			Во	GH-rev						
ATCAAGATGCTATAATG	GAGTTTAAACO 24	CC <mark>G</mark> CTGATCA 36	GCCTCGACT	rgtgccttc 48	TAGTTGC 60	bGH CA <mark>G</mark> CCA <mark>TCT</mark>	poly(A) signal GTTGTTTGC 72	ccc <mark>T</mark> cccc <mark>G</mark> 84	TGCCTTCCT:	TGA
CCCTGGAAGGTGCCACT	ı		AAAA <mark>TG</mark> A <mark>G</mark> O	oGH poly(A) si GAAA <mark>TTG</mark> CA	TCGCATT		GGTGTCATI	CTATTCTGGG	T	GGG
108 GCAGGACAGCAAGGGGG	120	bGH poly(A)			156 CGCTGGG	168	TTCTGAGG	180 C <mark>GG</mark> AAA <mark>G</mark> AACC	192	TCG
204 216		228	240	252 M13 rev		264	276	288		300
			M1:	3 Reverse	M13/p	OUC Reverse				
ATACCGTCGACCTCTAG	CTAGAGCTT 324	<mark>GGCGT</mark> AA <mark>T</mark> CA 336	6	348	T <mark>GTGTG</mark> AA 360		CC <mark>G</mark> C <mark>T</mark> CACA 372	AA <mark>TT</mark> CCACACA 384	ACA <mark>T</mark> AC <mark>G</mark> AG(396	CC <mark>G</mark>
lac promoter GAAGCATAAAGTGTAAA 408	A <mark>GCCTAGGGT</mark> O 420	GCCTAATGAG 432	CAP binding TGAGCTAAC 444	CTCACA <mark>TT</mark> A.	A <mark>TT</mark> GCGT	<mark>rgcg</mark> c <mark>t</mark> cac 468	<mark>TG</mark> CCC <mark>G</mark> C <mark>T1</mark>	TCCAGTCGGG 480	AAACC <mark>TGT</mark> C0	GTG
CCAGCTGCATTAATGAA	\ <mark>T</mark> C <mark>GG</mark> CCAAC(GCGCGGGGAG	A <mark>GG</mark> C <mark>GGTT1</mark>	GCGTATTG(GGCGCTC	<mark>TTCCG</mark> CTTC		440 C <mark>TG</mark> AC <mark>T</mark> CGC <mark>TG</mark>	C <mark>G</mark> C <mark>T</mark> C <mark>GGT</mark> C	GTT
504 516		528	540	552		564	576	588		600
CGGCTGCGGCGAGCGGT	TATCAGCTCAC 624	C <mark>T</mark> CAAA <mark>GG</mark> C <mark>G</mark> 636		648	AGAA <mark>T</mark> CAC)	GCAGGAAAC 672 ori	GAACA <mark>TGTGAG</mark> 684	CAAAA <mark>GG</mark> CC/ 696	A <mark>G</mark> C
AAAA <mark>GG</mark> CCA <mark>GG</mark> AACC <mark>GT</mark> 708	AAAAA <mark>GG</mark> CC <mark>(</mark> 720	GCGTTGCTGG 732	CGTTTTTCC 744		2 <mark>6</mark> 00000 756			AA <mark>TCGACGCT</mark> C 780	AA <mark>GT</mark> CA <mark>G</mark> AG0 792	GTG
		pBR322ori-F		ori						
GCGAAACCCGACAGGAC 804 816		ACCA <mark>GG</mark> C <mark>GTT</mark> 828	TCCCCCTGG 840	852	TCGTGCG	864	TCC <mark>G</mark> ACCC 876	GCCGCTTACC 888		900
GCCTTTCTCCCTTCGGG	GAAGCGTGGCC 924	GCTTTCTCAT 936		ori CTGTAGGTA 948	TCTCAGT		<mark>GTCGTTCG</mark> (972	TCCAA <mark>GCTGG</mark> 984	GCTGTGTGCA 996	AC <mark>G</mark>
AACCCCCGTTCAGCCC					ı	ı			ı	TGG
1008 TAACAGGATTAGCAGAG	1020	1032	TACAGAGTI	ori	1056	1068		1080	1092	Tee
1104 1116		1128	1140	1152		1164	1176	118		1200
GCTCTGCTGAAGCCAGT	TACCTTCGG/ 1224	AAAAG <mark>AGTT</mark> 123			CAAACAA/ 126		GGTAGCGG1 1272	GGTTTTTTG 1284	TTTGCAAGC 1296	
Ori AGATTACGCGCAGAAAA 1308	AAA <mark>GG</mark> ATCT(1320	CAA <mark>G</mark> AA <mark>G</mark> A <mark>T</mark> C 1332	CTTTGATC1 1344		<mark>GGGTCTG</mark> / 1356	AC <mark>G</mark> C <mark>T</mark> CA <mark>GT</mark> 1368		AAC <mark>T</mark> CAC <mark>GTT</mark> A 1380	A <mark>GGG</mark> ATTTT(1392	GGT
1300	1320	1332	1344	•	1330	1300		1360	1392	AmpR
CA <mark>TG</mark> AGATTATCAAAAA 1404 1416		CC <mark>T</mark> A <mark>G</mark> A <mark>TCCT</mark> 1428	TTTAAATTA 1 1440	AAAAA <mark>TG</mark> AA 1 1452	GTTTTAAA	A <mark>TCAA</mark> TCTA 1464	AA <mark>GT</mark> A <mark>T</mark> A <mark>T</mark> A 1476	A <mark>TGAGT</mark> AAAC <mark>T</mark> 148		AGT 1500
TACCAATGCTTAATCAG								CGTGTAGATAA		
1512 AGGGCTTACCATCTGGC	1524	153		AmpR	156		1572	1584	1596	
1608	1620	1632	1644		1656	1668		1680	1692	
GCGCAGAAGTGGTCCTG		CC <mark>G</mark> CC <mark>T</mark> CCA <mark>T</mark> 1728	CCA <mark>GTCTA</mark> 1 1740	TAA <mark>TTGTT</mark> 1752		A <mark>GCT</mark> AGAGT 1764	AA <mark>GT</mark> A <mark>GTT</mark> 0 1776	C <mark>G</mark> CCA <mark>GTT</mark> AA <mark>T</mark> 178		1800
GTTGTTGCCATTGCTAC	CA <mark>GG</mark> CA <mark>T</mark> C <mark>GT</mark> C	GGTGTCACGC 183		AmpR GGTATGGC 1848	TTCATTC/ 186		TCCCAACGA	TCAA <mark>GG</mark> CGAG 1884	TTACA <mark>TG</mark> AT(1896	
CCA <mark>TGTTG</mark> TGCAAAAA	(<mark>GCGGTT</mark> AGC	TCCTTC <mark>GGT</mark> C	C <mark>TCCG</mark> A <mark>TC</mark> C	AmpR STTGTCAGA	A <mark>GT</mark> AAGT	<mark>rgg</mark> cc <mark>g</mark> ca <mark>g</mark>	TGTTATCAC	O <mark>TCATGGTTAT</mark>	<mark>GG</mark> CA <mark>G</mark> CAC <mark>T</mark> (G C A
1908	1920	1932	1944	4 AmpR	1956	1968		1980	1992	
TAATTCTCTTACTGTCA 2004 2016		TAAGATGCTT 2028	2040	2052	AC <mark>T</mark> CAAC(2064	C <mark>TG</mark> AGAA <mark>T</mark> A 2076	A <mark>GTGTATGCGG</mark> 208		2100
TCTTGCCCGGCGTCAAT	AC <mark>GGG</mark> A <mark>T</mark> AA	Amp-R TACC <mark>G</mark> CGCCA	CA <mark>T</mark> AGCAGA	AmpR	A <mark>GTGCT</mark> Ç/	A <mark>TCATTGG</mark> A	AAAC <mark>GTT</mark> C1	TCGGGGCGAA	AAC <mark>T</mark> C <mark>T</mark> CAA(<mark>GG</mark> A
2112	2124	213		2148 AmpR	216		2172	2184	2196	
TCTTACCGCTGTTGAGA 2208	ATCCAGTTCGA 2220	A <mark>TGT</mark> AACCCA 2232 AmpR	CTCGTGCAC 2244		TCTTCAGO 2256	2268		AGCGTTTCTGG 2280 DR promoter	GTGAGCAAA 2292	AAC
AGGAAGGCAAAATGCCG 2304 2316			GACAC <mark>GG</mark> AA 2340	AA <mark>TGTTG</mark> AA 2352	TACTCATA	2364		ATTATTGAAGC 238		GG T 2400
TATTGTCTCATGAGCGG		T	TAGAAAAA <mark>1</mark>		T		ı			
2412 GATCGGGAGATCGATCT	2424 CCC <mark>G</mark> A <mark>T</mark> CCC	243 C <mark>T</mark> A <mark>GGGT</mark> CGA	<	2448 pRS-m			2472 Taagccag1	2484	2496	
2508	2520	2532	2544		2556	2568		2580	2592	
GGAGGTCGCTGAGTAGT 2604 2616		AAA <mark>TTT</mark> AA <mark>G</mark> C 2628	TACAACAAC 2640	GGCAA <mark>GG</mark> C <mark>T</mark> 2652	T <mark>G</mark> A C C <mark>G</mark> A (2664	2676	GCTTAGGGTT 268		<mark>GCG</mark> 2700
CTGCTTCGCGATGTACG	GGGCCAGA <mark>T</mark> A 2724	TACGCGTTGA 273		ATTGACTAG 2748	TTATTAA 276			OTCATTAGTTC 2784	A <mark>T</mark> AGCCCA <mark>T</mark> 2796	
TGGAGTTCCGCGTTACA				CMV enhance	er					
2808	2820	2832	2844	CMV enhand		2868		2880	2892	
AACGCCAATAGGGACTT 2904 2916		<mark>GTCAATGGGT</mark> 2928	<mark>GG</mark> A <mark>GTATT1</mark> 2940	2952		CTTGGCAGT 2964	ACA <mark>T</mark> CAA <mark>G</mark> 1 2976	GTATCATATG - 298		3000
CCTATTGACGTCAATGA	AC <mark>GGT</mark> AAA <mark>TG0</mark> 3024	GCCCGCC <mark>TGG</mark>		CMV enhand			CCTACTTGC 3072	CAGTACATCT 3084	AC <mark>GT</mark> ATTAG 3096	
CMV enhancer TCGCTATTACCATGGTG	ATGCGGTTT			GTGGATAG	I	AC <mark>T</mark> CAC <mark>GGG</mark>		AGTCTCCACCC	1	CAA
3108	3120	3132	3144	4 CMV promot	3156 er	3168		3180	3192	
TGGGAGTTTGTTTTGGC 3204 3216		AAC <mark>GGG</mark> AC <mark>TT</mark> 3228	TCCAAAATO 3240	GTCGTAACA 3252	AC <mark>T</mark> CC <mark>G</mark> C(CCCA <mark>TTG</mark> AC 3264		MV-F GCGGTAGGCGT 328		GAG 3300
CMV promoter	>					T7 T7 promoter	>			
GTCTATATAAGCAGAGC	C <mark>TGGTTTAGT</mark> C 3324	<mark>G</mark> AACC <mark>GT</mark> CA <mark>G</mark> 333		A <mark>G</mark> A <mark>T</mark> CC <mark>G</mark> C 1 3348	GGCCGCT/ 336		CAC <mark>T</mark> A <mark>T</mark> AGO 3372	GGAGAGCCGCC 3384	ACCA <mark>TGG</mark> CA(3396	
AA <mark>G</mark>										