

Fig S7. SHS iAs% Conditional Association Results

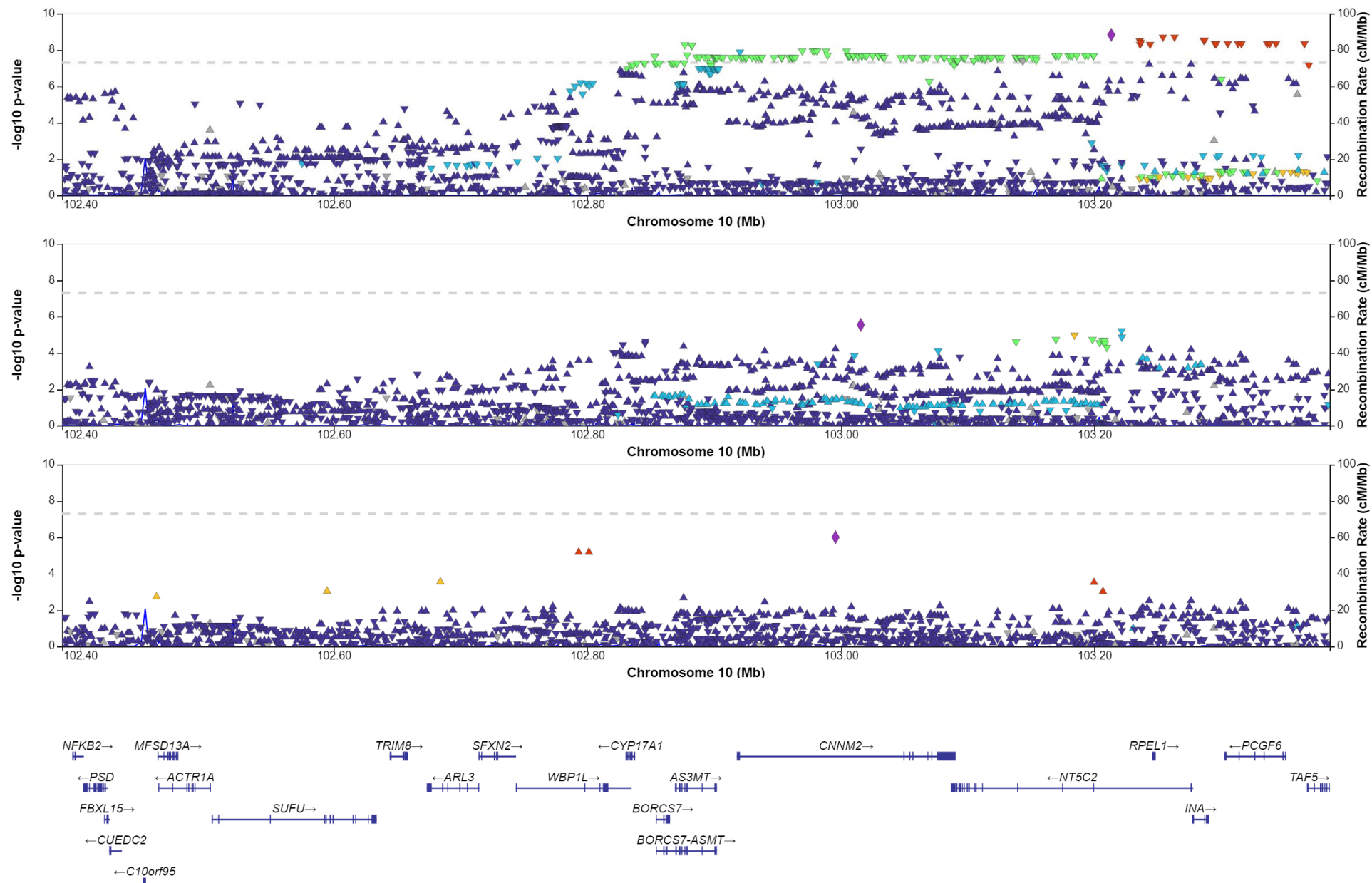


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Results of a post-imputation genetic association study of iAS% in the 10q24.32 region in SHS. P-values were generated with linear models adjusted for age, sex, and population structure. The top panel represents overall association results, the second shows p-values from models adjusted for the initial lead SNP, and the bottom panel shows the result of models adjusted for both previously identified variants. Two lead variants were identified in this analysis: chr10:103213304 and chr10:103015726.