

haplotype_ID

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graph TD; A(haplotype_ID) --> B(make_vcf_tt); B --> C(make_pfiles_tt); B --> D(make_cov_pheno_tt); C --> E(run_plink); D --> E; E --> F(combine_results); F --> G(cleanup); G --> H(all);
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The flowchart illustrates a sequential process starting with 'haplotype_ID' (yellow box). An arrow points down to 'make_vcf_tt' (blue box). From 'make_vcf_tt', two arrows branch out to 'make_pfiles_tt' (green box) and 'make_cov_pheno_tt' (green box). Arrows from both of these boxes point down to 'run_plink' (blue box). An arrow then points down to 'combine_results' (yellow box), followed by 'cleanup' (orange box), and finally 'all' (red box) at the bottom.

make_vcf_tt

make_pfiles_tt

make_cov_pheno_tt

run_plink

combine_results

cleanup

all