

Supplementary Table S10. The prediction of the transcription factors of ZEB1.

seqname	source	feature	start	end	score	strand	frame	attributes
ZEB1 NM_01323647	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323647	TF BS	TF BS	10 44	10 49	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323647	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323647	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323647	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptophan cluster factors;sequence=ACCGAAACT
ZEB1 NM_01323647	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C2H2 zinc finger factors;sequence=ATAGTAGGTG
ZEB1 NM_01128128	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01128128	TF BS	TF BS	14 77	14 83	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTTCCA
ZEB1 NM_01128128	TF BS	TF BS	84 4	85 1	1	-	.	TF=PITX1;class=Homeo domain factors;sequence=CTAATCCC
ZEB1 NM_01323641	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC

ZEB1 NM_01323641	TF BS	TF BS	14 42	14 47	1	-	.	CAC TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323641	TF BS	TF BS	75 4	81 1	1	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=TTG ACAG
ZEB1 NM_01323641	TF BS	TF BS	67 4	68 1	1	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323641	TF BS	TF BS	67 4	68 1	1	-	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323641	TF BS	TF BS	61 1	61 9	1	-	.	TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_01323641	TF BS	TF BS	10 14	10 23	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_01323638	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323638	TF BS	TF BS	38 8	39 3	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323638	TF BS	TF BS	18 43	18 48	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323638	TF BS	TF BS	47 6	48 2	1	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=TTG ACAG
ZEB1 NM_01323638	TF BS	TF BS	31 0	32 1	1	+	.	TF=ZNF384;class=C2H 2 zinc finger factors;sequence=TTTA

								AAAAAAAAA
ZEB1 NM_01323638	TF BS	TF BS	85 75	92 82	1	+	.	TF=GATA3;class=Other C4 zinc finger-type factors;sequence=AGATAAGA
ZEB1 NM_01323638	TF BS	TF BS	10 75	10 82	1	-	.	TF=HOXB4;class=Homeo domain factors;sequence=ATCATTAG
ZEB1 NM_01323638	TF BS	TF BS	39 5	40 2	1	-	.	TF=HOXC4;class=Homeo domain factors;sequence=GTCATTAA
ZEB1 NM_01323638	TF BS	TF BS	10 75	10 82	1	-	.	TF=HOXD4;class=Homeo domain factors;sequence=ATCATTAG
ZEB1 NM_01323638	TF BS	TF BS	10 12	10 20	1	-	.	TF=IRF6;class=Tryptophan cluster factors;sequence=ACCGAAACT
ZEB1 NM_01323638	TF BS	TF BS	14 15	14 24	1	-	.	TF=ZKSCAN1;class=C2H2 zinc finger factors;sequence=ATAGTAGGTG
ZEB1 NM_01323644	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCCAC
ZEB1 NM_01323644	TF BS	TF BS	14 42	14 47	1	-	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCCAC
ZEB1 NM_01323644	TF BS	TF BS	75	81	1	-	.	TF=MEIS1;class=Homeo domain factors;sequence=TTGACAG
ZEB1 NM_01323644	TF BS	TF BS	67 4	68 1	1	-	.	TF=HOXB4;class=Homeo domain factors;sequence=ATCATTAG
ZEB1 NM_01323644	TF BS	TF BS	67 4	68 1	1	-	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC

ZEB1 NM_01323644	TF BS	TF BS	61 1	61 9	1	-	.	ATTAG TF=IRF6;class=Tryptophan cluster factors;sequence=ACCGAAACT
ZEB1 NM_01323644	TF BS	TF BS	10 14	10 23	1	-	.	TF=ZKSCAN1;class=C2H2 zinc finger factors;sequence=ATAGTAGGTG
ZEB1 NM_01323643	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCACAC
ZEB1 NM_01323643	TF BS	TF BS	38 8	39 3	1	-	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCACAC
ZEB1 NM_01323643	TF BS	TF BS	18 43	18 48	1	-	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCACAC
ZEB1 NM_01323643	TF BS	TF BS	47 6	48 2	1	-	.	TF=MEIS1;class=Homeo domain factors;sequence=TTGACAG
ZEB1 NM_01323643	TF BS	TF BS	31 0	32 1	1	+	.	TF=ZNF384;class=C2H2 zinc finger factors;sequence=TTTAAAAA
ZEB1 NM_01323643	TF BS	TF BS	85	92	1	+	.	TF=GATA3;class=Other C4 zinc finger-type factors;sequence=AGATAAGA
ZEB1 NM_01323643	TF BS	TF BS	10 75	10 82	1	-	.	TF=HOXB4;class=Homeo domain factors;sequence=ATCATTAG
ZEB1 NM_01323643	TF BS	TF BS	39 5	40 2	1	-	.	TF=HOXC4;class=Homeo domain factors;sequence=GTCATTAA
ZEB1 NM_01323643	TF BS	TF BS	10 75	10 82	1	-	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC

ZEB1 NM_01323643	TF BS	TF BS	10 12	10 20	1	-	.	ATTAG TF=IRF6;class=Tryptophan cluster factors;sequence=ACCGAAACT
ZEB1 NM_01323643	TF BS	TF BS	14 15	14 24	1	-	.	TF=ZKSCAN1;class=C2H2 zinc finger factors;sequence=ATAGTAGGTG
ZEB1 NM_01323674	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCACAC
ZEB1 NM_01323674	TF BS	TF BS	10 44	10 49	1	-	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCACAC
ZEB1 NM_01323674	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Homeo domain factors;sequence=ATCATTAG
ZEB1 NM_01323674	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Homeo domain factors;sequence=ATCATTAG
ZEB1 NM_01323674	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptophan cluster factors;sequence=ACCGAAACT
ZEB1 NM_01323674	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C2H2 zinc finger factors;sequence=ATAGTAGGTG
ZEB1 NM_01323673	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCACAC
ZEB1 NM_01323673	TF BS	TF BS	10 18	10 24	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTCCA
ZEB1 NM_01323673	TF BS	TF BS	38 5	39 2	1	-	.	TF=PITX1;class=Homeo domain

ZEB1 NM_01323658	TF BS	TF BS	23 00	23 05	1	+	.	factors;sequence=CTA ATCCC TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323658	TF BS	TF BS	49 2	49 7	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323658	TF BS	TF BS	64	73	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_01323657	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323657	TF BS	TF BS	49 2	49 7	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323657	TF BS	TF BS	64	73	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_01323650	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323650	TF BS	TF BS	10 44	10 49	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323650	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323650	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323650	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptop han cluster

ZEB1 NM_01323650	TF BS	TF BS	61 6	62 5	1	-	.	factors;sequence=ACC GAAACT TF=ZKSCAN1;class=C 2H2 zinc finger
ZEB1 NM_01323648	TF BS	TF BS	23 00	23 05	1	+	.	factors;sequence=ATA GTAGGTG TF=ZNF354C;class=C2 H2 zinc finger
ZEB1 NM_01323648	TF BS	TF BS	10 44	10 49	1	-	.	factors;sequence=ATC CAC TF=ZNF354C;class=C2 H2 zinc finger
ZEB1 NM_01323648	TF BS	TF BS	27 6	28 3	1	-	.	factors;sequence=ATC CAC TF=HOXB4;class=Ho meo domain
ZEB1 NM_01323648	TF BS	TF BS	27 6	28 3	1	-	.	factors;sequence=ATC ATTAG TF=HOXD4;class=Ho meo domain
ZEB1 NM_01323648	TF BS	TF BS	21 3	22 1	1	-	.	factors;sequence=ATC ATTAG TF=IRF6;class=Tryptop han cluster
ZEB1 NM_01323648	TF BS	TF BS	61 6	62 5	1	-	.	factors;sequence=ACC GAAACT TF=ZKSCAN1;class=C 2H2 zinc finger
ZEB1 NM_01174095	TF BS	TF BS	23 00	23 05	1	+	.	factors;sequence=ATA GTAGGTG TF=ZNF354C;class=C2 H2 zinc finger
ZEB1 NM_01174095	TF BS	TF BS	10 44	10 49	1	-	.	factors;sequence=ATC CAC TF=ZNF354C;class=C2 H2 zinc finger
ZEB1 NM_01174095	TF BS	TF BS	27 6	28 3	1	-	.	factors;sequence=ATC CAC TF=HOXB4;class=Ho meo domain
ZEB1 NM_01174095	TF BS	TF BS	27 6	28 3	1	-	.	factors;sequence=ATC ATTAG TF=HOXD4;class=Ho meo domain

ZEB1 NM_001174095	TF BS	TF BS	21 3	22 1	1	-	.	factors;sequence=ATC ATTAG TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_001174095	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_001323646	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_001323646	TF BS	TF BS	14 42	14 47	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_001323646	TF BS	TF BS	75	81	1	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=TTG ACAG
ZEB1 NM_001323646	TF BS	TF BS	67 4	68 1	1	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_001323646	TF BS	TF BS	67 4	68 1	1	-	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_001323646	TF BS	TF BS	61 1	61 9	1	-	.	TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_001323646	TF BS	TF BS	10 14	10 23	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_001323645	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_001323645	TF BS	TF BS	14 42	14 47	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger

								factors;sequence=ATC CAC
ZEB1 NM_0 01323645	TF BS	TF BS	75 81		1 -	.		TF=MEIS1;class=Homo meo domain factors;sequence=TTG ACAG
ZEB1 NM_0 01323645	TF BS	TF BS	67 4	68 1	1 -	.		TF=HOXB4;class=Homo meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323645	TF BS	TF BS	67 4	68 1	1 -	.		TF=HOXD4;class=Homo meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323645	TF BS	TF BS	61 1	61 9	1 -	.		TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_0 01323645	TF BS	TF BS	10 14	10 23	1 -	.		TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_0 01174093	TF BS	TF BS	23 00	23 05	1 +	.		TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01174093	TF BS	TF BS	10 44	10 49	1 -	.		TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01174093	TF BS	TF BS	27 6	28 3	1 -	.		TF=HOXB4;class=Homo meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01174093	TF BS	TF BS	27 6	28 3	1 -	.		TF=HOXD4;class=Homo meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01174093	TF BS	TF BS	21 3	22 1	1 -	.		TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_0 01174093	TF BS	TF BS	61 6	62 5	1 -	.		TF=ZKSCAN1;class=C 2H2 zinc finger

ZEB1 NM_01323656	TF BS	TF BS	23 00	23 05	1	+	.	factors;sequence=ATA GTAGGTG TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323656	TF BS	TF BS	49 2	49 7	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323656	TF BS	TF BS	64	73	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_01323655	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323655	TF BS	TF BS	10 44	10 49	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323655	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323655	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323655	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_01323655	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_01323654	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323654	TF BS	TF BS	10 44	10 49	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger

								factors;sequence=ATC CAC
ZEB1 NM_0 01323654	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323654	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323654	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_0 01323654	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_0 01323653	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323653	TF BS	TF BS	10 44	10 49	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323653	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323653	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323653	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_0 01323653	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_0 01323652	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger

							factors;sequence=ATC CAC
ZEB1 NM_0 01323652	TF BS	TF BS	10 44	10 49	1	-	. TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323652	TF BS	TF BS	27 6	28 3	1	-	. TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323652	TF BS	TF BS	27 6	28 3	1	-	. TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323652	TF BS	TF BS	21 3	22 1	1	-	. TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_0 01323652	TF BS	TF BS	61 6	62 5	1	-	. TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_0 01323651	TF BS	TF BS	23 00	23 05	1	+	. TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323651	TF BS	TF BS	10 44	10 49	1	-	. TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323651	TF BS	TF BS	27 6	28 3	1	-	. TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323651	TF BS	TF BS	27 6	28 3	1	-	. TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_0 01323651	TF BS	TF BS	21 3	22 1	1	-	. TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_0 01323651	TF BS	TF BS	61 6	62 5	1	-	. TF=ZKSCAN1;class=C 2H2 zinc finger

ZEB1 NM_01323678	TF BS	TF BS	23 00	23 05	1	+	.	factors;sequence=ATAGGTG TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATCAC
ZEB1 NM_01323678	TF BS	TF BS	14 77	14 83	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTTCCA
ZEB1 NM_01323678	TF BS	TF BS	84 4	85 1	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTAATCCC
ZEB1 NM_01323677	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATCAC
ZEB1 NM_01323677	TF BS	TF BS	10 18	10 24	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTTCCA
ZEB1 NM_01323677	TF BS	TF BS	38 5	39 2	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTAATCCC
ZEB1 NM_01323676	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATCAC
ZEB1 NM_01323676	TF BS	TF BS	10 18	10 24	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTTCCA
ZEB1 NM_01323676	TF BS	TF BS	38 5	39 2	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTAATCCC
ZEB1 NM_01323649	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC

ZEB1 NM_01323649	TF BS	TF BS	10 44	10 49	1	-	.	CAC TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323649	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323649	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323649	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptop han cluster factors;sequence=ACC GAAACT
ZEB1 NM_01323649	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C 2H2 zinc finger factors;sequence=ATA GTAGGTG
ZEB1 NM_01323675	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323675	TF BS	TF BS	10 18	10 24	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTT CCA
ZEB1 NM_01323675	TF BS	TF BS	38 5	39 2	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323672	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323672	TF BS	TF BS	10 18	10 24	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTT CCA
ZEB1 NM_01323672	TF BS	TF BS	38 5	39 2	1	-	.	TF=PITX1;class=Home

01323672	BS	BS	5	2				o domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323671	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323671	TF BS	TF BS	10 18	10 24	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTT CCA
ZEB1 NM_0 01323671	TF BS	TF BS	38 5	39 2	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323666	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323666	TF BS	TF BS	14 77	14 83	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTT CCA
ZEB1 NM_0 01323666	TF BS	TF BS	84 4	85 1	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323642	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323642	TF BS	TF BS	14 42	14 47	1	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323642	TF BS	TF BS	75	81	1	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=TTG ACAG
ZEB1 NM_0 01323642	TF BS	TF BS	67 4	68 1	1	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATC

ZEB1 NM_01323642	TF BS	TF BS	67 4	68 1	1	-	.	ATTAG TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323642	TF BS	TF BS	61 1	61 9	1	-	.	TF=IRF6;class=Tryptophan cluster factors;sequence=ACCGAAACT
ZEB1 NM_01323642	TF BS	TF BS	10 14	10 23	1	-	.	TF=ZKSCAN1;class=C2H2 zinc finger factors;sequence=ATAGTAGGTG
ZEB1 NM_01174096	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCCAC
ZEB1 NM_01174096	TF BS	TF BS	10 44	10 49	1	-	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCCAC
ZEB1 NM_01174096	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01174096	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01174096	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptophan cluster factors;sequence=ACCGAAACT
ZEB1 NM_01174096	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C2H2 zinc finger factors;sequence=ATAGTAGGTG
ZEB1 NM_030751	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCCAC
ZEB1 NM_030751	TF BS	TF BS	10 44	10 49	1	-	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATC

								CAC
ZEB1 NM_030751	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_030751	TF BS	TF BS	27 6	28 3	1	-	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_030751	TF BS	TF BS	21 3	22 1	1	-	.	TF=IRF6;class=Tryptophan cluster factors;sequence=ACCGAAACT
ZEB1 NM_030751	TF BS	TF BS	61 6	62 5	1	-	.	TF=ZKSCAN1;class=C2H2 zinc finger factors;sequence=ATAGTAGGTG
ZEB1 NM_01323665	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCCAC
ZEB1 NM_01323665	TF BS	TF BS	14 77	14 83	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTCCA
ZEB1 NM_01323665	TF BS	TF BS	84 4	85 1	1	-	.	TF=PITX1;class=Homeo domain factors;sequence=CTAATCCC
ZEB1 NM_01323664	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger factors;sequence=ATCCAC
ZEB1 NM_01323664	TF BS	TF BS	14 77	14 83	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTCCA
ZEB1 NM_01323664	TF BS	TF BS	84 4	85 1	1	-	.	TF=PITX1;class=Homeo domain factors;sequence=CTAATCCC
ZEB1 NM_01323664	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2H2 zinc finger

01323663	BS	BS	00	05				H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323663	TF BS	TF BS	14 77	14 83	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTT CCA
ZEB1 NM_0 01323663	TF BS	TF BS	84 4	85 1	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323662	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323662	TF BS	TF BS	14 77	14 83	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTT CCA
ZEB1 NM_0 01323662	TF BS	TF BS	84 4	85 1	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323661	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_0 01323661	TF BS	TF BS	19 97	20 07	1	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAC
ZEB1 NM_0 01323661	TF BS	TF BS	19 97	20 06	1	-	.	TF=HOXD9;class=Ho meo domain factors;sequence=GCA ATAAAAC
ZEB1 NM_0 01323661	TF BS	TF BS	14 91	14 98	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323660	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC

								CAC
ZEB1 NM_01323660	TF BS	TF BS	19 97	20 07	1	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAC
ZEB1 NM_01323660	TF BS	TF BS	19 97	20 06	1	-	.	TF=HOXD9;class=Ho meo domain factors;sequence=GCA ATAAAAC
ZEB1 NM_01323660	TF BS	TF BS	14 91	14 98	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323659	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01323659	TF BS	TF BS	19 97	20 07	1	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAC
ZEB1 NM_01323659	TF BS	TF BS	19 97	20 06	1	-	.	TF=HOXD9;class=Ho meo domain factors;sequence=GCA ATAAAAC
ZEB1 NM_01323659	TF BS	TF BS	14 91	14 98	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01174094	TF BS	TF BS	23 00	23 05	1	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=ATC CAC
ZEB1 NM_01174094	TF BS	TF BS	14 77	14 83	1	+	.	TF=NFATC2;class=Rel homology region (RHR) factors;sequence=TTTT CCA
ZEB1 NM_01174094	TF BS	TF BS	84 4	85 1	1	-	.	TF=PITX1;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01128128	TF BS	TF BS	19 09	19 18	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger

								factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323673	TF BS	TF BS	14 50	14 59	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323678	TF BS	TF BS	19 09	19 18	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323677	TF BS	TF BS	14 50	14 59	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323676	TF BS	TF BS	14 50	14 59	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323675	TF BS	TF BS	14 50	14 59	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323672	TF BS	TF BS	14 50	14 59	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323671	TF BS	TF BS	14 50	14 59	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323666	TF BS	TF BS	19 09	19 18	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323665	TF BS	TF BS	19 09	19 18	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323664	TF BS	TF BS	19 09	19 18	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323663	TF BS	TF BS	19 09	19 18	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger

								factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323662	TF BS	TF BS	19 09	19 18	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01174094	TF BS	TF BS	19 09	19 18	0.9997 42203	+	.	TF=KLF5;class=C2H2 zinc finger factors;sequence=GCC CCACCCC
ZEB1 NM_0 01323647	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323674	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323658	TF BS	TF BS	13 74	13 84	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323657	TF BS	TF BS	13 74	13 84	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323650	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323648	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01174095	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01174093	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323656	TF BS	TF BS	13 74	13 84	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger

								factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323655	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323654	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323653	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323652	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323651	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323649	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01174096	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 30751	TF BS	TF BS	19 26	19 36	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323661	TF BS	TF BS	58 8	59 8	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323660	TF BS	TF BS	58 8	59 8	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323659	TF BS	TF BS	58 8	59 8	0.9997 38076	-	.	TF=MAZ;class=C2H2 zinc finger

								factors;sequence=CCC CCCTCCCC
ZEB1 NM_0 01323647	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323641	TF BS	TF BS	66 4	67 1	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323638	TF BS	TF BS	10 65	10 72	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323644	TF BS	TF BS	66 4	67 1	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323643	TF BS	TF BS	10 65	10 72	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323674	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323673	TF BS	TF BS	16 40	16 47	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323650	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323648	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01174095	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323646	TF BS	TF BS	66 4	67 1	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain

								factors;sequence=ATC ATTAA
ZEB1 NM_0 01323645	TF BS	TF BS	66 4	67 1	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01174093	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323655	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323654	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323653	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323652	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323651	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323677	TF BS	TF BS	16 40	16 47	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323676	TF BS	TF BS	16 40	16 47	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323649	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323675	TF BS	TF BS	16 40	16 47	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain

								factors;sequence=ATC ATTAA
ZEB1 NM_0 01323672	TF BS	TF BS	16 40	16 47	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323671	TF BS	TF BS	16 40	16 47	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323642	TF BS	TF BS	66 4	67 1	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01174096	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 30751	TF BS	TF BS	26 6	27 3	0.9995 65985	+	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323641	TF BS	TF BS	96	10 3	0.9994 17997	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AAG GTCAA
ZEB1 NM_0 01323638	TF BS	TF BS	49 7	50 4	0.9994 17997	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AAG GTCAA
ZEB1 NM_0 01323644	TF BS	TF BS	96	10 3	0.9994 17997	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AAG GTCAA
ZEB1 NM_0 01323643	TF BS	TF BS	49 7	50 4	0.9994 17997	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AAG GTCAA
ZEB1 NM_0 01323646	TF BS	TF BS	96	10 3	0.9994 17997	+	.	TF=NR2C2(var.2);class =Nuclear receptors with

								C4 zinc fingers;sequence=AAG GTCAA
ZEB1 NM_0 01323645	TF BS	TF BS	96	10 3	0.9994 17997	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AAG GTCAA
ZEB1 NM_0 01323642	TF BS	TF BS	96	10 3	0.9994 17997	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AAG GTCAA
ZEB1 NM_0 01323647	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323641	TF BS	TF BS	66 4	67 1	0.9993 31142	+	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323638	TF BS	TF BS	10 65	10 72	0.9993 31142	+	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323644	TF BS	TF BS	66 4	67 1	0.9993 31142	+	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323643	TF BS	TF BS	10 65	10 72	0.9993 31142	+	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323674	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323673	TF BS	TF BS	16 40	16 47	0.9993 31142	+	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323650	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Ho meo domain factors;sequence=ATC

								ATTAA
ZEB1 NM_01323648	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01174095	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323646	TF BS	TF BS	66 4	67 1	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323645	TF BS	TF BS	66 4	67 1	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01174093	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323655	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323654	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323653	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323652	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323651	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323677	TF BS	TF BS	16 40	16 47	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC

								ATTAA
ZEB1 NM_01323676	TF BS	TF BS	16 40	16 47	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323649	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323675	TF BS	TF BS	16 40	16 47	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323672	TF BS	TF BS	16 40	16 47	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323671	TF BS	TF BS	16 40	16 47	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323642	TF BS	TF BS	66 4	67 1	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01174096	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_030751	TF BS	TF BS	26 6	27 3	0.9993 31142	+	.	TF=HOXD4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323647	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323641	TF BS	TF BS	67 4	68 1	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323638	TF BS	TF BS	10 75	10 82	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC

								ATTAG
ZEB1 NM_01323644	TF BS	TF BS	67 4	68 1	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323643	TF BS	TF BS	10 75	10 82	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323674	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323650	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323648	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01174095	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323646	TF BS	TF BS	67 4	68 1	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323645	TF BS	TF BS	67 4	68 1	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01174093	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323655	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323654	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC

								ATTAG
ZEB1 NM_01323653	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323652	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323651	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323649	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323642	TF BS	TF BS	67 4	68 1	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01174096	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_30751	TF BS	TF BS	27 6	28 3	0.9992 12186	-	.	TF=HOXC4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323638	TF BS	TF BS	39 8	40 4	0.9991 27731	+	.	TF=MEIS1;class=Homeo domain factors;sequence=ATG ACAG
ZEB1 NM_01323643	TF BS	TF BS	39 8	40 4	0.9991 27731	+	.	TF=MEIS1;class=Homeo domain factors;sequence=ATG ACAG
ZEB1 NM_01323647	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323641	TF BS	TF BS	66 4	67 1	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC

ZEB1 NM_01323638	TF BS	TF BS	10 65	10 72	0.9990 53421	+	.	ATTAA TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323644	TF BS	TF BS	66 4	67 1	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323643	TF BS	TF BS	10 65	10 72	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323674	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323673	TF BS	TF BS	16 40	16 47	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323650	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323648	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01174095	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323646	TF BS	TF BS	66 4	67 1	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323645	TF BS	TF BS	66 4	67 1	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01174093	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC

ZEB1 NM_01323655	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	ATTAA TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323654	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323653	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323652	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323651	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323677	TF BS	TF BS	16 40	16 47	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323676	TF BS	TF BS	16 40	16 47	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323649	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323675	TF BS	TF BS	16 40	16 47	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323672	TF BS	TF BS	16 40	16 47	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323671	TF BS	TF BS	16 40	16 47	0.9990 53421	+	.	TF=HOXB4;class=Homeo domain factors;sequence=ATC

ZEB1 NM_01323642	TF BS	TF BS	66 4	67 1	0.9990 53421	+	.	ATTAA TF=HOXB4;class=Homeo domain factors;sequence=ATC
ZEB1 NM_01174096	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	ATTAA TF=HOXB4;class=Homeo domain factors;sequence=ATC
ZEB1 NM_030751	TF BS	TF BS	26 6	27 3	0.9990 53421	+	.	ATTAA TF=HOXB4;class=Homeo domain factors;sequence=ATC
ZEB1 NM_01323638	TF BS	TF BS	39 5	40 2	0.9988 88511	-	.	ATTAA TF=HOXD4;class=Homeo domain factors;sequence=GTC
ZEB1 NM_01323643	TF BS	TF BS	39 5	40 2	0.9988 88511	-	.	ATTAA TF=HOXD4;class=Homeo domain factors;sequence=GTC
ZEB1 NM_01128128	TF BS	TF BS	84 4	85 1	0.9984 78909	-	.	ATTAA TF=PITX2;class=Homeo domain factors;sequence=CTA
ZEB1 NM_01323673	TF BS	TF BS	38 5	39 2	0.9984 78909	-	.	ATCCC TF=PITX2;class=Homeo domain factors;sequence=CTA
ZEB1 NM_01323678	TF BS	TF BS	84 4	85 1	0.9984 78909	-	.	ATCCC TF=PITX2;class=Homeo domain factors;sequence=CTA
ZEB1 NM_01323677	TF BS	TF BS	38 5	39 2	0.9984 78909	-	.	ATCCC TF=PITX2;class=Homeo domain factors;sequence=CTA
ZEB1 NM_01323676	TF BS	TF BS	38 5	39 2	0.9984 78909	-	.	ATCCC TF=PITX2;class=Homeo domain factors;sequence=CTA
ZEB1 NM_01323675	TF BS	TF BS	38 5	39 2	0.9984 78909	-	.	ATCCC TF=PITX2;class=Homeo domain factors;sequence=CTA

									ATCCC
ZEB1 NM_01323672	TF BS	TF BS	38 5	39 2	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323671	TF BS	TF BS	38 5	39 2	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323666	TF BS	TF BS	84 4	85 1	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323665	TF BS	TF BS	84 4	85 1	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323664	TF BS	TF BS	84 4	85 1	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323663	TF BS	TF BS	84 4	85 1	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323662	TF BS	TF BS	84 4	85 1	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323661	TF BS	TF BS	14 91	14 98	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323660	TF BS	TF BS	14 91	14 98	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01323659	TF BS	TF BS	14 91	14 98	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA ATCCC
ZEB1 NM_01174094	TF BS	TF BS	84 4	85 1	0.9984 78909	-	.		TF=PITX2;class=Home o domain factors;sequence=CTA

								ATCCC
ZEB1 NM_01323638	TF BS	TF BS	83	93	0.9984 57723	-	.	TF=GATA2;class=Other C4 zinc finger-type factors;sequence=TTCTTATCTCT
ZEB1 NM_01323643	TF BS	TF BS	83	93	0.9984 57723	-	.	TF=GATA2;class=Other C4 zinc finger-type factors;sequence=TTCTTATCTCT
ZEB1 NM_01323638	TF BS	TF BS	39 5	40 2	0.9981 06843	-	.	TF=HOXB4;class=Homeo domain factors;sequence=GTCATTAA
ZEB1 NM_01323643	TF BS	TF BS	39 5	40 2	0.9981 06843	-	.	TF=HOXB4;class=Homeo domain factors;sequence=GTCATTAA
ZEB1 NM_01323647	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.	TF=HOXA9;class=Homeo domain factors;sequence=GTCGTAAACC
ZEB1 NM_01323674	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.	TF=HOXA9;class=Homeo domain factors;sequence=GTCGTAAACC
ZEB1 NM_01323658	TF BS	TF BS	13 55	13 64	0.9973 17418	+	.	TF=HOXA9;class=Homeo domain factors;sequence=GTCGTAAACC
ZEB1 NM_01323657	TF BS	TF BS	13 55	13 64	0.9973 17418	+	.	TF=HOXA9;class=Homeo domain factors;sequence=GTCGTAAACC
ZEB1 NM_01323650	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.	TF=HOXA9;class=Homeo domain factors;sequence=GTCGTAAACC
ZEB1 NM_01323648	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.	TF=HOXA9;class=Homeo domain factors;sequence=GTCGTAAACC
ZEB1 NM_01174095	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.	TF=HOXA9;class=Homeo domain factors;sequence=GTC

									GTAAACC
ZEB1 NM_001174093	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001323656	TF BS	TF BS	13 55	13 64	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001323655	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001323654	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001323653	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001323652	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001323651	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001323649	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001174096	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_030751	TF BS	TF BS	19 07	19 16	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_001323661	TF BS	TF BS	56 9	57 8	0.9973 17418	+	.		TF=HOXA9;class=Homeo domain factors;sequence=GTC

ZEB1 NM_01323660	TF BS	TF BS	56 9	57 8	0.9973 17418	+	.	GTAAACC TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_01323659	TF BS	TF BS	56 9	57 8	0.9973 17418	+	.	TF=HOXA9;class=Homeo domain factors;sequence=GTC GTAAACC
ZEB1 NM_01128128	TF BS	TF BS	12 90	13 00	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323673	TF BS	TF BS	83 1	84 1	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323678	TF BS	TF BS	12 90	13 00	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323677	TF BS	TF BS	83 1	84 1	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323676	TF BS	TF BS	83 1	84 1	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323675	TF BS	TF BS	83 1	84 1	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323672	TF BS	TF BS	83 1	84 1	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323671	TF BS	TF BS	83 1	84 1	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323666	TF BS	TF BS	12 90	13 00	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC

								CCCTCCCC
ZEB1 NM_01323665	TF BS	TF BS	12 90	13 00	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323664	TF BS	TF BS	12 90	13 00	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323663	TF BS	TF BS	12 90	13 00	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323662	TF BS	TF BS	12 90	13 00	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323661	TF BS	TF BS	19 37	19 47	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323660	TF BS	TF BS	19 37	19 47	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323659	TF BS	TF BS	19 37	19 47	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01174094	TF BS	TF BS	12 90	13 00	0.9969 87594	-	.	TF=MAZ;class=C2H2 zinc finger factors;sequence=GCC CCCTCCCC
ZEB1 NM_01323647	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01323641	TF BS	TF BS	48 4	49 1	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_0	TF	TF	88	89	0.9969	+	.	TF=NR2C2(var.2);class

01323638	BS	BS	5	2	78314			=Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01323644	TF BS	TF BS	48 4	49 1	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01323643	TF BS	TF BS	88 5	89 2	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01323674	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01323650	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01323648	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01174095	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01323646	TF BS	TF BS	48 4	49 1	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01323645	TF BS	TF BS	48 4	49 1	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA

ZEB1 NM_001174093	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_001323655	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_001323654	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_001323653	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_001323652	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_001323651	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_001323649	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_001323642	TF BS	TF BS	48 4	49 1	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_001174096	TF BS	TF BS	86	93	0.9969 78314	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG

ZEB1 NM_030751	TF BS	TF BS	86	93	0.9969 78314	+	.	GTCAA TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=GGG GTCAA
ZEB1 NM_01128128	TF BS	TF BS	84 4	85 1	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323673	TF BS	TF BS	38 5	39 2	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323678	TF BS	TF BS	84 4	85 1	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323677	TF BS	TF BS	38 5	39 2	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323676	TF BS	TF BS	38 5	39 2	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323675	TF BS	TF BS	38 5	39 2	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323672	TF BS	TF BS	38 5	39 2	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323671	TF BS	TF BS	38 5	39 2	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323666	TF BS	TF BS	84 4	85 1	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_01323665	TF BS	TF BS	84 4	85 1	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain

								factors;sequence=CTA ATCCC
ZEB1 NM_0 01323664	TF BS	TF BS	84 4	85 1	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323663	TF BS	TF BS	84 4	85 1	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323662	TF BS	TF BS	84 4	85 1	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323661	TF BS	TF BS	14 91	14 98	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323660	TF BS	TF BS	14 91	14 98	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01323659	TF BS	TF BS	14 91	14 98	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01174094	TF BS	TF BS	84 4	85 1	0.9969 68423	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=CTA ATCCC
ZEB1 NM_0 01128128	TF BS	TF BS	13 38	13 47	0.9964 48102	-	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_0 01323673	TF BS	TF BS	87 9	88 8	0.9964 48102	-	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_0 01323678	TF BS	TF BS	13 38	13 47	0.9964 48102	-	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=TGC

									AGCTGCG
ZEB1 NM_01323677	TF BS	TF BS	87 9	88 8	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323676	TF BS	TF BS	87 9	88 8	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323675	TF BS	TF BS	87 9	88 8	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323672	TF BS	TF BS	87 9	88 8	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323671	TF BS	TF BS	87 9	88 8	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323666	TF BS	TF BS	13 38	13 47	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323665	TF BS	TF BS	13 38	13 47	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323664	TF BS	TF BS	13 38	13 47	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323663	TF BS	TF BS	13 38	13 47	0.9964 48102	-	.		TF=BHLHE22(var.2);class=Basic helix-loop-helix factors

ZEB1 NM_01323662	TF BS	TF BS	13 38	13 47	0.9964 48102	-	.	(bHLH);sequence=TGC AGCTGCG TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323661	TF BS	TF BS	19 85	19 94	0.9964 48102	-	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323660	TF BS	TF BS	19 85	19 94	0.9964 48102	-	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01323659	TF BS	TF BS	19 85	19 94	0.9964 48102	-	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01174094	TF BS	TF BS	13 38	13 47	0.9964 48102	-	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=TGC AGCTGCG
ZEB1 NM_01128128	TF BS	TF BS	13 38	13 47	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_01323673	TF BS	TF BS	87 9	88 8	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_01323678	TF BS	TF BS	13 38	13 47	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_01323677	TF BS	TF BS	87 9	88 8	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic

								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323676	TF BS	TF BS	87 9	88 8	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic
								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323675	TF BS	TF BS	87 9	88 8	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic
								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323672	TF BS	TF BS	87 9	88 8	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic
								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323671	TF BS	TF BS	87 9	88 8	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic
								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323666	TF BS	TF BS	13 38	13 47	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic
								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323665	TF BS	TF BS	13 38	13 47	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic
								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323664	TF BS	TF BS	13 38	13 47	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic
								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323663	TF BS	TF BS	13 38	13 47	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic
								helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0	TF	TF	13	13	0.9964	+	.	TF=BHLHE22(var.2);cl

01323662	BS	BS	38	47	24556			ass=Basic helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323661	TF BS	TF BS	19 85	19 94	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323660	TF BS	TF BS	19 85	19 94	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323659	TF BS	TF BS	19 85	19 94	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01174094	TF BS	TF BS	13 38	13 47	0.9964 24556	+	.	TF=BHLHE22(var.2);cl ass=Basic helix-loop-helix factors (bHLH);sequence=CGC AGCTGCA
ZEB1 NM_0 01323647	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_0 01128128	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_0 01128128	TF BS	TF BS	13 49	13 55	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_0 01323641	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_0 01323641	TF BS	TF BS	12 2	12 8	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA

ZEB1 NM_01323638	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323638	TF BS	TF BS	52 3	52 9	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323644	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323644	TF BS	TF BS	12 2	12 8	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323643	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323643	TF BS	TF BS	52 3	52 9	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323674	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323673	TF BS	TF BS	16 79	16 85	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323673	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323673	TF BS	TF BS	89 0	89 6	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323658	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA

ZEB1 NM_01323657	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323650	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323648	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01174095	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323646	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323646	TF BS	TF BS	12 2	12 8	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323645	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323645	TF BS	TF BS	12 2	12 8	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01174093	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323656	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323655	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA

ZEB1 NM_01323654	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323653	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323652	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323651	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323678	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323678	TF BS	TF BS	13 49	13 55	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323677	TF BS	TF BS	16 79	16 85	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323677	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323677	TF BS	TF BS	89 0	89 6	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323676	TF BS	TF BS	16 79	16 85	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323676	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA

ZEB1 NM_01323676	TF BS	TF BS	89 0	89 6	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323649	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323675	TF BS	TF BS	16 79	16 85	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323675	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323675	TF BS	TF BS	89 0	89 6	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323672	TF BS	TF BS	16 79	16 85	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323672	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323672	TF BS	TF BS	89 0	89 6	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323671	TF BS	TF BS	16 79	16 85	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323671	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323671	TF BS	TF BS	89 0	89 6	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA

ZEB1 NM_01323666	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323666	TF BS	TF BS	13 49	13 55	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323642	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323642	TF BS	TF BS	12 2	12 8	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01174096	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_030751	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323665	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323665	TF BS	TF BS	13 49	13 55	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323664	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323664	TF BS	TF BS	13 49	13 55	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323663	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA

ZEB1 NM_01323663	TF BS	TF BS	13 49	13 55	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323662	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323662	TF BS	TF BS	13 49	13 55	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323661	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323660	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323659	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01174094	TF BS	TF BS	22 41	22 47	0.9963 05873	+	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01174094	TF BS	TF BS	13 49	13 55	0.9963 05873	-	.	TF=FOXL1;class=Fork head / winged helix factors;sequence=ATA AACA
ZEB1 NM_01323647	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323641	TF BS	TF BS	67 4	68 1	0.9962 0351	-	.	TF=HOXA4;class=Homeo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323638	TF BS	TF BS	10 75	10 82	0.9962 0351	-	.	TF=HOXA4;class=Homeo domain factors;sequence=ATC ATTAG

ZEB1 NM_01323644	TF BS	TF BS	67 4	68 1	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323643	TF BS	TF BS	10 75	10 82	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323674	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323650	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323648	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01174095	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323646	TF BS	TF BS	67 4	68 1	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323645	TF BS	TF BS	67 4	68 1	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01174093	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323655	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323654	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG

ZEB1 NM_01323653	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323652	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323651	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323649	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323642	TF BS	TF BS	67 4	68 1	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01174096	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_030751	TF BS	TF BS	27 6	28 3	0.9962 0351	-	.	TF=HOXA4;class=Homo domain factors;sequence=ATC ATTAG
ZEB1 NM_01323673	TF BS	TF BS	16 67	16 74	0.9961 79597	-	.	TF=HOXD4;class=Homo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323677	TF BS	TF BS	16 67	16 74	0.9961 79597	-	.	TF=HOXD4;class=Homo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323676	TF BS	TF BS	16 67	16 74	0.9961 79597	-	.	TF=HOXD4;class=Homo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323675	TF BS	TF BS	16 67	16 74	0.9961 79597	-	.	TF=HOXD4;class=Homo domain factors;sequence=ATA ATTAA

ZEB1 NM_01323672	TF BS	TF BS	16 67	16 74	0.9961 79597	-	.	TF=HOXD4;class=Homo domain factors;sequence=ATAATTAA
ZEB1 NM_01323671	TF BS	TF BS	16 67	16 74	0.9961 79597	-	.	TF=HOXD4;class=Homo domain factors;sequence=ATAATTAA
ZEB1 NM_01128128	TF BS	TF BS	19 97	20 07	0.9960 93368	-	.	TF=HOXA10;class=Homo domain factors;sequence=GGCAATAAAAA
ZEB1 NM_01323678	TF BS	TF BS	19 97	20 07	0.9960 93368	-	.	TF=HOXA10;class=Homo domain factors;sequence=GGCAATAAAAA
ZEB1 NM_01323666	TF BS	TF BS	19 97	20 07	0.9960 93368	-	.	TF=HOXA10;class=Homo domain factors;sequence=GGCAATAAAAA
ZEB1 NM_01323665	TF BS	TF BS	19 97	20 07	0.9960 93368	-	.	TF=HOXA10;class=Homo domain factors;sequence=GGCAATAAAAA
ZEB1 NM_01323664	TF BS	TF BS	19 97	20 07	0.9960 93368	-	.	TF=HOXA10;class=Homo domain factors;sequence=GGCAATAAAAA
ZEB1 NM_01323663	TF BS	TF BS	19 97	20 07	0.9960 93368	-	.	TF=HOXA10;class=Homo domain factors;sequence=GGCAATAAAAA
ZEB1 NM_01323662	TF BS	TF BS	19 97	20 07	0.9960 93368	-	.	TF=HOXA10;class=Homo domain factors;sequence=GGCAATAAAAA
ZEB1 NM_01174094	TF BS	TF BS	19 97	20 07	0.9960 93368	-	.	TF=HOXA10;class=Homo domain factors;sequence=GGCAATAAAAA
ZEB1 NM_01323647	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homo domain factors;sequence=GCAATAAAAT

ZEB1 NM_01323674	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01323650	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01323648	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01174095	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01174093	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01323655	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01323654	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01323653	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01323652	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01323651	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT
ZEB1 NM_01323649	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAT

ZEB1 NM_001174096	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAAT
ZEB1 NM_0030751	TF BS	TF BS	19 97	20 06	0.9957 77597	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAAT
ZEB1 NM_001128128	TF BS	TF BS	19 97	20 06	0.9954 82444	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAA
ZEB1 NM_001323678	TF BS	TF BS	19 97	20 06	0.9954 82444	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAA
ZEB1 NM_001323666	TF BS	TF BS	19 97	20 06	0.9954 82444	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAA
ZEB1 NM_001323665	TF BS	TF BS	19 97	20 06	0.9954 82444	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAA
ZEB1 NM_001323664	TF BS	TF BS	19 97	20 06	0.9954 82444	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAA
ZEB1 NM_001323663	TF BS	TF BS	19 97	20 06	0.9954 82444	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAA
ZEB1 NM_001323662	TF BS	TF BS	19 97	20 06	0.9954 82444	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAA
ZEB1 NM_001174094	TF BS	TF BS	19 97	20 06	0.9954 82444	-	.	TF=HOXD9;class=Homeo domain factors;sequence=GCAATAAAAA
ZEB1 NM_001128128	TF BS	TF BS	13 69	13 77	0.9953 88085	+	.	TF=NFIX;class=SMAD/NF-1 DNA-binding domain factors;sequence=TGT

ZEB1 NM_01323673	TF BS	TF BS	91 0	91 8	0.9953 88085	+	.	GCCAAG TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323678	TF BS	TF BS	13 69	13 77	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323677	TF BS	TF BS	91 0	91 8	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323676	TF BS	TF BS	91 0	91 8	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323675	TF BS	TF BS	91 0	91 8	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323672	TF BS	TF BS	91 0	91 8	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323671	TF BS	TF BS	91 0	91 8	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323666	TF BS	TF BS	13 69	13 77	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323665	TF BS	TF BS	13 69	13 77	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain

ZEB1 NM_01323664	TF BS	TF BS	13 69	13 77	0.9953 88085	+	.	factors;sequence=TGT GCCAAG TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323663	TF BS	TF BS	13 69	13 77	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323662	TF BS	TF BS	13 69	13 77	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01174094	TF BS	TF BS	13 69	13 77	0.9953 88085	+	.	TF=NFIX;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAG
ZEB1 NM_01323638	TF BS	TF BS	39 5	40 2	0.9953 249	-	.	TF=HOXA4;class=Ho meo domain factors;sequence=GTC ATTAA
ZEB1 NM_01323643	TF BS	TF BS	39 5	40 2	0.9953 249	-	.	TF=HOXA4;class=Ho meo domain factors;sequence=GTC ATTAA
ZEB1 NM_01323647	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_01323641	TF BS	TF BS	13 69	13 75	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_01323638	TF BS	TF BS	17 70	17 76	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_01323644	TF BS	TF BS	13 69	13 75	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain

								factors;sequence=CTG ACAG
ZEB1 NM_0 01323643	TF BS	TF BS	17 70	17 76	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323674	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323658	TF BS	TF BS	41 9	42 5	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323657	TF BS	TF BS	41 9	42 5	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323650	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323648	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01174095	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323646	TF BS	TF BS	13 69	13 75	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323645	TF BS	TF BS	13 69	13 75	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01174093	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323656	TF BS	TF BS	41 9	42 5	0.9949 89704	-	.	TF=MEIS1;class=Hom eo domain

									factors;sequence=CTG ACAG
ZEB1 NM_0 01323655	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323654	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323653	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323652	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323651	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323649	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323642	TF BS	TF BS	13 69	13 75	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01174096	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 30751	TF BS	TF BS	97 1	97 7	0.9949 89704	-	.		TF=MEIS1;class=Hom eo domain factors;sequence=CTG ACAG
ZEB1 NM_0 01323661	TF BS	TF BS	19 97	20 07	0.9949 04316	-	.		TF=CDX1;class=Home o domain factors;sequence=GGC AATAAAAC
ZEB1 NM_0 01323660	TF BS	TF BS	19 97	20 07	0.9949 04316	-	.		TF=CDX1;class=Home o domain

ZEB1 NM_01323659	TF BS	TF BS	19 97	20 07	0.9949 04316	-	.	factors;sequence=GGC AATAAAAC TF=CDX1;class=Home o domain factors;sequence=GGC AATAAAAC
ZEB1 NM_01323673	TF BS	TF BS	16 67	16 74	0.9940 71307	-	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323677	TF BS	TF BS	16 67	16 74	0.9940 71307	-	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323676	TF BS	TF BS	16 67	16 74	0.9940 71307	-	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323675	TF BS	TF BS	16 67	16 74	0.9940 71307	-	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323672	TF BS	TF BS	16 67	16 74	0.9940 71307	-	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323671	TF BS	TF BS	16 67	16 74	0.9940 71307	-	.	TF=HOXC4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_01323647	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_01128128	TF BS	TF BS	20 9	21 4	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_01128128	TF BS	TF BS	44 5	45 0	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_01323641	TF BS	TF BS	15 3	15 8	0.9940 48003	+	.	TF=ZNF354C;class=C2 H2 zinc finger

								factors;sequence=CTC CAC
ZEB1 NM_0 01323638	TF BS	TF BS	55 4	55 9	0.9940 48003	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323644	TF BS	TF BS	15 3	15 8	0.9940 48003	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323643	TF BS	TF BS	55 4	55 9	0.9940 48003	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323674	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323673	TF BS	TF BS	17 43	17 48	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323673	TF BS	TF BS	17 71	17 76	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323658	TF BS	TF BS	13 89	13 94	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323658	TF BS	TF BS	16 42	16 47	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323658	TF BS	TF BS	18 78	18 83	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323657	TF BS	TF BS	13 89	13 94	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323657	TF BS	TF BS	16 42	16 47	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger

								factors;sequence=CTC CAC
ZEB1 NM_0 01323657	TF BS	TF BS	18 78	18 83	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323650	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323648	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01174095	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323646	TF BS	TF BS	15 3	15 8	0.9940 48003	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323645	TF BS	TF BS	15 3	15 8	0.9940 48003	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01174093	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323656	TF BS	TF BS	13 89	13 94	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323656	TF BS	TF BS	16 42	16 47	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323656	TF BS	TF BS	18 78	18 83	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323655	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger

								factors;sequence=CTC CAC
ZEB1 NM_0 01323654	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323653	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323652	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323651	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323678	TF BS	TF BS	20 9	21 4	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323678	TF BS	TF BS	44 5	45 0	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323677	TF BS	TF BS	17 43	17 48	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323677	TF BS	TF BS	17 71	17 76	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323676	TF BS	TF BS	17 43	17 48	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323676	TF BS	TF BS	17 71	17 76	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323649	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger

								factors;sequence=CTC CAC
ZEB1 NM_0 01323675	TF BS	TF BS	17 43	17 48	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323675	TF BS	TF BS	17 71	17 76	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323672	TF BS	TF BS	17 43	17 48	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323672	TF BS	TF BS	17 71	17 76	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323671	TF BS	TF BS	17 43	17 48	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323671	TF BS	TF BS	17 71	17 76	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323666	TF BS	TF BS	20 9	21 4	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323666	TF BS	TF BS	44 5	45 0	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323642	TF BS	TF BS	15 3	15 8	0.9940 48003	+	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01174096	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 30751	TF BS	TF BS	19 41	19 46	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger

								factors;sequence=CTC CAC
ZEB1 NM_0 01323665	TF BS	TF BS	20 9	21 4	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323665	TF BS	TF BS	44 5	45 0	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323664	TF BS	TF BS	20 9	21 4	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323664	TF BS	TF BS	44 5	45 0	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323663	TF BS	TF BS	20 9	21 4	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323663	TF BS	TF BS	44 5	45 0	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323662	TF BS	TF BS	20 9	21 4	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323662	TF BS	TF BS	44 5	45 0	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323661	TF BS	TF BS	60 3	60 8	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323661	TF BS	TF BS	85 6	86 1	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323661	TF BS	TF BS	10 92	10 97	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger

								factors;sequence=CTC CAC
ZEB1 NM_0 01323660	TF BS	TF BS	60 3	60 8	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323660	TF BS	TF BS	85 6	86 1	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323660	TF BS	TF BS	10 92	10 97	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323659	TF BS	TF BS	60 3	60 8	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323659	TF BS	TF BS	85 6	86 1	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323659	TF BS	TF BS	10 92	10 97	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01174094	TF BS	TF BS	20 9	21 4	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01174094	TF BS	TF BS	44 5	45 0	0.9940 48003	-	.	TF=ZNF354C;class=C2 H2 zinc finger factors;sequence=CTC CAC
ZEB1 NM_0 01323661	TF BS	TF BS	19 97	20 07	0.9939 47523	-	.	TF=HOXD13;class=Ho meo domain factors;sequence=GGC AATAAAAC
ZEB1 NM_0 01323660	TF BS	TF BS	19 97	20 07	0.9939 47523	-	.	TF=HOXD13;class=Ho meo domain factors;sequence=GGC AATAAAAC
ZEB1 NM_0 01323659	TF BS	TF BS	19 97	20 07	0.9939 47523	-	.	TF=HOXD13;class=Ho meo domain

ZEB1 NM_001128128	TF BS	TF BS	29 6	30 8	0.9935 09177	+	.	factors;sequence=GGC AATAAAAC TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323658	TF BS	TF BS	17 29	17 41	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323657	TF BS	TF BS	17 29	17 41	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323656	TF BS	TF BS	17 29	17 41	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323678	TF BS	TF BS	29 6	30 8	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323666	TF BS	TF BS	29 6	30 8	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323665	TF BS	TF BS	29 6	30 8	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323664	TF BS	TF BS	29 6	30 8	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323663	TF BS	TF BS	29 6	30 8	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323662	TF BS	TF BS	29 6	30 8	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger factors;sequence=GAA CCGTTATGTC
ZEB1 NM_001323661	TF BS	TF BS	94 3	95 5	0.9935 09177	+	.	TF=OVOL2;class=C2H 2 zinc finger

ZEB1 NM_01323660	TF BS	TF BS	94 3	95 5	0.9935 09177	+	.	factors;sequence=GAA CCGTTATGTC TF=OVOL2;class=C2H 2 zinc finger
ZEB1 NM_01323659	TF BS	TF BS	94 3	95 5	0.9935 09177	+	.	factors;sequence=GAA CCGTTATGTC TF=OVOL2;class=C2H 2 zinc finger
ZEB1 NM_01174094	TF BS	TF BS	29 6	30 8	0.9935 09177	+	.	factors;sequence=GAA CCGTTATGTC TF=OVOL2;class=C2H 2 zinc finger
ZEB1 NM_01323673	TF BS	TF BS	19 59	19 68	0.9929 58964	-	.	factors;sequence=GAA CCGTTATGTC TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=CAC ACATGTC
ZEB1 NM_01323677	TF BS	TF BS	19 59	19 68	0.9929 58964	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=CAC ACATGTC
ZEB1 NM_01323676	TF BS	TF BS	19 59	19 68	0.9929 58964	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=CAC ACATGTC
ZEB1 NM_01323675	TF BS	TF BS	19 59	19 68	0.9929 58964	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=CAC ACATGTC
ZEB1 NM_01323672	TF BS	TF BS	19 59	19 68	0.9929 58964	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=CAC ACATGTC
ZEB1 NM_01323671	TF BS	TF BS	19 59	19 68	0.9929 58964	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=CAC ACATGTC
ZEB1 NM_01323638	TF BS	TF BS	39 5	40 2	0.9928 97745	+	.	TF=HOXB6;class=Ho meo domain factors;sequence=TTA ATGAC
ZEB1 NM_01323643	TF BS	TF BS	39 5	40 2	0.9928 97745	+	.	TF=HOXB6;class=Ho meo domain

								factors;sequence=TTA ATGAC
ZEB1 NM_0 01323673	TF BS	TF BS	16 67	16 74	0.9928 87021	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_0 01323677	TF BS	TF BS	16 67	16 74	0.9928 87021	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_0 01323676	TF BS	TF BS	16 67	16 74	0.9928 87021	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_0 01323675	TF BS	TF BS	16 67	16 74	0.9928 87021	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_0 01323672	TF BS	TF BS	16 67	16 74	0.9928 87021	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_0 01323671	TF BS	TF BS	16 67	16 74	0.9928 87021	-	.	TF=HOXB4;class=Ho meo domain factors;sequence=ATA ATTAA
ZEB1 NM_0 01128128	TF BS	TF BS	14 49	14 58	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_0 01323673	TF BS	TF BS	99 0	99 9	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_0 01323678	TF BS	TF BS	14 49	14 58	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_0 01323677	TF BS	TF BS	99 0	99 9	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_0 01323676	TF BS	TF BS	99 0	99 9	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors

ZEB1 NM_01323675	TF BS	TF BS	99 0	99 9	0.9925 53464	-	.	(bHLH);sequence=ACC ACATGTT TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01323672	TF BS	TF BS	99 0	99 9	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01323671	TF BS	TF BS	99 0	99 9	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01323666	TF BS	TF BS	14 49	14 58	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01323665	TF BS	TF BS	14 49	14 58	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01323664	TF BS	TF BS	14 49	14 58	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01323663	TF BS	TF BS	14 49	14 58	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01323662	TF BS	TF BS	14 49	14 58	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01174094	TF BS	TF BS	14 49	14 58	0.9925 53464	-	.	TF=MXI1;class=Basic helix-loop-helix factors (bHLH);sequence=ACC ACATGTT
ZEB1 NM_01323647	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0	TF	TF	48	49	0.9923	+	.	TF=NR2C1;class=Nucl

01323641	BS	BS	3	1	68199			ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0 01323638	TF BS	TF BS	88 4	89 2	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0 01323644	TF BS	TF BS	48 3	49 1	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0 01323643	TF BS	TF BS	88 4	89 2	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0 01323674	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0 01323650	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0 01323648	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0 01174095	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_0 01323646	TF BS	TF BS	48 3	49 1	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA

ZEB1 NM_01323645	TF BS	TF BS	48 3	49 1	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGGGTCAA
ZEB1 NM_01174093	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGGGTCAA
ZEB1 NM_01323655	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGGGTCAA
ZEB1 NM_01323654	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGGGTCAA
ZEB1 NM_01323653	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGGGTCAA
ZEB1 NM_01323652	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGGGTCAA
ZEB1 NM_01323651	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGGGTCAA
ZEB1 NM_01323649	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGGGTCAA
ZEB1 NM_01323642	TF BS	TF BS	48 3	49 1	0.9923 68199	+	.	TF=NR2C1;class=Nuclear receptors with C4 zinc fingers;sequence=TGG

								GGTCAA
ZEB1 NM_01174096	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_030751	TF BS	TF BS	85	93	0.9923 68199	+	.	TF=NR2C1;class=Nucl ear receptors with C4 zinc fingers;sequence=TGG GGTCAA
ZEB1 NM_01128128	TF BS	TF BS	29	36	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_01323658	TF BS	TF BS	14 62	14 69	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_01323657	TF BS	TF BS	14 62	14 69	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_01323656	TF BS	TF BS	14 62	14 69	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_01323678	TF BS	TF BS	29	36	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_01323666	TF BS	TF BS	29	36	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_01323665	TF BS	TF BS	29	36	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_01323664	TF BS	TF BS	29	36	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_0	TF	TF	29	36	0.9920	-	.	TF=RHOXF1;class=Ho

01323663	BS	BS			88812			meo domain factors;sequence=ATG ATCCT
ZEB1 NM_0 01323662	TF BS	TF BS	29 6	36 3	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_0 01323661	TF BS	TF BS	67 6	68 3	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_0 01323660	TF BS	TF BS	67 6	68 3	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_0 01323659	TF BS	TF BS	67 6	68 3	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_0 01174094	TF BS	TF BS	29 6	36 3	0.9920 88812	-	.	TF=RHOXF1;class=Ho meo domain factors;sequence=ATG ATCCT
ZEB1 NM_0 01128128	TF BS	TF BS	19 97	20 08	0.9920 60007	-	.	TF=CDX2;class=Home o domain factors;sequence=AGG CAATAAAAA
ZEB1 NM_0 01323678	TF BS	TF BS	19 97	20 08	0.9920 60007	-	.	TF=CDX2;class=Home o domain factors;sequence=AGG CAATAAAAA
ZEB1 NM_0 01323666	TF BS	TF BS	19 97	20 08	0.9920 60007	-	.	TF=CDX2;class=Home o domain factors;sequence=AGG CAATAAAAA
ZEB1 NM_0 01323665	TF BS	TF BS	19 97	20 08	0.9920 60007	-	.	TF=CDX2;class=Home o domain factors;sequence=AGG CAATAAAAA
ZEB1 NM_0 01323664	TF BS	TF BS	19 97	20 08	0.9920 60007	-	.	TF=CDX2;class=Home o domain factors;sequence=AGG CAATAAAAA
ZEB1 NM_0	TF	TF	19	20	0.9920	-	.	TF=CDX2;class=Home

01323663	BS	BS	97	08	60007			o domain factors;sequence=AGG CAATAAAAA
ZEB1 NM_0 01323662	TF BS	TF BS	19 97	20 08	0.9920 60007	-	.	TF=CDX2;class=Home o domain factors;sequence=AGG CAATAAAAA
ZEB1 NM_0 01174094	TF BS	TF BS	19 97	20 08	0.9920 60007	-	.	TF=CDX2;class=Home o domain factors;sequence=AGG CAATAAAAA
ZEB1 NM_0 01323641	TF BS	TF BS	86	93	0.9919 57077	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AGG GTCAG
ZEB1 NM_0 01323638	TF BS	TF BS	48 7	49 4	0.9919 57077	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AGG GTCAG
ZEB1 NM_0 01323644	TF BS	TF BS	86	93	0.9919 57077	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AGG GTCAG
ZEB1 NM_0 01323643	TF BS	TF BS	48 7	49 4	0.9919 57077	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AGG GTCAG
ZEB1 NM_0 01323646	TF BS	TF BS	86	93	0.9919 57077	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AGG GTCAG
ZEB1 NM_0 01323645	TF BS	TF BS	86	93	0.9919 57077	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc fingers;sequence=AGG GTCAG
ZEB1 NM_0 01323642	TF BS	TF BS	86	93	0.9919 57077	+	.	TF=NR2C2(var.2);class =Nuclear receptors with C4 zinc

								fingers;sequence=AGG GTCAG
ZEB1 NM_0 01323647	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01323674	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01323650	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01323648	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01174095	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01174093	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01323655	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01323654	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01323653	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01323652	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT
ZEB1 NM_0 01323651	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	TF=CDX4;class=Home o domain

ZEB1 NM_01323649	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	factors;sequence=GGC AATAAAAT TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT TF=CDX4;class=Home o domain
ZEB1 NM_01174096	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	factors;sequence=GGC AATAAAAT TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT TF=CDX4;class=Home o domain
ZEB1 NM_030751	TF BS	TF BS	19 97	20 07	0.9918 7081	-	.	factors;sequence=GGC AATAAAAT TF=CDX4;class=Home o domain factors;sequence=GGC AATAAAAT TF=CDX4;class=Home o domain
ZEB1 NM_01323647	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	factors;sequence=GGC AATAAAAT TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_01323641	TF BS	TF BS	66 4	67 1	0.9915 28411	+	.	factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain
ZEB1 NM_01323638	TF BS	TF BS	10 65	10 72	0.9915 28411	+	.	factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain
ZEB1 NM_01323644	TF BS	TF BS	66 4	67 1	0.9915 28411	+	.	factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain
ZEB1 NM_01323643	TF BS	TF BS	10 65	10 72	0.9915 28411	+	.	factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain
ZEB1 NM_01323674	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain
ZEB1 NM_01323673	TF BS	TF BS	16 40	16 47	0.9915 28411	+	.	factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain
ZEB1 NM_01323650	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	factors;sequence=ATC ATTAA TF=HOXA4;class=Ho meo domain

								factors;sequence=ATC ATTAA
ZEB1 NM_0 01323648	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01174095	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323646	TF BS	TF BS	66 4	67 1	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323645	TF BS	TF BS	66 4	67 1	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01174093	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323655	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323654	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323653	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323652	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323651	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323677	TF BS	TF BS	16 40	16 47	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain

								factors;sequence=ATC ATTAA
ZEB1 NM_0 01323676	TF BS	TF BS	16 40	16 47	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323649	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323675	TF BS	TF BS	16 40	16 47	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323672	TF BS	TF BS	16 40	16 47	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323671	TF BS	TF BS	16 40	16 47	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323642	TF BS	TF BS	66 4	67 1	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01174096	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 30751	TF BS	TF BS	26 6	27 3	0.9915 28411	+	.	TF=HOXA4;class=Ho meo domain factors;sequence=ATC ATTAA
ZEB1 NM_0 01323638	TF BS	TF BS	39 5	40 2	0.9915 24102	-	.	TF=GSX2;class=Home o domain factors;sequence=GTC ATTAA
ZEB1 NM_0 01323643	TF BS	TF BS	39 5	40 2	0.9915 24102	-	.	TF=GSX2;class=Home o domain factors;sequence=GTC ATTAA
ZEB1 NM_0 01323647	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc

									finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01128128	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323641	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323638	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323644	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323643	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323674	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323673	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323658	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0	TF	TF	23	23	0.9914	-	.		TF=THAP1;class=C2C

01323657	BS	BS	25	33	88497				H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323650	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323648	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01174095	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323646	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323645	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01174093	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323656	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323655	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA

ZEB1 NM_01323654	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323653	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323652	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323651	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323678	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323677	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323676	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323649	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323675	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG

									CCCTCA
ZEB1 NM_01323672	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323671	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323666	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323642	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01174096	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_30751	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323665	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323664	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_01323663	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.		TF=THAP1;class=C2C H THAP-type zinc finger

								factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323662	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323661	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323660	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323659	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01174094	TF BS	TF BS	23 25	23 33	0.9914 88497	-	.	TF=THAP1;class=C2C H THAP-type zinc finger factors;sequence=CTG CCCTCA
ZEB1 NM_0 01323673	TF BS	TF BS	16 66	16 75	0.9911 27936	-	.	TF=MNX1;class=Home o domain factors;sequence=AAT AATTAAA
ZEB1 NM_0 01323677	TF BS	TF BS	16 66	16 75	0.9911 27936	-	.	TF=MNX1;class=Home o domain factors;sequence=AAT AATTAAA
ZEB1 NM_0 01323676	TF BS	TF BS	16 66	16 75	0.9911 27936	-	.	TF=MNX1;class=Home o domain factors;sequence=AAT AATTAAA
ZEB1 NM_0 01323675	TF BS	TF BS	16 66	16 75	0.9911 27936	-	.	TF=MNX1;class=Home o domain factors;sequence=AAT AATTAAA
ZEB1 NM_0	TF	TF	16	16	0.9911	-	.	TF=MNX1;class=Home

01323672	BS	BS	66	75	27936			o domain factors;sequence=AAT AATTAAA
ZEB1 NM_0 01323671	TF BS	TF BS	16 66	16 75	0.9911 27936	-	.	TF=MNX1;class=Home o domain factors;sequence=AAT AATTAAA
ZEB1 NM_0 01128128	TF BS	TF BS	13 69	13 78	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323673	TF BS	TF BS	91 0	91 9	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323678	TF BS	TF BS	13 69	13 78	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323677	TF BS	TF BS	91 0	91 9	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323676	TF BS	TF BS	91 0	91 9	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323675	TF BS	TF BS	91 0	91 9	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323672	TF BS	TF BS	91 0	91 9	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323671	TF BS	TF BS	91 0	91 9	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding

								domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323666	TF BS	TF BS	13 69	13 78	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323665	TF BS	TF BS	13 69	13 78	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323664	TF BS	TF BS	13 69	13 78	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323663	TF BS	TF BS	13 69	13 78	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01323662	TF BS	TF BS	13 69	13 78	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01174094	TF BS	TF BS	13 69	13 78	0.9906 813	+	.	TF=NFIA;class=SMAD /NF-1 DNA-binding domain factors;sequence=TGT GCCAAGG
ZEB1 NM_0 01128128	TF BS	TF BS	19 08	19 19	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_0 01323673	TF BS	TF BS	14 49	14 60	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_0 01323678	TF BS	TF BS	19 08	19 19	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC

ZEB1 NM_01323677	TF BS	TF BS	14 49	14 60	0.9904 64788	+	.	CCCACCCCC TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323676	TF BS	TF BS	14 49	14 60	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323675	TF BS	TF BS	14 49	14 60	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323672	TF BS	TF BS	14 49	14 60	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323671	TF BS	TF BS	14 49	14 60	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323666	TF BS	TF BS	19 08	19 19	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323665	TF BS	TF BS	19 08	19 19	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323664	TF BS	TF BS	19 08	19 19	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323663	TF BS	TF BS	19 08	19 19	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01323662	TF BS	TF BS	19 08	19 19	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC CCCACCCCC
ZEB1 NM_01174094	TF BS	TF BS	19 08	19 19	0.9904 64788	+	.	TF=KLF4;class=C2H2 zinc finger factors;sequence=AGC

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