Obtain *hmaS* mutant fragment by PCR Adjust the concentration of Mn<sup>2+</sup> and Mg<sup>2+</sup> to determine the appropriate mutation rate

PCR system (50 µL)		PCR system (50 µL)			
Scpa1-F	2 µL	Scpa1-F	2 μL		
Hmas-R	2 µL	Hmas-R	2 µL		
gy9s(scpa1)-177-hmas3	2 ng	gy9s(scpa1)-177-hmas3	2 ng		
2 x Mix	25 μL	2 x Mix	25 µL		
Mn <sup>2+</sup> (1 mM)	2 µL	Mn <sup>2+</sup> (1 mM)	0.5 µL		
Mg <sup>2+</sup> (25 mM)	2.5 μL	Mg <sup>2+</sup> (25 mM)	2.5 μL		
dNTP	1 μL	dNTP	1 µL		
DDW	14.5 μL	DDW	16 µL		
PCR system (50 µL)		PCR system (50µ	PCR system (50µL)		
Scpa1-F	2 µL	Scpa1-F	2 μL		
Hmas-R	2 µL	Hmas-R	2 µL		
gy9s(scpa1)-177-hmas3	2 ng	gy9s(scpa1)-177-hmas3	2 ng		
2 x Mix	25 μL	2xMix	25 µL		
Mn <sup>2+</sup> (1 mM)	1 μL	Mn <sup>2+</sup> (1 mM)	1.5 μL		
Mg <sup>2+</sup> (25 mM)	2.5 μL	Mg <sup>2+</sup> (25 mM)	2.5 μL		
dNTP	1 µL	dNTP	1 μL		
DDW	15.5 μL	DDW	15 µL		

The target fragment is located on the Pccd-K-scpa1 plasmid, and the amplified target fragment is obtained by PCR.

The PCR products are detected by agarose gel electrophoresis, and the correct target fragment is 1248bp. We obtained the correct target fragment, and the sample is purified.

## 1. Obtain Bio177 vector fragment by PCR

The target fragment is located on the pYB1a-Bio177-Str plasmid, and the amplified target fragment is obtained by PCR. The PCR system and procedure are showed below.

PCR system (50 µL)				
Bio177-F	2 µL			
Bio177-R	2 µL			
gy9s(scpa1)-177-hmas3	2 ng			
2 x Mix	25 µL			
DDW	20 µL			

The PCR products are detected by agarose gel electrophoresis, and the correct target fragment is 5089bp. We obtained the correct target fragment, and the sample is purified.

## 2. Golden gate assembly

The *hmaS* mutant fragments and Bio177 vector fragments are connected by Golden gate assembly method, and the system is as follows.

Golden gate system (50 µL)			Golden gate		
hmaS mut	1 μL	37	°C	60 min	
Bio177	1 μL	55	°C	15 min	
T4 Buffer	1 µL	80	°C	15 min	
rCutsmart	1 µL				
T4 ligase	0.2 μL				
BsaI	0.5 μL				
DDW	5.3 μL				

## 3. Colony PCR

After cultured at 37°C for 12 hours, 30 colonies were selected on the plate. The colony PCR system and procedure were as follows.

PCR system (50 µL)

Bio177-R	2 µL
gy9s(scpa1)-177-hmas3	2 ng
2 x Mix	25 µL
DDW	20 µL

The PCR products were detected by agarose gel electrophoresis, and the results were as follows.



4. Sequencing for ensured appropriate mutation rate



