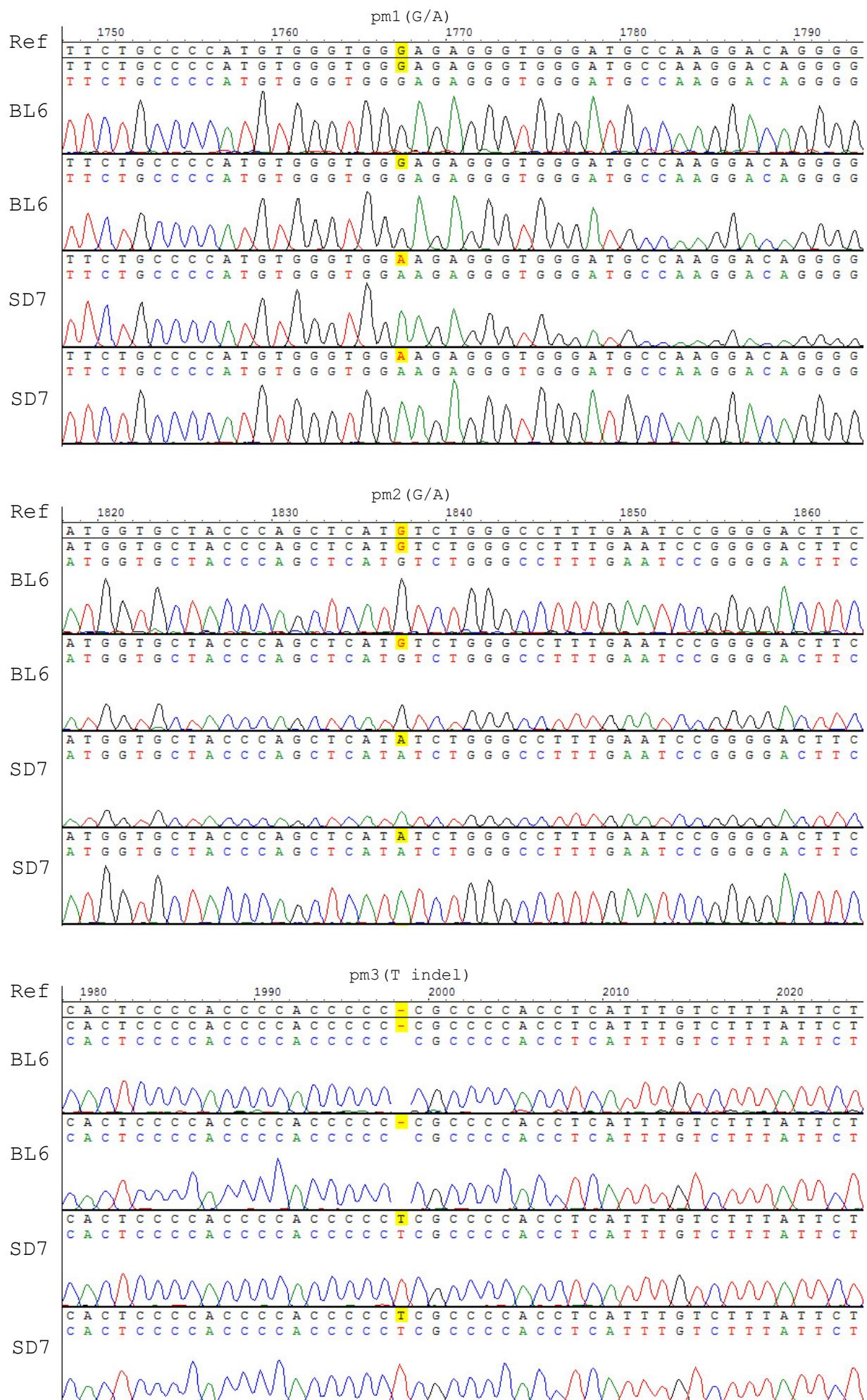


Figure S2 A



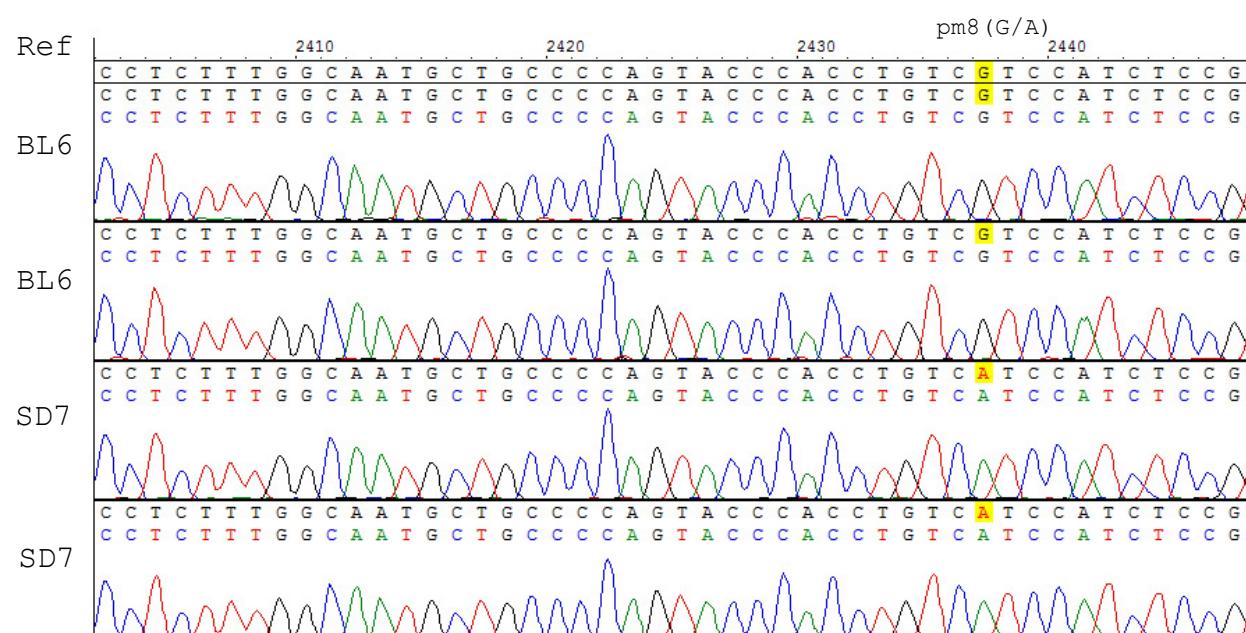
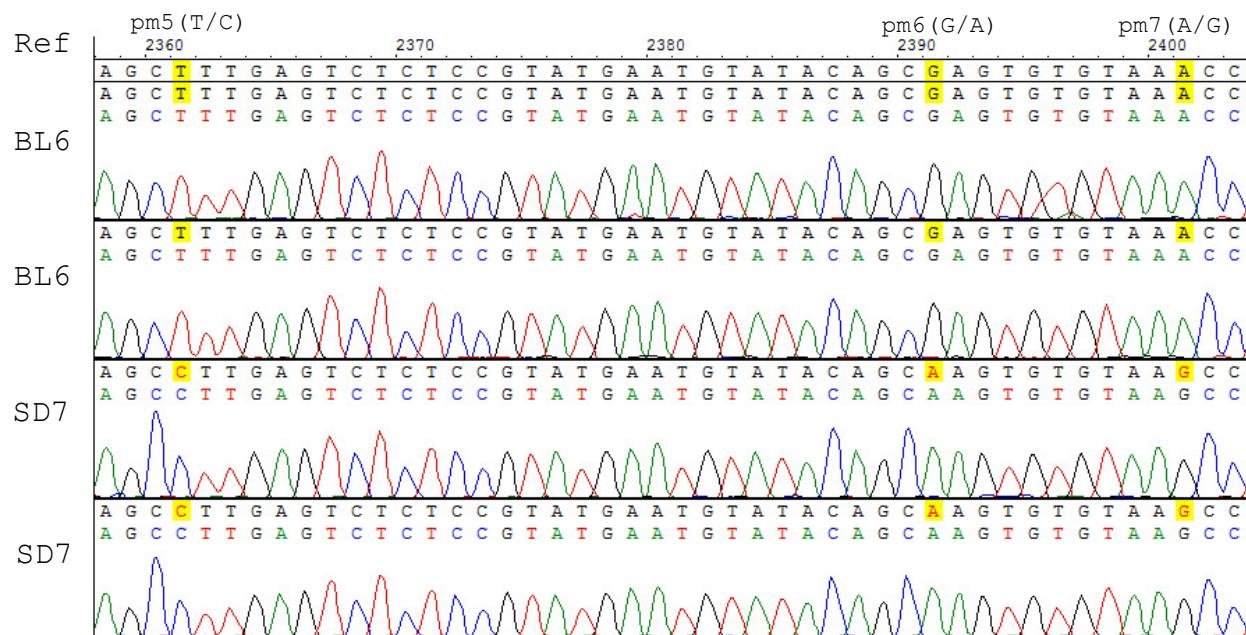
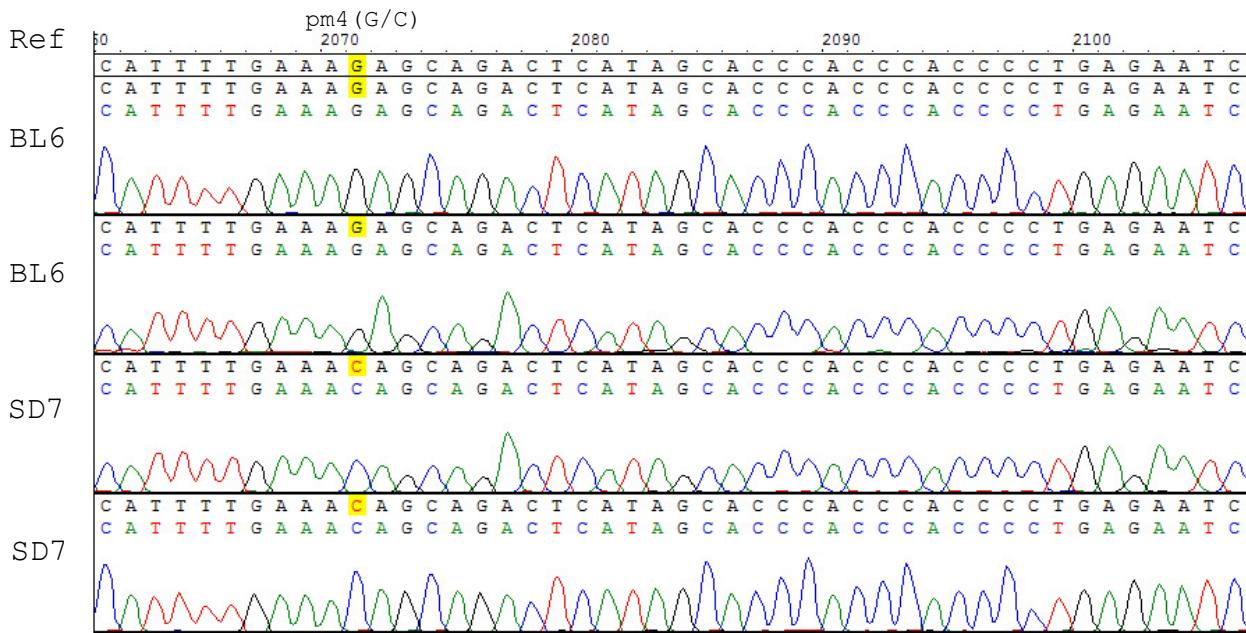


Figure S2 B

Thu Apr 26, 2012 18:59 +0200
H19_Ref_ENSMUST00000136359_NCBIM37_chr7_149761434-149764048_str_-1.str from 1 to 2615
Alignment to
H19_SD7_Seq.ape from 1 to 915

Matches():907
Mismatchs(#):7
Gaps():1702
Unattempted(.):0

Exon1

1	gggtgtggggggggggggggggggggggggatcgaaaaactggggaaatggggagactggaggagactgtgggggtccgaggacacctcg	100
1	~~~~~	1
101	gcatctggagtctggcaggaatgttaaggactgaggggctagctcaggcagagcaaaggcatcgcaaaaggctggaaaacatcgaggtaagctgaagg	200
1	~~~~~	1
201	cctgagctagggttggagagaatggggagccagacattcatcccggttactttggttacaggacgtggcggctggctggataaaggggagctgtgg	300
1	~~~~~	1
301	aagggttcaccccagacactggcagtgaaaggtagctggcagcgtggcagggtgaggaccgcgtctgtggcagggtgagttccctttctct	400
1	~~~~~	1
401	tggcctcgcttcaactgacttctaaacgaaggtaggtttagagagggggctgtgagaagaagcggctggctcgacgacaatggcacatagaaggcaga	500
1	~~~~~	1
501	tagtttagcaaaggagacatcgctcggggggagccgagacagaaggaggctggggaccattggcagccccaggtagggaaagagcttttagagagaagaaa	600
1	~~~~~	1
601	gaagaggtcagggttgcagtaaagactgaggccgtgcctccaggagggtataggagtcttggagacagtggcagagaccatggatccagcaaga	700
1	~~~~~	1
701	acagaaggcattctaggctgggtcaaacaggcaagatgggtcacaagacacagatggttccagccacaacatcccaccaccgttaattcactt	800
1	~~~~~	1
801	agaagaagggtcaagagtggtctggcaaagtcccaagttgccagagcctcaataactggagaatggaaaagaaggcagtgccagggtgtcaccagaag	900
1	~~~~~	1
901	gggagtgggctgcaggatcgactccagaggatttacagaaggaggctgcagtggtccagcctgcagacacaccattccatggactcg	1000
1	~~~~~	1
1001	gcccaggactgggtggaaaggccacagtggacttggatactgttatgccttaaccgctcagtggtccagcctgcagacacaccattccatggactcg	1100
1	~~~~~	1
1101	ggggagtcaggcacaggatgaaggccagacgaggcaggcaggcggggcagaatgaatgatgggtttctaggagggatgggtcaggttagagcgagta	1200
1	~~~~~	1

850 CTCTTTGGCAATGCTGCCAGTACCCACCTGTCCATCTCCGTCTGAGGGCAACTGGGTGTGG~~~~~ 915
* * * * * * * *
* * * * * * * * * * * * * *
2500 gtcttggaaaggcgtttccatataaagtgttcaacatgcctacttcattttggcccttcaccaggcccaccagagggtccatggtccatcaataaa 2599
915 ~~~~~ 915
* *
2600 tacagttacagtcatt 2615
915 ~~~~~ 915