

Supplementary data

1. Supplementary tables:

Supplementary table S1. *Sulfolobus* strains used in this work

Strains	Genotype and features	Reference
$\Delta\beta$ E233	Derived from <i>S. islandicus</i> Rey15A, carrying deletion of the <i>pyrEF</i> genes and the type III-B Cmr- β locus including 7 <i>cmr</i> - β genes	(1)
$\Delta\beta$ E233S1	Carrying deletion of the <i>pyrEF</i> genes, the <i>lacS</i> gene and the type III-B Cmr- β locus including 7 <i>cmr</i> - β genes	(2)
$\Delta\alpha\Delta\beta$ E233S1	Carrying deletion of the <i>pyrEF</i> genes, the <i>lacS</i> gene, the Cmr- α locus including 6 <i>cmr</i> - α genes and the Cmr- β locus including 7 <i>cmr</i> - β genes	(2)
Δ array $\Delta\beta$ E233	Derived from $\Delta\beta$ E233, carrying deletion of two CRISPR loci and the type I-A module	This work
Δ array $\Delta\beta$ E233S1	Derived from $\Delta\beta$ E233S1, carrying deletion of two CRISPR loci and the type I-A module; equal to MF1	(3)
4 α -H16A	Derived from $\Delta\beta$ E233, carrying a single mutation (H16A) of <i>cmr4</i> α	This work
4 α -D83A	Derived from $\Delta\beta$ E233, carrying a single mutation (D83A) of <i>cmr4</i> α	This work
4 α -K46/50A	Derived from $\Delta\beta$ E233, carrying a double mutation (K46A, K50A) of <i>cmr4</i> α	This work
4 α -W197A	Derived from $\Delta\beta$ E233, carrying a single mutation (W197A) of <i>cmr4</i> α	This work
4 α -E199/Y201A	Derived from $\Delta\beta$ E233, carrying a double mutation (E199A, Y201A) of <i>cmr4</i> α	This work
4 α -K251A	Derived from $\Delta\beta$ E233, carrying a single mutation (K251A) of <i>cmr4</i> α	This work
4 α -4G-A	Derived from $\Delta\beta$ E233, carrying a quadruple mutation (G244A, G245A, G250A, G252A) of <i>cmr4</i> α	This work
HD ^m	Derived from $\Delta\beta$ E233, carrying a double mutation (H14A, D15A) of <i>cmr2</i> α	This work
Palm ^m	Derived from $\Delta\beta$ E233, carrying a double mutation (D667A, D668A) of <i>cmr2</i> α	This work
HD ^m Palm ^m	Derived from $\Delta\beta$ E233, carrying a quadruple mutation (H14A, D15A, D667A, D668A) donor DNA of <i>cmr2</i> α	This work
HD ^m Palm ^m 4 α ^{D27A}	Derived from $\Delta\beta$ -HD ^m Palm ^m , carrying a single mutation (D27A) of <i>cmr4</i> α	This work

Supplementary table S2. Plasmids used in this work

Plasmids	Genotype and features	Reference
pSeSD1	A <i>Sulfolobus-E. coli</i> shuttle vector with an expression cassette controlled under ParaS-SD promoter	(4)
pSe-Rp	A <i>Sulfolobus</i> artificial mini-CRISPR cloning vector	(5)
pAC-SS1	An artificial mini-CRISPR locus plasmid derived from pSe-Rp, carrying one spacer matching the protospacer 1 (SS1) of the <i>S. islandicus</i> lacS gene	(5)
pAC-MS1	Derived from pSeSD1, carrying an artificial CRISPR locus with 10 copies of 43 nt SS1 spacer.	(6)
pAC-cmr6α-10His	Derived from pAC-MS1, both carrying an artificial CRISPR locus with 10 copies of SS1 spacer and expressing His-tagged Cmr6α	(6)
pS10i	An invader plasmid carrying a target sequence of spacer 10 in CRISPR locus 2 in <i>S. islandicus</i>	(7)
pSi_1125	A self-targeting plasmid expressing crRNA targeting the <i>SiRe_1125 (alba)</i> gene	This work
pCmr4α	Derived from pSeSD1, expressing His-tagged Cmr4α	This work
pCmr4α-D27A	Derived from pCmr4α with a single mutation(D27A) of the <i>cmr4α</i> gene, expressing His-tagged Cmr4α-D27A	This work
pAC-SS1-Cmr4α	Derived from pAC-SS1, both carrying an artificial CRISPR locus with 1 copy of SS1 spacer and expressing His-tagged Cmr4α	This work
pAC-SS1-Cmr4α-D27A	Derived from pAC-SS1, both carrying an artificial CRISPR locus with 1 copy of SS1 spacer and expressing His-tagged Cmr4α-D27A	This work
pGE-4α-D27A	A genome editing plasmid for single mutation(D27A) of the <i>cmr4α</i> gene in <i>S.islandicus</i>	This work
pGE-4α-H16A	A genome editing plasmid for single mutation(H16A) of the <i>cmr4α</i> gene in <i>S.islandicus</i>	This work
pGE-4α-D83A	A genome editing plasmid for single mutation(D83A) of the <i>cmr4α</i> gene in <i>S.islandicus</i>	This work
pGE-4α-K46/50A	A genome editing plasmid for double mutation (K46A, K50A) of the <i>cmr4α</i> gene in <i>S.islandicus</i>	This work
pGE-4α-W197A	A genome editing plasmid for single mutation(W197A) of the <i>cmr4α</i> gene in <i>S.islandicus</i>	This work
pGE-4α-E199/Y201A	A genome editing plasmid for double mutation (E199A, Y201A) of the <i>cmr4α</i> gene in <i>S.islandicus</i>	This work
pGE-4α-K251A	A genome editing plasmid for single mutation(K251A) of the <i>cmr4α</i> gene in <i>S.islandicus</i>	This work
pGE-4α-4G-A	A genome editing plasmid for quadruple mutation (G244A, G245A, G250A, G252A) of the <i>cmr4α</i> gene in <i>S.islandicus</i>	This work
pGE-2α-HD-A	A genome editing plasmid for double mutation (H14A, D15A) of the <i>cmr2α</i> gene in <i>S.islandicus</i>	This work
pGE-2α-DD-A	A genome editing plasmid for double mutation (D667A, D668A) of the <i>cmr1α</i> gene in <i>S.islandicus</i>	This work

Supplementary table S3. Oligonucleotides used in this work

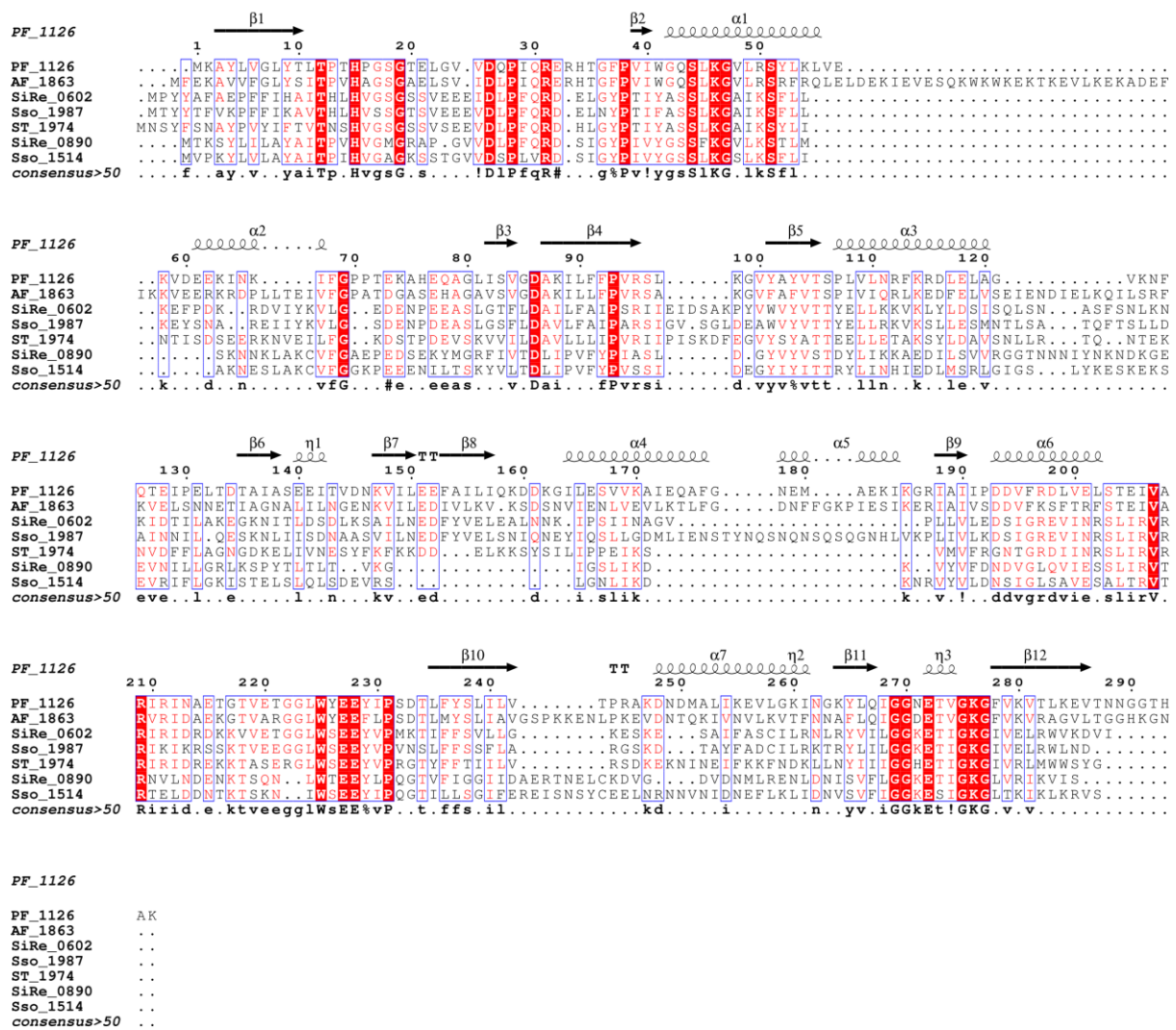
Oligonucleotide	Sequence (5'-3')
MRS-up	ATGCCCCGGGATGTAAACAAGTTAGG
MRS-dw	GGCACTCGAGAAAAAAGATTTTGCTTAATGGTG
Si_1125-up	AAAGAATGCAGCTAAGACATAGTTCATTACTGGTTTCTTTCCTA
Si_1125-dw	TAGCTAGGAAAGAAACCAGTAATGAAGTATGTCTTAGCTGCATT
Si_1581-up	AAAGATTATATAATTGGATAAGATTTTTCAAAGTATTGTTTTCA
Si_1581-dw	TAGCTGAAAACAATACTTTGAAAAATCTTATCCAATTATATAAT
Cmr4 α -fwd	GGGAATTCCATATGACCAAGAGTTATTTAATC
Cmr4 α -rev	GTCGTGCGACTGAAATCACCTTTATTCT
4 α -D27A-SpF	AAAG ACGTGGGAATGGGTAGAGCTCCAGGCGTAGTGGATTTACC
4 α -D27A-SpR	TAGC GGTAATCCACTACGCCTGGAGCTCTACCCATTCCCACGT
4 α -D27A-SOEF	GCATGTTGGTATGGGTAGAGCTCCAGGCGTAGTGGCTTTACCGTTC
4 α -D27A-SOER	GCCACTACGCCTGGAGCTCTACCCATACCAACATGCACTGGAGTTATTG
4 α -D27A-SalIF	ACGCGTCGACTCCAAAGTGTTGAATACC
4 α -D27A-NotIR	AAGGAAAAAAGCGGCCGCTTTAATACGTTCTCGTA
4 α -H16A-SpF	AAAG ACGTGGGAATGGGTAGAGCTCCAGGCGTAGTGGATTTACC
4 α -H16A-SpR	TAGC GGTAATCCACTACGCCTGGAGCTCTACCCATTCCCACGT
4 α -H16A-SOEF	CAATAACTCCAGTGGCGGTGGGAATGGGTAGAGCT
4 α -H16A-SOER	CCCATTCCCACCGCCACTGGAGTTATTGCGTAGGC
4 α -H16A-SalIF	ACGC GTCGAC TCCAAAGTGTTGAATACCAT
4 α -H16A-NotIR	AAGGAAAAA GCGGCCGC TTTAATACGTTCTCGTA
4 α -D83A-SpF	AAAG ATCTAGACTGGCAATAGGGTAAAAAACTGGTATGAGATCA
4 α -D83A-SpR	TAGC TGATCTCATACCAGTTTTTTACCCTATTGCCAGTCTAGAT
4 α -D83A-SOEF	CACTCATACCAGTTTTTTACCCTATTGCCAGCCTTGACGGTTACGTTTATGTTTC
4 α -D83A-SOER	ACCGTCAAGGCTGGCAATAGGGTAAAAAACTGGTATGAGTGCAGTAACTATAAAC
4 α -D83A-SalIF	ACGC GTCGAC GAAAGTTTTGTCAACTAAGG
4 α -D83A-NotIR	AAGGAAAAA GCGGCCGC AATTCGCTCATGAAATCACC
4 α -K46/50A-SpF	AAAG ATGGCTCTAGCTTTAAGGGAGTTTTAAAGTCAACATTAAT
4 α -K46/50A-SpR	TAGC ATTAATGTTGACTTTAAAACTCCCTTAAAGCTAGAGCCAT
4 α -K46/50A-SOEF	GTATGGCTCTAGCTTTGCAGGAGTTTTAGCATCAACATTAATGAGT
4 α -K46/50A-SOER	GATGCTAAAACCTCCTGCAAAGCTAGAGCCATACACTATTGGATAA
4 α -K46/50A-SalIF	ACGC GTCGAC GATAAGCAGGGAAGAAGGGT
4 α -K46/50A-NotIR	AAGGAAAAA GCGGCCGC GTCCACAAATTTTGAGACGT
4 α -W197A-SpF	AAAG AAATTTGTGGACTGAGGAGTATTTACCGCAAGGAACAGTA
4 α -W197A-SpR	TAGC TACTGTTCTTGCGGTAAATACTCCTCAGTCCACAAATTT
4 α -W197A-SOEF	AAACGTCACAAAATTTGGCGACTGAGGAGT
4 α -W197A-SOER	TCGCCAAATTTTGTGACGTTTTATTTTCAT
4 α -W197A-SalIF	ACGC GTCGAC AATAGGTTATCCAATAGTCT
4 α -W197A-NotIR	AAGGAAAAA GCGGCCGC GACCGAAATATTCTTGCCCTA
4 α -E199/Y201A-SpF	AAAG AAATTTGTGGACTGAGGAGTATTTACCGCAAGGAACAGTA

Oligonucleotide	Sequence (5'-3')
4α-E199/Y201A-SpR	TAGC TACTGTTTCCTTGCGGTAAATACTCCTCAGTCCACAAATTT
4α-E199/Y201A-SOEF	ATAAAAGACAAGGTTTATGTATTTGATAAC
4α-E199/Y201A-SOER	GTTTCCTTGCGGTAAAGCCTCTGCAGTCCAC
4α- E199/Y201A-SalIF	ACGC GTCGAC AATAGGTTATCCAATAGTCT
4α- E199/Y201A-NotIR	AAGGAAAAAA GCGGCCGC GACCGAAATATTCTTGCCTA
4α-K251A-SpF	AAAG TAGGAGGTAAGGAAACTATAGGTAAAGGACTTGTAAGAAT
4α-K251A-SpR	TAGC ATTCTTACAAGTCCTTTACCTATAGTTTCCTTACCTCCTA
4α-K251A-SOEF	GTAAGGAAACTATAGGTGCAGGACTTGTA
4α-K251A-SOER	CCTATAGTTTCCTTACCTCCTAGAAATACA
4α-K251A-SalIF	ACGC GTCGAC GTTTTTTACCCTATTGCCAG
4α-K251A-NotIR	AAGGAAAAAA GCGGCCGC ATTTTCTCACTCATATGGTA
4α-4G-A-SpF	AAAG TAGGAGGTAAGGAAACTATAGGTAAAGGACTTGTAAGAAT
4α-4G-A-SpR	TAGC ATTCTTACAAGTCCTTTACCTATAGTTTCCTTACCTCCTA
4α-4G-A-SOEF	CTAAGGAAACTATAGCTAAAGCACTTGTA
4α-4G-A-SOER	GCTATAGTTTCCTTAGCTGCTAGAAATACA
4α-4G-A-SalIF	ACGC GTCGAC GTTTTTTACCCTATTGCCAG
4α-4G-A-NotIR	AAGGAAAAAA GCGGCCGC ATTTTCTCACTCATATGGTA
2α-HD-A-SpF	AAAG CGACCCTCCTTGGAAGGCATGGGTAATTACAAGGAATATT
2α-HD-A-SpR	TAGC AATATTCCTTGTAATTACCCATGCCTTCCAAGGAGGGTCTG
2α-HD-A-SOEF	GCCTATTTTGCCGCCCTCCTTGGAAGGC
2α-HD-A-SOER	GGAGGGGCGGCAAAATAGGCTATTATTTTCTTATTTAAGAAC
2α-HD-A-SalIF	ACGC GTCGAC TCTGGCAAGATATTAGAC
2α-HD-A-NotIR	AAGGAAAAAA GCGGCCGC GTGGGGATAAGAAATAAATTAAGTGG
2α-DD-A-SpF	AAAG GTTAAACTTTATAGGCGATAACGCCAATATGTCGTCACCA
2α-DD-A-SpR	TAGC TGGTGACGACATATTGGCGTTATCGCCTATAAAGTTTAAC
2α-DD-A-SOEF	TTGGCGTTATCGCCTATAAAATTCAATGAAAAGTACGTAG
2α-DD-A-SOER	CTTTTCATTGAATTTTATAGGCGATAACGCCAAAATAGCAGCACCACCCAAAT ATATAG
2α-DD-A-SalIF	ACGC GTCGAC TCGGCCTGTAATATCAGTGAC
2α-DD-A-NotIR	AAGGAAAAAA GCGGCCGC GTCATATTCCCCGCTCTT

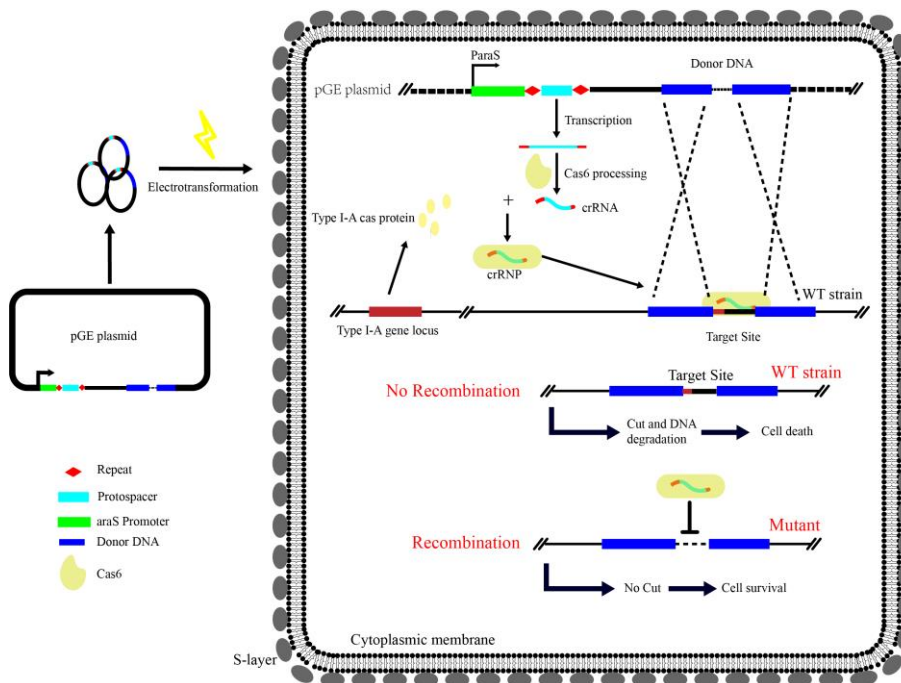
Supplementary table S4. Nucleic acid substrates used in this work

Name	Sequence (5'-3')	Size (mer)
RNA		
SS1-46	UGUUAAGUCUGGUUUCCCUCCAGGGUAUCUAAGCUUUGAAAAAAAAA	46
DNA		
S10-60	ACTATAGGGAGaATAGAATGCCCCCATTATACAATATCTACGTTTTAGATGAc ccccccc	60

2. Supplementary figures

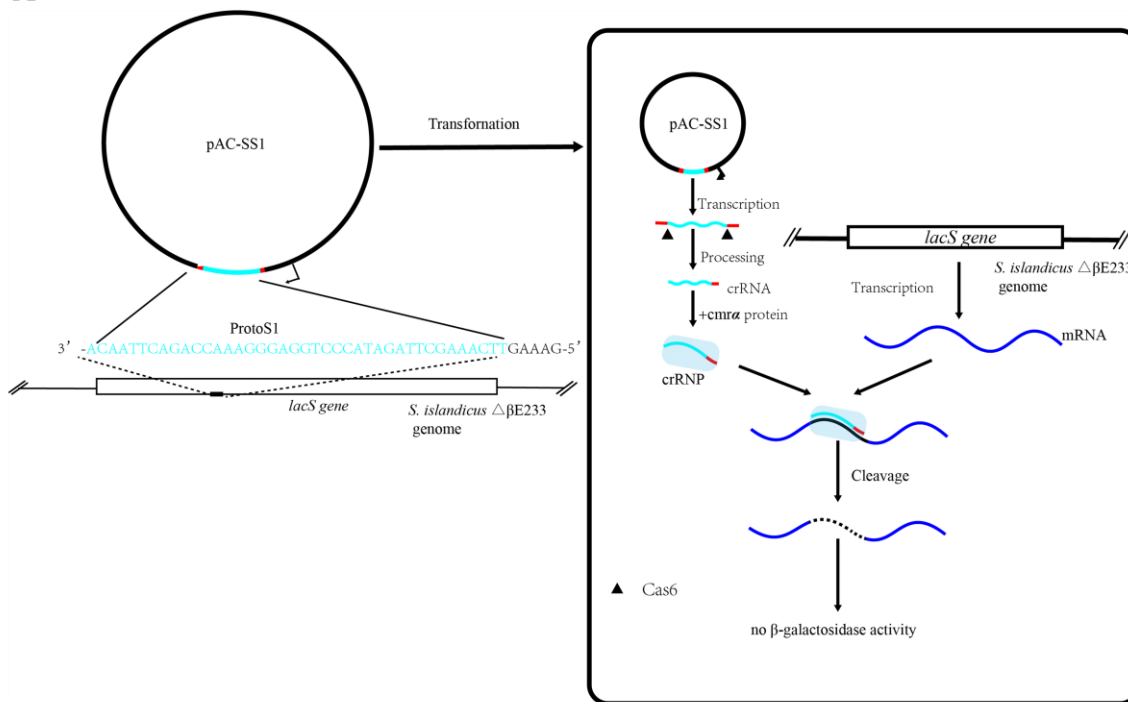


Supplementary figure S1. Conserved residues in *S. islandicus* Cmr4 α . Seven Cmr4 homologues (PF_1126, AF_1863, SiRe_0890, SSo1514, SSo_1987, SiRe_0602, ST_1974) were selected and aligned using ESPript 3.x (8). Thirteen highly conserved residues/motifs were chosen to produce eight Cmr4 α mutants as indicated below: Cmr4 α -H16A, Cmr4 α -D27A, Cmr4 α -D83A, Cmr4 α -K46/50A, Cmr4 α -W197A, Cmr4 α -E199/Y201A, Cmr4 α -K251A and Cmr4 α -4G-A.

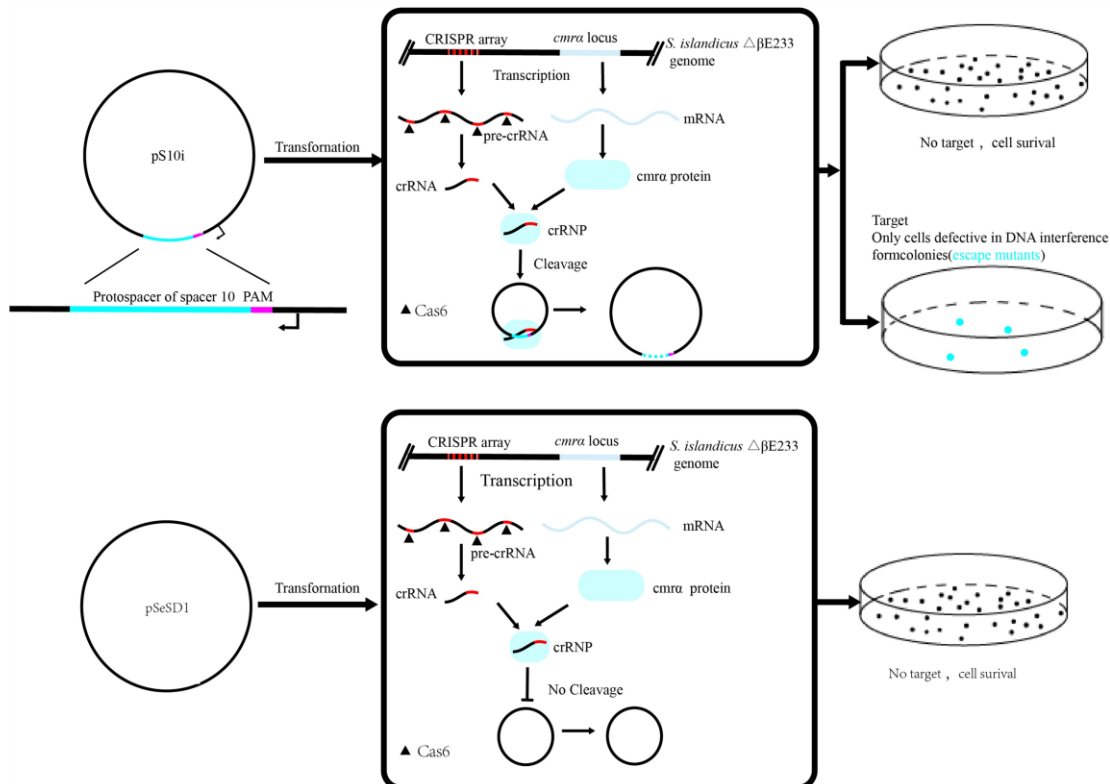


Supplementary figure S2. Mutagenesis was conducted by introduction of pGE plasmids into $\Delta\text{Cmr-}\beta$ cells by electroporation. The rationale is that crRNAs expressed from the CRISPR array in pGE plasmids guide the endogenous I-A immunity to specifically target individual fragment of the wild-type *cmr4 α* gene for degradation, and in the meantime, the plasmid-borne homologous arms carrying the mutated *cmr4 α* gene are allowed to recombine with the wild-type gene, giving the corresponding mutated *cmr4 α* gene in the host chromosome.

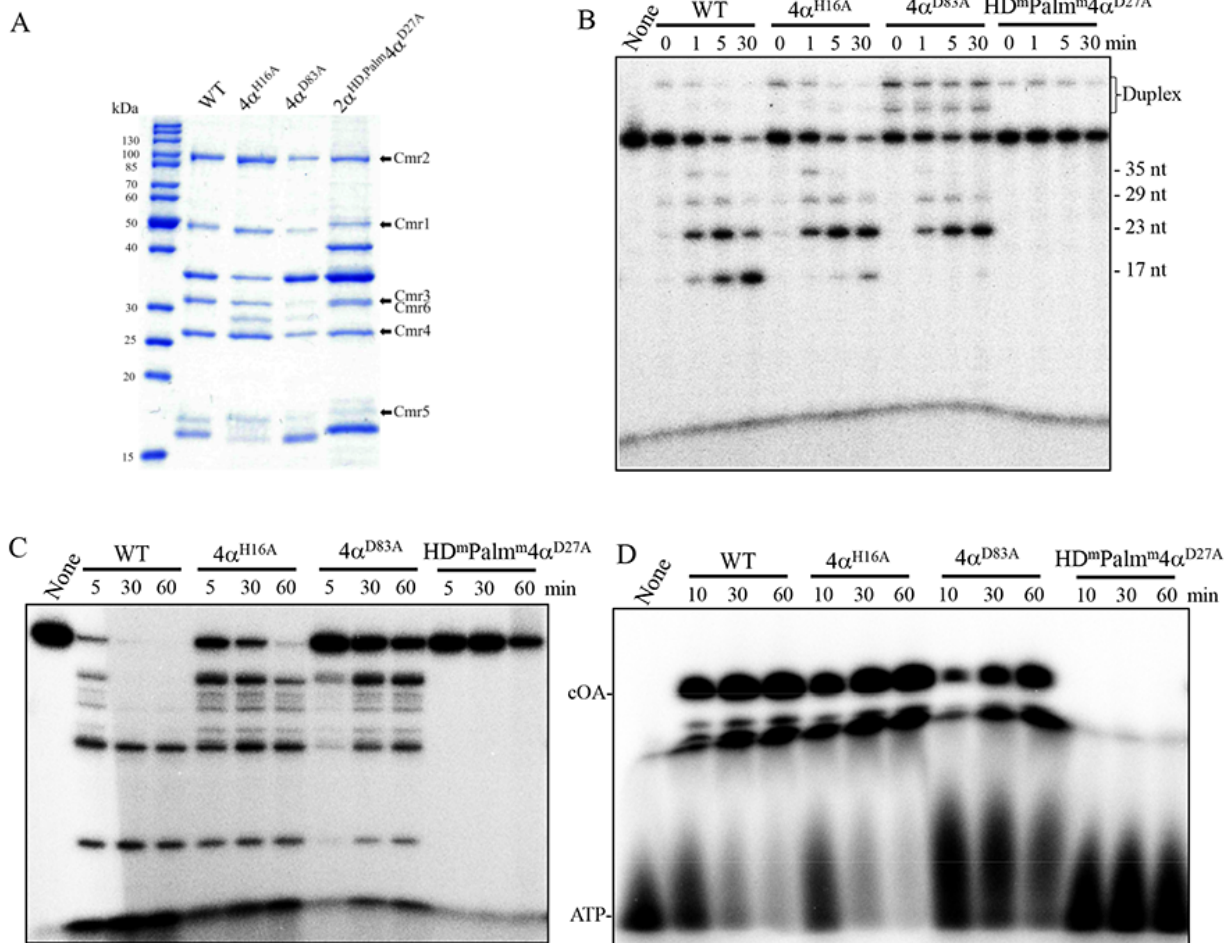
A



B



Supplementary figure S3. Two genetic assays developed for *S. islandicus*: the mini-CRISPR and reporter gene-based RNA interference assay (A) and the interference plasmid assay (B)



Supplementary figure S4. In vitro activities of wild type (WT) Cmr- α , Cmr- α _4 α^{H16A} , Cmr- α _4 α^{D83A} and Cmr- α _HD^m,Palm^m4 α^{D27A} . (A) SDS-PAGE analysis, (B) target RNA cleavage, (C) DNA cleavage and (D) cOA synthesis activity of the four complexes. None: Substrate only, no Cmr complex was added. Cleavage assay was conducted for the time periods indicated in the figures; Duplex: duplex of crRNA and substrate; cOA: cyclic tetra-adenylates.

References

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