Vector Summary

Vector ID	VB220105-1311kkn
Vector Name	pSPL3-hRPH3A_c.444Tmut
Vector Size	6403 bp
Vector Type	Mammalian Gene Expression Vector
Plasmid Copy Number	High
Antibiotic Resistance	Ampicillin
Cloning Host	VB UltraStable (or alternative strain)

Vector Map

Vector Components

Name	Position	Size (bp)	Type	Description	Application notes
SV40 promoter	10-339	330	promoter	*None*	note=Unknown feature type:Promoter color: #d84e4e; direction: RIGHT

Name	Position	Size (bp)	Type	Description	Application notes
XhoI	1022-1027	6	misc_feature	*None*	*None*
hRPH3A_c.444Tmut	1028-1411	384	misc_feature	*None*	note=hRPH3A_c.444Tmut
BamHI	1412-1417	6	misc_feature	*None*	*None*
RRE	2857-3090	234	misc_feature	*None*	full_name=HIV-1 Rev response element
gp41 peptide	3275-3319	45	ORF	*None*	codon_start=1 product=antigenic peptide corresponding to amino acids 655 to 669 of the HIV envelope protein gp41 (Lutje Hulsik et al., 2013)
SV40 poly(A) signal	3635-3769	135	polyA_signal	*None*	*None*
AmpR promoter	4443-4547	105	promoter	*None*	gene=bla note=Unknown feature type:Promoter color: #fd3434; direction: RIGHT
AmpR	4548-5408	861	ORF	*None*	codon_start=1 full_name=Ampicillin resistance gene gene=bla product=beta-lactamase
ori	5579-6167	589	rep_origin	*None*	*None*

Note: Components added by user are listed in **bold red** text.

Vector Sequence

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   1 CTGTGGAATG TGTGTCAGTT AGGGTGTGGA AAGTCCCCAG GCTCCCCAGC AGGCAGAAGT ATGCAAAGCA TGCATCTCAA
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 241 CTAATTTTTT TTATTTATGC AGAGGCCGAG GCCGCCTCGG CCTCTGAGCT ATTCCAGAAG TAGTGAGGAG GCTTTTTTGG
 321 AGGCCTAGGC TTTTGCAAAA AGCTTGGACT GTGTTTACTT GCAATCCCCC AAAACAGACA GAATGGTGCA TCTGTCCAGT
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 641 GCTAAGCTGA GTGAACTGCA CTGTGACAAG CTGCACGTGC TCTAGAGTCG ACCCAGCAGT AAGTAATACA TGTAATGCAA
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1041 TTGCTCATAG AAGGAAGTGG GAGAAATGGA AGCAAGTTCT TCCTCTTTAC ATCCACCAAG AGAAAGTGGG GGGCTACGTT
1121 GCCATGCCTC CCATGAGCAT GGCTCATCTG AGTGTGTTGG CTGTGTTTCA TCCACAGAAC GTCTGCACCA AGTGCGGAGT
1201 GGAGACCAAC AACCGCCTGC ATTCTGTGTG GCTCTGCAAA ATCTGCATTG AGCAGAGGGA TGTGAGTGCC CTGGTCCCAC
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6241 TATCCCCTGA TTCTGTGGAT AACCGTATTA CCGCCTTTGA GTGAGCTGAT ACCGCTCGCC GCAGCCGAAC GACCGAGCGC
6321 AGCGAGTCAG TGAGCGAGGA AGCGGAAGAG CGCCCAATAC GCAAACCGCC TCTCCCCGCG CGTTGGCCGA TTCATTAATG
6401 CAG
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Validation by Restriction Enzyme Digestion

Restriction Enzymes	Cutting Sites	DNA Fragments (bp)
XhoI	1023	6403
NdeI	1493	6403
ApaLI	1703, 4664, 5910	2961, 1246, 2196
EcoRV	1423	6403
BamHI	1413	6403
ApaLI+XhoI	1023, 1703, 4664, 5910	680, 2961, 1246, 1516
ApaLI+NdeI	1493, 1703, 4664, 5910	210, 2961, 1246, 1986
ApaLI+BamHI	1413, 1703, 4664, 5910	290, 2961, 1246, 1906
ApaLI+EcoRV	1423, 1703, 4664, 5910	280, 2961, 1246, 1916