

# Protocol for VTPHG

## 1. Oligo design for VTPHG

1.1 Find NGG in upstream -150 to -500 of a target genes TSS.

1.2 Select sgRNA: 20bp length upstream of NGG and do not have TTTT in it. Higher

CG number gives more efficient activation

1.3 add overhang to Sense: 5'- ttcg-N20

add overhang to Anti-sense: 5'- aaac-N20

eg: Target: 5'-AGCAATCGACATGCAAGCGGCTCGGAGCCAGGGGCACCTGC- 3'

Sense: ttcgCATGCAAGCGGCTCGGAGCC

Anti-sense: aaacGGCTCCGAGCCGCTTGCATG

1.4 Annealing: resuspend oligos with water to 20uM

Set up annealing reaction: primer-F 5ul

Primer-R 5ul

\*1x annealing buffer 40ul

95°C 5 min then close the PCR Amplifier and let it cool to RT

slowly

\*10x annealing buffer: 100mM Tris-HCl (pH 7.5), 10mM EDTA, 1M

NaCl

## 2. Backbone preparation

2.1 Use BbsI to digest VTPHG (see map).

Set up digestion reaction: BbsI(NEB) 1ul

10x Cutsmart buffer (NEB) 5ul

Plasmid	Xul (2ug)
Add water to	50ul
37°C	1.5h

2.2 Run agarose gel, we can see two bands, the small one is about 700bp ( this can facilitate us to select correct clone by PCR), the large one is about 14kb, cut the large band and extract the backbone by Qiagen kit. Measure the DNA concentration, shall be around 50ng/ul.

### 3. Ligation

#### 3.1 clone sgRNA into VTPHG

Set up ligation reaction: sgRNA	4ul
VTPHG	1ul
T4 DNA ligase (Roche)	0.5ul
T4 DNA ligase buffer (Roche)	1ul
Add water to	10ul
16°C	1h

#### 3.2 Transformation

Add 5ul ligation product into 20ul TOP10 competent cells, following standard transformation, then plate.

PCR select the correct clone

Primers used: U6BP-Fs: TCAACAAACGaacaataggacac

sgRNA2.0-Rs: aAAAAAGCACCGACTCGGTG

The correct clone will give around 300bp PCR product, while 1kb is incorrect.

### 3.3 Culture bacteria and miniprep

#### **StandaPurification and injection**