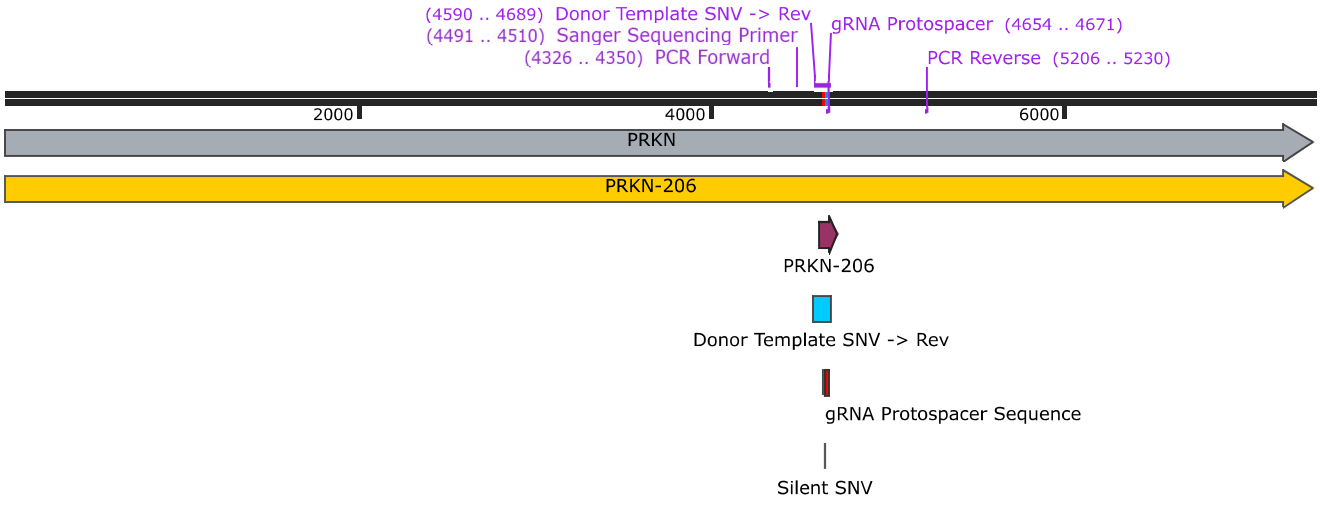

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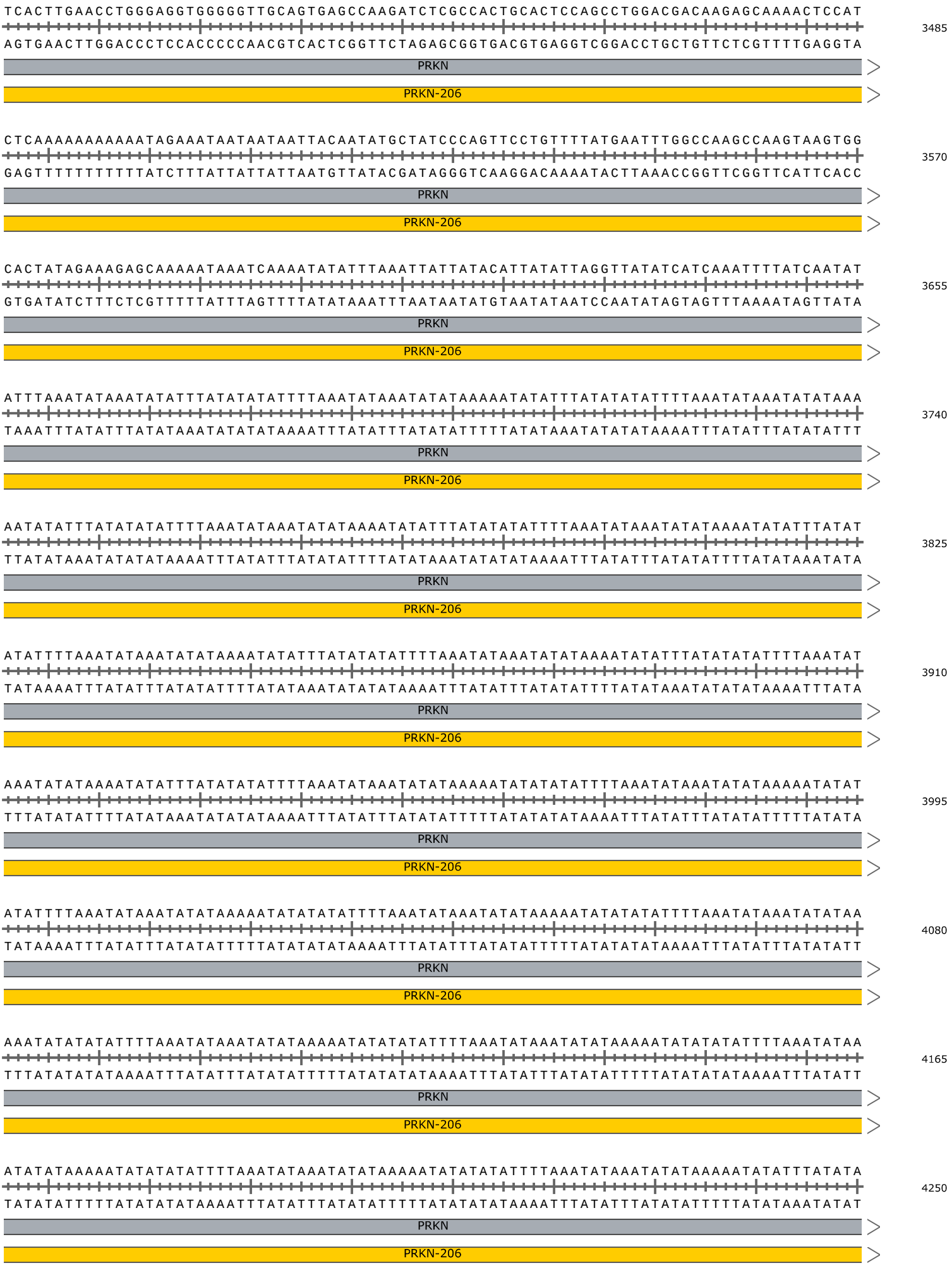
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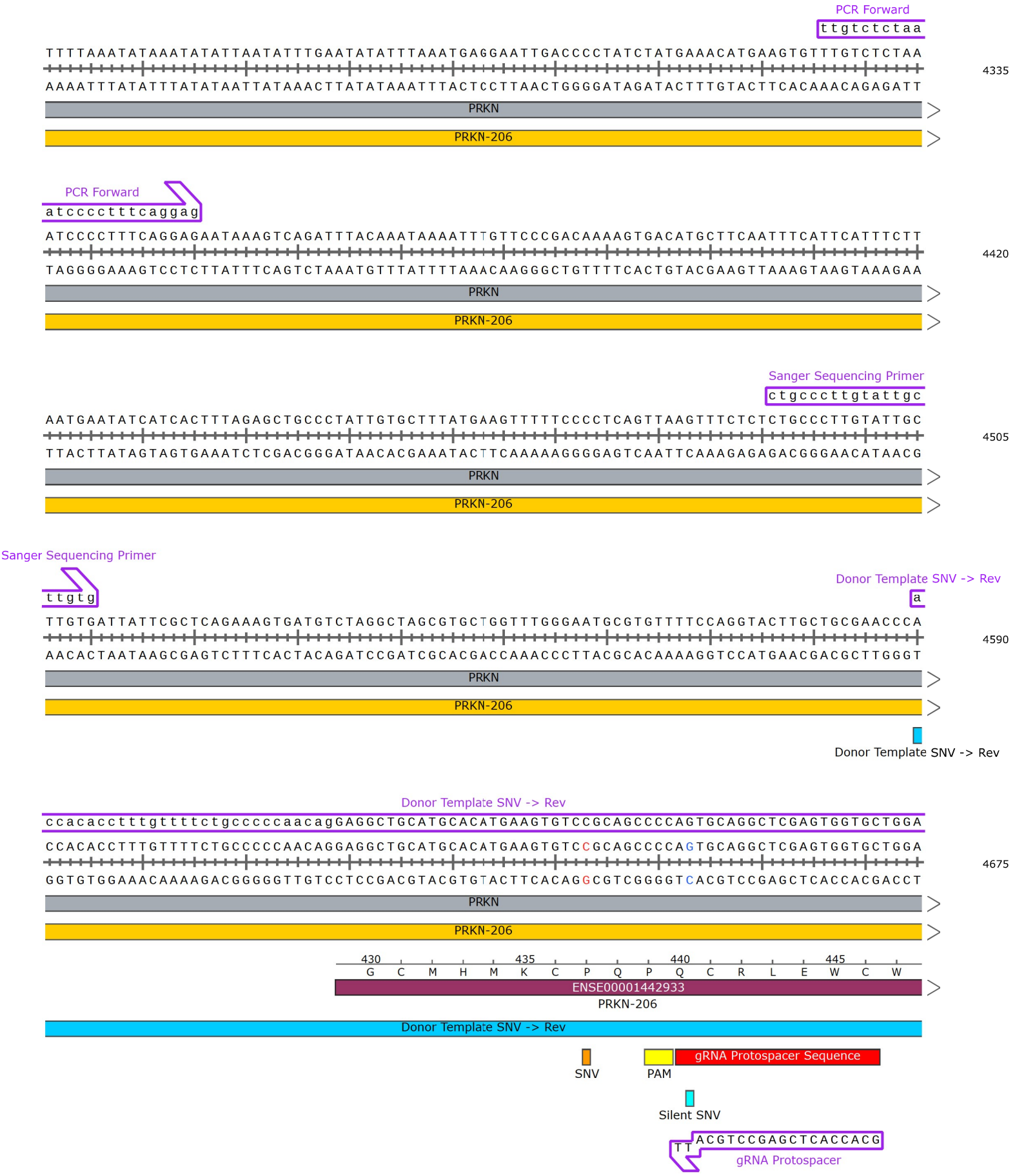
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PRKN
PRKN-206
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6715
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PRKN-206
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6800
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PRKN-206
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6885
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PRKN-206
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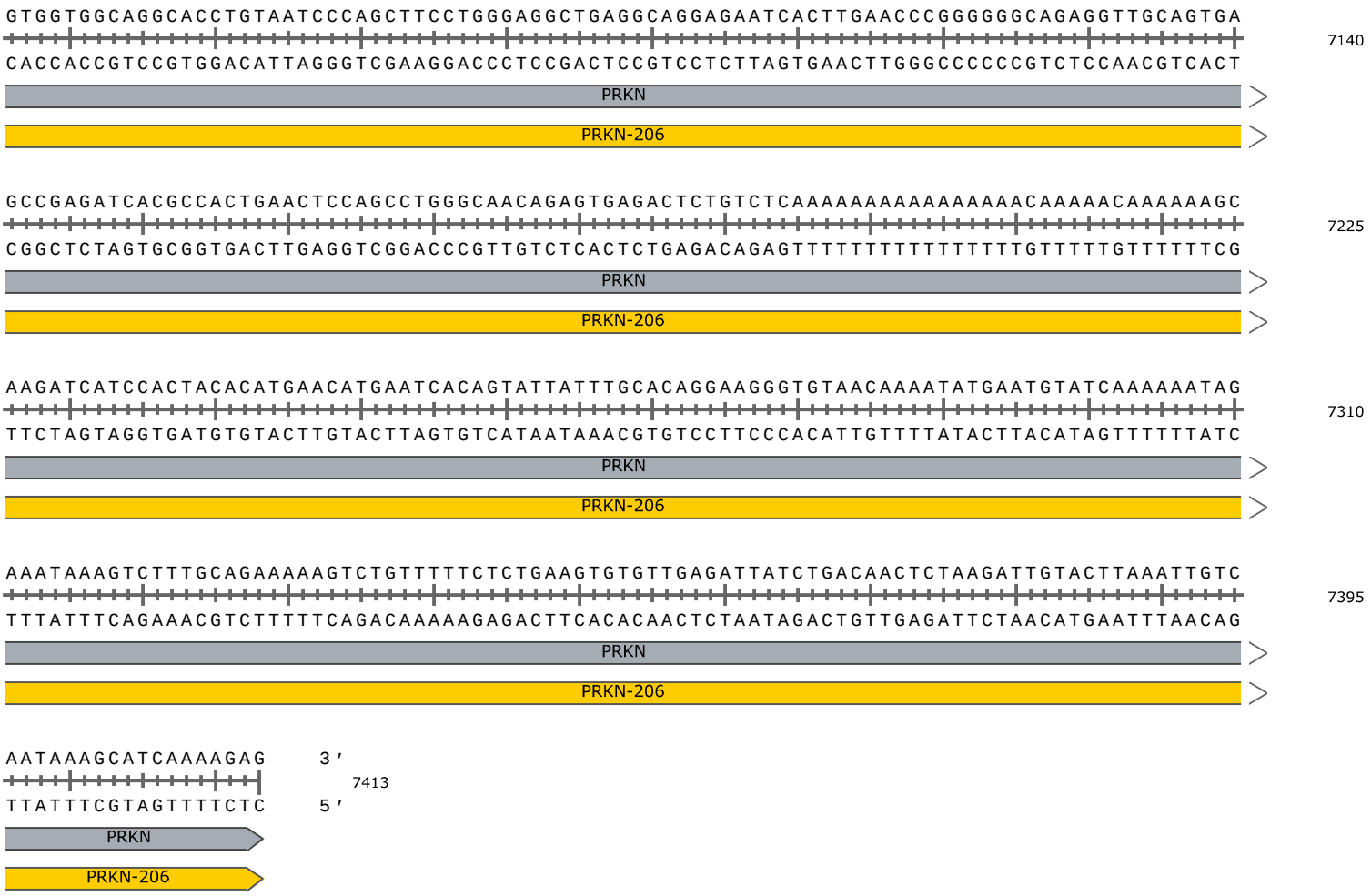
PRKN
PRKN-206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7225
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PRKN-206
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PRKN-206
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7395
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PRKN
PRKN-206
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7413
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PRKN
PRKN-206

	Feature			Location	Size			Type
	PACRG			1 .. 7413	7413 bp		←	gene
	/note	=	gene ENSG00000112530 Protein coding					
✓	**PRKN**			1 .. 7413	7413 bp		→	gene
	/note	=	gene ENSG00000185345 Protein coding					
	PACRG-201			1 .. 7413	7413 bp		←	prim_transcript
	/note	=	primary transcript ENST00000337019 Protein coding					
	PACRG-203			1 .. 7413	7413 bp		←	prim_transcript
	/note	=	primary transcript ENST00000366889 Protein coding					
✓	**PRKN-206**			1 .. 7413	7413 bp		→	prim_transcript
	/note	=	primary transcript ENST00000366898					
	PRKN-212			1 .. 7223	7223 bp		→	prim_transcript
	/note	=	primary transcript ENST00000673871 Nonsense mediated decay					
	PRKN-213			1 .. 7223	7223 bp		→	prim_transcript
	/note	=	primary transcript ENST00000674006 protein_coding_CDS_not_defined					
	PRKN-221			1 .. 7223	7223 bp		→	prim_transcript
	/note	=	primary transcript ENST00000674436 protein_coding_CDS_not_defined					
	PRKN-204			1 .. 6191	6191 bp		→	prim_transcript
	/note	=	primary transcript ENST00000366896					
	PRKN-205			1 .. 6191	6191 bp		→	prim_transcript
	/note	=	primary transcript ENST00000366897					
	PRKN-201			1 .. 4731	4731 bp		→	prim_transcript
	/note	=	primary transcript ENST00000338468 Nonsense mediated decay					
	PRKN-203			1 .. 4731	4731 bp		→	prim_transcript
	/note	=	primary transcript ENST00000366894 Nonsense mediated decay					
	PRKN-207			1 .. 4731	4731 bp		→	prim_transcript
	/note	=	primary transcript ENST00000479615 Nonsense mediated decay					
	PRKN-208			1 .. 4731	4731 bp		→	prim_transcript
	/note	=	primary transcript ENST00000610470					
				1 .. 1151	1151 bp		←	gene
	/note	=	gene ENSG00000286498 lncRNA					
				1 .. 1151	1151 bp		←	prim_transcript
	/note	=	primary transcript ENST00000660049 lncRNA					
✓	**Donor Template SNV -> Rev**			4590 .. 4689	100 bp		⊢	misc_feature
	PRKN-204			4619 .. 4731	113 bp		→	CDS
	/codon_start	=	1					
	/note	=	coding sequence ENSP00000355862					
	/translation	=	GCMHMKCPQPQCRLEWCWNCGCEWNRVCMGDHWFDV* 36 amino acids = 4.4 kDa					
	PRKN-205			4619 .. 4731	113 bp		→	CDS
	/codon_start	=	1					
	/note	=	coding sequence ENSP00000355863					
	/translation	=	GCMHMKCPQPQCRLEWCWNCGCEWNRVCMGDHWFDV* 36 amino acids = 4.4 kDa					

	Feature	Location	Size			Type
✓	**PRKN-206**	4619 .. 4731	113 bp		→	CDS
	/codon_start = 1					
	/note = coding sequence ENSP00000355865					
	/translation = GCMHMKCPQPQCRLEWCWNCGCEWNRVCMGDHWFDV*					
	36 amino acids = 4.4 kDa					
	PRKN-208	4619 .. 4731	113 bp		→	CDS
	/codon_start = 1					
	/note = coding sequence ENSP00000483773					
	/translation = GCMHMKCPQPQCRLEWCWNCGCEWNRVCMGDHWFDV*					
	36 amino acids = 4.4 kDa					
✓	**SNV**	4643 .. 4643	1 bp		⊢	misc_feature
	/note = SNV = T Rev = C					
✓	**PAM**	4649 .. 4651	3 bp		⊢	misc_feature
✓	**gRNA Protospacer Sequence**	4652 .. 4671	20 bp		⊢	misc_feature
✓	**Silent SNV**	4653 .. 4653	1 bp		⊢	misc_feature
	/note = Silent SNV = A Rev = G					

	Primer	Length	Binding Sites	Tm	Date Added
✓	**PCR Forward**	25-mer	4326 .. 4350	58°C	Mar 7, 2023
	/sequence = ttgtctctaaatccccttcaggag 44% GC / 7583.0 Da				
✓	**Sanger Sequencing Primer**	20-mer	4491 .. 4510	57°C	Mar 7, 2023
	/sequence = ctgcccttgtattgcttgtg 50% GC / 6081.0 Da				
✓	**Donor Template SNV -> Rev**	100-mer	4590 .. 4689	83°C	Mar 7, 2023
	/sequence = accacacctttgttttctgcccccaacagGAGGCTGCATGCACATGAAGTGTCCGCAGCCCCAGTGCAGGCTCGAGTGGTGCTGGAACTGTGGC 66% GC / 30,828.9 Da				
✓	**gRNA Protospacer**	20-mer	4654 .. 4671	62°C	Mar 7, 2023
	/sequence = GCACCACTCGAGCCTGCATT 60% GC / 6038.0 Da				
✓	**PCR Reverse**	25-mer	5206 .. 5230	58°C	Mar 7, 2023
	/sequence = GAACCCACTGCAGTTCATTTTACAG 44% GC / 7601.0 Da				