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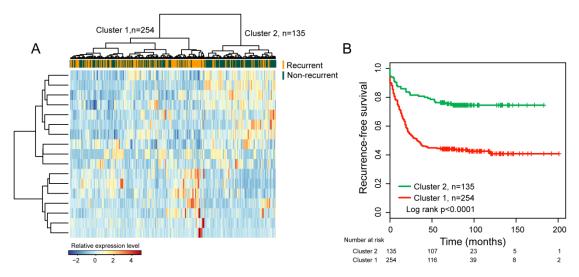
Supplemental Information

Recurrence-Associated Long Non-coding RNA

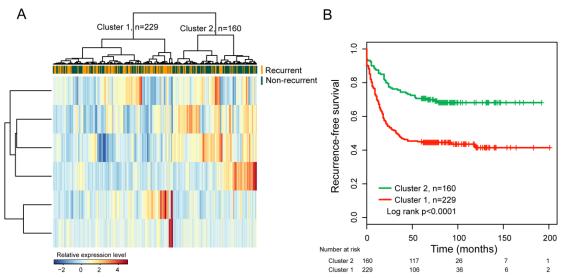
Signature for Determining the Risk of Recurrence

in Patients with Colon Cancer

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Supplemental Figure S1. (A) Unsupervised clustering of patients based on the expression pattern of 17 recurrence-associated lncRNAs. (B) Kaplan-Meier estimates of the recurrence-free survival of patients in the two sample clusters based on 17 recurrence-associated lncRNAs.



Supplemental Figure S2. (A) Unsupervised clustering of patients based on the expression pattern of six lncRNAs in the signature. Recurrence. (B) Kaplan-Meier estimates of the recurrence-free survival of patients in the two sample clusters based on six lncRNAs in the signature.

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Characteristics	Discovery	GSE39582	GSE14333	GSE17538	GSE33113	TCGA
	dataset	dataset	dataset	dataset	dataset	dataset
No. of patients	389	574	226	200	89	391
Age (years)						
Range	30.2-97	22-97	26-92	23-94	34-95	31-90
Median	68	68.1	67	66	73	67
Gender						
Female	175	257	120	98	47	181
Male	214	317	106	102	42	210
Stage						
I/II	196	304	135	134	89	227
III/IV	193	266	91	66	0	155
Unknown		4				9
Recurrence						
Yes	179	179	50	55	18	89
No	210	395	176	145	71	302
Recurrence-free						
survival (months)						
Range	0-201	0-201	0.9-142.6	0-142.6	1.8-120	0-147.9
Median	65	44.5	38.5	39	39.5	21.3

Supplemental Table S1. Clinical characteristics of patients with colon cancer in the six datasets