

Additional File 5. Search for the gene loss remnants of S100A7 and S100A15 from mouse and human, respectively.

Figure 1. Alignment of a partial sequences of S100A7 from mouse. The abbreviations correspond to the following species: H. sapiens - *Homo sapiens*; M. musculus - *Mus musculus*. Dots = identity with *H. sapiens* S100A7 sequence.

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          10   20   30   40   50   60   70   80   90   100
H. sapiens-S100A7 ATGAGCAACACTCAAGCTGAGAGGTCCATATAAGGCATGATCGACATGTTTACAAATACACCAAGACGTGATGACAAGATTGAGAAGCCAAGCCTGCTGA
M. musculus-Pseudo_S100A7 ...A.G...CA...A.A...GG.G...GGA...A.T...TG...C...GGGATAA...CA...T...CA...GATGG.AA...

          110  120  130  140  150  160  170  180  190  200
H. sapiens-S100A7 CGATGATGAAGGAGAACCTCCCCAACCTCCCTAGTGCCTGTGACAAAAAGGCCACAAATTACCTGCCGATGTCCTTGAGAAAAAGGACAAGAAATGAGGA
M. musculus-Pseudo_S100A7 A..AT...T.A...C.AA.T...TG..T.GAA.T.T.C..TT.ACATAATTGT.GGGCCTCTGT.A.ACAC.GA.GC.T

          210  220  230  240  250  260  270  280  290  300
H. sapiens-S100A7 TAAGAAGATTGATTTCTGAGATTCTGTCCTTGCTGGAGACATAGCCACAGACTACCACAAAGCAGGCCATGGAGCAGGCCCTGTCGGGGGCAAGC
M. musculus-Pseudo_S100A7 CT..TTT.ACAC.A.GCTGA.AAGGT

          ...
H. sapiens-S100A7 CAG
M. musculus-Pseudo_S100A7

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Figure 2. Genomic location of the partial S100A7 sequence from mouse. Data was obtained using S100A7 coding sequence from human as a reference against Ensembl database.

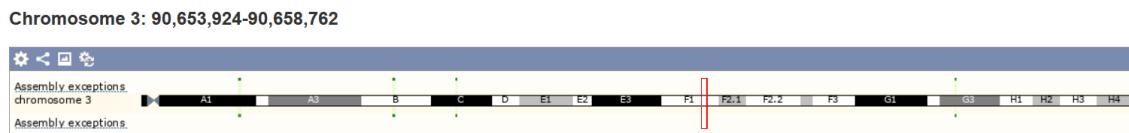


Figure 3. Alignment of a partial sequences of S100A15 from human. The abbreviations correspond to the following species: M. musculus - *Mus musculus*; P. troglodytes - *Pan troglodytes*; H. sapiens - *Homo sapiens*. Dots = identity with *M. musculus* S100A15 sequence.

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          10   20   30   40   50   60   70   80   90   100
M. musculus-S100A15 ATGCCAGACACCCAGTGGAGGACTCCCTCTTCAAATCATACACTGCTTCCATCAGTATGCTGCCGGAAAGGGGACAAGGAGACCTGTGTCCTGGAGG
P. troglodytes-S100A15 ...A.....C...A.....C.....C...C.G...A.....C...T...T.T.....C...
H. sapiens-Pseudo_S100A15 ----

          110  120  130  140  150  160  170  180  190  200
M. musculus-S100A15 AGTTGAAAGCTCTGCTTGGATAGTGCTCGCTTCATGGACACCTGGCCGGAGGCCATACATACAGAGCTGTTCCGGCAGGTGACAA
P. troglodytes-S100A15 ...C.G...C...A...C.AC...C.....G.....G.A.G.....T...T.....A...G...A...
H. sapiens-Pseudo_S100A15 ----TCA.....G.A.G.....T...T.....A...C...A...

          210  220  230  240  250  260  270  280  290  300
M. musculus-S100A15 AAACAAGGACACCCAGTCGCTTGGATAGTCCATGTCATCTGGCAAGCTGGTGAAGGACTACCATCTCCAGTTCACCCGGCAGTTGTGCAAC
P. troglodytes-S100A15 G.....C...C...C.....C.....C.....C.G...A...T...C.....C...
H. sapiens-Pseudo_S100A15 G.....C...C...C.....C.....C.G...C.A...A...C...C.G.

          310  320
M. musculus-S100A15 TACTGTACTGAGCACAGCCTCTACTAG
P. troglodytes-S100A15 ...G.CC.A.....C...
H. sapiens-Pseudo_S100A15 ...G.CC.A.....C...

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Figure 4. Genomic location of the partial *S100A15* sequence from human. Data was obtained using *S100A15* coding sequence from mouse as a reference against Ensembl database.

