	001	CGGGIGIIGG	CGGGIGICGG	GGCIGGCIIA	ACIAIGCGGC	AICAGAGCAG
	701	ATTGTACTGA	GAGTGCACCA	TATGCGGTGT	GAAATACCGC	ACAGATGCGT
	751	AAGG <u>AGAAAA</u>	TACCGCATCA	GGCGCCATTC	GCCATTCAGG	CTGCGCAACT
	801	GTTGGGAAGG	GCGATCGGTG	CGGGCCTCTT	CGCTATTACG	CCAGCTGGCG
	851	AAAGGGGGAT	GTGCTGCAAG	GCGATTAAGT	TGGGTAACGC	CAGGGTTTTC
p2RT:	901	CCAGTCACGA	CGTTGTAAAA	CGACGGCCAG	TGAATT <mark>CGAG</mark>	CTCGGTACCC
p2RT-CG14:					CCGC	GCGCGCGCGC
p2RT-MYC:					GCGG	GGAGGGGCGC
p2RT:	951	GGGGATCCTC	TAGAGTCGAC	CTGCAGGCAT	GCAAGCTTGG	CGTAATCATG
p2RT-CG14:		GCGCGCGCGC	GCGCG			
p2RT-MYC:		TTATGGGGAG	GGTTG			
-	1001	GTCATAGCTG	TTTCCTGTGT	GAAATTGTTA	TCCGCTCACA	ATTCCACACA
	1051	ACATACGAGC	CGGAAGCATA	AAGTGTAAAG	CCTGGGGTGC	CTAATGAGTG
	1101	AGCTAACTCA	CATTAATTGC	GTTGCGCTCA	CTGCCCGCTT	TCCAGTCGGG
	1151	AAACCTGTCG	TGCCAGCTGC	ATTAATGAAT	CGGCCAACGC	GCGGGGAGAG
	1201	GCGGTTTGCG	TATTGGGCGC	TCTTCCGCTT	CCTCGCTCAC	TGACTCGCTG
	1251	CGCTCGGTCG	TTCGGCTGCG	GCGAGCGGTA	TCAGCTCACT	CAAAGGCGGT
	1301	AATACGGTTA	TCCACAGAAT	CAGGGGATAA	CGCAGGAAAG	AACATGTTGT
	1351	GACACGATGC	AGCTTCAGGA	TCGGATCCGG	CTGTGGAATG	TGTGTCAGTT
	1401	AGGGTGTGGA	AAGTCCCCAG	GCTCCCCAGC	AGGCAGAAGT	ATGCAAAGCA
	1451	TGCATCTCAA	TTAGTCAGCA	ACCAGGTGTG	GAAAGTCCCC	AGGCTCCCCA
	1501	GCAGGCAGAA	GTATGCAAAG	CATGCATCTC	AATTAGTCAG	CAACCATAGT
	1551	CCCGCCCCTA	ACTCCGCCCA	TCC <i>C</i> GCCCCT	AACTCCGCCC	AGTTCCGCCC
	1601	ATTCTCCGCC	CCATGGCTGA	CTAATTTTTT	TTATTTATGC	AGAGGCCGAG
	1651	GCCGCCTCGG	CCTCTGAGCT	ATTCCAGAAG	TAGTGAGGAG	GCTTTTTTGG
	1701	AGGCCTAGGC	TTTTGCAAAA	AGCTTCACGC	TGCCGCAAGC	ACTCAGGGCG

601TTGTCTGTAA GCGGATGCCG GGAGCAGACA AGCCCGTCAG GGCGCGTCAG651CGGGTGTTGG CGGGTGTCGG GGCTGGCTTA ACTATGCGGC ATCAGAGCAG

Supplementary Figure 1