

Supplemental table 2. Number of identified proteins by iTRAQ experiments

	iTRAQ label				total	FDR ^a (%)
	114	115	116	117		
Data set	reference pool	cancer	cancer (metastasis)	polyp		
iTRAQ-1	ref. pool	C-1	Cm-1	P-1	4263	< 1
iTRAQ-2	ref. pool	C-2	Cm-2	P-2	4317	< 1
iTRAQ-3	ref. pool	C-3	Cm-3	P-3	4195	< 1
iTRAQ-4	ref. pool	C-4	Cm-4	P-4	4633	< 1
iTRAQ-5	ref. pool	C-5	Cm-5	P-5	4522	< 1
iTRAQ-6	ref. pool	C-6	Cm-6	P-6	4264	< 1
				average	4366	< 1

^a False discovery rate (FDR) was calculated by peptide sequence analysis using proteome discoverer