

## U6B-sgRNA2.0 map

5' CACCTAAATTGTAAGCGTTAATATTTTGTAAAATTCGCGTTAAATTTTTGTAAATCAGCTCATTTTTT  
 0 ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | 70  
 3' GTGGATTTAACATTCGCAATTATAAAACAATTTTAAAGCGCAATTTAAAAACAATTTAGTCGAGTAAAAAA  
 1 H L N C K R . Y F V K I R V K F L L N Q L I F  
 2 T . I V S V N I L L K F A L N F C . I S S F F  
 3 P P K L . A L I F C . N S R . I F V K S A H F L  
 0

5' AACCAATAGGCCGAAATCGGCAAAATCCCTTATAAATCAAAGAATAGACCGAGATAGGGTTGAGTGTG  
 0 ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | 140  
 3' TTGGTTATCCGGCTTTAGCCGTTTTAGGGAATATTTAGTTTTCTTATCTGGCTCTATCCCAACTCACAAC  
 1 . P I G R N R Q N P L . I K R I D R D R V E C C  
 2 N Q . A E I G K I P Y K S K E . T E I G L S V  
 3 T N R P K S A K S L I N Q K N R P R . G . V L  
 0

5' TTCCAGTTTGGAACAAGAGTCCACTATTAAGAACGTGGACTCCAACGTCAAAGGGCGAAAAACCGTCTA  
 0 ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | 210  
 3' AAGGTCAAACCTTGTCTCAGGTGATAATTTCTTGCACCTGAGGTTGCAGTTTCCCGCTTTTTGGCAGAT  
 1 S S L E Q E S T I K E R G L Q R Q R A K N R L  
 2 V P V W N K S P L L K N V D S N V K G R K T V Y  
 3 F Q F G T R V H Y . R T W T P T S K G E K P S  
 0

5' TCAGGGCGATGGCCCACTACGTGAACCATCACCTAATCAAGTTTTTTGGGGTCGAGGTGCCGTAAAGCA  
 0 ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | 280  
 3' AGTCCCGCTACCGGGTGATGCACTTGGTAGTGGGATTAGTTCAAAAAACCCAGCTCCACGGCATTTCGT  
 1 S G R W P T T . T I T L I K F F G V E V P . S  
 2 Q G D G P L R E P S P . S S F L G S R C R K A  
 3 I R A M A H Y V N H H P N Q V F W G R G A V K H  
 0

5' CTAAATCGGAACCCTAAAGGAGCCCCGATTTAGAGCTTGACGGGAAAGCCGGCGAACGTGGCGAGAA  
 0 ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | 350  
 3' GATTTAGCCTTGGGATTTCCCTCGGGGGCTAAATCTCGAACTGCCCTTTCGGCCGCTTGCACCGCTCTT  
 1 T K S E P . R E P P I . S A L T G K A G E R G E K  
 2 L N R N P K G S P P F R A . R G K P A N V A R  
 3 . I G T L K G A P D L E L D G E S R R T W R E  
 0

5' AGGAAGGGAAGAAAGCGAAAGGAGCGGGCGCTAGGGCGCTGGCAAGTGTAGCGGTCACGCTGCGCGTAAC  
 0 ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | 420  
 3' TCCTTCCCTTCTTTCGCTTTTCCCTCGCCCGGATCCCGCGACCGTTCACATCGCCAGTGCACGCGCATTG  
 1 G R E E S E R S G R . G A G K C S G H A A R N  
 2 K E G K K A K G A G A R A L A S V A V T L R V T  
 3 R K G R K R K E R A L G R W Q V . R S R C A .  
 0

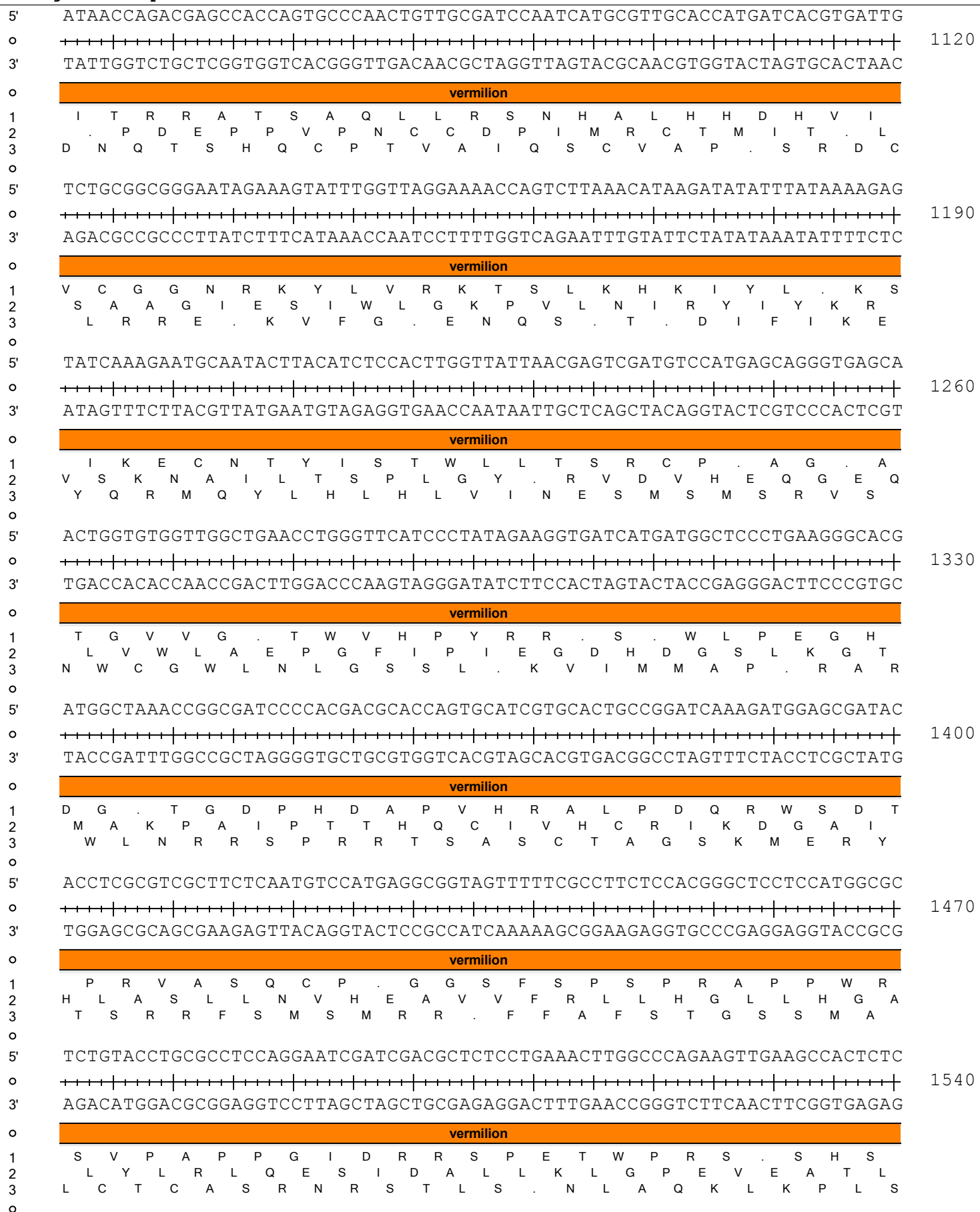
5' CACCACACCCGCGCGCTTAATGCGCCGCTACAGGGCGCGTCCCATTCGCCATTCAGGCTGCGCAACTGT  
 0 ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | 490  
 3' GTGGTGTGGGCGGCGGAATTACGCGGCGATGTCCCGCGCAGGGTAAGCGGTAAGTCCGACGCGTTGACA  
 1 H H T R R A . C A A T G R V P F A I Q A A Q L  
 2 T T P A A L N A P L Q G A S H S P F R L R N C  
 3 P P H P P R L M R R Y R A R P I R H S G C A T V  
 0

5' TGGGAAGGGCGATCGGTGCGGGCCTCTTCGCTATTACGCCAGCTGGCGAAAGGGGGATGTGCTGCAAGGC  
 0 ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | ++++++ | 560  
 3' ACCCTTCCCGCTAGCCACGCCCGGAGAAGCGATAATGCGGTCGACCGCTTTCCTCCCTACACGACGTTCCG  
 1 L G R A I G A G L F A I T P A G E R G M C C K A  
 2 W E G R S V R A S S L L R Q L A K G G C A A R  
 3 G K G D R C G P L R Y Y A S W R K G D V L Q G  
 0

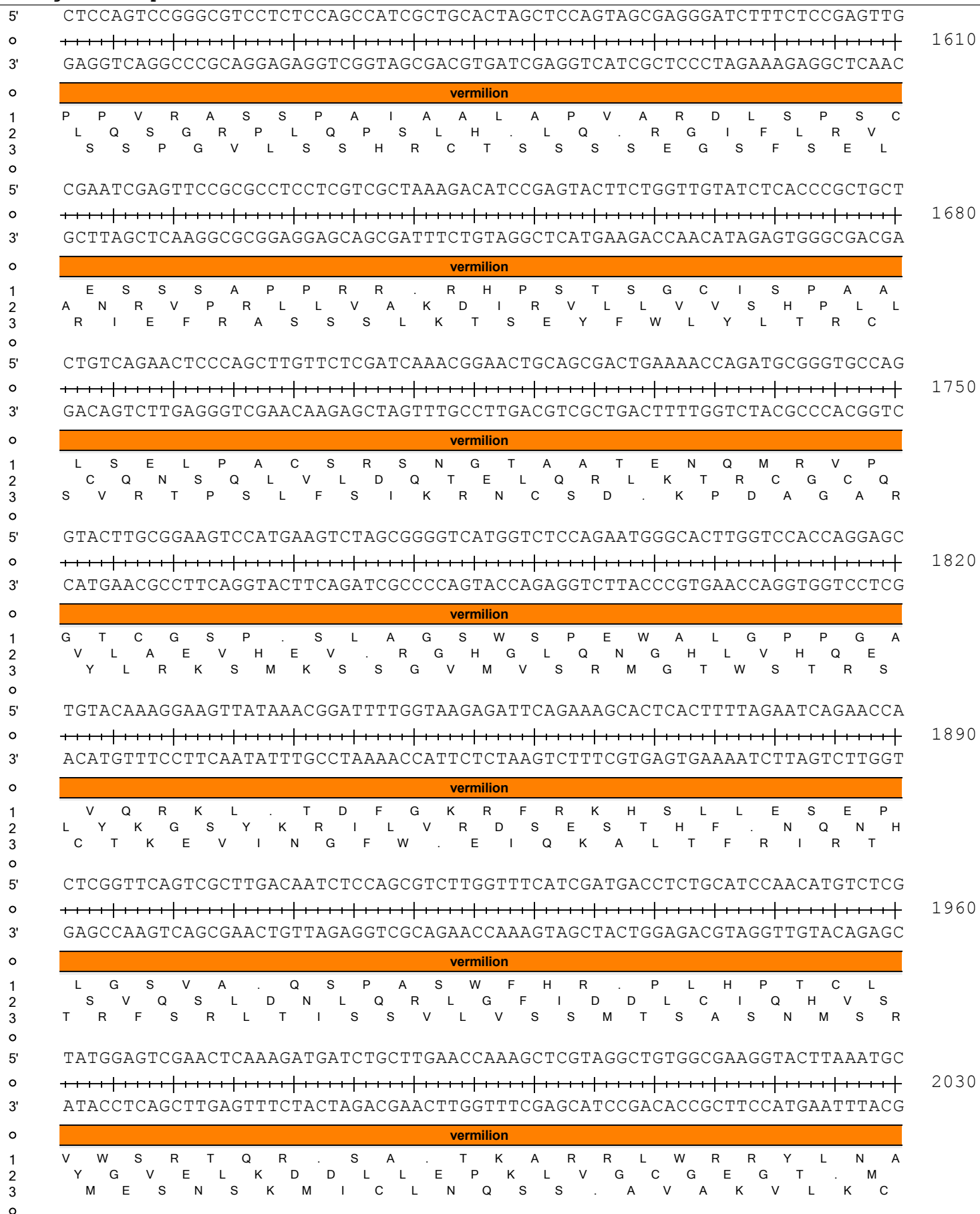
## U6B-sgRNA2.0 map

5'	GATTAAGTTGGGTAACGCCAGGGTTTTCCAGTCACGACGTTGTAAAACGACGGCCAGTGAATTGTAATA	
0	+++++	630
3'	CTAATTCAACCCATTGCGGTCCCAAAGGGTCAGTGCTGCAACATTTTGCTGCCGGTCACTTAACATTAT	
1	I K L G N A R V F P V T T L . N D G Q . I V I	
2	R L S W V T P G F S Q S R R C K T T A S E L . Y	
3	D . V G . R Q G F P S H D V V K R R P V N C N	
0		
5'	C G A C T C A C T A T A G G G C G A A T T G G G T A C A A G C T T A T T T A T T T T G T T A T G T T A T A T G T A T T A T A T G T C A G A C	
0	+++++	700
3'	GCTGAGTGATATCCCGCTTAACCCATGTTCTGAATAAAATAAAACAATAACAATATACATAATATACAGTCTG	
0		
	<b>vermilion</b>	
1	R L T I G R I G Y K L I Y F V M L Y V L Y V R	
2	D S L . G E L G T S L F I L L C Y M Y Y M S D	
3	T T H Y R A N W V Q A Y L F C Y V I C I I C Q T	
0		
5'	A T A A A G A A A A G G A A C A C A T C A A A T G T G A T A A C A A A G A C T A A A C A A G T A A T T T T A T T A C A C C A A A A C G A C A	
0	+++++	770
3'	T A T T T C T T T T C C T T G T G T A G T T T A C A C T A T T G T T T C T G A T T T G T T C A T T A A A A T A A T G T G G T T T T G C T G T	
0		
	<b>vermilion</b>	
1	H K E K E H I K C D N K D . T S N F I T P K R Q	
2	I K K R N T S N V I T K T K Q V I L L H Q N D	
3	. R K G T H Q M . . Q R L N K . F Y Y T K T T	
0		
5'	A A A C A G T A G G C A G A A C A A A C A C G C A T A G C C A A A C A T T G A C G A A T T G G A T A C C C T G C C G A T T G T C A G A C A	
0	+++++	840
3'	T T T G T C A T C C G T C T T G T T T G T T G C G T A T C G G T T T G T A A C T G C T T A A C C T A T G G G A C G G C T A A C A G T C T G T	
0		
	<b>vermilion</b>	
1	N S R Q N K Q R I A K H . R I G Y P A D C Q T	
2	K T V G R T N N A . P N I D E L D T L P I V R H	
3	K Q . A E Q T T H S Q T L T N W I P C R L S D	
0		
5'	C T T T T G T T G A T C A G T T T C T T G C G A A T G G T C T C G T C C A G C G G T G G A A T C G C C T C G C G G G A A T C A G A A A G	
0	+++++	910
3'	G A A A C A A C T A G T C A A A G A A C G C T T A C C A G A G C A G G T C G C C A C C T T A G C G G A G C G C C C C T T A G T C T T T T C	
0		
	<b>vermilion</b>	
1	L L L I S F L R M V S S S G G I A S R G I R K	
2	F C . S V S C E W S R P A V E S P R G E S E K	
3	T F V D Q F L A N G L V Q R W N R L A G N Q K S	
0		
5'	T G G A C A G A T T G A A C A G A T C C A G A A A C A C C T T G T A C C G A T C A C T G A A A C C A A A A A A A A C A A A G G G A G A A C	
0	+++++	980
3'	A C C T G T C T A A C T T G T C T A G G T C T T T G T G G A A C A T G G C T A G T G A C T T T G G T T T T T T T T T T G T T T C C C T C T T G	
0		
	<b>vermilion</b>	
1	V D R L N R S R N T P L Y R S L K P K K N K G R T	
2	W T D . T D P E T P C T D H . N Q K K T K G E	
3	G Q I E Q I Q K H L V P I T E T K K K Q R E N	
0		
5'	A G T T T G A G T T C A T T G A T C C C C G A T A T A A T C A C A T C T G C G A T G A T C A C C T G A G A G T G G A G C G C A G A T A T T G	
0	+++++	1050
3'	T C A A A C T C A A G T A A C T A G G G C T A T A T T A G T G T A G A C G C T A C T A G T G G A C T C T C A C C T C G C G T C T A T A A C	
0		
	<b>vermilion</b>	
1	V . V H . S P I . S H L R . S P E S G A Q I L	
2	Q F E F I D P R Y N H I C D D H L R V E R R Y .	
3	S L S S L I P D I I T S A M I T . E W S A D I	
0		

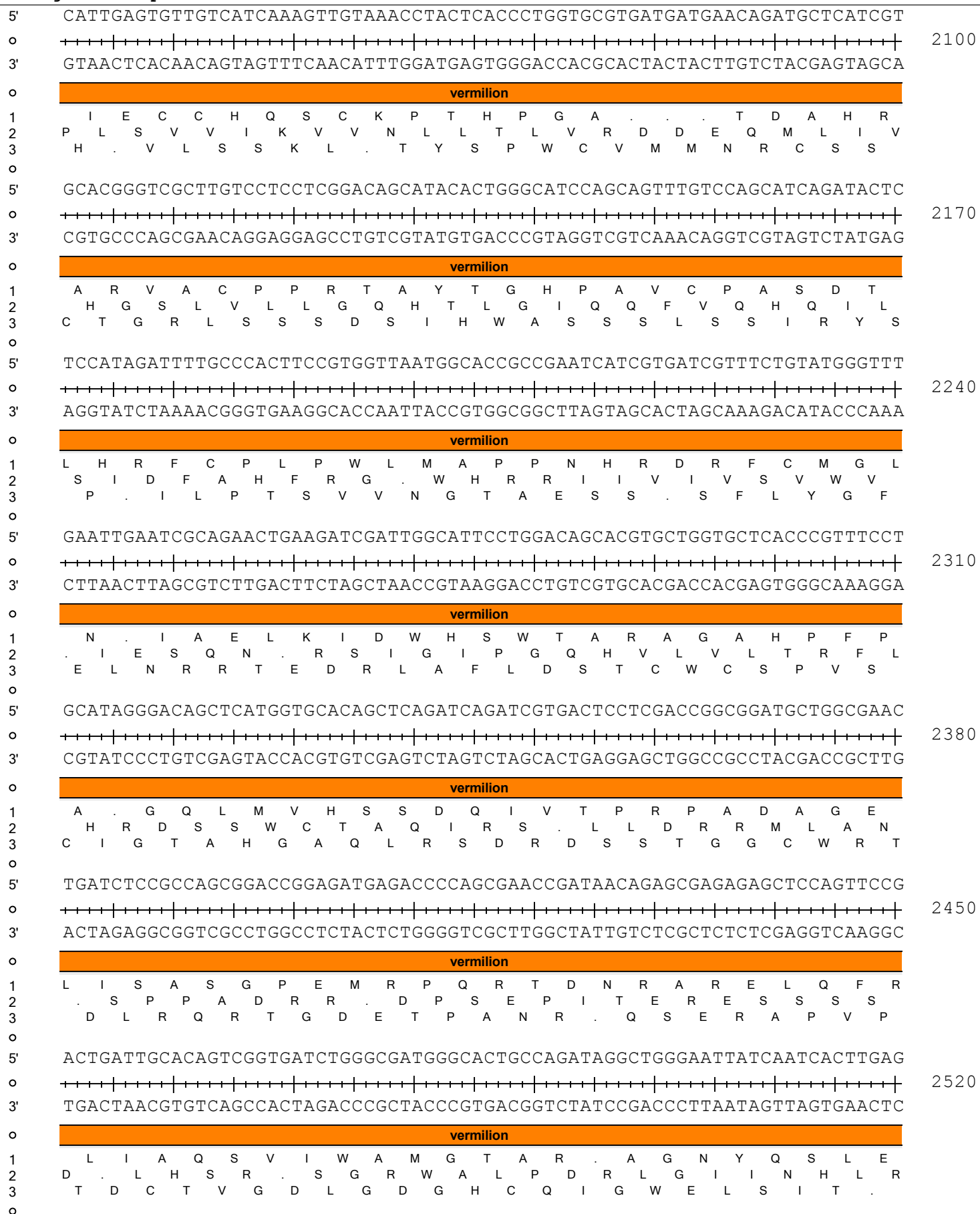
## U6B-sgRNA2.0 map



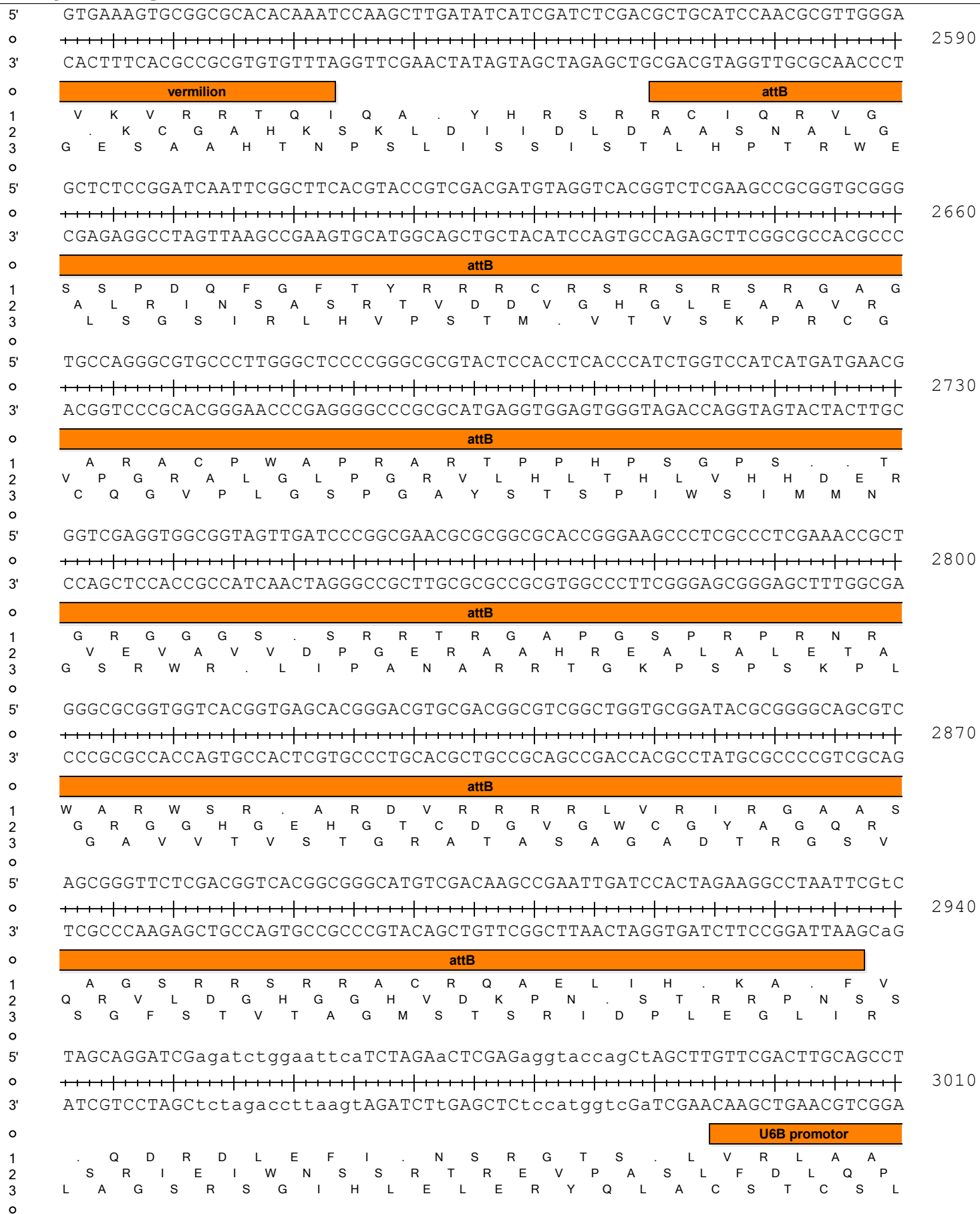
## U6B-sgRNA2.0 map



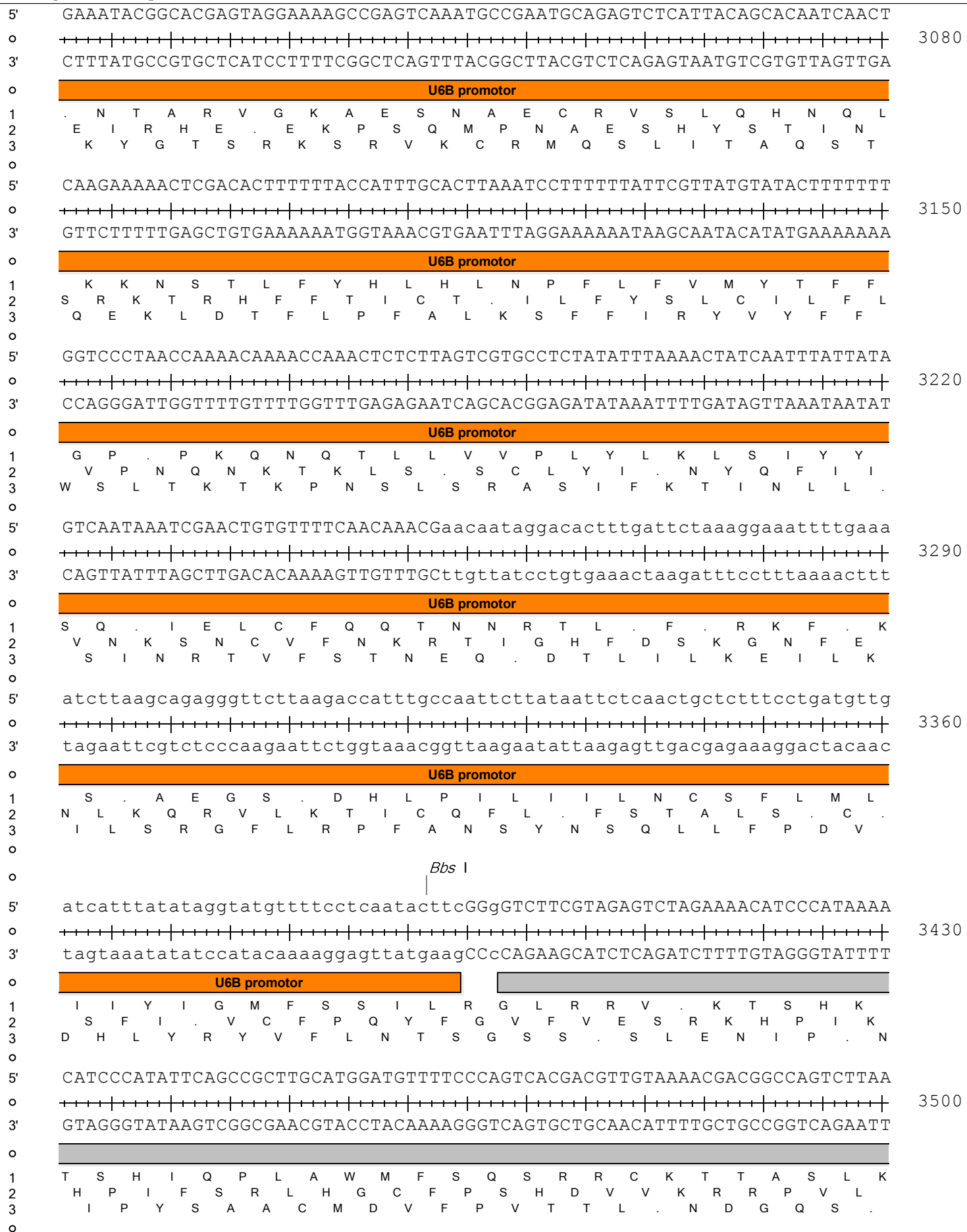
## U6B-sgRNA2.0 map



## U6B-sgRNA2.0 map



U6B-sgRNA2.0 map



## U6B-sgRNA2.0 map

5' GCTCGGGCCCCAAATAATGATTTTTATTTTACTGATAGTGACCTGTTTCGTTGCAACAAATTGATGAGCAA  
o ++++++  
3' CGAGCCCGGGGTTTATTACTAAAATAAACTGACTATCACTGGACAAGCAACGTTGTTTAACTACTCGTT  
o  
1 L G P Q I M I L F L I V T C S L Q Q I D E Q  
2 S S G P K . F Y F D . . P V R C N K L M S N  
3 A R A P N N D F I L T D S D L F V A T N . . A  
o  
5' TGCTTTTTTATAATGCCAACTTTGTACAAAAAGCAGGCTCCGCGGCCGCCCTTCACCCTAGAGGAG  
o ++++++  
3' ACGAAAAATATTACGGTTGAAACATGTTTTTTCGTCGGAGGCGCCGGCGGGGAAGTGGTGATCTCCTC  
o  
1 C F F I M P T L Y K K A G S A A A P F T T R G  
2 A F L . C Q L C T Y K K A Q A P R P P S P L E E  
3 M L F Y N A N F V Q K S R L R G R P L H H . R R  
o  
5' AGCAACTGCATAAGGCTATGAAGAGATACGCCCTGGTTCCTGGAACAATTAATTGCTTTTACAGATGCAC  
o ++++++  
3' TCGTTGACGTATTCCGATACTTCTCTATGCGGGACCAAGGACCTTGTTAATTAACGAAAATGTCTACGTG  
o  
1 E Q L H K A M K R Y A L V P G T I N C F Y R C T  
2 S N C I R L . R D T P W F L E Q L I A F T D A  
3 A T A . G Y E E I R P G S W N N . L L L Q M H  
o  
5' ATATCGAGGTGGACATCACTTACGCTGAGTACTTCGAAATGTCCGTTTCGGTTGGCAGAAGCTATGAAACG  
o ++++++  
3' TATAGCTCCACCTGTAGTGAATGCGACTCATGAAGCTTTACAGGCAAGCCAACCGTCTTCGATACTTTGC  
o  
1 Y R G G H H L R . V L R N V R S V G R S Y E T  
2 H I E V D I T Y A E Y F E M S V R L A E A M K R  
3 I S R W T S L T L S T S K C P F G W Q K L . N  
o  
5' ATATGGGCTGAATACAAATCACAGAATCGTCGTATGCAGTGAAAACCTCTTCAATTCTTTATGCCGGTG  
o ++++++  
3' TATACCCGACTTATGTTTAGTGTCTTAGCAGCATAACGTCACCTTTTGAGAGAAGTTAAGAAATACGGCCAC  
o  
1 I W A E Y K S Q N R R M Q . K L S S I L Y A G  
2 Y G L N T N H R I V V C S E N S L Q F F M P V  
3 D M G . I Q I T E S S Y A V K T L F N S L C R C  
o  
5' TTGGGCGCGTTATTTATCGGAGTTGCAGTTGCGCCCGCGAACGACATTTATAATGAACGTGAATTGCTCA  
o ++++++  
3' AACCCGCGCAATAAATAGCCTCAACGTCAACGCGGGCGCTTGCTGTAATATTACTTGCACCTAACGAGT  
o  
1 V G R V I Y R S C S C A R E R H L . . T . I A Q  
2 L G A L F I G V A V A P A N D I Y N E R E L L  
3 W A R Y L S E L Q L R P R T T F I M N V N C S  
o  
5' ACAGTATGGGCATTTTCGAGCCTACCGTGGTGTTCGTTTCCAAAAGGGGTGCAAAAATTTTGAACGT  
o ++++++  
3' TGTCATACCCGTAAAGCGTCGGATGGCACCACAAGCAAAGGTTTTTCCCAACGTTTTTTAAACTTGCA  
o  
1 Q Y G H F A A Y R G V R F Q K G V A K N F E R  
2 N S M G I S Q P T V V F V S K K G L Q K I L N V  
3 T V W A F R S L P W C S F P K R G C K K F . T  
o



U6B-sgRNA2.0 map

5' GCAAAAAAAGCTCCCAATCATCCAAAAAATTATTATCATGGATTCTAAAACGGATTACCAGGGATTTCAG  
 0 ++++++  
 3' CGTTTTTTTCGAGGGTTAGTAGGTTTTTAATAATAGTACCTAAGATTTTGCCTAATGGTCCCTAAAGTC  
 0  
 1 A K K A P N H P K N Y Y H G F . N G L P G I S  
 2 Q K K L P I I Q K I I I M D S K T D Y Q G F S Q  
 3 C K K S S Q S S K K L L S W I L K R I T R D F S

4060

Bbs I

5' TCGATGTGAATTGGAGAAGACCTGTTTTAGAGCTAGGCCAACATGAGGATCACCCATGTCTGCAGGGCCT  
 0 ++++++  
 3' AGCTACACTTAACCTCTTCTGGACAAAATCTCGATCCGGTTGTACTCCTAGTGGGTACAGACGTCCCGGA  
 0  
 1 V D V N W R R P V L E L G Q H E D H P C L Q G L  
 2 S M . I G E D L F . S . A N M R I T H V C R A  
 3 R C E L E K T C F R A R P T . G S P M S A G P

4130

gRNA core

5' AGCAAGTTAAAATAAGGCTAGTCCGTTATCAACTTGCCAACATGAGGATCACCCATGTCTGCAGGGCCA  
 0 ++++++  
 3' TCGTTCAATTTTATTCCGATCAGGCAATAGTTGAACCGGTTGTACTCCTAGTGGGTACAGACGTCCCGGT  
 0  
 1 A S . N K A S P L S T W P T . G S P M S A G P  
 2 Q V K I R L V R Y Q L G Q H E D H P C L Q G Q  
 3 S K L K . G . S V I N L A N M R I T H V C R A

4200

gRNA core

5' AGTGGCACCGAGTCGGTGCTTTTTttgctcacctgtgattgctcctactcaaatacaaaaacatcaaatt  
 0 ++++++  
 3' TCACCGTGGCTCAGCCACGAAAAaacgagtggacactaacgaggatgagtttatgtttttgtagtttaa  
 0  
 1 S G T E S V L F L L T C D C S Y S N T K T S N  
 2 V A P S R C F F C S P V I A P T Q I Q K H Q I  
 3 K W H R V G A F F A H L . L L L L K Y K N I K F

4270

gRNA core

U6B 3'UTR

5' ttctgtcaataaagcatatattttatattttttacaggaaagaattACtaGcGTAATATATAGACAA  
 0 ++++++  
 3' aagacagttatttcgtataaataaataaataaaatgctcctttcttaaTGatCgCATTATATATCTGTT  
 0  
 1 F L S I K H I Y L Y L F Y R K E L L A . Y I D N  
 2 F C Q . S I F I Y I Y F T G K N Y . R N I . T  
 3 S V N K A Y L F I F I L Q E R I T S V I Y R Q

4340

U6B 3'UTR

5' TGGTTTTCCGTTGACGTACATACATCTGACGTGTGTTTATTTAGACATAATAGTTATGTTTTACATCTT  
 0 ++++++  
 3' ACCAAAAGGCAACTGCATGTATGTAGACTGCACACAAAATAAATCTGTATTATCAATACAAAAGTGTAGAA  
 1 G F P L T Y I H L T C V Y L D I I V M F S H L  
 2 M V F R . R T Y I . R V F I . T . L C F H I F  
 3 W F S V D V H T S D V C L F R H N S Y V F T S

4410

5' TTTAATGTTTCGCTTAATGCGTATGCATACAAAATTTTAAATTTTCAACACAGTTGTTTTGTTTTCATCA  
 0 ++++++  
 3' AAATTACAAGCGAATTACGCATACGTATGTTTTAAAAATTTAAAGTTGTGTCAACAAAACAAAAGTAGT  
 1 F N V R L M R M H T K F L I F N T V V F V F I  
 2 L M F A . C V C I Q N F . F S T Q L F L F S S  
 3 F . C S L N A Y A Y K I F N F Q H S C F C F H H

4480

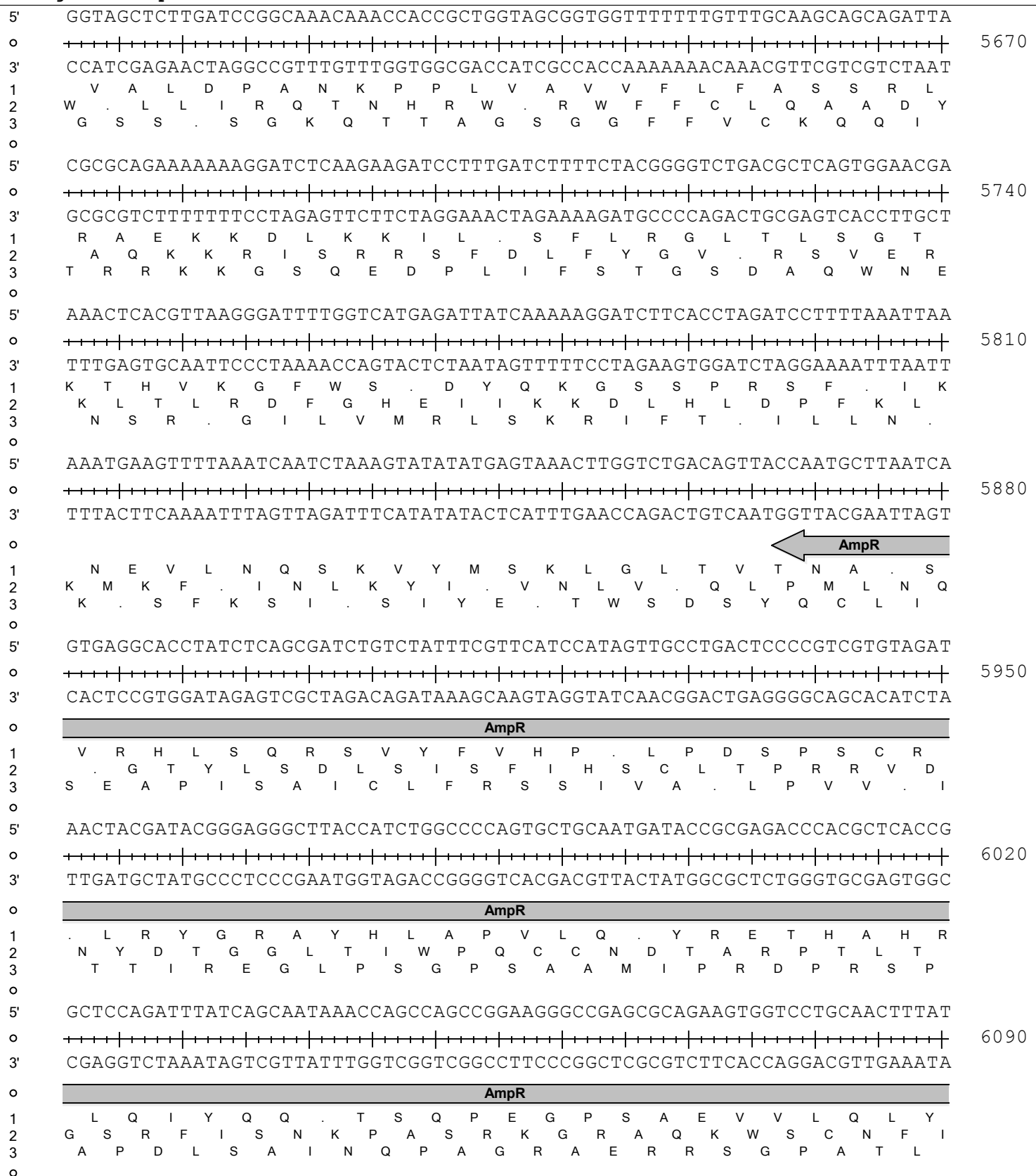
## U6B-sgRNA2.0 map

5' CtaGcGTAATATATAGACAATGGTTTTCCGTTGACGTACATACATCTGACGTGTGTTTATTTAGACATAA  
o ++++++  
4550  
3' GatCgCATTATATATCTGTTACCAAAGGCAACTGCATGTATGTAGACTGCACACAAATAAATCTGTATT  
1 T S V I Y R Q W F S V D V H T S D V C L F R H N  
2 L A . Y I D N G F P L T Y I H L T C V Y L D I  
3 . R N I . T M V F R . R T Y I . R V F I . T .  
o  
5' TAGTTATGTTTTCACATCTTTTTAATGTTTCGCTTAATGCGTATGCATACAAAATTTTTAATTTTCAACAC  
o ++++++  
4620  
3' ATCAATACAAAAGTGTAGAAAATTACAAGCGAATTACGCATACGTATGTTTTAAAATTTAAAAGTTGTG  
1 S Y V F T S F . C S L N A Y A Y K I F N F Q H  
2 I V M F S H L F N V R L M R M H T K F L I F N T  
3 . L C F H I F L M F A . C V C I Q N F . F S T  
o  
5' AGTTGTTTTTGTTCATCACTAGTGAGCTCCAGCTTTTGTTCCTTTAGTGAGGGTTAATTgCGcGCTT  
o ++++++  
4690  
3' TCAACAAAACAAAAGTAGTGATCACTCGAGGTCGAAAACAAGGAAATCACTCCAATTAACGCgCGAA  
1 S C F C F H H . . A P A F V P F S E G . L R A  
2 V V F V F I T S E L Q L L F P L V R V N C A L  
3 Q L F L F S S L V S S S F C S L . . G L I A R L  
o  
5' GCGTAATCATGGTCATAGCTGTTTCCTGTGTGAAATTGTTATCCGCTCACAATTCACACAACATACGA  
o ++++++  
4760  
3' CCGCATTAGTACCAGTATCGACAAAGGACACACTTTAACAATAGGCGAGTGTTAAGGTGTGTTGTATGCT  
1 W R N H G H S C F L C E I V I R S Q F H T T Y E  
2 G V I M V I A V S C V K L L S A H N S T Q H T  
3 A . S W S . L F P V . N C Y P L T I P H N I R  
o  
5' GCCGGAAGCATAAAGTGTAAAGCCTGGGGTGCCTAATGAGTGAGCTAACTCACATTAATTGCGTTGCGCT  
o ++++++  
4830  
3' CGGCCTTCGTATTTACATTTTCGGACCCCACGGATTACTCACTCGATTGAGTGTAATTAACGCAACGCGA  
1 P E A . S V K P G V P N E . A N S H . L R C A  
2 S R K H K V . S L G C L M S E L T H I N C V A L  
3 A G S I K C K A W G A . . V S . L T L I A L R  
o  
5' CACTGCCCCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTGCATTAATGAATCGGCCAACGCGCGGGGAG  
o ++++++  
4900  
3' GTGACGGGCGAAAAGTTCAGCCCTTTGGACAGCACGGTCGACGTAATTACTTAGCCGTTGCGCGCCCTC  
1 H C P L S S R E T C R A S C I N E S A N A R G  
2 T A R F P V G K P V V P A A L M N R P T R G E  
3 S L P A F Q S G N L S C Q L H . . I G Q R A G R  
o  
5' AGGCGGTTTGCGTATTGGGCGCTCTTCCGCTTCCTCGCTCACTGACTCGCTGCGCTCGGTTCGTTTCGGCTG  
o ++++++  
4970  
3' TCCGCCAAACGCATAACCCGCGAGAAGGCGAAGGAGCGAGTGAAGTGAAGCGACGCGAGCCAGCAAGCCGAC  
1 E A V C V L G A L P L P R S L T R C A R S F G C  
2 R R F A Y W A L F R F L A H . L A A L G R S A  
3 G G L R I G R S S A S S L T D S L R S V V R L  
o  
5' CGGCGAGCGGTATCAGCTCACTCAAAGGCGGTAATACGGTTATCCACAGAATCAGGGGATAACGCAGGAA  
o ++++++  
5040  
3' GCCGCTCGCCATAGTCGAGTGAGTTTCCGCCATTATGCCAATAGGTGTCTTAGTCCCCTATTGCGTCTCTT  
1 G E R Y Q L T Q R R . Y G Y P Q N Q G I T Q E  
2 A A S G I S S L K G G N T V I H R I R G . R R K  
3 R R A V S A H S K A V I R L S T E S G D N A G  
o

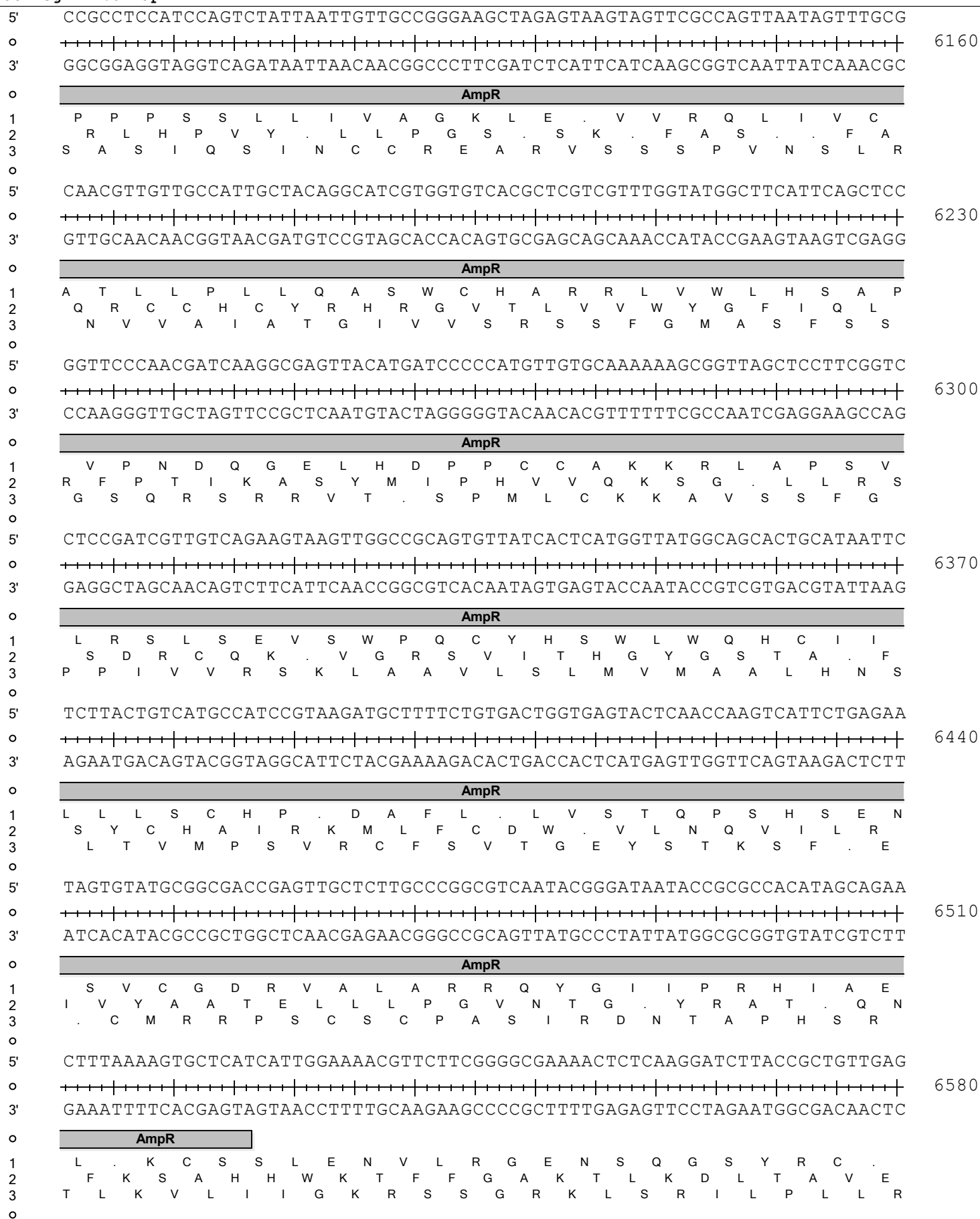
## U6B-sgRNA2.0 map

```
5' AGAACATGTGAGCAAAAAGGCCAGCAAAAAGGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTTTTTCCA
o ++++++
3' TCTTGTACACTCGTTTTCCGGTCGTTTTCCGGTCCTTGGCATTTCCTCCGGCGCAACGACCGCAAAAAGGT
1 R T C E Q K A S K R P G T V K R P R C W R F S
2 E H V S K R P A K G Q E P . K G R V A G V F P
3 K N M . A K G Q Q K A R N R K K A A L L A F F H
o
5' TAGGCTCCGCCCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGA
o ++++++
3' ATCCGAGGCGGGGGGACTGCTCGTAGTGTTTTTAGCTGCGAGTTCAGTCTCCACCGCTTTGGGCTGTCTT
1 I G S A P L T S I T K I D A Q V R G G E T R Q D
2 . A P P P . R A S Q K K S T L K S E V A K P D R
3 R L R P P D E H H K N R R S S Q R W R N P T G
o
5' CTATAAAGATAACCAGGCGTTTCCCCCTGGAAGCTCCCTCGTGCGCTCTCCTGTTCCGACCTGCCGCTTA
o ++++++
3' GATATTTCTATGGTCCGCAAAGGGGGACCTTCGAGGGAGCACGCGAGAGGACAAGGCTGGGACGGCGAAT
1 Y K D T R R F P L E A P S C A L L F R P C R L
2 T I K I P G V S P W K L P R A L S C S D P A A Y
3 L . R Y Q A F P P G S S L V R S P V P T L P L
o
5' CCGGATACCTGTCCGCCTTTCTCCCTTCGGGAAGCGTGGCGCTTTCTCATAGCTCACGCTGTAGGTATCT
o ++++++
3' GGCCTATGGACAGGCGGAAAGAGGGGAAGCCCTTCGCACCGCGAAAGAGTATCGAGTGCACATCCATAGA
1 P D T C P P F S L R E A W R F L I A H A V G I
2 R I P V R L S P F G K R G A F S . L T L . V S
3 T G Y L S A F L P S G S V A L S H S S R C R Y L
o
5' CAGTTCGGTGTAGGTCGTTTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCCGTTCAGCCCGACCGCTGC
o ++++++
3' GTCAAGCCACATCCAGCAAGCGAGGTTTCGACCCGACACACGTGCTTGGGGGGCAAGTCGGGCTGGCGACG
1 S V R C R S F A P S W A V C T N P P F S A P T A A
2 Q F G V R G R S L Q A G L C A R T P R S A R P L
3 S S V . V V R S K L G C V H E P P V Q P D R C
o
5' GCCTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCA
o ++++++
3' CGGAATAGGCCATTGATAGCAGAACTCAGGTTGGGCCATTCTGTGCTGAATAGCGGTGACCGTTCGTCGGT
1 P Y P V T I V L S P T R . D T T Y R H W Q Q P
2 R L I R . L S S . V Q P G K T R L I A T G S S H
3 A L S G N Y R L E S N P V R H D L S P L A A A
o
5' CTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTA
o ++++++
3' GACCATTGTCCTAATCGTCTCGCTCCATACATCCGCCACGATGTCTCAAGAACTTCACCACCGGATTGAT
1 L V T G L A E R G M . A V L Q S S . S G G L T
2 W . Q D . Q S E V C R R C Y R V L E V V A . L
3 T G N R I S R A R Y V G G A T E F L K W W P N Y
o
5' CGGCTACACTAGAAGGACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTT
o ++++++
3' GCCGATGTGATCTTCTGTGCATAAACCATAGACGCGAGACGACTTCGGTCAATGGAAGCCTTTTTTCTCAA
1 T A T L E G Q Y L V S A L C . S Q L P S E K E L
2 R L H . K D S I W Y L R S A E A S Y L R K K S
3 G Y T R R T V F G I C A L L K P V T F G K R V
o
```

## U6B-sgRNA2.0 map



## U6B-sgRNA2.0 map



U6B-sgRNA2.0 map

