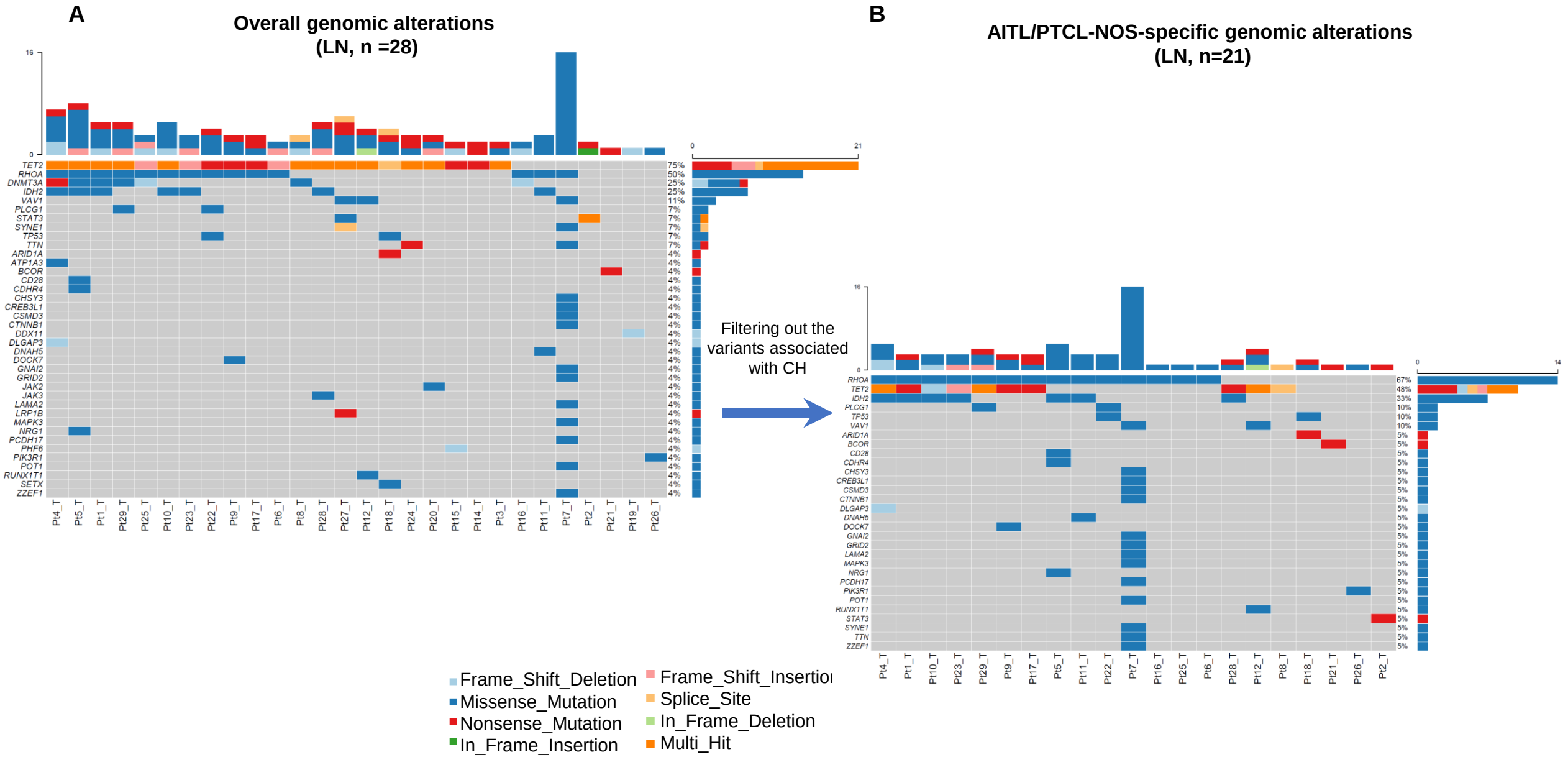
A
Overall genomic alterations
(LN, n =28)
B
AITL/PTCL-NOS-specific genomic alterations
(LN, n=21)
Filtering out the variants associated with CH
Frame_Shift_Deletion
Frame_Shift_Insertion
Missense_Mutation
Splice_Site
Nonsense_Mutation
In_Frame_Deletion
In_Frame_Insertion
Multi_Hit