

A. Mar-Hsa AAATAACA-- -TTTTTATAA -----ATTA TAAGATTCAA ATTTTAAAAG ATGGGGCTGG >>>
 Mar-Ptr AAATAACA-- -TTTTTATAA -----ATTA TAAGATTCAA ATTTTAAAAG ATGGGGCTGG >>>
 Mar-Gor AAATAACA-- -TTTTTATAA TTATAAATTA TAAGATTCAA ATTTTAAAAG ATGGGGCTGG >>>
 Mar-Ora AAATAACA-- -TTTTTATAA TTATAAATTA TAAGATTCAA ATTTTAAAAG ATGGGGCTGG >>>
 Mar-Sia AAATAACA-- -TTTTTATAA -----TTA TAAGATTCAA ATTTTAAAAG ATGGGGCTGG >>>
 Mar-Gre AAGTAAACA-- -TTTTTATAA -----TTA TAAGATTCAA ATTTTAAAAG ATGGGGCTGG >>>
 MAR-Rhe AAGTAAACA-- -TTTTTATAA -----TTA TAAGATTCAA ATTTTAAAAG ATGGGGCTGG >>>
 Mar-Owl AAATAACAAC ATCTTTATAA TTATAAATTA CAAGATTCTA ATTTTAAAAG ATGGGGCCAG >>>

Mar-Hsa GCACCTCCAGC TTGGGTGACA GAGCGAGACT --GTCTCAA AAAAAAAAAA-----
 Mar-Ptr GCACCTCCAGC TTGGGTGACA GAGGGAGACT CTGTCTCGAA AAAAAAAAAA AAAAAA
 Mar-Gor GCACCTCCAGC TTGGGTGACA GAGGGAGACT CTGTCTCAA AAAAAAAAAA-----
 Mar-Ora GCACCTCCAGC TTGGGTGACA GAGCGAGACT CTGTCTCAA AAAAAAAA-------
 Mar-Sia GCACCTCCAGC TTGGGTGACA GAGCAAGACT CTGTCTCAA AAAAAA---------
 Mar-Gre GCACCTCCAGC TTGGGTGACA GAGCGAGGCT CTGTCTCAA AAAAAA---------
 MAR-Rhe GCACCTCCAGC TTGGGTGACA GAGCGAGACT CTGTCTCAA AAAAA-----
 Mar-Owl GTACTCCAGC CAGGGTAACA GAGCAAACT CTGTCTTAAG AAAAA-----

Mar-Hsa AAAAGTA- --ACAGTTTT TGCATTGTTG GAATTTGGTA TTTGATATTG >>>
 Mar-Ptr AAAAGTAG TTACAGTTTT TGCATTGTTG GAATTTGGTA TTTGATATTG >>>
 Mar-Gor GAAAGTAG TTACAGTTTT TGCATTGTTG CAATTTGGTA TTTGATATTG >>>
 Mar-Ora AAAAGTAG TTACAGTTTT TGCATGGTTG GAATTTGGTA TTTGATATTG >>>
 Mar-Sia AAAAGTAG TTACAGTTTT TGCATTGTTG GAATTTGGTA TTTGATATTG >>>
 Mar-Gre AAAAGTAG TAACAGGTTT TGCATTGTTG GAATTTGGTG TTTAATATTG >>>
 MAR-Rhe AAAAGTAG TAACAGTTTT TGCATTGTTG GAATTTGGCA TTTAATATTG >>>
 Mar-Owl AAAAGTAG TTGCAGTTTT TGCATTGTTG GAATTCGCTA TTTGATATTG >>>

Mar-Hsa ACCGCAGTTA GTTTTGCACC AACCCTAAT CTTCATAG-A TTGAAATATA AATTAAAATT
 Mar-Ptr ACCGCAATTA GTTTTGCACC AGCCCAAT CTTCATAG-A TTGAAATATA AATTAAAATT
 Mar-Gor ACCGCAGTTA GTTTTGCACC AACCCTAAT CTTCATAG-A TTGAAATATA AATTAAAATT
 Mar-Ora ACCGCAATTA GTTTCACACC AACCCTAAT CTTCATAG-A TTGAAACATA AATTAAAATT
 Mar-Sia ACCGCAGTTA GTTTTGCATC AACCCTAAT CTTCATAG-A TTGAAACAGA AATTAAAATT
 Mar-Gre ACCGCAATTA GTTTTGCACC AACCCTAAT CTTCATAGAA TTGAAACAT- ----AAAATT
 MAR-Rhe ACCGCAATTA GTTTTGCACC AACCCTAAT CTTCATAGAA TTGAAACATA AATTAAAATT
 Mar-Owl ACCGCAGTTA GTTTTGCACC AACCCTAAT CTTCATAGTA TTAAAACATA AATTAAAATT

B.

MAR-Hsa	CACTC----	T	TTTGAGAGAA	AACATTTAAA	AATATACTTC	CACTTGACTA	TTATCCCATG	
MAR-Tar	CACGTCTATT	T-----	----	TTTAAA	AATATACTTC	TGCTTGACTA	TTATCCTATG	
MAR-Dog	CACTTCTGTT	TTTGAGAACA	AACATTTAAA	GATATA----	TATTTTTTTA	TTATCT----		
MAR-Hsa	ATAAAATAAC	AT-----	TTT	TAT-----	A	AATTATAAG-	-ATTCAAATT	TTAAAAGATG
MAR-Tar	GTAAAATAAT	ATAAAAGTTT	TAT---	TACA	AATTACAAG-	-ATTGAATT	TTTTAGAGTG	
MAR-Dog	-TAAAATA-T	ATATAAACTT	TACA	ACTAAA	AAGTACTGCA	AACTCGAAGT	TTAAAAGGTG	
MAR-Hsa	G----	GGCTG	GGTGCAGTGG	CTCACACCTG	TAATCCCAGC	ACTTTGGGAG	GCTGAGGCAG	>>>
MAR-Tar	GCATG	-----	-----	-----	-----	-----	-----	>>>
MAR-Dog	GCATA	-----	-----	-----	-----	-----	-----	>>>
MAR-Hsa	AGCCTAGTTA	TAATGATTTA	AAATTCACAG	TCCAAAACCG	CAGTTAGTTT	TGCACCAACC		
MAR-Tar	-----	-----	-----	-----	-----	-----		
MAR-Dog	-----	-----	-----	-----	-----	-----		
MAR-Hsa	CAATATCT--	-TCATAG-AT	TGAAATATAA	ATTA AAAATTG	CATTTGAAGT	AGAT		
MAR-Tar	--TATCTAG	CTCATGGAAT	TGAAACA--A	ATAAAAATTG	GCTTTAAAGT	AAAT		
MAR-Dog	--TATCT--	-TCATGAAAT	TAAAACATAA	A----AATTG	GCTTTGAAGT	CAAT		

C.

SETex2-Tar	ATCAATTAGG	CCGTTTCTCG	GCCTCTCTCA	GTGGGTGGGA	ATGAGGGGGA	ACTAAGCTTA	
SETex2-Owl	ATCAGTT	-----	-----	-----	GTGGAA	ATGAGAAGGA	ACCCAGCATG
SETex2-Rhe	ATCAGTT	-----	-----	-----	GTGGAA	ATGAGAAGGA	ACCCAGCATG
SETex2-Gre	ATCAGTT	-----	-----	-----	GTGGAA	ATGAGAAGGA	ACCCAGCATG
SETex2-Sia	ATCAGTT	-----	-----	-----	GTGGAA	ATGAGAAGGA	ACCCAGCATG
SETex2-Ora	ATCAGTT	-----	-----	-----	GTGGAA	ATGAGAAGGA	ACCCAGCATG
SETex2-Gor	ATCAGGT	-----	-----	-----	GTGGAA	ATGAGAAGGA	ACCCAGCATG
SETex2-Ptr	ATCAGTT	-----	-----	-----	GTGGAC	ATGAGAAGGA	ACCCAGCATG
SETex2-Hsa	ATCAGTT	-----	-----	-----	GTGGAA	ATGAGAAGGA	ACCCAGCATG
SETex2-Tar	AGTGGCTCAG	CCACTTCTGC	CTTCCCTCTCT	TGCAAGCAGT	TTATGCTCGA	GTAAGTCTG	
SETex2-Owl	TGTGGCTCAG	CCCCTTCTGT	GTTCCCTCTT	TGCAAGCGAT	TGACCTTGA	GTTGAGTCTG	
SETex2-Rhe	TGTGGCTCAG	CCCCTTCTGT	GTTCCCTCTT	TGCAAGCGAT	TGACCTTGA	GTTGAGTCTG	
SETex2-Gre	TGTGGCTCAG	CCCCTTCTGT	GTTCCCTCTT	TGCAAGCGAT	TGACCTTGA	GTTGAGTCTG	
SETex2-Sia	TGTGGCTCAG	CCCCTTCTGT	GTTCCCTCTT	TGCAAGCGAT	TGACCTTGA	GTTGAGTCTG	
SETex2-Ora	TGTGGCTCAG	CCCCTTCTGT	GTTCCCTCTT	TGCAAGCGAT	TGACCTTGA	GTTGAGTCTG	
SETex2-Gor	TGTGGCTCAG	CCCCTTCTGT	GTTCCCTCTT	TGCAAGCGAT	TGACCTTGA	GTTGAGTCTG	
SETex2-Ptr	TGTGGCTCAG	CCCCTTCTGT	GTTCCCTCTT	TGCAAGCGAT	TGACCTTGA	GTTGAGTCTG	
SETex2-Hsa	TGTGGCTCAG	CCCCTTCTGT	GTTCCCTCTT	TGCAAGCGAT	TGACCTTGA	GTTGAGTCTG	

Fig. 4. Alignment of the orthologous *MAR* region in nine primates and dog, in which the *Hsmar1* transposon is present (A) or absent (B), and alignment of the orthologous 3' end of the second exon of the *SET* gene in nine primate species (C). Hsa, human; Ptr, common chimpanzee; Gor, gorilla; Ora, orangutan; Sia, siamang; Gre, African green monkey; Rhe, rhesus macaque; Owl, owl monkey; Tar, tarsier; -, sequence gap; >>>, a portion of the sequence is not shown. The colored boxes above delimited alignment portions indicate the location of the *AluSx* insertion (red), the *Hsmar1* transposon insertion (green), the 3' end of SET exon 2 (orange), the 27-bp deletion in anthropoid primates relative to tarsier (blue), and the 77-bp exonized sequence linking the former SET exon 2 to the donor splice site of the second intron of *SETMAR* (yellow). The location of the SET stop codon in tarsier is shown in bold, and the start of intron 2 of *SETMAR* at dinucleotide GT is underlined in anthropoid primates.