

ANNEX IV

Model of Analytical Report

Note: please compile an annex form for each sample tested

Parameter to be reported:	Information provided:
Name and address of the test laboratory(*)	
Test report identification code(*)	<<000>>
Laboratory sample identification code(*)	<<000>>
Size of laboratory sample(*)	X kg
In case of sample division: Number and size of analytical samples	X analytical samples of Y g
Number and size of test portions analysed(*)	X test portions of Y mg
Total DNA amount analysed(*)	X ng/PCR
DNA sequence(s) tested for(*):	For each of the following provide reference to the method used <u>and</u> the average Ct number obtained Rice marker: 35S promoter: NOS terminator: CryIAb/CryIAc:
Other sequence(s) tested for:	Validation status: (e.g. inter-laboratory validated, in-house validated [please indicate according to which standard, guideline]) Description of DNA sequences detected (reference + target genes): Specificity of the method (screening, construct-specific or event-specific): Absolute Limit of Detection (copy number): Practical Limit of Detection (LOD related to the sample analysed), if determined:
Description of positive controls for target DNA, and reference materials(*)	Source and nature of the positive control and reference materials (e.g. plasmid, genomic DNA, CRM ...)
Information on the positive control(*)	Please indicate the amount (in ng DNA) of positive control analysed and the average Ct number obtained
Comments	

(*) Obligatory fields