Vector Map



pLVXS-sgRNA-mCherry-hyg Vector

Catalog No.

632649 (Not sold separately) Sold as a part of 632646





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Location of Features

- 5' LTR (5' long terminal repeat): 1–635
- Ψ (packaging signal): 685–822
- RRE (Rev-response element): 1,303–1,536
- cPPT/CTS (central polypurine tract and central termination sequence of HIV-1): 2,028–2,151
- *P*_{U6} (U6 promoter): 2,215–2,455
- P5 primer binding sequence: 2,443–2,464
- sgRNA Scaffold: 2,491–2,582
- P7 primer binding sequence: 2,593–2,623
- *P*_{PGK} (phosphoglycerate kinase promoter): 2,628–3,131
- mCherry (red fluorescent protein): 3,166–3,876
- IRES (encephalomyocarditis virus internal ribosome entry site): 3,895–4,493
- Hyg^R (hygromycin resistance gene): 4,494–5,519
- WPRE (woodchuck hepatitis virus posttranscriptional regulatory element): 5,567–6,158
- 3' SIN LTR (3' self-inactivating long terminal repeat): 6,366–6,602
- ori (high-copy-number ColE1/pMB1/pBR322/pUC origin of replication): 6,978–7,651 (complementary)
- Amp^R (ampicillin resistance gene): 7,796–8,656 (complementary)
- P_{AMP}^{R} (promoter for ampicillin resistance): 8,657–8,761 (complementary)

Note: The 35-base sequence between the two BsmBI sites will be replaced by your guide sequence. It is not meant as a control.

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This document has been reviewed and approved by the Quality Department.