

Supplementary information:

Primer name	Sequence	Usage
SY094	ACCAGGTTTCGTTCACTCATGGAAAATAGCG	Cloning of C-terminal Cre
SY095	GCGGCCGCCTAATCGCCATCTTCCAGCAGGCGCACCAT	Cloning of C-terminal Cre
SY096	GATATCGCCGCCACCATGCCCAAGAAGAAGAGGAAGGTG	Cloning of PhyB
SY097	GCATGCAACAGCTCCCAAACACTTC	Cloning of PhyB
SY118	CCGGAATTCATGCCCAAGAAGAAGAGGAAG	Subclone of PIF6CreN fragment
SY119	GCTCTCGAGTCAGTTCAGCTTGCACCAGG	Subclone of PIF6CreN fragment

Table S1. Primers

Addgene #	Name	Application
131780	pCS2+ PIF6CreN	For <i>in vitro</i> transcription of PIF6APBCreN
131781	pCS2+ PhyBΔCreC	For <i>in vitro</i> transcription of PhyBΔCreC

Table S2. Plasmids

Sequences

230 bp fragment

AGCCGGAGTCAGCCATGGGAAACTGCGGAAATGGATGCGATTCACTCGCTCCAGCT
TATTCTGAGAGACTCTTTTAAAGAATCTGGTGGTTCTGCCGGTGGCTCCGGTTCTGGC
TCCAGCGGTGGCAGCTCTGGTGCCTCCGGCACGGGTACTGCGGGTGGCACTGGCAG
CGGTTCCGGTACTGGCTCTGGCAACCGGAAATGGTTTCCCGCAGAACCTGAAGATGT
TCG

PhyBΔCreC

ATGCCAAGAAGAAGAGGAAGGTGTTTCCGGAGTCGGGGTAGTGGCGGTGGCCGTGGCAGTGGCCGTG
GCGGAGAAGAAGAACCGTCGTCAAGTCACACTCCTAATAACCGAAGAGGAGGAGAACAAGCTCAATCGTC
GGGAACGAAATCTCTCAGACCAAGAAGCAACACTGAATCAATGAGCAAAGCAATTCAACAGTACACCGTC
GACGCAAGACTCCACGCCGTTTTCGAACAATCCGGCGAATCAGGGAAATCATTGACTACTCACAATCAC
TCAAAACGACGACGTACGGTTCCTCTGTACCTGAGCAACAGATCACAGCTTATCTCTCTCGAATCCAGCG
AGGTGGTTACATTCAGCCTTTCGGATGTATGATCGCCGTCGATGAATCCAGTTTCCGGATCATCGGTTAC
AGTGAAAACGCCAGAGAAATGTTAGGGATTATGCCTCAATCTGTTCTACTCTTGAGAAACCTGAGATTC
TAGCTATGGAACTGATGTGAGATCTTGTTCACTTCTTCGAGCTCGATTCTACTCGAGCGTGCTTTCGT
TGCTCGAGAGATTACCTTGTTAAATCCGGTTTGGATCCATTCCAAGAATACTGGTAAACCGTTTTACGCC
ATTCTTCATAGGATTGATGTTGGTGTGTTATTGATTTAGAGCCAGCTAGAACTGAAGATCCTGCGCTTT
CTATTGCTGGTGTGTTCAATCGCAGAACTCGCGTTCGTGCGATTTCTCAGTTACAGGCTCTTCCTGG
TGGAGATATTAAGCTTTTGTGTGACACTGTCGTGGAAAGTGTGAGGGACTTGACTGGTTATGATCGTGTT
ATGGTTTATAAGTTTCATGAAGATGAGCATGGAGAAGTTGTAGCTGAGAGTAAACGAGATGATTTAGAGC
CTTATATTGGACTGCATTATCCTGCTACTGATATCCTCAAGCGTCAAGGTTCTTGTTAAGCAGAACCG
TGTCCGAATGATAGTAGATTGCAATGCCACACCTGTTCTTGTGGTCCAGGACGATAGGCTAACTCAGTCT
ATGTGCTTGGTTGGTTCTACTCTTAGGGCTCCTCATGGTTGTCACTCTCAGTATATGGCTAACATGGGAT
CTATTGCGTCTTAGCAATGGCGGTTATAATCAATGGAAATGAAGATGATGGGAGCAATGTAGCTAGTGG
AAGAAGCTCGATGAGGCTTGGGGTTGGTTGTTTGCATCACACTTCTTCTCGCTGCATACCGTTTCCG
CTAAGGTATGCTTGTGAGTTTTTGTGAGGCTTTCGGTTACAGTTAAACATGGAATTGCAGTTAGCTT
TGCAAATGTCAGAGAAACGCGTTTTGAGAACGCAGACACTGTTATGTGATATGCTTCTGCGTGACTCGCC
TGCTGGAATTGTTACACAGAGTCCAGTATCATGGACTTAGTGAAATGTGACGGTGCAGCATTTCTTTAC
CACGGGAAGTATTACCGTTGGGTGTTGCTCCTAGTGAAGTTCAGATAAAAGATGTTGTGGAGTGGTTGC
TTGCGAATCATGCGGATTCAACCGGATTAAGCACTGATAGTTTAGGCGATGCGGGGTATCCCGGTGCAGC
TGCGTTAGGGGATGCTGTGTGCGGTATGGCAGTTGCATATATCACAAAAGAGACTTTCTTTTTTGGTTT
CGATCTCACACTGCGAAAGAAATCAAATGGGGAGGCGCTAAGCATCATCCGGAGGATAAAGATGATGGGC
AACGAATGCATCCTCGTTCGTCCTTTCAGGCTTTTCTTGAAGTTGTTAAGAGCCGGAGTCAGCCATGGGA
AACTGCGGAAATGGATGCGATTCACTCGCTCCAGCTTATTCTGAGAGACTCTTTTAAAGAATCTGGTGGT
TCTGCCGGTGGCTCCGGTTCGGCTCCAGCGGTGGCAGCTCTGGTGCCTCCGGCACGGGTACTGCGGGTG
GCACTGGCAGCGGTTCCGGTACTGGCTCTGGCAACCGGAAATGGTTTCCCGCAGAACCTGAAGATGTTTCG

CGATTATCTTCTATATCTTCAGGCGCGGGTCTGGCAGTAAAACTATCCAGCAACATTTGGGCCAGCTA
AACATGCTTCATCGTCGGTCCGGGCTGCCACGACCAAGTGACAGCAATGCTGTTTCACTGGTTATGCGGC
GGATCCGAAAAGAAAACGTTGATGCCGGTGAACGTGAAAACAGGCTCTAGCGTTCGAACGCACTGATTT
CGACCAGTTTCGTTCACTCATGGAAAATAGCGATCGCTGCCAGGATATACGTAATCTGGCATTCTGGGG
ATTGCTTATAACACCCTGTTACGTATAGCCGAAATTGCCAGGATCAGGGTTAAAGATATCTCACGTA
ACGGTGGGAGAATGTTAATCCATATTGGCAGAACGAAAACGCTGGTTAGCACCCGAGGTGTAGAGAAGGC
ACTTAGCCTGGGGTA
ACTAACTGGTCGAGCGATGGATTTCCGTCTCTGGTGTAGCTGATGATCCGAAT
AACTACCTGTTTTGCCGGTCAGAAAAATGGTGTGCCGCGCCATCTGCCACCAGCCAGCTATCAACTC
GCGCCCTGGAAGGGATTTTTGAAGCAACTCATCGATTGATTTACGGCGCTAAGGATGACTCTGGTCAGAG
ATACCTGGCCTGGTCTGGACACAGTGCCCGTGTGCGAGCCGCGGAGATATGGCCCGCGCTGGAGTTTCA
ATACCGGAGATCATGCAAGCTGGTGGCTGGACCAATGTAATATTGTCATGAACTATATCCGTAACCTGG
ATAGTGAAACAGGGGCAATGGTGCCTGCTGGAAGATGGCGATTAG

NLS

PhyBΔ (1-621 a.a.)

GSAT linker (36 a.a.)

CreC (60 – 343 a.a.)

PIF6CreN

ATGCCAAGAAGAAGAGGAAGGTGATGTTCTTACCAACCGATTATTGTTGCAGGTTAAGCGATCAAGAGT
ATATGGAGCTTGTGTTGAGAATGGCCAGATTCTTGCAAAGGGCCAAAGATCCAACGTTTCTCTGCATAA
TCAACGTACCAAATCGATCATGGATTTGTATGAGGCAGAGTATAACGAGGATTTTCATGAAGAGTATCATC
CATGGTGGTGGTGGTGGCCATCACAATCTCGGGACACGCAGGTTGTTCCACAAAGTCATGTTGCTGCTG
CCCATGAAACAAACATGTTGAAAGCAATAAACATGTTGACGGTGGTTCTGCCGGTGGCTCCGGTTCTGG
CTCCAGCGGTGGCAGCTCTGGTGCCTCCGGCACGGGTA
CTGCGGTGGCACTGGCAGCGGTTCCGGTACT
GGCTCTGGCCTGACTGTGCACCAAACCTGCCTGCCCTCCCTGTGGATGCCACCTCTGATGAAGTCAGGA
AGAACCTGATGGACATGTTCAAGGACAGGCAGGCTTCTCTGAACACACCTGGAAGATGCTCCTGTCTGT
GTGCAGATCCTGGGCTGCCTGGTGCAAGCTGAAC

NLS

PIF6APB (1 – 100 a.a.)

GSAT linker (36 a.a.)

CreN (19 – 59 a.a.)

PhyBΔCreC-P2A-PIF6CreN

ATGCCCAAGAAGAAGAGGAAGGTGGTTTCCGGAGTCGGGGTAGTGGCGGTGGCCGTGGCGGTGGCCGTG
GCGGAGAAGAAGAACCCTCGTCAAGTCACACTCCTAATAACCGAAGAGGAGGAGAACAAGCTCAATCGTC
GGGAACGAAATCTCTCAGACCAAGAAGCAACACTGAATCAATGAGCAAAGCAATTCAACAGTACACCGTC
GACGCAAGACTCCACGCCGTTTTCGAACAAATCCGGCGAATCAGGGAAATCATTGACTACTCACAATCAC
TCAAAACGACGACGTACGGTTCCTCTGTACCTGAGCAACAGATCACAGCTTATCTCTCTCGAATCCAGCG
AGGTGGTTACATTCAGCCTTTCGGATGTATGATCGCCGTCGATGAATCCAGTTTCCGGATCATCGGTTAC
AGTGAAAACGCCAGAGAAATGTTAGGGATTATGCCCTCAATCTGTTCCCTACTCTTGAGAAACCTGAGATTC
TAGCTATGGAACTGATGTGAGATCTTTGTTCACTTCTTCGAGCTCGATTCTACTCGAGCGTGCTTTCGT
TGCTCGAGAGATTACCTTGTTAAATCCGGTTTGGATCCATTCCAAGAATACTGGTAAACCGTTTTACGCC
ATTCTTCATAGGATTGATGTTGGTGTGTTATTGATTTAGAGCCAGCTAGAACTGAAGATCCTGCGCTTT
CTATTGCTGGTGTGTTCAATCGCAGAACTCGCGGTTCTGTCGATTTCTCAGTTACAGGCTCTTCTCGG
TGGAGATATTAAGCTTTTGTGTGACACTGTCGTGGAAAGTGTGAGGGACTTGACTGGTTATGATCGTGT
ATGGTTTATAAGTTTTCATGAAGATGAGCATGGAGAAGTTGTAGCTGAGAGTAAACGAGATGATTTAGAGC
CTTATATTGGACTGCATTATCCTGCTACTGATATCCTCAAGCGTCAAGGTTCTTGTTAAGCAGAACCG
TGTCCGAATGATAGTAGATTGCAATGCCACACCTGTTCTTGTGGTCCAGGACGATAGGCTAACTCAGTCT
ATGTGCTTGGTTGGTCTACTCTTAGGGCTCCTCATGGTTGTCACTCTCAGTATATGGCTAACATGGGAT
CTATTGCGTCTTTAGCAATGGCGGTTATAATCAATGGAAATGAAGATGATGGGAGCAATGTAGCTAGTGG
AAGAAGCTCGATGAGGCTTTGGGGTTTGGTTGTTTGCCATCACACTTCTTCTCGCTGCATACCGTTTCCG
CTAAGGTATGCTTGTGAGTTTTTGTGATGCAGGCTTTTCGGTTTACAGTTAAACATGGAATTGCAGTTAGCTT
TGCAAATGTCAGAGAAACGCGTTTTGAGAACGCAGACACTGTTATGTGATATGCTTCTGCGTGACTCGCC
TGCTGGAATTGTTACACAGAGTCCCAGTATCATGGACTTAGTGAAATGTGACGGTGCAGCATTTCTTTAC
CACGGGAAGTATTACCCGTTGGGTGTTGCTCCTAGTGAAGTTCAGATAAAAGATGTTGTGGAGTGGTTGC
TTGCGAATCATGCGGATTCAACCGGATTAAGCACTGATAGTTTAGGCGATGCGGGGTATCCCGGTGCAGC
TGCGTTAGGGGATGCTGTGTGCGGTATGGCAGTTGCATATATCACAAAAGAGACTTTCTTTTTTGGTTT
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AACGAATGCATCCTCGTTCGTCCTTTCAGGCTTTTCTTGAAGTTGTTAAGAGCCGGAGTCAGCCATGGGA
AACTGCGGAAATGGATGCGATTCACTCGCTCCAGCTTATTCTGAGAGACTCTTTTAAAGAATCTGGTGGT
TCTGCCGGTGGCTCCGTTCTGGCTCCAGCGGTGGCAGCTCTGGTGCCTCCGGCACGGGTACTGCGGGTG
GCACTGGCAGCGGTTCCGGTACTGGCTCTGGCAACCGGAAATGGTTTCCCGCAGAACCTGAAGATGTTTCG
CGATTATCTTCTATATCTTACGGCGCGCGGTCTGGCAGTAAAACTATCCAGCAACATTTGGGCCAGCTA
AACATGCTTCATCGTCCGTTCCGGGCTGCCACGACCAAGTACAGCAATGCTGTTTCACTGGTTATGCGGC
GGATCCGAAAAGAAAACGTTGATGCCGGTGAACGTGCAAAACAGGCTCTAGCGTTTGAACGCACTGATTT
CGACCAGGTTTCTTCACTCATGGAAAATAGCGATCGCTGCCAGGATATACGTAATCTGGCATTCTGGGG
ATTGCTTATAACACCCTGTTACGTATAGCCGAAATGGCCAGGATCAGGGTTAAAGATATCTCACGTACTG
ACGGTGGGAGAATGTTAATCCATATTGGCAGAACGAAAACGCTGGTTAGCACCCGAGGTGTAGAGAAGGC

ACTTAGCCTGGGGTAACTAACTGGTCGAGCGATGGATTTCCGTCTCTGGTGTAGCTGATGATCCGAAT
AACTACCTGTTTTGCCGGGTCAGAAAAAATGGTGTGCGCGCCATCTGCCACCAGCCAGCTATCAACTC
GCGCCCTGGAAGGGATTTTTGAAGCAACTCATCGATTGATTTACGGCGCTAAGGATGACTCTGGTCAGAG
ATACCTGGCCTGGTCTGGACACAGTGCCCGTGTCTGGAGCCGCGCGAGATATGGCCCGCGCTGGAGTTTCA
ATACCGGAGATCATGCAAGCTGGTGGCTGGACCAATGTAAATATTGTCATGAACTATATCCGTAACCTGG
ATAGTGAAACAGGGGCAATGGTGCCTGCTGGAAGATGGCGATGGAAGCGGAGCTACTAACTTCAGCCT
GCTGAAGCAGGCTGGAGACGTGGAGGAGAACCCTGGACCTATGCCCAAGAAGAAGAGGAAGGTGATGTTC
TTACCAACCGATTATTGTTGCAGGTTAAGCGATCAAGAGTATATGGAGCTTGTGTTTGAGAATGGCCAGA
TTCTTGCAAAGGGCCAAAGATCCAACGTTTCTCTGCATAATCAACGTACCAAATCGATCATGGATTTGTA
TGAGGCAGAGTATAACGAGGATTTTCATGAAGAGTATCATCCATGGTGGTGGTGGTGGCCATCACAAATCTC
GGGGACACGCAGGTTGTTCCACAAAGTCATGTTGCTGCTGCCCATGAAACAAACATGTTGGAAAGCAATA
AACATGTTGACGGTGGTCTGCCGGTGGCTCCGGTCTGGCTCCAGCGGTGGCAGCTCTGGTGCCTCCGG
CACGGGTACTGCGGGTGGCACTGGCAGCGGTTCCGGTACTGGCTCTGGCCTGACTGTGCACCAAACCTG
CCTGCCCTCCCTGTGGATGCCACCTCTGATGAAGTCAGGAAGAACCTGATGGACATGTTCAGGGACAGGC
AGGCCTTCTCTGAACACACCTGGAAGATGCTCCTGTCTGTGTGCAGATCCTGGGCTGCCTGGTGCAGCT
GAACTGA

PhyB Δ CreC (60-343 a.a.)

GSG-P2A (Kim et al., 2011)

PIF6APBCreN