

NEBaseChanger Summary - Mon May 15 2023

Input sequence:BRho-GFP11-1D4

Type of mutagenesis: substitution

Mutagenesis region: 389 to 389

Replace/insert: A

Result

N Y F L L L W T T R L N R Q
\* L F S S A M D N S S \* P S K
V I I F F C Y G Q L V L T V K
GTAATTATTTCTTCTGCTATGGACA**ACTCGTCTTaACCGTCAAAG**
**CATTAATAAAAGAAGACGATACCTG**TTGAGCAGAATTGGCAGTTTC

Required Primers

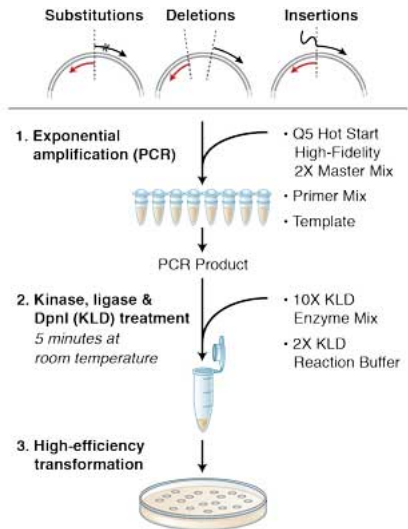
Name (F/R)	Oligo (Uppercase = target-specific primer)	Len	% GC	Tm	Ta *
Q5SDM_5/16/2023_F	A <b>ACTCGTCTTa</b> ACCGTCAAAG	21	43	60°C	61°C
Q5SDM_5/16/2023_R	GTCCATAGCAGAAGAAATAATTAC	25	32	60°C	

\* Ta (recommended annealing temperature)

PROTOCOL

The full Q5 Site-Directed Mutagenesis Protocol can be found in the manual, which can be downloaded from: [www.NEB.com/CN/E0554](http://www.NEB.com/CN/E0554)

For your convenience, a quick protocol is presented here:



Step I: Exponential Amplification (PCR)

	25 µl RXN	FINAL CONC.
Q5 Hot Start High-Fidelity 2X Master Mix	12.5 µl	1X
10 µM Forward Primer	1.25 µl	0.5 µM
10 µM Reverse Primer	1.25 µl	0.5 µM
Template DNA (1-25 ng/ µl)	1 µl	1-25 ng
Nuclease-free water	9.0 µl	

Cycling Conditions:

STEP	TEMP	TIME
Initial Denaturation	98 °C	30 seconds
	98 °C	10 sec
25 Cycles	61 °C*	10-30 sec
	72 °C	20-30 seconds/kb
Final Extension	72 °C	2 minutes
Hold	4-10 °C	∞

\*The recommended annealing temperature of 61 °C is specific for the mutagenic primers listed above when amplified with Q5 DNA Polymerase.

Step II: Kinase, Ligase & DpnI (KLD) Treatment

10 µl RXN	FINAL CONC.
-----------	-------------

PCR Product	1 µl	
2X KLD Reaction Buffer	5 µl	1X
10X KLD Enzyme Mix	1 µl	1X
Nuclease-free water	3 µl	

Mix well by pipetting gently up and down, and incubate at room temperature for 5 minutes.

Step III: Transformation

- 1
- Add 5 µl of the KLD reaction mixture directly from step II to 50 µl of chemically-competent cells.
- 2
- Incubate on ice for 30 minutes.
- 3
- Heat shock at 42 °C for 30 seconds.
- 4
- Incubate on ice for 5 minutes.
- 5
- Add 950 µl of SOC, gently shake at 37 °C for 1 hour.
- 6
- Spread 40-100 µl onto appropriate selection plate and incubate overnight at 37 °C (if necessary, make a 10-100 fold dilution of recovered cells before plating to avoid a lawn of colonies).

Sequence after Q5 SDM

TTACAGCTTATTTTGTCTTTGGGCCGACAGGGTGCAACCTCGAAGGGTTCTTCGCAACACTT  
GGAGGAGAGATCGCACTCTGGTCTCTTGTAGTACTTGCTATAGAGAGGTATGTGGTAGTGTG  
TAAGCCAATGTCAAACCTCAGGTTTGGTGAGAATCACGCAATTATGGGACTTGCCCTGACTT  
GGATAATGGCGATGGCGTGC CGCGGTGCACCTCTTGTGGGATGGTCAAGGTACATTCGGGAG  
GGGATGCAATGCAGTTGTGGAATCGATTATTACTTCTAGGCAAGAGGTAAACAACGAGTC  
TTTTGTGCATATATATGTTCTGTGTGCATTTTACGATCCCACTCGTAATTATTTTCTTCTGCT  
ATGGACAACCTCGTCTTAACCGTCAAAGAGGCGCGCGCGCAGCAGCAAGAGTCAGCGACAAC  
CAAAAAGCCGAGAAAGTAAGTAAGTTCGGATGGTAATCATAATGGTGGTCGCTTTCCTGATCTG  
CTGGGTCCCGTATGCCAGCGTCGCCTTTTATATTTTTACACATCAGGGGTCTGATTTTGGCC  
CGATCTTCATGACTATACCAAGTTTCTTTGCAAAAAGTCTTCCATTTACAACCCGTGAATT  
TACATAATGATGAACAACAGTTTCGCAATTGTATGCTGACAACTCTCTGCTCGGACGGAA  
CCCCCTTGGCGATGACGAAGCTAGTACAACCGCTAGCAAAACAGAGACATCCCAGGTGCGCAC  
CCGCAATAAGCGCGCGGACTACAAGGATGACGATGACAAGGATTACAAGACGACGATGATA  
AGGACTATAAGGATGATGACGACAATAATAGCAATTCTCGACGACTGCATAGGGTTACCC  
CCCTCTCCCTCCCCCCCCCTAACGTTACTGGCCGAAGCCGCTTGAATAAGGCCGGTGTG  
CGTTTGTCTATATGTTATTTTCCACCATATTGCCGTCTTTTGGCAATGTGAGGGCCCGGAAA  
CTGGCCCTGTCTTCTTGACGAGCATTCCTAGGGGTCTTTCCCTCTCGCCAAAGGAATGCA  
AGGTCTGTTGAATGTCGTGAAGGAAGCAGTTCTCTGGAAGCTTCTTGAAGACAAACAACGT  
CTGTAGGACCCCTTTGCAAGCAGCGGAACCCCCACCTGGCGACAGGTGCCTCTGCGGCCAA  
AAGCCAGCTGTATAAGATACACCTGCAAAAGCGGCAACAACCCAGTGCCACGTGTGAGTTG  
GATAGTTGTGAAAGAGTCAAAATGGCTCTCCTCAAGCGTATTCAACAAGGGGCTGAAGGATG  
CCCAAGAGGTACCCCATTTGATGGGATCTGATCTGGGGCTCGGTGCACATGCTTTACATGT  
GTTTAGTCAGAGTTAAAAAACGTCTAGGCCCCCCGAACCCAGGGGACGTGGTTTCTTTGA  
AAAACACGATGATAATGACCATGGTGAGCAAGCAGATCCTGAAGAACACCCGCTGCAGGAG  
ATCATGAGCTTCAAGGTGAACCTGGAGGGCGTGGTGAACAACACAGTGTTCACCATGGAGGG  
CTGGCGCAAGGGCAACATCCTGTTCGGCAACAGCTGGTGAGATCCGCGTGACCAAGGGCG  
CCCCCTCGCCTTCGCCTTCGACATCCTGAGCCCCGCTTCCAGTACGGCAACCGCACCTTC  
ACCAAGTACCCCGAGGACATCAGCGACTTCTTCATCCAGAGCTTCCCGCGCGGCTTCGTGTA  
CGAGCGACCTTCGCTACGAGGACGGCGGCTGGTGGAGATCCGACGACATCAACCTGA  
TCGAGGGGATGTTCTGTACCGCGTGGAGTACAAGGGCCGCAACTCCCCAACGACGGCCCC  
GTGATGAAGAAGACCATACCGGCTGCAGCCAGCTTCGAGGTGGTGTACATGAACGACGG  
CGTGGTGGTGGGCCAGGTGATCTGGTGTACCGCTGAACAGCGCAAGTTCTACAGCTGCC  
ACATGCGCACCTGATGAAGAGCAAGGCGTGGTGAAGGACTTCCCCGAGTACCACCTTCATC  
CAGCACCGCTGGGAAGACCTACGTGGAGGACGGCGGCTTCGTAGAGCAGCACGAGACCGC  
CATCGCCAGCTGACCAAGCTGGGCAAGCCCTTGGGAGCTGCACGAGTGGTGTAAAGCTC  
GAGCATGCATCTAGAGGGCCCTATTCCCTTTAGTGAGGGTTAATTGCTAGAGCTCGTGATC  
AGCCTCGACTGTGCCTTCTAGTTGCCAGCCATCTGTTGTTGCCCTCCCCCGTGCCTTCTCT  
TGACCTGGAAGGTGCCACTCTCACTGTCTTCTCAATAAAATGAGGAATTGCAATCGCAT  
TGTCTGAGTAGGTGTCATTCTATTCTGGGGGTGGGGTGGGGCAGGACGACAAGGGGAGGA  
TTGGGAAGACAATAGCAGGATGCTGGGGCGCGCGCGCTAAATTGTAAGCGTTAATATT  
TTGTTAAATTCGCGTTAAATTTTGTAAATCAGCTCATTTTTTAAACCAATAGGCCGAAAT  
CGGCAAAATCCCTTATAAATCAAAAGAATAGACCGAGATAGGGTTGAGTGTGTTTCCAGTTT  
GGAAACAAGAGTCCCATATTAAAGAAGTGGACTCCAACGTCAAAGGGCGAAAAACCGTCTAT  
CAGGGCGATGGCCACTACGTGAACCATCACCTTAATCAAGTTTTTGGGGTTCGAGGTGCGG  
TAAAGCACTAAATCGGAACCCCTAAAGGGAGCCCCGATTTAGAGCTTGACGGGAAAGCCGG  
CGAACGTGGCGAGAAAGGAAGGGAAGAAAGCGAAAGGAGCGGGCGCTAGGGCGCTGGCAAAT  
GTAGCGGTACAGCTGCGCGTAACCAACACACCCGCGCGCTTAATGCGCGCTACAGGGCGC  
GTCAGGTGGCACTTTTCGGGAAATGTGCGCGGAACCCCTATTGTTTATTTTCTAAATAC  
ATTCAAATATGTATCCGCTCATGAGACAATAACCTGATAAATGCTTCAATAATATTGAAAA  
AGGAAGAATCCTGAGGCGGAAAGAACAGCTGTGGAATGTGTGCAGTTAGGTGTGGAAG  
TCCCCAGGCTCCCCAGCAGGCAGAAGTATGAAAGCATGCATCTCAATTAGTCAGCAACCAG  
GTGTGGAAAGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGAAAGCATGCATCTCAATTAGT  
CAGCAACCATAGTCCCGCCCTAACTCCGCCAGTTCCGCCATTTCCGCCCATGGCTGA  
CTAATTTTTTTTATTTATGCAAGGCGGAGGCGGCTCGGCTCTGAGCTATTCCAGAAGTA  
GTGAGGAGGCTTTTGTGAGGCTAGGCTTTTGCAAGATCGATCAAGAGACAGGATGAGGA  
TCGTTTCGATGATTGAACAAGATGGATTGCACGAGGTTTCCGCGCGCTTGGGTGGAGAG  
GCTATTCCGCTATGACTGGGCACAACAGACAATCGGCTGCTCTGATGCCCGCTGTTCCGGC  
TGTACGCGCAGGGGCGCCCGGTTCTTTTGTCAAGACCGACCTGTCCGGTGCCTGAAATGAA  
CTGCAAGACGAGGACGCGCGCTATCGTGGCTGGCCACGACGCGGCTTCTTGCAGCTGT  
GCTCGAGCTTGTCACTGAAGCGGGAAGGACTGGCTGCTATTGGGCGAAGTGCCGGGCGAGG  
ATCTCCTGTCTATCTCACCTTGCTCTGCCAGAAAGTATCCATCATGGCTGATGCAATGCGG  
CGCTGCATACGCTTGATCCGGCTACCTGCCATTCGACCACCAAGCGAAACATCGCATCGA  
GCGAGCAGCTACTCGGATGGAAGCCGCTCTTGTGATCAGGATGATCTGGACGAAGAACATC  
AGGGGCTCGCGCCAGCCGAACGTGTCCCGAGGCTCAAGGCGAGCATGCCCGACGGCGAGGAT  
CTCGTGTGACCATGGCGATGCTGCTTGC CGAATATCATGGTGAAAAATGGCCGCTTTTC  
TGGATTTCATCGACTGTGGCCGCTGGGTGTGCGGACCGCTATCAGGACATAGCGTTGGCTA  
CCCGTGATATTGCTGAAGAACCTGGCGGCGAATGGGCTGACCGCTTCTCGTGTTCACGGT

ATCGCGCTCCCGATTTCGCAGCGCATCGCCTTCTATCGCCTTCTTGACGAGTTCTTCTGAGC  
GGGACTCTGGGGTTCGAAATGACCGACCAAGCGACGCCAACCTGCCATCACGAGATTTCGA  
TTCCACCCGCGCCTTCTATGAAAGGTTGGGCTTCGGAATCGTTTTCCGGGACGCCGGCTGGA  
TGATCCTCCAGCGCGGGGATCTCATGCTGGAGTTCTTCGCCACCCTAGGGGGAGGCTAACT  
GAAACACGGGAAGGAGACAATACCGGAAGGAACCCGCGCTATGACGGCAATAAAAAGACAGAA  
TAAACGACGCGGTGTTGGGTCGTTTGTTCATAAACGCGGGGTTCGGTCCCAGGGCTGGCACT  
CTGTCGATACCCCAACCGAGACCCCATTTGGGGCCAATACGCCCGCGGTTTCTTCTTTCCCCA  
CCCCACCCCAAGTTCGGGTGAAGGCCAGGGCTCGCAGCCAACGTGGGGCGGCAGGCC  
TGCCATAGCCTCAGGTTACTCATATATACTTTAGATTGATTTAAACTTCATTTTAAATTTA  
AAAGGATCTAGGTGAAGATCCTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTT  
TCGTTCCACTGAGCGTCAGACCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTTTTT  
TCTGCGCGTAATCTGCTGCTTGCAAAACAAAAAACACCGCTACCGCGGTGGTTTGTGTC  
CGGATCAAGAGCTACCAACTCTTTTCCGAAGGTAAC TGCGTT CAGCAGAGCGCAGATACCA  
AATACTGTTCTTCTAGTGTAGCCGTAGTTAGGCCACCACTTCAAGAACTCTGTAGCACCGCC  
TACATACCTCGCTCTGCTAATCCTGTTACCAGTGGCTGCTGCCAGTGGCGATAAGTCGTGTC  
TTACCGGTTGGACTCAAGACGATAGTTACCGGATAAGGCGCAGCGGTTCGGGCTGAACGGGG  
GGTTCGTGCACACAGCCAGCTTGGAGCGAACGACCTACACGAACTGAGATACCTACAGCG  
TGAGCTATGAGAAAGGCCACGCTTCCCGAAGGGAGAAAGCGGACAGGTATCCGGTAAGCG  
GCAGGTCGGAACAGGAGAGCGCACGAGGGAGCTTCCAGGGGAAACGCTGGTATCTTTAT  
AGTCCTGTCGGGTTTCGCCACCTCTGACTTGAGCGTCGATTTTGTGATGCTCGTCAGGGGG  
GCGGAGCCTATGGA AAAACGCCAGCAACGCGGCCCTTTTACGGTTCCTGGCCTTTTGCTGGC  
CTTTTGCTCACATGTTCTTTCTGCGTTATCCCCTGATTCTGTGGATAACCGTATTACCGCC  
AGGCGCGCCCGGATGTACGGGCCAGATTTACGCGTTGACATTGATTATTGACTAGTTATTAA  
TAGTAATCAATTACGGGGTCATTAGTTTCATAGCCCATATATGGAGTCCGCGTTACATAACT  
TACGGTAAATGGCCCGCCTGGCTGACCGCCCAACGACCCCGCCCATTGACGTCAATAATGA  
CGTATGTTCCCATAGTAACGCCAATAGGGACTTTCATTGACGTCAATGGGTGGAGTATTTA  
CGGTAAACTGCCACTTGGCAGTACATCAAGTGATCATATGCCAAGTACGCCCCCTATTGA  
CGTCAATGACGGTAAATGGCCCGCCTGGCATTATGCCCAGTACATGACCTTATGGGACTTTC  
CTACTTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGGTGATGCGGTTTTGGCAG  
TACATCAATGGGCGTGAGTAGCGGTTTGACTCACGGGGATTTC CAAGTCTCCACCCCATTTGA  
CGTCAATGGGAGTTTGTTTTGGCACCAAAATCAACGGGACTTTC AAAATGTCGTAACAACT  
CCGCCCCATTGAAGCAAATGGGCGGTAGGCGTGTACGGTGGGAGGCTATATAAGCAGAGCT  
CTCTGGCTAACTAGAGAACCCACTGCTTACTGGCTTATCGAAATTAATACGACTCACTATAG  
GGAGACCAAGCTTCTGGAGGCCCGGGCTTTCAGGGTACCGAAGAAGGATCCATGAATGGGA  
CGGAGGGA CTGAATTTCTATGTACCTTTTCAAACAAGACGGGAGTCGTCCGAAGTCCGTTT  
GAATATCCCCAGTATTACCTGGCAGAGCCGTGGCAATTCAGTGTGCTTGCCGCATATATGTT  
CCTTCTGATCGTATTGGGTTTCCCGATAAATTTCTTACCCTCTATGTACGGTGCAGCACA  
AGAAGCTGAGAACACCACTGAACTATATCCTCCTTAATCTGGCGGTTGCGAACCTTTTTCATG  
GTATTCGGCGGATTTACAACAACATTGTACACATCAC