

		20		40		60	
nad9GM10DNA new	ATGCTCTGTA	TAATACTTTT	CCCCGAGCGA	TGGTTTAGCG	TATTCGGAAT	TGTACCCAAG	60
nad9GM10control	60
nad9GM10 2H	60
nad9GM10 12H	60
		80		100		120	
nad9GM10DNA new	CCCTCCCATG	GGTTCATATC	CCGATTCATA	ACTAGAGCAT	GCAGCCGATC	CTGGATACAT	120
nad9GM10control	120
nad9GM10 2H	120
nad9GM10 12H	120
		140		160		180	
nad9GM10DNA new	AACTCTAAAA	AGTGTGTGTG	CAGTTTTGGA	TCTTTATTGG	TAGCCAGTCT	TTCACCTTCTG	180
nad9GM10control	180
nad9GM10 2HT.....	180
nad9GM10 12HT.....	180
		200		220		240	
nad9GM10DNA new	CCTCTCCACT	CCCATGCCTT	TCTTGGTCGG	ACCAACCCAA	CCGGCGATTT	CCGACAAGTC	240
nad9GM10controlT.....	240
nad9GM10 2HT.....	240
nad9GM10 12HT.....	240
		260		280		300	
nad9GM10DNA new	TTTCTGCTTA	GAGCAAGAAG	CGGAACCAAA	ATAAAGCTTT	CTTTATTTTC	ATTTATGGAT	300
nad9GM10control	300
nad9GM10 2H	300
nad9GM10 12H	300
		320		340		360	
nad9GM10DNA new	AACCAATCCA	TTTTCCAATA	TAGTTGGGAG	ATTTTACCCA	AGAAATGGGT	ACATAAAATG	360
nad9GM10controlT.....	360
nad9GM10 2HT.....	360
nad9GM10 12HT.....	360
		380		400		420	
nad9GM10DNA new	AAAAGATCGG	AACATGGGAA	TAGATCTTAT	ACCAATACTG	ACTACCCATT	TCCATTGTTG	420
nad9GM10controlT.....T.T.....	420
nad9GM10 2HT.....T.T.....	420
nad9GM10 12HT.....T.T.....	420
		440		460		480	
nad9GM10DNA new	TGCTTTCTAA	AATGGCATAc	CTATACAAGG	GTTCAAGTTT	CGATCGATAT	TTGCGGAGTG	480
nad9GM10control	T.....	480
nad9GM10 2H	T.....	480
nad9GM10 12H	T.....	480
		500		520		540	
nad9GM10DNA new	GATCATCCCT	CTCGAAAACG	AAGATTTGAA	GTTGTCCATA	ATTTACTGAG	TACTCGGTAT	540
nad9GM10controlT.....T.....	540
nad9GM10 2HT.....T.....	540
nad9GM10 12HT.....T.....	540
		560		580		600	
nad9GM10DNA new	AACTCACGCA	TTCGTGTACA	AACAAGTGCA	GACGAAGTAA	CACGAATATC	TCCGGTAGTC	600
nad9GM10controlT.....	600
nad9GM10 2HT.....	600
nad9GM10 12HT.....	600
		620		640		660	
nad9GM10DNA new	AGTCTATTTT	CATCAGCCGG	CCGGTGGGAG	CGAGAAGTAT	GGGATATGTC	TGGTGTITCT	660
nad9GM10controlT.....T.....	660
nad9GM10 2HT.....T.....	660
nad9GM10 12HT.....T.....	660
		680		700		720	
nad9GM10DNA new	TCCATCAATC	ATCCGGATTT	ACGCCGTATA	TCAACAGATT	ATGGTTTCGA	GGGTCATCCA	720
nad9GM10controlT.....	720
nad9GM10 2HT.....T.....	720
nad9GM10 12HT.....T.....	720
		740		760		780	
nad9GM10DNA new	TTACGAAAAG	ACTTTCCTCT	GAGTGGATAT	GTGGAAGTAC	GCTATGATGA	TCCAGAGAAA	780
nad9GM10control	780
nad9GM10 2H	780
nad9GM10 12HT.....	780
		800		820		840	
nad9GM10DNA new	CGTGTGGTTT	CTGAACCCAT	TGAGATGACC	CAAGAATTTT	GCTATTTCGA	TTTTGCTAGT	840
nad9GM10controlA.....T.....A.....	840
nad9GM10 2HA.....A.....	840
nad9GM10 12HA.....A.....	840
		860					
nad9GM10DNA new	CCTTGGGAAC	AGCGTAGCGA	CGGATAA	867			
nad9GM10controlC.....	T.....	867			
nad9GM10 2HC.....	T.....	867			
nad9GM10 12HC.....	T.....	867			