cancers

MDPI

Supplementary Materials: Liver Transplantation for Unresectable Intrahepatic Cholangiocarcinoma: The Role of Sequencing Genetic Profiling

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Table S1. Genes covered by SureSelect Cancer All-In-One (AIO) Solid Tumor Assay.

ABL1	*CDKN2A*	*FGFR4*	*MLH1*	*RAF1*
AKT1	*CDKN2B*	*FOXL2*	*MSH2*	*RB1*
ALK	*CIC*	*GNA11*	*MSH6*	*RET*
APC	*CSF1R*	*GNAQ*	*MTOR*	*RIT1*
AR	*CTNNB1*	*GNAS*	*MYC*	*ROS1*
ARAF	*DDR2*	*HNF1A*	*MYCN*	*SMAD4*
ARID1A	*DNMT3A*	*HRAS*	*MYD88*	*SMARCB1*
ATM	*EGFR*	*IDH1*	*NF1*	*SMO*
BCL2	*ERBB2*	*IDH2*	*NF2*	*SRC*
BCR	*ERBB3*	*JAK2*	*NFE2L2*	*STK11*
BRAF	*ERBB4*	*JAK3*	*NOTCH1*	*TERT*
BRCA1	*ESR1*	*KDR*	*NRAS*	*TMPRSS2*
BRCA2	*ETV1*	*KIT*	*NTRK1*	*TP53*
CCND1	*ETV4*	*KMT2A*	*PDGFRA*	*TSC1*
CCND2	*ETV6*	*KRAS*	*PDGFRB*	*TSC2*
CCNE1	*EZH2*	*MAP2K1*	*PIK3CA*	*VEGFA*
CD274	*FBXW7*	*MAP2K2*	*PIK3R1*	*VHL*
CDH1	*FGFR1*	*MAP2K4*	*PTCH1*	*WT1*
CDK4	*FGFR2*	*MDM2*	*PTEN*	
CDK6	*FGFR3*	*MET*	*PTPN11*	

Table S2. List of all pathogenic or likely-pathogenic mutations found in iCCA tissue samples from 12 patients who underwent liver transplantation.

Gene (Transcript)	cDNA (cNomen)	Patient	Pathway
KRAS(NM_033360.4)	c.183A>C	1	MAPK
KRAS(NM_033360.4)	c.35G>A	2	MAPK
KRAS(NM_033360.4)	c.35G>T	3	MAPK
KRAS(NM_033360.4)	c.35G>T	4	MAPK
KRAS(NM_033360.4)	c.35G>A	5	MAPK
KRAS(NM_033360.4)	c.35G>T	6	MAPK
KRAS(NM_033360.4)	c.35G>A	7	MAPK
KRAS(NM_033360.4)	c.35G>A	9	MAPK
KRAS(NM_033360.4)	c.216G>A	9	MAPK
KRAS(NM_033360.4)	c.216G>A	11	MAPK

NF1(NM_001042492.2)	c.3313A>T	2	MAPK
NF1(NM_001042492.2)	c.3144G>T	2	MAPK
NF1(NM_001042492.2)	c.808C>T	5	MAPK
NF1(NM_001042492.2)	c.4273G>T	6	MAPK
NF1(NM_001042492.2)	c.5788T>C	7	MAPK
BRAF(NM_001374258.1)	c.144G>A	1	MAPK
BRAF(NM_001374258.1)	c.1039C>T	4	MAPK
BRAF(NM_001374258.1)	c.1531G>T	9	MAPK
ERBB4(NM_005235.3)	c.1818G>T	9	MAPK
ERBB4(NM_005235.3)	c.3902A>G	12	MAPK
KIT(NM_000222.2)	c.811C>T	4	MAPK
KIT(NM_000222.2)	c.2008A>T	7	MAPK
FGFR2(NM_022970.3)	c.2063G>A	5	MAPK
FGFR3(NM_001354809.2)	c.1849A>G	12	MAPK
HRAS(NM_176795.4)	c.34G>A	5	MAPK
MAP2K4(NM_001281435.2)	c.779G>A	4	MAPK
MET(NM_000245.4)	c.3629G>A	12	MAPK
ATM(NM_000051.3)	c.283C>T	1	P53
ATM(NM_000051.3)	c.6448G>A	1	P53
ATM(NM_000051.3)	c.7472G>T	4	P53
ATM(NM_000051.3)	c.8254A>T	4	P53
ATM(NM_000051.3)	c.5908C>T	5	P53
ATM(NM_000051.3)	c.1715T>C	5	P53
ATM(NM_000051.3)	c.967A>G	7	P53
ATM(NM_000051.3)	c.5697C>A	9	P53
ATM(NM_000051.3)	c.4577C>T	9	P53
ATM(NM_000051.3)	c.4906C>T	12	P53
CDKN2A(NM_001195132.1)	c.194T>C	1	P53
CDKN2A(NM_001195132.1)	c.262G>T	2	P53
CDKN2A(NM_001195132.1)	c.194T>C	10	P53
CDKN2A(NM_001195132.1)	c.194T>C	12	P53
RB1(NM_000321.2)	c.1544C>T	6	P53
RB1(NM_000321.2)	c.2014G>T	11	P53
TP53(NM_001126114.2)	c.818G>A	10	P53
PIK3CA(NM_006218.2)	c.262C>T	2	PI3K-Akt
PIK3CA(NM_006218.2)	c.2113C>T	2	PI3K-Akt
PIK3CA(NM_006218.2)	c.2119G>A	6	PI3K-Akt
PIK3CA(NM_006218.2)	c.353G>A	7	PI3K-Akt
PIK3CA(NM_006218.2)	c.1624G>A	7	PI3K-Akt
PIK3CA(NM_006218.2)	c.1067T>C	7	PI3K-Akt
PIK3CA(NM_006218.2)	c.2119G>A	7	PI3K-Akt
PIK3CA(NM_006218.2)	c.1035T>A	9	PI3K-Akt
PTEN(NM_001304718.2)	c.435+1G>T	3	PI3K-Akt
PTEN(NM_000314.4)	c.493G>T	6	PI3K-Akt
PTEN(NM_000314.4)	c.640C>T	9	PI3K-Akt
PTEN(NM_001304717.5)	c.927G>A	12	PI3K-Akt
MTOR(NM_004958.3)	c.526G>A	5	PI3K-Akt
MTOR(NM_004958.3)	c.6577C>T	12	PI3K-Akt
APC(NM_001354895.2)	c.1500T>A	1	Wnt

APC(NM_001354895.2)	c.706C>T	3	Wnt
APC(NM_001354895.2)	c.1624C>T	3	Wnt
APC(NM_001354895.2)	c.1658G>A	4	Wnt
APC(NM_001354895.2)	c.4508C>A	4	Wnt
APC(NM_001354895.2)	c.471G>A	6	Wnt
CTNNB1(NM_001904.4)	c.1040T>C	4	Wnt
CTNNB1(NM_001904.4)	c.1628C>T	5	Wnt
CTNNB1(NM_001904.4)	c.1955C>T	12	Wnt
GNAS(NM_080425.3)	c.2531G>A	1	cAMP
GNAS(NM_080425.3)	c.2530C>T	3	cAMP
GNAS(NM_080425.3)	c.2531G>A	5	cAMP
GNAS(NM_080425.3)	c.2531G>A	9	cAMP
GNAS(NM_080425.3)	c.2531G>A	11	cAMP
GNA11(NM_002067.5)	c.196G>A	5	cAMP
GNA11(NM_002067.5)	c.412G>T	11	cAMP
GNA11(NM_002067.5)	c.785C>A	12	cAMP
GNAQ(NM_002072.5)	c.303C>A	2	cAMP
ARID1A(NM_006015.4)	c.5975C>T	5	Chromatin
ARID1A(NM_006015.4)	c.3219G>A	5	Chromatin
ARID1A(NM_006015.4)	c.3562C>T	5	Chromatin
ARID1A(NM_006015.4)	c.4024C>T	5	Chromatin
ARID1A(NM_006015.4)	c.880A>C	6	Chromatin
ARID1A(NM_006015.4)	c.5372C>A	12	Chromatin
IDH1(NM_001282387.1)	c.473C>T	2	EpigeneticRegulation
IDH1(NM_001282387.1)	c.481G>A	6	EpigeneticRegulation
IDH1(NM_001282387.1)	c.394C>T	8	EpigeneticRegulation
IDH1(NM_001282387.1)	c.1004C>T	9	EpigeneticRegulation
KMT2A(NM_001197104.1)	c.5383G>A	4	EpigeneticRegulation
KMT2A(NM_001197104.1)	c.5755G>A	12	EpigeneticRegulation
FBXW7(NM_001349798.2)	c.1160T>A	10	Notch
NOTCH1(NM_017617.5)	c.7171C>T	12	Notch
BRCA2(NM_000059.3)	c.3853G>T	7	DNARepair
SMAD4(NM_005359.5)	c.538C>T	12	TGF-Beta