













« GENERAL INFORMATION » FIELDS	FIELD MUST BE COMPLETED		 EXPORT	 EXPORT	ATTACHMENT 1 OR N FILE(S)	 DOC EXPORT	 DOC EXPORT
Line name	YES		YES	YES			
Secondary names	no		YES	YES			
Record date	no		no	YES			
Person	YES		no	YES			
Species	YES		YES	YES			
Ecotype	no		YES	YES			
Source	no		YES	YES	1 FILE	no	YES
MTA protection	no		YES	YES			
MTA details	no		no	YES	N FILES	no	YES
Article reference	no		YES	YES	1 FILE	no	YES
« GENETIC FEATURES » FIELDS	FIELD MUST BE COMPLETED		 EXPORT	 EXPORT	ATTACHMENT (1 or n)	 DOC EXPORT	 DOC EXPORT
Category	TRANS- GENESIS	ENDOGENOUS GENE MUTAG.					
Designation	YES	YES	YES	YES			
Short description	no	no	YES	YES			
Gene name	no	no	YES	YES			
Gene secondary names	no	no	YES	YES			
Method	no	no	YES	YES			
Method reference	no	no	YES	YES			
Transgene sequence	no	<i>n.a.</i>	YES	YES	1 FILE	YES	YES
Dose	<i>n.a.</i>	no	YES	YES			
Mutation sequence	<i>n.a.</i>	no	YES	YES	1 FILE	YES	YES
Sequencing realized by (+ Notebook ref)	no	no	no	YES			
Organism class	no	<i>n.a.</i>	YES	YES			
Pathogenicity targets	no	<i>n.a.</i>	YES	YES			
Selectable marker in plants	no	<i>n.a.</i>	YES	YES			
Strain	no	<i>n.a.</i>	YES	YES			
Genotyping protocol	no	no	YES	YES	1 FILE	YES	YES
Genotyping realized by (+ Notebook ref)	no	no	no	YES			

« PLANTS AND SEED BATCHES » FIELDS	FIELD MUST BE COMPLETED		 EXPORT	 EXPORT	ATTACHMENT 1 OR N FILE(S)	 DOC EXPORT	 DOC EXPORT
	PLANT	SEED BATCHES					
<b>TABLE</b>							
<b>ID</b>	<i>n.a.</i>	Automatic increment	no	YES			
<b>Identifier</b>	YES	YES	YES	YES			
<b>Generation</b>	no	no	YES	YES			
<b>Descent type (backcross/selfcross)</b>	<i>n.a.</i>	no	YES	YES			
<b>Mother plant</b>	<i>n.a.</i>	no	YES	YES			
<b>Comment</b>	no	no	YES	YES			
<b>Seed batches collected from a particular plant</b>	no	<i>n.a.</i>	YES	no			
<b>Phenotype</b>	no	no	YES	YES	N FILES	YES	YES
<b>Phenotyping realized by (+ Notebook ref)</b>	no	no	no	YES			
<b>Genotyping</b>	no	<i>n.a.</i>	YES	YES			
<b>Genotyping realized by (+ Notebook ref)</b>	no	<i>n.a.</i>	no	YES			
<b>Insertion loci sequences</b>	no	<i>n.a.</i>	YES	YES	1 FILE	YES	YES
<b>Insertion sequencing realized by (+ Notebook ref)</b>	no	<i>n.a.</i>	no	YES			
<b>Harvest date</b>	<i>n.a.</i>	no	YES	YES			
<b>Harvesting realized by (+ Notebook ref)</b>	<i>n.a.</i>	no	no	YES			
<b>Storage date</b>	<i>n.a.</i>	no	no	YES			
<b>Storage place and quantity</b>	<i>n.a.</i>	no	no	YES			
<b>Germination assay result</b>	<i>n.a.</i>	no	YES	YES			
<b>Germination assay date</b>	<i>n.a.</i>	no	YES	YES			
<b>Germination assay realized by (+ Notebook ref)</b>	<i>n.a.</i>	no	no	YES			
<b>Genetic feature segregation profile</b>	<i>n.a.</i>	no	YES	YES			
<b>Segregation profile realized by (+ Notebook ref)</b>	<i>n.a.</i>	no	no	YES			