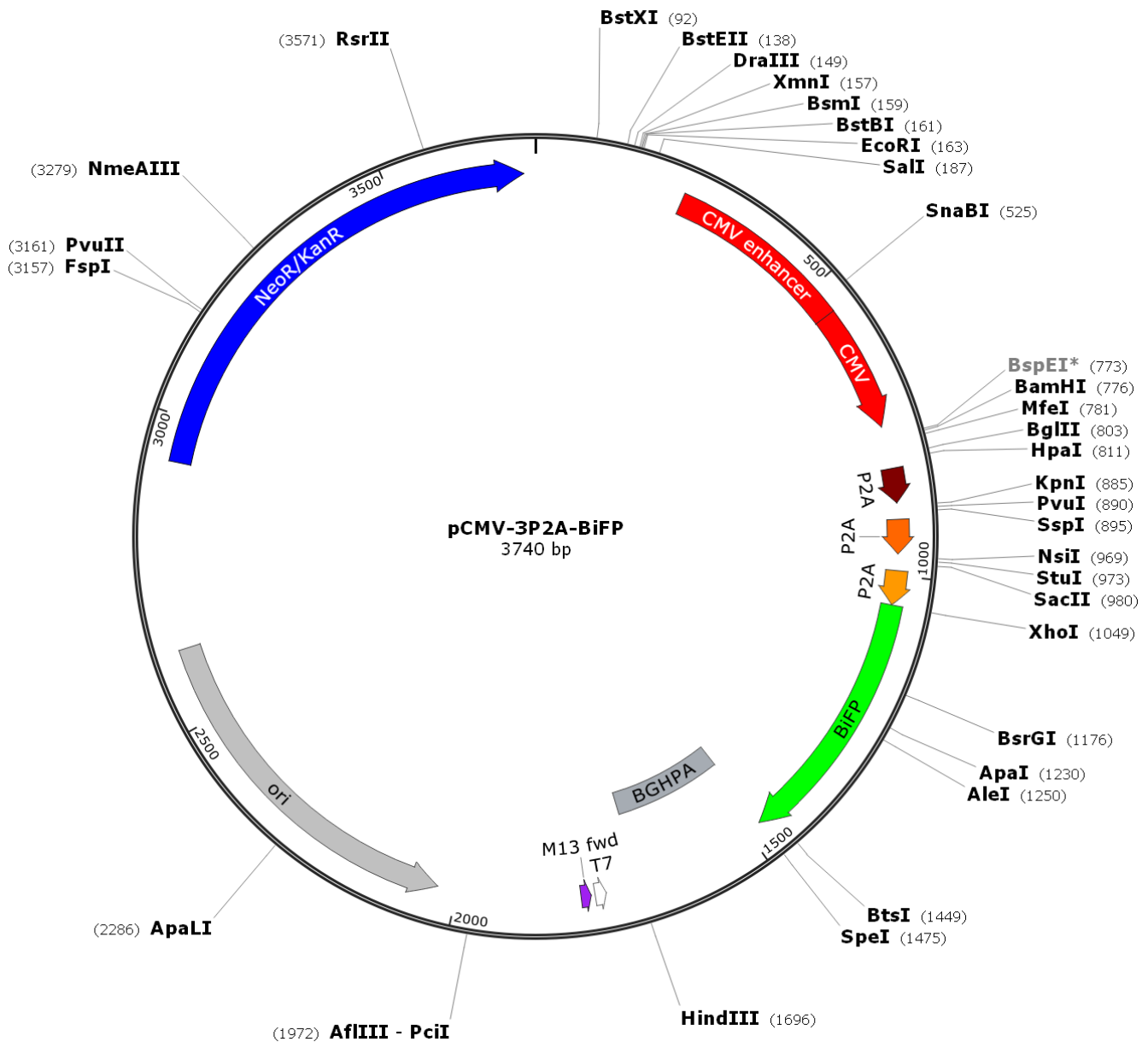


Vector: pCMV-3P2A-BiFP (based on pMOK Vector)

Antibiotic Selection: Kan

Creator(s): Hao Wang & Linjuan Huang @ Molecular Oncology Lab of The University of Chicago Medical Center

Date of Construction: April 2020



3xP2A Coding Regions and Restriction Enzyme Sites

1 GGATCC AATTGCAATGATCATCATGAC AGATCT GTTAAC ggaagc BamHI BglII HpaI
G S N C N D H H D R S V N G S

46 ggagctactaacttcagcctgctgaagcaggctggagacgtggag
G A T N F S L L K Q A G D V E

91 gagaaccctggacct GGTACC CGATCG AATATT ggatctggcgcc KpnI PvuI SspI
E N P G P G T R S N I G S G A

136 accaacttttccctcctgaaacaggctggcgatgtcgaagaaat
T N F S L L K Q A G D V E E N

181 cccggcctt ATGCAT AGGCCT CCGCGG ggctccggcgccaccaac NsiI StuI SacII
P G P M H R P P R G S G A T N

226 ttctccctgctgaagcaggccggcgacgtggaggagaacccccggc
F S L L K Q A G D V E E N P G

271 ccc CTCGAG atggtggagaaattcgtggaaactggaaaatcgca XhoI BiFP
P L E M V E K F V G T W K I A

316 gacagtcataactttggagagtacctaaggcaatcggagcccct
D S H N F G E Y L K A I G A P

361 aaggaactgtctgacggcggagatgccaccacaccaaccctgtac
K E L S D G G D A T T P T L Y

406 atcagccagaaggacggagataaaatgacagtgaagattgagaac
I S Q K D G D K M T V K I E N

451 gggccccctacattcctggacactcaggtgaagttcaaactgggc
G P P T F L D T Q V K F K L G

496 gaggagttcgacgagtttcccagcgataggagaaaggggtgaaa
E E F D E F P S D R R K G V K

541 tccgtggtcaatctggtcggcgaaaagctggtgtacgtccagaaa
S V V N L V G E K L V Y V Q K

586 tgggacggcaaggagactacctaagtgcgggaaatcaaggatgga
W D G K E T T Y V R E I K D G

631 aaactggtcgtgaccctgacaatgggagatgtggtggcagtgcg
K L V V T L T M G D V V A V R

676 agttacagacgggctacagaa TAA
S Y R R A T E *

pCMV-3P2A-BiFP Full-Length Sequence

GGAAACAGCTATGACCATGATTACGCCAAGCTCGAAATTACCCCTCACTAAAGGGAACAAAAGCTGGTACGAGGACAGGCTGG
 AGCCATGGGCATGGCTACTCAAGCTGATTTGATGGAGTTGGACATGGCCATGGCTGGTGACCACGTCGTGGAATGCCCTTC**GA**
ATTCAGCACCTGCACATGGGAC**CTCGAC****TAATAGTAATCAATTACGGGGTCATTAGTTTCATAGCCCATATATGGAGTTCCGC**
GTTACATAACTTACGGTAAATGGCCCGCTGGCTGACCGCCCAACGACCCCGCCCATTTGACGTCAATAATGACGTATGTT
 CCATAGTAACGCCAATAGGGACTTTCCATTGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTGGCAGTACATCA
 AGTGTATCATATGCCAAGTACGCCCCCTATTGACGTCAATGACGGTAAATGGCCCGCTGGCATTATGCCAGTACATGACC
 TTATGGGACTTTCCCTACTTGGCAGTACATCTACGTATTAGTCACTGCTATTACCATGGTGATGCGGTTTTGGCAGTACATCA
 ATGGGCGTGGATAGCGGTTTGACTCACGGGGATTTCCAAGTCTCCACCCCATTTGACGTCAATGGGAGTTTTGTTTTGGCACCA
 AAATCAACGGGACTTTCCAAAATGTCGTAACAACCTCCGCCCATTTGACGCAAAATGGGCGGTAGGCGTGTACGGTGGGAGGTC
 TATATAAGCAGAGCTGGTTTAGTGAACCGTCAGATCC**GGATCC**AATTGCAATGATCATCATGAC**AGATCT**GTTAAC**ggaagc**
ggagctactaacttcagcctgctgaagcaggctggagacgtggaggagaaccctggacct**GGTACC**CGATCG**AATATT**ggat
ctggcgccaccaacttttccctcctgaaacaggctggcgatgtcgaagaaaatcccggccct**ATGCAT**AGGCCCTCCGGG**gg**
ctccggcgccaccaacttctccctgctgaagcaggccggcgacgtggaggagaacccccggcccc**CTCGAG**atggtggagaaa
 ttctggtggaacctggaaaatcgagacagtcataactttggagagtacctgaaggcaatcgagcccctaaggaactgtctg
 acggcgagatgccaccacaccaacctgtacatcagccagaaggacggagataaaaatgacagtgaagattgagaacggggcc
 ccctacattcctggacactcagtggaagtt**ca**aactgggagagttcgacgagtttccagcgataggagaaaggggtg
 aatccgtggtcaatctggtcggcgaaaagctggtgtacgtccagaaatgggacggcaaggagactacctatgtcgggaaa
 tcaaggatggaaaactggtcgtgaccctgacaatgggagatgtggtggcagtgcgagttacagacgggctacagaataa**AC**
TAGTcctcgaactgtgccttctagtgtgccagccatctggtgtttgccctccccctgaccttccctgacctggaagtgcca
 ctcccactgtcctttcctaataaaatgaggaattgcatcgcattgtctgagttaggtgtcattctattctgggggtggggt
 ggggcaggacagcaaggggaggattgggaagacaatagcaggcatgctggggat**AAGCTT**TTAAATAA**GGAGGAATAACAT**
ATGACCATGATTAC**GCC**AAGCT**CCAATTC**GCCCTATAGTGAGTCGTATTACAATTCAGTGGCCGTC**GTTTTACTATGCGGTTG**
TGAAATACCGCACAGATGCGTAAGGAGAAAAATACCGCATCAGGCGCTCTTCCGCTTCCCTCGCTCACTGACTCGCTGCGCTCG
 GTCGTTCCGGCTGCGGCGAGCGGTATCAGCTCAAAAGCGGTAATACGGTTATCCACAGAATCAGGGGATAACCGCAGAA
 AGAACATGTGAGCAAAAAGCCAGCAAAAAGCCAGGAACCGTAAAAAGCCCGCTTGCTGGCGTTTTTCCATAGGCTCCGCC
 CCTGACGAGCATCACAAAAATCGACGCTCAAGTCAAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGGCGTTTTCCC
 CCTGGAAGCTCCCTCGTGCCTCTCCTGTTCCGACCCGCGCTTACCGGATACCTGTCCGCCTTTTCTCCCTTCGGGAAGCG
 TGGCGCTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCCGCTCCAAGCTGGGCTGTGTGCACGAACC
 CCCCCTCAGCCGACCGCTGCGCCTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGCCACTG
 GCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGTACAGAGTTCCTGAAGTGGTGGCCTAACACG
 GCTACACTAGAAGGACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTTGATC
 CGGCAAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTTGTTTGAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAA
 GATCCTTTGATCTTTTCTACGGGTCTGACGCTCAGTGAACGAAAACCTCACGTTAAGGGATTTTGGTCATGAGATTATCAA
 AAAGGATCTTCACCTAGATCCTTTTAAATTAATAAATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAAACTTGGTCTGA
 CAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCTGACTCCCCGTC
 ATTCAAATATGTATCCGCTCATGAGACAATAACCCGATAAAATGCTTCAATAATAT**ATG**attgaacaagatggattgcacgc
 aggttctccggccgcttgggtggagaggctattcggctatgactgggcacaacagacaatcggctgctctgatgccgccgtg
 ttccggctgtcagcgcaggggcccgggttcttttgtcaagaccgacctgtccggtgccctgaatgaactgcaagacgagg
 cagcgcggctatcgtggctggccacgacgggcttccctgctcagctgtgctcagcgttgtcactgaagcgggaagggactg
 gctgctattgggcgaagtgccggggcaggatctcctgtcatctcacctgtcctgcccagaaaagtatccatcatggctgat
 gcaatgcggcggtgcatacgttgatccggctacctgccattcagaccaccaagcgaacatcgcatcgagcagacagcga
 ctcgatggaagccggtcttctgcatcaggatgatctggacgaagagcatcaggggctcgcgccagccgaactgttcgccag
 gctcaaggcagcagatgcccagcggcgaggatctcgtcgtgacctggtcgatgctgcttgcgcaatatcatggtggaaaaat
 ggccgctttctgattcctcagactgtggcggctgggtgtggcggaccgctatcaggacatagcgttggctaccctgata
 ttgctgaagagcttggcggcgaatgggctgaccgcttccctcgtgctttacggtatcgccgctcccgattcgcagcgcacgc
 cttctatcgccttcttgacgagttcttc**TGA**CCTTTCGTCTTCAAGaatt

Zero Cutters:

#	Enzyme	Specificity			
1	AclI	AACGTT	13	BmtI	GCTAGC
2	AfeI	AGCGCT	14	Bpu10I	CCTNAGC
3	AflIII	CTTAAG	15	BsaI	GGTCTCNNNNN
4	AgeI	ACCGGT	16	BsiWI	CGTACG
5	AscI	GGCGCGCC	17	BspDI	ATCGAT
6	AseI	ATTAAT	18	BssHII	GCGCGC
7	AsiSI	GCGATCGC	19	BstAPI	GCANNNNTGC
8	AvrII	CCTAGG	20	BstZ17I	GTATAC
9	BaeI	(N) 5 (N) 10ACNNGTAYC (N) 7 (N) 5	21	ClaI	ATCGAT
10	BbvCI	CCTCAGC	22	Eco53kI	GAGCTC
11	BcgI	NN (N) 10CGA (N) 6TGC (N) 10NN	23	EcoRV	GATATC
12	BlpI	GCTNAGC	24	FseI	GGCCGGCC
			25	MluI	ACGCGT
			26	NheI	GCTAGC

27 NotI GCGGCCGC
 28 NruI TCGCGA
 29 PacI TTAATTA
 30 PmeI GTTTAAAC
 31 PmlI CACGTG
 32 PpuMI RGGWCCY
 33 PshAI GACNNNGTC
 34 PsiI TTATAA
 35 PspXI VCTCGAGB
 36 PstI CTGCAG
 37 SacI GAGCTC

38 SbfI CCTGCAGG
 39 ScaI AGTACT
 40 SfiI GGCCNNNNGGCC
 41 SgrAI CRCCGGYG
 42 SmaI CCCGGG
 43 SrfI GCCC GGCC
 44 SwaI ATTTAAAT
 45 TspMI CCCGGG
 46 XbaI TCTAGA
 47 XcmI CCANNNNNNTGG
 48 XmaI CCCGGG

One-Cutters:

#	Enzyme	Specificity	Sites &
flanks	Cut positions		
(blunt - 5' ext. - 3' ext.)			
1	Acc65I	GGTACC	list
2	AccI	GTMKAC	list
3	AflIII	ACRYGT	list
4	AleI	CACNNNGTG	list
5	ApaI	GGGCCC	list
6	ApaLI	GTGCAC	list
7	AvaI	CYCGRG	list
8	BamHI	GGATCC	list
9	BclI	TGATCA	list
10	BglII	AGATCT	list
11	BsaBI	GATNNNNATC	list
12	BsaXI	NNN (N) 9AC (N) 5CTCC (N) 7NNN	list
13	BsgI	GTGCAG (N) 14NN	list
14	BsmBI	CGTCTCNNNNN	list
15	BsmI	GAATGCN	list
16	BsoBI	CYCGRG	list
17	BspEI	TCCGGA	list
18	BsrGI	TGTACA	list
19	BstBI	TTCGAA	list
20	BstEII	GGTNACC	list
21	BstXI	CCANNNNNNTGG	list
22	Bsu36I	CCTNAGG	list

23	BtsI	GCAGTGNN	list
24	CspCI	NN (N) 11CAA (N) 5GTGG (N) 10NN	list
25	DraIII	CACNNNGTG	list
26	EagI	CGGCCG	list
27	EcoNI	CCTNNNNNAGG	list
28	EcoO109I	RGGNCCY	list
29	EcoRI	GAATTC	list
30	Esp3I	CGTCTCNNNNN	list
31	FspI	TGCGCA	list
32	HindIII	AAGCTT	list
33	HpaI	GTTAAC	list
34	KpnI	GGTACC	list
35	MfeI	CAATTG	list
36	NmeAIII	GCCGAG (N) 19NN	list
37	NsiI	ATGCAT	list
38	Paer7I	CTCGAG	list
39	PciI	ACATGT	list
40	PflMI	CCANNNNNNTGG	list
41	PspOMI	GGGCC	list
42	PvuI	CGATCG	list
43	PvuII	CAGCTG	list
44	RsrII	CGGWCCG	list
45	SacII	CCGCGG	list
46	SalI	GTGCAC	list
47	SexAI	ACCWGGT	list
48	SnaBI	TACGTA	list
49	SpeI	ACTAGT	list