



MEMORANDUM

Application	APP204075
Applicant	The Malaghan Institute for Medical Research
From	Dr Tim Strabala, Acting Manager and Principal Scientist, New Organisms
To	Dr Clark Ehlers, Acting General Manager, Hazardous Substances and New Organisms
Purpose of the Memorandum	Section 42A Pathway assessment for APP204075
Date of Advice	24 July 2020

Purpose

1. This memo provides my pathway assessment and recommendation for your consideration of application APP204075 to develop genetically modify human cell lines that package 3rd generation self-inactivating lentiviral vector particles, and to generate genetically modified human T cells expressing genes that regulate the activity of human immune cells, using the aforementioned lentiviral vectors.
2. At this stage, no application has been formally submitted, pending your decision on the application pathway.

Background

3. Application APP204075, from the Malaghan Institute of Medical research (the applicant), intends to seek approval to develop genetically modify human cell lines that package 3rd generation self-inactivating lentiviral vector particles, and to generate genetically modified human T cells expressing genes that regulate the activity of human immune cells, using the aforementioned lentiviral vectors. It is intended that the application will be submitted under sections 40(1) and 42A of the Hazardous Substances and New Organisms (HSNO) Act 1996 ('the HSNO Act').

Statutory criteria for pathway assessment – s42A

4. Under section 42A of the HSNO Act, an application made under section 40(1) to develop a new organism may, instead of specifying the information required by or under section 40(2), describe:
 - (a) a project for the development of genetically modified organisms; and
 - (b) the identity of the host organisms; and
 - (c) the nature and range of the proposed genetic modifications.
5. If an application provides all of the above information, the EPA may decide to undertake a rapid assessment of the adverse effects of carrying out the project if it is satisfied that:
 - (a) any host organism specified for the project meets the criteria for host organisms prescribed in the Hazardous Substances and New Organisms (Low-Risk Genetic Modification) Regulations 2003 (Low-Risk Regulations) (i.e. are either “**category 1 host organisms**” or “**category 2 host organisms**”); and
 - (b) any genetic modification specified for the project meets the criteria for genetic modification procedures prescribed in the Low-Risk Regulations (i.e. are either “**category A genetic modifications**” or “**category B genetic modifications**”).
6. The decision to undertake a rapid assessment under section 42A of the HSNO Act is currently sub-delegated to the Manager, New Organisms, by the Chief Executive¹ under the instrument of delegation effective as of 6 March 2020. However, as I am currently the Acting Manager, New Organisms, but I am conducting this assessment under my usual role of Principal Scientist, New Organisms, the decision must be undertaken by the General Manager of Hazardous Substances and New Organisms.

Assessment of the application against statutory criteria

7. I have assessed the application against the statutory criteria and my findings are set out in the table below.

Checklist	Yes	No	Details
<u>Information Requirements (s 42A(1))</u>			
Does the application describe a project for the development of genetically modified organisms?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<i>To develop genetically modify human cell lines that package 3rd generation self-inactivating lentiviral vector particles, and to generate genetically modified human T cells expressing</i>

¹ Per the sub-delegation dated 6 March 2020 from the Chief Executive to the Manager, New Organisms, under delegated authority dated 6 May 2016 from the EPA to the Chief Executive pursuant to section 19 of the Act.

Checklist	Yes	No	Details
			<i>genes that regulate the activity of human immune cells, using the aforementioned lentiviral vectors.</i>
Does the application describe the identity of the host organisms?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<i>Human (Homo sapiens) cell lines(See Schedule)</i>
Does the application describe the nature and range of the proposed genetic modifications?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<i>Packaging of 3rd generation self-inactivating lentiviral vector particles in human cell lines, and generation of genetically modified human T cells expressing genes that regulate the activity of human immune cells, using the aforementioned lentiviral vectors.</i>
<u>Host Organisms</u> (s 42A(2)(a) & regs 6 & 7)			
Are the host organisms clearly identifiable and classifiable according to genus, species, and strain or other sub-specific category as appropriate?	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<i>Human (Homo sapiens) cell lines (See Schedule)</i>
Category 1 host organisms			
Do the host organisms satisfy <u>all</u> of the following:			
(a) are not normally able to (or contain infectious agents normally able to) cause disease in humans, animals, plants, or fungi;	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
(b) do not produce desiccation-resistant structures, such as spores or cysts, that can normally be disseminated in the air;	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
(c) are characterised to the extent that their main biological characteristics are known; and	<input checked="" type="checkbox"/>	<input type="checkbox"/>	
(d) do not normally infect, colonise, or establish in humans.	<input checked="" type="checkbox"/>	<input type="checkbox"/>	

Checklist	Yes	No	Details
Category 2 host organisms			
Are the host organisms <u>either</u> :			
(a) micro-organisms of risk group 1 ² or risk group 2 ³ that <u>either</u> :			
(i) are or contain an infectious agent pathogenic to humans, animals, plants, or fungi; or	<input type="checkbox"/>	<input type="checkbox"/>	
(ii) produce desiccation-resistant structures, such as spores or cysts, that may normally be disseminated in the air; or	<input type="checkbox"/>	<input type="checkbox"/>	
(iii) are not characterised to the extent that its main biological characteristics are known; or	<input type="checkbox"/>	<input type="checkbox"/>	
(iv) normally infect, colonise, or establish in humans;	<input type="checkbox"/>	<input type="checkbox"/>	
(b) a mammalian cell line containing active viruses or infectious agents normally able to cause disease in humans;	<input type="checkbox"/>	<input type="checkbox"/>	
(c) a whole animal, vertebrate or invertebrate, including oocytes, zygotes, early embryos, and other cells able to grow without human intervention into a whole animal; or	<input type="checkbox"/>	<input type="checkbox"/>	
(d) a whole plant <u>either</u> :			
(i) with a reproductive structure and that is not kept in a closed container; or	<input type="checkbox"/>	<input type="checkbox"/>	
(ii) with a reproductive structure and that is kept in a closed container; or	<input type="checkbox"/>	<input type="checkbox"/>	
(iii) without a reproductive structure and that is not kept in a closed container.	<input type="checkbox"/>	<input type="checkbox"/>	

² Risk group 1 means micro-organisms that are unlikely to cause diseases in humans, animals, plants, or fungi.

³ Risk group 2 means micro-organism that: (a) may cause disease in humans, animals, plants, or fungi but are unlikely to be a serious hazard to laboratory personnel, the community, animals, or the environment; and (b) have effective treatment and preventive measures with respect to any infections that they may cause; and (c) present a limited risk of the spread of infection.

Checklist	Yes	No	Details
<u>Proposed genetic modifications (s42A(2(b) & regs 4 & 5)</u>			
Do the proposed genetic modifications involve any of the developments specified in the Schedule to the Low-Risk Regulations (being developments that are considered not to be low-risk modifications)?	<input type="checkbox"/>	<input checked="" type="checkbox"/>	
Category A genetic modification			
Do the proposed genetic modifications satisfy <u>all</u> of the following:			
(a) involve a category 1 host organism;	<input type="checkbox"/>	<input type="checkbox"/>	
(b) carried out under a minimum of PC1 containment ⁴ ;	<input type="checkbox"/>	<input type="checkbox"/>	
(c) do not increase the pathogenicity, virulence, or infectivity of the host organism to laboratory personnel, the community, or the environment; and	<input type="checkbox"/>	<input type="checkbox"/>	
(d) do not result in the genetically modified organism having a greater ability to escape from containment than the unmodified host organism.	<input type="checkbox"/>	<input type="checkbox"/>	

⁴ PC1 containment means (a) the conditions for the physical containment of organisms described as Physical Containment Level 1 (PC1) in AS/NZ containment standard 2243.3:2002 (Safety in Laboratories Part 3: Microbiological Aspects and Containment Facilities); and (b) the modifications referred to in the following MAF Biosecurity Authority containment standards: (i) 154.03.02 (31 October 2002) (containment facilities for micro-organisms); (ii) 154.03.03 (31 October 2002) (containment facilities for vertebrate laboratory animals); (iii) 154.02.08 (31 October 2002) (transitional and containment facilities for invertebrates); (iv) 155.04.09 (24 March 2003) (containment facilities for new organisms, including genetically modified organisms, of plant species).

Category B genetic modification

Do the proposed genetic modifications satisfy the following:

- | | | |
|--|-------------------------------------|-------------------------------------|
| (a) carried out under a minimum of PC2 containment ⁵ ; <u>and either</u> | <input checked="" type="checkbox"/> | <input type="checkbox"/> |
| (b) if a category 1 host organism is used <u>both</u> of the following are satisfied: | | |
| (i) the nucleic acid that is introduced is be characterised to the extent that <u>either</u> : | | |
| (A) its sequence is known; or | <input checked="" type="checkbox"/> | <input type="checkbox"/> |
| (B) its gene function is understood; and | <input checked="" type="checkbox"/> | <input type="checkbox"/> |
| (ii) the modification <u>does not result in either</u> of the following: | | |
| (A) a genetically modified organism that is more pathogenic, virulent, or infectious to laboratory personnel, the community, or the environment than a category 2 host organism; and | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| (B) the genetically modified organism having a greater ability to escape from containment than the unmodified host organism. | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| (c) if a category 2 host organism is used <u>both</u> of the following criteria are met: | | |
| (i) the modification involves <u>either</u> : | | |
| (A) a host organism that is not normally able to cause disease in humans, animals, plants, or fungi; or | <input type="checkbox"/> | <input type="checkbox"/> |
| (B) a host organism that is normally able to cause disease in humans, animals, plants, or fungi provided that the nucleic | <input type="checkbox"/> | <input type="checkbox"/> |

⁵ PC2 containment means (a) the conditions for the physical containment of organisms described as Physical Containment Level 2 (PC2) in AS/NZ containment standard 2243.3:2002 (Safety in Laboratories Part 3: Microbiological Aspects and Containment Facilities); and (b) the modifications referred to in the following MAF Biosecurity Authority containment standards: (i) 154.03.02 (31 October 2002) (containment facilities for micro-organisms); (ii) 154.03.03 (31 October 2002) (containment facilities for vertebrate laboratory animals); (iii) 154.02.08 (31 October 2002) (transitional and containment facilities for invertebrates); (iv) 155.04.09 (24 March 2003) (containment facilities for new organisms, including genetically modified organisms, of plant species).

acid that is introduced is characterised to the extent that satisfies <u>all</u> of the following:		
1. its sequence is known;	<input type="checkbox"/>	<input type="checkbox"/>
2. its gene function is known; and	<input type="checkbox"/>	<input type="checkbox"/>
3. its potential gene products are understood; and	<input type="checkbox"/>	<input type="checkbox"/>
(ii) the modification <u>does not either</u> :		
(C) increase the pathogenicity, virulence, or infectivity of the host organism to laboratory personnel, the community, or the environment; and	<input type="checkbox"/>	<input type="checkbox"/>
(D) result in the genetically modified organism having a greater ability to escape from containment than the unmodified host organism.	<input type="checkbox"/>	<input type="checkbox"/>

8. In summary, I have concluded that the information requirements in section 42A(1) are satisfied and the host organisms and proposed genetic modifications conform to the requirements for host organism and genetic modification in the Low-Risk Regulations as required by section 42A(2).

Other Considerations

9. The decision to carry out a rapid assessment under section 42A is a discretionary decision. Therefore, you are not obliged to carry out a rapid assessment just because the criteria in section 42A are met. In deciding whether to carry out a rapid assessment of the application, you should be aware of the following:

- The applicant already holds an approval for the development of lentiviral vectors and the development of genetically modified T cells, which is limited in its purpose solely to CD8+ T cells. The current application broadens the approved T cells beyond the CD8+ T cells allowed under their current approval to all human T cells.

Recommendation

10. I have assessed the application against the statutory criteria and consider that it meets the requirements set out in section 42A of the Act, and the Low-Risk Regulations.

11. I therefore recommend that a rapid assessment of this application is undertaken in accordance with section 42A of the HSNO Act.



24 July 2020

Dr Tim Strabala

Date

Acting Manager and Principal Scientist, New Organisms

Decision

Rapid assessment criteria

I agree that application APP204075 meets the requirements of section 42A of the HSNO Act.

I do not agree that application APP204075 meets the requirements of section 42A of the HSNO Act.

Decision whether to undertake rapid assessment

I have decided that application APP204075 should be subject to a rapid assessment of the adverse effects of carrying out this project under section 42A of the HSNO Act.

I have decided that application APP204075 should not be subject to a rapid assessment of the adverse effects of carrying out this project under section 42A of the HSNO Act.



Dr Clark Ehlers

Date 27/07/2020

General Manager (Acting), Hazardous Substances and New Organisms

Schedule: List of organisms, risk categorisations and genetic modifications proposed for development under APP204075

Organism	Taxonomic name	Organism and genetic modification risk categorisations
Human viral vector packaging cell lines and human cancer patient-derived primary T cell lines	<i>Homo sapiens</i>	Category 1 host organism Category B genetic modification