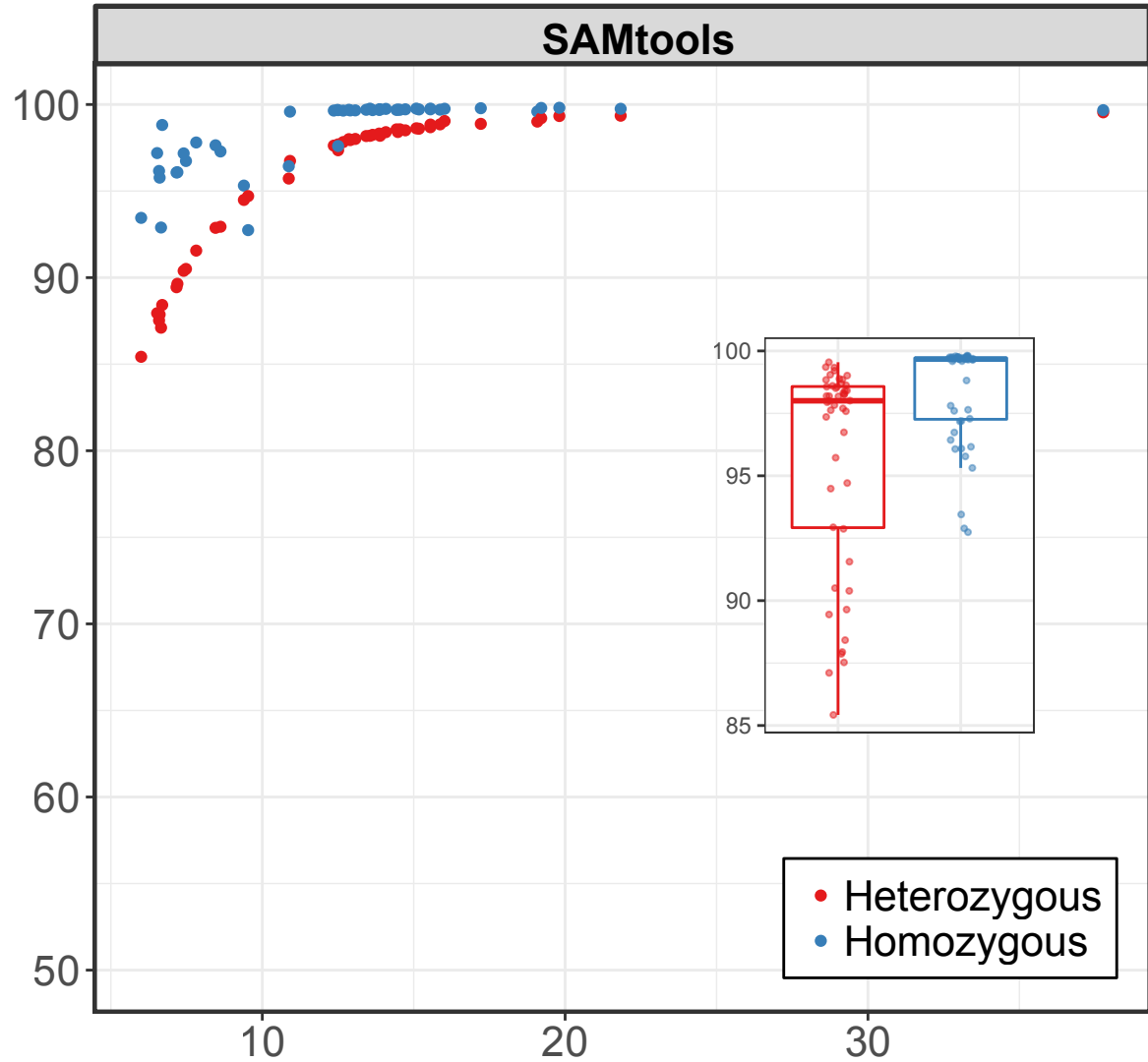
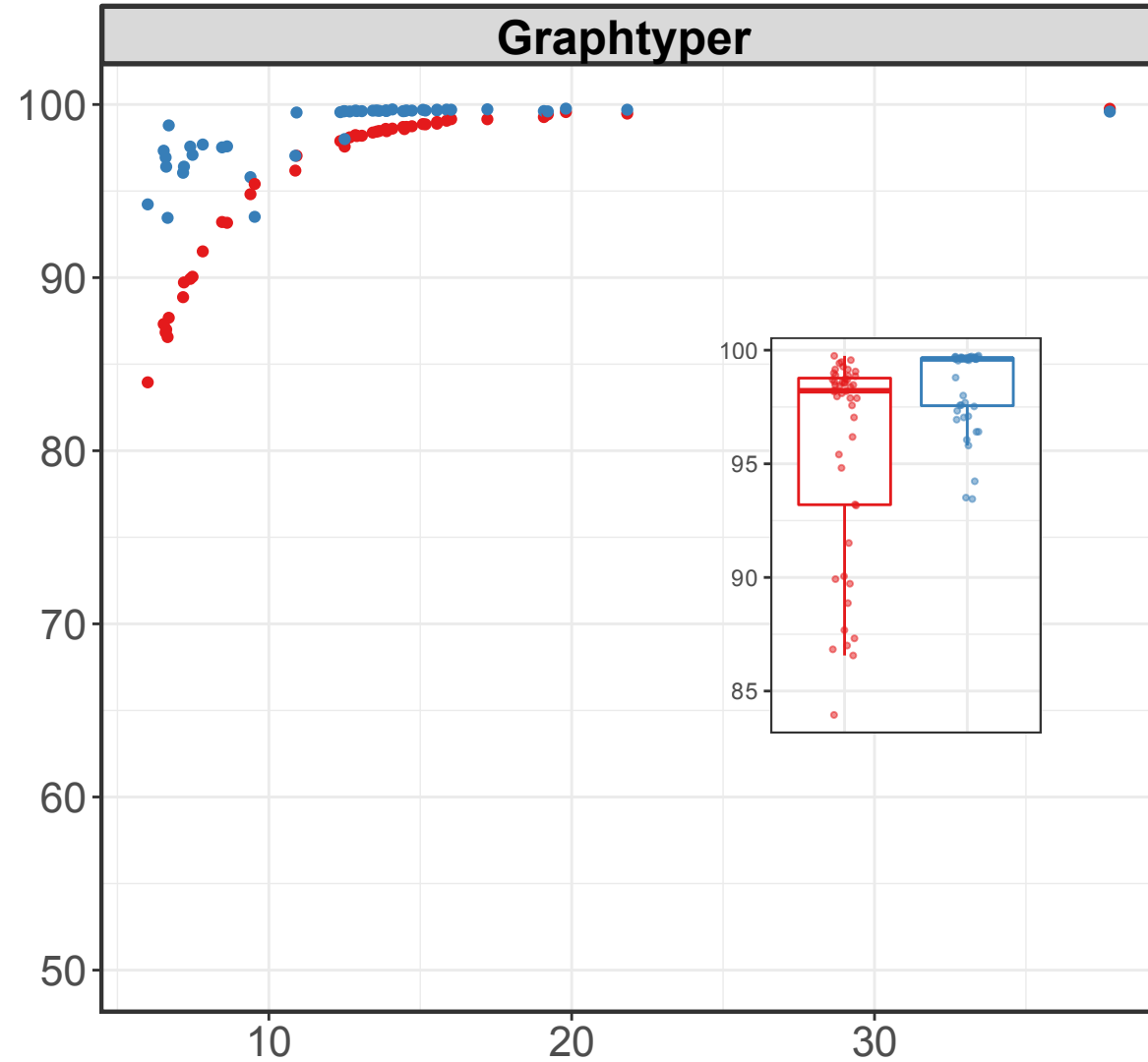
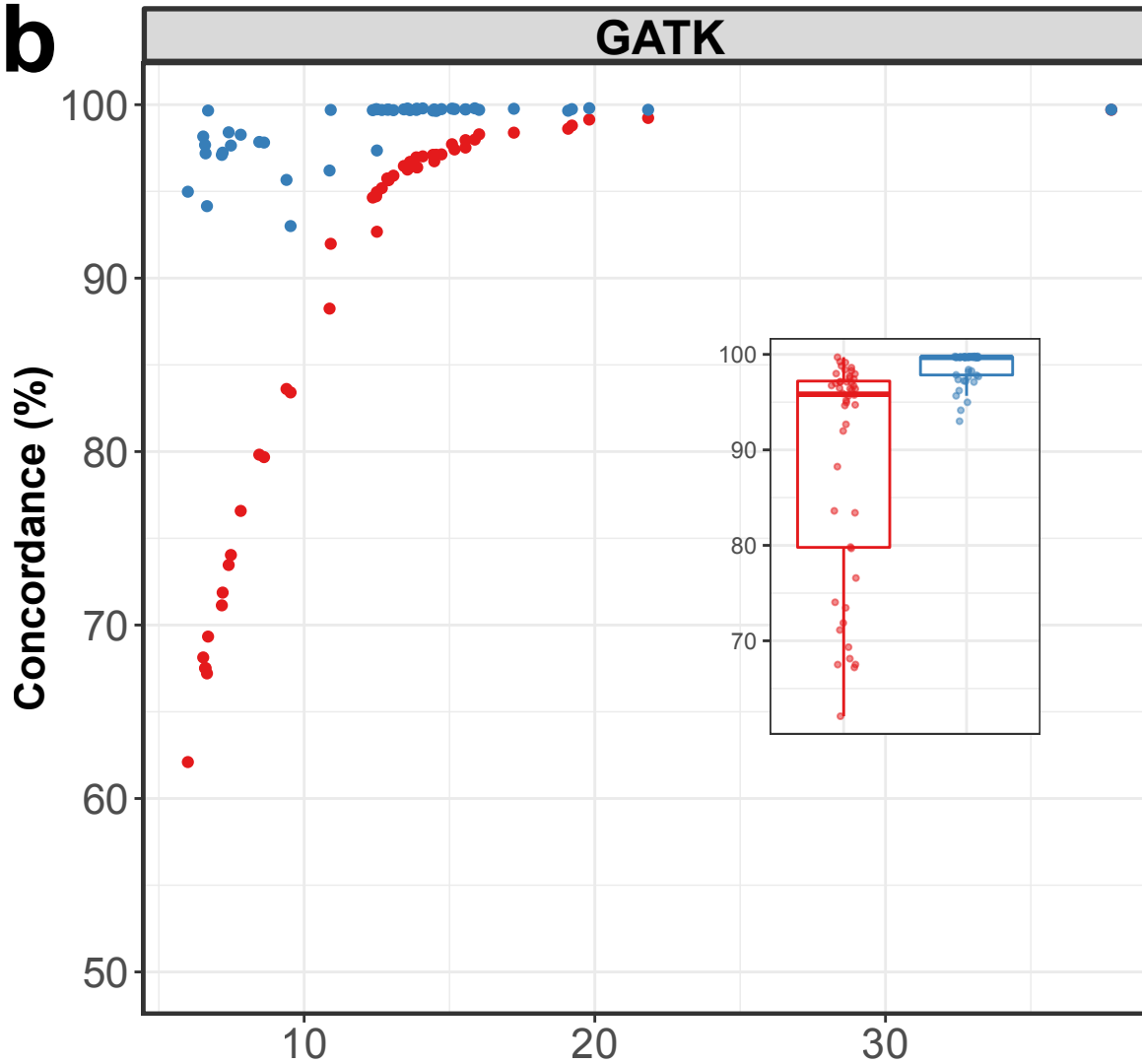


a

	Heterozygous concordance				Homozygous concordance			
	full		filtered		full		filtered	
	raw	imp	raw	imp	raw	imp	raw	imp
<i>GATK</i>	89.17	99.11	89.24	99.21	98.74	99.18	98.75	99.27
<i>Graph typer</i>	95.79	99.36	95.82	99.44	98.55	99.51	98.59	99.57
<i>SAMtools</i>	95.73	98.91	95.77	98.99	98.46	99.37	98.48	99.41

b

C