Date: 2020-08-13 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: 5128 | Length: 127 agaaa cat gtttgtctac gaggtagtatatctggtatttctg gaggaattcggttattata $\tt gggaggtaccatgaaagaacattcctactttagaaatagatattctctgttcaaacctga$ $\verb|cttgaaca| actggttttatgtgggtgtcaccagtgctttataaaagtaaactgcatcgcc|$ l-69 l*-*59 l*-*49 l-39 1-29 1-19 $\tt CCGAGTTCAGTCATACTGCACCAGCTGAGCAATGCATGGAGTGGACCTGTAGGCGACTTG$ 1-9 |1 |11 |21 |31 CATCGTCTTCAACATGAAGATAGCCACAGTGTCAGTGCTTCTGCCCTTGGCTCTTTGCCT M K I A T V S V L L P L A L C L

Gene: SPINK5 - Sequence: NG_009633.1 Transcript: NM_006846.3 - Protein: NP_006837.2 LRG_110t1 -

111

11

$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
tgctaaat

Exon 2 Start: 6376 End: 6401 Length: 25
$. \\ \ . \\ \ . \\ \ . \\ \ . \\ \ . \\ \ . \\ \ $
61 71 81 ATGCTGCCAGTAAGAATGAAGATCAGgttagtcctgctttttctgttcattgaattcatt A A S K N E D Q 21

aggad	catgag	cctct	ctctc	acaca	cacac	acaca	cacac	acaca	ctcaa	catat	taatg
											_
	·		•				•				
atgai	tggatg	gtctgc	aagag	gcaat	ttatc	atgtt	aaagg	agcat	tttaa	laattt	cataa
		•									

 ${\tt acaaaaaataaatcaactggaaaact}$

Exon 3 Start: 11352 End: 11479 Length: 127
91 101 111 121 131 141
GAAATGTGCCATGAATTTCAGGCATTTATGAAAAATGGAAAACTGTTCTGTCCCCAGGAT
EMCHEFQAFMKNGKLFCPQD 31
151
AAGAAATTTTTTCAAAGTCTTGATGGAATAATGTTCATCAATAAATGTGCCACGTGCAAA K K F F Q S L D G I M F I N K C A T C K 51 61
$\begin{tabular}{ll} ATGATACT gt gag taa agg ttt cttt cttt cttt ccaat gttt gag ttaa cag ctag tct \\ M & I & L \end{tabular}$

ctgaactggtaaatgtattctttttctttcaagtgcatttttctaaaccgaaatggttaa
ttcctggt

Exon 4 Start: 13177 End: 13249 Length: 72
211 221 231 241 251 261
${\tt GGAAAAAGAAGCAAAATCACAGAAGAGGGCCAGGCATTTAGCAAGAGCTCCCAAGGCTAC}$
EKEAKS QKRARHLARAPKAT 71 81
271 281

gtcctgtcttagc

gattttat

Exon 6 Start: 29571 End: 29634 Length: 63
411 421 431 441 451 461 GAAAACCGGGTCCCAAATTGGTGTAAAAAGTGAAGGGGAATGTAAGAGCAGTAATCCAGA K T G S Q I G V K S E G E C K S S N P E 141 151
471

	.a
	.a.
	t
cagt	

Exc	on '	7 S	tar	t: 3	3052	3	En	d:	306	50	Le	ngt	h:	127	•			
aaa	atg [.]	tctct	ttt	tato	acc	tag	gag	att	ggg	agc	aagt	tga	tgg	caa	ıgga	act	gtg	ctct
gt	ctga	agtto	ctgg	caca	ıtag	tag	gtt	atc	tgt	tta	ctaa	ata	aat	gaa	ıtaa	aac	aag	taaa
aaa	agt [.]	tttat	att	taad	cac	atg	aat	tat	aga	.gta	ctta	.ctg	agc	tad	ato	cta	cac	agct
gg¹	tac [.]	tgcto	:taa;	gtgg	gece	ata	gca	.atg	tca	.gag	ggac	tga	gtt	caa	ıtaa	aaa	ttt	ctga
aaa	acaį	gtgtt	gtc	agca	ıtta	.caa	tct	tgg	taa	gtt.	tcaa	gtt	ctt	tto	cct	tgt	tct	tcag
	ΓGΤ. V		GTG(CTTI	491 TCG R	GCC	CTT	50 TGT V	TAG	AGA	51 TGGA G 17	AGA R				CAC.	AAG R	531 GGAA E
AAT N	ΓGA' D	541 TCCTC P V 181	TTC L			TGA		GAA	GAC	GCA		AAT N		TGT			GTG C	591 TGCT A
GA(GCT(L	601 GTTgt F	aag	tago	catc	atc	ccc	agg	tgg	act	tgat	gat	gat	gca	ctt	gg	ttg	ctgt

cccgagaatcactcagcagagagataaaatccttttcatgaagcatgcaattctttgtct
taaaattt

Exon 8 Start: 32194 End: 32257 Length: 63
611 621 631 641 651 665 TTTAAAAGAAGCTGAAAATGCCAAGCGAGAGGGTGAAACTAGAATTCGACGAAATGCTGA L K E A E N A K R E G E T R I R R N A E 211 225
$. \\$ at gcctaaaccttt gagtaat gaaataa agt catt gtcttt tattat tatat agat at tt

ttcttaatatcaaaattccccaaattgactattcctagaatattaatcattagattcaga
$\verb tttaaaacagcatttcctacctaaggcccaaaatgtgcagtttccacctgtgtttctcca \\$
$\tt gtcctctagccccaaaagaagatataagatagaaagtttgtataaaccaaattctgctt$
aatt

Exon 9 Start: 35383 End: 35510 Length: 127
671 681 691 701 711 721
GATTTTTGCAAGGAATATGAAAAACAAGTGAGAAATGGAAGGCTTTTTTGTACACGGGAG
D F C K E Y E K Q V R N G R L F C T R E 231 241
731 741 751 761 771 781
AGTGATCCAGTCCGTGGCCCTGACGGCAGGATGCATGGCAACAAATGTGCCCTGTGTGCT S D P V R G P D G R M H G N K C A L C A
251 261

tgattggattgatgagtaaaaataactttgaaaggaagctttgttgttgagaaccatctg
ctagtata

Exon 10 Start: 36847 End: 36934 Length: 87
801 811 821 831 841 851 842 831 842 831 842 832 832 833 833 834 835
861 871 881

aaacattgtctgaattcaaagtgtgata

Exon 11 Start: 38896 End: 39023 Length: 127
891 901 911 921 931 941
AAACTCTGCAGTCAATATCAAAATCAGGCAAAGAATGGAATACTTTTCTGTACCAGAGAA
K L C S Q Y Q N Q A K N G I L F C T R E
951 961 971 981 991 100
N D P I R G P D G K M H G N L C S M C Q
321 331
GCCTACTTgtgagtatagagttttagaatgtcaaagaagggatcttgcaggtaatt
A Y F

ttagcaat

Exon 12 Start: 40263 End: 40344 Length: 81
1011 1021 1031 1041 1051 1061 CCAAGCAGAAAATGAAGAAAAGGAAAAAGCCTGAAGCACGAGCTAGAAACAAAAGAGAATC Q A E N E E K K K A E A R A R N K R E S 341 351
1071 1081 1091

ttccagagaaattctacatgta

Exon 13 Start: 41483 End: 41610 Length: 127
1101 1111 1121 1131 1141 1151 GAGCTTTGCAGTGAATATCGAAAGCTTGTGAGGAACGGAAAACTTGCTTG
1161 1171 1181 1191 1201 1211 AACGATCCTATCCAGGGCCCAGATGGGAAAGTGCATGGCAACACCTGCTCCATGTGTGAG N D P I Q G P D G K V H G N T C S M C E 391 401

$\hbox{\tt tcatttccatgttcaattatgggagggccacttcaacataaaaatgaaagaattaaggca}$
agtgggcc

Exon 14 Start: 42384 End: 42465 Length: 81 BE AWARE: Flanking intron is shared with the following exon
1221 1231 1241 1251 1261 1271
CCAAGCAGAAGAAGAAAAGAAAGAAGGAAGGTAAATCAAGAAACAAAAGACAATC
Q A E E E E K K K E G K S R N K R Q S 411 421
1281 1291 1301
${\tt TAAGAGTACAGCTTCCTTTGAGgtgagtttatatcctccagcaactcagagggatatggc}$
K S T A S F E 431

gactatttcttggt

Exon 15 Start: 42810 End: 42937 Length: 127 BE AWARE: Flanking intron is shared with the previous exon
1311
1371 1381 1391 1401 1411 1421 CATCCAGGGCCCAGATGGAAAAATGCATGGCAACACCTGCTCCATGTGTGAGGCCTTCTT

	•		•		•							
at'	tatgat	ataaa	gagat	gcaga	ataag	ccaga	cacaga	aaaaa	catata	aatgt	gtgat	ζt
Ct	$\verb ctcttatatgtgggatcta a a a a a a a gcta a ctcatagtag aggatggtgctta ca $											

Exon 16 Start: 45981 End: 46029 Length: 48
1431 1441 1451 1461 1471 TCAACAAGAAGAAGAGCAAGAGCAAAGGCTAAAAGAGAGCTGCAAAGgtaatattctc Q Q E E R A R A K A K R E A A K 481 491

•		•	•	•	•	•	•			•	•
ctgaaca	aaaag	ggtcc	tgata	agaag	tgtcc	agagt	accaa	tttta	tttca	caatt	att
caaattt											
	0 00				0 00	Ü		Ü			
	•										
catattt	ggctt	llatt	agatt	cccgc	igita	gagta	llaac	rrate	SCC		

Exon 17 Start: 48066 End: 48193 Length: 127
1481 1491 1501 1511 1521 1531 GAAATCTGCAGTGAATTTCGGGACCAAGTGAGGAATGGAACACTTATATGCACCAGGGAG E I C S E F R D Q V R N G T L I C T R E 501 511
1541 1551 1561 1571 1581 1591 CATAATCCTGTCCGTGGCCCAGATGGCAAAATGCATGGAAACAAGTGTGCCATGTGTGCC H N P V R G P D G K M H G N K C A M C A 521 531

tgagtctcagatccttcatgcatgtgtagagtatagaccgtgagttatatattagaaagg
tggattgg

Exon 18 Start: 49782 End: 49866 Length: 84
1611 1621 1631 1641 1651 1661 CAAACTTGAAGAAGAAGAGAAAAATGATAAAGAAGAAAAAGGGAAAGTCGAGGCTGA K L E E E E K K N D K E E K G K V E A E 541 551
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$

•	•	•	•	•	•	•	•	•	•	•	•
gtttctttg	gcaaac	tggga	aaatg	gtgttt	ggato	cccat	ataca	agagt	actco	ccttt	c
	_										
	•	•	•	•			•		•		
cttattca	atttat	tcaca	ttaaa	tctat	taatt	gctcg	tgcag	gagag	gagato	aatca	эc
									, 0		
•	•	•	•	•	•	•	•		•	•	•
cattttct	ccctg	gattgt	gtcaa	ittata	ttacc	taaga	aagaa	agtata	tatag	attgt	Ξŧ
•	•	•	•	•							
atcttact	gtctga	atgca	gacta	ıa							

Exc	n	19		St	art	t:	527	97]	End:	: 5	529	24	1	Len	gth	: 1:	27				
tgt	at	ac	cad	cag	gatt	tcc	ttt	acc	aa [.]	ttta	aag	gag	gc	cat	aca	.ttt	ttc	ttct	ct	gtai	tctt	
tca	ıca	gtį	gco	cta	ıgtg	gca	.gaa	taa	ta [.]	tttg	ga c	cct	gaa	agg	ttt	tct	aat	gaga	ataa	aata	aaca	
gca	ıat	cca	atg	gtt	cca	aaa	.tgt	gtg	ac	ctag	gtt	cct	ctį	gaa	aag	cat	ttt;	ggtt	act	tat	caac	
ctt	ca	gca	ato	cad	ctt	ttt	cca	tgc	aa [.]	tcat	tta	atg	;tti	tta	gtt	ttt	aaa	gcag	gtat	ttgʻ	tgga	
gga	ıag	ati	tto	cta	ıgtg	gtt	tag	;tta	.ttį	ggad	cto	ctt	aaa	aac	ctg	ctt	ctg	ctt	ati	ttgį	gcag	•
GAG E			GC/	170 AGT	GA		TCG R	TCA	TT	ATGT	ΓG <i>I</i>	AGG	AA'	ГGG	ACG	ACT	CCC P	CTGT	TAC(1 AGAG E	751
		TC(P	CTA	176 ATT	_	GGG G	TCT L	771 AGA D	TG	GGA/	۸A <i>I</i>	ATC			CAA	CAC	CTG(M		1 ΓGAA E	811
GCC	тт	CT:	Γgt	tga	igte	ggg	cgg	cag	cca	acte	gct	gc	ta	ctg	agt	gtg	gga	gaag	gato	cago	catc	

gggtgggcaagaggggtgacattggaagttttctccaggagatagat
taggtcac

Exon 20 Start: 53897 End: 53963 Length: 66
1821 1831 1841 1851 1861 1871 CCAGCAAGAAGCAAAAGAAAAGAAAGAAGCTGAACCCAGAGCAAAAGTCAAAAGAGAAGC Q Q E A K E K E R A E P R A K V K R E A 611 621

tagtccaggggttgagaaactttttcttaacagaacaga
acggctg

Exon 21 Start: 55391 End: 55518 Length: 127
1891 1901 1911 1921 1931 1941 GAGACATGCGATGAATTTCGGAGACTTTTTGCAAAATGGAAAACTTTTCTGCACAAGAGAA E T C D E F R R L L Q N G K L F C T R E 631 641
1951 1961 1971 1981 1991 2001 AATGATCCTGTGCGTGGCCAGATGGCAAGACCCATGGCAACAAGTGTGCCATGTGTAAG N D P V R G P D G K T H G N K C A M C K 651 661
2011

tgagccaggcaaataatatgtattgtgtttgtcctttccttacatgtaatagaaacagaa
tctctgga

Exon 22 Start: 57399 End: 57495 Length: 96
2021 2031 2041 2051 2061 2071 CCAGAAAGAAATGAGGAAAGAAGAAGAAGAAGAAGAAGAAGA
2081 2091 2101 2111 ACATGGTTCCAGTGGTGGAGGAGGAAACACTCAGgtgagagcaacctctaatttcag H G S S G G G G N T Q 701

taaaggcagtgctaagtccttgagagaacagagaagtttctccttaaggttgcagggtaa
tcatatcaataatcgttctcatcagtatcacatgctc

Exon 23 Start: 59466 End: 59593 Length: 127 BE AWARE: Flanking intron is shared with the following exon
2121 2131 2141 2151 2161 2171 GACGAATGTGCTGAGTATCGGGAACAAATGAAAAATGGAAGACTCAGCTGTACTCGGGAG D E C A E Y R E Q M K N G R L S C T R E 711 721
2181 2191 2201 2211 2221 2231 AGTGATCCTGTACGTGATGCTGATGGCAAATCGTACAACAATCAGTGTACCATGTGTAAA
S D P V R D A D G K S Y N N Q C T M C K 731 741

•	•	•	•	•	•	•	•	•	•	•	•
atgaat	ggctat	tttctc	cattte	gctate	gctco	ttcca	tgcaa	attat	tcttt	ttgct	aat
				•					•	•	•
ttccaa	atatt	ctattt	ttttc	tcttg	gcgttc	ctctaa	ıggaad	caatga	gcctt	agcaa	ıggg
				•							
cagcta	accaa	gtggct	gattt	ctatt	gtatt	ttcta	ıt				

Exon 24 Start: 60015 End: 60087 Length: 72 BE AWARE: Flanking intron is shared with the previous exon
2271
$. \qquad . \\ tgactctcactgataacttgttcagtgaagaaacctaatgccaagtggacatctttaagt$

 $\verb|tcaaatttggccttttaaccttatttcgtagatagggaaaatga|\\$

Exon 25 Start: 61038 End: 61165 Length: 127 BE AWARE: Flanking intron is shared with the following exon
aacctttgtgattacatgggccccacttggataatccagggtaacttccccattttaaat
2321 2331 2341 2351 2361 2371
GATACATGTGATGAGTTTAGAAGCCAAATGAAAAATGGAAAACTCATCTGCACTCGAGAA D T C D E F R S Q M K N G K L I C T R E 781 791
2381 2391 2401 2411 2421 2431
AGTGACCCTGTCCGGGGTCCAGATGGCAAGACACATGGCAATAAGTGTACTATGTGTAAG S D P V R G P D G K T H G N K C T M C K
801 811
$\begin{tabular}{lllllllllllllllllllllllllllllllllll$
catttcttaccagtttgggaaaatga

Exon 26 Start: 61324 End: 61420 Length: 96 BE AWARE: Flanking intron is shared with the previous exon	
2451 2461 2471 2 tttcattgttttccccccagGGAAAGGGAAGCAGCTGAAAAAAAAAAAAAAAAAAAAAA	2481
2491 2501 2511 2521 2531 GACAGGAGCAATACAGGAGAAAGGAGCAATGACAAAGAGGTA D R S N T G E R S N D K E 831 841	

Exon 27 S	Start: 64862	! End: 6	4989 Leng	gth: 127	
atatatatata					
	atatatataatc	ttgcttgca	atatatatata	atatagttttta	aaaggaat
gggctgcttta					
	agtgaaattcca	ttagggcaa	gacaatcatt	tgatagagatgo	caatggag
aagattcaaad					
	cttcttaggtag	agattgaat	aaagtgcctt	taaaacatatt1	ccttctct
gaaatttcact					
	ttctctgttttt	ttcctgtgt	tatgagttta	tatctaatcggt	ccaatcat
gttatcaggti					
	ttgaaagattat	accatgaca	gtaacaactt	tttctgctactg	gttggtag
2541 GATCTGTGTCC D L C R	GTGAATTTCGAA	2561 GCATGCAGA M Q R	GAAATGGAAA(GCTTATCTGCAC	
2601 AATAACCCTGT N N P V	2611 TTCGAGGCCCAT R G P Y 871	2621 ATGGCAAGA G K M			
2661 AGCATCTTgta	acgtaaaaaggt	 ttatcaata	aatttgatag	· ttgtgcctgtt1	.gctagaa

actgaaca

Exon 28 Start: 65794 End: 65866 Length: 72	
	tttc
	ıttat
	tttt
	;ttaa
	gcag
$\tt TGATCGAGAAGCTAATGAAAGAAAAAAAAAGAAAGATGAAGAAAATCAAGTAGCAAGCA$	2721 CCCTC S
2731	cttc:
	tcaa

 aactcaactgcat

Exon 29 Start: 66752 End: 66879 Length: 127
2741 2751 2761 2771 2781 2791 GATGAGTGCAGTGAATTTCGAAACTATATAAGGAACAATGAACTCATCTGCCCTAGAGAG D E C S E F R N Y I R N N E L I C P R E 921 931
2801 2811 2821 2831 2841 2851 AATGACCCAGTGCACGGTGCTGATGGAAAGTTCTATACAAACAA
2861

agaaactgctgcttgaataagtttgtatatttatgaaccccatgggattttaaaaataag
acaactag

Exon 30 Start: 68012 End: 68108 Length: 96
2871 2881 2891 2901 2911 2921 TCTAACAGAAGCTTTGGAAAGGGCAAAGCTTCAAGAAAAGCCATCCCATGTTAGAGCTTC L T E A L E R A K L Q E K P S H V R A S 961 971
2931 2941 2951 2961 TCAAGAGGAAGACAGCCCAGACTCTTTCAGTTCTCTGgtaaggaggactatttctgaaaa Q E E D S P D S F S S L 981

gtgactaatttcacatatggtaaaatacatagtcagg

Exc	on (31 S	tart	5: 7	'228	38	E:	nd:	72	418		Len	gth:	1	30			
aga	att [.]	 tcttct	tate	gggc	agg	gtaa	att	gtg	tca	cata	att	ata	aagg	gata	att ¹	tgc	ttt	ctta
ttį	gct [.]	 ttgcta	ctco	cttg	ttt	tctt	ctg	ttt	act	tac	ctt	gtc:	aagt	ta	aaa	att	gcc	aaat
aca	agg	 aagtta	aato	ccac	cto	cata	atg	gtg	caa	aat	caa	tct	ttga	ngt	ttg	aat	aac	atat
ato	cac	 actcct	ttta	acaa	LCC8	attt	tat	tcaa	agt	tgga	att	aagį	gaad	ctc	aaga	agg	ttt	tctt
aag	gcc	 cacccc	tctt	cctt	gaa	atgo	cca	taa	agt	acg	tct	gct [.]	ttat	;tt	tttį	gct	tct	tcag
		2971 TGAGAT E M 991	GTG		GAC		CCG	AGT	ATT	GCC	CAG R	GAT		TA'	TCT"	TTG'	TCC	
		3031 AAAGCC K P 1011	TGT(V	CTGT	'GG'	ΓGΑC	CGA	TGG	CCA	AAC	CTA Y	CAA	CAAT	CC'	TTG	CAT	GCT	3081 CTGT C
	E	3091 AAACCT N L	gtaa	agta	ttc	caag	gtt	gcc	cca	tcai	tat	ctt	ccag	gtt	taga	aat	ttc	tcag

ctagagtgttaacccatagtaatgcactgatataaattcgaaatgtgttgcaagtaattt
tgtacagtcat

Exon 32 Start: 74826 End: 74916 Length: 90
3101
3161 3171 3181
gttacgttgtgaggagcactgggttctggttttgttctcttccactgagtaatggacatt

${\tt aaggataattaagtaattactgaaataagtggaaataatgtagcagttaatattttaaaa}$
${\tt tatttaatgggtccattaatagattgtttatgagcaatgtctccaacataacgttactta}$
tttcaggataggttttgttttctgtgatcta

Exon 33 Start: 78012 End: 78391 Length: 379
3191 *1 *11 *21 *31 *41 *51 GACGAATGACAGGAAGATTGTTGAAAGCCATGAGGGAAAAAATAAACCCCAGTTCTGAAT D E *
*61 *71 *81 *91 *101 *111 CACCTACCTTCACCATCTGTATATACAAAGAATTCTTCGGAGCTTGTCTTATTTGCTATA
*121 *131 *141 *151 *161 *171 GAAAACAATACAGAGCTTTTGGGAATGGACTCACTGATTTTCAGTCTTTTCCATCTCTTT

*181 *191 *201 *211 *221 *231 *201 *211 *221 *231
*241 *251 *261 *271 *281 *291 ATGTCCTGATTACAATGCTGTCTGTCCAACTGCCTGTTCAATAAAAGTAAACTCAGCAGA
*301 *311 *321 *331 *341 *351 ACACCCTTTCTGGGATTTCTTTGTCACTATCTGGATAATAGATATTTGCTTTTAAAGAAA
*361 *371
tattcattaagttacctcct

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RerenceTypeSetter: Version: 2.0, Version Date: 06/08/2020