies from releasing trade secrets and commercial or financial information that is privileged or confidential (5 U.S.C.552(b)(4)). Yield10 Bioscience considers certain information in this document as trade secret or commercial information that is privileged and confidential. Disclosure of this information would cause substantial competitive harm to Yield10 Bioscience by allowing other companies to unfairly compete with it.

First, Yield10 Bioscience must protect the nature of traits which it has selected to be important, that, if imparted to a new plant variety, would represent a competitive advantage in the marketplace. Disclosure of this information would reveal the company's marketing strategy, which identifies targets of potential commercial opportunity. Second, Yield10 Bioscience must

keep some aspects of its research confidential: what it is doing, how it is doing it, and how far along it is. Disclosure of this information would enable competitors to duplicate Yield10 Bioscience research and development without incurring the investment of time and money expended by the company. This information would also provide competitors with commercially valuable knowledge about the specific products Yield10 Bioscience is interested in commercializing and the likely time for commercialization. Moreover, Yield10 Bioscience must protect its intellectual property. Yield10 Bioscience must keep research information strictly confidential because in some cases, patent applications have not been filed or patents are pending and have not been published.

Specifically, Yield10 Bioscience designates the following as Confidential Business Information:

Genetic elements used to create trait. Yield10 Bioscience biotechnology traits consist of vectors transferred into plants, which comprise genes for the expression of traits and regulatory sequences such as promoters, enhancers, signaling peptides and terminators. Disclosure of genetic elements comprising the genotype of the resulting plants would provide the company's competitors with the knowledge of what genetic sequences the company is using. Disclosure of this information may also reveal the specific modifications the company made in assembling the DNA constructs and enhance their usefulness. It is in Yield10 Bioscience's commercial interest that these trade secrets are not publicly disclosed.

Construct Identity. Yield10 Bioscience technology traits are encoded by genetic constructs that the company identifies with unique nomenclature. Disclosure of the construct identity could enable competitors to determine the status of the company's proprietary research and development programs, if construct identity is linked to the genetic elements contained in these constructs.

Phenotype Descriptions. Yield10 Bioscience seeks to improve certain characteristics of plants that are economically important to its customers. Disclosure of the phenotypes that the company is pursuing would enable competitors to determine the company's R&D objectives and the status of its R&D programs.

Accordingly, Yield10 Bioscience requests all FOIA protections for the CBI submitted herein.

This information is both customarily and actually treated as private information by Yield10 Bioscience and was provided to the government under an assurance of privacy. We consider this information as confidential within the meaning of 5 U.S.C §552(b)(4), the Freedom of Information Act's Exemption 4.

#### 3. Description of Comparator Plant

Scientific Name: *Camelina sativa* Common Name: Camelina, False Flax, Gold of Pleasure Cultivar and/or Breeding Line: [ used to produce the transgenic herbicide tolerant plants.

]. These breeding lines will be **CBI-deleted** 

There are two types of Camelina, spring and winter, named for their respective growth habit. Both are promising crops that produce high levels of seed oil (Vollmann and Eynck, 2015; Berti et al. 2016.; Malik et al. 2018). The spring growth habit is dominant in Camelina. Winter lines are better adapted as cover crops in colder, temperate climates. Winter lines require vernalization, or exposure to cold, with temperatures below 8°C for 2-3 weeks at seedling to rosette stage to induce bolting and flowering, whereas spring lines do not (Anderson et. al. 2018). Camelina plants are highly sensitive to herbicides including glufosinate, glyphosate, sulfonylureas, and imidazolonones. Yield10 Bioscience is engineering glufosinate tolerance [\_\_\_\_\_\_]. CBI-deleted

4. Genotype of the Modified Plant

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a. Sequence of the Insertion

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b. Table 1: Annotation of the Inserted Genetic Material	]	CBI-deleted
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The expression cassette for <i>bar</i> in plasmid [	CBI-deleted
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c. Information about insertion site. Plants containing single copy inserts of the T-DNA have been isolated	d
after Agrobacterium-mediated transformation of Camelina lines [	CBI-deleted
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]. After identification of the best events for commercial advancement in field trials, **CBI-deleted** whole genome sequencing will be carried out to ensure the presence of the desired expression cassette for *bar* and the absence of unintended DNA fragments derived from the vector (plasmid) backbone. Sequence analysis of flanking genomic DNA regions will also be used to determine the loci of the T-DNA insert and its junctions with the plant genome to confirm insert integrity and location.

#### 5. Description of New Trait

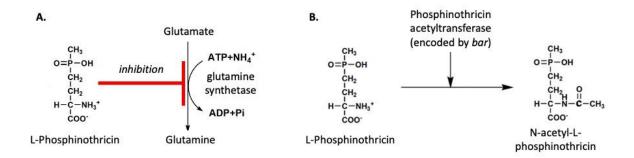
**a. Intended trait.** Herbicide tolerance. The inserted genetic material imparts Camelina plants with tolerance to glufosinate, a broad spectrum post-emergent broadleaf herbicide.

**b.** Intended phenotype. Tolerance to commercial herbicides containing the active ingredient glufosinate.

c. Description of the Mechanism of Action (MOA). The mechanism of action of the PAT protein has been well studied (for review see Takano and Dayan, 2020). The herbicide glufosinate (or L-phosphinothricin) inhibits glutamine synthase (FIG. 1.A) when applied to plants such that high levels of ammonia accumulate and inhibition of the photorespiratory pathway and photosynthesis occur, killing the plant (Takano and Dayan, 2020). Expression of the *bar* gene [ ] in transgenic plants results in the production of phosphinothricin N-acetyltransferase (PAT), a protein which catalyzes the conversion of L-phosphinothricin (L-PPT) to the non-phytotoxic form, N-acetyl-L-phosphinothricin, by acetylation (FIG. 1B). Plants engineered with a genetic construct for expression of the *bar* gene are thus tolerant to post-emergent applications of glufosinate containing herbicides (Thompson et al., 1987; De Block et al., 1987; Wehrmann et al., 1996). [ CBI-deleted

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**Figure 1. Glufosinate detoxification by expression of the** *bar* gene encoding phosphinothricin acetyltransferase. A. Glufosinate (also known as L-phosphinothricin) exhibits its herbicidal action through inhibition of glutamine synthetase, resulting in the accumulation of ammonia and inhibition of photosynthesis. B. Expression of the *bar* gene in transgenic plants, encoding phosphinothricin acetyltransferase, converts L-phosphinothricin to N-acetyl-L-phosphinothricin such that the transgenic plant is tolerant to glufosinate.

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We have provided the following elements which support our RSR request:		
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2.		CBI-deleted
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3. The mechanism of action of genetically engineered glufosinate tolerance in plants is	well studied	
(Figure 1) [		CBI-deleted
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Please let us know if you have any further questions or need additional information as you re request. Sincerely,	eview our	

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### References

	Anderson, J. V., Horvath, D. P., Doğramaci, M., Dorn, K. M., Chao, W. S., Watkin, E. E., Hernandez, A. G., Marks, M. D. and Gesch, R. 2018. Expression of FLOWERING LOCUS C and a frameshift mutation of this gene on chromosome 20 differentiate a summer and winter annual biotype of Camelina sativa. <i>Plant Direct</i> <b>2</b> , e00060. Berti, M., Gesch, R., Eynck, C., Anderson, J. and Cermak, S. 2016. Camelina uses, genetics, genomics, production, and management. <i>Ind Crop Prod</i> <b>94</b> , 690-710.	
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7.	Takano, H. K. and Dayan, F. E. 2020. Glufosinate-ammonium: a review of the current state of knowledge. Pest Manag Sci. 76: 3911-3925.	
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