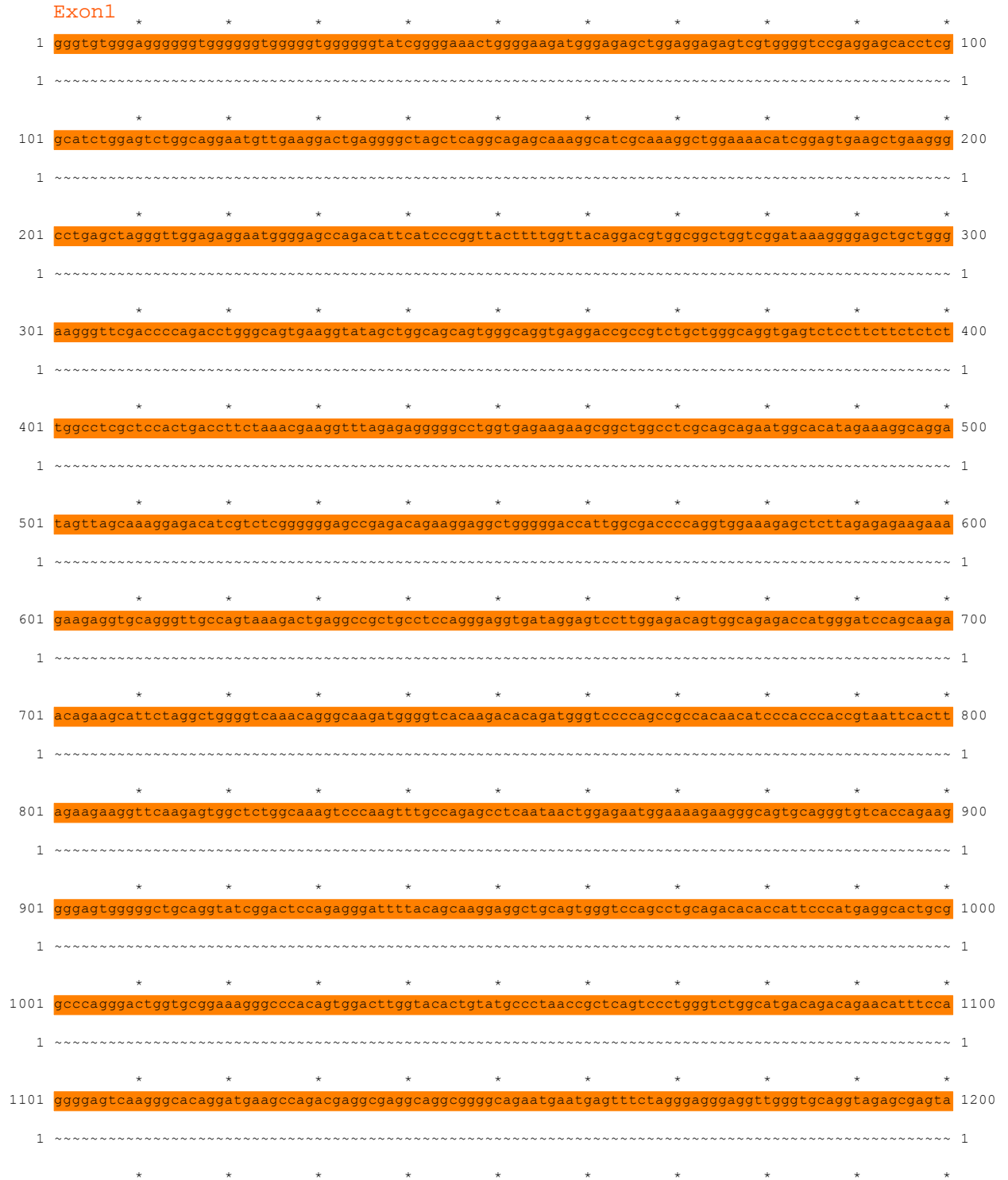


Figure S2 B

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
Mismatches (#):7
Gaps ():1702
Unattempted(.):0



850 CTCTTTGGCAATGCTGCCCCAGTACCCACCTGTCATCCATCTCCGTCTGAGGGCAACTGGGTGTGG~~~~~ 915

* * * * *
* * * * *

2500 gtctggaagcagttccatcataaagtgttcaaatgcctacttcatcctttgccctcctcaccagggcctcaccagaggtcctgggtccatcaataaa 2599

915 ~~~~~ 915

* *
2600 tacagttacagtcatt 2615

915 ~~~~~ 915