

10 20 30 40 50 60 70

Top
 AlaAlaAlaIleTrpLeuAla . LeuValAspSerLysTrpValAlaAspProAsnSerCysProThrVal
 ArgGlnProTyrGly . HisAspTrpTrpThrAlaAsnGlySerArgIleArgIleHisAlaGlnGln .
 GlySerHisMetAlaSerMetThrGlyGlyGlnGlnMetGlyArgGlySerGluPheMetProAsnSerGlu

pET-28a_WRS(1-48)-2-T7ter.seq(1>1279) ← GCGGCAGCCATATGGCTAGCATGACTGGTGGACAGCAAATGGGTCGCGGATCCGAATTCATGCCCAACAGTG
 pET-28a_WRS(1-48)-2-T7.seq(1>1286) → ATGGGTCGCGGATCCGAATTCATGCCCAACAGTG
 WRS(1-48).seq(1>144) → EcoR atgcccaacagtg

80 90 100 110 120 130 140

Top
 SerProHisLeuCysTrpSerCysSerThrAlaSerProHisLysGlySerSer . GlyProSerLysArg
 AlaArgIleSerAlaGlyAlaValGlnGlnHisArgHisThrArgGlyAlaArgLysValProGlnSerGly
 ProAlaSerLeuLeuGluLeuPheAsnSerIleAlaThrGlnGlyGluLeuValArgSerLeuLysAlaGly

pET-28a_WRS(1-48)-2-T7ter.seq(1>1279) ← AGCCCGCATCTCTGCTGGAGCTGTTCAACAGCATCGCCACACAAGGGGAGCTCGTAAGGTCCCTCAAAGCGG
 pET-28a_WRS(1-48)-2-T7.seq(1>1286) → AGCCCGCATCTCTGCTGGAGCTGTTCAACAGCATCGCCACACAAGGGGAGCTCGTAAGGTCCCTCAAAGCGG
 WRS(1-48).seq(1>144) → agcccgcatctctgctggagctgttcaacagcatcgccacacaaggggagctcgtaaggtccctcaaagcgg

150 160 170 180 190 200 210

Top
 GluMetArgGlnArgMetLysLeuIleLeuGln . ArgCysTrpCysHis . LysCysSerSerSerThr
 LysCysValLysGly . Asn . PheCysSerLysAspValGlyValIleLysAsnValAlaArgAlaPro
 AsnAlaSerLysAspGluIleAspSerAlaValLysMetLeuValSerLeuLysMet . LeuGluHisHis

pET-28a_WRS(1-48)-2-T7ter.seq(1>1279) ← GAAATGCGTCAAAGGATGAAATTGATTCTGCAGTAAAGATGTTGGTGTTCATTAATAATGTAGCTCGAGCACC
 pET-28a_WRS(1-48)-2-T7.seq(1>1286) → GAAATGCGTCAAAGGATGAAATTGATTCTGCAGTAAAGATGTTGGTGTTCATTAATAATGTAGCTCGAGCACC
 WRS(1-48).seq(1>144) → gaaatgcgtaaaggatgaaattgattctgcagtaaagatgttggtgtcattaaaaatgtagctcgagcacc Xho

220 230 240 250

Top
 ThrThrThrThrThrGluIleArgLeuLeuThrLysProGlu
 ProProProProLeuArgSerGlyCys . GlnSerProLys
 HisHisHisHis . AspProAlaAlaAsnLysAlaArg

pET-28a_WRS(1-48)-2-T7ter.seq(1>1279) ← ACCACCACCACCACTGAGATCCGGCTGCTAACAAGCCCGAAA
 pET-28a_WRS(1-48)-2-T7.seq(1>1286) → ACCACCACCACCACTGAGATCCGGCTGCTAACAAGCCCGAAA