

Supplementary Table 1 Primer sequences of colorectal cancer 14-gene panel

Oligonucleotide	Sequence 5'↔3'	T _m
<i>CHST11</i>		
Forward primer	GTTTCGCGTTTTTCGGGTTTC	
Reverse primer	AACTTTATCCTAACTACGAAAACCG	
Probe	FAM-CGCGGCGGGGTTTTTGTTTTGCG-BHQ1	
PCO	CGCAAAATTAAAAATCCTGCTGCG-PHO	49.00
<i>ZNF671</i>		
Forward primer	GTTCGTTAGTTTCGTTATAGGATC	
Reverse primer	AATACCTACGTCCCCGATCG	
Probe	HEX-CTACCGCTCCCGACAACCTATCCGC-BHQ1	
PCO	GCGATTAGTTGTTGGGAGTGATAG-PHO	48.70
<i>VAV3-AS1</i>		
Forward primer	GATCGAGGGAGTAGGAGTC	
Reverse primer	CTACAACGACGACCGAATCG	
Probe	ROX-CGACGCCCCGCGACCCGTCAAC-BHQ2	
PCO	GTTAATGGGACGCGGGTGTCG-PHO	49.60
<i>ZEB2</i>		
Forward primer	CGATTTTATCGGATTTTCGCGC	
Reverse primer	ATAACGATCCGCGAACAACG	
Probe	FAM-CCCATCGCGCCCAAACCCCGAAA-BHQ1	
PCO	TTTCGAAGTTTGGATGTGAAGGG-PHO	35.91
<i>IKZF1</i>		
Forward primer	TTTCGTTTTTTGTATCGGAGTAGC	
Reverse primer	TACCCCGCAACCGACTCG	
Probe	HEX-CGCGCACCTCTCGACCGCCTC-BHQ1	
PCO	GAGGTGGATGAGAAGTATGCG-PHO	33.96
<i>NDRG4</i>		
Forward primer	ATTTAGTATAGTTCGCGCGGC	
Reverse primer	ACCTAAAAAACACCCCGATCG	
Probe	HEX-CCCCCGCCGACTTCTCACCCG-BHQ1	

PCO	CGGGAGAGAAGTCGGTGAGGG-PHO	57.29
<i>ITGA4</i>		
Forward primer	GGTTTCGTTTTTGTTCACGCGC	
Reverse primer	CTAAAAAACCCGAACCAAAACGC	
Probe	FAM-CAACCGAAATTCCCCAACGCCCCCT-BQH1	
PCO	AGGGAGTGTTGGGGAATTATGGTTG-PHO	55.56
<i>C9orf50</i>		
Forward primer	GATTTTTTAGTTTTTCGGAGTTTCGC	
Reverse primer	CCTAACGACGAAAACCTCCG	
Probe	HEX-CGCCCCGCGCTCCGAACGACTC-BHQ1	
PCO	GAGACGTTTGGAGTGTGGGCG-PHO	55.73
<i>BCAN</i>		
Forward primer	TACGTTCGCGTAGTTTTCGTC	
Reverse primer	CGAAATTTATCGCCTCCTCCG	
Probe	ROX-CTCCTCCGAATACACGACTCAATCCTACAA	
PCO	TTGAAGGATTGAGTTGTGTATTTGGAAGAG-PHO	55.33
<i>BMP3</i>		
Forward primer	CAACTAATTTAAAATTCAACCCTCG	
Reverse primer	GCGTCGTTGCGGGATATTTC	
Probe	ROX-AGGAGTCGGCGGCGGAGTCG-BHQ2	
PCO	CGATTCCGTTGCCGATACCT-PHO	41.07
<i>SDC2</i>		
Forward primer	CCCGAACCCCGAACCCG	
Reverse primer	TTCGGCGTAGTTCGCGTATAC	
Probe	FAM-CCCGAACCTAAACCGCAATCGCTACGA-BH	
PCO	TCGAAGCGATTGTAGTTTAGGTTTGGG-PHO	55.16
<i>SEPT9-1</i>		
Forward primer	TGGATGGGATTATTTTCGGATTTC	
Reverse primer	CGCAAATCCTCTCCAACACG	
Probe	HEX-CGACCGCAACAACCAACCCAACAC-BHQ1	
PCO	GTGTAAAGTTGATTGTAGTGGTTG-PHO	37.87

SEPT9-2

Forward primer	AGGGCGCGTAGGGTTTAC	
Reverse primer	TAATCCAAACGCCCTCCCG	
Probe	FAM-CGCCGCGCGCCCCCGAC-BHQ1	
PCO	GTCAGGGGTGTGTGGCG-PHO	45.09

SFMBT2

Forward primer	TAGAAGGGCGGGAGCGC	
Reverse primer	CCACCTTCCCTAAAACGCG	
Probe	ROX-CGTAGGGTCGAGGCGAGGGGTTG-BHQ2	
PCO	CAATTCCTCGTTTCGACCTGACG-PHO	44.80

ACTB

Forward primer	GTGATGGAGGAGGTTTAGTAAGTT	
Reverse primer	CCAATAAAACCTACTCCTCCCTTAA	
Probe	CY5-ACCACCACCCAACACACAATAACAAACAC	

Supplementary Table 2 Primer sequences used in cancerous and paracancerous tissues

Oligonucleotide	Sequence 5'↔3'
<i>CHST11</i>	
Forward primer	GTTTCGCGTTTTTCGGGTTTC
Reverse primer	AACTTTATCCTAACTACGAAAACCG
Probe	FAM-CGCGGCGGGGTTTTTGTTTTGCG-BHQ1
<i>VAV3-AS1</i>	
Forward primer	GATCGAGGGAGTAGGAGTC
Reverse primer	CTACAACGACGACCGAATCG
Probe	ROX-CGACGCCCCGCGACCCGTCAAC-BHQ2
<i>ZEB2</i>	
Forward primer	CGATTTTATCGGATTTTCGCGC
Reverse primer	ATAACGATCCGCGAACAACG
Probe	FAM-CCCATCGCGCCCAAACCCCGAAA-BHQ1
<i>IKZF1</i>	

Forward primer	TTTCGTTTTTTGTATCGGAGTAGC
Reverse primer	TACCCCGCAACCGACTCG
Probe	HEX-CGCGCACCTCTCGACCGCCTC-BHQ1
<i>ITGA4</i>	
Forward primer	GGTTTCGTTTTTTGTTTACGCGC
Reverse primer	CTAAAAAACCCGAACCAAAACGC
Probe	FAM-CAACCGAAATTCCCCAACGCCCCCT-BQH1
<i>C9orf50</i>	
Forward primer	GATTTTTTAGTTTTTCGGAGTTTCGC
Reverse primer	CCTAACGACGAAAACCTCCG
Probe	HEX-CGCCCCGCGCTCCGAACGACTC-BHQ1
<i>SEPT9-2</i>	
Forward primer	AGGGCGCGTAGGGTTTAC
Reverse primer	TAATCCAAACGCCCTCCCG
Probe	ROX-CGCCGCGCGCCCCCGAC-BHQ2
<i>SFMBT2</i>	
Forward primer	TAGAAGGGCGGGAGCGC
Reverse primer	CCACCTTCCCTAAAACGCG
Probe	ROX-CGTAGGGTTCGAGGCGAGGGGTTG-BHQ2
<i>ACTB</i>	
Forward primer	GTGATGGAGGAGGTTTAGTAAGTT
Reverse primer	CCAATAAAACCTACTCCTCCCTTAA
Probe	CY5-ACCACCACCCAACACACAATAACAAACACA-BHQ2

Supplementary Table 3 Primer sequences for colorectal cancer 8-gene outer and inner panel

Oligonucleotide	Sequence 5'↔3'
<i>CHST11</i>	
Outer forward primer	TTGTTTAGTTTTGTTTCGCGTTTTTC
Outer reverse primer	AACTTTATCCTAACTACGAAAACCG

Inner forward primer	GTTTCGCGTTTTTCGGGTTTC
Inner reverse primer	TAACTACGAAAACCGAAATATCCG
Probe	FAM-CGCGGCCGGGGTTTTTGTTTTGCG-BHQ1

VAV3-AS1

Outer forward primer	TTTTTCGCGCGGGATCGAG
Outer reverse primer	CTACAACGACGACCGAATCG
Inner forward primer	GATCGAGGGAGTAGGAGTC
Inner reverse primer	GACGACCGAATCGCGACG
Probe	ROX-CGACGCCCGCGACCCGTCAAC-BHQ2

ZEB2

Outer forward primer	GTTTTCGAGTTATTATCGATTTTATC
Outer reverse primer	ATAACGATCCGCGAACAACG
Inner forward primer	CGATTTTATCGGATTTTCGCGC
Inner reverse primer	CCGCGAACAACGAAACCCG
Probe	FAM-CCCATCGCGCCCAAACCCCGAAA-BHQ1

IKZF1

Outer forward primer	TTTGCGCGTTTCGTTTTTTGTATC
Outer reverse primer	TACCCCGCAACCGACTCG
Inner forward primer	TTTCGTTTTTTGTATCGGAGTAGC
Inner reverse primer	CAACCGACTCGACCCCG
Probe	HEX-CGCGCACCTCTCGACCGCCTC-BHQ1

ITGA4

Outer forward primer	GGTCGGGTTTCGTTTTTGTTTAC
Outer reverse primer	CTAAAAAACCCGAACCAAAACGC
Inner forward primer	GGTTTCGTTTTTGTTTACGCGC
Inner reverse primer	GAACCAAAACGCGAATCCTAC
Probe	FAM-CAACCGAAATTCCCCAACGCCCCCT-BHQ1

C9orf50

Outer forward primer	TCGTTTTTCGGGATTTTTTAGTTTTTC
Outer reverse primer	CCTAACGACGAAAACCTTCCG

Inner forward primer GATTTTTTAGTTTTTCGGAGTTTCGC
 Inner reverse primer CGACGCAACAACGACCCG
 Probe HEX-CGCCCCGCGCTCCGAACGACTC-BHQ1

SEPT9-2

Outer forward primer AGGGCGCGTAGGGTTTAC
 Outer reverse primer TAATCCAAACGCCCTCCCG
 Inner forward primer TTTTACGTAGGCGGCGCG
 Inner reverse primer CAAACGCCCTCCCGCTAAA
 Probe ROX-CGCCGCGCGCCCCCGAC-BHQ2

SFMBT2

Outer forward primer GTGGGGTAGAAGGGCGG
 Outer reverse primer CCACCTTCCCTAAAACGCG
 Inner forward primer TAGAAGGGCGGGAGCGC
 Inner reverse primer TTCCCTAAAACGCGAAACCCAAA
 Probe ROX-CGTAGGGTTCGAGGCGAGGGGTTG-BHQ2

ACTB

Forward primer GTGATGGAGGAGGTTTAGTAAGTT
 Reverse primer CCAATAAAACCTACTCCTCCCTTAA
 Probe CY5-ACCACCACCCAACACACAATAACAAACACA-BHQ2

Supplementary Table 4 2 × 2 contingency tables for the three-marker methylation panel across different subgroups

Subgroup	TP	FN	FP	TN
Overall cohort (117 CRC <i>vs</i> 128 combined controls: 68 healthy + 60 benign)	98	19	6	122
CRC <i>vs</i> healthy-only controls (117 CRC <i>vs</i> 68 healthy individuals)	98	19	0	68
CRC <i>vs</i> benign-only controls (117 CRC <i>vs</i> 60 benign gastrointestinal patients)	98	19	6	54

TP: True positive; FN: False negative; FP: False positive; TN: True negative; CRC: Colorectal cancer.