

**Supplementary Table S1.** *Primers* used for amplification and isolation of repetitive sequences

Primer name	Sequence (5'-3')	Size (bp)	Ta (°C)	GC (%)	Fragment (bp)
OgreCL39Ctg378F	TAGTGAATGTCTTTGGG	23	62,9	47,8	548
	CCTACC			3	
OgreCL39Ctg378R	TGCTTCAGACACGAAG	23	61,1	43,4	
	ATCTTGA			8	284
SIRECL22Ctg385F	TGCAAGAAGAAAACCC	23	59,3	39,1	
	AAGTGAA			4	
SIRECL22Ctg385R	AGGACATGCATCTTTTG	23	61,1	43,4	698
	ACCATC			8	
TarCL121Ctg47F	GCTTGAGTCTCCTATTT	23	62,9	47,8	
	TGTGGC			3	498
TarCL121Ctg47R	CACCCCCACATTTTGAG	23	62,9	47,8	
	GTAATG			3	
ChromoCL32Ctg253F	AATCTCTGGTCCCAAA	23	62,9	47,8	512
	AGACTCC			3	
ChromoCL32Ctg253R	GGCATTACCTCGTAGTC	23	64,7	52,1	
	CTAAGG			7	583
AthilaCL9Ctg216F	CTTTTGGTGATCCGAGC	23	62,9	47,8	
	ATTCTC			3	
AthilaCL9Ctg216R	CTGTGCCAACTAAGAC	23	62,9	47,8	171
	GAATGAC			3	
TorkCL10Ctg699F	AAACGACCTGTCTAAT	23	61,1	43,4	
	CCTGCAT			8	171
TorkCL10Ctg699R	TTTCAGATCAGGTGGA	23	62,9	47,8	
	GACTTCG			3	
SatCL3Ctg1182F	CTAGATTGAAACCCGA	24	61,8	41,6	171
	AACAAGGT			7	
SatCL3Ctg1182R	ACTTTCAACCCCAATTA	23	61,1	43,4	
	GCCTGA			8	