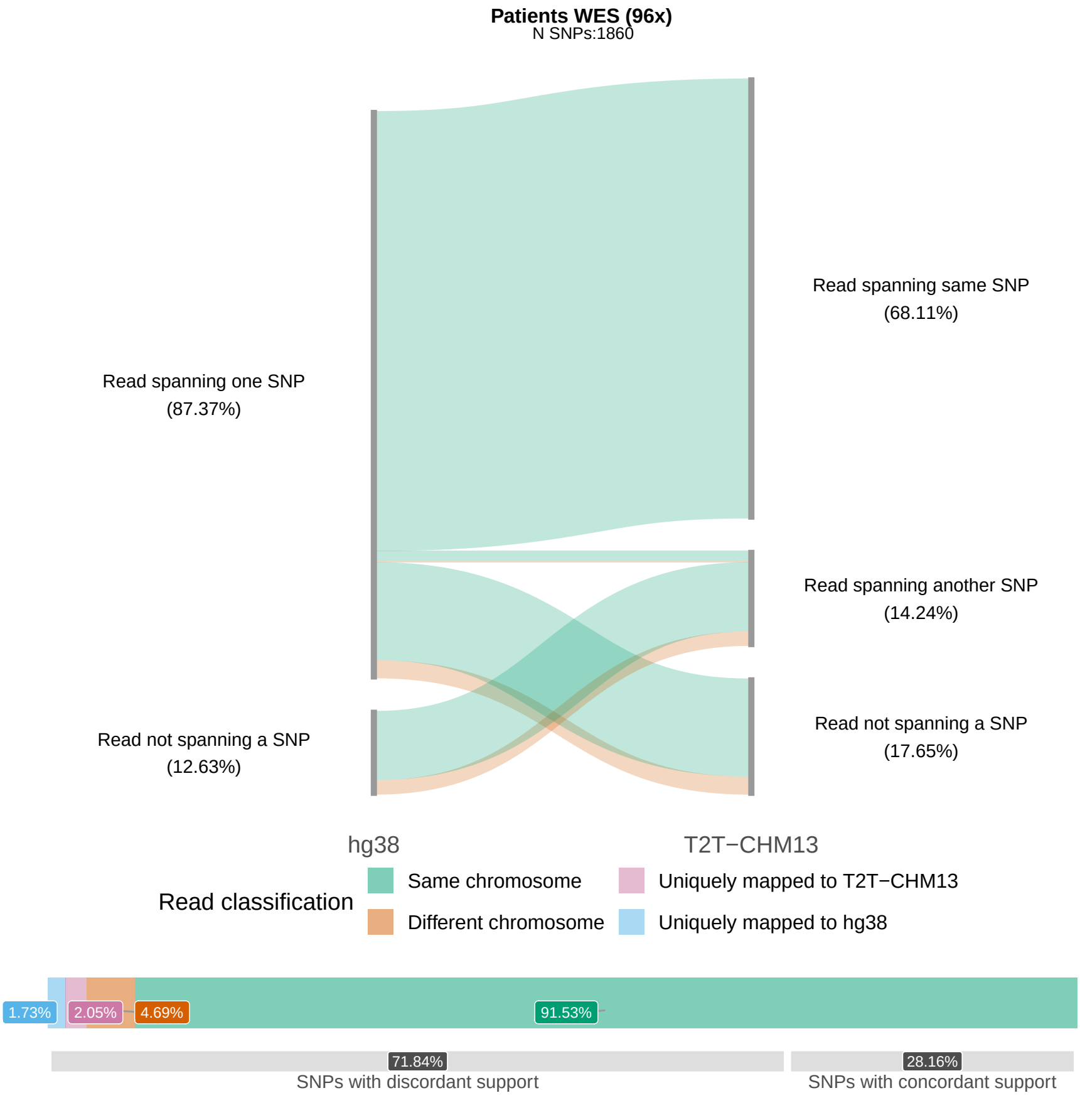
Patients WES (96x)
N SNPs:1860
Read spanning one SNP
(87.37%)
Read not spanning a SNP
(12.63%)
Read spanning same SNP
(68.11%)
Read spanning another SNP
(14.24%)
Read not spanning a SNP
(17.65%)
hg38
T2T−CHM13
Read classification
Same chromosome
Different chromosome
Uniquely mapped to T2T−CHM13
Uniquely mapped to hg38
1.73%
2.05%
4.69%
91.53%
71.84%
SNPs with discordant support
28.16%
SNPs with concordant support