Scale	50 bases mm39
chr13:	48,689,420 48,689,430 48,689,440 48,689,450 48,689,450 48,689,450 48,689,540 4
>CA	CTTACGAGAACCACTGCATGAACAACTCTTGCTTACACGAACCCAAGGCAGAAGGAACGTATGCTGGTATAATAAGGCCCTAAGAAAGGCAGGTCGTATAAGTCCTTGTGGGCAAAATCTCTGCCCTAAAACTATGCAACCT
	User Supplied Track
User Track	
6720427J07Rik	GENCODE VM27
07204271071CIK	Mide 7d
	RefSeq genes from NCBI
RefSeq Curated	
	RefSeg mRNAs mapped to this assembly
AK042687	Mouse mRNAs from GenBank
KY467470	
AK160312	
	AJ459695
4.28 _	35 vertebrates Basewise Conservation by PhyloP
Cons 35 Verts	
-4.1 _	
Gans	Multiz Alignments of 35 Vertebrates
	TTAGGAGAACCACTGCATGAAGAACAACTCTTGCTTACACGAACCCAAGGCAGAAGGCAAAGGAAAGGTATGCTGGTATAATAAGGCCCTAAGAAAGGCAGGTCGTATAGTTAACTCCTTGTGGGCAAAATCTCTGCCCTAAAACTATGCAACCT
	CTTACGAGAACCACTGCACGAAGAAC TTTTGCTTACACGCAT GGAACTATATGCTGGTATGATATGCCCCTAAGAAAGGCCCTAAGAAAGGCAGGTCGTATAGTTAACTCCTTGTGGGCAAAATCTCTGCCCTAAAACTATGCAACCT
	CGT - TA AGAAA A T C TGCAA ACAAAC TC - TGCTTACACCACA GCCAAGCAAGCAAGCAATCGCACC CTAGAACTATAGCAACCTTAGCACCTTAGAAACTATGCAACCTTAGAAACTATGAAGAAAGTAATGAAGAAAGTAATGAAGAAAGTAATGAAAACTATGAAAACTATGAAGAAAACTATGAAGAAAACTATGAAGAAAACTATGAACTATGAAAACTATGAAGAAAACTATGAAAACTATGAAAACTATGAAAACTATGAAAAACTATGAAAAAACTATGAAGAAAACTATGAAAACTATGAAAAACTATGAAGAAAAGAAAAAACTATGAAGAAAAGAAAAAACTATGAAAAAACTATGAAGAAAAACTATGAAGAAAAACTATGAAGAAAAACTATGAAGAAAAACTATGAAGAAAACTATGAAGAAAAACTATGAAGAAAACTATGAAGAAAACTATGAAAACTATGAAAACTATGAAAAACTATGAAAACTATGAAAAACTATGAAAACTATGAAAAACTATGAAAAAAAA
	- TTTGAGAAAACCTGCAGTAAGTGCTTACACCTGAATAA-ATAACAAGGAAACCTTAGAATAATAAGGCCCTAAGAAAGGCAGCAGGTCGTATAGTTACCTCCTTGTGGGCAAAATCCCTGCCTAAAACTATGCAACCT
	TTANGAGANACTTGCAGGNAGAAC - TA TA TTACACCAAAGAAGAAGAAGAAACATACACGTGGATAATAAGGCCCTAAGAAAGGCAGAGGTGGTATAGTTACCTCCTTGTGGGCAAAATCCCTGCCCTAAAACTATGCACCTG - TGAGAAAATTTGCAGGAGAGGAAGACA- GC - GC - TTACACCAAAGGAAGGAGAGAATTACCTGCCCTAAAACTATGCAACCTT
	TENGRAPH TENGRAPH AND THE CONTROL OF
	TT- GAGAAAACCTGCAGAAAGAAC - TC-TGCTTACACCAAAGCAAAGTAGCAAAGTAGCAAACTTACCCTTGAATAATAAGGCCCTAAGAAAGGCAGGTCGTATAGTTACCTCCTTGTGGGCAAAATCCCTGCCTAAAACTATGCAACCT
	STTATGAGAAAACCTGCAGAAGGAACTC-TGCTTACACCAAAGCACGGAAGGAAACTGACCATGAACTATGCAACCT
	TTCTAAGGAAACTTGCAGAAAGAAC-++CA-TGCTTACACCAAAACAAGGTGA-AAAG-AGCTTACCGTGGAGTAATAAGGCCCTAAGAAAGGCAGGCCGTATAGTTACCTCCTTGTGGGCAAAATCCCTGCCTAAAACTATGCAACCT
	CTCGTGAGAAACTCTCCAGAAAGATG-TGTTTACACCAAAGGAAAGG
Opossum	Repeating Elements by RepeatMasker
RepeatMasker	
	Short Genetic Variants from European Variant Archive Release 3
EVA SNP Release 3	