

Table 9.					
Glucocorticoid Receptor DNA Footprints					
5' Position	3' Position	Footprint Sequence	DNA Source	Method	Reference
		DR3 TGTTCTNNNTGTTCT			
		IR3 TGTTCTNNNAGAACA			
		ER3 TCTTGTTNNNACAAGA			
-187	-163	ATGGTTACA AACTGTTCT TAAACA	MMTV	DNAase	Scheidereit et al. 1983
-122	-105	AA TGTTCT GATCTGAGC	MMTV	DNAase	Scheidereit et al. 1983
-102	-89	TTAGTGTCT ATTT	MMTV	DNAase	Scheidereit et al. 1983
-83	-71	TGTTCT TTTGAA	MMTV	DNAase	Scheidereit et al. 1983
-305	-289	CCAAGGAGGGGACAGTGG	MMTV	DNAase	Payvar et al. 1983
-283	-269	ACTAAT AGAACA TTA	MMTV	DNAase	Payvar et al. 1983
-189	-166	TT TTGGTT ACA AACTGTTCT TAA	MMTV	DNAase	Payvar et al. 1983
-159	-135	GATGTGAGACAAGTGGTTCCCTGAC	MMTV	DNAase	Payvar et al. 1983
-127	-84	GGTATCAAAGGTTCTGATCTGAGCTCT AGTGTCT ATTTTCCT	MMTV	DNAase	Payvar et al. 1983
4313	4343	TTGGCACACGCCTTTTTGTGCTGAATC	MMTV	DNAase	Payvar et al. 1983
4442	4471	GGGTCCTGGAAAGGACGATGTCCTAAT	MMTV	DNAase	Payvar et al. 1983
5345	5367	GTGTAGATCAGTCAGATCAGATT	MMTV	DNAase	Payvar et al. 1983
5372	5393	GCAAAAAGGATCTATTTGGAAA	MMTV	DNAase	Payvar et al. 1983
-265	-245	CCCGGTACACTGTGCCTCCC	HMT-IIA	DNAase	Karin et al. 1984
-180	-164	CAA ACTGTTCT TAAAAC	MMTV	MI, MP	Scheidereit & Beato 1984
-2708	-2676	GTGTACAG TCTTGTCT CCCTCCGTGCAAGC	RUG	DNAase	Cato et al. 1984
-2674	-2652	AACCGGACACGGAGTCCTTTTGC	RUG	DNAase	Cato et al. 1984
-2644	-2619	CACTCTGTTCAC TCTGTTCT ATGCCT	RUG	DNAase	Cato et al. 1984
86	113	CCCTTTGGGCACAATGTGCTCTGAGGGG	HGHPEI1	ExonucI	Slater et al. 1985
-357	-326	AAGTTCAGATCAAGGTCAGGAACAGAGAGACA	MOMSVLTR	DNAase	Miksicek et al. 1986
-267	-250	CA AGAACA GATGGAACAG	MOMSVLTR	DNAase	Miksicek et al. 1986
-194	-172	CCA AGAACA GATGGTCCCCAGAT	MOMSVLTR	DNAase	Miksicek et al. 1986
-349	-331	ATCAAGGTCAGGAACAGAG	MOMSVLTR	DNAase	Defranco & Yamamoto 1986
-271	-247	GGGCCA AGAACA GATGGAACAGCTG	MOMSVLTR	DNAase	Defranco & Yamamoto 1986
-198	-176	AGGGCCA AGAACA GATGGTCCCC	MOMSVLTR	DNAase	Defranco & Yamamoto 1986
-202	-174	CAACAGACTATAAAATTCCTCTGTGGCTT	CLYSOZYM	DNAase	von der Ahe et al. 1986
-76	-49	TTGACA ACTGTAGAACA GAGGAATCAA	CLYSOZYM	DNAase	von der Ahe et al. 1986
-1196	-1156	CCAGGACATCATGAAAGGGAGCAGTGAACATACCCTTCCCG	RTO	DNAase	Danesch et al. 1987
-433	-389	TGCACAGCGAGTTCTAGTGAGTTTCATTTGCCAAGGACATGCCAA	RTO	DNAase	Danesch et al. 1987

DR3 - Direct Repeat with three base pair inter-half site spacing, IR3 - Inverted repeat with three base pair inter-half site spacing, ER3 - Everted repeat with three base pair inter-half site spacing, MMTV - murine mammary tumor virus, HMT-IIA - human metallothionein IIA gene, RUG - rabbit uteroglobin gene, HGHPEI1 - human growth hormone (promoter, exon 1, intron1), MOMSVLTR - Moloney murine sarcoma virus long terminal repeat, CLYSOZYM - chicken lysozyme gene, RTO - rat tryptophan oxygenase gene, MP - methylation protection, MI - methylation interference, ExonucIII - exonuclease III, MMTV bases highlighted in blue indicate sequence differences between studies.