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1 The Case for Reforming the EU Regulatory System for GMOs.

- 2 Christopher J. Pollock* and Rosemary S. Hails**.
- 3
- 4 *Institute of Biological, Environmental and Rural Sciences, Aberystwyth
- 5 University, Penglais, Aberystwyth SY23 3FG and **NERC Centre for Ecology

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- 6 and Hydrology, Wallingford OX10 8BB, UK.
- 7
- 8 Corresponding Author: Pollock, C.J (cip@aber.ac.uk)
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1	Background. The first iteration of the EU Directive that controls the
2	deliberate release of GMOs into the environment was adopted in 1990, and
3	the first commercial planting of GM crops occurred in 1996 in the USA (or
4	1998 in the EU). Since then, the global cultivated area has risen to exceed
5	170 M ha and the majority of global production of soya and cotton is via GM
6	varieties [1]. Over this period, cultivation within the EU has been restricted to
7	ca 0.1M ha. Although the main barrier to release of GM food crops for
8	cultivation in the EU lies at the political level, the UK Advisory Committee on
9	Releases to the Environment (ACRE) has recently raised concerns both about
10	the regulatory approach adopted by the EU and the implementation of the
11	current current processes for carrying out Environmental Risk assessments
12	(ERA) <u>process and about their future fitness for purpose (ACRE</u>
13	http://www.*wp1-3). This article summarises these concerns and suggests a
14	way forward.
15	Addressing current challenges. These include the lack of precision with
16	which the concept of adverse effects (harm) is used within ERA and the
17	challenges of developing a proportionate approach to unanticipated effects
18	Adverse Effects. There is no consensus within the EU as to what constitutes an
19	adverse environmental effect. This has led to increasing data requirements and
20	to inappropriate recommendations for risk management options. The, and this
21	leads to difficulties asre illustrated by GM herbicide tolerant crops. These
22	systems allow farmers to practice efficient weed control to enhance crop
23	productivity, potentially at the expense of biodiversity that depends upon weeds
24	at the base of the food chain. This negative impact could be compensated for by

Comment [RSH1]: Actually the example we give, with GMHT crops, is about absence of defined risk management because harm has not been defined. This more general statement is more relevant to GMIR crops.

1 setting aside field margins or other areas for farmland biodiversity, but in the 2 absence of clear guidelines on what does or does not constitute harm, appropriate compensatory measures cannot be usefully defined. Defining harm 3 would also facilitate a more structured approach to ERA [2], thus reducing the 4 inclusion of extraneous information and improving the value of the ERA in 5 decision-making. 6

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8	Monitoring and dDealing with the unexpected. In addition to assessing the risks	
9	associated with the intended changes made to a GM crop, ERA must also	
10	address the possibility that any identified unintended changes could cause	Comment [RSH2]: I think its is in this section to distinguish betwee
11	harm. In the absence of a plausible link between a characteristic of a GM crop	identified unintended changes and 'unknown unknowns'
12	(and its use) and such harm, there is the potential for open-ended evidence-	
13	gathering that does not add value to the risk assessment. It is not possible	
14	practical to test for ERA to identify unintended effects in advance by testing	
15	every characteristic of a GM crop and its use under every conceivable scenario.	
16	It is important therefore, that the EU adopts a proportionate approach that	
17	makes optimal use of existing evidence. There is a significant 'weight of	
18	evidence' available from the cultivation of GM crops outside of the EU as well as	3
19	from existing information requirements. Where this evidence indicates changes	
20	to a GM crop, or its use ,are identified and potentially that can be linked to	
21	adverse effects, the ERA should follow a structured and proportionate process	
22	to characterise the risk to the environment. Where identified but unintended	
23	changes cannot be plausibly linked to adverse effects, further data should not	
24	be required.	Comment [RSH3]: I think this conclusion is required for clarity

(SH2]: I think its important o distinguish between ended changes and owns'

1	Finally, some countries, including those in the EU, require post-market	
2	monitoring to detect any unanticipated adverse effects from the commercial	
3	growing of GM crops. In this context, WhereIn terms of unintended effects,	
4	cannot be defined nor linked to a GMO, ACRE supports the use of effective	
5	questionnaires_to detect significant changes aimed at users of the product but	
6	is doubtful of the value of large-scale environmental monitoring to attribute	
7	causality to the cultivation of GM crops	Comment [RSH paragraph is needed
8	(http://www.defra.gov.uk/acre/files/pmem-final-report.pdf; ACRE http://* WP2).	
9	Emerging Challenges For Process-based Regulation. The current EU	Comment [RSH Louise about this se
10	Directive lists techniques that can lead to the generation of GMOs and those	' ACRE has already understanding how
11	which are deemed not to. However, these lists are not exhaustive, which leads	it's important to get The original draft ta
12	to confusion in interpretation of the regulations as to whether the legislation	based approach then whether the product without connecting
13	applies to novel organisms developed by techniques that were not envisaged	text from the end to clear that the definit there are problems
14	when the legislation was adopted in 1990. This approach to regulating	process –based bit of with transgenics and is nonsensical giver
15	organisms based on how they were produced (so-called "process-based") was	about genomes. I amended the text s
16	adopted in 1990 and at this time, many of the techniques being used and	confusing and adde element in the last s
17	developed today were not envisaged. Consequently, it is not clear whether the	
18	organisms produced by them are captured by the GMO legislation. This results	
19	in regulatory uncertainty – a major block to innovation and to effective ERA.	Comment [RSH Louise: 'Why? Onc
20	Adoption of a process-based regulatory system dditionally, basing a regulatory	carry out an effectiv
21	system on how organisms are produced has resulted in also results in	
22	organisms with the same phenotype being dealt with differently (e.g.	
23	phenotypically identical herbicide-tolerant plants produced by transgenesis or	
24	traditional mutagenesis). This disparity may increase become more evident	
25	depending on whether regulators consider organisms produced by <u>a suite of</u>	

Comment [RSH4]: And |I think this paragraph is needed.

comment [RSH5]: Comment from ouise about this section:

ACRE has already been criticised for not understanding how the legislation works so it's important to get this right.

The original draft talks about a processbased approach then, at the end talks about whether the product occurs naturally without connecting the 2.. I've moved the text from the end to the front to make it clear that the definition has 2 parts and there are problems with both parts i.e. process –based bit only categorically deals with transgenics and the genotype element is nonsensical given what we now know about genomes.

I amended the text so that it wasn't so confusing and added the 'genotype' element in the last section.'

Comment [RSH6]: Comment from Louise: 'Why? Once captured we could carry out an effective ERA'.

I	new techniques as GMOs. Regulation of process-based approaches is both
2	open to interpretation and difficult to future-proof. PFor example, plants
3	developed using chemical or radiation mutagenesis are exempt from existing
4	EU regulation. However, the same trait in the same organism caused by the
5	same sequence alteration but using a different mutagenic process (such as zinc
6	finger nucleases, TALENS or oligonucleotides) may be covered, depending on
7	the interpretation of the EU Commission and member-state regulators.
8	Furthermore, Ttheir views are likely to differ as reflected in It is likely that the
9	conclusions will reflect those set out in the EU Commission's working group
10	report on New Technologies [3]. Other emerging technologies involveclude GM
11	techniques that are captured by the definition of a GMO during the as part of the
12	breeding process but that result in an organisms that does not contain any
13	inserted or recombinant DNA (e.g. reverse breeding). As the process requires
14	the use of a GM intermediate, some may argue that the resultant organism is
15	captured by the directive-GMO legislation whereas others may considernot.
16	that it is not because the product could have occurred naturally. Comment [m7]: This concept hasn't been introduced.
17	
18	We conclude that a process-based approach to regulation is difficult to interpret
19	and to future-proof and this will become more apparent as the technology and
20	its application advances.problematic now, and likely to become more so in the
21	future. This is also consistent with current views of genome organisation that do
22	not align with the EU definition of a GMO ("an organismin which the genetic
23	material has been altered in a way that does not occur naturally by mating
24	and/or natural recombination.") When this was written, genomes were
25	considered to be relatively uniform and stable and the definition was intended to

1	capture the range of artificially induced alterations to an organism's genetic	
2	material. Now that the extent of natural genetic and epigenetic variation is more	
3	fully understood, identifying alterations that do not occur naturally is much more	
4	challenging. Within an individual and between individuals of the same species,	Comment [m8]: Confuses the story line
5	the genetic material can exist in naturally altered forms whilst the phenotype	
6	remains unchanged, through the effects of genetic redundancy between	
7	multigene family members [4,5] Equally, a single nucleotide change can result	
8	in a significant phenotype change (ACRE http://www.wp3*).	
9		
10	Options for Change. A major reason for regulating organisms based on how	
11	they were developed is to address the concern that the technologies may cause	
12	unintended alterations to the organisms' genetic material that could not have	
13	occurred naturally or through conventional breeding. Since the GMO legislation	
14	was adopted in 1990 we have learned a great deal from genomic studies. We	
15	now understand that this notion that genomes are relatively uniform and stable	
16	entities is incorrect. Within an individual and between individuals of the same	
17	species, the genetic material can exist in naturally altered forms whilst the	
18	phenotype remains unchanged, through the effects of genetic redundancy	
19	between multigene family members [4,5] Equally, a single nucleotide change	
20	can result in a significant phenotype change (ACRE http://www.wp3*). We have	
21	also gained significant evidence from the widespread use of GMOs. From this	
22	we conclude that We propose that there is little scientific justification for a	
23	regulatory system that is triggered by the process by which new organisms are	
24	developed (i.e. recombinant DNA technology) rather than by their novelty or	
25	potential for harm (i.e. their phenotype).	

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2	Additionally, we suggest that a regulatory system that takes account of potential
3	benefits and includes potential compensatory measures (thus encouraging a
4	more explicit cost-benefit approach) would better align with other regulatory
5	frameworks and result in greater potential benefits for the environment and
6	human health than the existing system. This approach would also improve the
7	efficiency of the process to ensure appropriate risk-assessment without stifling
8	innovation.technological . It and would obviate the pressure to develop need for
9	technological approaches that are developed purely to deliver material that-lies
10	outside current regulations (ACRE http://www.wp2*). Shifting to a phenotype-
11	based system would-also permit resolution of the wider issues discussed above
12	and promote proportionate and effective ERA-assessments backed by
10	enprepriete peet merket europillence menitering (ACDE http://www.up2t)
13	appropriate post-market surveillance <u>monitoring</u> (ACRE http://www.wp3_) .
13 14	Further details of such a system are given in (ACRE http://www.wp1*)
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13 14 15	Acknowledgements. The authors are grateful to Les Firbank, University
13 14 15 16	appropriate post-market surveinance-monitoring (ACRE http://www.wp3.). Further details of such a system are given in (ACRE http://www.wp1.) Acknowledgements. The authors are grateful to Les Firbank, University of Leeds and Keith Lindsey, Durham University, who were lead authors
13 14 15 16 17	Further details of such a system are given in (ACRE http://www.wp1*) Acknowledgements. The authors are grateful to Les Firbank, University of Leeds and Keith Lindsey, Durham University, who were lead authors for two of the ACRE documents that formed the basis for the article; to
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 13 14 15 16 17 18 19 20 21 22 23 24 	appropriate post-market surveinance intermediate intermediate intermediate surveinance intermediate interm

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1 References

- 2 1 James, C. (2012) Global Status of Commercialized Biotech/GM Crops:
- 2012. ISAAA Brief No. 44. ISAAA: Ithaca, NY. 3
- 2 Wolt J.D., Keese P., Raybould A.F., Fitzpatrick J.W., Burachick M., Gray A.J., Schiemann J., Sears M., Wu F. (2010). Problem formulation in the 4
- 5
- environmental risk assessment of genetically modified plants. Transgenic 6 Research 19: 425 - 436. 7
- 3 European Commission (2011) EU Commission's New Techniques Working 8
- 9 Group Final Report.
- 4 Bolle C et al. (2013) GABI-DUPLO: a collection of double mutants to 10
- overcome genetic redundancy in Arabidopsis thaliana. Plant J. 75, 157-171. 11
- 12 5 Patterson AH et al. (2010) Insights from the comparison of plant genome
- 13 sequences. Ann. Rev. Plant Biol. 61, 349-372.