

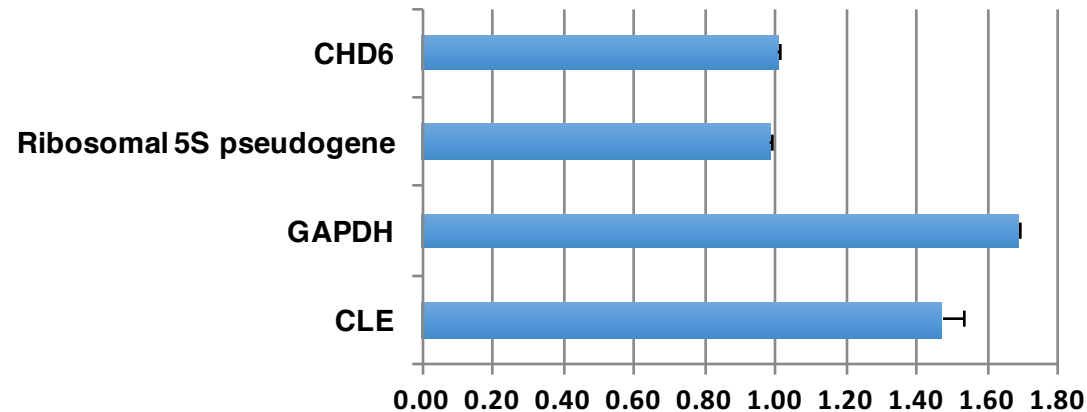
Supp. Fig. S3

A

	CHD6		Ribosomal 5S pseudogene		GAPDH		CLE	
	Mean	STDEV	Mean	STDEV	Mean	STDEV	Mean	STDEV
CBD (Cts)	26.983	0.007	18.704	0.046	11.893	0.027	24.151	0.058
CLE-CBD (Cts)	26.642	0.204	18.887	0.029	7.027	0.059	16.898	0.004
Ct Difference	0.341		-0.183		4.865		7.253	

B

Relative mRNA abundance hCLE-CBD (1/Ct) / CBD (1/Ct)



Supp. Fig. S3. Validation of RNAs bound to purified hCLE-CBD. RNAs bound to CBD or hCLE-CBD were used for qRT-PCR to detect the abundance of non-associated (CHD6 and Ribosomal 5S pseudogene) or associated (GAPDH and hCLE) mRNAs to purified CLE-CBD. A). Table showing the Ct amplification of the selected mRNAs in CBD and CLE-CBD RNAs. Mean and standard deviation of three independent measures is represented. B). Diagram showing relative abundance of each mRNAs taking as 1.00 the 1/Ct values found in the CBD sample.