

Table 4 ESTs related to transcriptional regulation

NO.	GenBank ID of homology	Putative function
E01	NP_850477.1	small nuclear ribonucleoprotein D2
E02	ABD32582.1	Integrase, catalytic region; Zinc finger, CCHC-type
E40	XP_002340257.1	C6 transcription factor, putative
E41	YP_004227811.1	LysR family transcriptional regulator
E79	NP_001133511.1	FYN-binding protein
E105	XP_003225943.1	zinc finger protein 420-like
E120	EFN78085.1	Zinc finger and BTB domain-containing protein 17
E134	AAN63154.1	transcription factor MYBS3
E141	ABG65911.1	Zinc finger, C3HC4 type family protein, expressed
E144	ZP_00371933.1	Ribbon-helix-helix protein, copG family domain protein
E157	XP_003614387.1	rRNA intron-encoded homing endonuclease
E159	ABG65969.1	retrotransposon protein, putative, Ty3-gypsy subclass
E185/ E186	AAO46041.1	zinc finger protein ZFP252
E201	AAD15818.1	transcription factor MYC7E
E207	XP_002530460.1	transcription cofactor, putative
E213	XP_002135694.1	GA25504 gene product from transcript GA25504-RA
E216	XP_002768057.1	DEAD box ATP-dependent RNA helicase
E220	XP_003606916.1	Glycine-rich RNA binding protein