

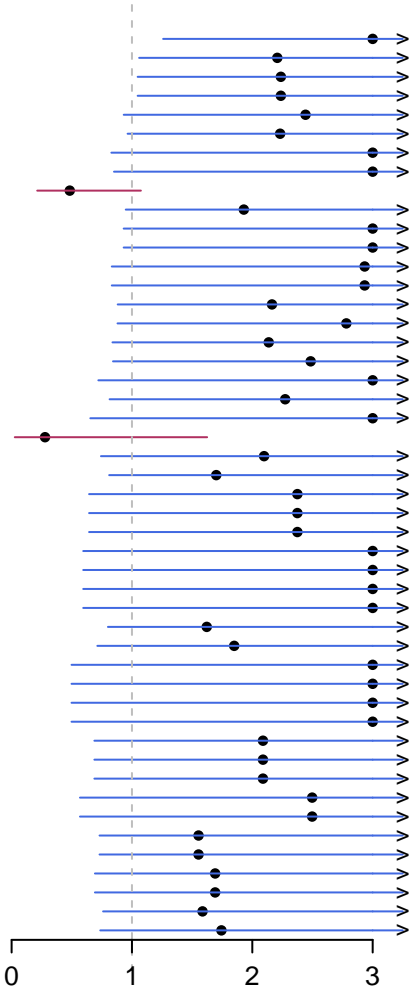
Relapse (n = 68) v/s Primary (n = 77)

Relapse

Primary

OR

P-value



<i>NTRK2</i>	5	18	3.811	*
<i>FAT3</i>	20	37	2.208	*
<i>APC</i>	17	33	2.237	*
<i>PKHD1</i>	17	33	2.237	*
<i>ATRX</i>	8	19	2.442	*
<i>MAGI2</i>	12	25	2.231	
<i>CDK8</i>	0	5	Inf	
<i>FGFR1</i>	2	9	4.329	
<i>MAP3K1</i>	24	16	0.483	
<i>KMT2C</i>	26	42	1.93	
<i>BRIP1</i>	4	13	3.226	
<i>PIK3CG</i>	4	13	3.226	
<i>PHLPP2</i>	4	12	2.933	
<i>XBP1</i>	4	12	2.933	
<i>LET2</i>	10	21	2.164	
<i>EPLA6</i>	5	14	2.781	
<i>GRIN2A</i>	9	19	2.136	
<i>INPP4B</i>	6	15	2.485	
<i>ROBO2</i>	2	8	3.795	
<i>ARID1A</i>	7	16	2.273	
<i>MLAP</i>	1	6	5.607	
<i>PTEN</i>	6	25	0.278	
<i>PTPRD</i>	7	15	2.098	
<i>AMER1</i>	20	32	1.7	
<i>MET</i>	4	10	2.375	
<i>PMS2</i>	4	10	2.375	
<i>PNRC1</i>	4	10	2.375	
<i>CSE2RB</i>	2	7	3.276	
<i>FGF12</i>	2	7	3.276	
<i>RAD17</i>	2	7	3.276	
<i>SF3A1</i>	7	7	3.276	
<i>FOXO1</i>	28	41	1.621	
<i>EPHB6</i>	9	17	1.85	
<i>AGTR2</i>	1	5	4.612	
<i>CRYL</i>	1	5	4.612	
<i>EPHA5</i>	1	5	4.612	
<i>FGF2</i>	1	5	4.612	
<i>ATR</i>	6	13	2.089	
<i>MYO18A</i>	6	13	2.089	
<i>NOTCH1</i>	6	13	2.089	
<i>PLCG2</i>	3	8	2.497	
<i>SEMA3A</i>	3	8	2.497	
<i>GNAO1</i>	19	29	1.553	
<i>PALB2</i>	19	29	1.553	
<i>ASXL1</i>	11	19	1.691	
<i>PARP4</i>	11	19	1.691	
<i>DICER1</i>	21	32	1.586	
<i>SEMA3E</i>	12	21	1.743	

Odds ratio with 95% CI
(1 = no effect, < 1 Relapse has more mutants)