

Figure S16. Human-mouse-fugu alignment and predicted TFBS of CNE9.

TFBS on the forward strand are shown in blue and TFBS on the reverse strand are shown in red below the human-mouse-fugu sequence alignment. The binding TF and position weight matrix similarity score are listed next to each predicted site.

human	CTGCTCATTACTTCTTCCTTCTGTATCGAGACCCTTATTTACATTTTCAGATTA	53
mouse	CTGCTCATTACTTCTTCTTTCTGTATCAAGACCCTTATTTACATTTTCAGATTA	53
fugu	CTGCTCATTAGTCTATCGCTTGGATCAAACGCTGTATTTACATATCAAATTA	53
	acccttaTTAcatttc 0.948 XFD1.01	