

1011.188	ACAAGATCGTGTATCCTTGGCCGCT.....AAATGGAATGACCTTAACTGGGGTTCTAGCCACITTTCT	Hsap NM_000379 i03 Rnor NM_017154 i03 Ggal chr3_4307816_4308333	11 1 2 OK 11 1 2 OK 11 1 2 OK
	ACAAGATCGTGTATCCTTGGCCGCT.....CAATGAAAACACCTTAACTGTGGTTGTAGTCAATTTTCT		
	CAAAATCGTGTATCCTTGGAGTCT.....AAATGAAATGACCTTAACTAATCATTTAGCCAGCAATCT		
1011.190	TTTGGAAAGGTATCCTTGTTCAC.....GAGAAATGTGATCTTGACTGGAAAAATAGATGTGTCGTG	Hsap NM_000968 i03 Rnor NM_022510 i03 Ggal chr10_1716825_1717468	11 1 1 OK 11 1 1 OK 11 1 1 OK
	CITTTGAAAGGTATCCTTGTGGTG.....AAAGAAAAATCTTGAGAAAAGAAATCAGATGTGTCGTG		
	CITTTGAAATGTATCCTTGTATCT.....GAAAGGTTGTCTTAACTGTAGATACTAGATGTGTCGTG		
1011.193	AGTTTGTGGAATATCCTTGGGTTTC.....CCTTGGCTAGGCTCCTAAACTCAGGGACAGCTACAACAAG	Hsap NM_004573 i17 Rnor NM_053478 i17 Ggal chr5_2788180_2788814	11 2 1 OK 11 2 2 OK 10 0 0 OK
	AGTTTGTGGAATATCCTTGGGCTTT.....GCTTGGCTAGCTCCTATACCACAACAGCTATAAATAA		
	AGTTTGTGGAATATCCTTGGGAGT.....CCAGTTGGCTTCAATGAGCAATTATAGATACAACAAG		
1011.195	TGCCTCAGTGTATCCTTGGGCGCT.....GAATGATTTTCTTGAACAGATTGATTCAGTGGAGCTGC	Hsap NM_006141 i06 Rnor NM_031026 i06 Ggal chr11_11675855_11676572	11 1 2 OK 11 1 1 OK 11 1 1 OK
	TGCCTCAGTGTATCCTTGGGCGCT.....GAATGATTTTCTTGAACCAATGATTCAGTGGAGCTGC		
	TGCTTACAGTGTATCCTTCCCTGCTC.....CTATGTATTTTCTTAAATAAATGATTCAGTGGAGCTGC		