

Majority	10	20	30	40	50	60	70	80	90	100	110	120	130	140	150	160
Md 1.03A-MDP00029542.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.03B-MDP00029931.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.03E-MDP00029543.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.03J-MDP000081519.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.03F-MDP0000533838.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.08-MDP0000174018.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.09-MDP0000260110.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.03C-MDP0000611199.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.03H-MDP0000418277.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.03D-MDP000081519.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.06D-MDP0000288293.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.06A-MDP0000399350.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.14-MDP0000195610.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.01-MDP0000103621.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.06B-MDP0000216907.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.13A-MDP0000611200.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.13C-MDP0000750556.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.05-MDP0000149794.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.02-MDP00000942516.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.12-MDP0000199487.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.11B-MDP0000203197.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											
Md 1.11A-MDP0000789174.seq	AT TGGG TTTT TCACATACGAA	CCGAGTTCACCTCGGTCAT	CCCCCTCTAGTGTGTT	CAATGCCTTT	TTTCTTGATGC											

Decoration Decoration #: Shade (with solid black) residues that match the Consensus exactly.