

**SUPPLEMENTARY TABLE 5**  
**HOUSEKEEPING GENE PROMOTERS**

<b>GENE</b>	<b>EPD NUMBER ep--</b>
Hs superoxide dismutase 1 ( <i>SOD-1</i> )	07053
Hs dihydrofolate reductase ( <i>DHFR</i> )	07056
Mm hypoxanthine phosphoribosyltransferase ( <i>HPRT</i> )	07058
Hs adenosine deaminase ( <i>ADA</i> )	11113
Mm thymidylate synthetase ( <i>TS</i> )	15031
Mm adenosine deaminase ( <i>ADA</i> )	15032
Mm DNA polymerase b'	16049
Mm dihydrofolate reductase P1 ( <i>DHFR</i> )	24032
Hs thymidine kinase ( <i>TK</i> )	25035
Mm RNA polymerase II ( <i>RP215</i> )	28001
Hs glucose-6-phosphate dehydrogenase ( <i>G6PD</i> )	30014
Mm malate dehydrogenase cytosolic ( <i>MDH1</i> )	33012
Hs nucleolin	36018
Hs ADP-ribosylation factor 1 ( <i>ARF1</i> )	39001
Mm p97 ATPase	59010
Hs cyclin D1	60011
Hs DNA nucleotidylexotransferase ( <i>TdT</i> )	64003
Hs topoisomerase 3 alpha ( <i>TOP3A</i> )	64007
Hs single stranded DNA binding protein 1	73019
Hs glyceraldehyde-3-phosphate dehydrogenase ( <i>GAPD</i> )	73651

The – 200 to +100 regions of these promoters were scanned for potential GABP, Sp1, YY1 and BoxA factor binding sites (as defined for nonaligned motifs in Figure 2).