

Molecular Biology

Entry Clone Accession:

Entry Clone Source: Invitrogen

SGC Construct ID: MTHFRA-c042

Protein Region: D38-N644

Vector: pFB-CT10HF-LIC

Tag: C-TEV;C-10HIS;C-FLAG

Host: DH10Bac

Sequence (with tag(s)):

MDPERHERLREKMRRRLES GDKWFSLEFFPPRTAEGAVNLSRFDRMAAGGPLYIDVTW
HPAGDPGSDKETSSMMIASTAVNYCGLETILHMTCCRQRLEEITGHLHKAKQLGLKNIM
ALRGDPIGDQWEEEEEGGFNYAVDLVKHIRSEFGDYFDICVAGYPKGHP EAGSFEADLKHL
KEKVSAGADFIITQLFFEADTFFRFVKACTDMGITCPIVPGIFPIQGYHSLRQLVKLSKLEV
PQEIKDVIEPIKDNDAAIRNYGIELAVSLCQELLASGLVPGLHFYTLNREMATTEVLKRLG
MWTEDP RRPLPWALSAHPKRREEDVRPIFWASRPKSYIYRTQEWDEFPNGRWGNSSSPAF
GELKDYYLFY LKSKSPKEELLKMWGEELTSEASVFEV FVLYLSGEPN RNNGHKVTCLPWN
DEPLAAETSL LKEELLRVNRQGILTINSQPNINGKPSSDPIVGWGPSGGYVFQKAYLEFFTS
RETAEALLQVLKKYELRVNYHLNVKGENITNAPELQPN AVTWGIFPGREIIQPTVVDPV
SFMFWKDEAFALWIEQWGKLYEEESPSRTIIQYIHDNYFLVNLVDNDFPLDNCLWQVVED
TLELLNAENLYFQSHHHHHHHHHHDYKDDDDK

Sequence after tag cleavage:

MDPERHERLREKMRRRLES GDKWFSLEFFPPRTAEGAVNLSRFDRMAAGGPLYIDVTW
HPAGDPGSDKETSSMMIASTAVNYCGLETILHMTCCRQRLEEITGHLHKAKQLGLKNIM
ALRGDPIGDQWEEEEEGGFNYAVDLVKHIRSEFGDYFDICVAGYPKGHP EAGSFEADLKHL
KEKVSAGADFIITQLFFEADTFFRFVKACTDMGITCPIVPGIFPIQGYHSLRQLVKLSKLEV
PQEIKDVIEPIKDNDAAIRNYGIELAVSLCQELLASGLVPGLHFYTLNREMATTEVLKRLG
MWTEDP RRPLPWALSAHPKRREEDVRPIFWASRPKSYIYRTQEWDEFPNGRWGNSSSPAF
GELKDYYLFY LKSKSPKEELLKMWGEELTSEASVFEV FVLYLSGEPN RNNGHKVTCLPWN
DEPLAAETSL LKEELLRVNRQGILTINSQPNINGKPSSDPIVGWGPSGGYVFQKAYLEFFTS
RETAEALLQVLKKYELRVNYHLNVKGENITNAPELQPN AVTWGIFPGREIIQPTVVDPV
SFMFWKDEAFALWIEQWGKLYEEESPSRTIIQYIHDNYFLVNLVDNDFPLDNCLWQVVED
TLELLNAENLYFQ

DNA Sequence:

TTAAGAAGGAGATATACTATGGACCCTGAGCGGCATGAGAGACTCCGGGAGAAGATG
AGGCGGCGATTGGAATCTGGTGACAAGTGGTTCTCCCTGGAATTCTTCCCTCCTCGAA
CTGCTGAGGGAGCTGTCAATCTCATCTCAAGGTTTGACCGGATGGCAGCAGGTGGCC
CCCTCTACATAGACGTGACCTGGCACCCAGCAGGTGACCCTGGCTCAGACAAGGAGA
CCTCCTCCATGATGATCGCCAGCACCGCCGTGAACTACTGTGGCCTGGAGACCATCCT
GCACATGACCTGCTGCCGTCAGCGCCTGGAGGAGATCACGGGCCATCTGCACAAAGC
TAAGCAGCTGGGCCTGAAGAACATCATGGCGCTGCGGGGAGACCCAATAGGTGACCA
GTGGGAAGAGGAGGAGGGAGGCTTCAACTACGCAGTGGACCTGGTGAAGCACATCC
GAAGTGAGTTTGGTGACTACTTTGACATCTGTGTGGCAGGTTACCCCAAAGGCCACCC
CGAAGCAGGGAGCTTTGAGGCTGACCTGAAGCACTTGAAGGAGAAGGTGTCTGCGG
GAGCCGATTTCATCATCACGCAGCTTTTCTTTGAGGCTGACACATTCTTCCGCTTTGTG
AAGGCATGCACCGACATGGGCATCACTTGCCCCATCGTCCCCGGGATCTTTCCCATCC
AGGGCTACCACTCCCTTCGGCAGCTTGTGAAGCTGTCCAAGCTGGAGGTGCCACAGG
AGATCAAGGACGTGATTGAGCCAATCAAAGACAACGATGCTGCCATCCGCAACTATG
GCATCGAGCTGGCCGTGAGCCTGTGCCAGGAGCTTCTGGCCAGTGGCTTGGTGCCAG
GCCTCCACTTCTACACCCTCAACCGCGAGATGGCTACCACAGAGGTGCTGAAGCGCC
TGGGGATGTGGACTGAGGACCC CAGGCGTCCCCTACCCTGGGCTCTCAGTGCCCACC
CCAAGCGCCGAGAGGAAGATGTACGTCCCATCTTCTGGGCCTCCAGACCAAAGAGTT

ACATCTACCGTACCCAGGAGTGGGACGAGTTCCCTAACGGCCGCTGGGGCAATTCCTC
TCCCCCTGCCTTTGGGGAGCTGAAGGACTACTACCTCTTCTACCTGAAGAGCAAGTCC
CCCAAGGAGGAGCTGCTGAAGATGTGGGGGGAGGAGCTGACCAGTGAAGCAAGTGT
CTTTGAAGTCTTTGTTCTTTACCTCTCGGGAGAACCAAACCGGAATGGTCACAAAGTG
ACTTGCCTGCCCTGGAACGATGAGCCCCCTGGCGGCTGAGACCAGCCTGCTGAAGGAG
GAGCTGCTGCGGGTGAACCGCCAGGGCATCCTCACCATCAACTCACAGCCCAACATC
AACGGGAAGCCGTCCTCCGACCCCATCGTGGGCTGGGGCCCCAGCGGGGGCTATGTC
TTCCAGAAGGCCTACTTAGAGTTTTTCACTTCCCGCGAGACAGCGGAAGCACTTCTGC
AAGTGCTGAAGAAGTACGAGCTCCGGGTAAATTACCACCTTGTCATGTGAAGGGTG
AAAACATCACCAATGCCCCTGAACTGCAGCCGAATGCTGTCACTTGGGGCATCTTCCC
TGGGCGAGAGATCATCCAGCCCACCGTAGTGGATCCCGTCAGCTTCATGTTCTGGAAG
GACGAGGCCTTTGCCCTGTGGATTGAGCAGTGGGGAAAGCTGTATGAGGAGGAGTCC
CCGTCCCGCACCATCATCCAGTACATCCACGACAACCTACTTCCTGGTCAACCTGGTGG
ACAATGACTTCCCCTGGACAACCTGCCTCTGGCAGGTGGTGGAAGACACATTGGAGC
TTCTCAACGCAGAGAACCTCTACTTCCAATCGCACCATCATCACCATCACCATCACC
CCATGATTACAAGGATGACGACGATAAGTGAGGATCC

Protein Expression

Procedure: Protein was expressed in insect cells in 6L Sf9 media (ThermoFisher), infected with 3mL/L P2 virus.

Protein Purification

Procedure: Protein was purified by affinity (Ni-NTA, Qiagen) and size-exclusion (Superdex 200) chromatography, followed by cleavage of the C-terminal tag by His-tagged tobacco etch virus protease (1:20 mass ratio) overnight at 4°C and re-passage over Ni-NTA resin. Selenomethionine(SeMet)-derivatized proteins were expressed using SelenoMethionine Medium Complete (Molecular Dimensions) and purified as above.

Concentration: 21.0 mg/ml

Mass-spec Verification: yes

Structure Determination

Crystallization: SeMet-derivatized and native *HsMTHFR*₃₈₋₆₄₄ were concentrated to 15-20 mg/ml, and crystals were grown by sitting drop vapour diffusion at 20°C, in mother liquor containing 0.1M Na citrate tribasic, 22.5% PEG4K, 5% 2-propanol.

Data Collection: *Beamline:* Dmnd I04-1; *Resolution:* 2.5 Å

Data Processing: X-ray diffraction data were collected at the Diamond Light Source and processed using XIA2. The *HsMTHFR*₃₈₋₆₄₄ structure was solved by selenium multi-wavelength anomalous diffraction phasing using autoSHARP, and subjected to automated building with BUCCANEER.