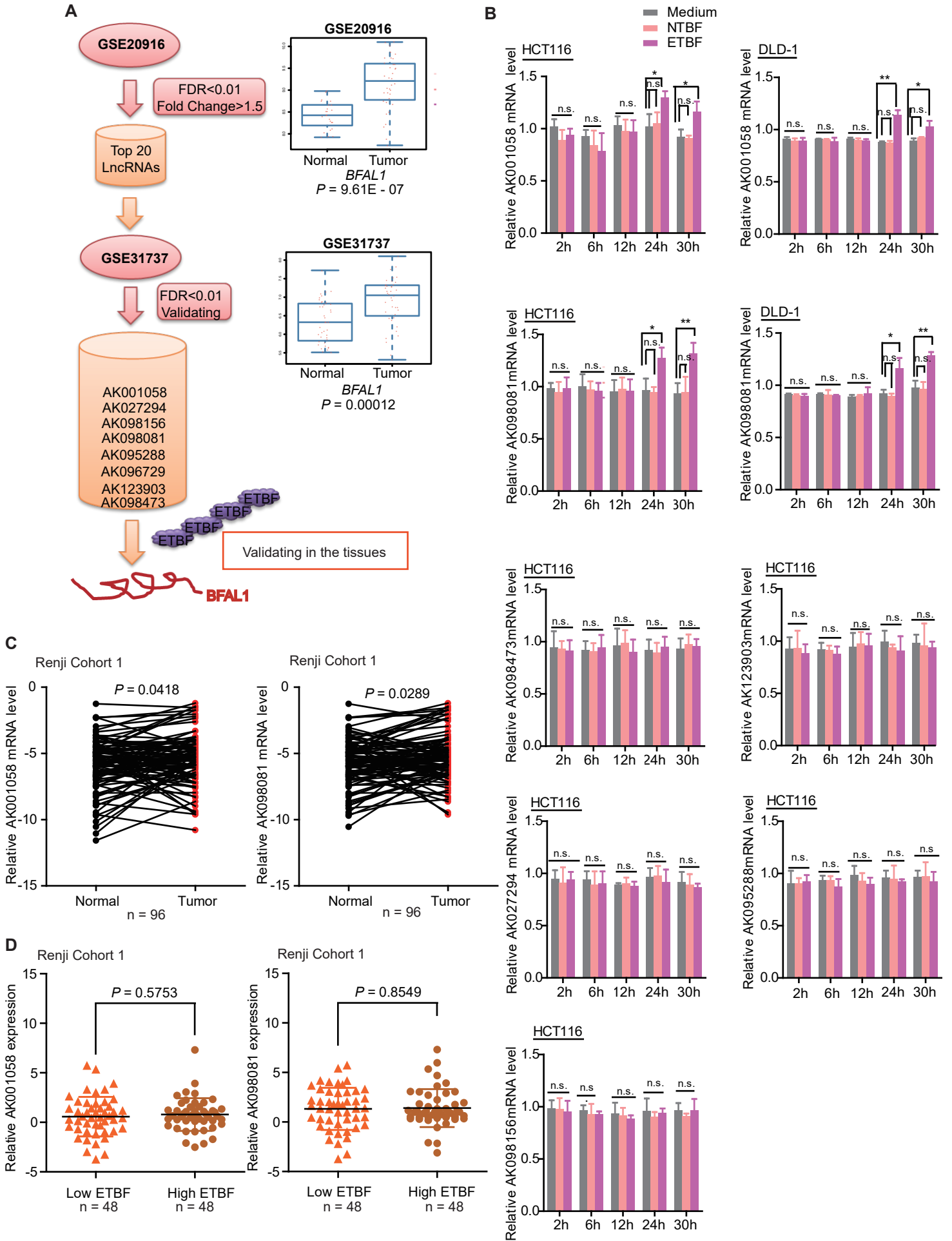


Supplementary Figure S1

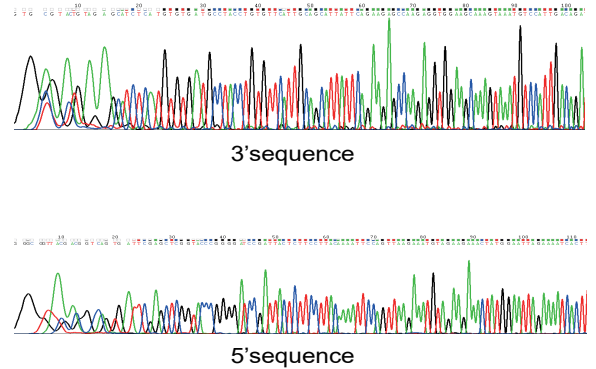
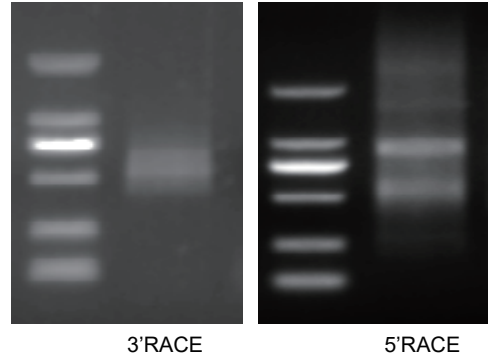


Supplementary Figure S1

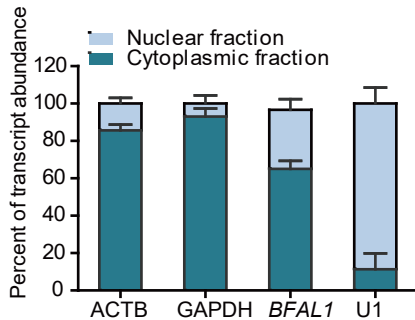
E

BFAL1 sequence

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AATAAAC CAG
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F



G

Ab Inito ncRNA transcriptome predictor

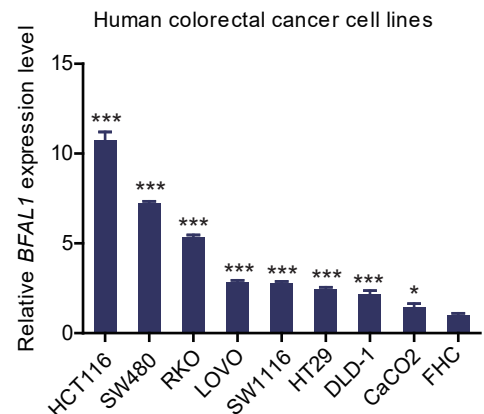
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 Predicting ncRNA.../work/ncRNA/web/ncRNA_180611744340829/Input_180611744340829.tx

Sequence name	Coding Probability	Non-coding Probability
AK096729.1	4.78%	95.22%

H

HOMOLOGY FEATURES	HIT NUM	3	
	HIT SCORE	11.365636671002	
	FRAME SCORE	43.059232312413	
ORF_FRAMEFINDER	COVERAGE	5.74 %	
	LOG-ODDS SCORE	29.77	
	TYPE	Full	
	Legend: non-coding		

I



J

Renji Cohort 1

