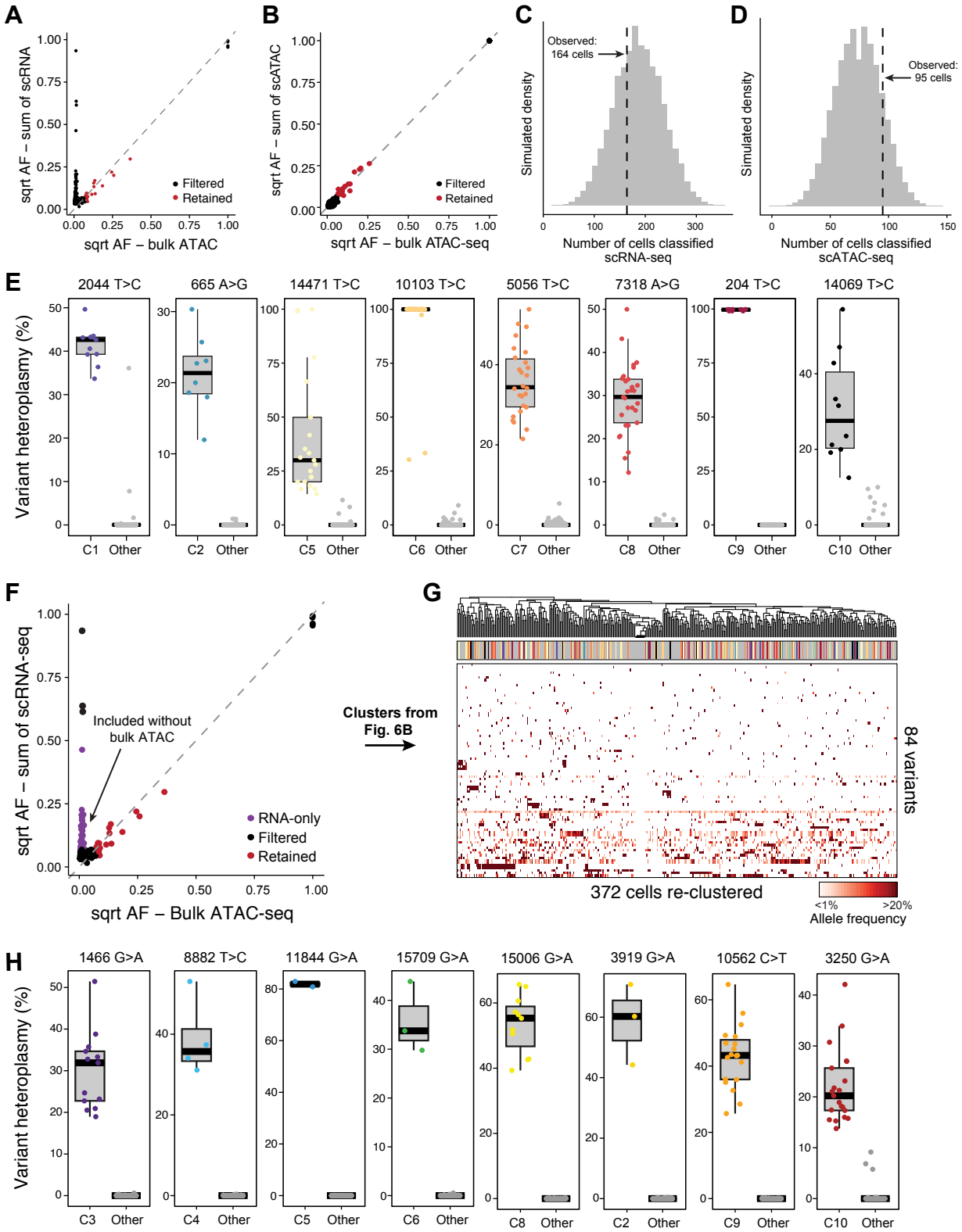
Figure S6


A
sqrt AF – sum of scRNA
sqrt AF – bulk ATAC
Filtered
Retained
B
sqrt AF – sum of scATAC
sqrt AF – bulk ATAC-seq
Filtered
Retained
C
Observed: 164 cells
Simulated density
Number of cells classified scRNA-seq
D
Observed: 95 cells
Simulated density
Number of cells classified scATAC-seq
E
2044 T>C
665 A>G
14471 T>C
10103 T>C
5056 T>C
7318 A>G
204 T>C
14069 T>C
Variant heteroplasmy (%)
C1 Other
C2 Other
C5 Other
C6 Other
C7 Other
C8 Other
C9 Other
C10 Other
F
sqrt AF – sum of scRNA-seq
sqrt AF – Bulk ATAC-seq
Included without bulk ATAC
RNA-only
Filtered
Retained
Clusters from Fig. 6B
G
84 variants
372 cells re-clustered
<1%
>20%
Allele frequency
H
1466 G>A
8882 T>C
11844 G>A
15709 G>A
15006 G>A
3919 G>A
10562 C>T
3250 G>A
Variant heteroplasmy (%)
C3 Other
C4 Other
C5 Other
C6 Other
C8 Other
C2 Other
C9 Other
C10 Other