Table S1. Mutational Analysis of FL Samples

MSKCC dataset 28 samples
Michigan data set 113 samples GRCh37/hg19

MKSCC DATA SET (28 samples)	Amino Acid	Type	Transcript position: alteration	Amino acid position: alteration	Ensemble Position
1	1	Missense	A>T	M>L	2488104
2	110	Missense	A>C	N>I	2491286
3	122	Missense	C>A	S>R	2491323
4	126	Missense	T>A	F>L	2491333
5	157	Missense	G>A	S>N	2492072

MICHIGAN DATA SET (113 samples)	Amino Acid	Type	Transcript position: alteration	Amino acid position: alteration	Ensemble Position
1	1	Missense	T>A	M>K	2488105
2	1	Missense	G>A	M>L	2488106
3	1	Missense	G>A	M>L	2488106
4	12	Truncation	G>A	W>X	2488138
5	12	Truncation	G>A	W>X	2488139
6	36	Missense	C>A	A>D	2489202
7	42	Missense	G>T	C>F	2489220
8	53	Missense	G>A	C>Y	2489253
9	54	Truncation	C>A	C>X	2489257
10	57	Missense	C>G	C>W	2489266
11	57	Missense	G>C	C>S	2489265
12	61	Truncation	T>G	Y>X	2489786
13	62	Missense	G>C	R>P	2489788
14	63	Missense	T>G	V>G	2489791
15	68	Missense	G>A	G>R	2489805
16	69	Truncation	G>T	E>X	2489808
17	72	Missense	G>T	G>V	2489818
18	73	Missense	C>A	T>K	2489821
19	82	Missense	A>C	T>P	2489847
20	83	Missense	T>G	Y>D	2489850
21	88	Missense	A>T	N>I	2489865
22	95	Truncation	C>T	Q>X	2489886
23	96	Truncation	C>A	C>X	2489891
24	97	Truncation	C>T	Q>X	2489892
25	97	Truncation	C>T	Q>X	2489892
26	97	Truncation	C>T	Q>X	2489892
27	102	Missense	G>C	A>P	2489907
28	111	Missense	G>A	C>Y	2491289
29	112	Missense	C>T	S>F	2491292
30	135	Missense	G>A	C>Y	2491361
31	158	Truncation	C>T	Q>X	2492074
32	186	Frameshift	A>GCTGGCT(	S>FS	2493116
33	187	Truncation	G>A	W>X	2493120
34	204	Truncation	G>A	W>X	2493171
35	266	Missense	C>T	T>M	2494657