S8 Table. Comparison of somatic CNV gain between patients with detectable and undetectable MRD

Gene	Detectable MRD			Undetectable MRD			p-value	OR
	Gain (n)	**Wt (n)**	**Frequency (%)**	**Gain (n)**	**Wt (n)**	**Frequency (%)**		
ERBB2	7	11	38.9	17	45	27.4	0.3886	1.673
CDK12	6	12	33.3	10	52	16.1	0.1767	2.564
GATA3	2	16	11.1	1	61	1.6	0.1254	7.368
GNAS	2	16	11.1	3	59	4.8	0.3131	2.425
BRIP1	3	15	16.7	2	60	3.2	0.0724	5.821
CDKN2A	2	16	11.1	0	62	0.0	**0.0484**	Inf
HOXB13	3	15	16.7	0	62	0.0	**0.0099**	Inf
MYC	3	15	16.7	16	46	25.8	0.5394	0.579
PPM1D	2	16	11.1	0	62	0.0	**0.0484**	Inf
SPOP	2	16	11.1	6	56	9.7	1	1.164
EGFR	2	16	11.1	2	60	3.2	0.2169	3.671
KRAS	2	16	11.1	2	60	3.2	0.2169	3.671
MDM4	2	16	11.1	3	59	4.8	0.3131	2.425
MPL	2	16	11.1	0	62	0.0	**0.0484**	Inf
RAD51C	2	16	11.1	2	60	3.2	0.2169	3.671
VHL	2	16	11.1	0	62	0.0	**0.0484**	Inf

CNV, copy number variant; MRD, molecular residual disease; OR, odds ratio.