PROPOSAL FORM FOR ASSESSMENT OF GENETIC MANIPULATION WORK

		GMAC Ref No.: (For official use only)		
Name	of Principal Investigator:			
Name	of Institution :			
Exper	Experiment Risk Group (please check the appropriate box):			
	Category A Category B	Category C		
Α.	A. Experimental detail (attach separate sheet if necessary)			
1.	Project title (Please provide reference number title.)	ers for projects with the same		
2.	Research unit involved			
3.	Experimental objective			
4.	Rationale for the experiment			

5. Scope of experiment – involvement of			
		Microorganisms and/or viruses	
		Toxins	
		Animals	
		Cells	
		Others, please specify:	
	Note	e: For experiments that involve animals, cells and/or others, <u>please</u> <u>proceed to 6 and the rest</u> , but <u>skip 7 & 8</u> . For experiments that involve microorganisms, viruses and/or toxins, <u>please skip 6</u> and proceed to the rest of the form. For experiments that involve multiple experimental organisms (i.e. animals/cells together with microorganisms/viruses/toxins), please proceed to fill up <u>all</u> relevant questions. For experiments that involve HIV-based lentiviral vectors, please <u>also</u> fill up the table in <u>Appendix I</u> .	
6.	6. Project with experiment involving animals/cells/others		
a.	Can t	he modification result in a predictable change of the following:	
	i) Increased oncogenicity: □Yes □No		
	ii) Potential to change natural microbiome/ecology of the organism: □Yes □No		
	Deta	ails, if "yes":	
b.		cription of gene(s) involved, gene construct(s) and intended erimental host system.	
C.		hod of gene delivery (bacteriophage, vectors, breeding, injection, ogical delivery vehicle/carrier etc.)	

7. Project with experiment involving microorganisms/viruses/toxins			
a.	Nam	e of microorganism/virus/toxin:	
b.	Age i)	e microorganism/virus/toxin listed under the BATA List of Biological ents and Toxins, and/or a potential human pathogen? If Yes, provide the BATA Schedule: If No, provide the risk grouping (for biological agent):	
C.	Brief	description of gene modification on the microorganism/virus:	
	i)	Gene(s) involved and gene construct(s) and intended experimental host system (if chimeric microorganism is created, please specify the backbone and the inserted genes):	
	ii)	Natural host of microorganism/virus:	
	iii)	Method of gene delivery (transformation, conjugation, vectors, breeding, injection, biological delivery vehicle/carrier etc. For retroviral vectors, please specify the (viral) origin, and also note all safety features included in the constructs. For HIV-based lentiviral vectors, please specify the BATA Schedule of HIV lentiviral vectors):	

8. For dual use research of concern (DURC) (This section is to screen for DURC relating to research work involving GMOs-microorganisms/viruses/toxins. Kindly fill up the following questions to the best of your abilities.)			
Is there a reasonable possibility that the modification might result in a change of the following?			
(If unsure, choose "Yes" and explain under Details).			
i) Increase in host range: Yes			
Details, if "yes" to any of the above: (e.g. if the resultant product has increased drug resistance, please provide info on the extent of the resistance, and if there is still effective drug or treatment for infected individuals)			
9. Duration of the experiment			
□1 year □2 years □3 years			

10.	The proposed work will be performed in the following biocontainment level:		
	□BSL1/ABSL1 □BSL2/ABSL2 □BSL2+/ABSL2+ □BSL3/ABSL3		
11.	Measures to ensure containment, safe handling, storage and disposal		
12. For organism/microorganism			
a.	Experimental GMO Material to be obtained from (please note requirements for import permit for importation of biological agents and/or toxins regulated under the BATA):		
b.	a. Anticipated date of transfer or receipt:		
13.	Pl's declaration:		
	I declare that the above information is accurate and complete based on risk assessment, and to the best of my knowledge.		
	I recognise that the actual risk may defer from the assessed risk. I will continue to monitor the risk of the project and the genetically modified organism. Should the risk assessment change with respect to 8 and/or 10, I will stop work immediately and notify the IBC, which will then notify MOH and GMAC.		

Submitted by PI:		
PI's Name and Signature	Appointment / Laboratory	Date
Contact Details		
Address :		
Business Tel Number :	Fax Numb	
Business Email :		
Reviewed by IBC:		
	I Signature pplicable for Category A experim	
	has been sought from relevant reg se/possession/import/transport of th	-
If yes, please provide copy	of document or reference.	
If no, please explain why.		
*(highlight where applicable)		
For a list of regulatory contact points, please refer to Section 6.6 and 6.7 (page 26-28).		

Appendix I

Please complete the table below to assess if the HIV-based lentiviral vector system (HIV LVS) used suffices the revised criteria of HIV LVS under the BATA Fourth Schedule.

A HIV LVS that possesses <u>at least 2 of the following features</u> is classified under the <u>Fourth Schedule of the BATA</u>:

- **I.** the U3 region of the 3'LTR in the transfer vector is absent or altered, which results in a stable self-inactivating (SIN) configuration;
- II. the HIV genes for packaging function are split to a minimum of 2 packaging plasmids (excluding the env plasmid);
- III. the vpr, vpu, vif and nef genes are either absent or altered to be non-functional; and
- IV. the vector system requires minimally 4-recombination to achieve Replication Competent Lentivirus (RCL)

S/N	HIV-based Lentiviral Vector System (include vector names)	Features incorporated in vector system	Supporting Documents	Suffice HIV LVS criteria regulated under BATA Fourth Schedule
1	Example- XXX Vector System Transfer vector – ABC plasmid Packaging vector 1 – 123 plasmid Envelope vector – XYZ plasmid	Feature III	Attachments	Υ
2	Example- BBB Vector System Transfer vector – XYZ plasmid Packaging vector 1 – 456 plasmid Envelope vector – RST plasmid	Feature I only	NA	N