

IVA cloning: A single-tube universal cloning system exploiting bacterial *In Vivo* Assembly

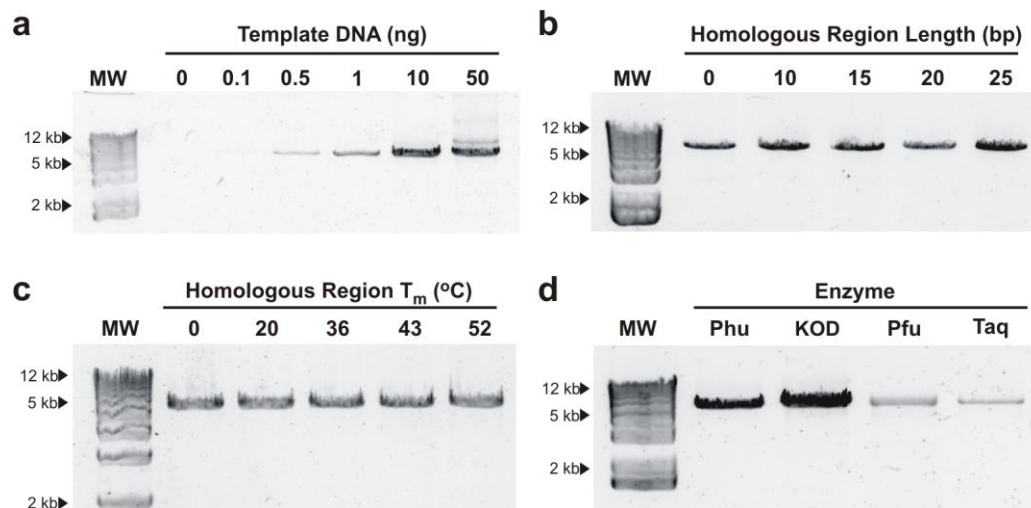
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SUPPLEMENTARY MATERIAL



Supplementary Figure S1. Method and primer optimisation. Agarose gels showing PCR amplifications with increasing amount of template DNA (**A**). Using different homologous region lengths (**B**) and T_m (**C**), equivalent amplifications are seen in all cases, attributing the differences in colony formation exclusively to recombination efficiency. (**D**) Agarose gel showing difference in amplification efficiency for four polymerases performing the same deletion. (MW = 1 kb Plus Ladder).

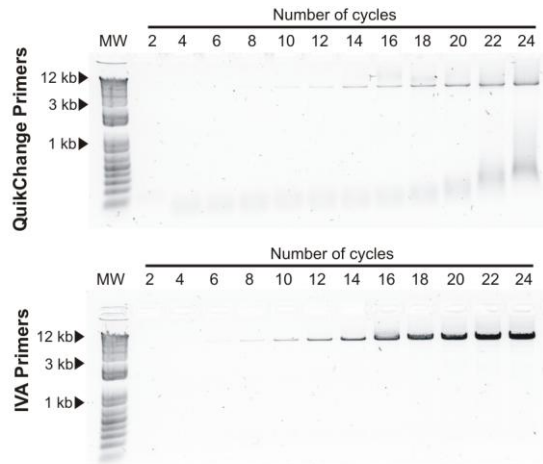
a QuikChange primer design



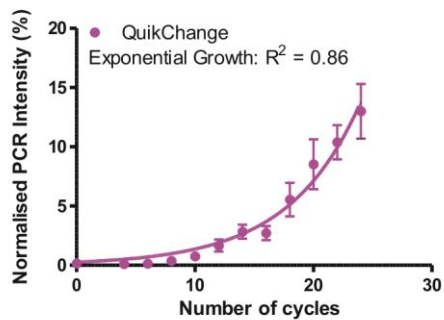
IVA primer design



b Mutagenesis



c



Supplementary Figure S2. Insertion, deletion and mutagenesis amplifications. (A) A comparison of primer design for mutagenesis using QuikChange™ (upper - blue) and IVA (lower - green). **(B)** Comparison of IVA and QuikChange™ amplification rates by examining band intensity every two PCR cycles. Higher intensity and no low molecular weight smearing are seen using IVA primers. (MW = 1 kb Plus Ladder). **(C)** When plotting intensity against cycle number, an exponential curve can be fitted ($R^2 = 0.86$) when using QuikChange™ mutagenesis primers (normalised to IVA cloning).

Supplementary Table S1. List of primers. Homologous regions are shown in italics and mutation codons in bold.

Optimisation				Function	
Template opt.					
OPT1-F	AATGACAGCTCATCCTCAGAGAACCGG			Amplify pRK5 without recombination	
OPT1-R	TCCTCCGTGAGAATGACCCAAAAGCC				
Length opt.			T _m (°C)	Length (bp)	
OPT2-F	<i>CGCCCGGCGG</i> AATGACAGCTCATCCTCAGAGAACCGG		40	10	Delete GluA3 NTD region in pRK5
OPT2-R	<i>CCGCCGGGCG</i> TCCTCCGTGAGAATGACCCAAAAGCC		40	10	
OPT3-F	<i>CACGTCCAGAAGAAT</i> AATGACAGCTCATCCTCAGAGAACCGG		40	15	
OPT3-R	<i>ATTCTTCTGGACGTG</i> TCCTCCGTGAGAATGACCCAAAAGCC		40	15	
OPT4-F	<i>ATTCTTATGGACATTAATTA</i> AATGACAGCTCATCCTCAGAGAACCGG		40	20	
OPT4-R	<i>TAATTAATGTCCATAAGAAT</i> TCCTCCGTGAGAATGACCCAAAAGCC		40	20	
OPT5-F	<i>ATTATAATATTTACTATATATTATT</i> AATGACAGCTCATCCTCAGAGAACCGG		40	25	
OPT5-R	<i>AATAATATATAGTAAATATTATAAT</i> TCCTCCGTGAGAATGACCCAAAAGCC		40	25	
T_m opt.			GC%	T _m (°C)	
OPT6-F	<i>ATTATTAATTATTTA</i> AATGACAGCTCATCCTCAGAGAACCGG		0	20	Delete GluA3 NTD region in pRK5
OPT6-R	<i>TAAATAATTAATAAT</i> TCCTCCGTGAGAATGACCCAAAAGCC		0	20	
OPT7-F	<i>GTCATCAGTTCTTTC</i> AATGACAGCTCATCCTCAGAGAACCGG		40	36	
OPT7-R	<i>GAAAGAAGTATGAC</i> TCCTCCGTGAGAATGACCCAAAAGCC		40	36	
OPT8-F	<i>GACGTCAGCGTGGTA</i> AATGACAGCTCATCCTCAGAGAACCGG		60	45	
OPT8-R	<i>TACCACGCTGACGTC</i> TCCTCCGTGAGAATGACCCAAAAGCC		60	45	
OPT9-F	<i>GGCGTCAGCGCGGTC</i> AATGACAGCTCATCCTCAGAGAACCGG		80	53	
OPT9-R	<i>GACCGCGTGACGCC</i> TCCTCCGTGAGAATGACCCAAAAGCC		80	53	
Delete/Insert					
INS1-F	<i>GAACAAAACTCATCTCAGAAGAGGATCTG</i> TTCCCAACACCATCAGCATAGGTGG				Inserting myc tag at GluA3 N-terminus in pRK5-GluA3 (IVA)
INS1-R	<i>TCTTCTGAGATGAGTTTTGTTC</i> TCCTCCGTGAGAATGACCCAAAAGCC				
INS2-F	P -TCAGAAGAGGATCTG TTCCCAACACCATCAGCATAGGTGG				Inserting myc tag at GluA3 N-terminus (phosphorylated primers)
INS2-R	P -GATGAGTTTTGTTC TCCTCCGTGAGAATGACCCAAAAGCC				
DEL1-F	<i>TGGGTCATTCTACGGAGGA</i> AATGACAGCTCATCCTCAGAGAACCGG				Deleting GluA3 N-terminal domain in pRK5-GluA3
DEL1-R	<i>TCCTCCGTGAGAATGACCCAAAAGCC</i>				
DEL2-F	<i>CTGATTTTTGGTGTC</i> TCTTCTAACAGCATAAGATAGGGGGG				Deleting myc tag at GluA2 N-terminus in pIRES-GluA2
DEL2-R	<i>GACACCAAAAATCAGTCCCATAAAACAG</i>				
DELINS1-F	<i>GAAAACCTGTACTTCCAGTCC</i> ATG GTG AGC AAG GGC GAG GAG CTG				Replacing IRES with linker in pIRES-GluA2 EGFP
DELINS1-R	<i>GGACTGGAAGTACAGGTTTTT</i> CTCATTTCGTTTCGCTCGGCCCTTG				

Mutagenesis**IVA cloning**

MUT1-F	CCAGATCGTGAAGCTA TGC AAGAATGGCATCGGGTACCACTACATCC	GluA1 E202C mutation in pIRES-
MUT1-R	GCATAGCTT CACGATCTGGCCAGGATG	GluA1
MUT2-F	ACTGAAGCATTCCGT TCCCTT CGGAAGCAGAGGATTGAAATATCCCG	GluA2 N292S mutation in pIRES-
MUT2-R	GGAACGGAA TGCTTCAGTCATCACTTGGACAG	GluA2
MUT3-F	ACAAATTGTGAGTGTT TGC AAGCATGTCAAAGGCTACCATTATATCATC	GluA4 G208C mutation in pRK5-
MUT3-R	GCAAACTC CACAATTTGTTCTAAAATGTTTTGAAGC	GluA4
MUT4-F	ACCGACTACCTCCAG TAGT CCGCCATCACCCGCATCCCC	y2 A219STOP mutation in pGW1-y2
MUT4-R	CTACTGGAGG TAGTCGGTGGCACGG	

QuikChange mutagenesis

MUT5-F	CCAGATCGTGAAGCTAT GCA AAGAATGGCATCG	GluA1 E202C mutation in pIRES-
MUT5-R	CGATGCCATTCTT GCAT AGCTTCACGATCTGG	GluA1
MUT6-R	GACTGAAGCATTCCGTT TCCCTT CGGAAGCAGAGG	GluA2 N292S mutation in pIRES-
MUT6-F	CCTCTGCTTCCGAAG GGAAC GGAATGCTTCAGTC	GluA2
MUT7-R	GAACAAATTGTGAGTGTT TGCA AAGCATGTCAAAGGCTAC	GluA4 G208C mutation in pRK5-
MUT7-F	GTAGCCTTTGACATGCTT GCAAACTC CACAATTTGTTT	GluA4
MUT8-F	CCGACTACCTCCAG TAGT CCGCCATCACCCG	y2 A219STOP mutation in pGW1-y2
MUT8-R	CGGGTGATGGCG ACTACT GGAGGTAGTCGG	

Subcloning

SUB1-F	ACCGTCAGATCCGCTAGC ATGAAGACGAGCCGCCGCGGC	Amplifies GSG1L for pIRES vector
SUB1-R	GATCTGAGTCCGGTAGC TCACACCCAGTGCCCCAGGACCC	
SUB2-F	GCTACCGGACTCAGATCTCGAGC	Amplifies pIRES vector for GSG1L
SUB2-R	GCTAGCGGATCTGACGGTTCACTAAAC	
SUB3-F	TGGTACCGAGCTCGGATCC ATGCAAAAGATTATGCATATTTCTGTCTCCTTTCTC	Amplifies GluA2 for pcDNA4/TO
SUB3-R	GTGCTGGATATCTGCAGAATTC CTAATTTTAACTCTCGATGCCATATACGTTGTAAC	
SUB4-F	GAATTCTGCAGATATCCAGCACAGTGGC	Amplifies pcDNA4/TO vector for GluA2
SUB4-R	GGATCCGAGCTCGGTACCAAGCTTAAG	
SUB5-F	ATCGATAAGCTTGATTGAGCTAGCC ACCATGGTGAGCAAGGGCG	Amplifies EGFP-Homer1c for AAV-CW3SL vector
SUB5-R	CTCGAGATAATCAACCTCTGGATTA TTAGCTGCATTCTAGTAGCTTGGCCAAATTATCC	

Multi-site

INS3-F	GACTACAAGGACGACGATGACAAG TCTTCTAACAGCATAACAGATAGGGGGC	Inserts FLAG at GluA2 N-terminus in pCustom vector
INS3-R	TCATCGTCGTCCTTGTAGTC GACACCAAAAATCAGTCCCATAAAACAGGAGA	
DEL3-F	TTCCAGAATTTTGCAACTTAT AAGGAAGGTTACAACGTATATGGCATCGAGAG	Deletes FLAG at GluA2 C-terminus in pCustom vector
DEL3-R	ATAAGTTGCAAAATTCTGGGAATTCTGCGAGGAAG	

DELINS2-F	AAGGATGACGACGATAAG AATGACAGCTCATCTCAGAGAACCGG	Exchanges GluA3 NTD for FLAG at in pRK5-GluA3
DELINS2-R	ATCGTCGTCATCCTTGAATCTCCGTGAGA	
SUB3-F	AGATCGGAAGCGGAAGCGGC GGGCTGTTTGATCGAGGTGTTCAAATGC	Amplifies γ 2 for GluA3- γ 2 tandem construct
SUB3-R	CTTCTGGTGGGAAGGGATCC TCATACGGGCGTGGTCCGGCGG	
SUB4-F	GGATCCCTTCCCACCAGAAGCATG	Amplifies pRK5-GluA3 vector for GluA3- γ 2 tandem construct
SUB4-R	GCCGCTTCCGCTTCC GATCTTAACACTTTCTGTTCCATACACGTTGTAG	
XhoI sites		
MUT9-F	ATGGGGCAAAGCGTGCTC GAG GCGGTCTTCTTTTTAGTCTGGGGC	XhoI site 1 in pRK5-GluA3
MUT9-R	CTC GAGCACGCTTTGCCCATTTTCTTCTG	
MUT10-F	ACTGGAAAGAGTCATGCATG CTC GAGCCAACATTACAGTTTTCCAGATTGTCAAC	XhoI site 2 in pRK5-GluA3
MUT10-R	GAG CATGCATGACTCTTTCCAGTAAAATGTCAGTAAAAC	
MUT11-F	GCAAGGATGTGATATTTCTC GAG GATCACTTTCTGGGCGCATTGTTGGAG	XhoI site 3 in pRK5-GluA3
MUT11-R	CTC GAGAAATATCACATCCTTGCTGCATGAAAGCA	
MUT12-R	CCTGATGCGGTATTTTCTC GAG ACGCATCTGTGCGGTATTTACACCG	XhoI site 4 in pRK5-GluA3
MUT12-F	CTC GAGAAAATACCGCATCAGGCGCCATTC	
MUT13-R	ACAAGCTGTGACCGT CTC GAGCTGCATGTGTCAGAGGTTTTACCC	XhoI site 5 in pRK5-GluA3
MUT13-F	GAG ACGGTCACAGCTTGCTGTAAGC	
Assembly		
ASS1-F	GCAGTGAGCGCAACGCAA TGCTTAGGGTTAGGCGTTTTGCGC	Amplifies CMVtet for Assembly
ASS1-R	CTATGGAGGTCAAACAGCG TCTCTATCACTGATAGGGAGATCTCTATCAC	
ASS2-F	CGCTGTTTTGACCTCCATAG ATGGTGAGCAAGGGCGAGGAGC	Amplifies EGFP for Assembly
ASS2-R	GATGGTGTGGGGAATCC CTTGTACAGCTCGTCCATGCCGAG	
ASS3-F	GGATTCCCAACACCATCAGCATAGG	Amplifies GluA3 for Assembly
ASS3-R	GCCGCTTCCGCTTCC GATCTTAACACTTTCTGTTCCATACACGTTGTAG	
ASS4-F	CGAAGCTTGAGCTCGAG TCATACGGGCGTGGTCCGGC	Amplifies γ 2 for Assembly
ASS4-R	AGATCGGAAGCGGAAGCGGC GGGCTGTTTGATCGAGGTGTTCAAATGC	
ASS5-F	TGGCCGCCATGGCCCAACTTG	Amplifies pRK5-GluA4 vector for Assembly
ASS5-R	ATTGCGTTGCGCTCACTGCCCG	
Library		
LIB1-F	TGGCCGCCATGGCCCAACTTG	Amplifies pRK5-GluA3 vector for Library
LIB1-R	ATTGCGTTGCGCTCACTGCCCG	
LIB2-F	GCAGTGAGCGCAACGCAAT GCTCGCCCGACATTGATTATTGACTAG	Amplifies CMV promoter for Library
LIB2-R	CTATGGAGGTCAAACAGCG AGCTCTGCTTATATAGACCTCCCACCG	

LIB3-F	GCAGTGAGCGCAACGCAAT CACTTGTGGACTAAGTTTGTTCGCATCC	Amplifies CamKII promoter for Library
LIB3-F	CTATGGAGGTCAAAACAGCG GCTGCCCCCAGAACTAGGGG	
LIB4-F	CGCTGTTTGACCTCCATAG CGAATTCGAATATGCCGTACATCTTTGCC	Amplifies GluA1 for Library from pIRES-GluA1
LIB4-F	GGGCCATGGCGGCCA TTACAATCCTGTGGCTCCAAGGGC	
LIB5-F	CGCTGTTTGACCTCCATAG GCTAGCGGATTCTTCTGCCTTCACTTC	Amplifies GluA2 for Library from pIRES-GluA2
LIB5-F	GGGCCATGGCGGCCA CTCGAGGCACTCAGAAGGTTCTATC	
LIB6-F	CGCTGTTTGACCTCCATAG GAATTCGGCACGAGGTTGCGCC	Amplifies GluA3 for Library from pIRES-GluA3
LIB6-F	GGGCCATGGCGGCCA GGATCCCTAGATCTTAACACTTTCTGTTCC	