100X coverage, run 5

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

Filter criterion	DNMs	miscalled variants	sequencing errors
AB_{max}	49	627	132
AB_{min}	49	305	60
AD	49	286	109
DP_{max}	49	662	133
DP_{min}	49	82	23
$GATK_{Best\ Practice}$	49	497	87
no known	49	637	133
GQ	49	24	9
QUAL	49	88	1
reassembly	42	497	111
DNM candidates	49	662	133

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