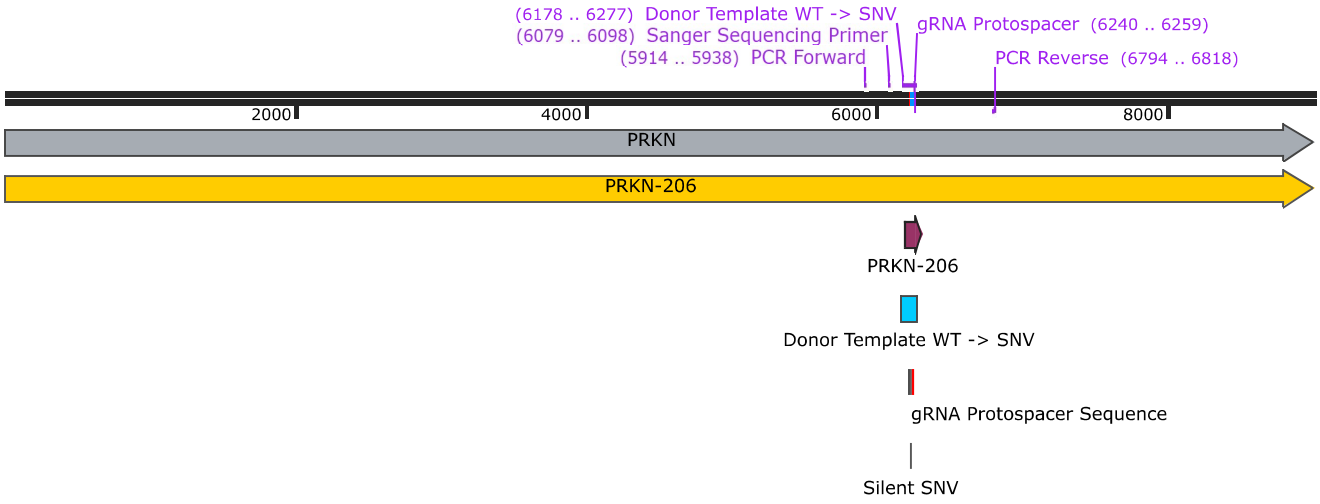

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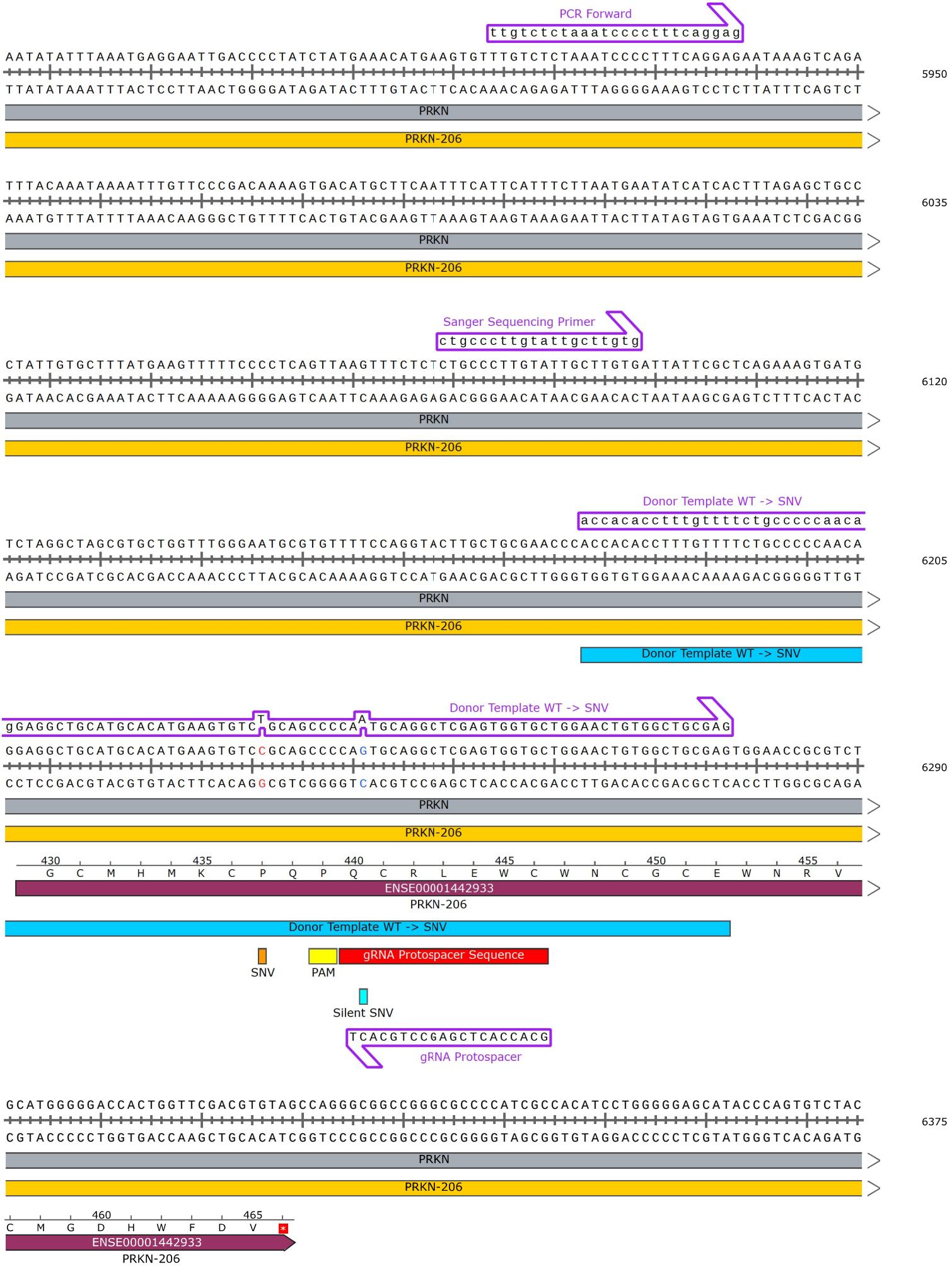
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GACATTT
PCR Reverse
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6885
PRKN
PRKN-206
TACTTGACGTCACCCAAG
PCR Reverse
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TTCTTGGGGTCCGTTATCAGTTTTTGTAAACAAATAGGAATTCTTAAGGTAGATATATTTAGCGTAATTACTTTATGGTTGATAC
6970
PRKN
PRKN-206
CGTAAATCAACTTGTCACAAAGTGAGAAATTATGAAAGTTAATTTGAATGTTGAATGTTTGAATTACAGGGAAGAAATCAAGTTA
GCATTTAGTTGAACAGTGTTTCACTCTTTAATACTTTCAATTAAACTTACAACTTACAAACTTAATGTCCCTTCTTTAGTTCAAT
7055
PRKN
PRKN-206
ATGTACTTTCATTCCCTTTCATGATTTGCAACTTTAGAAAGAAATTGTTTTTCTGAAAGTATCACCAAAAAATCTATAGTTTGAT
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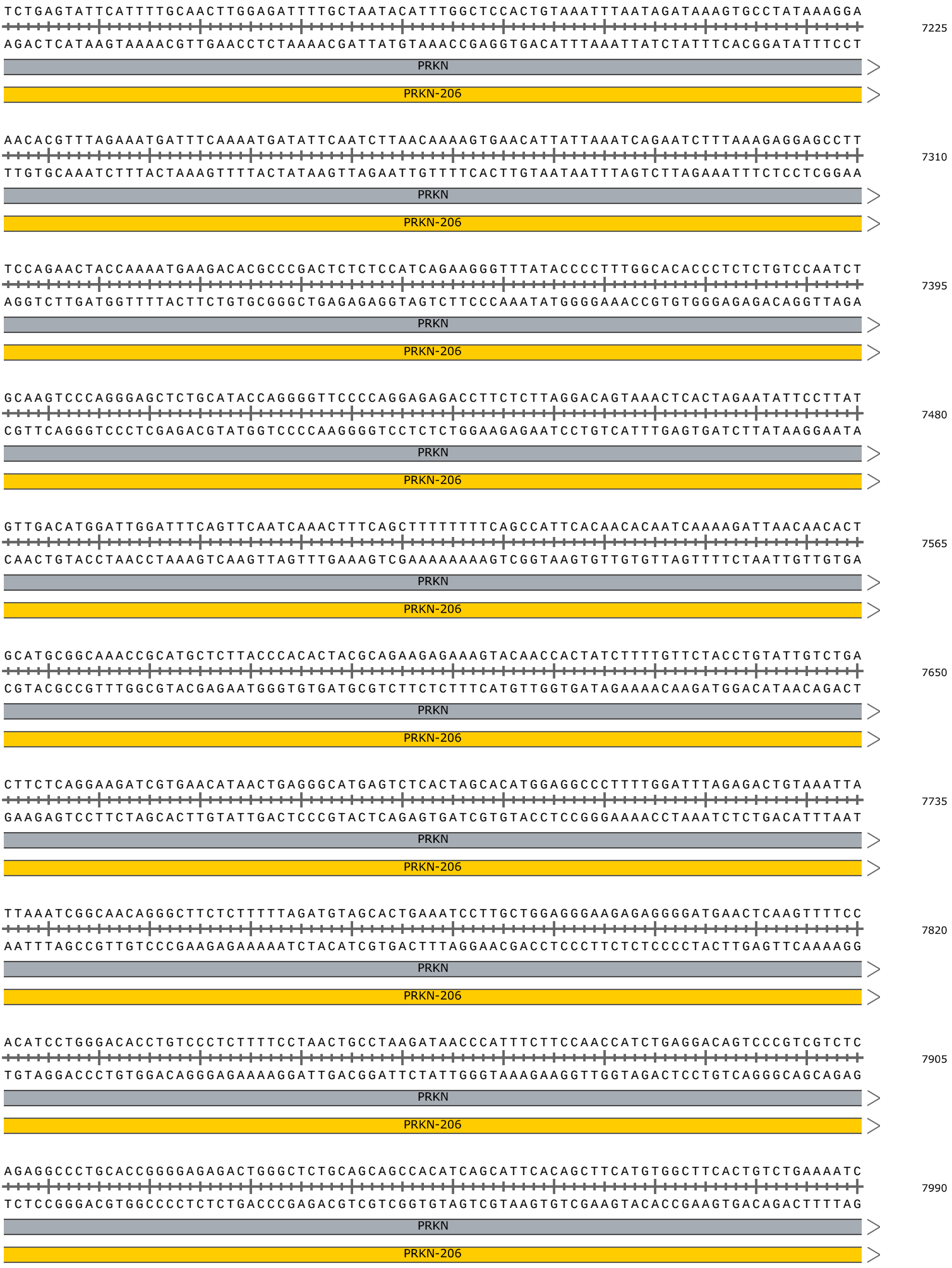
7140
PRKN
PRKN-206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PRKN
PRKN-206
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7310
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PRKN
PRKN-206
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7395
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PRKN-206
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7480
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PRKN
PRKN-206
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7565
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PRKN
PRKN-206
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7650
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PRKN
PRKN-206
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7735
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7820
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PRKN
PRKN-206
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7905
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PRKN
PRKN-206
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7990
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PRKN
PRKN-206


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PRKN-206
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PRKN
PRKN-206
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8245
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PRKN-206
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8330
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PRKN-206
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8500
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PRKN
PRKN-206
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8585
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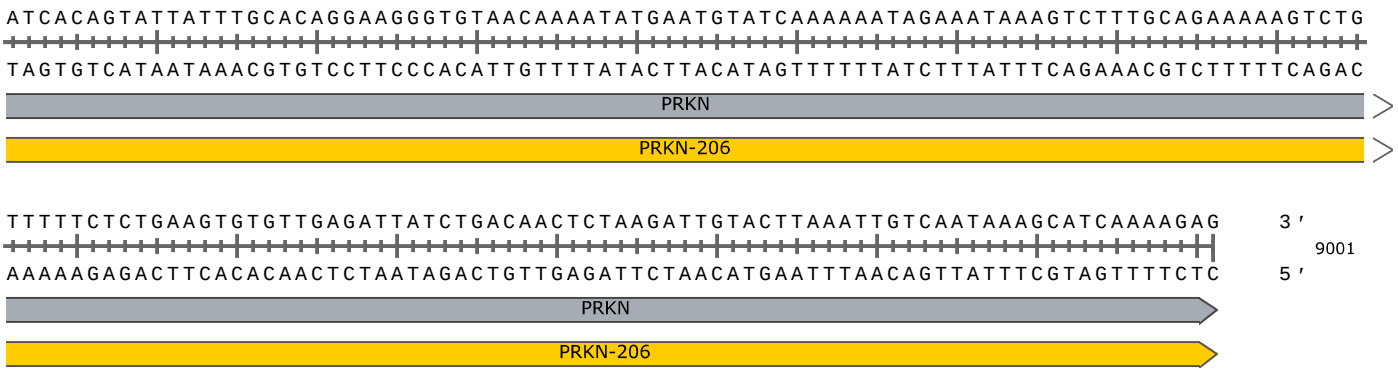
PRKN
PRKN-206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PRKN
PRKN-206

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3′
9001
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5′
PRKN
PRKN-206

	Feature			Location			Size			Type
	PACRG			1	..	9001	9001 bp		←	gene
	/note	=	gene ENSG00000112530 Protein coding							
✓	**PRKN**			1	..	9001	9001 bp		→	gene
	/note	=	gene ENSG00000185345 Protein coding							
	PACRG-201			1	..	9001	9001 bp		←	prim_transcript
	/note	=	primary transcript ENST00000337019 Protein coding							
	PACRG-203			1	..	9001	9001 bp		←	prim_transcript
	/note	=	primary transcript ENST00000366889 Protein coding							
✓	**PRKN-206**			1	..	9001	9001 bp		→	prim_transcript
	/note	=	primary transcript ENST00000366898							
	PRKN-212			1	..	8811	8811 bp		→	prim_transcript
	/note	=	primary transcript ENST00000673871 Nonsense mediated decay							
	PRKN-213			1	..	8811	8811 bp		→	prim_transcript
	/note	=	primary transcript ENST00000674006 protein_coding_CDS_not_defined							
	PRKN-221			1	..	8811	8811 bp		→	prim_transcript
	/note	=	primary transcript ENST00000674436 protein_coding_CDS_not_defined							
	PRKN-204			1	..	7779	7779 bp		→	prim_transcript
	/note	=	primary transcript ENST00000366896							
	PRKN-205			1	..	7779	7779 bp		→	prim_transcript
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	PRKN-201			1	..	6319	6319 bp		→	prim_transcript
	/note	=	primary transcript ENST00000338468 Nonsense mediated decay							
	PRKN-203			1	..	6319	6319 bp		→	prim_transcript
	/note	=	primary transcript ENST00000366894 Nonsense mediated decay							
	PRKN-207			1	..	6319	6319 bp		→	prim_transcript
	/note	=	primary transcript ENST00000479615 Nonsense mediated decay							
	PRKN-208			1	..	6319	6319 bp		→	prim_transcript
	/note	=	primary transcript ENST00000610470							
				1068	..	2739	1672 bp		←	gene
	/note	=	gene ENSG00000286498 lncRNA							
				1068	..	2739	1672 bp		←	prim_transcript
	/note	=	primary transcript ENST00000660049 lncRNA							
✓	**Donor Template WT -> SNV**			6178	..	6277	100 bp		⊢	misc_feature
	PRKN-204			6207	..	6319	113 bp		→	CDS
	/codon_start	=	1							
	/note	=	coding sequence ENSP00000355862							
	/translation	=	GCMHMKCPQPQCRLEWCWNCGCEWNRVCMGDHWFDV* 36 amino acids = 4.4 kDa							
	PRKN-205			6207	..	6319	113 bp		→	CDS
	/codon_start	=	1							
	/note	=	coding sequence ENSP00000355863							
	/translation	=	GCMHMKCPQPQCRLEWCWNCGCEWNRVCMGDHWFDV* 36 amino acids = 4.4 kDa							

	Feature	Location	Size			Type
✓	**PRKN-206**	6207 .. 6319	113 bp		→	CDS
	/codon_start = 1					
	/note = coding sequence ENSP00000355865					
	/translation = GCMHMKCPQPQCRLEWCWNCGCEWNRVCMGDHWFDV*					
	36 amino acids = 4.4 kDa					
	PRKN-208	6207 .. 6319	113 bp		→	CDS
	/codon_start = 1					
	/note = coding sequence ENSP00000483773					
	/translation = GCMHMKCPQPQCRLEWCWNCGCEWNRVCMGDHWFDV*					
	36 amino acids = 4.4 kDa					
✓	**SNV**	6231 .. 6231	1 bp		⊢	misc_feature
	/note = WT = C SNV = T					
✓	**PAM**	6236 .. 6238	3 bp		⊢	misc_feature
✓	**gRNA Protospacer Sequence**	6239 .. 6259	21 bp		⊢	misc_feature
✓	**Silent SNV**	6241 .. 6241	1 bp		⊢	misc_feature
	/note = WT = G Silent SNV = A					

	Primer	Length	Binding Sites		Tm	Date Added
✓	**PCR Forward**	25-mer	5914 .. 5938	→	58°C	Mar 7, 2023
	/sequence = ttgtctctaaatccccttcaggag 44% GC / 7583.0 Da					
✓	**Sanger Sequencing Primer**	20-mer	6079 .. 6098	→	57°C	Mar 7, 2023
	/sequence = ctgcccttgtattgcttgtg 50% GC / 6081.0 Da					
✓	**Donor Template WT -> SNV**	100-mer	6178 .. 6277	→	81°C	Mar 7, 2023
	/sequence = accacacctttgttttctgcccccaacagGAGGCTGCATGCACATGAAGTGTCTGCAGCCCCAATGCAGGCTCGAGTGGTGCTGGAACTGTGGCT 68% GC / 30,828.0 Da					
✓	**gRNA Protospacer**	20-mer	6240 .. 6259	←	64°C	Mar 7, 2023
	/sequence = GCACCACTCGAGCCTGCACT 65% GC / 6022.9 Da					
✓	**PCR Reverse**	25-mer	6794 .. 6818	←	58°C	Mar 7, 2023
	/sequence = GAACCCACTGCAGTTCATTTTACAG 44% GC / 7601.0 Da					