

Article

Establishment of highly Transplantable Cholangiocarcinoma Cell Lines from Patient Derived Xenograft Mouse Model

Table 1. Primers for CCA cell line characterization.

Genes	Primer sequence (5'→3')		Products (bp)
	Sense	Antisense	
<i>AFP</i>	TTTTGGGACCCGAACCTTCC	CTCCTGGTATCCTTTAGCAACTCT	451
<i>ALB</i>	GGTGTGATTGCCTTTGCTC	CCCTTCATCCC GAAGTTCAT	502
<i>CK7</i>	CCCTCAATGAGACGGAGTTG	GTAGGTGGCGATCTCGATGT	497
<i>CK19</i>	TCCCGCGACTACAGCCACTAC TACACGACC	CGCGACTTGATGTCCATGAGCC GCTGGTAC	746
<i>GGT</i>	ACAACAGCACCACACGAAAA	TCAGCTCAGCACGGTAGTTG	501
<i>αSMA</i>	CTTCAGCTTTCAGCTTCCCTGA	TTGCTCTGTGCTTCGTCACC	274

Table 2. Durations of the F0 tumor developments, patient survival times, percent of CK19 positive area and percent of Ki-67 positive nuclei.

Code	Survival* (days)	Duration** (days)	%CK19	%Ki-67
D049	257	58	27	40
D058	105	123	17	30
D068	212	28	37	5
D070	208	144	46	10
D078	193	163	2	10
D088	165	106	39	5
D090	161	163	39	0.1
D096	144	39	19	0.1
D113	94	152	25	0
D119	71	194	15	5
D131	31	47	63	10
D138	23	111	32	5

*Survival was calculated from the surgery to death; **Duration was calculated from implantation to removal.

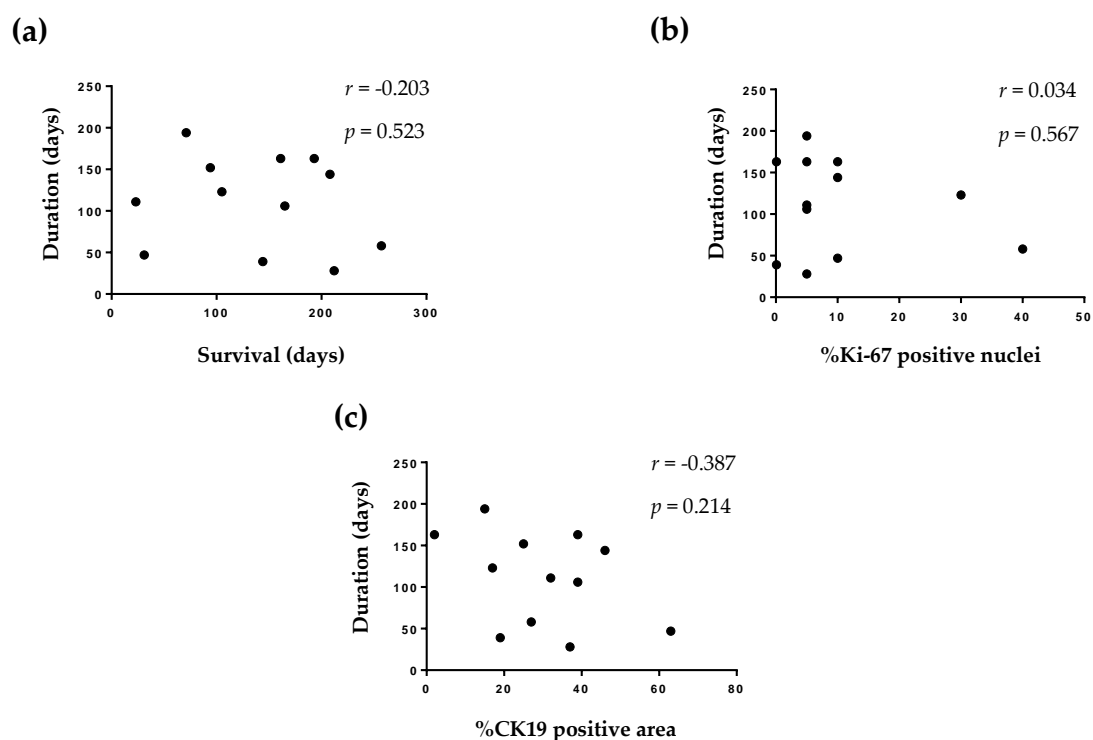


Figure 1. The correlation of the F0 tumor development and patient survival, percentage of Ki-67 positive nuclei and percentage of CK19 positive area.

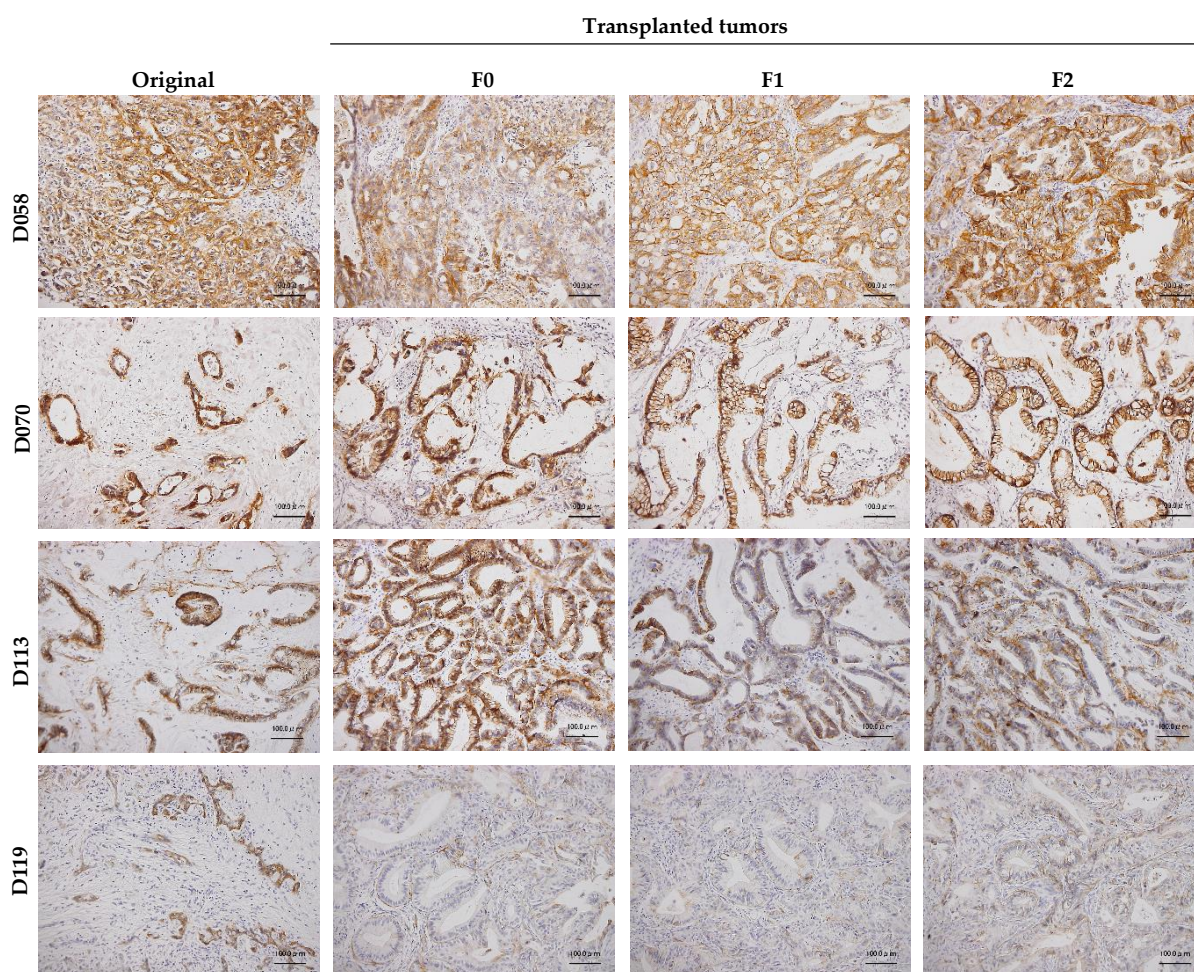


Figure 2. Comparison of EpCAM expressions between original tumor tissues from patient (original) and serially transplanted tissues (F0, F1 and F2). Bar = 100 μ m.