

Supporting information

RNA Aptamer-Mediated Gene Activation Systems for Inducible Transgene Expression in Animal Cells

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Table S1. Sequences of trigger RNAs

Trigger RNA	Sequence (5' → 3')
MP _{x1}	GGUGAGUAGAGCGAGGAGCAUCAGCCCUCGCGUGUGACGAGCAGACCAUAU GGGGUCGCUCG
MP _{x2}	GGUCCACCAGAGUAGAGCAUCAGCCUACUCCGGACAAGCAGCAGAGGAUAU GGCCUCGCGUCGAGUAGAGCGAGGAGCAUCAGCCCUCGCGUGUGACGAGCA GACCAUAUGGGGUCGCUCG
MP _{x4}	GGUUCUGCCAUGAGGAAUGACCAUCAGGCAUUCCGAUCCGAGGAGCAGACG AUAUGGCGUCGCUCGCGCAGAAUGAGACGAUCACGCACUUCGGAGUGAC CAGCAGAGCAUAUGGGCUCGCGUGGGCCACCAGAGUAGAGCAUCAGCCUACU CCGGACAAGCAGCAGAGGAUAUGGCCUCGCGAGUAGAGCGAGGAGCAU CAGCCCUCGCGUGUGACGAGCAGACCAUAUGGGGUCGCUCG
MP _{x1S}	GGUGAGUAGAGCGAGGAGCAUCAGCCCUCGCGUGUGACGAGCAGACCAUAU GGGGUCGCUCGUCACACAAACUACUCUU
MP _{x2S}	GGUCCACCAGAGUAGAGCAUCAGCCUACUCCGGACAAGCAGCAGAGGAUAU GGCCUCGCGUCGAGUAGAGCGAGGAGCAUCAGCCCUCGCGUGUGACGAGCA GACCAUAUGGGGUCGCUCGUCACACAAACUACUCAAUGUCCGAAAGGUGG A
MP _{x4S-α}	GGUUCUGCCAUGAGGAAUGACCAUCAGGCAUUCCGAUCCGAGGAGCAGACG AUAUGGCGUCGCUCGCGCAGAAUGAGACGAUCACGCACUUCGGAGUGAC CAGCAGAGCAUAUGGGCUCGCGUGGGCCACCAGAGUAGAGCAUCAGCCUACU CCGGACAAGCAGCAGAGGAUAUGGCCUCGCGAGUAGAGCGAGGAGCAU CAGCCCUCGCGUGUGACGAGCAGACCAUAUGGGGUCGCUCGUCACACAAAC UACUCAAUGUCCGAAAGGUGGCAAACACUCCAAAGCAGCCAAACGGAUCA ACAUGGCAGCGGUGCUU
MP _{x4S-β}	GGUUCUGCCAUGAGGAAUGACCAUCAGGCAUUCCGAUCCGAGAAGUGACGA UCACGCACUUCGGCUCGAGGAGCAGACGAUAUGGCGUCGCUCGCGAGUGAC CAGCAGAGCAUAUGGGCUCGCGUGGGCCACCAGAGUAGAGCAUCAGCCUACU CCGGACAAGCGAGGAGCAUCAGCCCUCGCGAGUAGAGCAGCAGAGGAUAUG GCCUCGCGUCGUGUGACGAGCAGACCAUAUGGGGUCGCUCGUCACACAAAC UACUCAAUGUCCGAAAGGUGGCAAACACUCCAAAGCAGCCAAACGGAUCA ACAUGGCAGCGGUGCUU
MP _{x4S-γ}	GGUUCUGCCAUGAGGAAUGACCAUCAGGCAUUCCGAUCCGAGAAGUGACGA UCACGCACUUCGGCUCGAGAGUAGAGCAUCAGCCUACUCCGAGUGAGCGAG GAGCAUCAGCCCUCGCGCCACCAGGAGCAGACGAUAUGGCGUCGCUCGCGG ACAACCAGCAGAGCAUAUGGGCUCGCGUGGGAGUAGAGCAGCAGAGGAUAUG GCCUCGCGUCGUGUGACGAGCAGACCAUAUGGGGUCGCUCGUCACACAAAC UACUCAAUGUCCGAAAGGUGGCAAACACUCCAAAGCAGCCAAACGGAUCA ACAUGGCAGCGGUGCUU
MC _{x4S}	GGUUCUGCCAUGAGGAAUGACCAUCAGGCAUUCCGAUCCGACUGAAUGCCU GCGAGCAUCGGCUCGAGAAUGAGACGAUCACGCACUUCGGAGUGACUGAAUG CCUGCGAGCAUCGCCACCAGAGUAGAGCAUCAGCCUACUCCGGACAACUGA AUGCCUCGAGCAUCGAGUAGAGCGAGGAGCAUCAGCCCUCGCGUGUGACU GAAUGCCUCGAGCAUCUCACACAAACUACUCAAUGUCCGAAAGGUGGCA AACACUCCAAAGCAGCCAAACGGAUCAACAUGGCAGCGGUGCUU

BPx4S

GGUUCUGCCAUGAGGGCCCUGAAGAAGGGCCCGAUCCGAGGAGCAGACGAU
AUGGCGUCGCUCCCGGCUGCACGGCCCUGAAGAAGGGCCGGGAGUGACCAGC
AGAGCAUAUGGGCUCGCUGGGCCACCACCGCCUGAAGAAGGGCGGCGGAC
AAGCAGCAGAGGAUAUGGCCUCGCUGCGAGUAGAGGGCCUGAAGAAGGGC
CCGUGUGACGAGCAGACCAUAUGGGGUCGCUCGUCACACAAACUACUAAA
UGUCCGAAAGGUGGCAAACACUCCAAAGCAGCCAAACGGAUCAAAACAUGGC
AGCGGUGCUUU

Table S2. Oligonucleotide sequences for real-time PCR

Sequence (5' → 3')		
GFP	FW primer	CGAGGACAGCGTGATCTTC
	RV primer	CCACGGTGGCGTTGCT
	Taqman probe	CCGACAAGATCATCC
RFP	FW primer	TCAAGGAGGCCGACAAAGAG
	RV primer	GTA CTTGGCCACAGCCATCTC
	Taqman probe	CCTACGTCGAGCAGCA

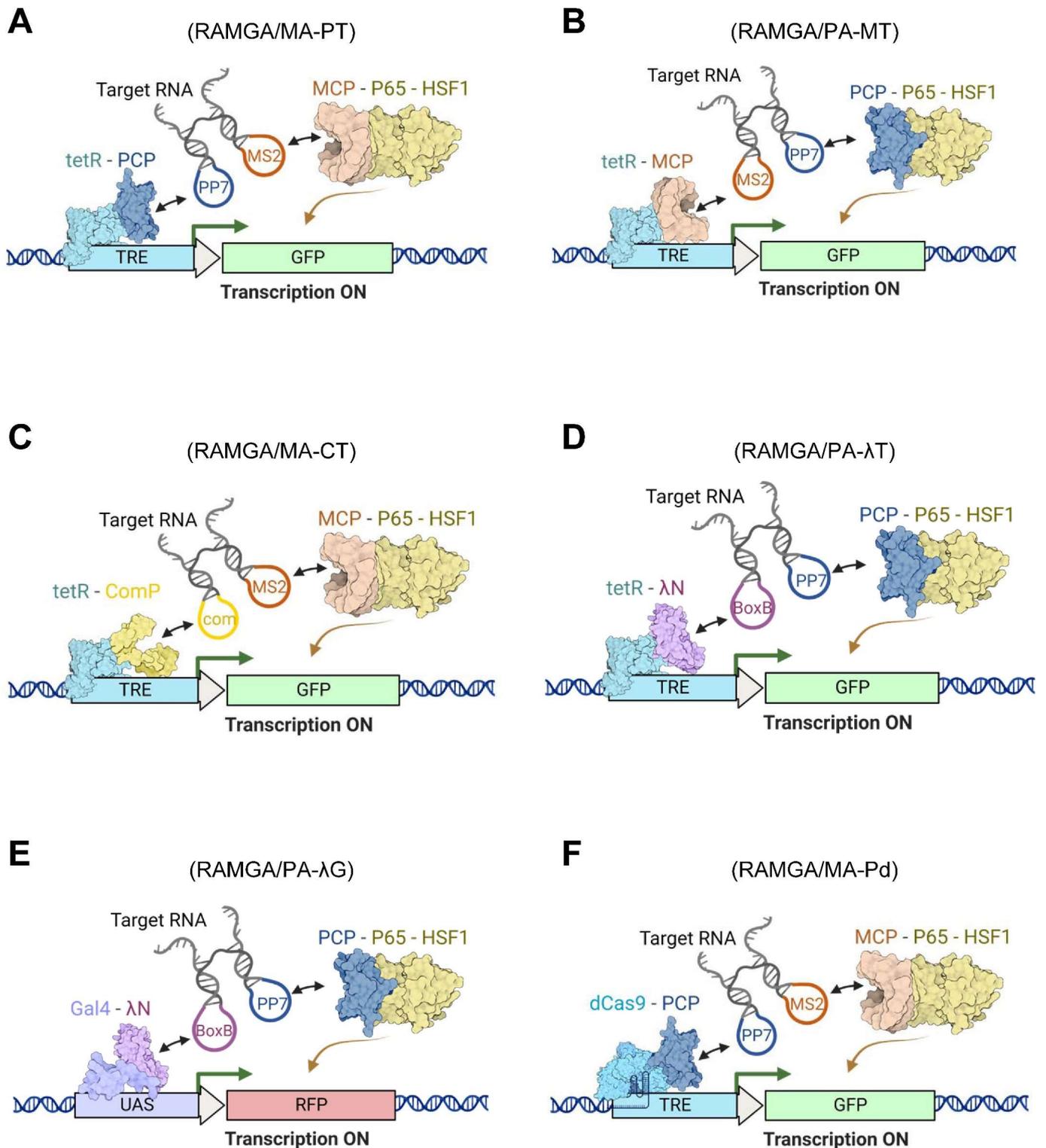


Figure S1. Schematic illustrations of transgene expression based on the RNA Aptamer-Mediated Gene Activation (RAMGA) system. (A) The RAMGA/MA-PT system consists of tetR-PCP and MCP-P65-HSF1, which can recognize and respond to RNA containing MS2-PP7 aptamers. (B) The RAMGA/PA-MT system consists of MCP-tetR and PCP-P65-HSF1, which can recognize and respond to RNA containing MS2-PP7 aptamers. (C) The RAMGA/MA-CT system consists of MCP-P65-HSF1 and COM-tetR, which can recognize and respond to RNA containing MS2-com aptamers. (D) The RAMGA/PA-λT system consists of PCP-P65-HSF1 and λN-tetR, which can recognize and respond to RNA containing PP7-BoxB aptamers. (E) The RAMGA/PA-λG system consists of PCP-P65-HSF1 and λN-Gal4, which can recognize and respond to RNA containing PP7-BoxB aptamers. (F) The RAMGA/MA-Pd system consists of MCP-P65-HSF1 and PCP-dCas9, which can recognize and respond to RNA containing MS2-PP7 aptamers.

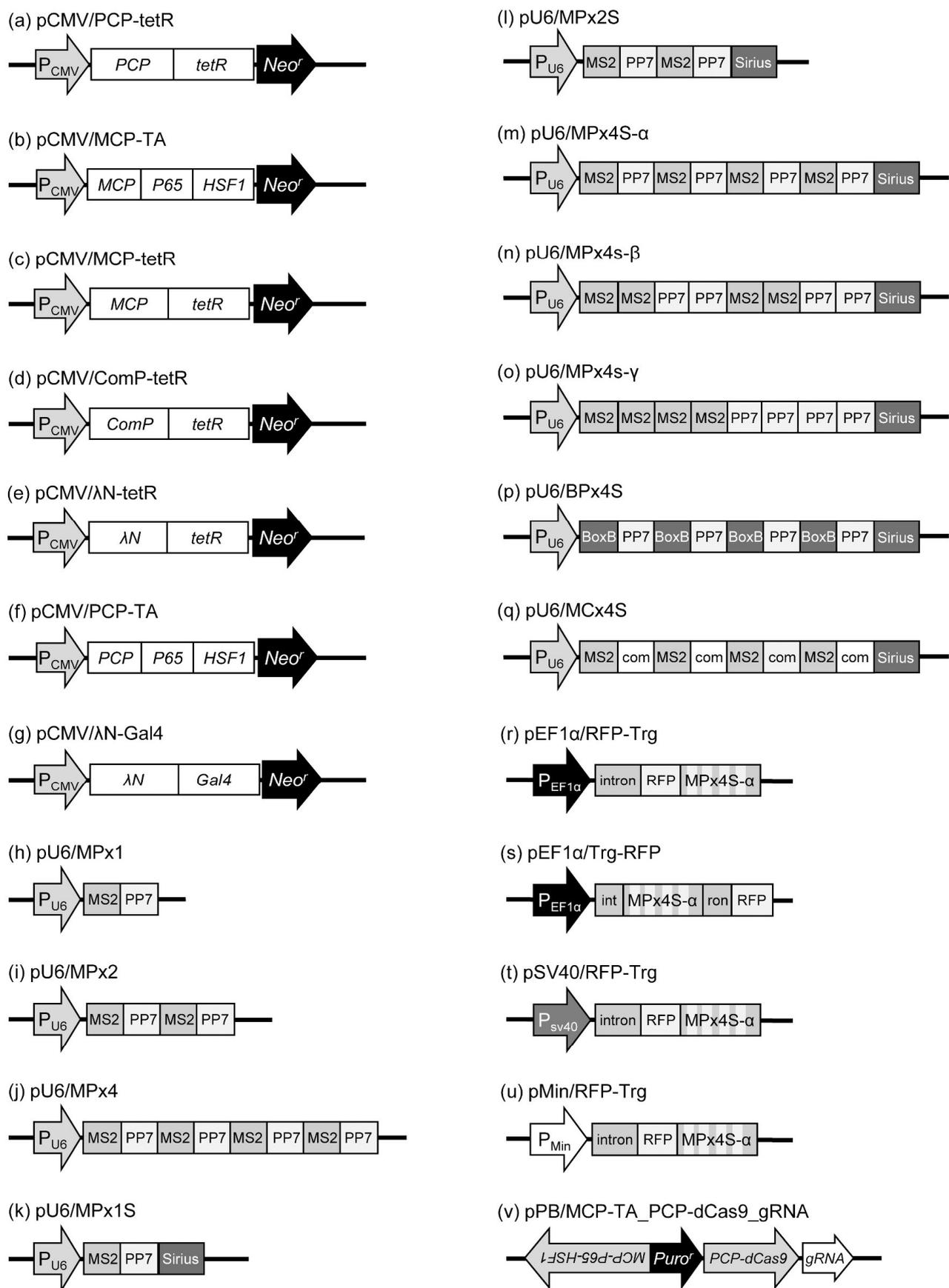
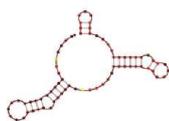


Figure S2. Schematic diagrams of plasmid structures. (a–g) Fusion protein expression vectors. (h–q) Trigger RNA expression vectors with a U6 promoter. (r–u) mRNA expression vectors with promoters EF1 α , SV40, and minimal promoter. (v) All-in-one vector for expressing MCP-P65-HSF1, PCP-dCas9, and gRNA.

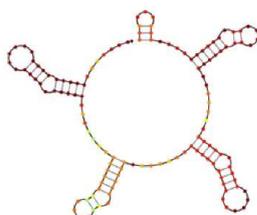
(a) MPx1

Free energy of secondary structure:
-26.20 kcal/mol



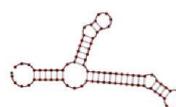
(b) MPx2

-49.10 kcal/mol



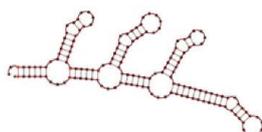
(c) MPx1S

-42.50 kcal/mol



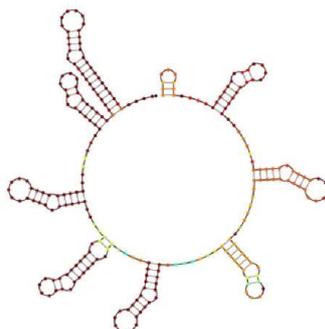
(d) MPx2S

-77.80 kcal/mol



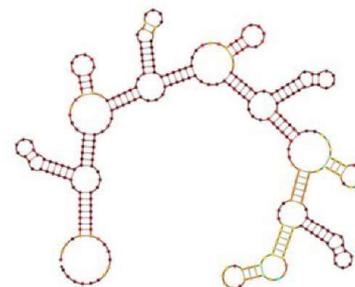
(e) MPx4

-112.10 kcal/mol



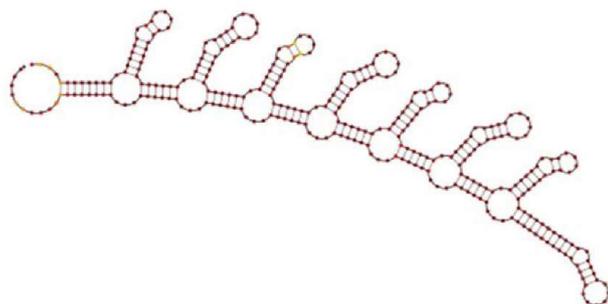
(f) MCx4S

-103.90 kcal/mol



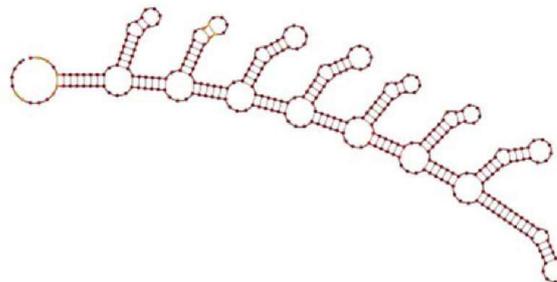
(g) MPx4S- α

-154.40 kcal/mol



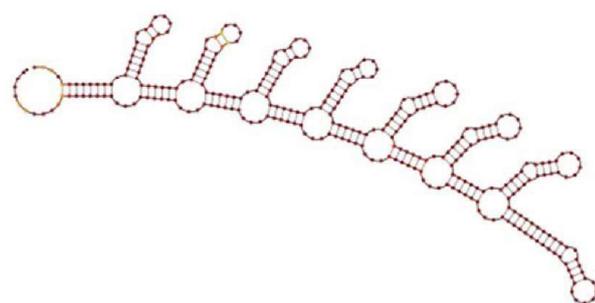
(h) MPx4S- β

-154.40 kcal/mol



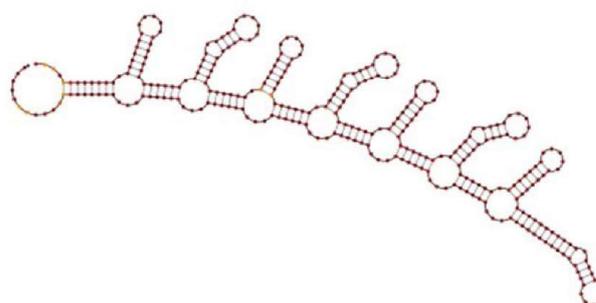
(i) MPx4S- γ

-154.40 kcal/mol



(j) BPx4S

-175.80 kcal/mol



Equilibrium probability at 37.0 °C



Figure S3. Predicted secondary structures of trigger RNAs along with their free energy calculations. The predictions were performed using NUPACK software.

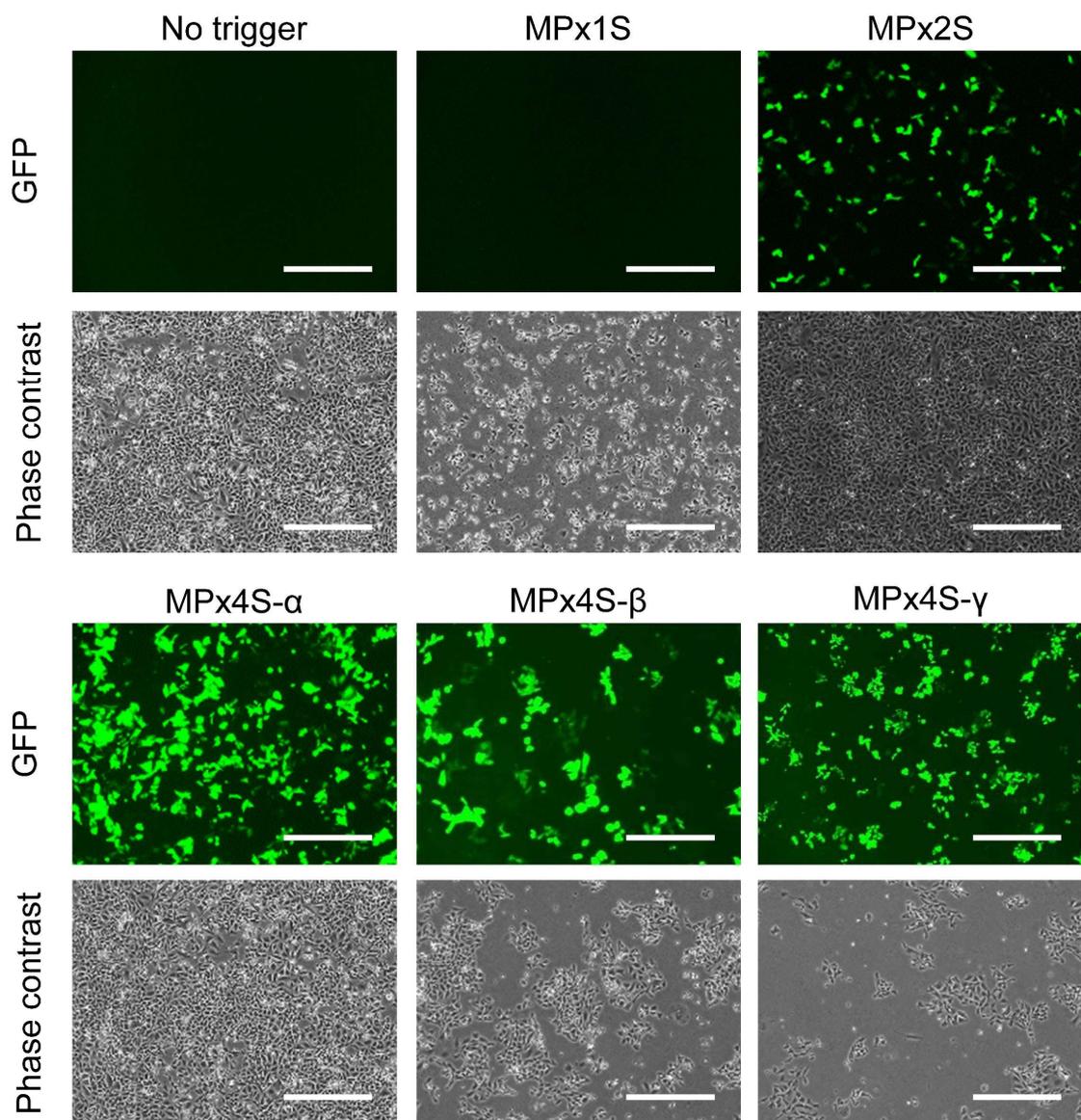


Figure S4. Phase-contrast and fluorescence images of CHO cells transfected with the RAMGA/MA-PT system, either with or without the presence of MS2-PP7 trigger RNA variants, which include MPx1S, MPx2S, MPx4S- α , MPx4S- β , and MPx4S- γ . GFP, Green Fluorescent Protein. Scale bars = 500 μ m.

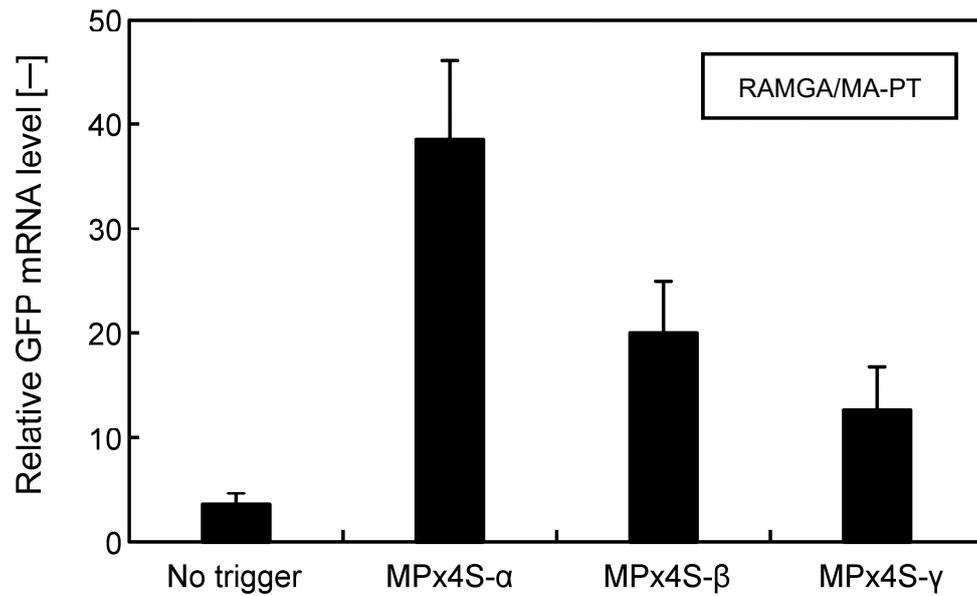


Figure S5. Relative GFP mRNA levels in CHO cells transfected with the RAMGA/MA-PT system in the presence or absence of MS2-PP7 trigger RNA variants containing MPx4S- α , MPx4S- β , and MPx4S- γ . Data represent the mean \pm SD.

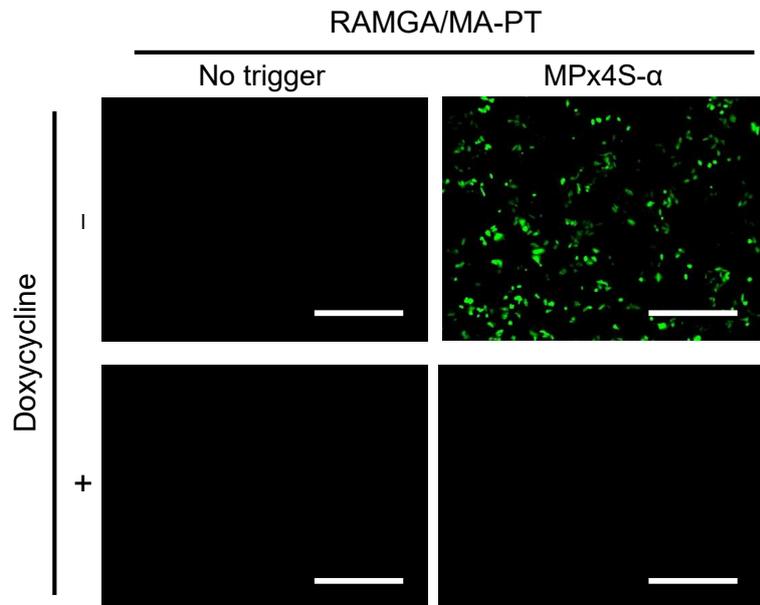


Figure S6. Phase-contrast and fluorescence images of CHO cells for post-transfection using RAMGA/MA-PT system with or without MPx4S- α as a trigger, in the absence (-) or presence (+) of doxycycline. Scale bars = 500 μ m.

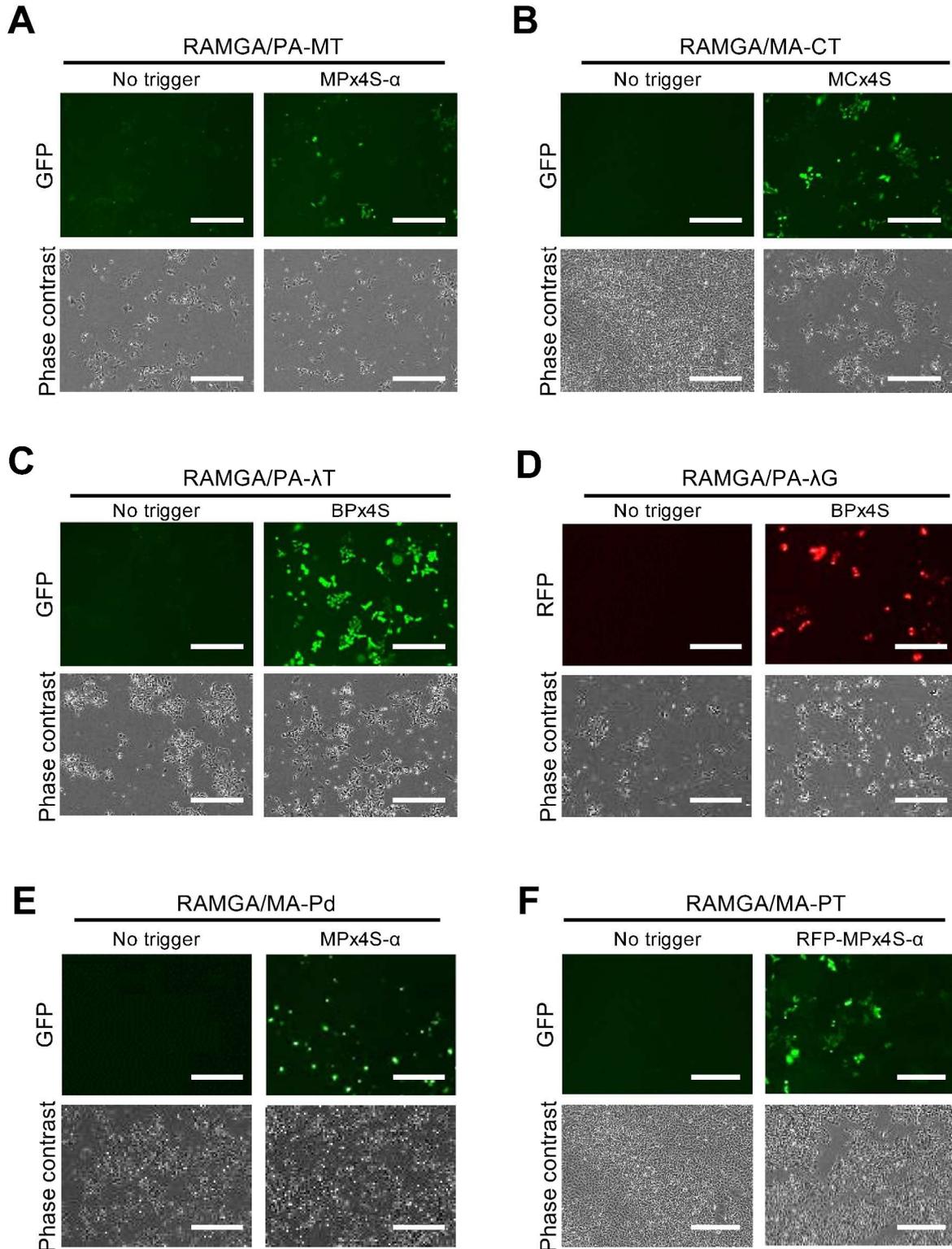


Figure S7. Phase-contrast and fluorescence images of CHO cells for post-transfection using various RAMGA systems. (A) Response of the RAMGA/PA-MT system to MPx4S- α . (B) Response of the RAMGA/MA-CT system to MCx4S. (C) Response of the RAMGA/PA- λ T system to BPx4S. (D) Response of the RAMGA/PA- λ G system to BPx4S. (E) Response of the RAMGA/MA-Pd system to MPx4S- α . (F) Response of the RAMGA/MA-PT system to RFP mRNA containing MPx4S- α . Scale bars = 500 μ m.