<u>ANNEX</u>

Identi- fication number of the additive	Additive	Composition, chemical formula, description, analytical method	Species or category of animal	Maximum age	Minimum content	Maximum content	Other provisions	End of period of
					CFU/kg of fresh material		other provisions	authorisa- tion
Category	: technological additiv	es. Functional group: silage additives						
1k20721	Lactiplantibacillus plantarum LMG P- 21295	Additive composition Preparation of Lactiplantibacillus plantarum LMG P-21295 containing a minimum of 5 × 10 ¹⁰ CFU/g additive Solid form Characterisation of the active substance Viable cells of Lactiplantibacillus plantarum LMG P-21295 Analytical method ¹ Enumeration in the feed additive of Lactiplantibacillus plantarum LMG P-21295: - Spread plate method using MRS Agar (EN 15787) Identification of Lactiplantibacillus plantarum LMG P-21295:	All animal species	-		-	 In the directions for use of the additive and premixtures, the storage conditions shall be indicated. Minimum content of the additive when it is not used in combination with other micro- organisms as silage additives: 1x10⁸ CFU/kg fresh material. For users of the additive and premixtures, feed business operators shall establish operational procedures and organisational measures to address potential risks resulting from their use. Where those risks cannot be eliminated by such procedures and measures, the additive and premixtures shall be used with personal skin and breathing protective equipment. 	[10 years from the date of entry into force of this Regulation. To be completed by the OP]

¹ Details of the analytical methods are available at the following address of the Reference Laboratory: <u>https://joint-research-centre.ec.europa.eu/eurl-fa-eurl-fa-eurl-fa-eurl-fa-authorisation/eurl-fa-evaluation-reports_en.</u>

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					CFU/kg of fresh material		Other provisions	authorisa- tion					
Category:	Category: technological additives. Functional group: silage additives												
		 Pulsed Field Gel Electrophoresis (PFGE) - CEN/TS 17697 or DNA sequencing methods 											