

		high - 0.33x		high - 1x		low - 0.33x		low - 1x	
		total sites	% real	total sites	% real	total sites	% real	total sites	% real
Poolparty	PPalign ^a	70,067	99.5%	91,509	99.8%	345,223	99.8%	452,827	99.8%
	PPanalyze ^b	68,292	99.5%	86,036	99.8%	334,664	99.8%	422,029	99.8%
ANGSD^c	$p < 0.0$	527,762	17.2%	920,208	12.0%	864,530	51.4%	1,323,894	41.1%
	$p < 0.01$	94,104	69.6%	209,617	43.5%	344,606	92.6%	411,213	98.1%
	$p < 0.001$	57,027	98.7%	127,423	67.5%	275,355	99.8%	373,603	99.6%
	10 reads + 3 inds/pop ^d	56,060	98.7%	127,415	67.5%	271,175	99.8%	373,473	99.6%
Poolparty	20 global reads ^e	67,233	99.5%	86,011	99.8%	328,654	99.8%	421,737	99.8%
to ANGSD	10 reads + 3 inds/pop ^e	66,168	99.5%	85,987	99.8%	323,806	99.8%	421,568	99.8%

- a min. 20 global reads, MAF 0.001
- b sites specified by PPalign; min. 3 ind+10 reads/pop
- c min. 20 global reads, MAF 0.001, min. 6 global individuals
- d filtering of sites specified by ANGSD $p < 0.001$
- e no additional p-value or MAF filter