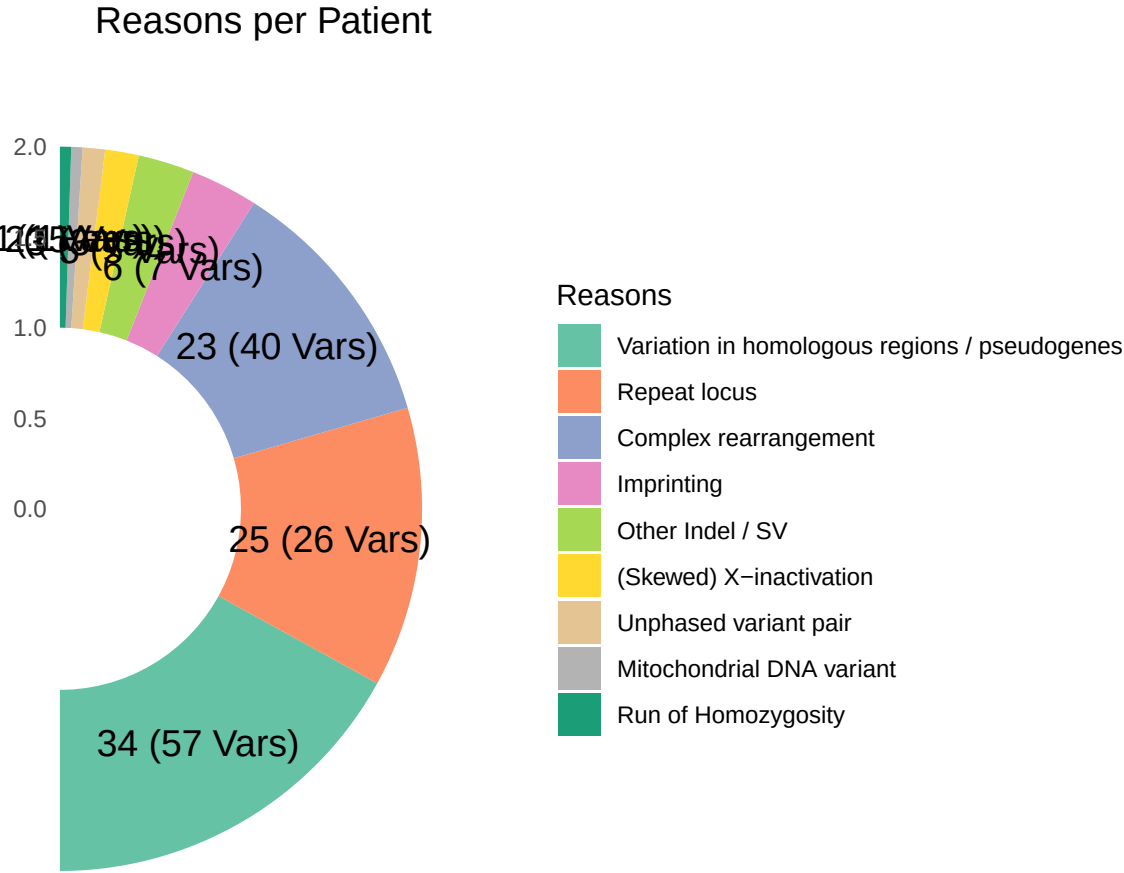
Reasons per Patient
2.0
1.0
0.5
0.0
6 (7 Vars)
23 (40 Vars)
25 (26 Vars)
34 (57 Vars)
Reasons
Variation in homologous regions / pseudogenes
Repeat locus
Complex rearrangement
Imprinting
Other Indel / SV
(Skewed) X−inactivation
Unphased variant pair
Mitochondrial DNA variant
Run of Homozygosity