

Name	Gene Sequence	Translation
<i>Streptomyces catenulae</i> strain NRRL B-2342 large terminase subunit gene	>NZ_JODY01000031.1:c42687-41037 Streptomyces catenulae strain NRRL B-2342 GTGAGTCCCATTCCAGCCCTCACAGCAGAAGAGATCGACGACCTTGAACCGACCTTCCTCGGCCCGACGT GGCAGAAGGGAGCCGATGGTTCTGGAAGCTCCCCGAGCGCACCCCTCGGATGGCAGATCGCAGGCTGGGC TGCTGAGTTCTCAAGGCCGAGGACGGCGGCCCTGGCGTTTCACGAGCGAGCAACTGCGCTTTGTCTTG TGGTGGTACGCGGTCGATGAGGCCGGTTCGTTTTCGTTTACCGCAAAGGTGTTTTGCAGCGTCTAAAGGGCT GGGGC TTA_TAGC AAGGACCTCTTCTCGCCGTAATAAGCGTGGTTCGAGTTCGTGGGGCCCTCACGCTTTT CGCACTGGGGCCAGACGGACAGCCGTTGGCGTCCCGCACCCGCAGGCGTGGGTGCAGATCGCGGCCGT CAGCCGTGATCAGACGCGCAACACGATGACGCTGATGCCGTCTTTGCTCTCAGACAACTCATCGAGACG TACGGCATCAAGGCTGGGGCCGAAGTATCCGCGCGAACGGCGGCCGAGAGGCTCGAAGCAGTCACCT CATCGTTCCGCGCCCTTGAGGGCGGCCGAGTCACTTTCACAGTGCTCAACGAAACGCACCATTGGGTGAG TGGCAATAACGGCGACCGTATGTACGAGACGATCGACGGTAACGCGACGAAAAAGGACTCGCGCTATCTG GCGATCACGAACGCCTTTCTCCCCGGCGAGGATTCCGTCGGGGAGAAGATGCGTGAGGCGTACGAGAAAA TCCTTGAAGGCCGTGCGGCCGACGTGGGCTTCATGTACGACTCCGTCGAGGCCACCCCGCGACTCCGCT GACGCCGGAGGCAATCCGGATCGTCCTACCGAAGATCCGCGGCGACGCCATCTGGCTGAAGGTCGAGACG ATCCTCCAGTCCATCCTGGACACGACCATCGCACCTTCGCGCTCGCGCCGGATGTGGCTGAACCAGATCG TCGCCCTCCGAGGATGCCCTCTACGGCCCCGGCCGAGTGGGACGTCTGCGCGACGAGTCTCTCCACCTCGA CCCCGGCGACGAGATCACCTCGGCTTCGACGGCGGCAAGACGGACGACTCGACTTGCCTGGTGGCCGTC CGCACCTCGGACATGGCTGCTTTCGTCTCGGCCCTGTGGGAGAAGCCGGATGGCCCGAAGGGTGACGGCT GGATCGTTCCCCGGGCTCAGGTCGACAGCGCCGTGCACGACGCGTTCCGCGTCTTCGACGTGACGGCGTT CTACGCCGACGTGGCCCTGTGGGAGAGCTACATCTCCGAGTGGGACGCCGCGTACGGCGAGGGCCTGGCC GTGAAGGCACCGGGCAAGGACGCGATCGGCTGGGACATGCGCTCCAGCCTGAAGCTCTCCACCATGGCCC ACGAGCGGCTGATGCGGTTCGATCTTCGATCGCAAGCTGAAGCACGACGGTGACCGCAAGCTCCGCCGGCA CACGCTGAACGCGATGCGCCGCTCGACCAACTACGGCGTCTCCTTCGGCAAGGAGAGCCGCGAGTCGCCG AAGAAGGTCGACGCGTACGCCGCGCTGATGCTCGCGCACGAGGCCCTCGTCGACTTCCGCGCCCGAGGCA AGAAGGTCCGCAAGCGGACCGGCCGGGGCTATTTCTTTTAA	>Streptomyces_cat_g6 VSPIPALTAEEIDDLEPTFLGPTWQKGADGSWKLPERTLGWQIAGWAAEFLKAEDGGPWRFTSEQLRFVLW WYAVDEAGRFBVYRKGVLRQLKGGW TTA_TAGC >Streptomyces_cat_g78 KDPLLAVISVVEFVGPSRFSHWGPDGQPVGVPHQPQAWVQIAAVSRDQTRNTMTLMPSLLSDKLIETYGIKAG AELIRANGGRQRLEAVTSSFRALEGGRVTFVLNETHHWVSGNNGDRMYETIDGNATKKDSRYLAITNAFLP GEDSVGEKMRAYEKILEGRAADVGFMYDSVEAHPATPLTPEAIRIVLPKIRGDALWLKVETILQSILDTTI APSRSRMWLNQIVASEDALYGPAEWDVLRDESLHLDPGDEITLGFDDGGKTDSTCLVAVRTSDMAAFVLGL WEKPDGPKGDGWIVPRAQVDSAVHDAFRVFDVTAIFYADVALWESYISEWDAAYGEGVLAVKAPGKDAIGWDMR SSLKLSTMAHERLMRSIFDRKLKHDGDRKLRRHTLNAMRRSTNYGVSFCKESRESPPKKVDAYAAALMLAHEAL VDFRARGKKVRKRTGRGYFL