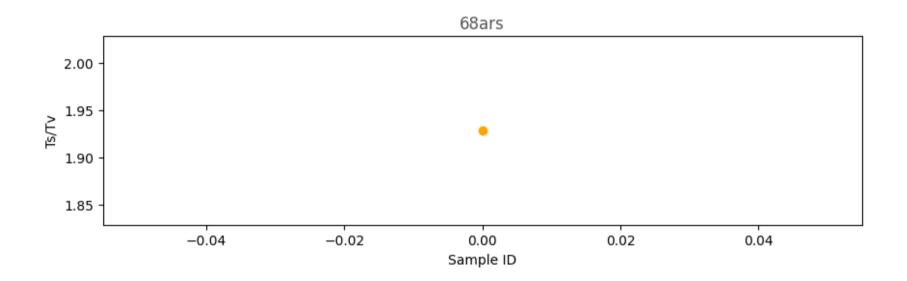
#### **Summary Numbers**

	SNPs			indels		MNPs	others		
Callset	n	ts/tv	(1st ALT)	n	frm*				
68ars	4,089,501	1.93	1.93	948,588	1	0	0		
* frameshift ratio: out/(out+in)									

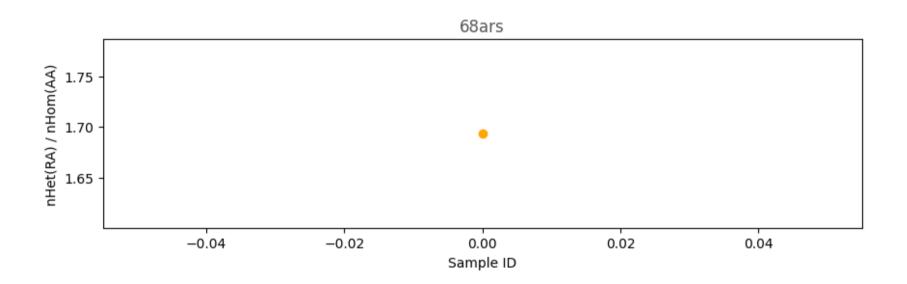
	singl	etons (A	multiallelic		
Callset	SNPs	ts/tv	indels	sites	SNPs
68ars	62.9%	1.90	67.8%	98,827	2,312

- 68ars../ngc/projects2/gm/data/archive/2022/variants/snv/68arseprf-103830872610-Normal\_B lood\_noinfo-WGS\_v1\_IlluminaDNAPCRFree\_X-220126\_A01411\_BH2CL7DSX3-RHGM\_LABKA\_WGS-WGS03718\_snv\_germline\_raw.haplotype\_caller.vcf.gz

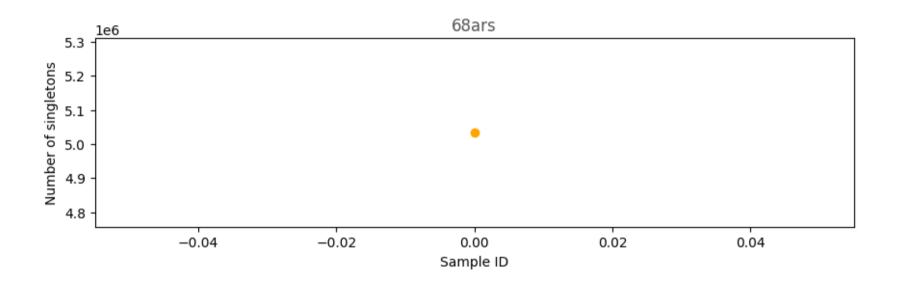
# Ts/Tv by sample



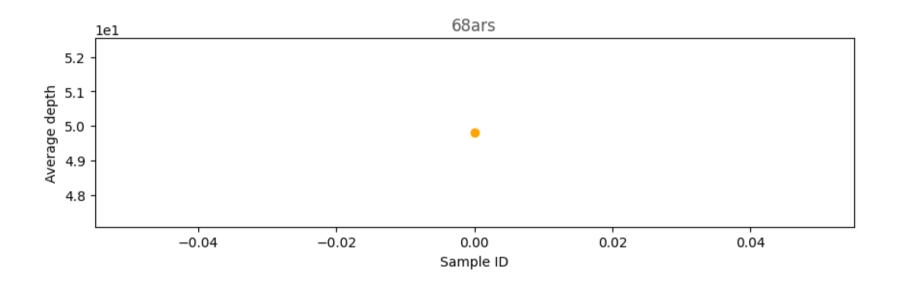
### Hets vs non-ref Homs by sample



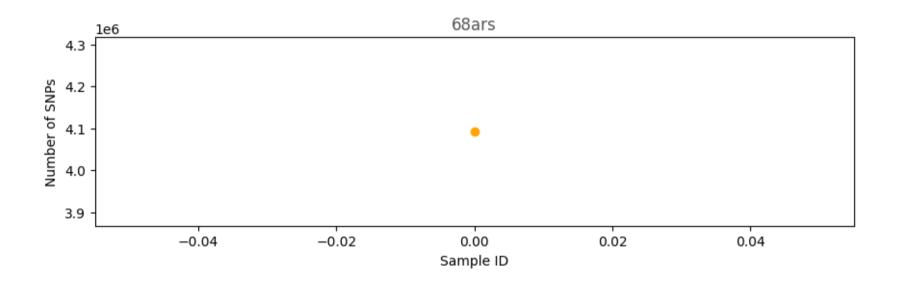
### Singletons by sample (hets and homs)



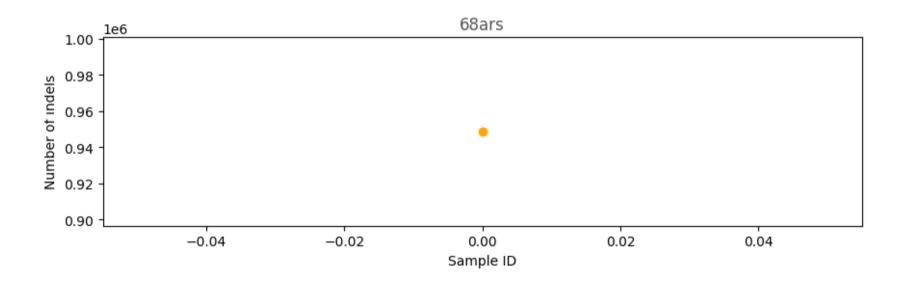
## Average depth by sample



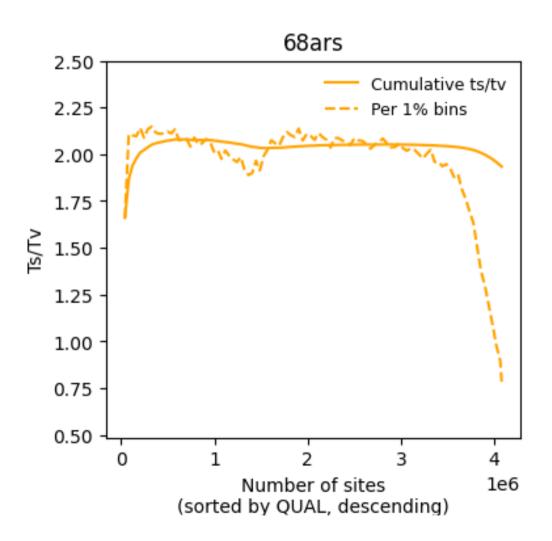
## Number of SNPs by sample



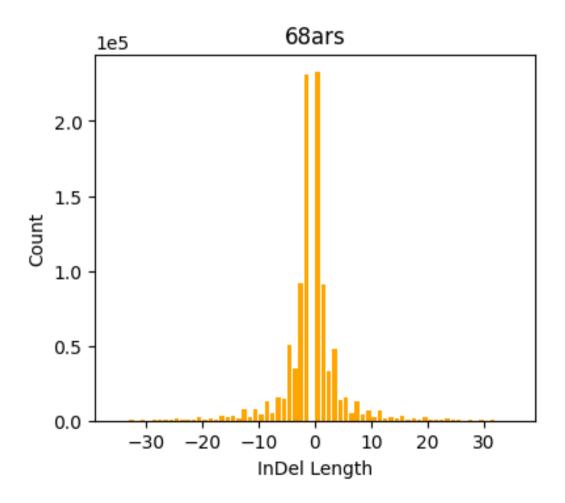
### Number of indels by sample



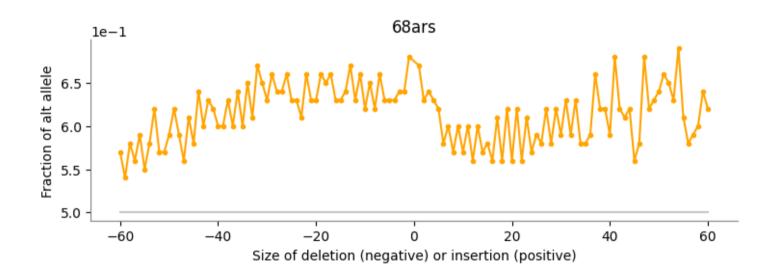
#### Ts/Tv stratified by QUAL



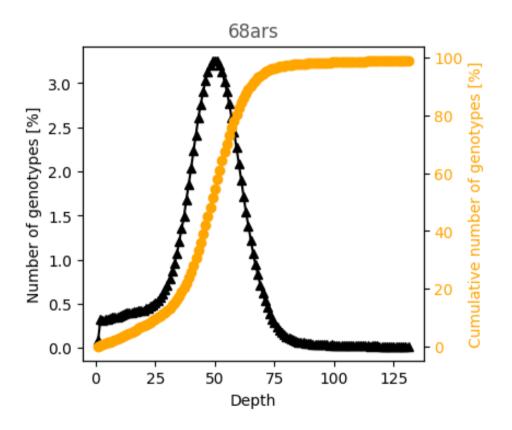
#### Indel distribution



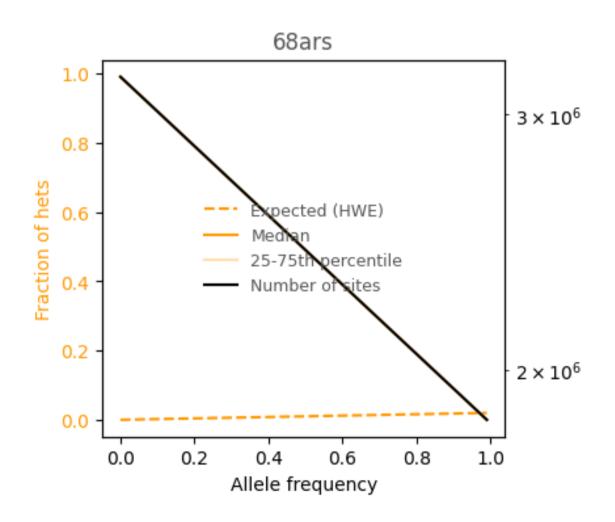
#### Fraction of alternate indel allele



## Depth distribution



#### Number of HETs by AF



### Substitution types

