Strain conservation

Date: 2022-08-17

Tags: 1_Wetlab 3_Microbiology
Created by: Stefanie Brands

1/2

(_Written by Stefanie Brands_)

(_Last update: 2022.08.18_)

After cloning or receiving of a new genetic construct, every strain is conserved in the {strain collection|aim}. In order to confirm the DNA sequence of the construct, sequencing is performed before conservation.

{27.04.2022|Date of experiment}

The plasmid DNA of a $\{5 \text{ mL LB-Kan}|:\text{preculture:}\}\$ is extracted with the $\{\text{Macherey-Nagel NucleoSpin}|:\text{Plasmid isolation kit:}\}\$, following the protocol for $\{\text{low-copy plasmids}|\text{protocol}\}\$. Elution was performed in $\{50|\mu\text{L}|\text{H2O}|\text{elution volume}\}\$ at $\{70|^{\circ}\text{C}|\text{elution temperature}\}\$. The DNA concentration was measured with an $\{\text{Implen N60 Nanophotometer}|\text{device}\}\$. Plasmid DNA of three colonies was isolated. The concentrations were $\{1199.9|\text{ng}/\mu\text{L}|(:\text{for:})\text{ clone 1}|\text{DNA}\}\$, $\{999.3|\text{ng}/\mu\text{L}|(:\text{for:})\text{ clone 2}|\text{DNA}\}\$, and $\{1185.9|\text{ng}/\mu\text{L}|(:\text{for:})\text{ clone 3}|\text{DNA}\}\$.

Clone	Concentration [ng/μl]	260/280	260/230	
1	1199.8	1.95	2.33	
2	999.3	1.94	2.32	
3	1185.9	1.94	2.31	

Sequencing was performed at {Eurofins Genomics|sequencing service provider} on plasmids of clones 1-3 to confirm the DNA sequence and the success of the mutagenesis. The sequencing primer {TATAAGCAGAGCTGGTTTAGTGAACC|sequencing primer} was used in the premix. According to

Strain conservation

Date: 2022-08-17

Tags: 1_Wetlab 3_Microbiology
Created by: Stefanie Brands

2/2

Eurofins instructions, the premix was set up with $\{5|\mu L|: plasmid DNA - diluted to 100 ng/\mu L -: |premix component\} and <math>\{5|\mu L| (:of:) 5 \mu M \text{ sequencing primer}| premix component\}.$

Seq. ID	Primer	Sequence	Length [bp]	Tm [°C]	Target	Result
FMA451	FXR_Seq_fw	TATAAGCAGAGCTGGTTTAGTGAACC	26	57	pnoCherry pCMV	G002A
FMA455	FXR_Seq_fw	TATAAGCAGAGCTGGTTTAGTGAACC	26	57	pnoCherry pCMV	G002A
FMA454	FXR_Seq_fw	TATAAGCAGAGCTGGTTTAGTGAACC	26	57	pnoCherry pCMV	G002A
FMA452	FXR_Seq_fw	TATAAGCAGAGCTGGTTTAGTGAACC	26	57	pnoCherry pCMV	WT

Because sequencing was successful for all clones, clone 2 was chosen randomly for cryoconservation.

{05.05.2022|Date of experiment}

A $\{5 \text{ mL LB-Kan}|: overnight preculture:}$ was grown over night for cryoconservation. $\{\text{Three aliquots per strain} | \text{number of cryotubes} \}$ of $\{500 | \mu L | \text{preculture} | \text{cryoconservation mix} \}$ were mixed with $\{500 | \mu L | \text{sterile ice cold } 50\% \text{ (w/v) glycerol} | \text{cryoconservation mix} \}$ each and stored at $\{-80 \text{ °C} | \text{storage temperature} \}$ in cryotubes. Additionally, $\{\text{three minipreps of } 100 \text{ ng/}\mu L | \text{:plasmid DNA:} \}$ in $\{\text{TE buffer} | \text{eluent} \}$ originating from the same colony were stored at -80 °C in the plasmid collection.