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Biotechnology - Microorganisms - Report on the criteria used to classify Group I genetically modified microorganisms

Biotechnik - Mikroorganismen - Bericht über gebräuchliche Einstufungskriterien bei gentechnisch veränderten Mikroorganismen der Gruppe FVFFW

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Biotechnologie - Microorganismes - Rapport sur les criteres utilisés pour classer les microorganismes génétiquement modifiés du Groupe I

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Biotechnologie - Microorganismes - Rapport sur les critères utilisés pour classer les microorganismes génétiquement modifiés du Groupe I

This CEN Report was approved by CEN on 14 October 1998. It has been drawn up by the Technical Committee CEN/TC 233.

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EUROPEAN COMMITTEE FOR STANDARDIZATION COMITÉ EUROPÉEN DE NORMALISATION EUROPÄISCHES KOMITEE FÜR NORMUNG

Central Secretariat: rue de Stassart, 36 B-1050 Brussels

Page 2 CR 13426:1998

Contents

Foreword	3
Introduction	
1 Scope	
2 Safety criteria for Group I GMM's	
3 Conclusions and recommendations	
Annex A (informative) List of national competent authorities	
Annex B (informative) Bibliography	

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SIST CR 13426:1999

Page 3 CR 13426:1998

Foreword

This CEN Report has been prepared by the Technical Committee CEN/TC 233 "Biotechnology", the secretariat of which is held by AFNOR.

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Introduction

From the beginning of the application of genetic modification of microorganisms there has been concern among users as well as authorities about the potential dangers of this new technique, especially with respect to the creation of new strains that could be hazardous to man, animals, plants or the environment in general. Many national authorities, advisory committees and also the European Commission, set up a legal framework to ensure the safe application of techniques of genetic modification. In order to be able to determine what, if any, safety precautions should be taken in individual biotechnological operations, safety criteria were set up to supplement the legislation. The NIH (see annex B [1]) and the OECD (see annex B [2]) were among the first to publish widely accepted safety criteria. Both publications have been regularly updated. They have had an important harmonizing impact and also strongly influenced the European Directive 90/219/EEC on contained use of genetically modified microorganisms (GMM's), issued in 1990 (see annex B [3]). This Directive aims at regulating the use of GMM's in such a way that no untoward effects are to be expected for man and environment.

Many microorganisms currently in use in modern biotechnology, whether it be research, development or production, have a long record of safe use. They have been employed for centuries in classical biotechnology without creating untoward effects to the environment and they have not been seen to be hazardous to humans. Therefore, in giving safety guidelines on how to use GMM's in general making a distinction between safe and potentially unsafe organisms is logical.

General safety criteria for hosts, vectors and inserts that can be used to construct Group I GMM's and criteria for such GMM's themselves are given in an annex of the Directive. These criteria were somewhat detailed originally but were amended into more general terms later (see annex B [4]).

At present Directive 90/219/EEC is being revised in which the distinction of the two groups of microorganisms is abandoned. However, assessment of the risks that GMM's may pose to man and environment will stay obligatory and, as a consequence thereof, the need for biological safety criteria for microorganisms.

Given the rather general quality of the Group I safety criteria, guidelines on their interpretation were issued (see annex B [5], [6]). These leave room, however, for different decisions on whether hosts, vectors, inserts and GMM's qualify for Group I or not. As a consequence, and also because European Union Member States differ in the kind of activities with GMM's that are carried out, national interpretations of the safety guidelines and the accompanying lists of Group I hosts and vectors have diverged to some extent.

1 Scope

This CEN Report reviews the safety criteria for hosts, vectors, inserts and genetically modified microorganisms (GMM's) used by the competent authorities of Member States of the European Union to classify GMM's as intrinsically safe, i.e. as Group I GMM's. The report

Page 5 CR 13426:1998

may assist both users and regulatory authorities in classifying GMM's, including mammalian and plant cells, as safe for human health and the environment.

This CEN Report covers Group I GMM's that are applied in research and development activities and those applied in industrial applications, which are frequently activities on a larger scale. Although most of the information available relates to GMM's that are to be used for research and development, the safety criteria for GMM's to be used for the latter kind of activities show much overlap. No listings of hosts and vectors are given, because such lists are readily available and are inevitably incomplete due to the continuous addition of newly classified biological agents.

2 Safety criteria for Group I GMM's

The safety criteria used by authoritative bodies of several European countries to classify hosts, vectors, inserts and GMM's as Group I are summarized in table I. The table relates them to the general safety criteria given in Directive 90/219/EEC and the most recent official guidelines thereof (see annex B [6]). More detailed information on the national criteria can be found in the publications of the respective authoritative bodies (see annex A).

Several national authoritative bodies also publish lists of Group I hosts and vectors or host-vector systems classified as such. These lists however are rather incomplete because of infrequent updates and, of far more importance, because they do contain only those microorganisms of which a notifier requested official recognition. In many cases such a request is not made because of reasons of confidentiality, which means that the hosts and vectors are only mentioned in the notification, but not incorporated in the lists of Group I microorganisms.

SIST CR 13426:1999

Not all countries where activities with Group PGMM's occur do have predefined specific national safety criteria available. b06d69923fa5/sist-cr-13426-1999

In Sweden for several Group I hosts and vectors used in industrial applications simply the relevant criteria of the NIH Guidelines were used (e.g. for hosts like K12 strains of *Escherichia coli* and Chinese Hamster Ovary cells and vectors like pBR322-derivatives, pKGE327 and pKGE439) (see annex B [14]).

In Belgium the still rather general guidelines of Directive 96/134/EEC are an integral part of the national legislation with some minor modifications. Although their scope is broadened to encompass animals and plants as well, no additional specifications are available (see annex B [7]).

Also in Norway these guidelines, as well as those on the general safety criteria given in the original Directive are the only official criteria available (see annex B [15]).

For several countries, notably Germany, it is difficult to represent the national safety criteria used in table I, because their legislation gives comprehensive safety criteria for microorganisms, but not specifically for Group I. Points that need to be considered when carrying out the risk assessment to decide on the relevant containment level are given instead, sometimes differentiated into risk assessment made for research applications and industrial applications. A guide to such risk assessment including typical examples has been issued by the German Occupational Safety and Benefit Organization of Chemical Industry (see annex B [8]), while in the United-Kingdom for large scale applications at the lowest containment level several criteria are available (see annex B [10]).

²age 6 3R 13426:1998

Table 1: Overview of safety criteria for Group I genetically modified microorganisms

Table 1.a: Host and parental organism

SAFETY CRITERIA							CEN	MEME	CEN MEMBER STATES	TATE	S						
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LEVEL 1: CRITERIA ACCORDING TO DIRECTIVE 90/219/EEC		ps://st		iΊ													
Not likely to cause disease in humans, animals and plants.		tan		<u>'e</u>													
EVEL2: CRITERIA ACCORDING TO DIRECTIVE 96/134 (GUIDELINES)		dar		<u>eh</u>													
A non virulent strains of a pathogenic species can be considered to fulfill this criterion if:	·	ds.i	(S										·····			-
 it has a long history of safe use without having detrimental effects on humans, animals or plants; and/or 	DU		SI										_				
it contains only a small quantity of genetic material that encodes for virulence; or	Joak	ni/ca	[a]	A						_							
it does contain mutations that diminish virulence to an acceptable level.	595	SI tal	n	N													
EVEL 3 : CRITERIA ACCORDING TO MEMBER STATES	9231	ST og/st 923 f	da														
Not pathogenic (to humans, animals and plants).	15/	CR and	9	X	NE NE	-		_				z	Ž			,	¥
No potentially harmful associated biological agents.	SISI	lar en	e	X T	E NE	Ø						z	į			MS	5 '
থ০ self mobilisable plasmids present.	[-C]	3 <u>42</u> ds/	S	R	•								Z				Ţ.
.ow horizontal gene transfer with biological agents associated with animals or plants.	-1	16: Sis	.1	D	-	ŋ						Ŀ					Ţ.
n case of a non virulent strain of a pathogenic species: irreversibly deficient in genes	342	199 1/79 3 4 2	te		<u>'</u>	'		_				ŀ	'				
o case of a non virulant strain of a nothononia condian on little of acceptant	0-	- 7b	n) 													
letermining genes to avoid reversion to pathogenicity.	199	91e		R	-	1		t				z —	•				,
n case of use of viral vectors: no helper functions that will generate the production of	y	_	P	£ `	<u>'</u>	-		ŀ					Z				\
turonomous replicating infectious virus.		380	1	V	_												
i case of raige scale applications : having a long history of safe use under GII SP conditions or lower : or if not		0-4		L	Ž.			_					-				
having genomic traits that are harmless to the environment and that do diminish the		c08	<u> </u>	<u> </u>	Ē							z z	ZZ			1 1	
chances of survival or reproduction of the host in the environment (biological		-84.		V													
n case of large scale applications: no risk to himan health and the environment	+	39-	2		Į.	+	_	-					-				T
n case of large scale applications :			2	' -		5	_	-					,			-	
experimentally shown safe use;			•			g						z				N.O.	
long history of safe use;		_	품	<u>~</u>	H	Ø		•				z	,			;	
 built in biological containment, e.g. diminishing chance of survival or replication, without negative effects on the environment. 					ı	ڻ ص						z	ı				<u> </u>
n case of large scale applications : Not containing known adventitious agents.			'	'		5						,					1
						,]						•	5

Table 1.b : Vector

SAFETY CRITERIA							ပ	CEN MEMBER STATES	MBEF	STA	TES							
•	А	В	£	X	ш	N N	g	GR	_	ICF I	RE		z	Ĭ Z	Д.	S	SW L	¥
_EVEL 1: CRITERIA ACCORDING TO DIRECTIVE 90/219/EEC																		
Not endowing the GMM with phenotype likely to cause disease to humans, animals and plants or adverse effects in the environment.																		
EVEL 2: CRITERIA ACCORDING TO DIRECTIVE 96/134 (GUIDELINES)		htt									·							
No genes present that encode for an active protein (e.g. virulence determinants, toxins) at a		ps://:			i										-			
Vot self transferable or mobilisable in case of presence steep the determine fertimental traits in certain micro-oranisms, but that do not endow the GMM with the		standa			 Гel												 -	
shenotype aforementioned.		rds			1		-								_		_	
n case of large scale applications: not self transferable.		.itel		(5	S													
n case of large scale applications : not consisting or transferance reference controlled.		i.ai/ b06		sta	<u>Γ</u>								-					
n case of large scale applications : sufficient genomic characterization or a genome limited		cata d69	<u>,</u>	ar	4]						_							
o those gene sequences required for its intended function, in case it seems probable that		alog 992:	SIS	ıd	N]			•							 -			
THE GAMINI WILL DEVELOP THE PLIESTING TO MEMBER STATES		/sta 3fa5	Г <u>С</u>	a	D											<u>.</u>		
EVEL 3: CHILEMIA ACCORDING TO MEMBER STATES		nda 5/si	R	r	A	+			-	\perp		+	\dashv	+	$\frac{1}{1}$		1	
Of known type (plasmid, cosmid, phage, minichromosome, viral vector).		ard ist-	134	d	j	+	, ;	+	_	+	+	+	z	<u> </u>	-			
Pestricted host-range.		ls/s ·cr	42	S.	8]		(2)		,	1				_				,
Well characterized.		ist 13	5:1	ž	Ð		5		_	-			,	-			_	乡
Sufficient genomic characterization means knowledge of :		797 426	999	e	P									-				
size of vector;		7 b9 5-1)	Ž	F					-								
, origin of each fragment defived from a naturally occurring generic element, mothod of construction.		1e 8		ă	RI		,		,				 . z	뉟				_
function and relative position of the following genetic elements: a) structural genes, b)		3-2:		¥)	<u>.</u>				•					≓				
regulating sequences, c) synthetic sequences, d) transposons (provirus sequences included) and e) replicons relevant to the GMM.		380- -				•									-			
imited in size to relevant sequences as much as possible.		lc0		-	E'		G2)						z				>	놀
Vo presence of transposon or provirus sequences potentially relevant for transposition or negration in the host.		8-84.			W									J _N		·		
Vo presence of transposon or provirus sequences with harmful sequences, unless these are assential to the vector. In the latter case, under no circumstances pathogenic GMM's may		}9-		1	•		1								-	•		ı
No capability of autonomous transfer of genetic elements to other organisms.							g						z	Z	-	-	\vdash	Ţ.,
.ow level of co-transfer.		,		-	ı		G1)											
																00,	"continued"	"pe

Biologically safe strains only
 In case of industrial application