

Supplemental Table 1

Patient number	MET exon 14 mutation	Histologic subtype	IHC MET	MET amplif.	Ratio/ mean of copy number	Other molecular alterations	IHC HGF
CHUL #2	c.3076_3082+2del	ADC	3+	No	1.2/3.1		60
CHUL #7	c.3082+2T>C	ADC	2+	No	0.36/1.67		60
CHUL #15	c.2942-31_2942-4del	ADC	2+	No	1.04/2.91		120
CHUL #31	c.2942-36_2942-4del	ADC	2+	No	0.4/1.82		70
CHUL #33	c.3082G>T	ADC	2+	No	0.62/3.97	TP53 c.809T>C	90
CHUL #63	c.3082+2T>C	ADC	ND	ND			60
CHUL #66	c.2942-1G>C	ADC	ND	ND		PIK3CA c.1636C>A (Q546K)	60
CHUL #68	c.3080A>G	ADC	3+	ND		KRAS c.35G>A (G12D)	50
CHUL #45	3077_3082+1del	ADC	3+	Yes	clusters in 67% of cells	PIK3CA c.1262C>T; TP53 c.396G>C	0
CHUL #57	c.3082+1G>A	ADC	2+	ND		GNAS c.601C>T; MDM2 and CDK4 amplification	0
CHUL #60	c.3082+3A>T	ADC	3+	ND			0
CHUL #61	c.2942-21_2942-3del	ADC	2+	ND			0
CHUL #9	c.2942-14_2942-4del	ADC	1+	No	0.99/1.84	PTEN loss	NI
CHUL #14	c.2942-47_2949del	ADC	1+	No	0.95/1.5	PTEN loss	NI
CHUL #37	c.2942-19_2942-4del	ADC	ND	ND		TP53 c.610G>T	NI
CHUL #51	c.2942-21_2945del	ADC	2+	ND			NI
CHUL #69	c.2942-21_2942-8del	ADC	ND	ND			NI
CHUL #70	c.3082G>C	ADC	3+	ND			NI