100X coverage, run 7

filter criterion (y-axis), proportion (x-axis)

filter criterion	DNMs	miscalled variants	sequencing errors
AB_{max}	47	574	112
AB_{min}	47	294	44
AD	47	366	92
DP_{max}	47	635	125
DP_{min}	47	383	43
$GATK_{Best Practice}$	47	484	62
no known	47	601	124
GQ	47	14	12
QUAL	47	92	1
reassembly	37	470	112
DNM candidates	47	635	125

Legend: DNMs (green), miscalled variants (orange), sequencing errors (red)