

Supplementary Table SII Exon-intron boundaries of the rat *NMS* gene

Exon number	Exon size	Sequence at exon/intron junction		Intron size
		5'-splice donor	3'-splice acceptor	
1	N.D.	CCTCCTCAGgtaaggg roSerSerG	gtgatagGAGCTTCTC lyAlaSerP	2058 bp
2	56 bp	GACCCAGAGgtatgct AspProGlu	ctttcagCGACTGGCA ArgLeuAla	305 bp
3	51 bp	CAACCTAAGgtaatat GlnProLys	cttccagGAAAGCCGG GluSerArg	2169 bp
4	24 bp	TACAAAAGGgtgagga TyrLysArg	cgttcagTTTTTATTT PheLeuPhe	90 bp
5	54 bp	AACTCCGAGgtctgtg AsnSerGlu	gttgcagTTTGCTCCC PheAlaPro	1788 bp
6	84 bp	CTGCACACagtaggtg LeuHisThr	ttcccagGATTCCAGG AspSerArg	932 bp
7	36 bp	CCTAAGAAGgtatgtg ProLysLys	gttacagGATCCTACC AspProThr	626 bp
8	42 bp	CTTTTCAGGgtatagc LeuPheArg	N.D. CCTAGGAAT ProArgAsn	>0.5 kb
9	44 bp	GACGGCAAG N.D.	N.D. AATCCTATG	>0.5 kb
10	N.D.	N.D.		

The exon and intron sequences are shown in uppercase and lowercase letters, respectively. The genomic sequence of the *NMS* gene was identified in one genomic contig of rat chromosome 9 (RefSeq accession number NW_047814). N.D., not determined.