

Quantity	average	median	min	max	stdev
No. mutations, WGS	844	328	110	8,192	1,306
No. mutations, WGA	1,531	694	23	10,966	2,230
% mutations in overlap, WGS	31%	31%	1%	74%	20%
% mutations in overlap, WGA	44%	45%	3%	71%	13%
No. putative mutations removed, VAQ	309	164	17	11,439	1,372
No. putative mutations removed, LOH	539	169	3	8,538	1,051
No. putative mutations removed, 10bp-SNV	46	19	0	1,515	115
No. putative mutations removed, 10bp-INDEL	28	16	0	535	45
No. putative mutations removed, dbSNP	16	8	0	348	33
No. putative mutations removed, <10%	1	1	1	36	3
% overlap removed, VAQ	53%	57%	0%	100%	28%
% overlap removed, LOH	51%	52%	0%	99%	29%
% overlap removed, 10bp-SNV	3%	2%	0%	20%	3%
% overlap removed, 10bp-INDEL	4%	4%	0%	22%	4%
% overlap removed, db-SNP	3%	2%	0%	47%	7%