

INK2S00107R_GRN_Q125X_E08_AA
7861 bp

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GCAGGTAGGAGAGCGGCCGCGCAGACCTCTCGCCTGCTCCTGCCCAGGGGCCCGCCAGGGCCATGTGAGCTTGAGGTTCCCCTGG
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GRN

AGTCTCAGCCGGAGACAACAGAAGAACCGCTTACTGAAACTCCTTGGGGGTTCTGATACACTAGGGGGAGTTTTATGGGAAAGAG
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GAAGCAGTAATTGCAGTGACGCCCCGTTAGAAGGGGCTTTCTACCTCCCCAGCATTCCCCCAAAGCAGGGACCACACCATTCTTG
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| GRN |
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ACCCAGCTCCACCCCTGTCGGTAGGTGCTGGCTTCTTCCCCTCTCCTGGTGGTGGTGGGTGGTTCCCGCGGCGGCCTGGAGCCGG
 TGGGTCGAGGTGGGGACAGCCATCCACGACCGAAGAAGGGGAGAGGACCACCACCACCCACCAAGGGCGCCGCCGGACCTCGGCC

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AGGGGCGCGCGACCCTGGGCTGGGAGCTCCGAGGGCCTGGGAACGAGACCTGAGACCTTGGCTTCTCGAAGGTAGTAGGGACTTG
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GGAGTGGTGACTGAACCTGGTCTGGCTCCTCCTTACTTCCTCTTGTTGCGGGTGGGACGAGCTAGCTTCCGCCTCTCCCAGCCAC
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| $\square$ GRN |
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CGGACAGCCTGCTGAGAACACCCAGGAAGCAGGCGGTGCCAGCTGCAGGTGCTTTGCCTGGGAGCTGTGGGGCTGAGGAGAGGGT
 GCCTGTCGGACGACTCTTGTGGGTCCTTCGTCCGCCACGGTCGACGTCCACGAAACGGACCCTCGACACCCCGACTCCTCTCCCA


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GGAAAGCCAGGTGGAGCAGAGAGGATGTGAGTGACTGGGTGGGTGAGATTTCCTGCCCCTCCCCCCGCAGTGGTATCCACACCTA
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GACTCGTGGGGTAACTGAGGCACAGACAGAGAGCAACTTCTCAGGCCCTCACAGTTGGCAATTCTAGGATTAGGACCCAAGTGCG بНبНبН ب CTGAGCACCCCATTGACTCCGTGTCTGTCTCTCGTTGAAGAGTCCGGGAGTGTCAACCGTTAAGATCCTAATCCTGGGTTCACGC

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ATTTTCAGGCAGTCCCTGTACCCTGTTTCTGTTGTACCTGTTGCACCATTCCCAGGCACTGCCCATCGTGCCACTAGTGATATGA بץ ب ب ب ب TAAAAGTCCGTCAGGGACATGGGACAAAGACAACATGGACAACGTGGTAAGGGTCCGTGACGGGTAGCACGGTGATCACTATACT


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| GRN-201 |

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CTCCACCTCCTGGGCTCAAGCGATTCTCCTGCCTCAGCCTCTTGAGTAGCTGGGATTGCAGGTGTGCGCTACCACGCATGGCTAA
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 AATTGTAGAGATTCGAAGTCAAAGGAGGAAATTTTATTTCCACACCGACCCACACCACCAAGTTCGGACATTAGGGTCGTGAATC

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GGAGGCTGAGGTGGGTGGATCAGCTGAGGTCAGGAGTTCAAGACCAGCCTGACCAATATGGTGAAACCCCCTCTCTGCTAAAAAT

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GATCGTGCCACTGAACTCGAGCATGGGCAACAGAGCAAGACTGTCTCAAAAAAAAAAAAAAAAAGGGGGTGAGCAGACGTGGTGG
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CACGCTCCCACAGTCCCAGCTACTTAGTAGGAGGCCAAGGTTGGAGGATTGCTTGATCCCAGGAGTCTGAGTCCAGCCTGGGCAA

GTGCGAGGGTGTCAGGGTCGATGAATCATCCTCCGGTTCCAACCTCCTAACGAACTAGGGTCCTCAGACTCAGGTCGGACCCGTT

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AAAATAATGGAGAGTCTTGTAACTGGCTCCCAAGAGGCTCAACAGACATTACTGTTTTTGCTTCTTCATTATGAGTTACCTCTCT HبHHHH TTTTATTACCTCTCAGAACATTGACCGAGGGTTCTCCGAGTTGTCTGTAATGACAAAAACGAAGAAGTAATACTCAATGGAGAGA $\begin{array}{r}\hline \text { GRN } \\ \hline \text { GRN-201 } \\ \hline\end{array}$ GGCCACCCCACTGAACTAGCTGGGCTAGCTGAGCCTGGGAGAAGAGTTGTTTAGGAAGTGAGAGGCTGCTCTCCACAGAGACTCA
 CCGGTGGGGTGACTTGATCGACCCGATCGACTCGGACCCTCTTCTCAACAAATCCTTCACTCTCCGACGAGAGGTGTCTCTGAGT

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GAATAGATGGAGAGGCAAGGGCAGGGTTTAGCATGCTTGAGGAATCTCAGAGGGCCCTGGTGGTGTGGGGGACCCTCAGAACACA
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GGTGTCTCAAGGGCTGACCCAGCTTCTGTGTCCTTTTCTCTGGGTGAGGAGGGGACATTCATGGGCAGATGGTGACCTCTGGGGA НبץبץبНبН CCACAGAGTTCCCGACTGGGTCGAAGACACAGGAAAAGAGACCCACTCCTCCCCTGTAAGTACCCGTCTACCACTGGAGACCCCT

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AGGCAGCCCAGACTCCACTGGCCACCATATTTCCTTTTTCACAACTTTCTCACCCCTGTGGTTTCCCATGTCATCATGTGGCCGC


TTCCCGCAAGGCCTTAGCGGGGTGCAGGTATGAACATAGTGTCAGGCAAGGAGGCATCTGGAGGGGAACCCTGGCTTTTCCTGGG بץبН AAGGGCGTTCCGGAATCGCCCCACGTCCATACTTGTATCACAGTCCGTTCCTCCGTAGACCTCCCCTTGGGACCGAAAAGGACCC

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GCCCTGTGGCCTGCTGCCTGGACCCCGGAGGAGCCAGCTACAGCTGCTGCCGTCCCCTTCTGGTGAGTGCCCCTCAGCCTAGGCA世بНبНبНب世 CGGGACACCGGACGACGGACCTGGGGCCTCCTCGGTCGATGTCGACGACGGCAGGGGAAGACCACTCACGGGGAGTCGGATCCGT



TCCCTGTCTTTCTAGGACAAATGGCCCACAACACTGAGCAGGCATCTGGGTGGCCCCTGCCAGGTTGATGCCCACTGCTCTGCCG
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GCCACTCCTGCATCTTTACCGTCTCAGGGACTTCCAGTTGCTGCCCCTTCCCAGAGGTGAGCGTGCCATCAGCCCAGTGGAGGGG
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GGCCGTGGCATGCGGGGATGGCCATCACTGCTGCCCACGGGGCTTCCACTGCAGTGCAGACGGGCGATCCTGCTTCCAAAGATCA بץبץبץ CCGGCACCGTACGCCCCTACCGGTAGTGACGACGGGTGCCCCGAAGGTGACGTCACGTCTGCCCGCTAGGACGAAGGTTTCTAGT
 GRN-201

GGTGCAGCTGGGGTGTGGGTGCAGGGCAGGCAGACGGGCAGCATGTGGAGTCTGGAACCCAGGAGCCCAGCTGGCGGGGGCAGCC
 CCACGTCGACCCCACACCCACGTCCCGTCCGTCTGCCCGTCGTACACCTCAGACCTTGGGTCCTCGGGTCGACCGCCCCCGTCGG


PCR Forward

AgGAGGGTGCGGGAGAAAGTGCAAGACTCCAGGTCCAGGCGTTGTGGGGGTGGGGAGAGGTCGAGCTGGGCCGGTCTAATACCAA
 TCCTCCCACGCCCTCTTTCACGTTCTGAGGTCCAGGTCCGCAACACCCCCACCCCTCTCCAGCTCGACCCGGCCAGATTATGGTT


CCCATGGTCAGTGGGTGCCCCTTCCCCATGCCATCTTGCTGAGGGAGGGACTGGATTGTGAGGAGGGTGAGTTAGGCCTGCCTAG Н- $+\boldsymbol{+}$ GGGTACCAGTCACCCACGGGGAAGGGGTACGGTAGAACGACTCCCTCCCTGACCTAACACTCCTCCCACTCAATCCGGACGGATC


GAGATCACTGAGCCTTAGTGTCACCCTCAAACCCCAGTAGCTGGGCTTGCAGGCCCTGGTGCCACCAGCTCCTTGTGTGATGGGG
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GRN-201

Donor Template SNV -> REV
cctgagtgggctggtagtatcctgggtcatcttgtccacagGTAACAACTCCGTGGGTGCCATCCAGTGCCCTG
GAGTCACCTTCCCTGAGTGGGCTGGTAGTATCCTGGGTCATCTTGTCCACAGGTAACAACTCCGTGGGTGCCATCCAGTGCCCTG



ATAGTCAGTTCGAATGCCCGGACTTCTCCACGTGCTGTGTTATGGTCGATGGCTCCTGGGGGTGCTGCCCCATGCCCCAGGTACA

TATCAGTCAAGCTTACGGGCCTGAAGAGGTGCACGACACAATACCAGCTACCGAGGACCCCCACGACGGGGTACGGGGTCCATGT

Donor Tem
Protospacer Sequence
TATCAGTC
gRNA Protospacer

AATCTGGGGGAGATGGGGGTATGTGGAGGGAAGTGGGGGCAGAGTTGGGGGCCAGGGGCAGGGGGTGAAGACGGAGTCAGGACCA
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GRN-201

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GCCAGGGGCTGGGGCGGGGCCTCATTGACTCCAAGTGTAGGAAAAAGTTTCCTCCATCCTGGCTGCCCCTCACGTTTGCTCCTCT
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CTCCAAGGAGAACGCTACCACGGACCTCCTCACTAAGCTGCCTGCGCACACAGGTACCAGAGGCAGGGTGCAGATACAGGGGTGG廿нみН GAGGTTCCTCTTGCGATGGTGCCTGGAGGAGTGATTCGACGGACGCGTGTGTCCATGGTCTCCGTCCCACGTCTATGTCCCCACC


 CCGGGGGAAAGGAGGGAAAATCCGGACCGGAATCCTAGTGACGTTCCACCACATtCGCCATGGGAGGTAGAAGTTGTGGACCAAG



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 AAAATGGGTCCATGGGTCCCCACCGCCCACCCACCCGACTCGTGTCACACCGTCCGTCGGCCCGGGGTCACGGGTGGACGGGAAG

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GRN-201
ttcatctaccctaggctgtgtactgtgaggaccacatacactgctgtcccacgagatttacgtgtgacacgcagangegtacctg
 AAGTAGACGGGATCCGACACACGACACTCCTGGTGTATGTGACGACAGGGCGCCCCAAATGCACACTGTGCGTCTTCCCATGGAC


TGAACAGGGGCCCCACCAGGTGCCCTGGATGGAGAAGGCCCCAGCTCACCTCAGCCTGCCAGACCCACAAGCCTTGAAGAGAGAT

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gtcccctatgatantgtcagcagctgtccctcctccgatacctactaccaictcacgtctagagagtgagactactgtccantcc +HبH+H caggagacactattacagtcgtcgacagg aggaggctatggacgacgattgagtgcagaccoctcaccccgacgacaggttagg


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GRN-201

TAGGTGATACCCAGCTCTGACAGATTCGTCCCCAGCTGGAGGTGCTGTAAGCAGGAGAGGCGGGCTGGAGTAGGTAGGGGCTCGG
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CCGGGCTTCCTTATCCCACCCCAGAGACATCGGCTGTGACCAGCACACCAGCTGCCCGGTGGGGCAGACCTGCTGCCCGAGCCTG HبН H GGCCCGAAGGAATAGGGTGGGGTCTCTGTAGCCGACACTGGTCGTGTGGTCGACGGGCCACCCCGTCTGGACGACGGGCTCGGAC


GGTGGGAGCTGGGCCTGCTGCCAGTTGCCCCATGTGAGTGCCTCCCTGCCTGCCCCTGGATAGGGGAGCTAAGCCCAGTGAGGGG



ACAGGAACATAATGCCATTCTGTGCTCCCTTCCCCGCCAGGCTGTGTGCTGCGAGGATCGCCAGCACTGCTGCCCGGCTGGCTAC ННН TGTCCTTGTATTACGGTAAGACACGAGGGAAGGGGCGGTCCGACACACGACGCTCCTAGCGGTCGTGACGACGGGCCGACCGATG


ACCTGCAACGTGAAGGCTCGATCCTGCGAGAAGGAAGTGGTCTCTGCCCAGCCTGCCACCTTCCTGGCCCGTAGCCCTCACGTGG
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GCCCCCCTCTGACCATCCAGGGCGTCTGTTGTGCTGATCGGCGCCACTGCTGTCCTGCTGGCTTCCGCTGCGCAGCCAGGGGTAC بнبНبн CGGGGGGAGACTGGTAGGTCCCGCAGACAACACGACTAGCCGCGGTGACGACAGGACGACCGAAGGCGACGCGTCGGTCCCCATG


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CAAGTGTTTGCGCAGGGAGGCCCCGCGCTGGGACGCCCCTTTGAGGGACCCAGCCTTGAGACAGCTGCTGTGAGGGACAGTACTG
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| GRN-201 |  |  |
| CCACATTACAAGCTGCCATCCCCTCCCCGTTTCAGTGGACC <br>  GGTGTAATGTTCGACGGTAGGGGAGGGGCAAAGTCACCTGG | gtgaccagatgcttttccctatccacaggagtatttgigtat <br>  ACCGGTCCACGAAAAGGGATAGGTGTCCCCACAAACACACA | 7820 |
| GRN |  |  |
| GRN-201 |  |  |
| GTGCGCGTGTGCGTTTCAATAAAGTTTGTACACTTTCTTAA <br>  cacgCgCacacgcanagttatt tcanacatgtganagant | $\begin{aligned} & 3^{\prime} \\ & 5^{\prime} \end{aligned}$ |  |
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/note $\quad=\quad$| gene ENSG000000030582 |
| :--- |
|  |
| Protein coding |

| GRN-212 |  |  | 1 | .. | 4649 | 4649 bp | $\square$ | $\rightarrow$ | prim_transcript |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| /note |  | primary transcript ENST00000588170 Retained intron |  |  |  |  |  |  |  |
| GRN-215 |  |  | 17 | .. | 4491 | 4475 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note |  | primary transcript ENST00000589536 |  |  |  |  |  |  |  |
| GRN-209 |  |  | 20 | .. | 5042 | 5023 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000587518 |  |  |  |  |  |  |  |
| GRN-207 |  |  | 20 | .. | 4229 | 4210 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000587109 |  |  |  |  |  |  |  |
| GRN-211 |  |  | 37 | .. | 5224 | 5188 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000588143 |  |  |  |  |  |  |  |
| GRN-221 |  |  | 53 | .. | 5080 | 5028 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000593167 |  |  |  |  |  |  |  |
| GRN-201 |  |  | 57 | .. | 7861 | 7805 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000053867 |  |  |  |  |  |  |  |
| GRN-213 |  |  | 57 | .. | 5799 | 5743 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000588237 |  |  |  |  |  |  |  |
| GRN-208 |  |  | 57 | .. | 5266 | 5210 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000587387 |  |  |  |  |  |  |  |
| GRN-220 |  |  | 57 | .. | 5059 | 5003 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000592783 |  |  |  |  |  |  |  |
| GRN-218 |  |  | 57 | .. | 5038 | 4982 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENST00000591740 |  |  |  |  |  |  |  |
| GRN-219 |  |  | 57 |  | 4544 | 4488 bp | $\square$ | $\rightarrow$ | prim_transcript |
| /note | $=$ | primary transcript ENSTOOOO0592323 Retained intron |  |  |  |  |  |  |  |
| GRN-203 |  |  | 57 | .. | 4030 | 3974 bp | $\square$ | $\rightarrow$ | prim_transcript |


/note
= primary transcript ENST00000586782 Nonsense mediated decay
GRN-201

* 12 segments $=1782 \mathrm{bp}$
/note $\quad=$ coding sequence ENSP00000053867
/translation $=$ MWTLVSWVALTAGLVAGTRCPDGQFCPVACCLDPGGASYSCCRPLL,,DKWPTTLSRHLGGPCQVDAHCSAGHSCIFTVSGTSSCCPFPE,, AVACGDGHHCCPRGFHCSADGRSCFQRS,,GNNSVGAIQCPDSQFECPDFSTCCVMVDGSWGCCPMPQ,,ASCCEDRVHCCPHGAFCDLV HTRCITPTGTHPLAKKLPAQRTNRA ,,VALSSSVMCPDARSRCPDGSTCCELPSGKYGCCPMPN,,ATCCSDHLHCCPQDTVCDLIQSKCLSK ENATTDLLTKLPAHT,,VGDVKCDMEVSCPDGYTCCRLQSGA WGCCPFTQ ,,AVCCEDHIHCCPAGFTCDTQKGTCEQGPHQVPWMEKAPA HLSLPDPQALKRDVPCDNVSSCPSSDTCCQLTSGEWGCCPIPE,,AVCCSDHQHCCPQGYTCVAEGQCQRGSEIVAGLEKMPARRASLSH PRDIGCDQ HTSCPVGQTCCPSLGGSWACCQLPH ,,AVCCEDRQHCCPAGYTCNVKARSCEKEVVSAQPATFLARSPHVGVKDVECGEGHF Gوibl

| /note | $=$ coding sequence ENSP00000467616 |
| :--- | :--- |
| /translation | $=$ MWTLVSWVALTAGLVAGTRCPDGQFCPVACCLDPGGASYSCCRPLL,,DKWPTTLSRHLGGPCQVDAHCSAGHSCIFTVSGTSSCCPFPE,, | AVACGDGHHCCPRGFHCSADGRSCFQRS,,GNNSVGAIQCPDSQFECPDFSTCCVMVDGSWGCCPMPQ,,AVCCEDHIHCCPAGFTCDTQ KGTCEQGPHQVPWMEKAPAHLSLPDPQALKRDVPCDNVSSCPSSDTCCQLTSGEWGCCPIPE,,AVCCSDHQHCCPQGYTCVAEGQCQR GSEIV A GLEKMPARRASLSHPRDIGCDQHTSCPVGQTCCPSLGGSWACCQLPH,,AVCCEDRQHCCPAGYTCNVKARSCEKEVVSAQPAT FLARSPHVGVKDVECGEGHFCHDNQTCCRDNRQGWACCPYRQ,,GVCCADRRHCCPAGFRCAARGTKCLRREAPRWDAPLRDPALRQLL* 436 amino acids $=46.8 \mathrm{kDa}$

GRN-222 $3920 . .7550 \quad 3631 \mathrm{bp} \quad \square \quad \rightarrow \quad$ CDS

* 11 segments $=1314 \mathrm{bp}$
/note $\quad=$ coding sequence ENSP00000492014
/translation $=$ MWTLVSWVALTAGLVAGTRCPDGQFCPVACCLDPGGASYSCCRPLL,,DKWPTTLSRHLGGPCQVDAHCSAGHSCIFTVSGTSSCCPFPE,, AVACGDGHHCCPRGFHCSADGRSCFQRS,,GNNSVGAIQCPDSQFECPDFSTCCVMVDGSWGCCPMPQ,,ASCCEDRVHCCPHGAFCDLV HTRCITPTGTHPLAKKLPAQRTNRA,,VALSSSVMCPDARSRCPDGSTCCELPSGKYGCCPMPN,,ATCCSDHLHCCPQDTVCDLIQSKCLSK ENATTDLLTKLPAHT,,VGDVKCDMEVSCPDGYTCCRLQSGA WGCCPFTQ,,AVCCEDHIHCCPAGFTCDTQKGTCEQGPHQVPWMEKAPA HLSLPDPQALKRDVPCDNVSSCPSSDTCC ,,RDNRQGWACCPYRQ,,GVCCADRRHCCPAGFRCAARGTKCLRREAPRWDAPLRDPALRQLL 438 amino acids $=47.0 \mathrm{kDa}$


| /note $=$ | coding sequence ENSP00000467431 |
| :--- | :--- |
| /translation $=$ | MWTLVSWVALTAGLVAGTRCPDGQFCPVACCLDPGGASYSCCRPLL,,DKWPTTLSRHLGGPCQVDAHCSAGHSCIFTVSGTSSCCPFPE,,I |
|  | KGPCQCRFLCVPQAVACGDGHHCCPRGFHCSADGRSCFQRS, GNNSVGAIQCPDSQFECPDFSTCCVMVDGSWGCCPMPQ,,ASCCEDR | Y出CGRHGAEFABL¥ HTOR:3 kDa

GRN-211
( 5 segments $=490 \mathrm{bp}$


```
Feature
    Location
    Size
```



```
GRN-218
/note \(\quad=\) coding sequence ENSP00000467022
/translation = MWTLVSWVALTAGLVAGTRCPDGQFCPVACCLDPGGASYSCCRPLL,,DKWPTTLSRHLGGPCQVDAHCSAGHSCIFTVSGTSSCCPFPE,,
                AVACGDGHHCCPRGFHCSADGRSCFQRS,,GNNSVGAIQCPDSQFECPD
                135 amino acids = 14.1 kDa
```



| /note | $=$ coding sequence ENSP00000466271 |
| :--- | :--- |
| /translation | $=$ MWTLVSWVALTA GLVAGTRCPDGQFCPVACCLDPGGASYSCCRPLL,,DKWPTTLSRHLGGPCQ | 62 amino acids $=6.6 \mathrm{kDa}$



| Donor Template SNV -> REV | 4942 | . .5041 | 100 bp | $\square$ | $\mapsto$ | misc_feature |
| :--- | ---: | ---: | ---: | ---: | ---: | ---: |
| PAM | 5001 | . .5003 | 3 bp | $\square$ | $\mapsto$ | misc_feature |
| Protospacer Sequence | 5004 | . .5023 | 20 bp | $\square$ | $\mapsto$ | misc_feature |
| SNV | 5006 | . .5006 | 1 bp | $\square$ | $\mapsto$ | misc_feature |

/note $\quad=\quad$|  |
| :--- |
|  |
|  |
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|  |
|  |
|  |

GRN-217 5044 .. $6055 \quad 1012 \mathrm{bp} \quad \square \quad \rightarrow \quad$ prim_transcript
/note $\quad=\quad$ primary transcript ENST000000590984
Retained intron

( 7 segments $=1166 \mathrm{bp}$




