

31.1.2024

A9-0014/138

Amendment 138

Benoît Biteau

on behalf of the Verts/ALE Group

Report

A9-0014/2024

Jessica Polfjärd

Plants obtained by certain new genomic techniques and their food and feed
(COM(2023)0411 – C9-0238/2023 – 2023/0226(COD))

Proposal for a regulation

Chapter II

Text proposed by the Commission

Amendment

[...]

deleted

Or. en

31.1.2024

A9-0014/139

Amendment 139

Benoît Biteau

on behalf of the Verts/ALE Group

Report

A9-0014/2024

Jessica Polfjärd

Plants obtained by certain new genomic techniques and their food and feed
(COM(2023)0411 – C9-0238/2023 – 2023/0226(COD))

Proposal for a regulation

Annex I

Text proposed by the Commission

Amendment

Criteria of equivalence of NGT plants to conventional plants

deleted

A NGT plant is considered equivalent to conventional plants when it differs from the recipient/parental plant by no more than 20 genetic modifications of the types referred to in points 1 to 5, in any DNA sequence sharing sequence similarity with the targeted site that can be predicted by bioinformatic tools.

(1) substitution or insertion of no more than 20 nucleotides;

(2) deletion of any number of nucleotides;

(3) on the condition that the genetic modification does not interrupt an endogenous gene:

(a) targeted insertion of a contiguous DNA sequence existing in the breeder's gene pool;

(b) targeted substitution of an endogenous DNA sequence with a contiguous DNA sequence existing in the breeder's gene pool;

(4) targeted inversion of a sequence of any number of nucleotides;

(5) any other targeted modification of any size, on the condition that the resulting DNA sequences already occur (possibly with modifications as accepted under points (1) and/or (2)) in a species from the

AM\P9_AMA(2024)0014(138-139)EN.docx

PE756.833v01-00

breeders' gene pool.

Or. en