

Table SI. Primers used for reverse transcription-quantitative PCR.

Gene name	Forward primer	Reverse primer
ERCC1	5'-CTCAAGGAGCTGGCTAAGATGT-3'	5'-CATAGGCCTTGTTAGGTCTCCAG-3'
GSTP1	5'-CCCTGTACCAGTCCAATACCATC-3'	5'-GCCCCGCCTCATAGTTGGTGT-3'
ATP7B	5'-CCCTTCTTGTACCAGCCCAG-3'	5'-TCGGCTCCATCAGGAAGAGA-3'
ANXA3	5'-CCCATCAGTGGATGCTGAAG-3'	5'-TCACTAGGGCCACCATGAGA-3'
GAPDH	5'-GGTATCGTGGAAGGACTCATGAC-3'	5'-ATGCCAGTGAGCTTCCCGTTCAG-3'

ANXA3, annexin A3; ERCC1, excision repair cross-complementing 1; GSTP1, glutathione S-transferase P1.

Table SII. ANXA3 expression, as determined by IHC.

Variable	Cases, n	ANXA3 expression score				P-value ^a
		0	1	2	3	
Tissues						<0.0001
CRC	169	8	66	81	14	
NT	169	77	58	33	1	0.003
Stage						
I	13	1	10	1	1	
II	66	4	33	25	4	
III	71	3	21	43	4	
IV	19	0	2	12	5	

^aData are presented as the number of cases with specific IHC scores. For the comparison between CRC and NT groups, the Bowker's test was used. For the comparison among different clinical stages (stage I-IV), the Fisher's exact test was used. P<0.05 was considered to indicate a statistically significant difference. ANXA3, annexin A3; CRC, colorectal cancer; IHC, immunohistochemistry; NT, paracancerous normal tissues.