

## Consensus Alignment: S. cerevisiae MUD1, modified MUD1, Untitl...

GAAACAAAAAAAATGGAAAGTCAGGGTGTGGTGTAAAGCTGCTATCACACAAAGACGGCCTCAGGCACTCAAACCTAAAGAAACCATGTCAGCG  
 consensus sequence Untitled Consensus

GAAACAAAAAAAATGGAA-----AAAGCTGCTATCACACAAAGACGGCCTCA-----TCAAACCTAAAGAAACCATGTCAGCG  
 aligned sequence S. cerevisiae MUD1

GAAACAAAAAAAATGGAAAGTCAGGGTGTGGTGTAAAGNNNNNNNTACGTTAACAAATACGAGGCACTTACTCCGNNNNNNNNNTCAGCA  
 Cloning & Seq homology Index Library specific primer Barcode  
 modified\_MUD1  
 aligned sequence modified MUD1

TATGTATATACTTGTAAATTACGTTCTAAATCTTCTTCTACTAACGTTTCATTATTCTATACTCTATGACCAATAAACAGACTGTACTTCA  
 consensus sequence Untitled Consensus

TATGTATATACTTGTAAATTACGTTCTAAATCTTCTTCTACTAACGTTTCATTATTCTATACTCTATGACCAATAAACAGACTGTACTTCA  
 aligned sequence S. cerevisiae MUD1

AAGTCATATACTTGTAAATTACGTTCTAAATCTTCTTCTGTAGTTGTTTCATTATTCTATACTCTATGACCAATAAACAGGACTGTACTTCA  
 Library variable region  
 modified\_MUD1  
 aligned sequence modified MUD1

AAATTTACCCAGTAGGCCAGCAAATAAGAAAATTATACCAAGATTACTTCTGAAACACATTAATCCAAACAAGTATGCCATTAATCCGTCGCTACCC  
 consensus sequence Untitled Consensus

AAATTTACCCAGTAGGCCAGCAAATAAGAAAATTATACCAAGATTACTTCTGAAACACATTAATCCAAACAAGTATGCCATTAATCCGTCGCTACCC  
 aligned sequence S. cerevisiae MUD1

AAATTTACCCAGTAGGCCAGCAAATAAGAAAATTATACCAAGATTACTTCTGCCTGGAGTTCGCTATTCTATTGTAGTTTCAATGTGACTGCGTTCC  
 Library variable region Library specific primer Cloning & ... homology  
 modified\_MUD1  
 aligned sequence modified MUD1

CTTCCCCACAATAAATTACAAATATCATCACACCCTGATGTTACTAGATGATCAAATGGGCCTCTTGAAGTTCTATTCAAGATCATCAAAGATGA  
consensus sequence Untitled Consensus

CTTCCCCACAATAAATTACAAATATCATCACACCCTGATGTTACTAGATGATCAAATGGGCCTCTTGAAGTTCTATTCAAGATCATCAAAGATGA  
aligned sequence S. cerevisiae MUD1

ACTCCCCACAATAAATTACAAATATCATCACACCCTGATCTTACTAGGTGATCAAAGGGCCTCTTGAAGTTCTATTCAAGATCATCAAAGCTGA  
modified\_MUD1  
aligned sequence modified MUD1

CTAACCAAGCATTAAACGTTGTCACTCAAGAAGAACGACCGGTTCTAGAAAAATACACGACAACAGCATTAAAAGTTCAAGGTCGAAAGTGAG  
consensus sequence Untitled Consensus

CTAACCAAGCATTAAACGTTGTCACTCAAGAAGAACGACCGGTTCTAGAAAAATACACGACAACAGCATTAAAAGTTCAAGGTCGAAAGTGAG  
aligned sequence S. cerevisiae MUD1

CTAACCAAGCATTAAACGTTGTCACTCAAGAAGAACGACCGGTTCTAGAAAAATACACGACAACAGCATTAAAAGTTCAAGGTCGAAAGTGAG  
modified\_MUD1  
aligned sequence modified MUD1

AATGGGAAAGGCTCGAACAAATTGTTATTGGTCTTCAATAGAAATGCAAAAAAAAAAGGTAATGACGAAACGTACAACCTGATATAAGAAGGTG  
consensus sequence Untitled Consensus

AATGGGAAAGGCTCGAACAAATTGTTATTGGTCTTCAATAGAAATGCAAAAAAAAAAGGTAATGACGAAACGTACAACCTGATATAAGAAGGTG  
aligned sequence S. cerevisiae MUD1

ACTGGGAAAGACTCGAACAAATTGTTATTGGTCTTCAATAGAAATCAAAAAAAAAAGGTAATTACGAAACGTACAACCTGATATAAGAAGGTG  
modified\_MUD1  
aligned sequence modified MUD1



CTTAAAGCAAGGAAACTAACGCAAGTTACGTAGTGATGATATGCGCCAAAAAGTTCAAGGCTAAAAGGCAAATAAGACGCCGAAGCATAAGCTGA  
aligned sequence S. cerevisiae MUD1

CTTAAAGCAAGGAAACTAACGCAAGTTACGTAGTGATATTGCGCCACAAAGTTCAAGGCTAAAAGGCAAATAAGACGCCGAAGCATAAGCTGA  
modified\_MUD1  
aligned sequence modified MUD1

GATCAAGAAAGGTAGAGGGAGGCTGAGATTGATAGGATTGTAAAAGAATTGAAACCCGTAGATTGGAAAACATGAAGTCTCAGCAAGAAAATCTAAAACA  
consensus sequence Untitled Consensus

GATCAAGAAAGGTAGAGGGAGGCTGAGATTGATAGGATTGTAAAAGAATTGAAACCCGTAGATTGGAAAACATGAAGTCTCAGCAAGAAAATCTAAAACA  
aligned sequence S. cerevisiae MUD1

GATCAAGAAAGGTAGAGGGAGGCTGAGATTGATAGGATTGTAAAAGAATTGAAACCCGTAGATTGGAAAACATCAAGTCTCAACCAGAAAATCTAAAACA  
modified\_MUD1  
aligned sequence modified MUD1

ATCGCAGAACCTCTTAAGCGGGCTAAAGTGTCCAATACAATGGAAAATCCACCGAACAAAGTCCTCTTATACAAAATTGCCAAGCGGCCTACCGAG  
consensus sequence Untitled Consensus

ATCGCAGAACCTCTTAAGCGGGCTAAAGTGTCCAATACAATGGAAAATCCACCGAACAAAGTCCTCTTATACAAAATTGCCAAGCGGCCTACCGAG  
aligned sequence S. cerevisiae MUD1

ATCGCAGAACCTCTTAAGCGGGCTAAAGTGTCCAATACAAGGAAAATACACCGAACAAAGGTCTTATACAAAATTGCCCTAGCGGCCTACCGAG  
modified\_MUD1  
aligned sequence modified MUD1

CAATTATTGTCGCAAATACTTGGCAATGAGGCTTAGTTGAAATCAGATTAGTTAGCGTTCGTAAACCTAGCTTCGTGGAATACGAGACCGTTGCTGATG  
consensus sequence Untitled Consensus

CAATTATTGTCGCAAATACTTGGCAATGAGGCTTAGTTGAAATCAGATTAGTTAGCGTTCGTAAACCTAGCTTCGTGGAATACGAGACCGTTGCTGATG  
aligned sequence S. cerevisiae MUD1

CAATTATCCTCGCAAATACTTGGCAAGGAGGTTTAGTTGAAATCAGTTAGTTAGCGTTCGTACCTAGCTTCGTGGAATACGAGACCGTTGCTGTTG  
modified\_MUD1  
aligned sequence modified MUD1

CTACGAAAATCAAGAACAGTTAGGCTCCACTTACAAGCTACAAAACAATGACGTTACCATAGGATTGCTAAGTAGAATTCCCTTGCGGAAGTATAACC  
consensus sequence Untitled Consensus

CTACGAAAATCAAGAACAGTTAGGCTCCACTTACAAGCTACAAAACAATGACGTTACCATAGGATTGCTAAGTAGAATTCCCTTGCGGAAGTATAACC  
aligned sequence S. cerevisiae MUD1

CTACGAAAATCAAGAACAGTTAGGCTCCACTTACAAGCTACAAAACAATAACGTTACCATAGGATTGCTAAGTAGAATTCCCTTGCGGAAGTATAACC  
modified\_MUD1  
aligned sequence modified MUD1

TCGAGTAAAGAAATTACAGATAAATTGAATAACGTTCTCCATTATGAATAAAATATGTATTACAAGCTTAGAAAATAATGTGCTTTATTAC  
consensus sequence Untitled Consensus

TCGAGTAAAGAAATTACAGATAAATTGAATAACGTTCTCCATTATGAATAAAATATGTATTACAAGCTTAGAAAATAATGTGCTTTATTAC  
aligned sequence S. cerevisiae MUD1

TCGAGTAAAGAAATTACAGATAAATTGAATAACGTTCTCCATTATTAAATAAAATATTATTACAAGCTTAGAAAATAAGGTGCTTTATTAC  
modified\_MUD1  
aligned sequence modified MUD1