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Near-Consensus Position TTGACAAAACGGTTAAATTTATG<mark>TATAAT</mark>G Change = -0.015Sequence A TTGACAAACGGTTAAATTTATGTACAATG < Consensus Position TTGACAAACGGTTAAATTT<mark>TATAAT</mark>AAAG Change = -0.005Consensus Position TGACAAACCGGTTTATTCTATACAAT Change = -0.052Sequence B TGACAAACCGGTTTATTCTATATAATAAAT Near-Consensus Position TGACAAACCGGTTTATTCTAATGTACAATT Change = -0.131