Wnt activated β -catenin and YAP proteins enhance the expression of non-coding RNA component of RNase MRP in colon cancer cells

Supplementary Material

Supplementary Table 1. List of RT-PCR primers

Target	Туре	Sequence		
RMRP	For	5' – CGTGCTGAAGGCCTGTATCC – 3'		
	Rev	5' – GCTTCTTGGCGGACTTTGG – 3'		
β-catenin	For	5' – CGGGATCCACAAGAAACGGCTTTCA – 3'		
	Rev	5' – GAGAATTCCAGGTCAGTATCAAACCA – 3'		
YAP	For	5' – AGCAGGATGGTGGGACTCAA – 3'		
	Rev	5' – TGAGGGCAGGGTGCTTTGG – 3'		
TBX5	For	5' – CCAGCCTAGATTACACATCGTG – 3'		
	Rev	5' – TGTGGTTCTGGTAGGAAGTCAC – 3'		
TCF4	For	5' – CAAAGACGAGGGCGAACAGG – 3'		
	Rev	5' – TGGAGATAGGTTCGGGCGGT – 3'		
Axin-2	For	5' – GCTAC AGGAAGACCCGCAG – 3'		
	Rev	5' – TAGTGGTGG TGGACATGCTTCG – 3'		
CTGF	For	5' – AAGACCTGTGGGATGGGCA – 3'		
	Rev	5' – CAAACTTGATAGGCTTGGAGATT – 3'		
TAZ	For	5' – CACCCACGATGACCCCAGA – 3'		
	Rev	5' – GCCAGGTTAGAAAGGGCTCAC – 3'		
c-myc	For	5' – AGGCTCCTGGCAAAAGGTCA – 3'		
	Rev	5' – ACATTCTCCTCGGTGTCCGA – 3'		
hTERT	For	5' – CGTCCAGTGCCAGGGG – 3'		
	Rev	5' – CGTGGGTGAGGTGAGGTGT – 3'		
GAPDH	For	5' – TGACATCAAGAAGGTGGTGA – 3'		
	Rev	5' – TCCACCACCCTGTTGCTGTA – 3'		
β2M	For	5' – CATCCATCCGACATTGAAGTTG – 3'		
	Rev	5' – ATGGTTCACACGGCAGGCA – 3'		

Supplementary Table 2. List of northern probes

RNA	Location	Length	C omunes	
name	(nt)	(nt)	Sequence	
RMRP	5 - 266	262	CGTGCTGAAGGCCTGTATCCTAGGCTACACA	
			CTGAGGACTCTGTTCCTCCCCTTTCCGCCTAG	
			GGGAAAGTCCCCGGACCTCGGGCAGAGAGT	
			GCCACGTGCATACGCACGTAGACATTCCCCG	
			CTTCCCACTCCAAAGTCCGCCAAGAAGCGTAT	
			CCCGCTGAGCGGCGTGGCGCGGGGGGCGTCA	
			TCCGTCAGCTCCCTCTAGTTACGCAGGCAGT	
			GCGTGTCCGCGCACCAACCACGGGGGCTCA	
			TTCTCAGCGCGGCT	
5.8S rRNA	2 - 156	155	ACTCTTAGCGGTGGATCACTCGGCTCGTGCG	
			TCGATGAAGAACGCAGCTAGCTGCGAGAATTA	
			ATGTGAATTGCAGGACACATTGATCATCGACA	
			CTTCGAACGCACTTGCGGCCCCGGGTTCCTC	
			CCGGGGCTACGCCTGTCTGAGCGTCGCTT	
U6 snRNA	1 - 106	106	GTGCTCGCTTCGGCAGCACATATACTAAAATT	
			GGAACGATACAGAGAAGATTAGCATGGCCCC	
			TGCGCAAGGATGACACGCAAATTCGTGAAGC	
			GTTCCATATTTT	
5S rRNA	1 - 120	120	GTCTACGGCCATACCACCCTGAACGCGCCCG	
			ATCTCGTCTGATCTCGGAAGCTAAGCAGGGTC	
			GGGCCTGGTTAGTACTTGGATGGGAGACCGC	
			CTGGGAATACCGGGTGCTGTAGGCTT	



Supplementary Figure 1. *RMRP* expression in liver cancer patient tissues.

RT-PCR analysis of *RMRP* expression levels in liver tumor patient tissues. A bar graph shows the relative level in tumor tissues versus normal tissues. P: patient, N: normal tissue, T: tumor tissue.



Supplementary Figure 2. Predicted β -catenin/TCF sites in *RMRP* promoter and Wnt or LiCI activated *RMRP* expression level.

(A) Diagram of the *RMRP* promoter with three putative TBEs (TCF Binding Elements) upstream of the *RMRP* promoter. An arrow indicates the transcription start site of the *RMRP* gene. (B) Quantitative real-time PCR analysis of *RMRP* level in HEK293 cells following Wnt3a treatment for 1 hour. A bar graph shows *RMRP* level from three independent experiments. Mean and standard deviation is shown (*p < 0.01). (C) RT-PCR analysis of *RMRP* expression following lithium chloride (20 mM) treatment for 0, 0.5, 1, 3 and 6 hrs in Beas-2B cells. (D) Northern blot analysis of *RMRP* expression in HEK293 cells following lithium chloride (20 mM) treatment for 0, 0.5, 1 and 3 hrs. (E) RT-PCR analysis of *Axin-2* mRNA expression from the samples of (D).



Supplementary Figure 3. β -catenin and YAP associate to TBX5 binding sites and TATA box of *RMRP* promoter.

(A) Chromatin immunoprecipitation (ChIP) analysis using anti- β -catenin and anti-YAP antibodies in SW480 cells. TBX5 binding sites (TB5) and the TATA box of the *RMRP* promoter were tested for the recruitment of β -catenin or YAP. Recruitment of β -catenin or YAP is shown by ChIP on TEAD binding site in the *CTGF* promoter and on TBE3 (TCF Binding Elements 3) in the VEGFA promoter, respectively. (B) ChIP analysis using anti-FLAG antibodies in HEK293 cells following transfection of FLAG- β -catenin and FLAG-YAP in HEK293 cells



Supplementary Figure 4. Wnt signaling could enhance mitochondrial DNA replication.

(A) Fluorescence in situ hybridization (FISH) analysis of *RMRP* (green) after Wnt3a treatment for 1 hr in H1299 cells. Yellow arrows indicate the nucleolus of cells. (B) FISH analysis of *RMRP* (green) after Wnt3a treatment for 1 or 6 hrs in HeLa cells.

Yellow lines indicate nuclear boundaries. (C) Diagram of mitochondrial genome. PCR primers designed for 16S ribosomal RNA gene were used.