

## Supplemental Figure for:

Next-Generation Sequencing of Circulating Tumor DNA Reveals Frequent Alterations in Advanced Hepatocellular Carcinoma Sadakatsu Ikeda et al.

Figure S1. Genes covered by Guardant 360

## Genes with Complete Exon and Partial Intron Coverage

Gene	LOD	Gene	LOD	Gene	LOD	Gene	LOD
APC	<0.1%	AR	<0.1%	ARID1A	<0.1%	BRAF	<0.1%
BRCA1	<0.1%	BRCA2	<0.1%	CCND1	<0.1%	CCND2	<0.1%
CCNE1	<0.1%	CDK4	<0.1%	CDK6	<0.1%	CDKN2A	<0.1%
CDKN2B	<0.1%	EGFR#	<0.1%	ERBB2	<0.1%	FGFR1	<0.1%
FGFR2	<0.1%	HRAS	<0.1%	KIT	<0.1%	KRAS	<0.1%
MET	<0.1%	MYC	<0.1%	NF1	<0.1%	NRAS	<0.1%
<b>PDGFRA</b>	<0.1%	PIK3CA	<0.1%	PTEN	<0.1%	RAF1	<0.1%
TP53	<0.1%						

Genes in bold are those that are also analyzed for copy numer variations (CNVs). # EFGR indels in exon 19 and 20.

## Genes with Critical Exon Coverage

Gene	LOD	Gene	LOD	Gene	LOD	Gene	LOD
AKT1	<0.1%	ALK	<0.1%	ARAF	<0.1%	ATM	<0.1%
CDH1	<0.1%	CTNNB1	<0.1%	ESR1	<0.1%	EZH2	<0.1%
FBXW7	<0.1%	FGFR3	<0.1%	GATA3	<0.1%	GNA11	<0.1%
GNAQ	<0.1%	GNAS	<0.1%	HNF1A	<0.1%	IDH1	<0.1%
IDH2	<0.1%	JAK2	0.1%	JAK3	<0.1%	MAP2K1	<0.1%
MAP2K2	<0.1%	MLH1	<0.1%	MPL	<0.1%	NFE2L2	<0.1%
NOTCH1	<0.1%	NPM1	0.2%	NTRK1	<0.1%	PTPN11	<0.1%
RET	<0.1%	RHEB	0.1%	RHOA	<0.1%	RIT1	<0.1%
ROS1	0.1%	SMAD4	<0.1%	SMO	<0.1%	SRC	<0.1%
STK11	<0.1%	TERT	<0.1%	VHL	<0.1%		

## Rearrangements

Gene	LOD	Gene	LOD	Gene	LOD	Gene	LOD	
ALK	<0.1%	NTRK1	<0.1%	RET	<0.1%	ROS1	<0.1%	