

# pRL-CMV Vector



Technical Bulletin No. 237

INSTRUCTIONS FOR USE OF PRODUCT E2261. PLEASE DISCARD PREVIOUS VERSIONS.

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## I. Description

The pRL-CMV Vector<sup>(a,b,c)</sup> (Figure 1) is intended for use as an internal control reporter and may be used in combination with any experimental reporter vector to co-transfect mammalian cells. All of Promega's pRL Reporter Vectors contain a cDNA (*Rluc*)<sup>(a)</sup> encoding *Renilla* luciferase, which was originally cloned from the marine organism *Renilla reniformis* (sea pansy; 1). As described below, the *Renilla* luciferase cDNA contained within the pRL Vectors has been modified slightly to provide greater utility.

The pRL-CMV Vector contains the CMV<sup>(b)</sup> enhancer and early promoter elements to provide high-level expression of *Renilla* luciferase in co-transfected mammalian cells. *Renilla* luciferase is a 36kDa monomeric protein that does not require post-translational modification for activity (2). Therefore, like firefly luciferase, the enzyme may function as a genetic reporter immediately following translation. For information about the use of this plasmid in conjunction with a reporter vector containing the firefly luciferase gene, refer to the *Dual-Luciferase® Reporter Assay System*<sup>(c,d,e)</sup> Technical Manual (#TM040).

To avoid DNA methylation, all pRL Vectors are isolated from a *dam-/dcm-* *E. coli* K host strain. Therefore, initial propagation of the pRL-CMV Vector should be conducted in an appropriate *E. coli* host strain lacking endogenous restriction endonuclease activity (e.g., JM109).



## II. Product Components

Product	Size	Cat.#
pRL-CMV Vector	20µg	E2261

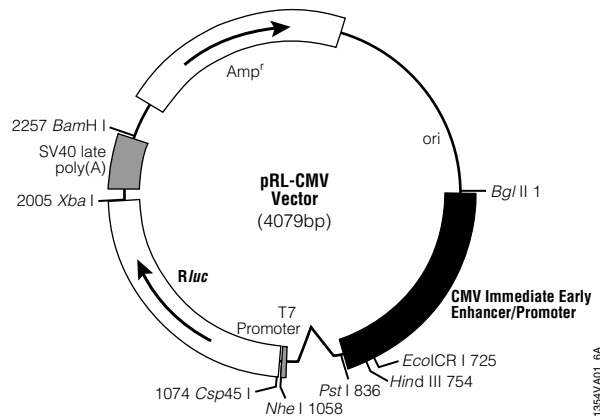
All pRL Vectors are supplied in TE buffer (pH 7.4) and are provided with a glycerol stock of bacterial strain JM109.

**Storage Conditions:** Store vector DNA at  $-20^{\circ}\text{C}$ . Store glycerol stock of JM109 at  $-70^{\circ}\text{C}$ .

## III. Features of the pRL-CMV Vector

### A. CMV Enhancer/Promoter Regions

The pRL-CMV Vector contains the CMV immediate-early enhancer/promoter region, which provides strong, constitutive expression of the *Renilla* luciferase cDNA in a variety of cell types. The promiscuous nature of the CMV enhancer/promoter has been demonstrated in transgenic mice, where its transcriptional activity was observed in 24 of the 28 murine tissues examined (3).



### pRL-CMV Vector circle map and sequence reference points:

CMV enhancer and immediate early promoter	7–803
Chimeric intron	860–996
T7 promoter (–17 to +2)	1040–1058
T7 promoter transcription start site	1057
<i>RLuc</i> reporter gene	1068–2003
SV40 late polyadenylation signal	2045–2246
$\beta$ -lactamase ( <i>Amp<sup>r</sup></i> ) coding region	2393–3253

**Note:** —, position of intron; *RLuc*, cDNA encoding the *Renilla* luciferase enzyme; *Amp<sup>r</sup>*, gene conferring ampicillin resistance in *E. coli*; ori, origin of plasmid replication in *E. coli*. Arrows within the *RLuc* and *Amp<sup>r</sup>* gene indicate the direction of transcription.

## B. Chimeric Intron

Downstream of the CMV enhancer/promoter region of the pRL-CMV Vector is a chimeric intron comprised of the 5'-donor splice site from the first intron of the human  $\beta$ -globin gene, and the branch and 3'-acceptor splice site from an intron preceding an immunoglobulin gene heavy chain variable region (4). The sequences of the donor and acceptor splice sites, along with the branchpoint site, have been modified to match the consensus sequences for optimal splicing (5).

Transfection studies have demonstrated that the presence of an intron flanking a cDNA insert frequently increases the level of gene expression (6–9). In the pRL-CMV Vector, the intron is positioned 5' to *Rluc* to minimize the utilization of cryptic 5'-donor splice sites that may reside within the reporter gene sequence (10).

## C. T7 Promoter

A T7 promoter is located downstream of the chimeric intron, and immediately precedes the *Rluc* reporter gene. This T7 promoter can be used to synthesize RNA transcripts in vitro using T7 RNA Polymerase (Cat.# P2075). T7 RNA Polymerase can also be used to synthesize active *Renilla* luciferase in a cell-free coupled eukaryotic in vitro transcription/translation reaction (e.g., Promega's TNT<sup>®</sup> Reticulocyte Lysate<sup>(c,d,f,g)</sup> [Cat.# L4610] or Wheat Germ Extract [Cat.# L4140] Systems).

## D. *Renilla* Luciferase Reporter Gene (*Rluc*)

The *Renilla* luciferase cDNA inserted into all of the pRL Vectors is derived from the anthozoan coelenterate *Renilla reniformis* (1) but contains nucleotide changes that were engineered during the construction of the individual vectors. The following bases were altered in the pRL-CMV Vector: base 1298 (T→C), to eliminate an internal *Bgl* II site; base 1841 (T→C), to eliminate an internal *Bam* H I site; base 1874 (C→T), to eliminate internal *Nar* I, *Kas* I, *Ban* I and *Acy* I sites. These nucleotide substitutions do not alter the amino acid sequence of the encoded *Renilla* luciferase reporter enzyme.

## E. SV40 Late Polyadenylation Signal

Polyadenylation signals cause the termination of transcription by RNA polymerase II and signal the addition of approximately 200–250 adenosine residues to the 3'-end of the RNA transcript (11). Polyadenylation has been shown to enhance RNA stability and translation (12,13). The late SV40 polyadenylation signal, which is extremely efficient and has been shown to increase the steady-state level of RNA approximately 5-fold more than the early SV40 polyadenylation signal (14), has been positioned 3' to the *Rluc* gene in the pRL-CMV Vector to increase the level of *Renilla* luciferase expression.

**Note:** The T7 Promoter Primer offered by Promega (Cat.# Q5021) cannot be used for sequencing this vector because of a mismatch between the 3' end of the primer and the vector DNA.

#### IV. Transfection of Mammalian Cells with pRL-CMV Vector

The pRL-CMV Vector may be used in combination with any experimental reporter vector to co-transfect mammalian cells. However, it is important to realize that *trans* effects between promoters on co-transfected plasmids can potentially affect reporter gene expression (15). Primarily, this is of concern when either the control or experimental reporter vector, or both, contain very strong promoter/enhancer elements (such as CMV). The occurrence and magnitude of such effects will depend on several factors: a) the combination and activities of the genetic regulatory elements present on the co-transfected vectors, b) the relative ratio of experimental vector to control vector introduced into the cells, and c) the cell type transfected.

To help ensure independent genetic expression between experimental and control reporter genes, preliminary co-transfection experiments should be performed to optimize both the **amount** of vector DNA and the **ratio** of the co-reporter vectors added to the transfection mixture. Similar to the firefly luciferase assay, the *Renilla* luciferase assay is extremely sensitive, providing accurate measurement of  $\leq 10$  femtograms of *Renilla* luciferase, with linearity over 7 orders of enzyme concentration. Therefore, it is possible to use relatively small quantities of pRL-CMV Vector to provide low-level, constitutive co-expression of *Renilla* luciferase control activity. Ratios of 10:1 to 50:1 (or greater) for experimental vector:pRL-CMV Vector combinations are feasible and may aid greatly in suppressing the occurrence of *trans* effects between promoter elements.

The pRL-CMV Vector can be used for both transient and stable expression of genes. For stable expression, the pRL-CMV Vector must be co-transfected with an expression vector containing a selectable gene in mammalian cells. Transfection of DNA into mammalian cells may be mediated by cationic lipids (16,17), calcium phosphate (18,19), DEAE-Dextran (20–22), polybrene-DMSO (23,24), or electroporation (25,26).

Transfection systems based on cationic lipid compounds (Transfectam<sup>®</sup> Reagent<sup>(h)</sup>, TransFast<sup>™</sup> Transfection Reagent<sup>(i)</sup> and Tfx<sup>™</sup>-10, Tfx<sup>™</sup>-20 and Tfx<sup>™</sup>-50 Reagents<sup>(j)</sup>), Calcium Phosphate and DEAE-Dextran are available from Promega. For more information and a protocol for the Transfectam<sup>®</sup> Reagent, please request the *Transfectam<sup>®</sup> Reagent for the Transfection of Eukaryotic Cells Technical Bulletin* (#TB116). Information about the TransFast<sup>™</sup> Transfection Reagent can be found in the *TransFast<sup>™</sup> Transfection Reagent Technical Bulletin* (#TB260). Protocols for the use of the Tfx<sup>™</sup> Reagents can be found in the *Tfx<sup>™</sup>-10, Tfx<sup>™</sup>-20 and Tfx<sup>™</sup>-50 Reagents for the Transfection of Eukaryotic Cells Technical Bulletin* (#TB216). For transfection procedures using calcium phosphate or DEAE-Dextran, please request the *ProFection<sup>®</sup> Mammalian Transfection Systems Technical Manual* (#TM012).

**Note:** All of Promega's Technical Bulletins and Technical Manuals are available online at [www.promega.com](http://www.promega.com).

**Note:** For assistance in determining transfection conditions for different cell lines, Promega offers the Transfection Assistant available online at [www.promega.com/transfectionasst/](http://www.promega.com/transfectionasst/).

## V. pRL-CMV Vector Restriction Sites and Vector Sequence

### A. pRL-CMV Vector Restriction Sites

The following restriction enzyme tables were constructed using DNASTAR® sequence analysis software. Please note that we have not verified this information by restriction digestion with each enzyme listed. The location given specifies the 3' end of the cut DNA (the base to the left of the cut site). For more information on the cut sites of these enzymes, or if you identify a discrepancy, please contact your local Promega Branch Office or Distributor. In the U.S., contact Promega Technical Services at 800-356-9526. Vector sequences are also available in the GenBank® database (GenBank®/EMBL Accession Number AF025843) and on the Internet at [www.promega.com/vectors/](http://www.promega.com/vectors/).

**Table 1. Restriction Enzymes That Cut the pRL-CMV Vector Between 1 and 5 Times.**

Enzyme	# of Sites	Location	Enzyme	# of Sites	Location
<b>Aat II</b>	4	284, 337, 420, 606	<b>BstZ I</b>	1	2012
<b>Acy I</b>	5	281, 334, 417, 603 2640	<b>Cla I</b>	1	2250
<i>Afl II</i>	2	826, 1023	<b>Csp45 I</b>	1	1074
<i>Afl III</i>	1	1250	<b>Dde I</b>	4	2679, 3219, 3385, 3794
<b>Alw44 I</b>	2	2508, 3754	<b>Dra I</b>	4	2216, 2602, 3294, 3313
<i>AlwN I</i>	1	3659	<i>Drd I</i>	2	815, 3966
<i>AspH I</i>	4	727, 2512, 2597, 3758	<i>Dsa I</i>	1	519
<b>Ava II</b>	4	1116, 1838, 2816, 3038	<i>Eae I</i>	5	14, 68, 1418, 2012 2787
<b>Bal I</b>	2	16, 70	<i>Eag I</i>	1	2012
<b>BamH I</b>	1	2257	<i>Ear I</i>	2	1238, 2381
<b>Ban I</b>	4	624, 949, 1872, 3227	<b>EclHK I</b>	1	3180
<b>Ban II</b>	1	727	<b>Eco52 I</b>	1	2012
<i>Bbs I</i>	2	934, 1908	<b>EcoI CR I</b>	1	725
<b>Bcl I</b>	2	1352, 1561	<i>Fsp I</i>	1	2957
<b>Bgl I</b>	5	142, 249, 371, 442 3062	<b>Hae II</b>	1	3828
<b>Bgl II</b>	1	1	<i>Hga I</i>	4	685, 2648, 3378, 3956
<i>Bsa I</i>	2	888, 3114	<b>Hinc II</b>	2	675, 2155
<b>BsaO I</b>	4	2015, 2662, 2811, 3734	<i>Hind II</i>	2	675, 2155
<i>BsaA I</i>	2	499, 1800	<b>Hind III</b>	1	754
<i>BsaB I</i>	1	2256	<b>Hpa I</b>	1	2155
<i>BsaH I</i>	5	281, 334, 417, 603 2640	<b>Hsp92 I</b>	5	281, 334, 417, 603 2640
<i>BsaJ I</i>	3	519, 1841, 3908	<b>MspA1 I</b>	3	2544, 3485, 3730
<b>BsaM I</b>	2	2076, 2169	<b>Nci I</b>	3	2644, 2995, 3691
<i>Bsm I</i>	2	2076, 2169	<b>Nco I</b>	1	519
<b>Bsp1286 I</b>	4	727, 2512, 2597, 3758	<b>Nde I</b>	1	393
<i>BspH I</i>	3	1636, 2340, 3348	<b>Nhe I</b>	1	1058
<i>BspM I</i>	1	850	<b>Not I</b>	1	2012
<b>BsrBR I</b>	1	2256	<i>Nsp I</i>	2	1194, 1254
<i>BsrG I</i>	2	102, 1766	<i>PaeR7</i>	2	916, 4074
<i>BssS I</i>	3	1726, 2511, 3895	<i>Ple I</i>	5	563, 924, 1040, 3189, 3692
<b>Bst98 I</b>	2	826, 1023	<b>Pst I</b>	1	836
			<b>Pvu I</b>	1	2811
			<b>Sac I</b>	1	727

(continued)

**Note:** The enzymes listed in boldface type are available from Promega.

**Table 1. Restriction Enzymes That Cut the pRL-CMV Vector Between 1 and 5 Times (continued).**

Enzyme	# of Sites	Location	Enzyme	# of Sites	Location
<b>Sca I</b>	2	1036, 2699	<b>Tfi I</b>	5	1158, 1179, 1518, 1833, 1846
<b>Sin I</b>	4	1116, 1838, 2816, 3038	<b>Vsp I</b>	3	166, 1168, 3005
<b>SnaB I</b>	1	499	<b>Xba I</b>	1	2005
<b>Spe I</b>	1	158	<b>Xcm I</b>	1	1717
<b>Ssp I</b>	3	11, 58, 2375	<b>Xmn I</b>	2	1602, 2580
<b>Sty I</b>	1	519			

**Table 2. Restriction Enzymes That Do Not Cut the pRL-CMV Vector.**

<b>AccB7 I</b>	<b>Bbu I</b>	<b>BssH II</b>	<b>EcoR I</b>	<b>Nru I</b>	<b>Sgf I</b> <sup>(k)</sup>
<b>Acc I</b>	<b>BfrB I</b>	<b>Bst1107 I</b>	<b>EcoR V</b>	<b>Nsi I</b>	<b>SgrA I</b>
<b>Acc III</b>	<b>Bln I</b>	<b>Bst29 I</b>	<b>EcoR124 I</b>	<b>Pac I</b>	<b>Sma I</b>
<b>Acc65 I</b>	<b>Blp I</b>	<b>BstE II</b>	<b>EcoR124 II</b>	<b>Pau I</b>	<b>Sna I</b>
<b>Acr I</b>	<b>Bmg I</b>	<b>BstX I</b>	<b>Ecoprr I</b>	<b>PfIM I</b>	<b>Sph I</b>
<b>Acs1371 I</b>	<b>Bpl I</b>	<b>Bsu36 I</b>	<b>Ehe I</b>	<b>Pfu I</b>	<b>SpI I</b>
<b>Afa24R I</b>	<b>Bpu1102 I</b>	<b>Csp I</b>	<b>Esp16 I</b>	<b>PinA I</b>	<b>Srf I</b>
<b>Age I</b>	<b>Bpu10 I</b>	<b>Dra II</b>	<b>Esp3 I</b>	<b>Pme I</b>	<b>Sse8387 I</b>
<b>AhyA I</b>	<b>Bpu1102 I</b>	<b>Dra III</b>	<b>Fse I</b>	<b>Pml I</b>	<b>Sse8647 I</b>
<b>Ama I</b>	<b>Bpu1268 I</b>	<b>Dsa VI</b>	<b>Fsu I</b>	<b>Ppu10 I</b>	<b>Stu I</b>
<b>Aos III</b>	<b>BsaX I</b>	<b>EciE I</b>	<b>Gsp I</b>	<b>PpuM I</b>	<b>StySJ</b>
<b>Apa I</b>	<b>Bsb I</b>	<b>Eco0109 I</b>	<b>Hine I</b>	<b>PshA I</b>	<b>StySQ</b>
<b>Ape I</b>	<b>BscE I</b>	<b>Eco47 III</b>	<b>I-Ppo I</b>	<b>Psp5 II</b>	<b>Swa I</b>
<b>Asc I</b>	<b>BscJ I</b>	<b>Eco72 I</b>	<b>Kas I</b>	<b>PspA I</b>	<b>Tth I</b>
<b>Asp5H I</b>	<b>Bse59 I</b>	<b>Eco81 I</b>	<b>Kpn I</b>	<b>Pss I</b>	<b>Tth111 I</b>
<b>Asp78 I</b>	<b>BseR I</b>	<b>Eco82 I</b>	<b>Kpn2 I</b>	<b>Pvu II</b>	<b>Uba1220 I</b>
<b>Ava I</b>	<b>Bsg I</b>	<b>EcoA I</b>	<b>Mlu I</b>	<b>Rsr II</b>	<b>Uba1221 I</b>
<b>Ava III</b>	<b>BshL I</b>	<b>EcoB I</b>	<b>Mlu1106 I</b>	<b>Sac II</b>	<b>Uba1326 I</b>
<b>Avr II</b>	<b>BsiW I</b>	<b>EcoD I</b>	<b>Mlu113 I</b>	<b>Sal I</b>	<b>UbaD I</b>
<b>Bae I</b>	<b>Bsmb1</b>	<b>EcoDR2</b>	<b>Nae I</b>	<b>SanD I</b>	<b>Van91 I</b>
<b>Bbe I</b>	<b>Bsp120 I</b>	<b>EcoDR3</b>	<b>Nar I</b>	<b>Sap I</b>	<b>Xho I</b>
<b>BbeA I</b>	<b>Bsp87 I</b>	<b>EcoE I</b>	<b>NgoA IV</b>	<b>Sci I</b>	<b>Xma I</b>
<b>Bbf7411 I</b>	<b>BspG I</b>	<b>EcoN I</b>	<b>NgoM IV</b>	<b>SexA I</b>	
<b>BbrP I</b>	<b>BspJ106 I</b>	<b>EcoO109 I</b>	<b>Nli3877 I</b>	<b>Sfi I</b>	

**Table 3. Restriction Enzymes That Cut the pRL-CMV Vector 6 or More Times.**

<b>Aci I</b>	<b>BstO I</b>	<b>Hae III</b>	<b>Mae II</b>	<b>Nde II</b>	<b>SfaN I</b>
<b>Alu I</b>	<b>BstU I</b>	<b>Hha I</b>	<b>Mae III</b>	<b>Nla III</b>	<b>Taq I</b>
<b>Alw26 I</b>	<b>Cfo I</b>	<b>Hinf I</b>	<b>Mbo I</b>	<b>Nla IV</b>	<b>Tru9 I</b>
<b>Bbv I</b>	<b>Dpn I</b>	<b>Hpa II</b>	<b>Mbo II</b>	<b>Rsa I</b>	<b>Xho II</b>
<b>Bsr I</b>	<b>Dpn II</b>	<b>Hph I</b>	<b>Mnl I</b>	<b>Sau3A I</b>	
<b>BsrS I</b>	<b>Fnu4H I</b>	<b>Hsp92 II</b>	<b>Mse I</b>	<b>Sau96 I</b>	
<b>Bst71 I</b>	<b>Fok I</b>	<b>Mae I</b>	<b>Msp I</b>	<b>ScrF I</b>	

**Note:** The enzymes listed in boldface type are available from Promega.

## B. pRL-CMV Vector Sequence

The sequence shown corresponds to the mRNA synthesized from the *Renilla* luciferase gene from the CMV promoter.

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1  AGATCTTCAA TATTGGCCAT TAGCCATATT ATTCATTGGT TATATAGCAT
51  AAATCAATAT TGGCTATTGG CCATTGCATA CGTTGTATCT ATATCATAAT
101 ATGTACATTT ATATTGGCTC ATGTCCAATA TGACCGCCAT GTTGGCATTG
151 ATTATTGACT AGTTATTAAT AGTAATCAAT TACGGGGTCA TTAGTTCATA
201 GCCCATATAT GGAGTTCCGC GTTACATAAC TTACGGTAAA TGGCCCGCCT
251 GGCTGACCGC CCAACGACCC CCGCCCATTG ACGTCAATAA TGACGTATGT
301 TCCCATAGTA ACGCCAATAG GGACTTTCCA TTGACGTCAA TGGGTGGAGT
351 ATTTACGGTA AACTGCCAC TGGCAGTAC ATCAAGTGTA TCATATGCCA
401 AGTCCGCCCC CTATTGACGT CAATGACGGT AAATGGCCCG CCTGGCATTG
451 TGCCAGTAC ATGACCTTAC GGGACTTTCC TACTTGGCAG TACATCTACG
501 TATTAGTCAT CGCTATTACC ATGGTGATGC GGTTTTGGCA GTACACCAAT
551 GGGCGTGGAT AGCGGTTTGA CTCACGGGGA TTTCCAAGTC TCCACCCCAT
601 TGACGTCAAT GGGAGTTTGT TTTGGCACCA AAATCAACGG GACTTTCCAA
651 AATGTCGTAA TAACCCCGCC CCGTTGACGC AAATGGGCGG TAGGCGTGTA
701 CGGTGGGAGG TCTATATAAG CAGAGCTCGT TTAGTGAACC GTCAGATCAC
751 TAGAAGCTTT ATTGCGGTAG TTTATCACAG TTAATTGCT AACGCAGTCA
801 GTGCTTCTGA CACAACAGTC TCGAACTTAA GCTGCAGAAG TTGGTCGTGA
851 GGCCTGGGC AGGTAAGTAT CAAGGTTACA AGACAGGTTT AAGGAGACCA
901 ATAGAACTG GGCTTGTCGA GACAGAGAAG ACTCTTGCGT TTCTGATAGG
951 CACCTATTGG TCTTACTGAC ATCCACTTTG CCTTTCTCTC CACAGGTGTC
1001 CACTCCAGT TCAATTACAG CTCTTAAGGC TAGAGTACTT AATACGACTC
1051 ACTATAGGCT AGCCACCATG ACTTCGAAAG TTTATGATCC AGAACAAAGG
1101 AAACGGATGA TAACTGGTCC GCAGTGGTGG GCCAGATGTA AACAAATGAA
1151 TGTTCTTGAT TCATTTATTA ATTATTATGA TTCAGAAAAA CATGCAGAAA
1201 ATGCTGTTAT TTTTTTACAT GGTAACGCGG CCTCTTCTTA TTTATGGCGA
1251 CATGTTGTGC CACATATTGA GCCAGTAGCG CGGTGTATTA TACCAGACCT
1301 TATTGGTATG GGCAAATCAG GCAAATCTGG TAATGGTTCT TATAGGTTAC
1351 TTGATCATTG CAAATATCTT ACTGCATGGT TTGAACTTCT TAATTTACCA
1401 AAGAAGATCA TTTTGTGCGG CCATGATTGG GGTGCTTGTT TGGCATTTCG
1451 TTATAGCTAT GAGCATCAAG ATAAGATCAA AGCAATAGTT CACGCTGAAA
1501 GTGTAGTAGA TGTGATTGAA TCATGGGATG AATGGCCTGA TATTGAAGAA
1551 GATATTGCGT TGATCAAATC TGAAGAAGGA GAAAAAATGG TTTTGGAGAA
1601 TAACTTCTTC GTGGAAACCA TGTGCCATC AAAAATCATG AGAAAGTTAG
1651 AACCAGAAGA ATTTGCAGCA TATCTTGAAC CATTCAAAGA GAAAGGTGAA

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1701 GTTCGTCGTC CAACATTATC ATGGCCTCGT GAAATCCCGT TAGTAAAAGG
1751 TGGTAAACCT GACGTTGTAC AAATTGTTAG GAATTATAAT GCTTATCTAC
1801 GTGCAAGTGA TGATTTACCA AAAATGTTTA TTGAATCGGA CCCAGGATTC
1851 TTTTCCAATG CTATTGTTGA AGGTGCCAAG AAGTTTCCTA ATACTGAATT
1901 TGTCAAAGTA AAAGGTCTTC ATTTTTTCGCA AGAAGATGCA CCTGATGAAA
1951 TGGGAAAATA TATCAAATCG TTCGTTGAGC GAGTTCTCAA AAATGAACAA
2001 TAATTCTAGA GCGGCCGCTT CGAGCAGACA TGATAAGATA CATTGATGAG
2051 TTTGGACAAA CCACAAC TAG AATGCAGTGA AAAAAATGCT TTATTTGTGA
2101 AATTTGTGAT GCTATTGCTT TATTTGTAAC CATTATAAGC TGCAATAAAC
2151 AAGTTAACAA CAACAATTGC ATTCATTTTA TGTTTCAGGT TCAGGGGGAG
2201 GTGTGGGAGG TTTTTTAAAG CAAGTAAAAAC CTCTACAAAT GTGGTAAAAAT
2251 CGATAAGGAT CCAGGTGGCA CTTTTCGGGG AAATGTGCGC GGAACCCCTA
2301 TTTGTTTTATT TTTCTAAATA CATTCAAATA TGTATCCGCT CATGAGACAA
2351 TAACCCTGAT AAATGCTTCA ATAATATTGA AAAAGGAAGA GTATGAGTAT
2401 TCAACATTTT CGTGTCGCCC TTATTCCTT TTTTGC GGCA TTTTGCCTTC
2451 CTGTTTTTTC TCACCCAGAA ACGCTGGTGA AAGTAAAAGA TGCTGAAGAT
2501 CAGTTGGGTG CACGAGTGGG TTACATCGAA CTGGATCTCA ACAGCGGTAA
2551 GATCCTTGAG AGTTTTTCGCC CCGAAGAACG TTTTCCAATG ATGAGCACTT
2601 TTAAAGTTCT GCTATGTGGC GCGGTATTAT CCCGTATTGA CGCCGGGCAA
2651 GAGCAACTCG GTCGCCGCAT ACACTATTCT CAGAATGACT TGGTTGAGTA
2701 CTCACCAGTC ACAGAAAAGC ATCTTACGGA TGGCATGACA GTAAGAGAAT
2751 TATGCAGTGC TGCCATAACC ATGAGTGATA ACACTGCGGC CAACTTACTT
2801 CTGACAACGA TCGGAGGACC GAAGGAGCTA ACCGCTTTTT TGCAACAACAT
2851 GGGGGATCAT GTAACTCGCC TTGATCGTTG GGAACCGGAG CTGAATGAAG
2901 CCATACCAA CGACGAGCGT GACACCACGA TGCCTGTAGC AATGGCAACA
2951 ACGTTGCGCA AACTATTAAC TGGCGAACTA CTTACTCTAG CTTCCCGGCA
3001 ACAATTAATA GACTGGATGG AGGCGGATAA AGTTGCAGGA CCACTTCTGC
3051 GCTCGGCCCT TCCGGCTGGC TGGTTTATTG CTGATAAATC TGGAGCCGGT
3101 GAGCGTGGGT CTCGCGGTAT CATTGCAGCA CTGGGGCCAG ATGGTAAGCC
3151 CTCCCGTATC GTAGTTATCT ACACGACGGG GAGTCAGGCA ACTATGGATG
3201 AACGAAATAG ACAGATCGCT GAGATAGGTG CCTCACTGAT TAAGCATTGG
3251 TAACTGTCAG ACCAAGTTTA CTCATATATA CTTTAGATTG ATTTAAAACT
3301 TCATTTTTTAA TTTAAAAGGA TCTAGGTGAA GATCCTTTTT GATAATCTCA
3351 TGACCAAAAT CCCTTAACGT GAGTTTTTCGT TCCACTGAGC GTCAGACCCC
3401 GTAGAAAAGA TCAAAGGATC TTCTTGAGAT CCTTTTTTTC TGCGCGTAAT
3451 CTGCTGCTTG CAAACAAAAA AACCACCGCT ACCAGCGGTG GTTTGTGTTGC
3501 CGGATCAAGA GCTACCAACT CTTTTTCCGA AGGTAACTGG CTTCAGCAGA
3551 GCGCAGATAC CAAATACTGT TCTTCTAGTG TAGCCGTAGT TAGGCCACCA

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3601  CTTCAAGAAC  TCTGTAGCAC  CGCCTACATA  CCTCGCTCTG  CTAATCCTGT
3651  TACCAGTGGC  TGCTGCCAGT  GCGGATAAGT  CGTGTCTTAC  CGGGTTGGAC
3701  TCAAGACGAT  AGTTACCGGA  TAAGGCGCAG  CGGTCGGGCT  GAACGGGGGG
3751  TTCGTGCACA  CAGCCCAGCT  TGGAGCGAAC  GACCTACACC  GAACTGAGAT
3801  ACCTACAGCG  TGAGCTATGA  GAAAGCGCCA  CGCTTCCCGA  AGGGAGAAAG
3851  GCGGACAGGT  ATCCGGTAAG  CGGCAGGGTC  GGAACAGGAG  AGCGCACGAG
3901  GGAGCTTCCA  GGGGGAAACG  CCTGGTATCT  TTATAGTCCT  GTCGGGTTC
3951  GCCACCTCTG  ACTTGAGCGT  CGATTTTGT  GATGCTCGTC  AGGGGGGCGG
4001  AGCCTATGGA  AAAACGCCAG  CAACGCGGCC  TTTTACGGT  TCCTGGCCTT
4051  TTGCTGGCCT  TTTGCTCACA  TGGCTCGAC

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## VI. Related Products

### pRL Family of *Renilla* Luciferase Vectors for Co-Reporter Applications

Product	Size	Cat.#
pRL-TK Vector <sup>(a,c)</sup>	20µg	E2241
pRL-SV40 Vector <sup>(a,c)</sup>	20µg	E2231
pRL-null Vector <sup>(a,c)</sup>	20µg	E2271

To inquire about the availability of bulk packaging and pricing for individual pRL Vectors, please contact Promega. Please visit our Internet site ([www.promega.com](http://www.promega.com)) or call Promega Technical Services to inquire about the availability of new promoter variations within the pRL family of co-reporter vectors.

### Luciferase Assay

Product	Size	Cat.#
Dual-Luciferase™ Reporter Assay System <sup>(c,d,e)</sup>	1 each	E1910
Dual-Luciferase™ Reporter Assay 10-Pack <sup>(c,d,e)</sup>	1,000 assays	E1960
Dual-Luciferase™ Reporter 1000 Assay System <sup>(c,d,e)</sup>	1,000 assays	E1980

### Transfection Systems

Product	Size	Cat.#
Transfectam® Reagent <sup>(h)</sup> for the Transfection of Eukaryotic Cells	1mg	E1231
	0.5mg	E1232
Tfx™-50 Reagent <sup>(i)</sup>	2.1mg	E1811
Tfx™-10 Reagent <sup>(i)</sup>	9.3mg	E2381
Tfx™-20 Reagent <sup>(i)</sup>	4.8mg	E2391
Tfx™ Reagents Transfection Trio <sup>(i)</sup>	5.4mg	E2400
TransFast™ Transfection Reagent <sup>(i)</sup>	1.2mg	E2431
ProFection® Mammalian Transfection System - Calcium Phosphate	1 system	E1200
ProFection® Mammalian Transfection System - DEAE-Dextran	1 system	E1210

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<sup>(b)</sup>The CMV promoter and its use are covered under U.S. Pat. Nos. 5,168,062 and 5,385,839 owned by the University of Iowa Research Foundation, Iowa City, Iowa and licensed FOR RESEARCH USE ONLY. Commercial users must obtain a license to these patents directly from the University of Iowa Research Foundation.

<sup>(c)</sup>Certain applications of this product may require licenses from others.

<sup>(d)</sup>U.S. Pat. Nos. 5,283,179, 5,641,641, 5,650,289, Australian Pat. No. 649289, and European Pat. No. 0 553 234, have been issued to Promega Corporation for a firefly luciferase assay method, which affords greater light output with improved kinetics as compared to the conventional assay.

<sup>(e)</sup>U.S. Pat. No. 5,744,320 has been issued to Promega Corporation for quenching reagents and assays for enzyme-mediated luminescence.

<sup>(f)</sup>U.S. Pat. Nos. 5,324,637, 5,492,817 and 5,665,563, European Pat. No. 0 566 714 B1, Australian Pat. No. 660329, and Japanese Pat. No. 2,904,583 have been issued to Promega Corporation for coupled transcription/translation systems that use RNA polymerases and eukaryotic lysates.

<sup>(g)</sup>The method of recombinant expression of *Coleoptera* luciferase is covered by U.S. Pat. Nos. 5,583,024, 5,674,713 and 5,700,673.

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