

The Fleischman Lab

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Title	Site-Directed Mutagenesis via QuikChange II XL		
Introduction	DNA. We frequently use this technique to generate plasmic Jak2 ^{v617F} , Calr ^{lns5} , Calr ^{Del52}), but it may also be used for other or FLAG) or restriction enzymes. The protocol described here has been optimized for our lab halved to reduce costs, and the reaction and cycling param for large plasmids. To save time, we use Z-competent (a.k.a XL-10 Gold cells supplied with the kit. While we have descriprotocol, other <i>E. coli</i> strains and transformation methods require the use of ccdB Survival cells). As validation of the rwe have included brief procedures for plasmid propagation. Please note that insertions and deletions are limited to rou recommend sequential mutagenesis reactions. Additionally QuikChange tool may not be specific enough for all sequen such as Pfu Ultra have the potential to introduce unintended mutated plasmids must be screened for unintended mutated interest. For more information, especially if you are new to site-dire protocol for the QuikChange II XL kit: www.agilent.com/cs/library/usermanuals/public/200521.p	cribed here has been optimized for our lab. The mutagenesis reaction volume is costs, and the reaction and cycling parameters have been altered to improve yield is. To save time, we use Z-competent (a.k.a. Mix & Go) DH5α cells as opposed to the upplied with the kit. While we have described the Z-competent cell method in this coli strains and transformation methods may be used (e.g. some of our plasmids f ccdB Survival cells). As validation of the mutated plasmid is extremely important, brief procedures for plasmid propagation and sequencing. Insertions and deletions are limited to roughly 20-25 bp. For larger indels, we ential mutagenesis reactions. Additionally, the primers designed with the may not be specific enough for all sequences, and even high-fidelity polymerases have the potential to introduce unintended base pair changes. For these reasons, is must be screened for unintended mutations by fully sequencing the gene of union, especially if you are new to site-directed mutagenesis, please read the full QuikChange II XL kit: Nocs/library/usermanuals/public/200521.pdf	
Materials	 Plasmid DNA Site-directed mutagenesis primers (100 ng/μl), des design tool at http://www.genomics.agilent.com/p Molecular-grade water Agilent QuikChange II XL site-directed mutagenesis PCR tubes or tube strips Thermal cycler Competent bacterial cells (Z-competent or heat shown in the second in the sec	rimerDesignProgram.jsp kit ock)	

Protocol	A.		Notes
1.	Prepare reactions site-directed mu	ıtagenesis:	The Quiksolution (DMSO) thaws very
	Reagent	Volume	slowly. Remove from the freezer well
	10x reaction buffer	2.5 μl	before preparing the reactions.
	F primer (100 ng/μl)	0.625 μΙ	
	R primer (100 ng/μl)	0.625 μΙ	Do not allow the dNTP mix or Pfu
	dNTP mix	0.5 μΙ	Ultra to stay out of the freezer for
	Quiksolution	1.5 μΙ	longer than is necessary.
	H ₂ O	16.25 μΙ	
	_ Plasmid DNA (10 ng/μl)	2.5 μΙ	
	Pfu Ultra	0.5 μΙ	
		1 : (1 - 1)	5 " 1 (511)
2.	Flick the tubes to mix and spin dow	•	For small plasmids (<5 kb), extension
	thermal cycler and set the cycling 1x 95°C	2.5 min	time may be reduced to 1 min/kb.
	18x 95°C	1 min	
	60°C	1 min	
	68°C	2 min/kb	
	1x 68°C	7 min	
	1x 4°C	∞	
3.	Add 0.5 μl Dpnl to each tube. Flick		Do not allow the DpnI to stay out of
	spin down briefly. Incubate in the	thermal cycler at 37°C for	the freezer for longer than is
	1 hour.		necessary.
	B. Transform bacteria (Z-con	nnetent method)	
1.	Pre-heat two LB-agar plates contains		Plates should be agar-side up to
	antibiotic(s) per transformation at		prevent condensation from dripping
	agar is warm to the touch.		onto the agar surface.
2.	Transfer 2 μl of plasmid to a 1.5 m	I microcentrifuge tube	
	and incubate on ice for 15 minutes	S.	
3.	Thaw an aliquot of Z-competent D		Not all plasmids are compatible with
	mix and spin down briefly, then ge	ently pipette up and down	DH5 α cells. Be sure you are using the
	2-3 times to resuspend cells.		correct strain and transformation
	Transfer 25 of a factor in the first	ا باد الناد مسم مطهم	method for your plasmid.
4.	Transfer 25 µl of competent cells t	•	Return unused competent cells to the
	containing the plasmid and incuba	ite on ice for 5 minutes.	-80°C freezer.
5.	Pipette 10 μl of the transformation	n mixture into the center	Sterilize spreader by dipping in 100%
J.	of a pre-warmed LB-agar plate. Us		EtOH and lighting on fire. Allow to
	gently and evenly spread the cells	·	cool for 20 seconds before using.
	the plate.	over the chine surface of	coorjor 20 seconds before using.
	•		
6.	Sterilize the spreader and repeat s	tep 5 with the remaining	
	transformation mixture (≈17 μl) or	•	
	, , ,	•	
7.	Return plates to the incubator. Inc	ubate overnight. Do not	Plates should be agar-side up to
	incubate more than ~16-18 hours	to prevent formation of	prevent condensation from dripping
	satellite colonies.		onto the agar surface.

	C. Isolate and analyze plasmids	
1.	Prepare LB media containing the appropriate antibiotic(s).	
	Prepare 14 ml culture tubes by adding 3 ml LB+antibiotic	
	per tube.	
2.	Pick one colony per tube by using a sterile 200 μl pipette tip	Only pick isolated colonies.
	to gently touch the colony. Eject the pipette tip into the	
	culture tube. Cap loosely and incubate in the shaker at 37°C	Longer culture times generally
	250 rpm for 8-16 hours or until media becomes cloudy.	produce more concentrated
		minipreps.
3.	Miniprep 1-2 ml of bacterial culture using the Zymo ZR	The protocol is posted above the
	Plasmid Miniprep-Classic kit.	bacteria bench.
4.	Measure DNA concentration on the NanoDrop. Prepare	Plasmid DNA concentration must be
	sequencing reactions for up to 6 colonies per mutagenesis	60 ng/μl or greater for sequencing.
	reaction:	
	Plasmid DNA 600 ng	Choose a sequencing primer that
	Primer (10 μM) 2 μl	reads the region of your intended
	H ₂ O to 12 μl	mutation.
	- '	
5.	Analyze sequencing results for a) successful mutagenesis,	Tip: Align the sequencing results to
	and b) absence of unintended base pair changes. If	the intended sequence using BLAST to
	mutagenesis was successful, choose one plasmid for further	search for unintended base pair
	validation. Prepare sequencing reactions as in step 4 for	changes.
	additional sequencing primers as needed to read the	
	complete region of interest. Ensure no unintended base	
	pair changes were made before using the altered plasmid in	
	downstream applications.	