

Title: DNA Elements for Constitutive Androstane Receptor- and Pregnane X Receptor-mediated Regulation of Bovine CYP3A28 Gene

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S4 Table. Oligonucleotides used for the internal site-directed mutagenesis of transcription factor binding sites.

Target motif	5'→3' sequences	Restriction enzyme site	Length (bp)	%GC
ER6	<i>F</i> : TTATTGATTCCTACAGAATA <u>actAgTCa</u> AGGGAtGTaAAACTGAAGTGCATGTTAT	BcuI	57	31.6
	<i>R</i> : ATAACATGCACTTCAGTTTtACaTCCCTtG <u>AcTagt</u> TATTCTGTAGGAAATCAATAA		57	31.6
DR1	<i>F</i> : GGCTTTATTTTATTTTCTGTtAC <u>gccgcGGt</u> TTACTGATAAACTGATTG	Cfr42I	50	38.0
	<i>R</i> : CAATCAGTGTTATCAGTAAa <u>CCgcggc</u> GTaACAGAAAATAAAATAAAGCC		50	38.0
DR5	<i>F</i> : AGGGAATTCCTGGGTGAaa <u>CtAGTCTAGc</u> GTgAGATAAAACTCCTTAGAAGGTT	BcuI	55	41.8
	<i>R</i> : AACCTTCTAAGGAGTTTTATCTcACgCTAG <u>ACTaGtt</u> TCACCCAGGAAATTCCT		55	41.8

Primers used for mutagenesis of ER6, DR1 present in the *CYP3A28* proximal promoter and DR5 located in F3 are indicated and lowercase letters refer to mutated bases. A selected restriction enzyme site was inserted in both primers (underlined in the 5'→3' sequence) for the diagnosis of successful base changing.

