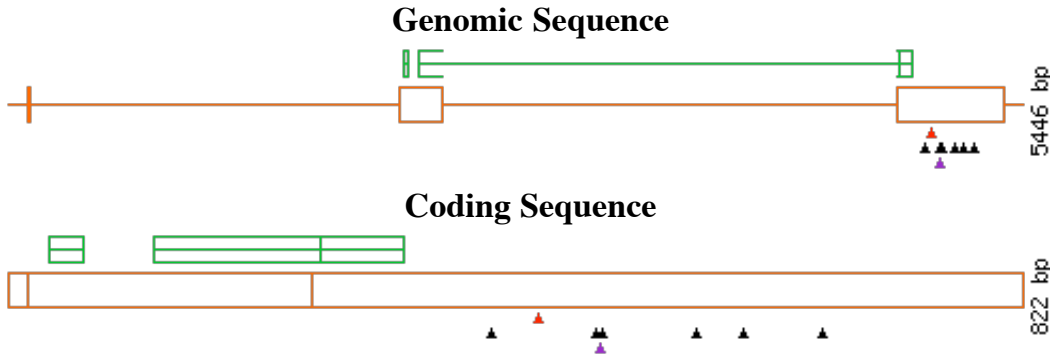


tgif

IPB005540, IPB005542, IPB003893, IPB007689, IPB003350, IPB003106, IPB000327, IPB008888, and IPB006712



For help interpreting these results, view the [PARSENP Introduction](#) page.

#	View On Sequence	Nucleotide Change	Effect	Restriction Enzyme Differences from REBASE		PSSM Difference	Description	Zygoty
				Gained in Variant	Lost from Reference			
1	G C	C4915A	H131N				2229	Homo
2	G C	C4953A	Y143*	DdeI			3287	Homo
3	G C	G5000A	G159D		BstEII , MaeIII , Tsp4CI		1202	Homo
4	G C	C5004T	Y160=		BsiYI , BstEII , MaeIII		4875	Homo
5	G C	C5006T	P161L	EcoNI , MnII			4689	Homo
6	G C	C5081T	T186I				8407	Homo
7	G C	C5120T	T199M	MslI	FnuDII		7947	Homo
8	G C	G5183T	C220F				489	Homo

[Download Tab-Separated table](#)

No [protein homology model](#) was submitted. You may add one using any or all of the fields below.

Blocks Families:

Blocks File: no file selected

Sequence Alignment: no file selected

-OR-

Reverse PSI-Blast/SIFT for Models

Genomic Sequence

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	6
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P K H G L L A N G E D R N S Y E P G	146
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I C H T T I T T A T Q G I N T T T R 200
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