

Additional file 4. Distribution of allele loss in tumors with LOH

	Expected		Observed		Chi-Square p value
	LOH ^{mut}	LOH ^{wt}	LOH ^{mut}	LOH ^{wt}	
All tumors with LOH (n=72)	36	36	7	65	< 0.001
ER- tumors with LOH (n=34)	17	17	3	31	< 0.001
ER+ tumors with LOH (n=38)	19	19	4	34	< 0.001