Figure S1: schematic alignment of brain-derived cDNAs with specific CASC1 duplicate loci

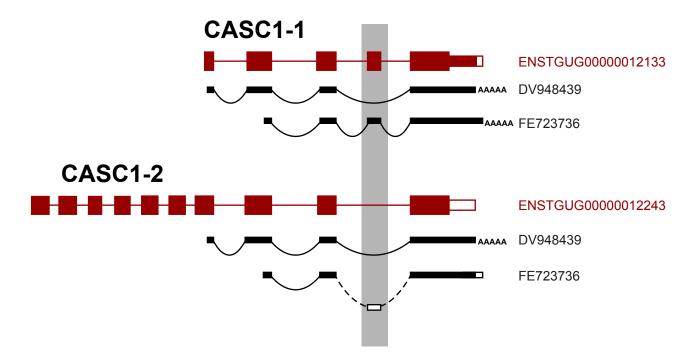


Figure S2: percent identity and schematic alignment of brain-derived cDNAs with specific YTHDC2 expanded loci

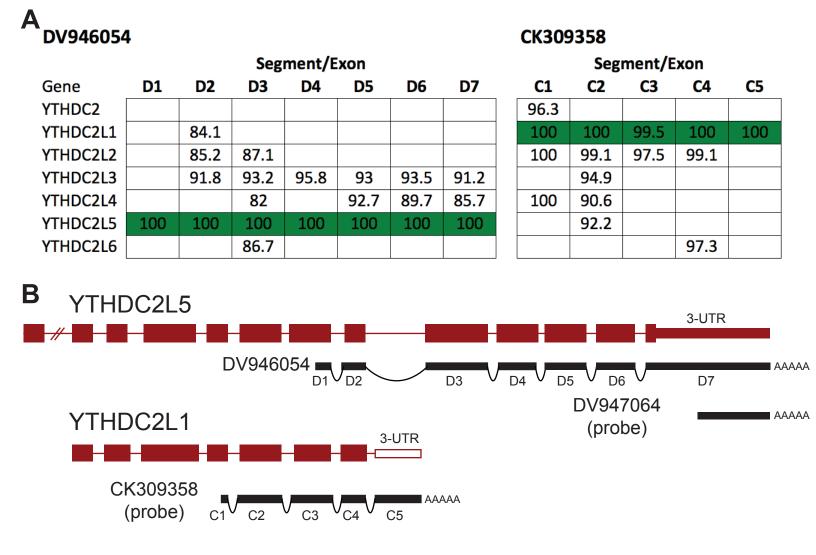


Figure S3A: example of an Ensembl gene prediction which splits a gene across two partial models

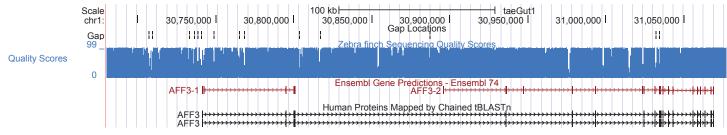


Figure S3B: example of a misannotated Ensembl gene prediction

Quality Scores	Scale chr2: Gap 99_	45,150,000 l	45,200,000 l Gap Locations Zebra finch Sequencing Quality Scores	Gut1 45,250,000	
,	0	MUTED-1	Ensembl Gene Predictions - Ensembl 74 MUTED-2	· · · · ·	
			Human Proteins Mapped by Chained tBLASTn		
		TXNDC5	MUTED K******		