		Best TRANSFAC Match			
SeqSet	Sequence Logo	Sequence Logo	Accession	Factor	Divergence
EPD-liver	TTAATÇATTA	TGGTTAATAATTAACAGT	M00790	HNF1	0.32
LSPS	TTAATGATTA	GCTTAATGATTACCA	M00132	HNF1	0.24
WF	<u> GGGGEEGGGGE</u>	<u><u>e</u>GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG</u>	M00932	Sp-1	0.78
WF	EGGG EGGGGG	EGGG	M00986	Churchill	0.60
EPD-muscle	<u>CCATATATGG</u>	GCCATATATGGEGAG	M00215	SRF	0.20
EPD-muscle		ACCORCECTOR	M00491	MAZR	0.72
EPD-muscle	CCATATATGG	CATCTCCTTATATCCCAT GGGGCCATTATATCCCCAT	M01007	SRF	0.25
EPD-muscle	<u>CCATATATGG</u>	GTTTACATATATAG	M00392	AGL3	0.27
EPD-muscle		GGGGAGGG	M00649	MAZ	0.69
EPD-muscle	<u> </u>	<u><u><u></u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u><u></u></u>	M00083	MZF1	0.60

Table 9. Top motifs identified by MDscan after masking simple repeats