

## Generation of Str-KDEL-IRES-MannosidaseII-mScarlet-i

Str-KDEL\_ManII-SBP-mCherry (Plasmid #65253) from Franck Perez was used as a template.

The insert EcoRI\_mScarlet-i\_FseI was synthesised as gBlocks® Gene Fragment by Integrated DNA technologies (724bp):

```
taagcagaattcATGGTGAGCAAGGGCGAGGCAAGTCAAGGAGTTCATGCGGTTCAAGGTGCACATGGA  
GGGCTCCATGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCCGCCCTACGAGGGCACCC  
AGACCGCCAAGCTGAAGGTGACCAAGGGTGGCCCCCTGCCCTTCTCCTGGGACATCCTGTCCCCTCAG  
TTCATGTACGGCTCCAGGGCCTTCATCAAGCACCCCGCCGACATCCCCGACTACTATAAGCAGTCCTT  
CCCCGAGGGCTTCAAGTGGGAGCGCGTGATGAACTTCGAGGACGGCGGGCCCGTGACCGTGACCCAGG  
ACACCTCCCTGGAGGACGGCACCTGATCTACAAGGTGAAGCTCCGCGGCACCAACTTCCCTCCTGAC  
GGCCCCGTAATGCAGAAGAAGACAATGGGCTGGGAAGCGTCCACCGAGCGGTTGTACCCCGAGGACGG  
CGTGCTGAAGGGCGACATTAAGATGGCCCTGCGCCTGAAGGACGGCGGCCGCTACCTGGCGGACTTCA  
AGACCACCTACAAGGCCAAGAAGCCCGTGCAGATGCCCGGCGCCTACAACGTCGACCGCAAGTTGGAC  
ATCACCTCCACAACGAGGACTACACCGTGGTGAACAGTACGAACGCTCCGAGGGCCGCCACTCCAC  
CGGCGGCATGGACGAGCTGTACAAaggccggcctTAAGca
```

### Cloning procedure:

- Double digest of template and insert with EcoRI-HF and FseI (to remove the SBP and mCherry-tags from the template)
- digested template was dephosphorylated (antarctic phosphatase NEB), and gel purified
- both fragments were ligated using T4 ligase (Promega) into the final product of 6146bp.

Sequencing revealed the complete sequence as the following:

```
[...] tggttttcctttgaaaaacacgatgataagcttgccacaaccgggagggcgccATGAAGTTAA  
GTCGCCAGTTCACCGTGTTTTGGCAGCGGATCTTCTGCGTCGTAATCTTCTCACTCTACCTGATGCTG  
GACAGGGGTCACTTGGACTACCTTCGGGGCCGCGCCAGGAGGGTCTTTTCCGCAGGGCCAGCTTTC  
AATATTGCAAGAAAAGATTGACCATTTGGAGCGTTTGCTCGCTGAGAACAACGAGATTATCTCAAATA  
TCAGAGACTCAGTCATCAACCTGAGCGAGTCTGTGGAGGACGGCCCGGGGGTCAACCAGGCAACGCC  
AGCCAAGGCTCCATCCACCTCCACTCGCCACAGTTGGCCCTGCAGGCTGACCCCAGAGACTGTTTTGA  
TCCCACCGGTGCGCACCGcaattcATGGTGAGCAAGGGCGAGGCAAGTCAAGGAGTTCATGCGGT  
TCAAGGTGCACATGGAGGGTCCATGAACGGCCACGAGTTCGAGATCGAGGGCGAGGGCGAGGGCCGC  
CCCTACGAGGGCACCCAGACCGCCAAGCTGAAGGTGACCAAGGGTGGCCCCCTGCCCTTCTCCTGGGA  
CATCCTGTCCCCTCAGTTCATGTACGGCTCCAGGGCCTTCATCAAGCACCCCGCCGACATCCCCGACT  
ACTATAAGCAGTCCTTCCCCGAGGGCTTCAAGTGGGAGCGCGTGATGAACTTCGAGGACGGCGGGCC  
GTGACCGTGACCCAGGACACCTCCCTGGAGGACGGCACCTGATCTACAAGGTGAAGCTCCGCGGCAC  
CAACTTCCCTCCTGACGGCCCCGTAATGCAGAAGAAGACAATGGGCTGGGAAGCGTCCACCGAGCGGT  
TGTACCCCGAGGACGGCGTGCTGAAGGGCGACATTAAGATGGCCCTGCGCCTGAAGGACGGCGGCCG  
TACCTGGCGGACTTCAAGACCACCTACAAGGCCAAGAAGCCCGTGCAGATGCCCGGCGCCTACAACGT  
CGACCGCAAGTTGGACATCACCTCCACAACGAGGACTACACCGTGGTGAACAGTACGAACGCTCCG  
AGGGCCGCACTCCACCGGGCGCATGGACGAGCTGTACAAaggtgcttaaGGccggccttaaggcctc  
gagggccaattaattaactctagataactgatcataa [...]
```

Seq. Primers (5' -> 3'):

IRES_fwd	TGGCTCTCCTCAAGCGTATT	mScarlet-i_m2_fwd	CCCCGACTACTATAAGC
ManII_m1_fwd	ACCCAGAGACTGTTTGG		