

Additions and Corrections

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Response elements of the androgen-regulated C3 gene.

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Page 4460, Table I: There are two errors in this table. At nucleotide –258, sequence should read ACAGTG and at nucleotide 2559, sequence should read ACAGTG. The corrected table is shown below.

TABLE I

Potential regulatory sequences in prostatein C3 gene

Potential regulatory sequences within the C3 gene and its 5'- and 3'-flanking DNA. Nucleotide numbering is according to Hurst and Parker (21) and refers to the 5' nucleotide. Regulatory proteins are abbreviated and referenced. Homology relates to the consensus DNA sequence for regulatory factor binding.

Position	Nucleotide number	Sequence	Regulatory protein	% Homology	
5'-Flanking	-2098 ^a	ATGTAAAT	OCT 1 (87)	87.5	
	-2091 ^a	ATGCAAGT	OCT 1	87.5	
	-1773	CACTGT	c-myc (86)	100	
	-1731 ^a	GGGTGTGAT	CACCC (72)	78	
	-1718 ^a	GTGCAAAT	OCT 1	87.5	
	-776	AAATCTTAGAGAGAAAA	TOPO 1 (84)	87.5	
	-698 ^a	TGAGTCA	AP 1 (85)	100	
	-676 ^a	CTTTCCAAA	C/EBP (87)	89	
	-472 ^a	AAGCAAAT	OCT 1	87.5	
	-258 ^a	ACAGTG	c-myc	100	
	-124 ^a	TCGCCC	SP 1 (87)	83	
	First intron	845	TGATGTCA	CREB (87)	87.5
		1070	GTGTGGTTTG	SV40 Core enhancer (88)	100
1176		TGACTAG	AP 1	86	
1287 ^a		ATGCAAAT	OCT 1	100	
1456 ^a		AGGTGTGGC	CACCC	89	
Second intron	1920	AGGCGG	SP 1	83	
Third exon	2559 ^a	ACAGTG	c-myc	100	
3'-Flanking	3032 ^a	CCCTGGGG	AP 2 (87)	87.5	
	3444	CCCCAGCC	AP 2	87.5	
	3515	CCCCAGCC	AP 2	87.5	
	3867	CACTGT	c-myc	100	
	3951	CACTGT	c-myc	100	
	4060 ^a	GCCTGGGA	AP 2	87.5	
	4163	GGGCGG	SP 1	100	
	4281 ^a	GCCTGGTG	AP 2	87.5	

^a Sequence shown is in the antisense strand.

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