

Conserved noncoding elements associated with ENSG00000125798 (FOXA2)

No.	CNE ID	Sequence 1 (fugu - FUGU4)	Sequence 2 (human - NCBI35)	Flanking genes	Location	Length (bp)	Identity (%)	Sequence	VISTA plot	Alignment	Binding sites
1	N2994	scaffold_23:403983-404135:-1	chr20:22507230-22507374:-1	ENSG00000170422; FOXA2	5'upstream 36 bp; 3'downstream 2449 bp	145	76.1	VIEW	VIEW	VIEW	VIEW
2	N2995	scaffold_23:404596-404693:-1	chr20:22507880-22507974:-1	ENSG00000170422; FOXA2	5'upstream 686 bp; 3'downstream 1849 bp	95	65.7	VIEW	VIEW	VIEW	VIEW
3	N2996	scaffold_23:405458-405555:-1	chr20:22509813-22509920:-1	ENSG00000170422; FOXA2	5'upstream 2619 bp; 3'UTR, 3'downstream 1 bp	108	65.7	VIEW	VIEW	VIEW	VIEW
4	N2997	scaffold_23:407405-407520:-1	chr20:22512555-22512669:-1	FOXA2	Intron 1	115	81.9	VIEW	VIEW	VIEW	VIEW
5	N2998	scaffold_23:407848-407936:-1	chr20:22513400-22513487:-1	FOXA2; SSTR4	5'upstream 299 bp; 5'upstream 450634 bp	88	68.5	VIEW	VIEW	VIEW	VIEW
6	N2999	scaffold_23:411020-411185:-1	chr20:22582340-22582504:-1	FOXA2; SSTR4	5'upstream 69239 bp; 5'upstream 381617 bp	165	69.9	VIEW	VIEW	VIEW	VIEW