

1) Prepare 6 ml of 5X ISO Buffer in a 15 ml falcon tube as follows:

3 ml 1 M Tris-HCl pH 7.5  
+ 150  $\mu$ l 2 M MgCl<sub>2</sub>  
+ 240  $\mu$ l 100 mM dNTP mix (25 mM each: dGTP, dCTP, dATP, dTTP)  
+ 300  $\mu$ l 1 M DTT  
+ 1.5 g PEG-8000  
+ 300  $\mu$ l 100 mM NAD  
+ \_\_\_\_\_ dH<sub>2</sub>O to  
6 ml

Store at -20 C in 320  $\mu$ l aliquots.

2) Prepare 1.2 ml of Gibson assembly master mix as follows:

320  $\mu$ l 5X ISO Buffer  
+ 0.64  $\mu$ l 10 U/ $\mu$ l T5 exonuclease\*  
+ 20  $\mu$ l 2 U/ $\mu$ l Phusion polymerase  
+ 160  $\mu$ l 40 U/ $\mu$ l Taq ligase  
+ \_\_\_\_\_ dH<sub>2</sub>O to  
1.2 ml

Store at -20 C in 15  $\mu$ l aliquots.

\*This is optimized for 20-150 bp sequence homology overlaps

3) Thaw a 15  $\mu$ l aliquot of the Gibson assembly master mix, and keep on ice until use.

4) Measure the DNA concentration (ng/ $\mu$ l) of each assembly piece.

5) Add 100 ng of the linearized vector backbone and equimolar amounts of the other assembly pieces to the thawed 15  $\mu$ l master mix in a 20  $\mu$ l total volume assembly reaction mixture as follows:

linearized vector backbone (100 ng)  
+ each additional assembly piece (to equimolar with backbone)  
+ 15  $\mu$ l Gibson assembly master mix  
+ \_\_\_\_\_ dH<sub>2</sub>O to  
20  $\mu$ l

6) Incubate the assembly reaction at 50 C for 60 minutes, and then place on ice.

7) Transform 5  $\mu$ l of the assembly reaction into 100  $\mu$ l of competent *E. coli* and/or run a diagnostic agarose gel to check for successful assembly.