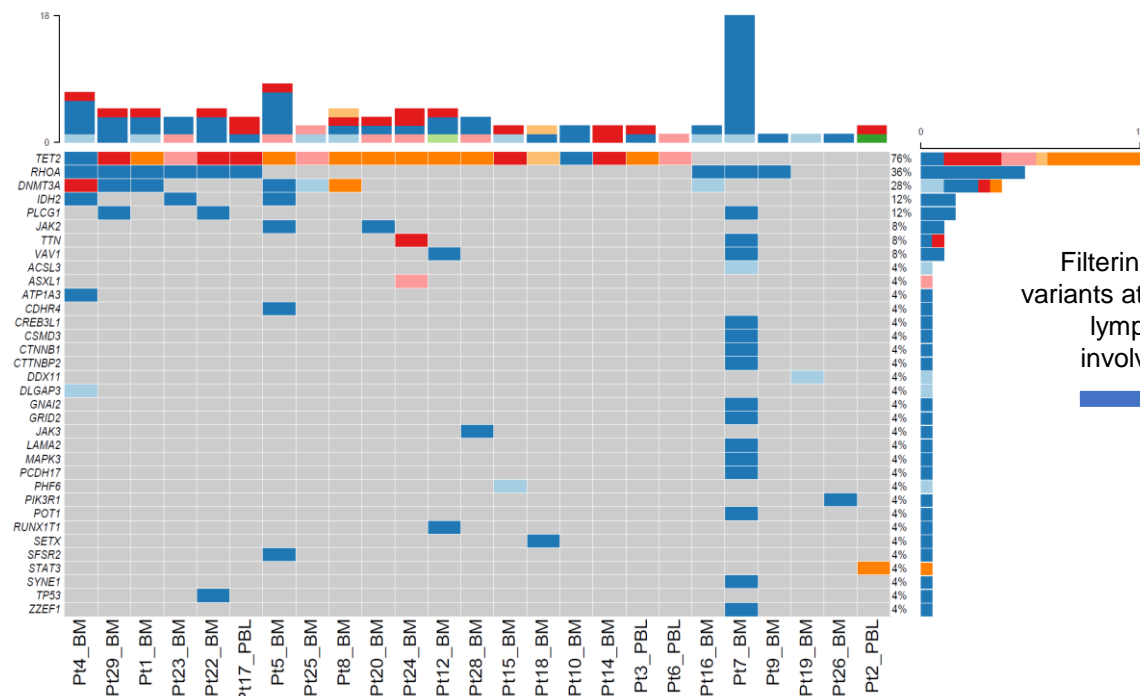


A

Overall genomic alterations (paired BM/PB, n =25)

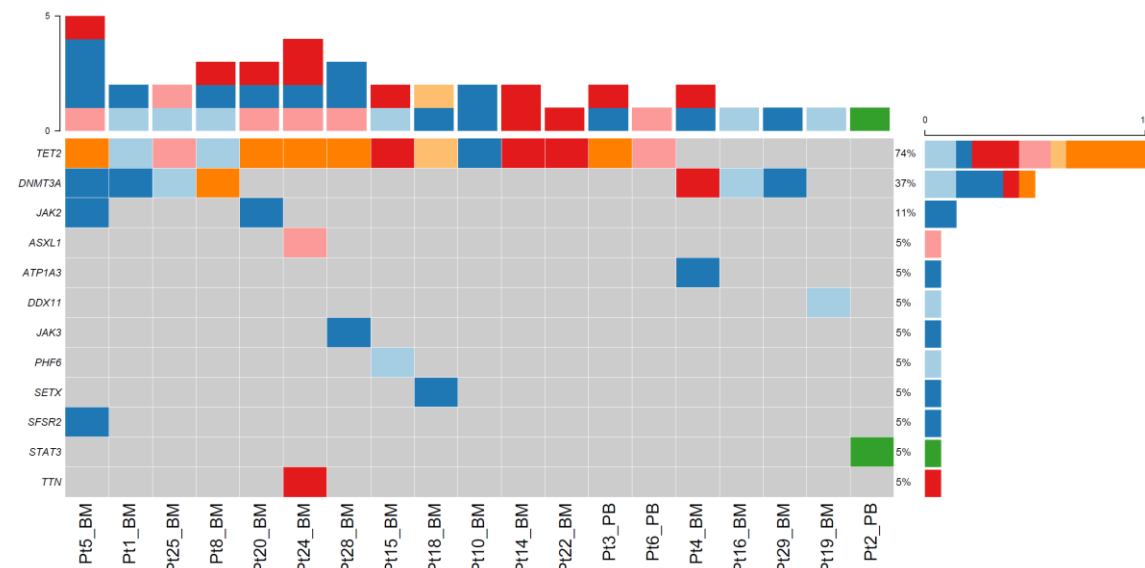


Filtering out the
variants attributable to
lymphoma
involvement



B

CH-associated genomic alterations (Paired BM/PB, n =19)



- Frame_Shift_Deletion
- Missense_Mutation
- Nonsense_Mutation
- In_Frame_Insertion
- Frame_Shift_Insertion
- Splice_Site
- In_Frame_Deletion
- Multi_Hit