

SUPPLEMENTARY TABLE S1. SEQUENCES OF OLIGONUCLEOTIDES USED FOR ANALYSIS
OF THE FULL INTEGRITY OF α MHCNEOPGKHYGRO^a

<i>Primer name</i>	<i>Position</i>	<i>Annealing temperature (°C)</i>	<i>Product size (bp)</i>	<i>Sequence (5' → 3')</i>
α MHC_neo sense	5157	55	419	cgg cac tct tag caa acc tc
α MHC-neo ras I	5576			ctc gct ctg cag ttc att ca
α MHC_neo sense	5157	55	717	cgg cac tct tag caa acc tc
α MHC-neo ras II	5874			gat gct ctt cgt cca gat ca
PGK-hygro sense	6669	55	665	tag cag ctt tgc tcc ttc g
PGK-hygro ras I	7334			act gac ggt gtc gtc cat c
PGK-hygro sense	6669	55	679	tag cag ctt tgc tcc ttc g
PGK-hygro ras II	7348			tcc atc aca gtt tgc cag tg

^aPCR program used together with primers listed in Supplementary Table S1: 94°C for 1 min; 35 cycles of 94°C for 1 min, 55°C for 1 min, and 72°C for 90 sec; 72°C for 10 min; and 4°C (endlessly).