

Suppl. Table S3. Detailed comparisons between MMR genomic status and IHC.

	MLH1			PMS2			MSH2			MSH6			MLH1/PMS2			MSH2/MSH6		
		Alt	No		Alt	No		Alt	No		Alt	No		Alt	No		Alt	No
	Lost	28	10	Lost	28	10	Lost	3	6	Lost	7	7	Lost	32	19	Lost	10	6
	Pres	7	77	Pres	7	77	Pres	4	110	Pres	11	98	Pres	6	65	Pres	14	93
Correct classification proportion (%)	86.1			65.0			91.9			85.4			79.5			83.7		
Correlation (%)	0.7			0.3			0.3			0.4			0.6			0.4		
p	4.9e-13			0.00661			0.00833			0.00083			1.3e-10			5.0e-5		

Alt, Altered; Pres, preserved. p-values calculated using Fisher’s exact independence test.