Gene: SDHB - Sequence: NG_012340.1 Transcript: NM_003000.2 - Protein: NP_002991.2 Date : February 26, 2015

1^{st} line: Base numbering. Full stops for intronic $+/-$ 5, 10, 15 2^{nd} line: Base sequence. lower case Introns, upper case Exons 3^{rd} line: Amino acid sequence. Printed on FIRST base of codon 4^{th} line: Amino acid numbering. Numbered on 1^{st} and increments of 10
Exon 1 Start: 5001 End: 5223 Length: 222
-149 -139 -129 -119 -109 -99 ACCCGGGAAACCGGAAGCCGCCTCCCACTTGGTTGCTCGTACGCGGCTAGTGGGTCCTCA
-89 -79 -69 -59 -49 -39 GTGGATGTAGGCTGGGCGCGCGATGTTCGACGGGGACACCGGCGGAGAGCGACCTCGGGG
-29 -19 -9 1 11 21 TTAAGGGGTGGGCTGACGTCAGGAGCCAAGATGGCGGCGGTGGTCGCCCTCTCCTTGAG M A A V V A L S L R 1
31 41 51 61 71 GCGCCGGTTGCCGGCCACAACCCTTGGCGGAGCCTGCCTG
agagaggaaaagtcaggaaagccacagagactgcctggagctgatggggctgcggagag

actgagg	gaagca	ataggg	gcaagg	gcctca	igggag	gatgga	uggaco	ctcctc	cccct	ggtgc	agt
gctttgg	gtttgg	gcttcg		ccctg		gttato			ctcgca	igcago	ttg
tgagtta	agggca	acgggo	:aattt	tttac			gcggg	ggggg	ggggao	gttca	.gaa
agggga		attgct					natgao	ct			

Ex	on	2	St	art	: 14	1283	E:	nd:	144	10	L	eng	th:	127	7			
tc	ttt	ata	ittc	atc	gtga	aatgt	tta	attt	aac	tct [.]	tgt	tca	ctad	caaa	agtaa	agt	tct	gtg
aa	gat	gaa	ıggt	ttt.	tgtg	gtatt	aag	tgct	gtg	tca	cta	gcc	ccta	aata	accat	tgt	ctg	gata
ta	tta	gca	itat	gat	agg†	tgctt	aat	actt	gtt	gga	tat	tga:	atgo	cct	gcct	ttt	cta	aga
aga	att	atg	gtgt	cag [.]	taa†	tgtgt	gag	ttta	ıtat	cca	gcg	tta	cato	ctg1	ttgtį	gcc	ago	caaa
at	gga	att	atc	ttg	tati	ttcta	att [.]	tttt	ttt	cct [.]	ttt	tgt:	gaad	ctt1	taaaa	aaa	ttt	cag
GC(CTC S		81 G G		CCA(Q	91 GACAG T A		CAGC	101 CCAC T		TCC P			CAA(K	12: GAAA: K 41	ГТТ	GCC A	131 CATC I
TA' Y	ГСG R		14 GGA D		AGA(D	151 CAAGG K A 51	CTG	GAGA	161 CAA K	ACC'		17: TAT(M		GACT	18: TAT(Y 61	GAA	.GTT V	191 TGAC D
CT' L	ΓΑΑ N	TAA K	ıgtg	agt:	atci	tctgt	gaa	agcc	:agc	tat	tga	agga	agag	gtto	cttga	att	tga	nttt
ag	gga	cat	gct	ttt	caca	atcct	tgg	aagg	gctt	aaa	aat	ctg	agat	cat	tccga	atg	gct	cctg
tt [.]	ttt	aaa	ıaca	aga	atga	agctt	ggg	ccgg	gca	cgg	tgg	ctc	acgo	ccta	ataa	tcc	cag	gcac
tt [.]	tgg	gag	gct	cag	gcgg	ggcag	gatc	atct	gag	gtt	ggg	agt [.]	tcaa	agad	ccaat	tgt	ggo	ccaa
ca ⁻	tgg	tga	aac	tcc	cct	ctcta	icta	aaaa	ıtac	aga	aat	tag	ctgg	ggta	atagi	tgg	cto	catg
cc.	tgt	aat	;															

Exon	3	Start:	260	26	End:	2611	l1 I	Length:	85			
												gtc
gagct	ttcag	gaagtaa	iaaaa	ıcaaa	aacto	:ggcca	actgta	atggaag	ttgtg	;tcata	cttg	
												ag
tacta	actac	tttago	ctttc	catt	tccta	igaaat	:gtgti	ttgcact	gtgat	catca	aaat	
												ega
aactg	gtgca	ıgtactt	aaaa	itgta	agaat	ctgca	naaaa	caaaact	acaga	acaaa	ttcc	
												ta
aggt	gacct	gagaag	gacca	laatg	gataa	igctaa	ataca	tccaggt	gtctc	cgatt:	atat	
												ag
tgata	aaagt	gtaggg	gaggt	tgaa	.cgtta	acataa	aatac	cactgga	tattt	ttctt	tgtt	
	GGCCC G P	211 CATGGT M V		22 GATG D A	CTTTA	23 ATCA <i>I</i> I K	AGATT	241 AAGAATG K N E	AAGTT			
	J P	71	ь			ıĸ	1 1		v v	ם ע	1	L
	TTCCG F R	271 GAAGATO R S 91	CATGC C	28 AGAG R E	AAGgt	gagca	atttca	attcctg	ttggg	;ctcca	gata	ıct
tgtgg	gtctt	ccaaag	gaggo	ttgg	gctgg	gccaaa	agctga	atagaga	.catct	ggaaa	acag	gct
												at
gcago	ctctc	tgctgg	ggtga	actt	gctgt	gcaad	cctaat	ttccttt	tcttg	;ggtaa	atgo	
												att
ccaaa	aatto	taacca	itggt	gcaa	gttct	ggttt	cctgag	gtttact	ttcct	tcagg	agca	
												ag
ctato	cctgt	ggttac	:tgca	itctg	gcggt	ccctg	gtgag	ctgctcc	acgct	tcaag	gcct	
ctage	gacct	gctggc	ctggc	acct	tcc							

Exon 4 Start:	30435 End:	30571 Length	: 136
	 cgcccaggcctctg	 ggcctcctccagtgc	
	 .gccacatggtcat		 aactcaagccattagaa
	gaagaaggaaggaa	agggagaaaagccaa	
	agtgctgccctga	 atggcacagcaaggaį	ggatccagaagaaagta
		 gtgaggatgtgttaaa	atgtgtgtgtctctttcag
GCATCTGTGGCTCTTC I C G S C	301 311 GTGCAATGAACATC A M N I 101		331 341 FCTAGCTTGCACCCGAA L A C T R R 111
GGATTGACACCAACCT		381 AAAAATCTACCCTCT KIYPL	391 401 CCACACATGTATGTGA P H M Y V I 131
TAAAGGATCTTGTTCC K D L V P	121 CCgtgagtttctgc		gtttttatttgcatggg
gggcagtgctatgtgt	 .gttacgctattag	gtttttcagggcagga	 atttgtttgagtattta
aggctcctcttctagt	 cagatagtaaata	 agatgggactgtttco	
gaggaccacagtgtca	 actgagagattagg	gaaaatctccagatc	
agtttaattaaagaat	ttgaatggcagae	agataaacgcctagaa	 aaatcctccagcagctc
atttggtgaggtttgt	ct		

Exon 5 Start: 31306 End: 31422 Length: 116
431 441 451 461 471 483 484 485
491 501 511 521 531 GAATCTCAGGAAGGCAGCAGCAGTATCTGCAGTCCATAGAAGAGCGTGAGAAACTGgtc E S Q E G K Q Q Y L Q S I E E R E K L 171

Exon	6	St	art	: 3	3509	7	Er	ıd:	35198	3 I	eng	th:	101	L			
ttgtt	tttį	gtg	ttc	ttg	gttt	tct	att	tct	ttttt	agac	:aag	tgaį	gagg	gtt	cttg	gttt	att
aaagg	gaa	agt	gca	cta	ıtac	gtg	gat	tta	aagaac	cctg	ggc:	agat	tgtt	ta	catt	tca	ectt
gcaag	atc	ttt	att	aat	ctg	cta	aat	ato	ctgcaa	latca		cttg	ggat	tt.	tgct	:aag	gaaa
attca	cat	gca	.agt	agg	gcac	ttt	gtt	cat	tgcact	gaco	cca	aagg	gtaa	aca [.]	ttta	wato	agc
tgaag	aca	tag	cag	agt	ctc	tcc	cgt	cao	caagct	cctg	gcct	ctc	tttt	ct	cccc	ata	acag
541 GACGG D G 181		CTA	551 .CGA E			56 TCT L	CTC			GTAC	CAC				591 CAGC S		CTGG W
601 TGGAA W N 201	CGG. G	AGA		ATA		62 GGG G	GCC			TTAT M		641 Ggt	gagg	gtg	ctcc	tta	att
gcttt	aaga	aga	aat	tga	ıaac	tca	ago	cato	ccagaa	igtco	aaga	agga	acaa	ata	gatt	gct	gta
agcca	gcc	att	ctg	agg	gggc	ctg	gtt	tc	cttcat	tgcc	atc	cagt	tcca	aaa	cagg	ŗgta	att
ggttg	cat	tgt	att	cac	ttg	ctt	tcg	ggta	aaatct	gggg	gccti	tgt	ctgo	caa	attg	godt	ttg
tttca	gtg	cag	tga	ggg	gtca	.ccc	caa	nagg	gggtgo	aaac	cat	gcc1	ttgo	cag	atgt	tta;	aaaa
tgtat	ttc	cgt	ttg	cct	ggt	ggt	cag	gagg	gctccc	cctc	tcc	C					

Ex	on	/	St	art	: 3	6441	1	Łna	l: 3	3650	63	ΙL	eng	th:	122	2				
ct	ggg	tag	gaa	gct	att	gagc	aatį	gtg	aaa	ata	att	agg	gaga	tga	tac	ccag	tgt	tgta	ataa	
tt	aag	cac	ccg	ggt	acc	aggt	gcaį	gta	tgg	gat	caa	agg	gttt	gca	ggga	acca	ıtta	aato	caag	
tg	aag	cta	caa	gtg	ttg	ggaa	cct	ttc	cti	tga:	aag	ate	gac	aag	aatt	taaa	ıgag	gtai	tgga	
ga	tga	ggc	atc	taa	gct	gagt	gaat	ttc	cct	ttt	cct	ctg	gcac	tcc	caga	agct	ttg	gagt	ttga	
gc	cag	ggt	gcc	gct	ctg	gcct	gttį	gat	tgg	gca	gct	cag	gcta	atc	atco	ccte	gtt	ttt	ccag	
GC A	CTA Y	TCG R	65 CTG		GAT'	66 TGAC D 22	TCC.			671 ГGА(D		CAC T	68 CAGA E		GCG(R	69 CCTG L 23	GC(A	CAA(K	70: GCTG L	1
CA Q	GGA D		71 ATT F		TCT. L	72 ATAC Y 24	CGC R (731 CAC T		CAT M	74 GAA N		CACA T	75 AAGG R 25	ACC T	CTG1 C	76: ICCT P	1
AA K	Ggt	acg	tgg	ggc	tgg	tgac	cct	cag	;ct	ctg	cct	cag	gagc	tga	cac	gcca	ıgaa	agta	agca	
tg	tgc	tca	caa	aga	ggt	gatt	ggc:	aga	.gag	gca	acc	cta	ıgtc	att	tgag	gaag	gaa	aagg	ggac	
ca	gca	tat	gcc	tgt	cct [.]	ttca	gtgʻ	tga	.gc	ctaį	ggg	gct	agt	ttg	gcto	ctta	ıgtt	gca	aagg	
ag	tag	act	gac	tcg	ggc	cacc	tcaį	gga	Igca	agg	ctt	tat	ttg	tag	gcta	acac	agg	ggao	catt	
gt	ggg					acag												cca	agga	
aa	C																			

Exon 8 Start: 40213 End: 40449 Length: 236	
tctttgagcagtcagaggggcacctctgccctgggcctgtttacctgttagagtgagcc	
${\tt acacaagaaggaattcaacgtggcagagctcttgccagagggagctgaccatcttcaaat}$	
gttaaaaaaggtagacagaagagagcccttagcccaacccctatggttttgagggacttg	
atttgcattcaccttgcttggacactgaaccagctgaggaagga	
ann phonon and a suppression of the suppression	
tgggttttccctttcagtttcagttatcatttgttattgattcttttctcttttttcaag	
771 781 791 801 811 821	
GGTCTGAATCCAGGGAAAGCTATTGCAGAGATCAAGAAAATGATGGCAACCTATAAGGAG	
G L N P G K A I A E I K K M M A T Y K E 261 271	
1201	
831 841 * 11 * 21 * 31 * 41	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V * 281	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V * 281 *51 *61 *71 *81 *91 *101	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V * 281	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	
AAGAAAGCTTCAGTTTAACTGTTTCCATGCTAAACATGATTTATAACCAGCTCAGAGCTG K K A S V *	

 $\tt gggatccata attgtgcctacttcattcatctggtatgagaatgcctggcacttatt$

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Reader: Version: 1, Version Date: 11/02/2015 Writer: Version: 1, Version Date: 11/02/2015 Control: Version: 1, Version Date: 11/02/2015