

**pKCC was obtained from pECFP-N1 (Clontech #6900-1) after insertion of a new mutiple cloning site in NheI-NotI and mutation of the 3' end of *Xba*I site to prevent methylation ( $T_{690} \rightarrow A$ , see \* in MCS below)**

- Human cytomegalovirus (CMV) immediate early promoter: 1–589  
Enhancer region: 59–465  
TATA box: 554–560  
Transcription start point: 583  
C→G mutation to remove *Sac*I site: 569
- MCS: 591–689
- SV40 early mRNA polyadenylation signal  
Polyadenylation signals: 825–830 & 854–859  
mRNA 3' ends: 863 & 875
- f1 single-strand DNA origin: 922–1377
- Bacterial promoter for expression of Kan<sup>r</sup> gene:  
–35 region: 1439–1444; –10 region: 1462–1467  
Transcription start point: 1474
- SV40 origin of replication: 1517–1853
- SV40 early promoter  
Enhancer (72-bp tandem repeats): 1551–1622 & 1623–1694  
21-bp repeats: 1698–1718, 1719–1739 & 1741–1761  
Early promoter element: 1774–1780  
Major transcription start points: 1770, 1808, 1814 & 1819
- Kanamycin/neomycin resistance gene  
Neomycin phosphotransferase coding sequence: 1902–2696  
G→A mutation to remove *Pst*I site: 2084  
C→A (Arg to Ser) mutation to remove *Bss*H II site: 2430
- Herpes simplex virus (HSV) thymidine kinase (TK) polyadenylation signal  
Polyadenylation signals: 2932–2937 & 2945–2950
- pUC plasmid replication origin: 3281–3924

