

Genome editing: scientific opportunities, public interests and policy options in the EU

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What is genome editing?

- The deliberate alteration of a selected DNA sequence in a cell, in particular using site-specific nucleases
- Vitally important tool in basic research to help understand biological functions and disease mechanisms
- Opportunities enhanced by CRISPR-Cas9, which is relatively easy to design, produce and use
- Expected wide range of innovative applications in human and animal health, agriculture and food systems, microbial biotechnology and the bioeconomy



Genome editing evokes enthusiasm but also controversy

- Safety, ethical and other issues have been raised. For example:
 - "Not natural"
 - "Too many gaps in knowledge"
 - "Impacts are uncertain and may be inequitable"
 - "Regulation cannot keep pace with speed of technology innovation"
- What needs to be done to realise the potential of genome editing and take account of societal concerns?



EASAC Working Group

- Working Group started in early 2016, consensus output peer reviewed in early 2017, endorsed by member academies and published 23 March
- Our purpose:
 - To take a broad perspective on research advances and potential applicability to innovation in different sectors
 - To advise on options to ensure appropriate frameworks for managing innovation
- It is our view that policy considerations should primarily concentrate on sector-specific product regulation and not on the general principles of genome editing as a technology



How should the applications of genome editing be assessed and regulated?

- We emphasise the importance of ensuring that regulation is evidencebased, takes into account likely benefits as well as hypothetical risks, is proportionate and sufficiently flexible to cope with future advances in science
- Our report covers applications in
 - Plant breeding
 - Animal breeding in agriculture
 - Xenotransplantation
 - Gene drive to modify populations in the wild
 - Micro-organisms
 - Human cell genome editing building also on the work of FEAM



Plant breeding in agriculture

- Genome editing brings additional precision in research to increase crop quality, e.g. for improved water and nitrogen use efficiency, better resistance to pests and diseases, reduced allergens
- Novel crops are now appearing, e.g. CRISPR-Cas9-edited mushrooms
- Should such products be considered as GMOs?
 - We ask that regulators confirm that if crops do not contain DNA from an unrelated organism, they do not fall within the scope of GMO legislation
 - The aim should be to regulate the trait/product not the technology



Animal research and development

- Already subject to stringent regulation of laboratory animal research and to the principles of the 3Rs (replacement, reduction, refinement)
- Significant opportunities for genome editing in livestock and aquaculture

 to enhance animal health and welfare as well as increase agricultural
 production, e.g. protections from porcine reproductive and respiratory
 syndrome and African swine fever
- We recommend that animal breeding in agriculture should be governed by the same principles as proposed for crop breeding:
 - To regulate the trait not the technology
 - Be open and explicit about what is being done



Xenotransplantation

- Genome editing is transforming prospects by tackling barriers to transfer of tissues and organs from animals to treat loss or dysfunction in patients – removing endogenous retroviruses and xenoreactive epitopes that trigger rejection
- Also research advances in human-animal (pig) chimeras, e.g. to produce pancreas for transplantation
- Scientific community has important role to assist regulators to prepare for the potential opportunities now coming within range



Gene drive to modify populations in the wild

- Process of biased inheritance that enables genes to be transmitted from parent to offspring at increased rate
- Opportunities for insect-vector disease control to address public health challenges e.g. malaria, Zika, dengue fever
- "Self-sustaining" efficacy and safety questions including ecological consequences and safeguards to manage risk of escape during research
- EASAC supports recommendations by US NAS for a phased approach to research to enable responsible development and allow sufficient time to consider what amendments are needed to regulatory frameworks



Micro-organisms and the bioeconomy

- Genome editing in microbes has not raised new issues for regulatory frameworks
- Wide range of potential applications pharmaceuticals and other highvalue chemicals, biofuels, biosensors, bioremediation, and the food chain
- Concerns raised elsewhere about potential for genome editing research to expand outside regulated laboratory settings. We recommended that Global Young Academy should assess issues raised by activities of Do-It-Yourself biology community
- Concerns also raised elsewhere about potential biosecurity implications of misuse (microbes and other species). EASAC is organising a workshop later in 2017 with NAS, IAP, Leopoldina Academy and Volkswagen Foundation



Human cell genome editing: EASAC recommendations

- *Basic research:* intensive research is needed and should proceed subject to appropriate rules and standardised practices
- *Clinical use in somatic gene editing:* should proceed within existing regulatory frameworks (EMA and national agencies)
- *Clinical use in germline interventions:* not acceptable to proceed unless and until scientific, ethical, safety and efficacy issues have been resolved and there is broad societal consensus
- EASAC work draws on previous work by FEAM and International Summit on Gene Editing. Issues will be discussed in greater detail by Professor Lovell-Badge



In conclusion

- In addition to sector-specific elements for assessment and regulation, there are cross-cutting issues for public engagement and for enhancing equity of access to benefits
- EASAC report is on <u>http://www.easac.eu/home/reports-and-</u> <u>statements/detail-view/article/genome-editi.html</u>
- eLife Feature article is on <u>http://www.easac.eu/home/journal-publications/detail-view/article/elife-1.html</u>
- Thank you to EASAC Working Group, our peer reviewers, colleagues in FEAM, Professors Lovell-Badge, Schiemann, Rima and Tanner and Dr Kessler
- We welcome wide-ranging discussion today and on subsequent follow-up