> AUauTPM1ab-i2, id, ib ( 788 bp: 735 common) GACTCTTCGATTGTGCTCAGTTCGGCAAGTAGCTATCCTCAAATATTCCTTTCCTCGCGCTTTTCGGCACCAAATAAACCTTCAAAATGGAGCAGATTAAGAAG~AAGATGGCCAACTTGAAGAAGCAACAAGATGAAGCTGAAGAAAAAGCAAGGAAAGCTGAAGAAGAGCTCGCTGAAACCAACGCCAAGGCCCAAGCTGCTGAAGAAGATGTCACCAGATTGATTCAAGAAATGGAAAAATTAGAAGAGGAACTAGATTCTACTGAGTCAAAGCTGTCCACAACCATGCAAAAACTGTCAGAGGCTGAAAAACAAGCTGATGAGAGTGAAAGGGCAAGGAAAGTTTTGGAGAACAGAGGAATTACCGATGATGAAAGGCTGACCAGGCTCGAAACTGAACTTGGAGAGCTTACTTCCAAAAATGAAAAGGTTGAAGCAGAGTACGAAGAGACCTGCAATGAAATTAATGATCTAGAGCAAAGACTTGATGAGGAAGAAAGCAAAAGTGAAGAATATGAAGGGAGAGTGAAGGAGCTTGAGGCTGAGGTCATGTTAGTTGGCAACAACCTAAGGAGCTTGGGAATCAGTGAAGGCAAAGCATCTGAGAGAGAGGACACCTATCAATCTAAACTGAACGAACTCAGCGAGAAATTCCAAGAGACTGACGCACAAGCAGAAAGCCTCGAAAACCGCGTGAAGGAACTTGAAAACCAGCTCGCTGACCTCGAAGACCAAAATAGCCAAGACAAAGATGCTTACAATACTGCTAAACGTGACTATGATTC

> AUauTPM2ab-i2, i4 (786 pbases: all common to i2 and i4) CACTCCAAGGCGTTAAAGCTCGTTTAGCTAAAATTCAAGAAGACATCGACGAAGCAGATAACAGGGCGGTAAATGCAAACCATGAAATCCGTGAGGCCCAAACAAGATTGGAGAAGGCTGAAGGAGAATTGGGAACTCATGAACGAAGGCAAAAGCTCCTGGAACTTGAGTTAGAGAATTATAAGGATAGATACACAGAGGCTGCAGATAAGCTGGAAAAGGTCCTGTCCGAATCCGAGGGAATCGAAGTAAAACGTCGTGAACTCGAGGAAACTGAAGCTCATGGAGATGAACGCCTTCAAGAGCTTGAGGAGCAAGTCAAAGAGGCACGACATTTGCTTGAGGAAAATGAGGTGAAGCTGGTTGAAGTAAAGCGCAAAGCTGTTGTATTGAAGCGAGACATCGAAAGAGTGTCCGAAAAGGCCGAAACCCACGAAAAGAGAATGAATGTTTTAGAGGAAACCATCCAATCAACTGGCGAATCTCTCCGCGTCCTCGAAGATCGTGAGCTTGAATCGTCTGACAAGGAGGCTCTCAACGAAGAGAAAATTGCCTTCTTGGAGGGGCAAATGAAAGAAGCTGAAATTCGCTTAGAAGCAGCAGACAGACAAGCTCAAAATTTGGAGAGAGTGATGTTGGACACCGACACAGAAAGAAACGTATGGATCCAAAAGAGAGAAGGAATCGAAAGGGAAATGGCAGATATCGAAGCAATGGACAGCTTGGAGGAAGAATAAAATGAAACCAAAAACATTTCATTCAAACCAATGTGCCTTGTTTGTAGTT

> AUauTPM1ab-i2 (442 bp: 91 common to i2,id and ib) CGAGAAATTCCAAGAGACTGACGCACAAGCAGAAAGCCTCGAAAACCGCGTGAAGGAACTTGAAAACCAGCTCGCTGACCTCGAAGACGACCTACAGAAAGAGAAGGACGACTACGACAAAGTCAAACAAGATTACGACTCAACGATCGCTGAACTCGCTGAAATGTAAACAAAGAACAATTGTTTCGACTCAACTTAGTTAAAGACTGCACCGGATTTAAACTGGATAAATTTCACCTTTTAGGATACTTATTTTTGTAAACGTCTTGTTACCAATAACCTTTAGCTTGCCGATTACTCCTCAAAATGTAAATATGAACTGCTCGTTCAACGTCTTGACAACTGGCGTAGCTTCATTCCGTTCCGTGACAGCTTTTGGCACAAGTATCTTTGATGACTTATTTCCAATGTAGTTGTAGGACTGTCGTAACTGTTTGCGAAA

> AUauTPM1ab-id ( 281 bp: 91 common i2,id, ib) CGAGAAATTCCAAGAGACTGACGCACAAGCAGAAAGCCTCGAAAACCGCGTGAAGGAACTTGAAAACCAGCTCGCTGACCTCGAAGAGGAAGTGACTCGAAAGAAAGATTCCTATCAGAAAATTAAGCATGACTATGATGGTGCTCTAATTGAACTCAGCGACATGTAATAAGTATGCAGGTTTCAAAAACATTGTAGATGCATTTAGATTCTTATGATATAGAATAGAACATGGTATTGTTTCTTTGTTGGAACTCGATGGCCATGTGTCCATTTTCACT

> AUauTPM1ab-ib ( 390 bp: 91 common to i2,id and ib) CGAGAAATTCCAAGAGACTGACGCACAAGCAGAAAGCCTCGAAAACCGCGTGAAGGAACTTGAAAACCAGCTCGCTGACCTCGAAGACCAAAATAGCCAAGACAAAGATGCTTACAATACTGCTAAACGTGACTATGATTCAATGCTTATCGAGATCGTAGAAATGTAAACATTGCAAAGTCATACAACCAAAGGACTTGACAACCGACATTGACCAAACCAGCGTTAACTTGGAGTGTATTTTATAACTAGTGTTAGATTAACTAGAAATTTAACCAGCAGCTAACTGGATTTAGAATAACTCTATTTATCTAACCAGAGTTAGACTAACTAGCGTAAGACTTGAAGCCAAGCATCGAACAGAAAACAGTGTCAACGCTTTACCCTTAA