

Suppl Table S5. Panels for detailed comparisons between MMR genomic status and MSI by PCR: a) classical and b) alternative groups.

a)

	MLH1		PMS2		MSH2		MSH6		Total
	No	Alt	No	Alt	No	Alt	No	Alt	
MSS	69	8	74	3	75	2	69	8	77
MSI-L+MSI-H	10	24	29	5	32	2	27	7	34
Total	79	32	103	8	107	4	96	15	111
Coincidences proportion (%)	83.8		71.2		69.4		68.5		
Correlation (%)	61.3		19.3		8.1		13.8		
p-value	3.859e-10		0.05599		0.5847		0.2262		

b)

	MLH1		PMS2		MSH2		MSH6		Total
	No	Alt	No	Alt	No	Alt	No	Alt	
MSS + MSI-L	75	12	84	3	85	2	75	12	87
MSI-H	4	20	19	5	22	2	21	3	24
Total	79	32	103	8	107	4	96	15	111
Coincidences Proportion (%)	85.6		80.2		78.4		70.3		
Correlation (%)	63.2		27.7		13.3		-1.6		
p-value	2.594e-10		0.01141		0.2036		1		

MSI-L, microsatellite instability - low; MSI-H, microsatellite instability - high; MSS, microsatellite stable.

p-values calculated using Fisher's exact independence test.