

Supplementary Figures

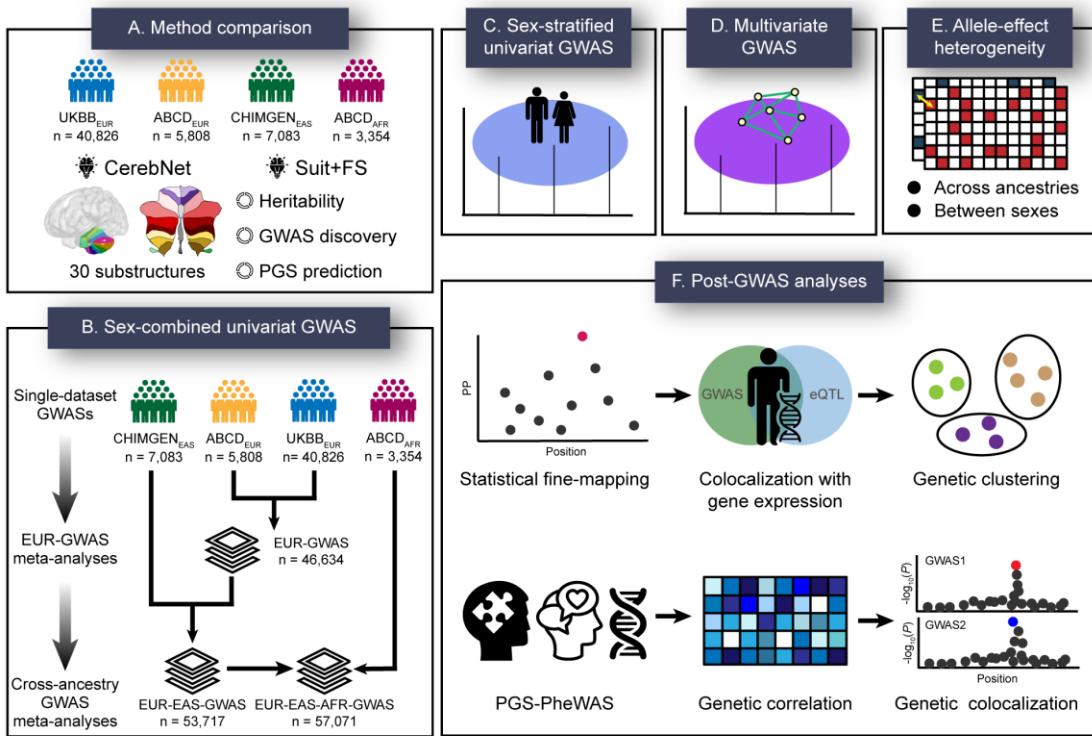


Fig. S1 | A schematic summary of study design. This study includes four independent subgroups with raw genetic and neuroimaging data (UKBB_{EUR}, ABCD_{EUR}, CHIMGEN_{EAS}, and ABCD_{AFR}). **(A)** To compare the cerebellar segmentation methods, we calculate the volumes of 30 cerebellar substructures using CerebNet and SUIT+FS, and we evaluate their differences in heritability estimation, GWAS discovery, and PGS prediction. **(B)** In the sex-combined univariate analyses, we further include total cerebellar volume as a trait and perform GWASs for the 31 cerebellar volumetric traits across the four subgroups, respectively. We also conduct one EUR-GWAS and two cross-ancestry (EUR-EAS and EUR-EAS-AFR) GWAS meta-analyses. **(C)** In the sex-stratified univariate analyses, we repeat the seven sets of GWASs in males and females, respectively. **(D)** Multivariate GWASs are performed in both sex-combined and sex-stratified populations. **(E)** We identify allelic-effect heterogeneity across ancestries

based on sex-combined EUR-GWASs, CHIMGEN_{EAS}-GWASs, and ABCD_{AFR}-GWASs, while we detect allelic-effect heterogeneity between sexes based on sex-stratified EUR-GWASs, CHIMGEN_{EAS}-GWASs, and ABCD_{AFR}-GWASs. (F) The post-GWAS analyses include statistical fine-mapping, genetic colocalization with cerebellar gene expression, genetic clustering for cerebellar subregions, PGS-PheWAS, and genetic correlation and colocalization with neuropsychiatric disorders. Abbreviations: ABCD, the Adolescent Brain Cognitive Development study; AFR, African ancestry; CHIMGEN, the Chinese Imaging Genetics study; EAS, East Asian ancestry; EUR, European ancestry; FS, FreeSurfer; GWAS, genome-wide association study; PGS, polygenic score; PheWAS, phenome-wide association study; SUIT, spatially unbiased infra-tentorial; FS, FreeSurfer; UKBB, the UK Biobank study.

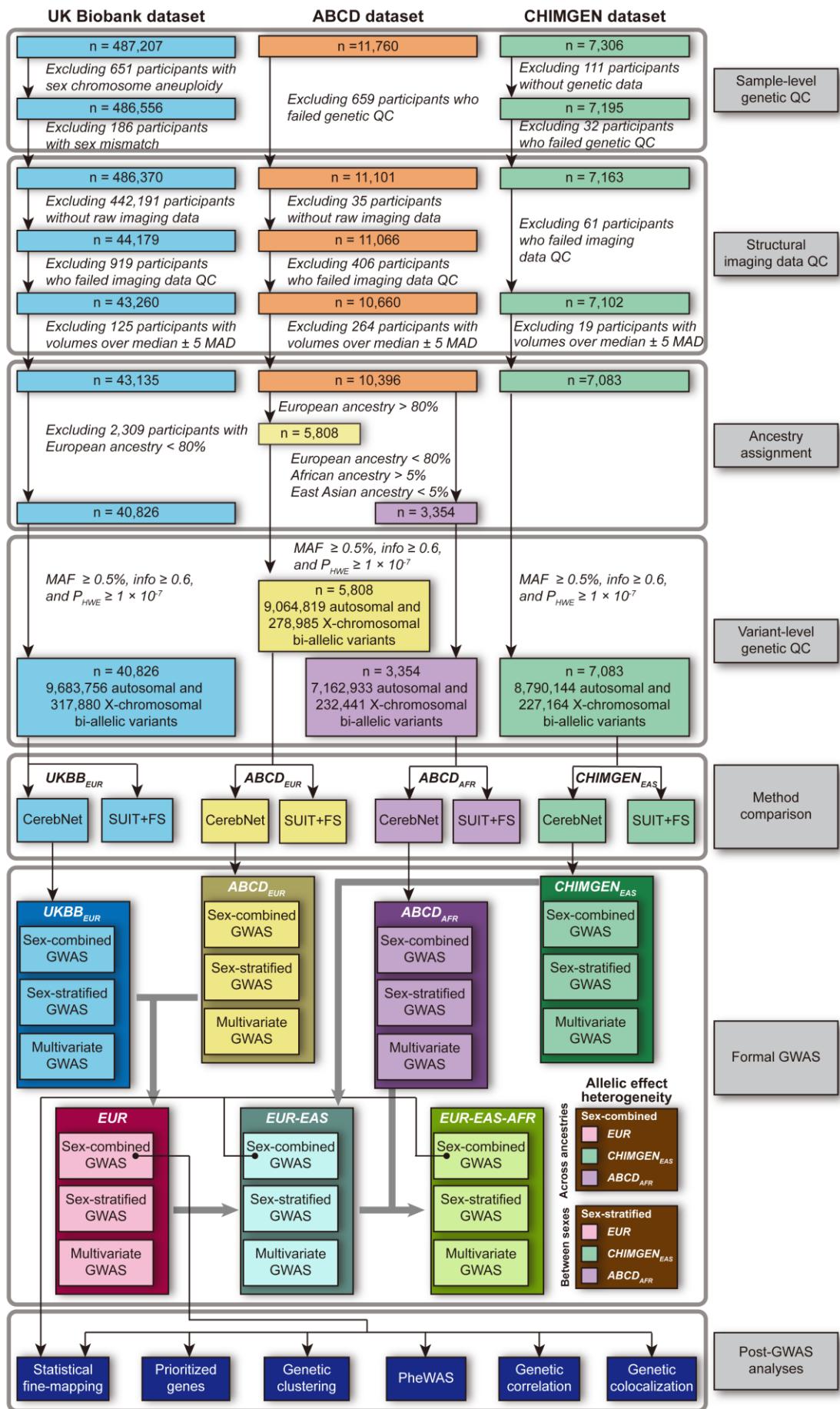
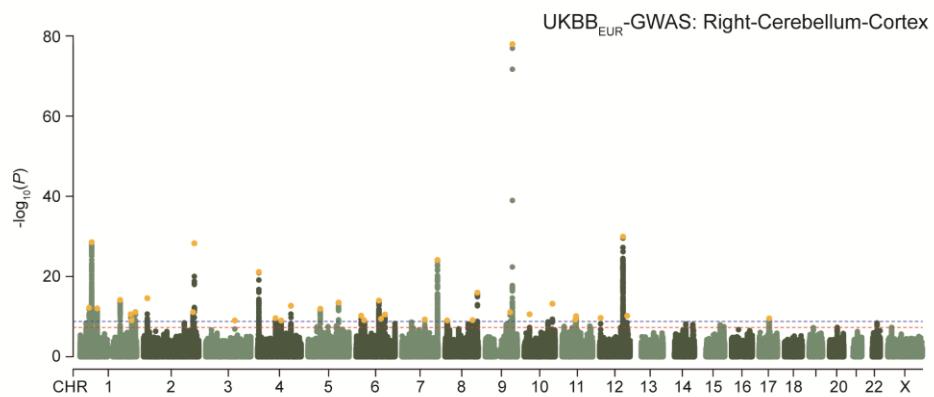
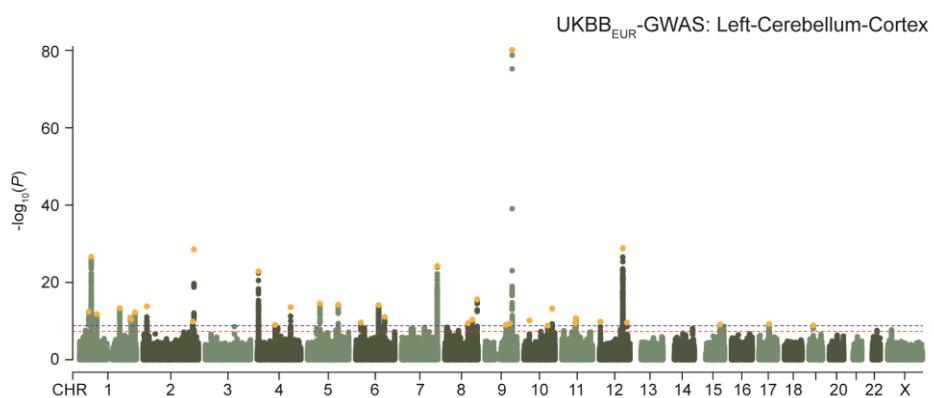
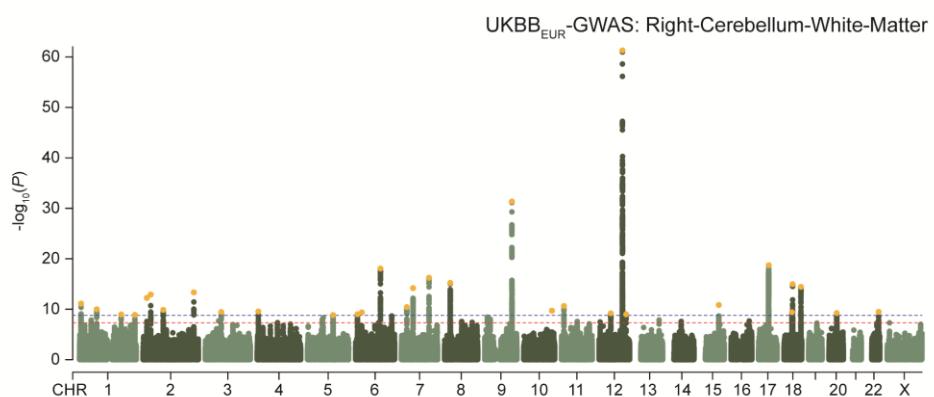
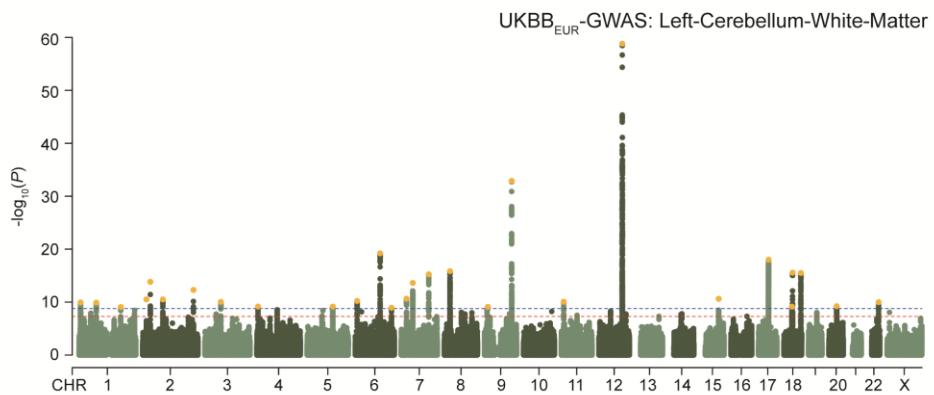
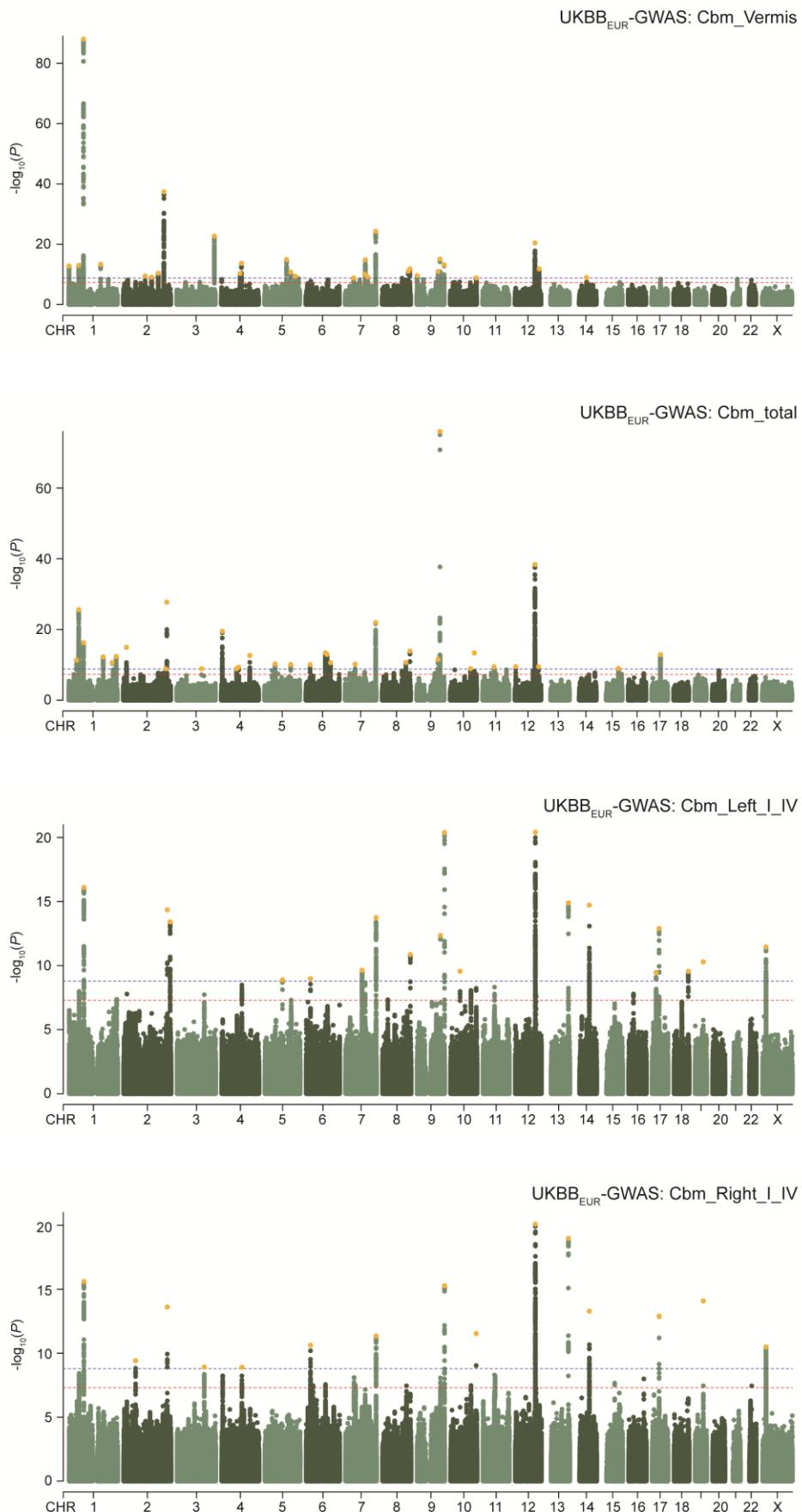
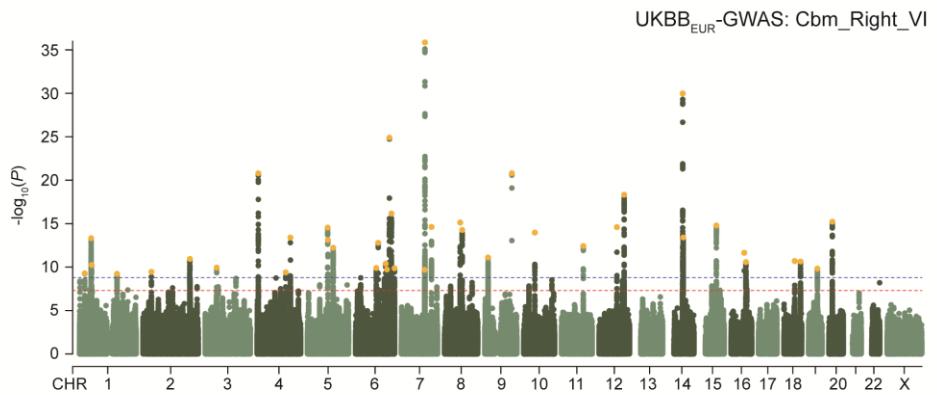
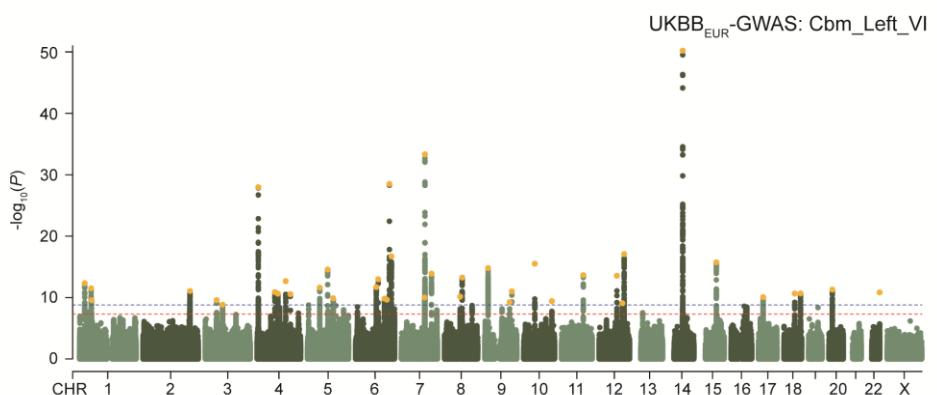
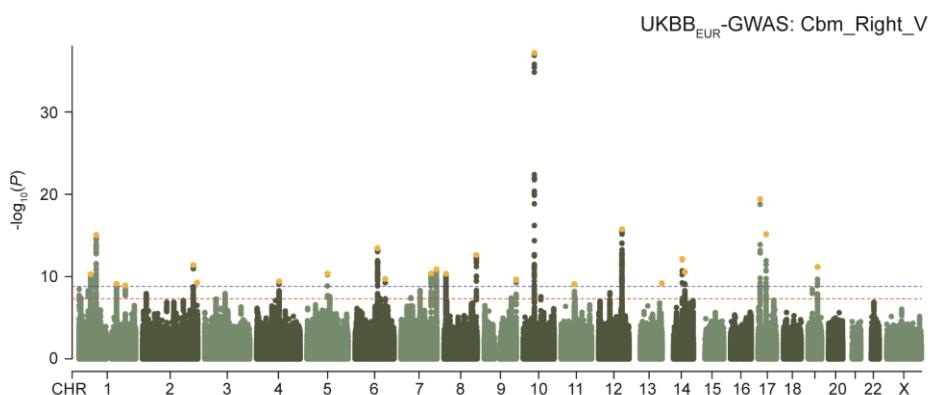
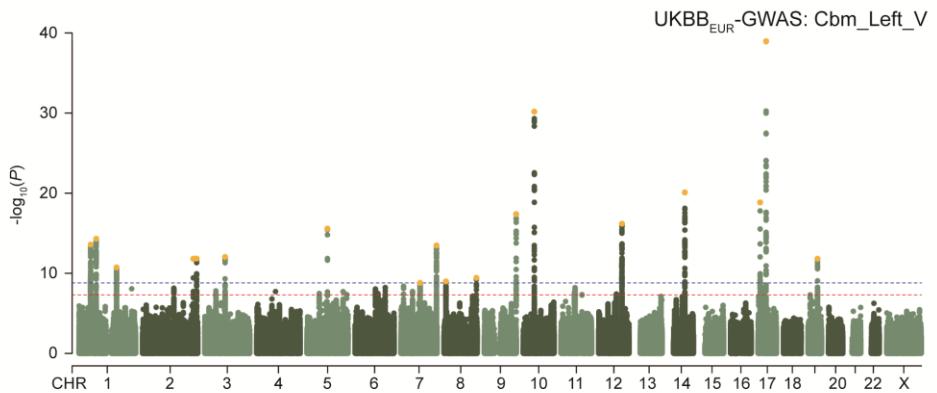


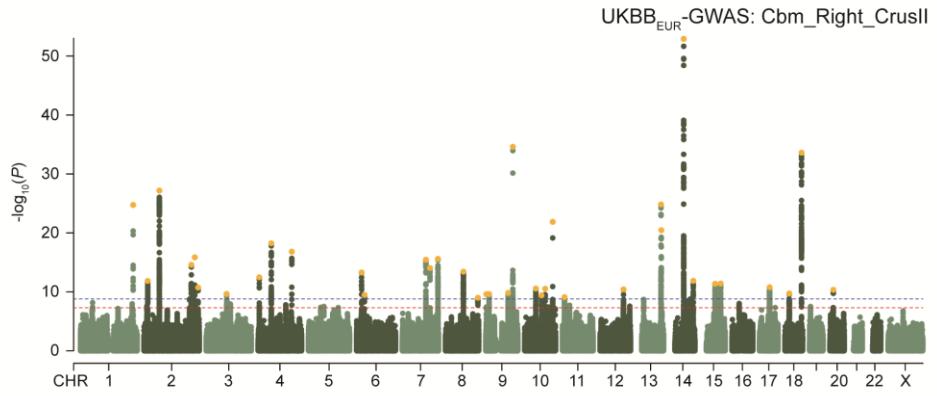
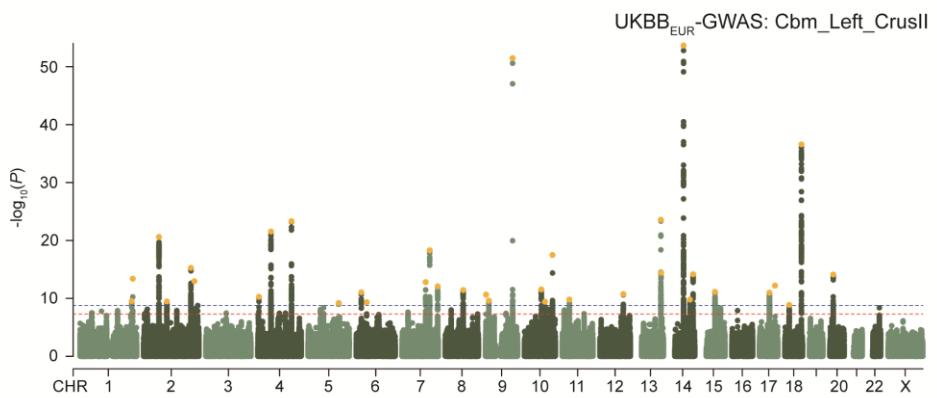
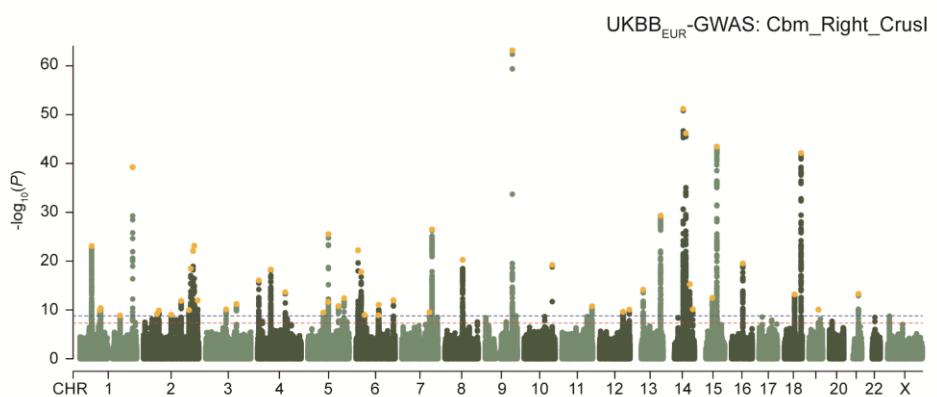
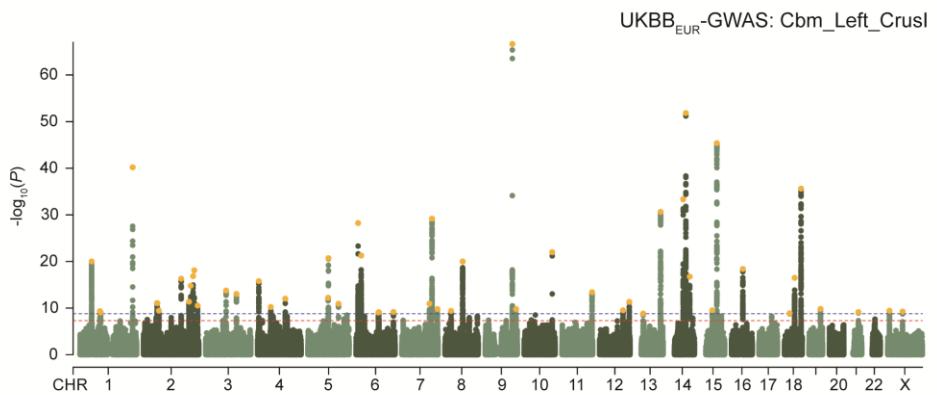
Fig. S2 | Participant selection, quality control, and analysis workflow.

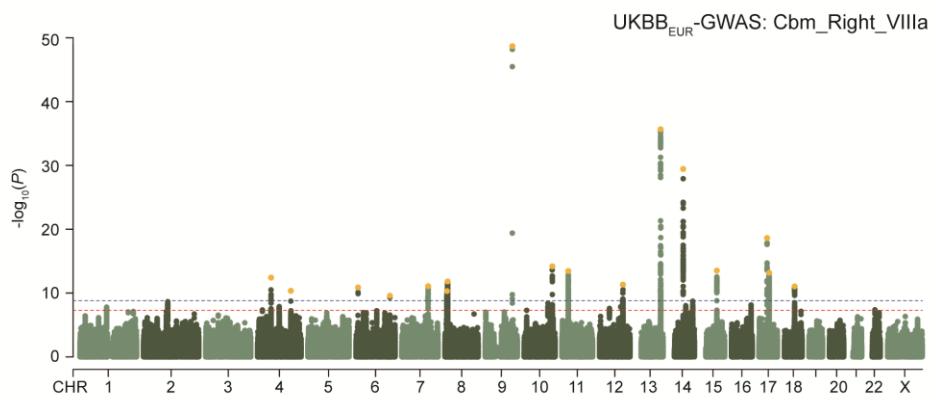
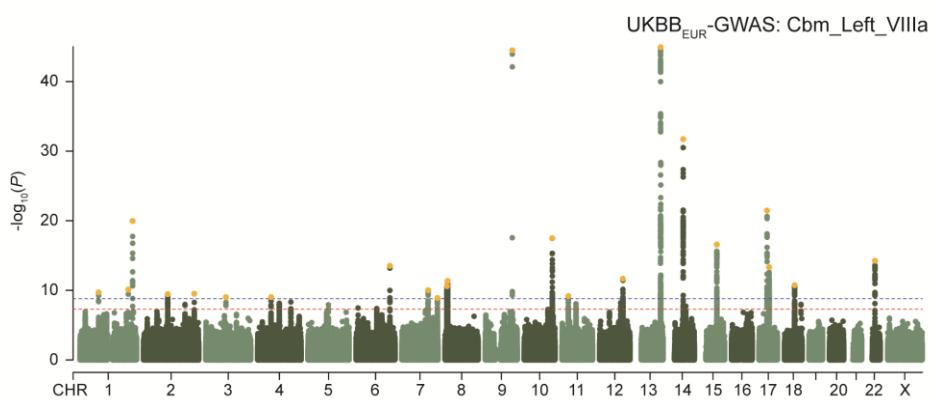
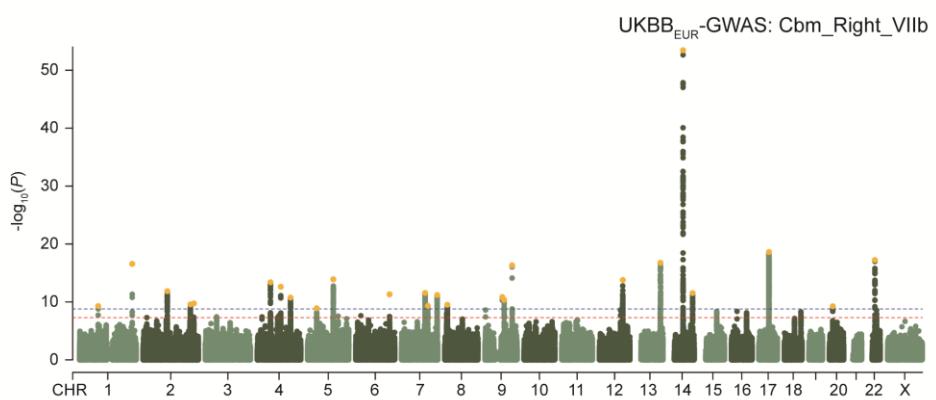
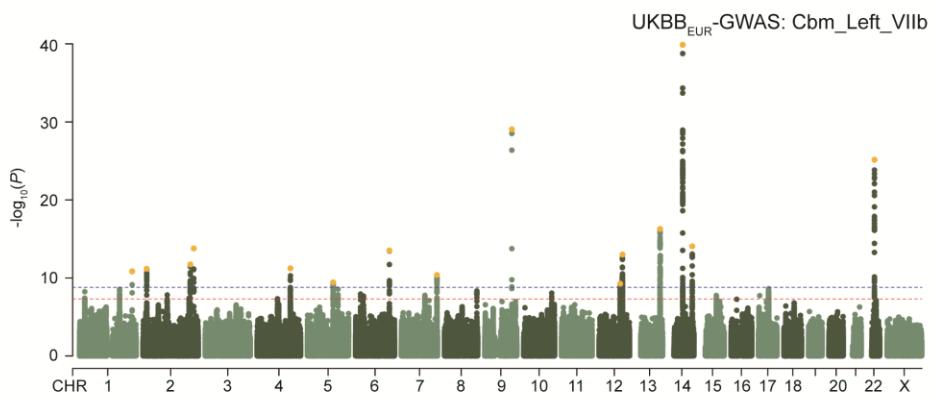
Abbreviations: ABCD, the Adolescent Brain Cognitive Development study; AFR, African ancestry; CHIMGEN, the Chinese Imaging Genetics study; EAS, East Asian ancestry; EUR, European ancestry; FS, FreeSurfer; GWAS, genome-wide association study; HWE, Hardy-Weinberg equilibrium; info, imputation information score; MAD, median absolute deviation; MAF, minor allele frequency; PheWAS, phenome-wide association study; QC, quality control; SUIT, spatially unbiased infra-tentorial; UKBB, the UK Biobank study.

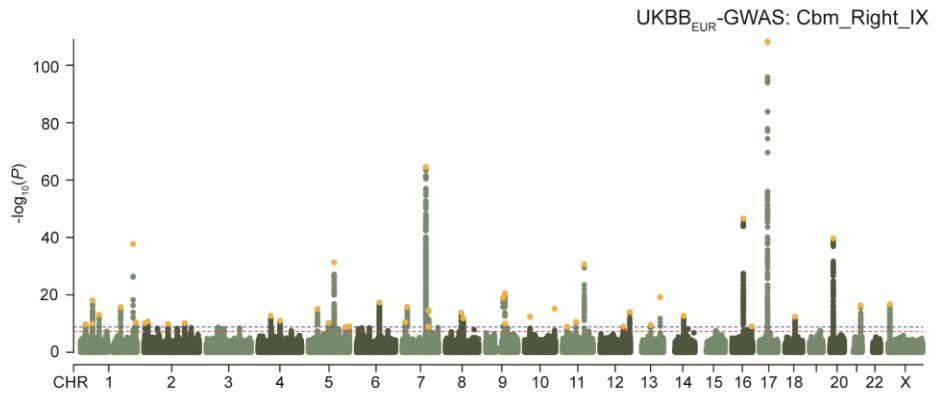
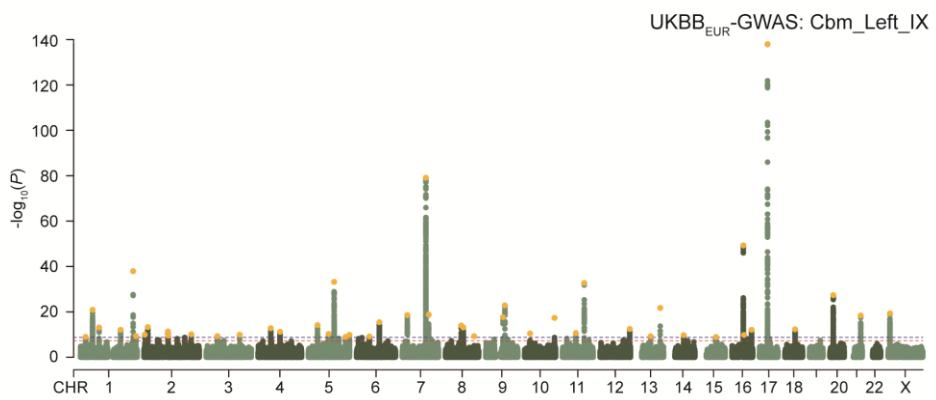
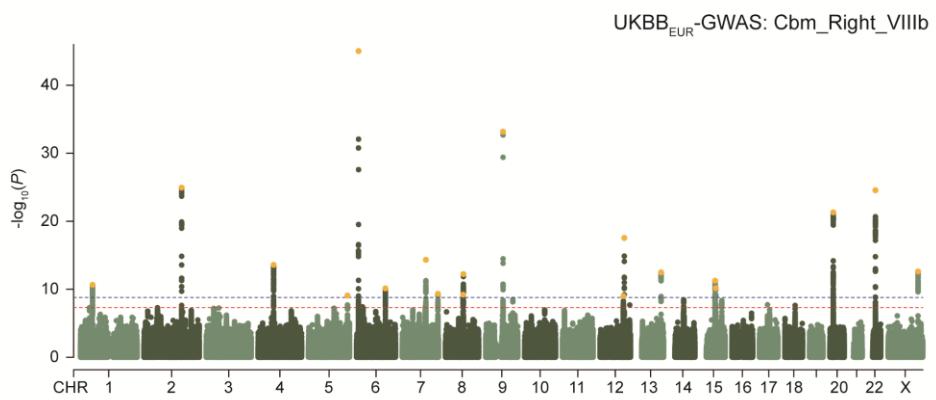
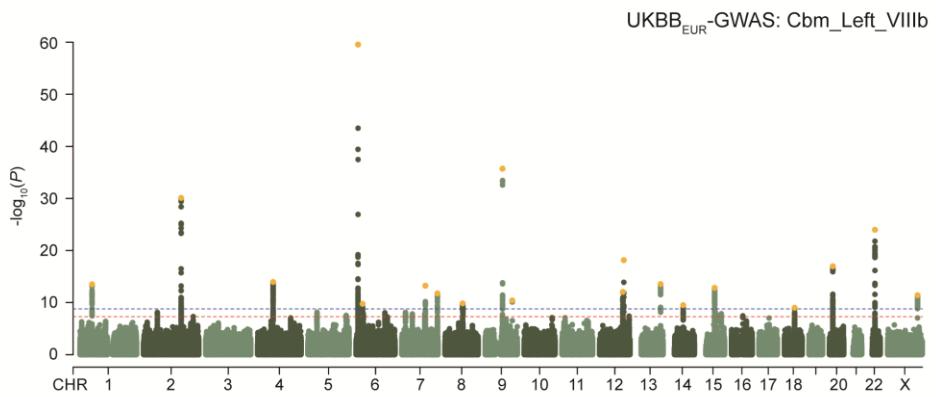


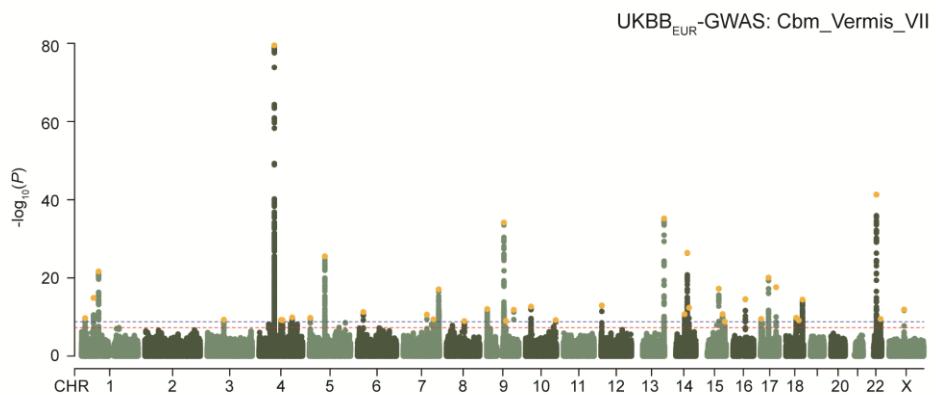
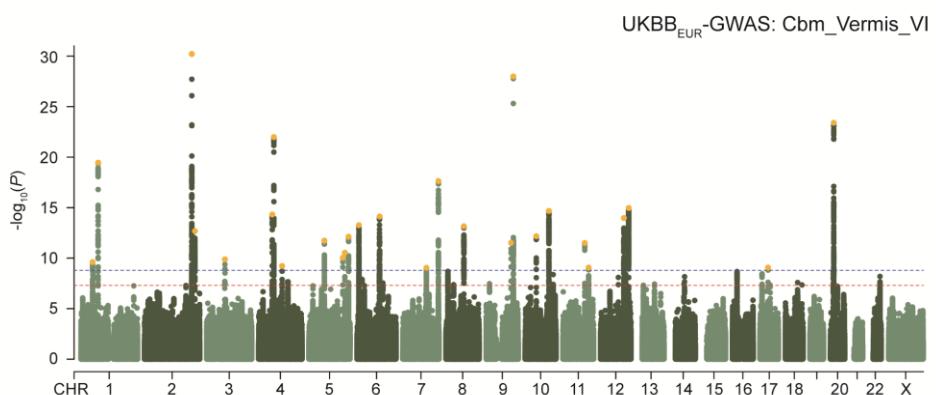
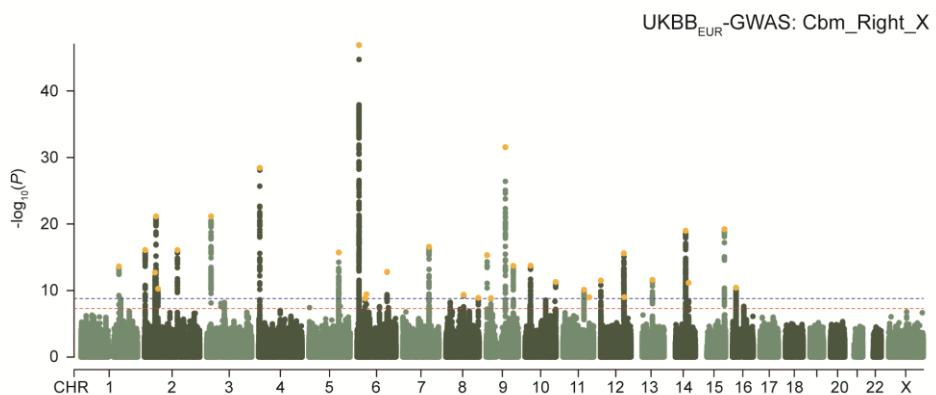
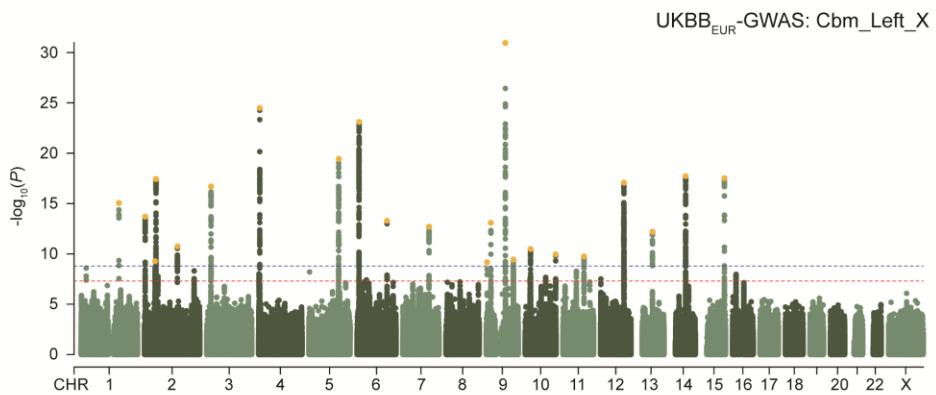


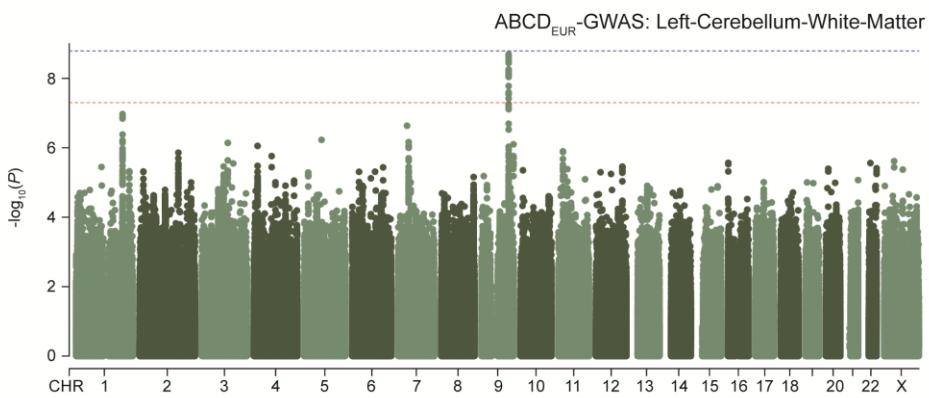
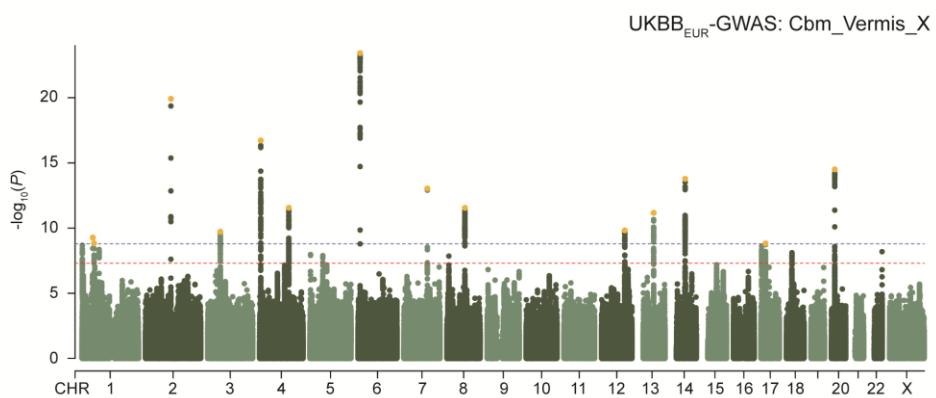
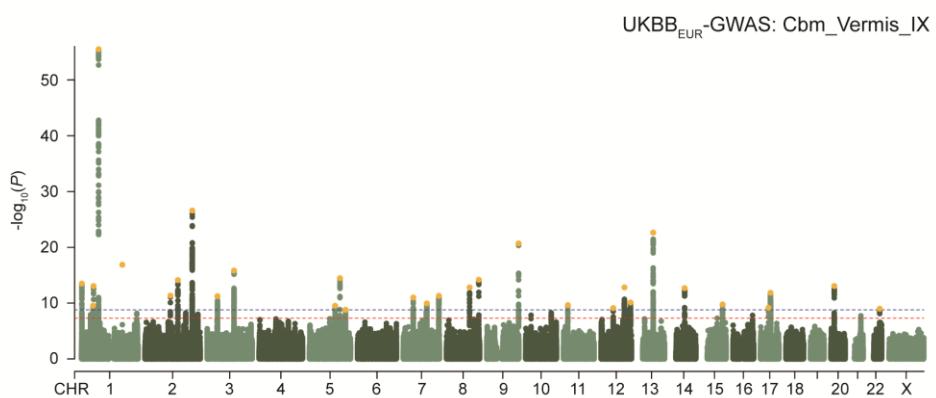
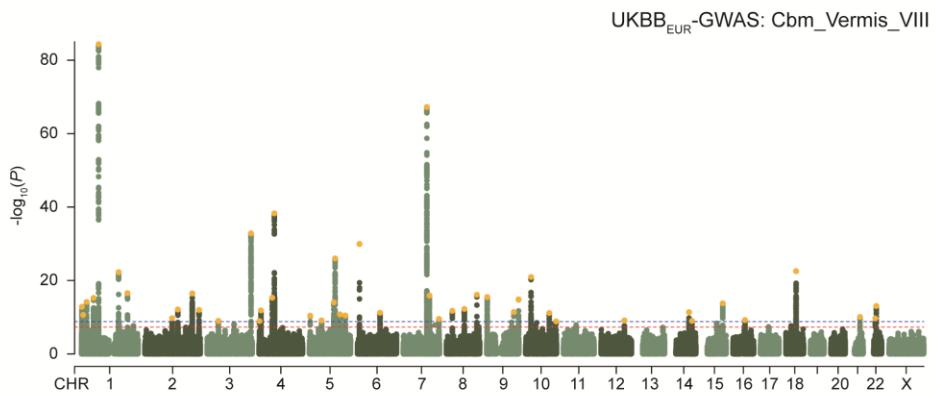


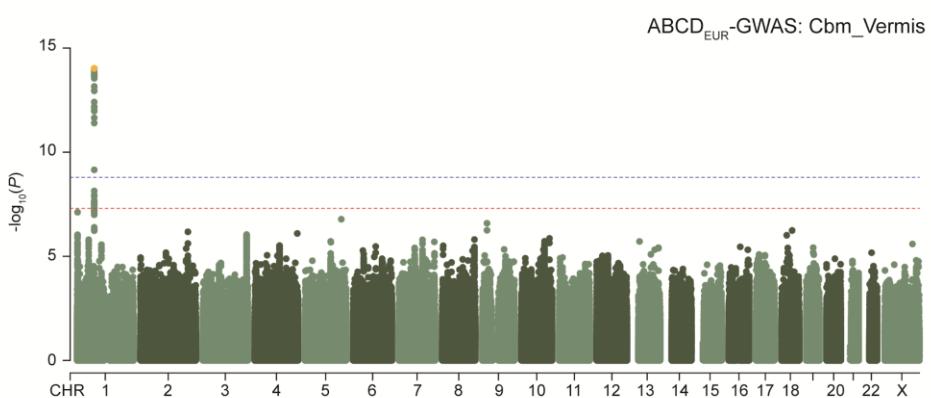
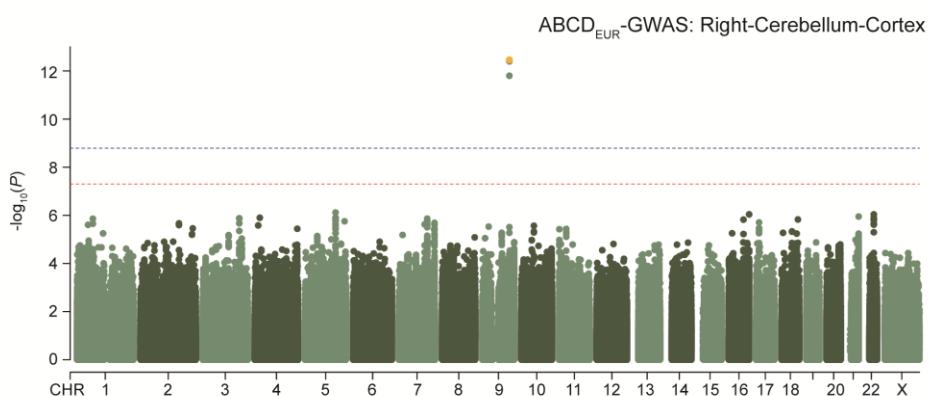
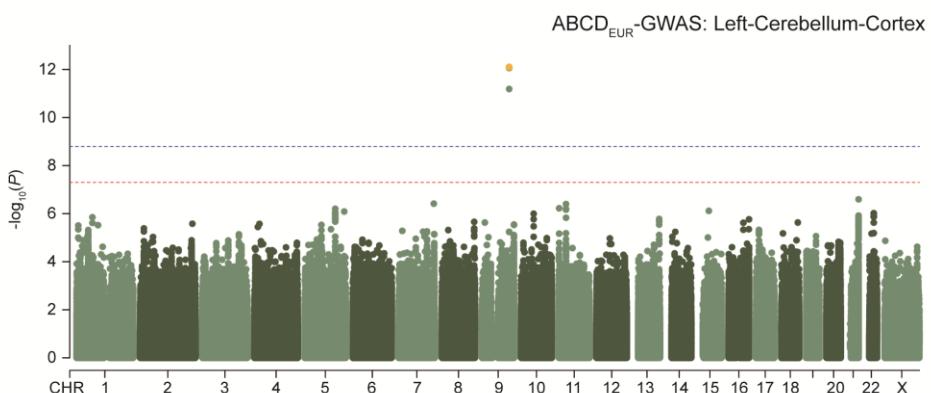
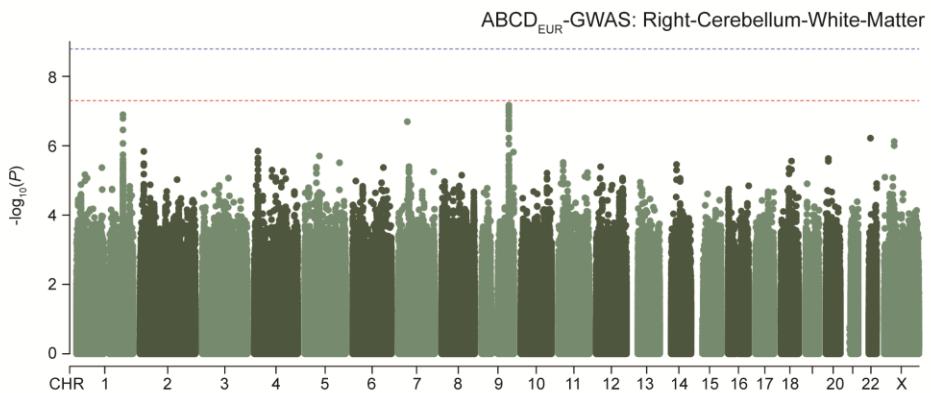


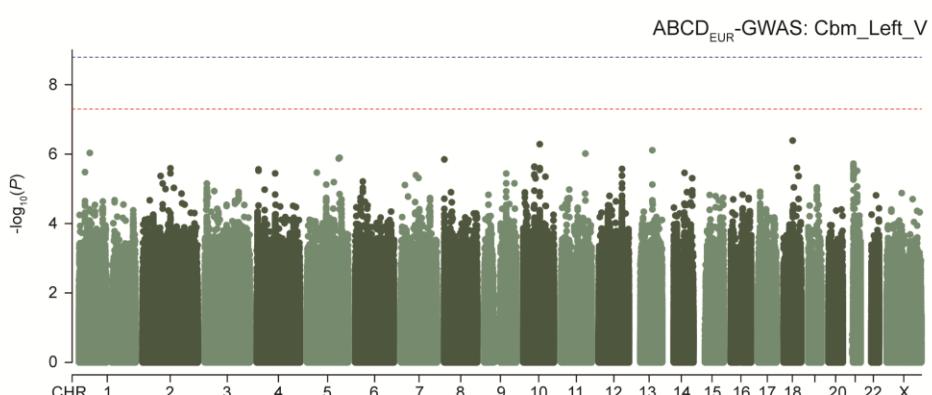
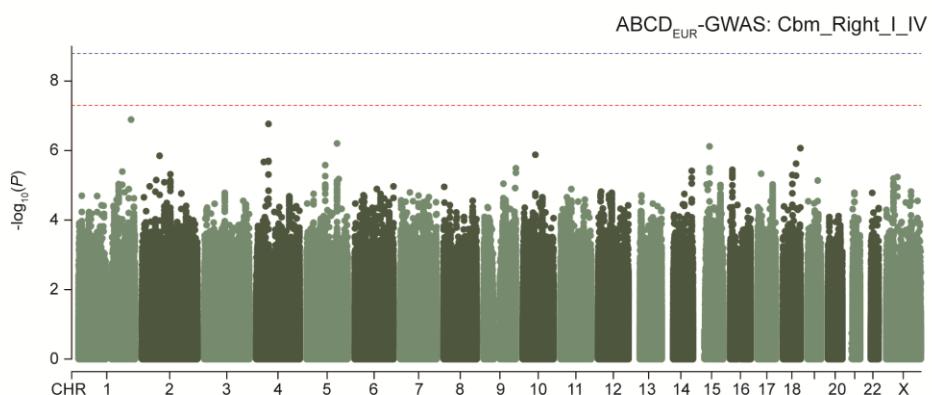
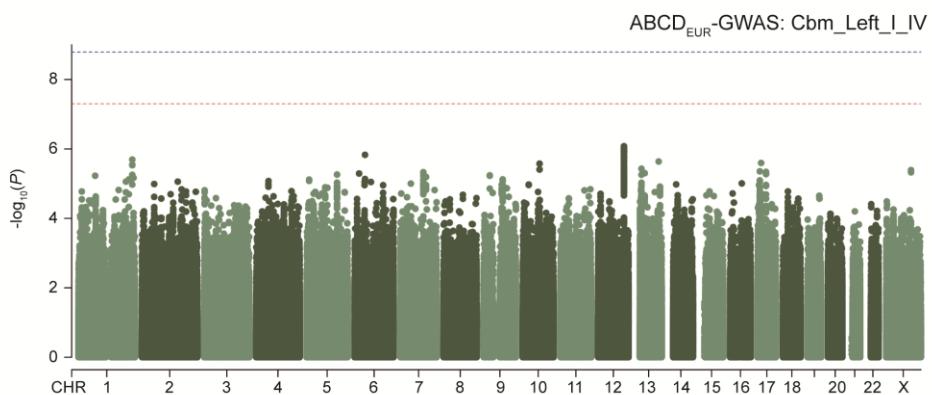
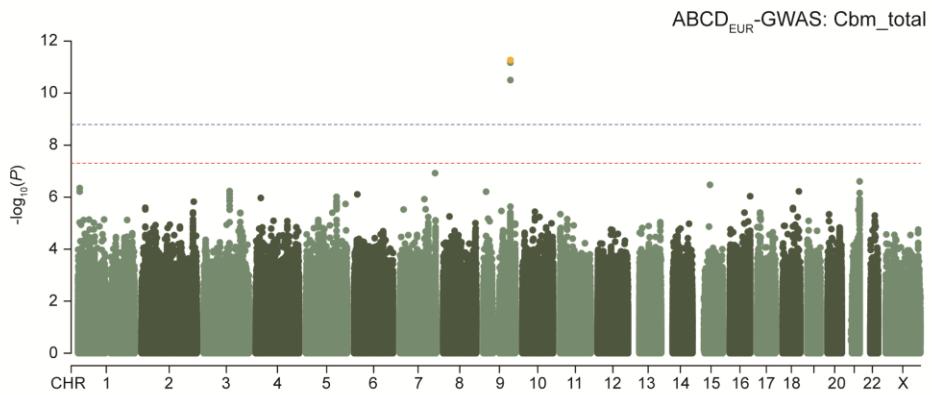


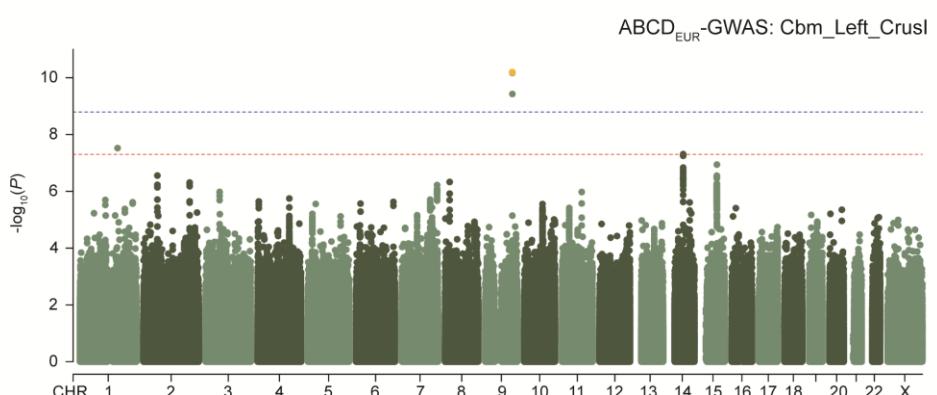
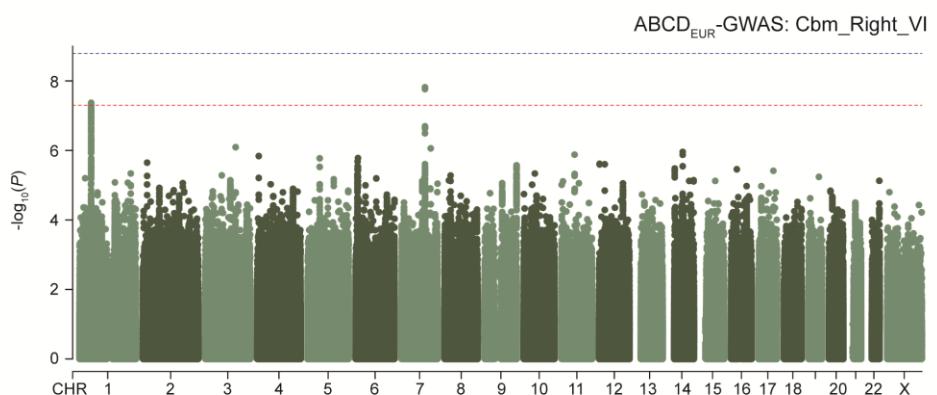
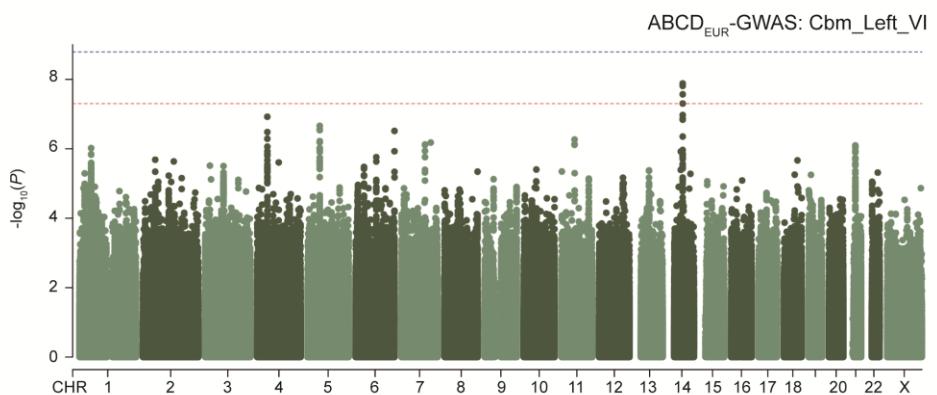
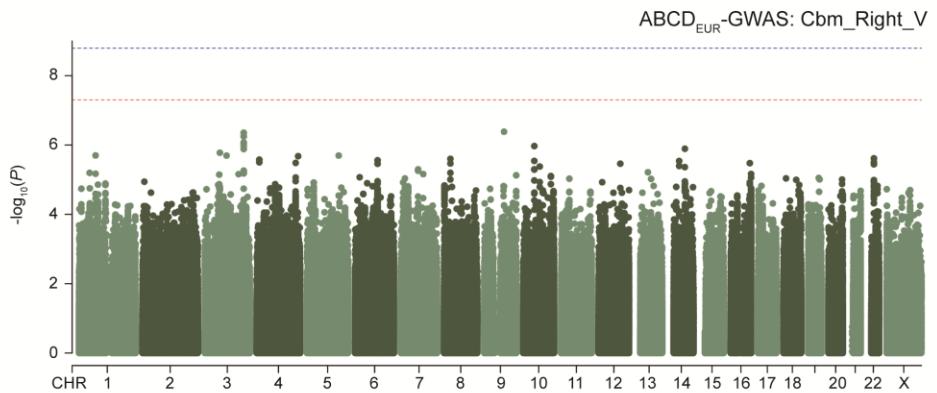


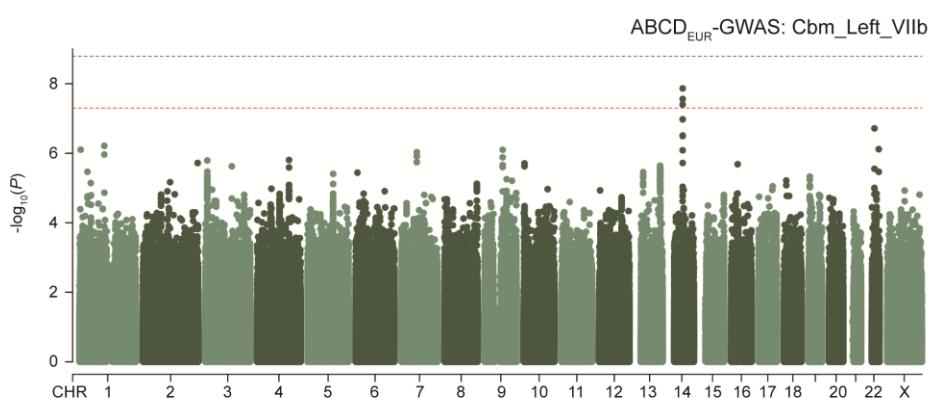
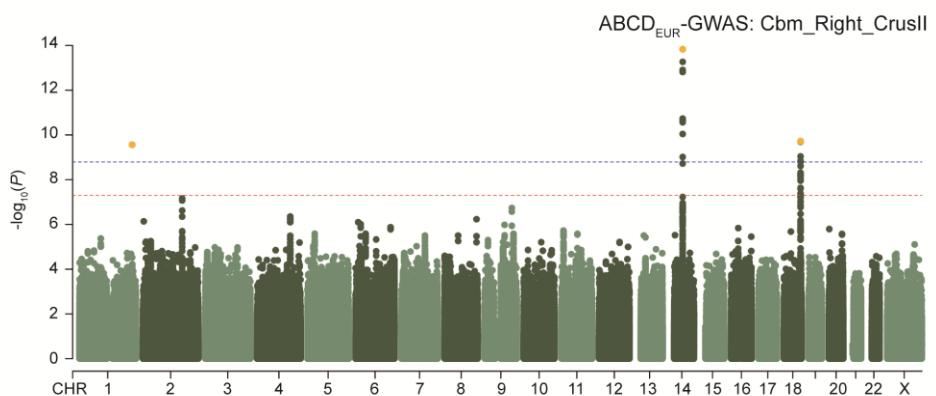
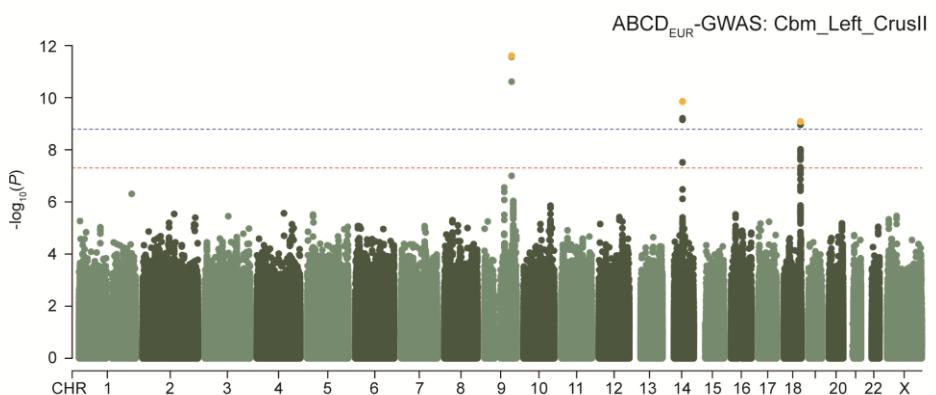
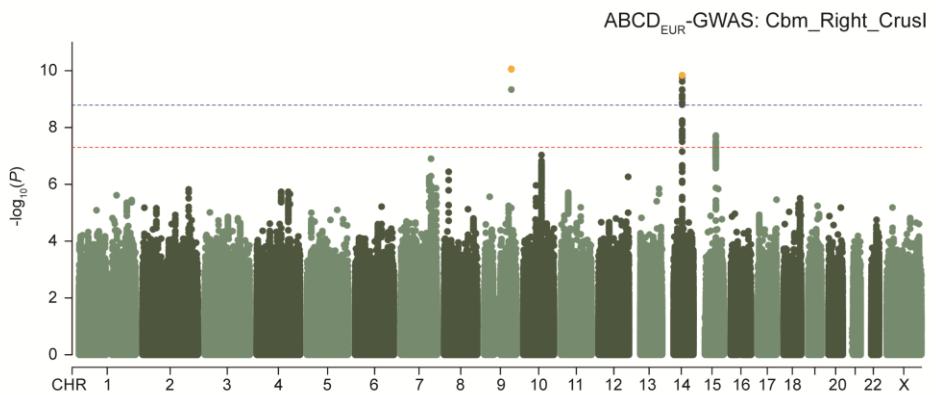


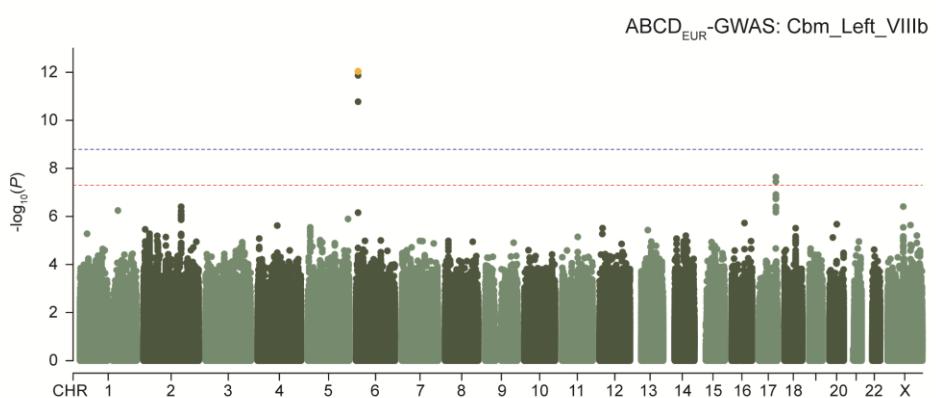
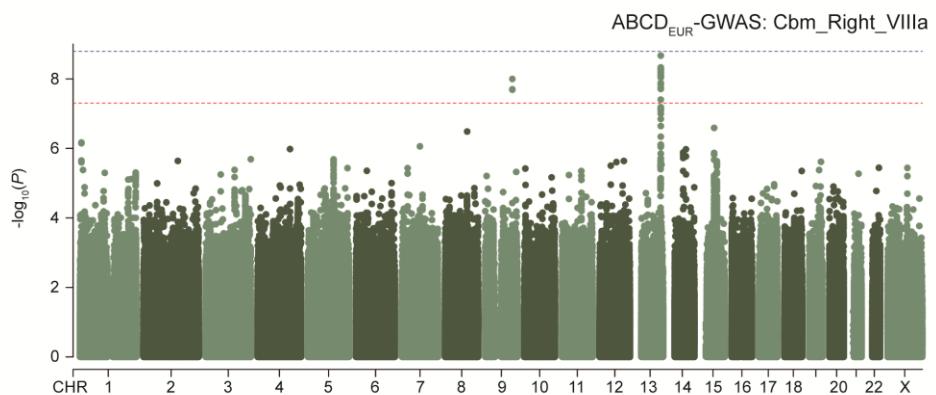
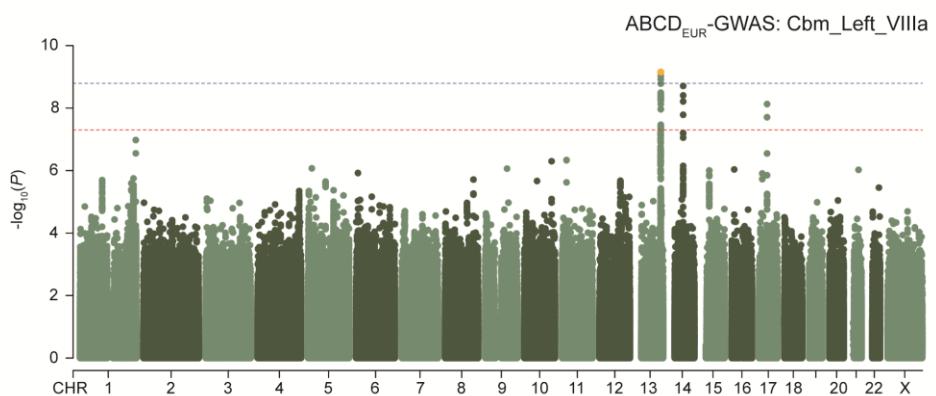
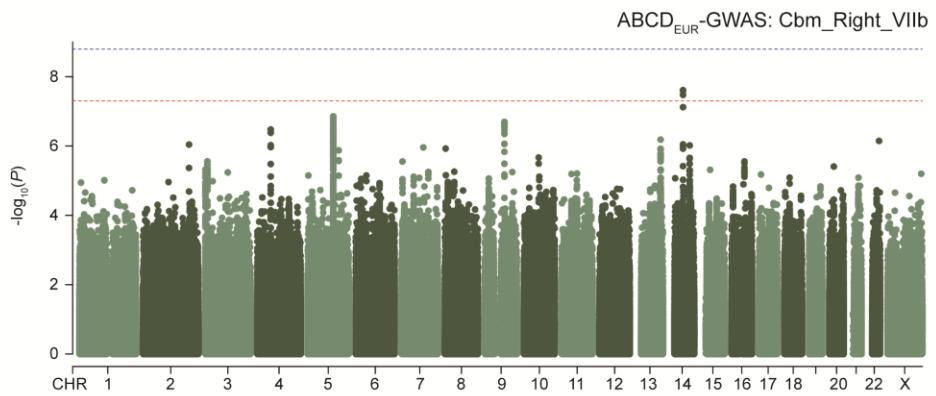


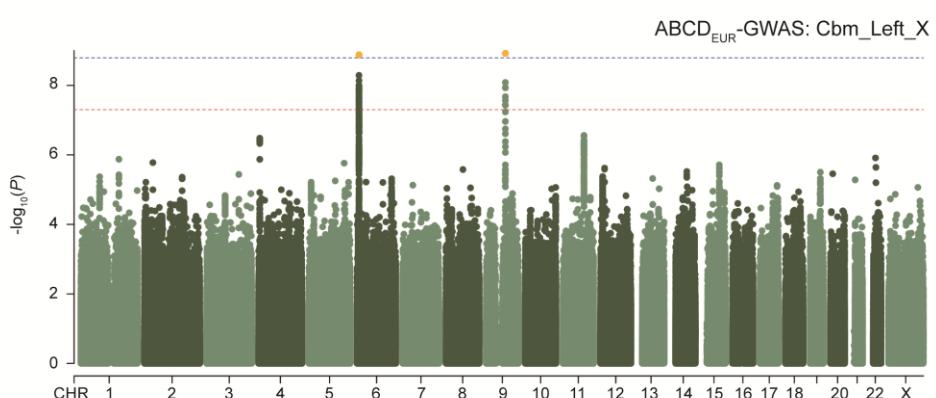
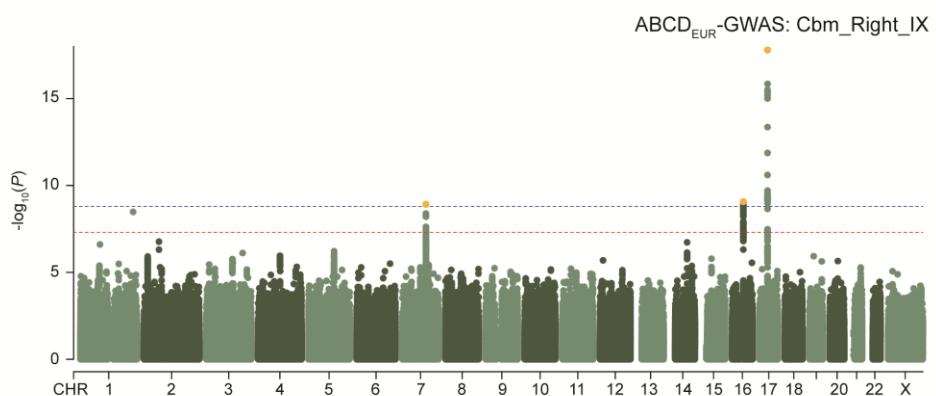
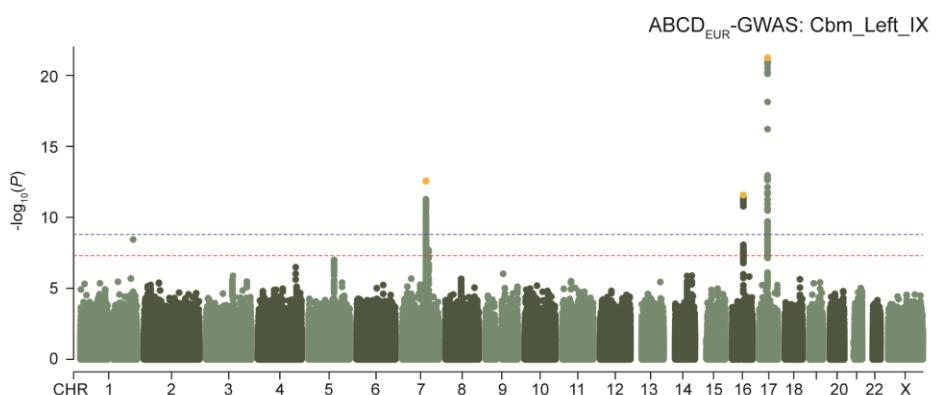
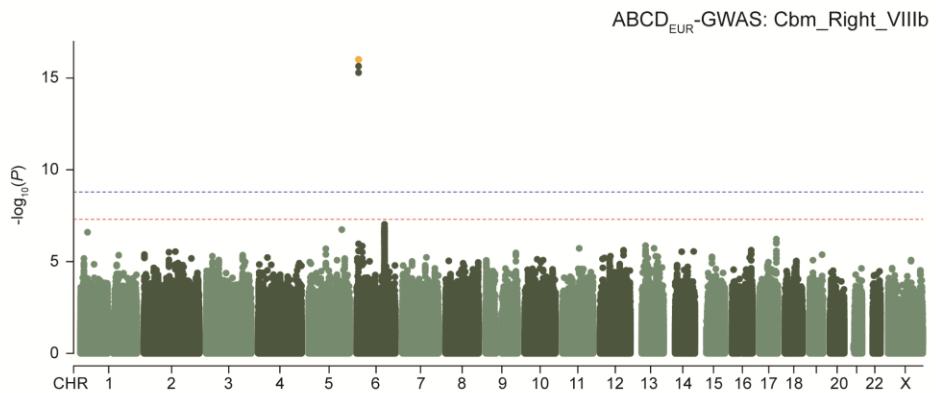


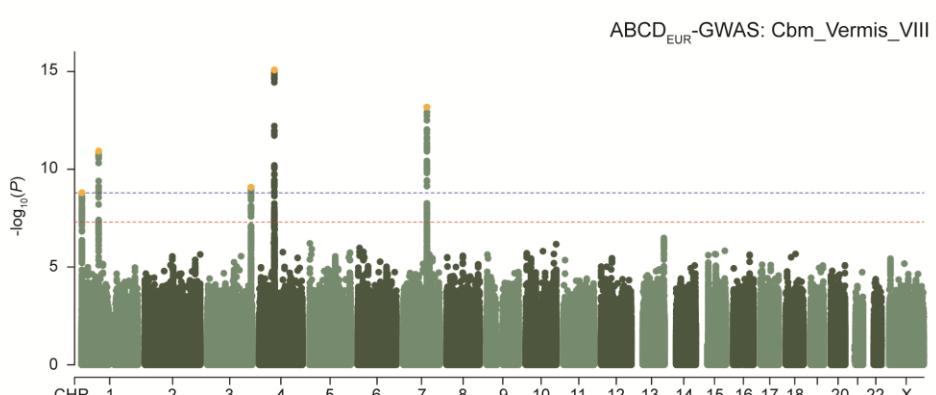
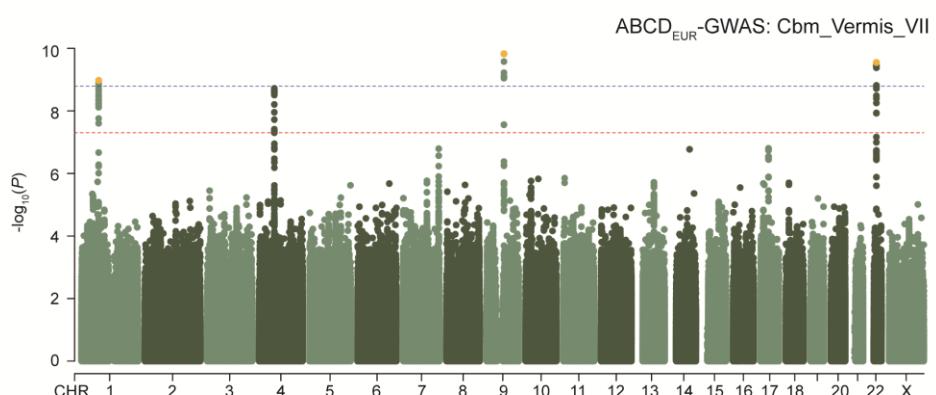
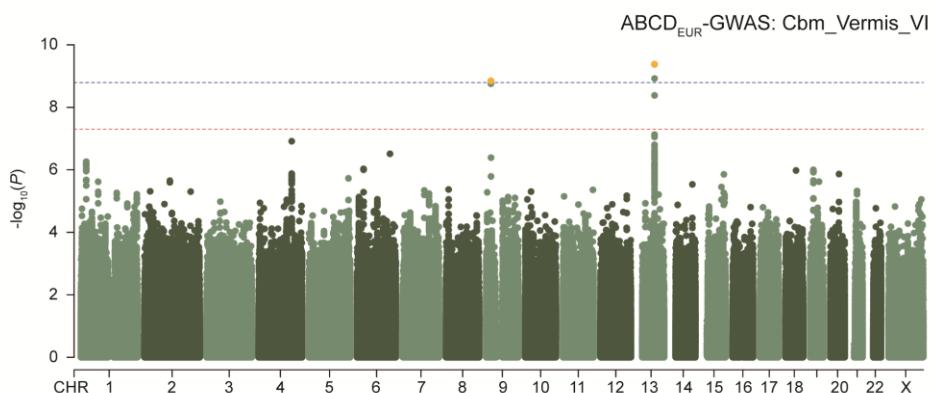
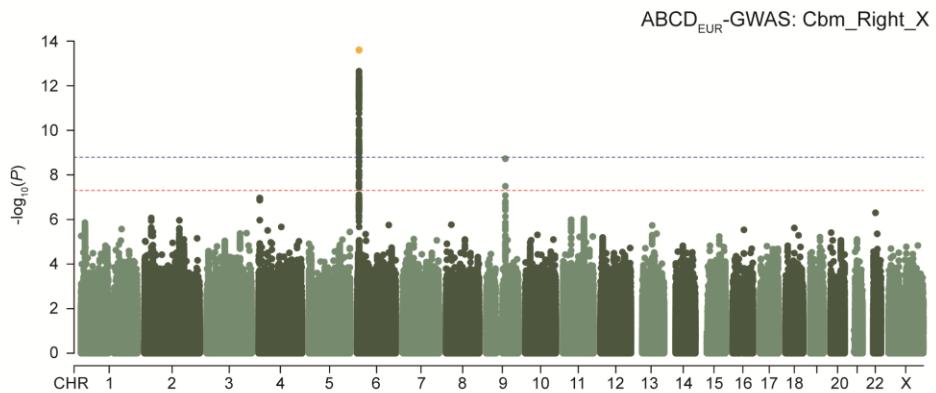


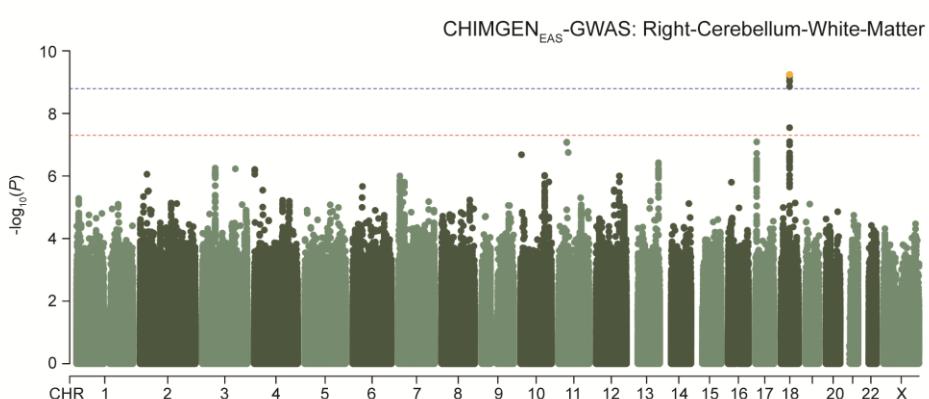
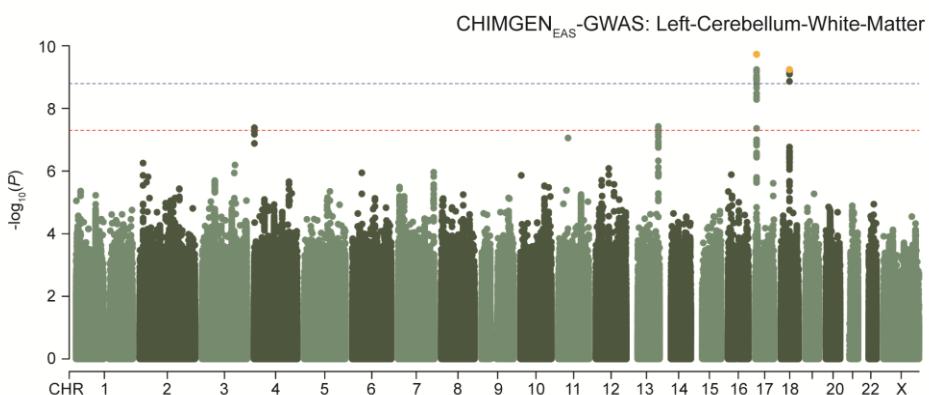
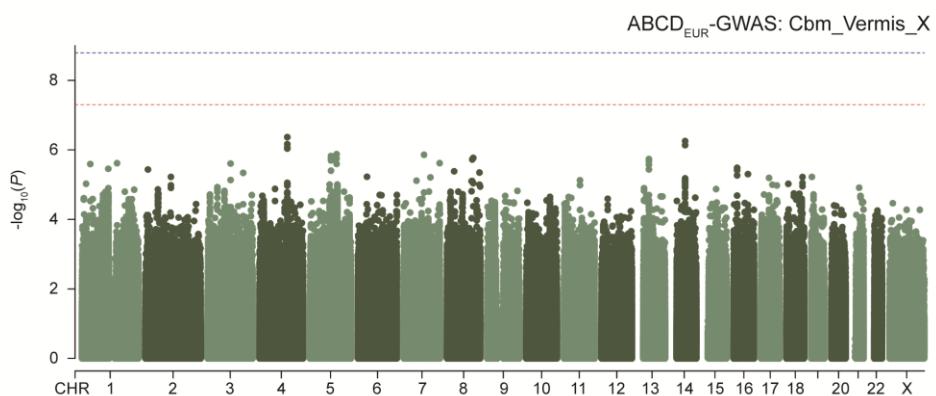
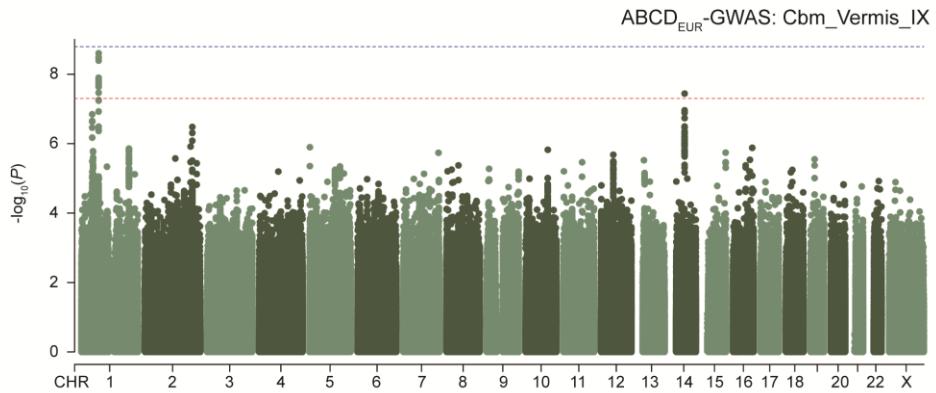


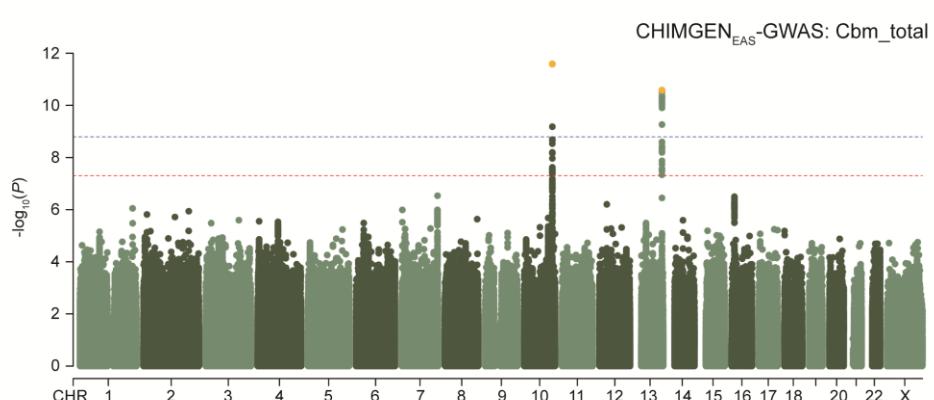
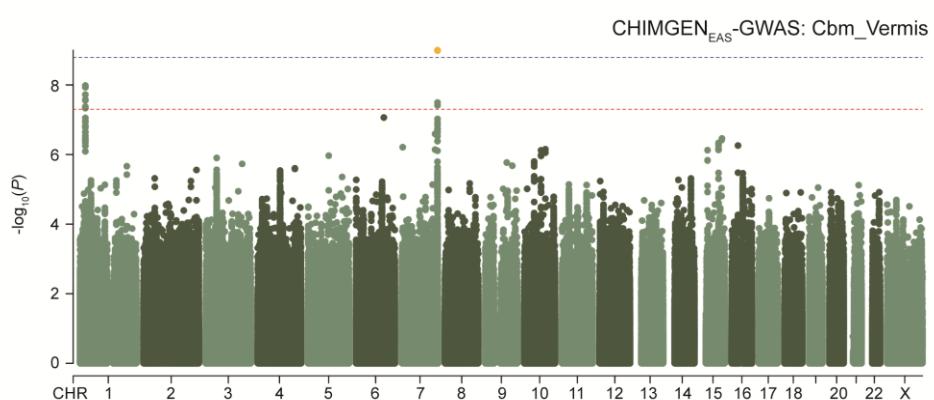
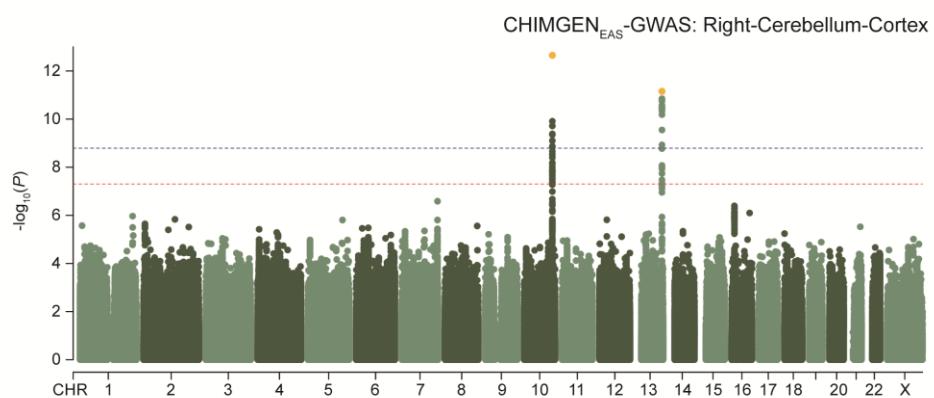
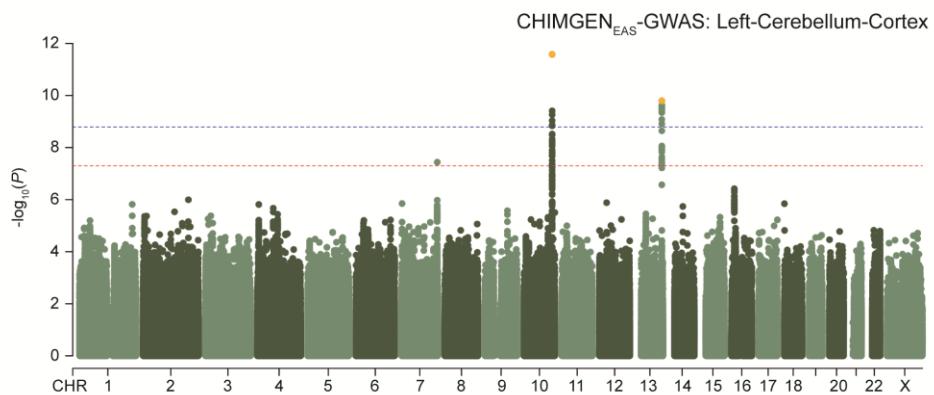


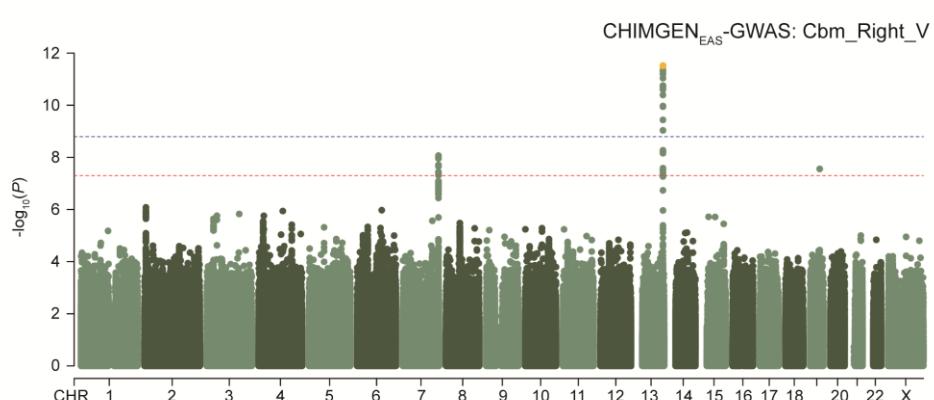
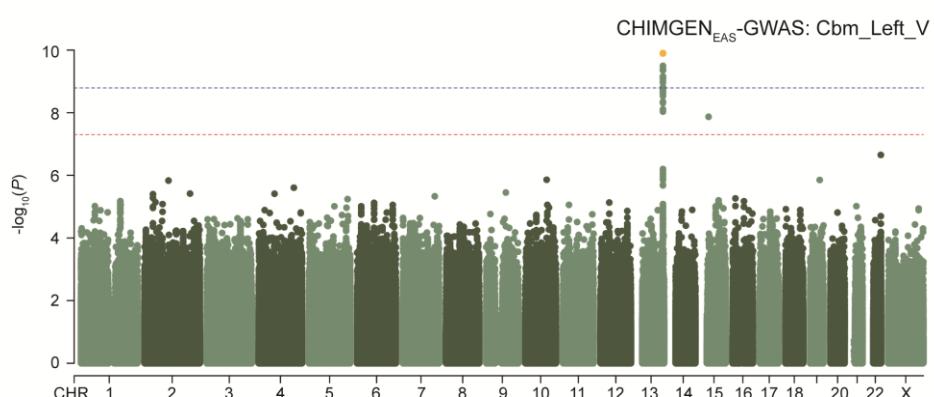
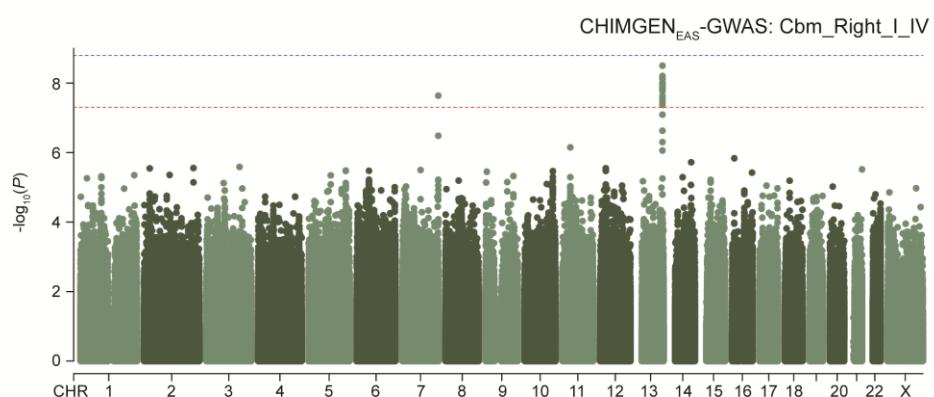
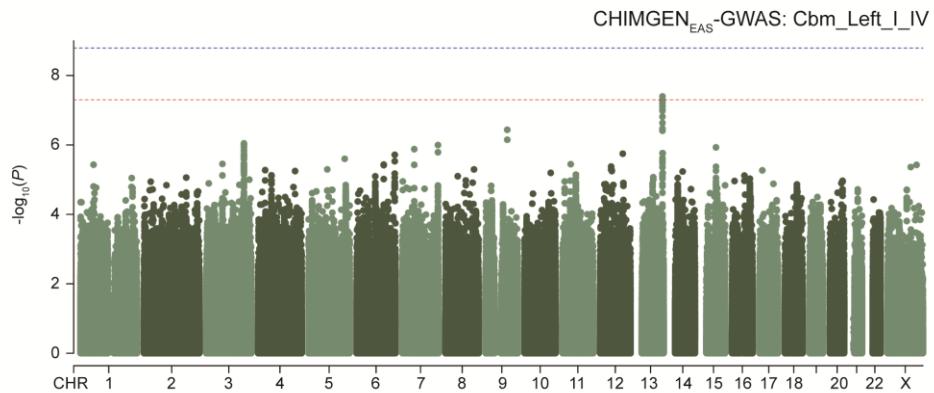


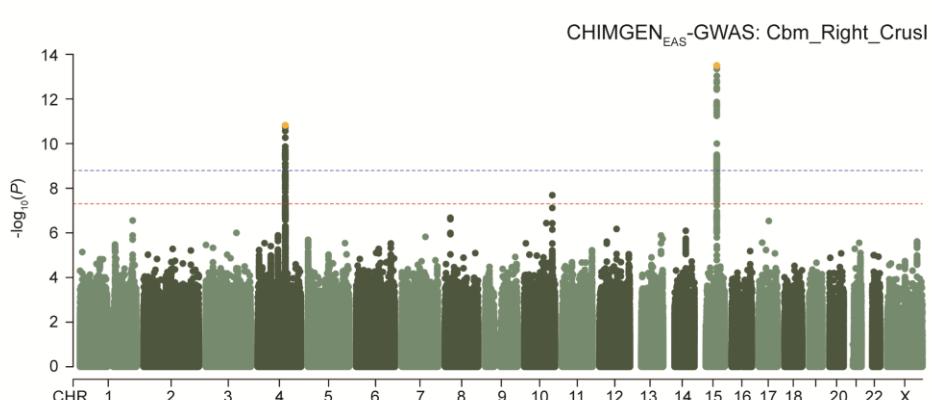
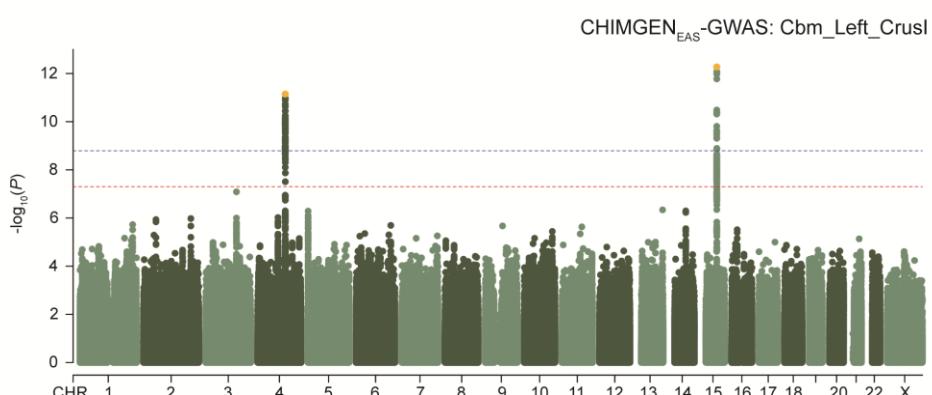
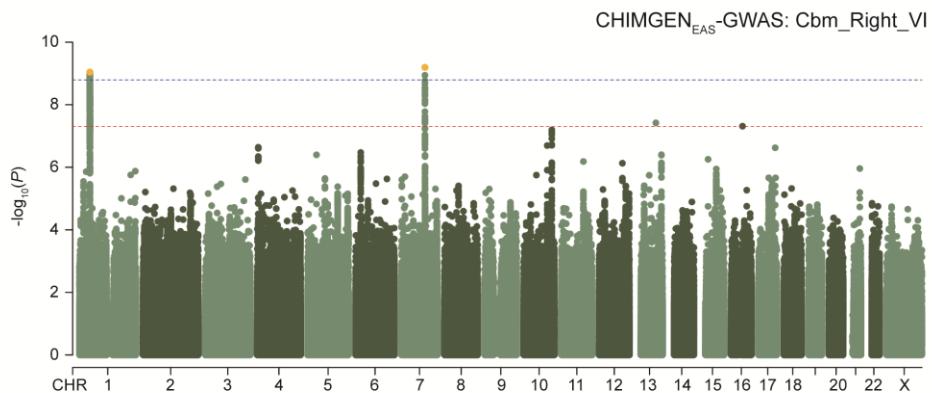
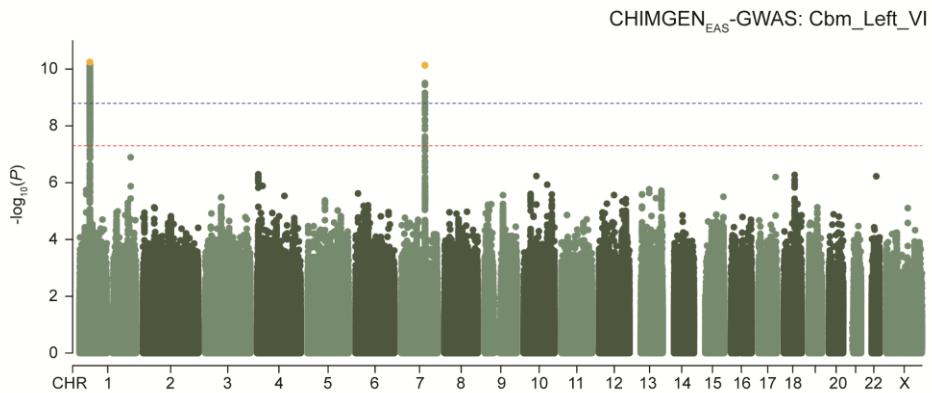


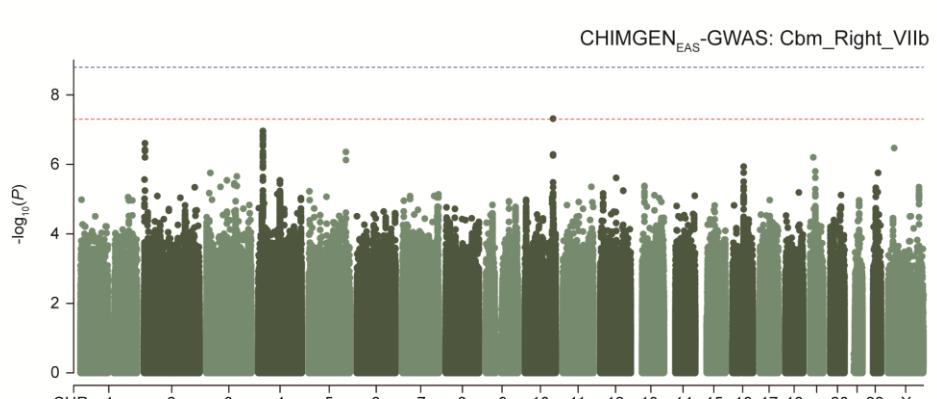
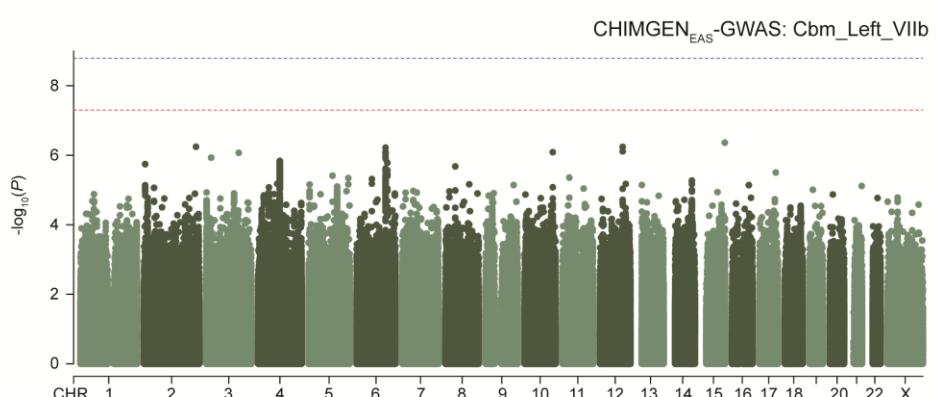
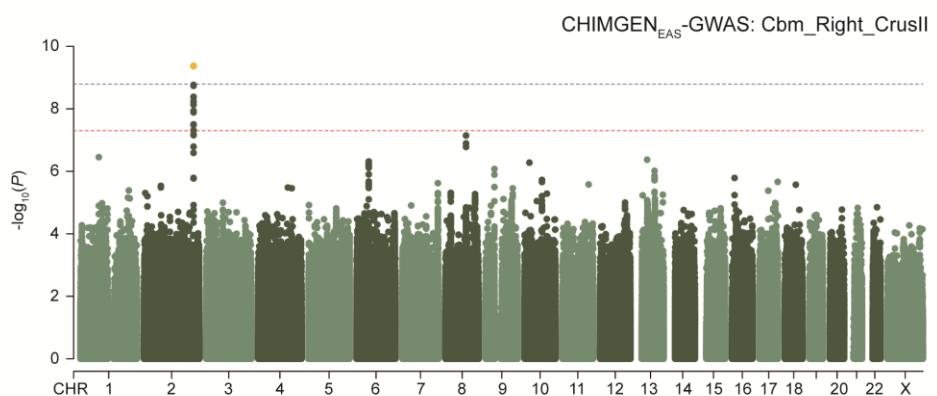
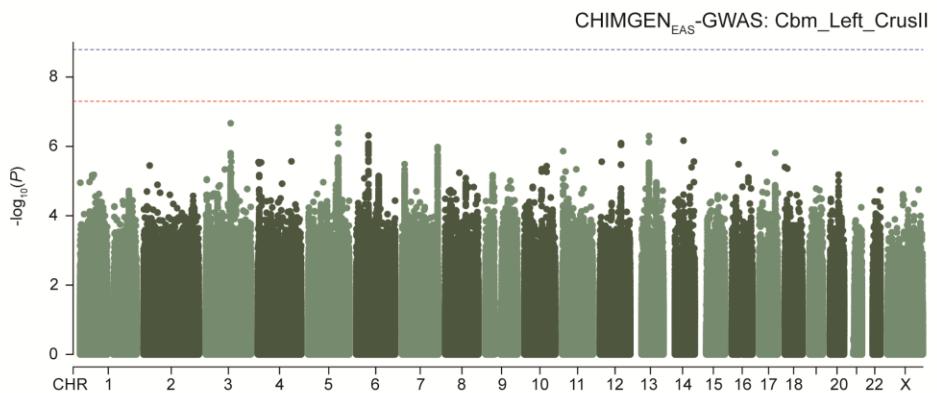


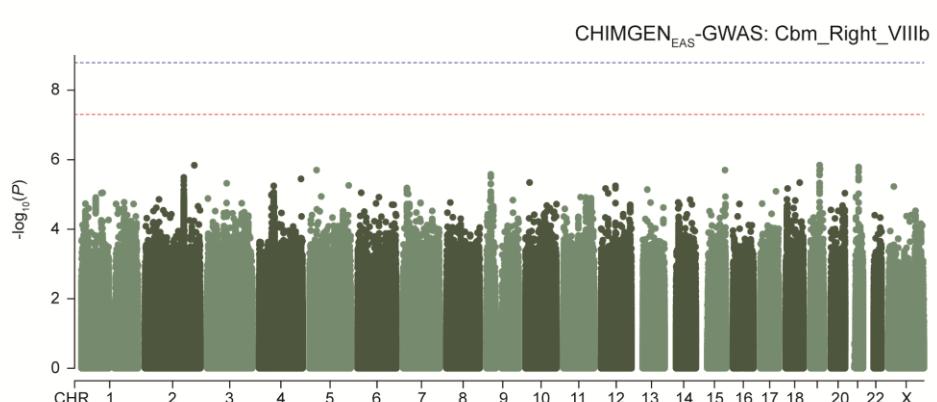
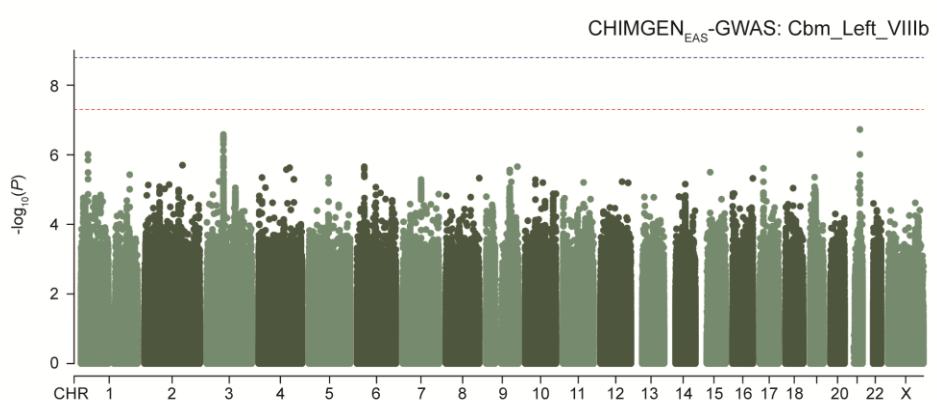
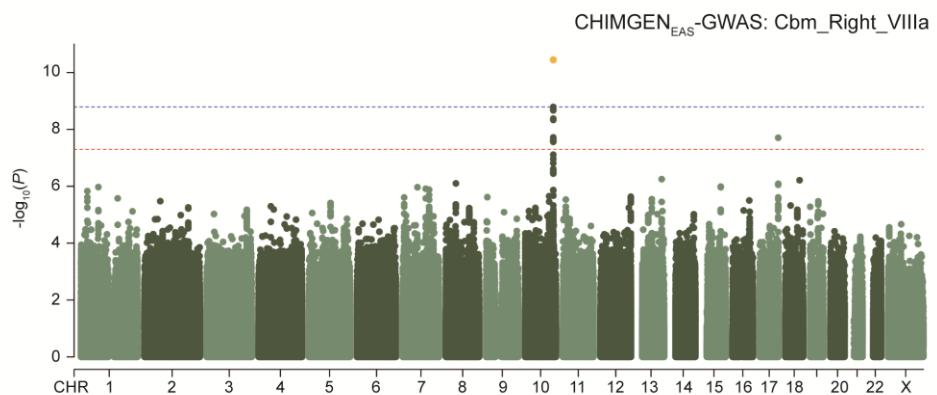
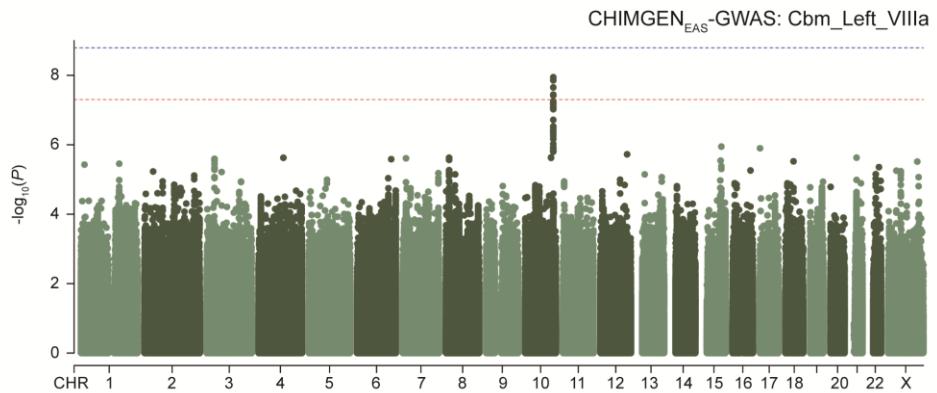


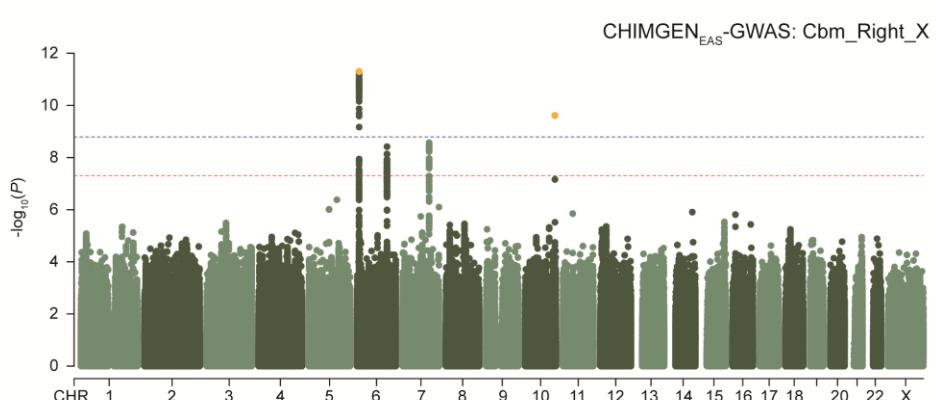
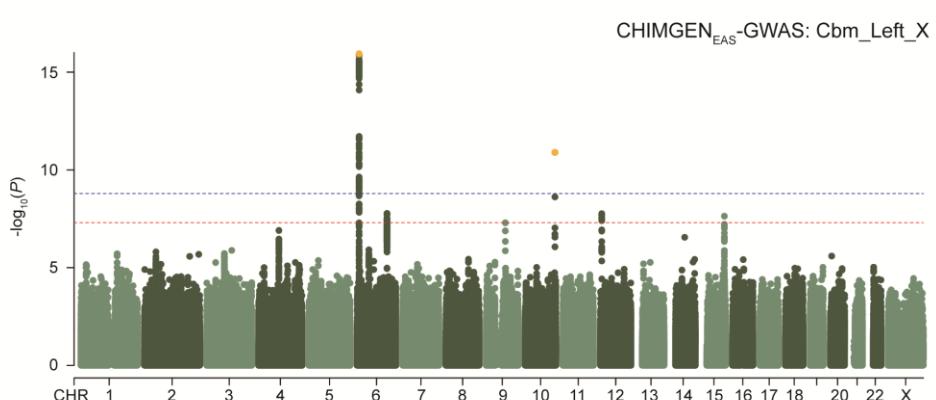
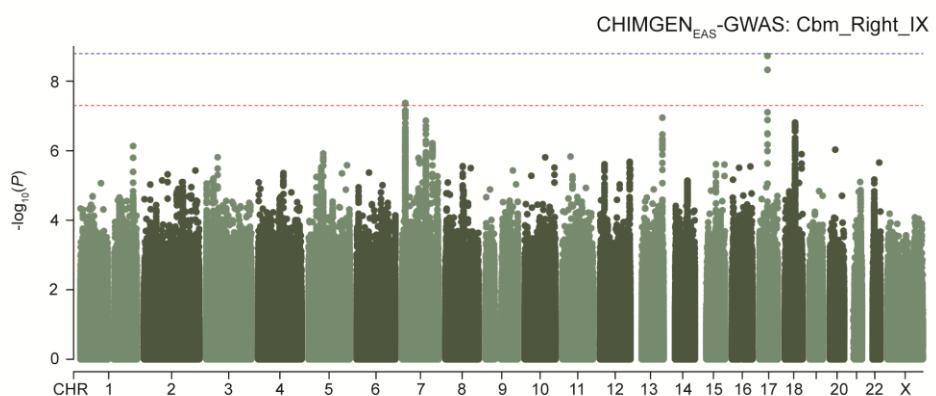
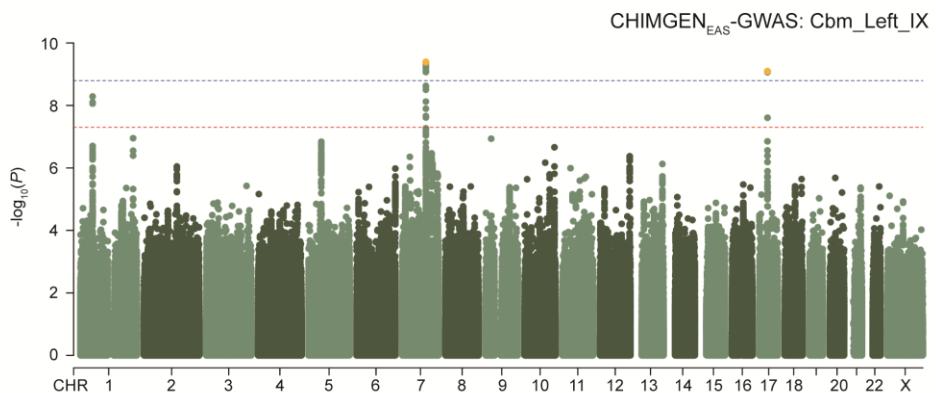


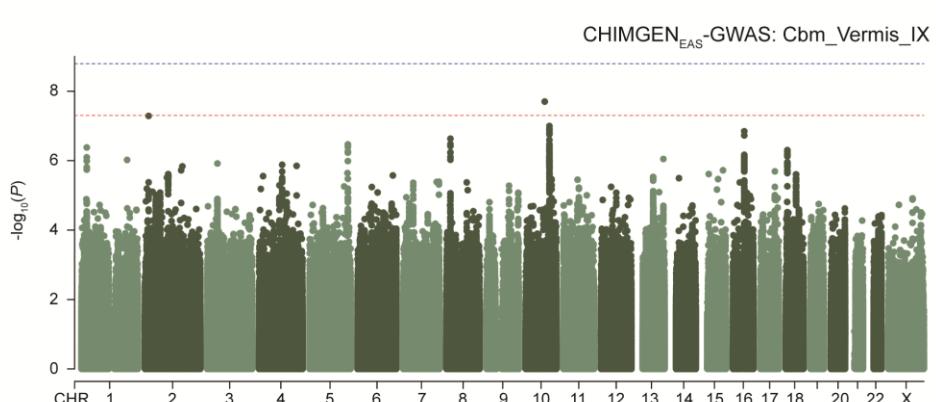
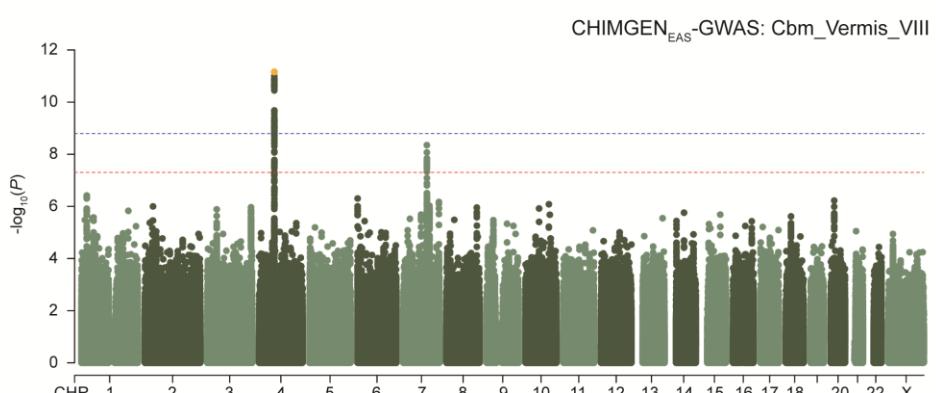
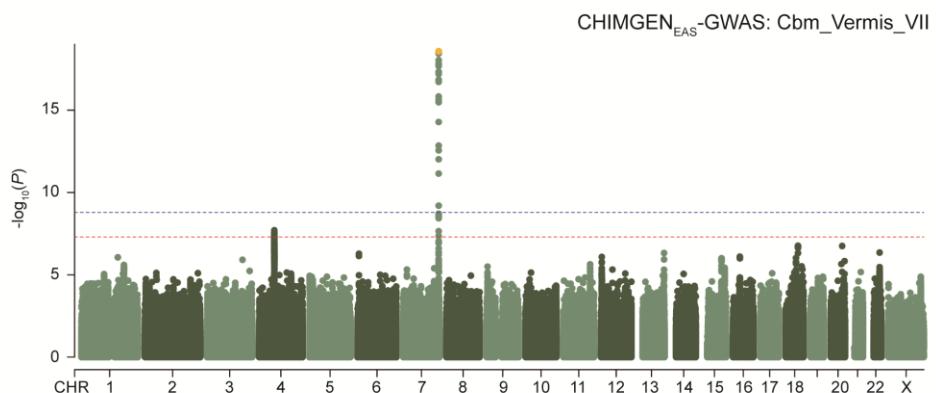
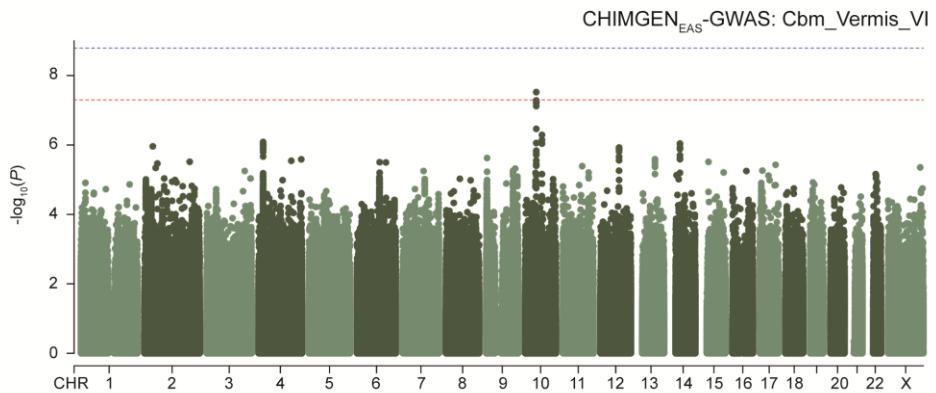


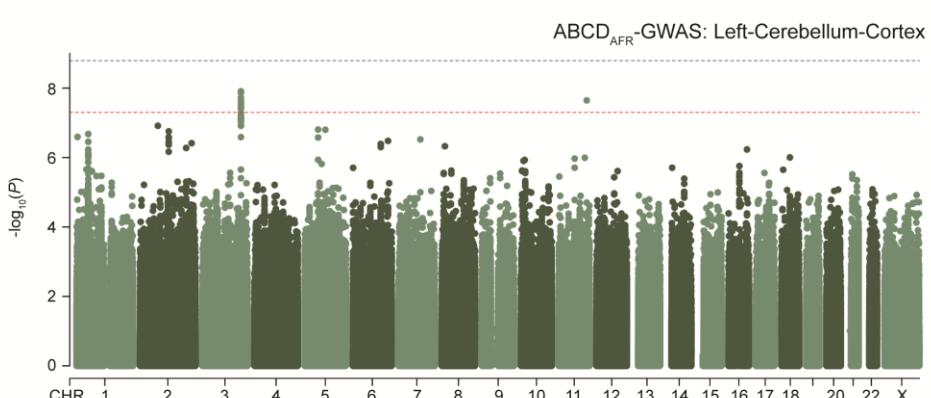
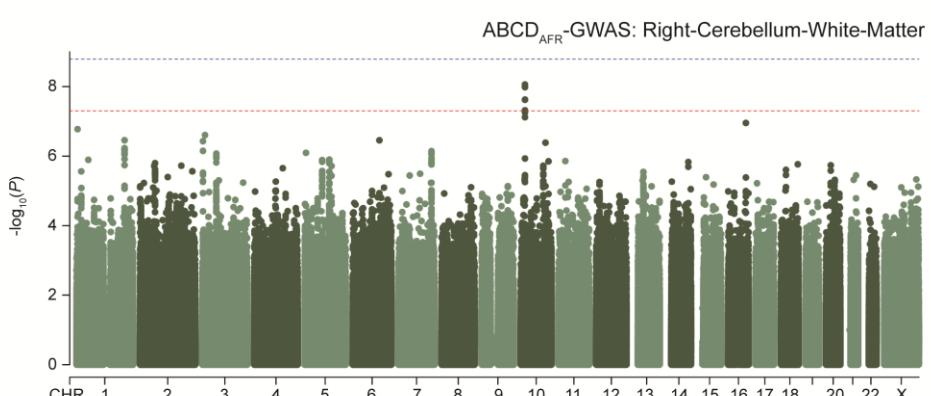
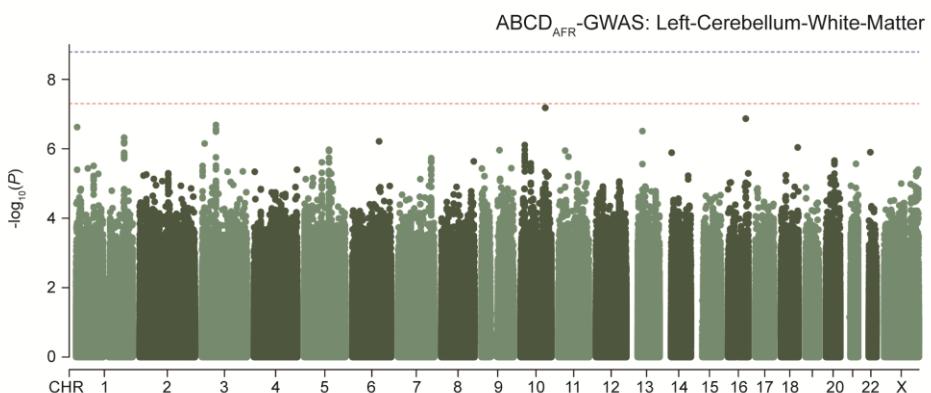
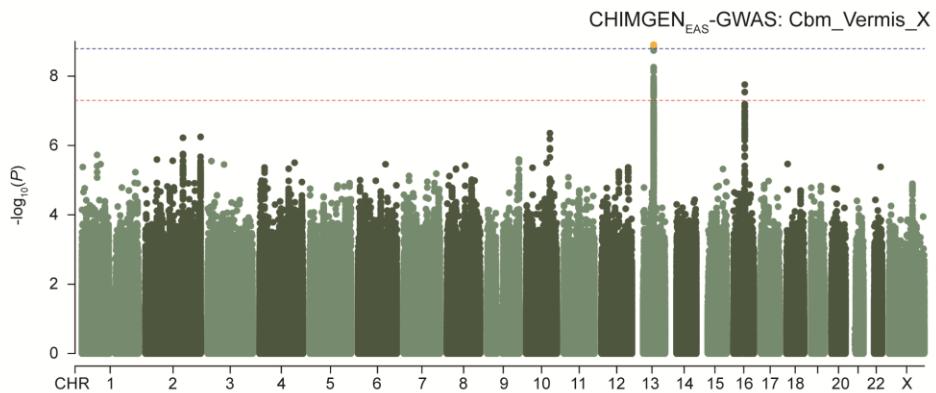


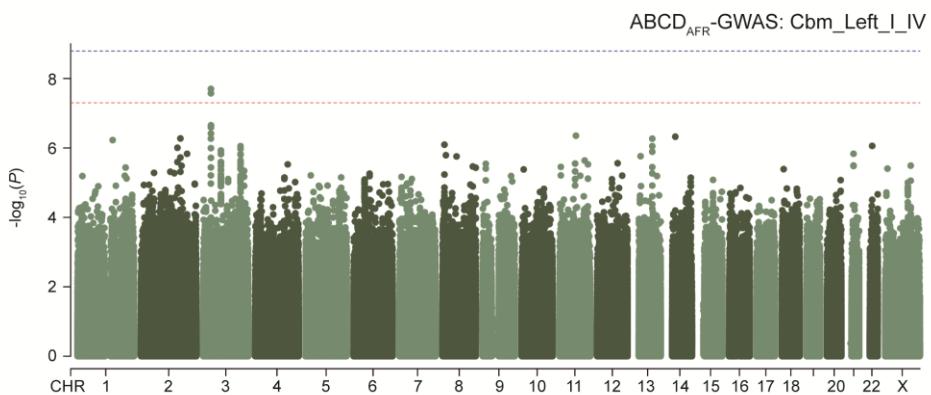
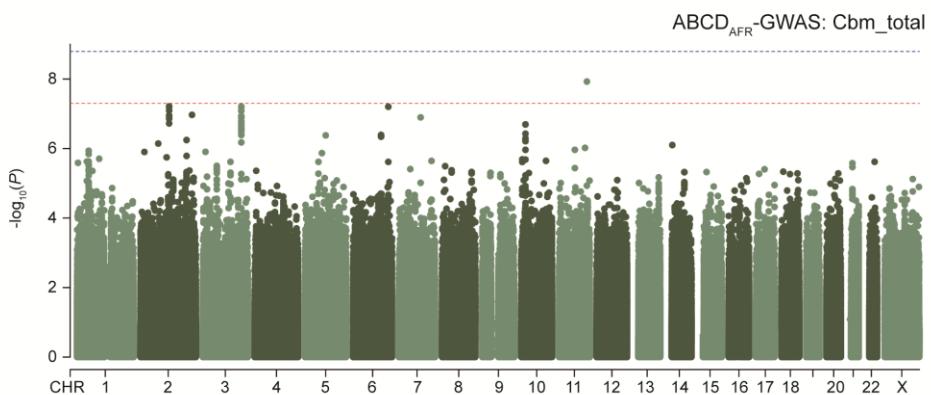
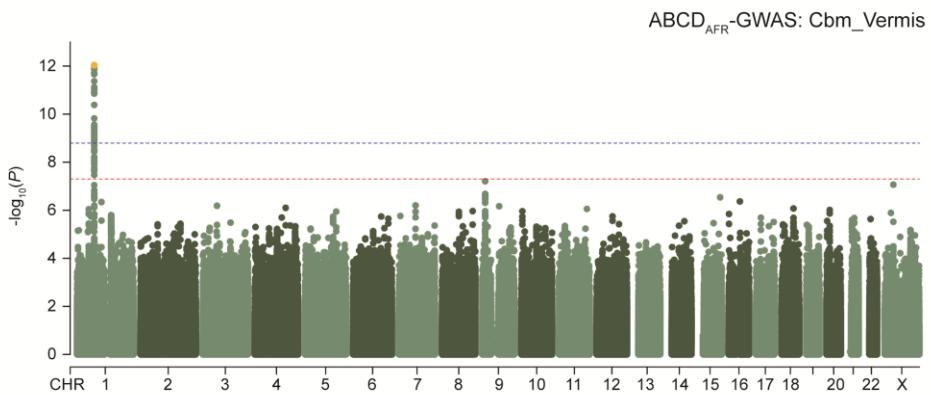
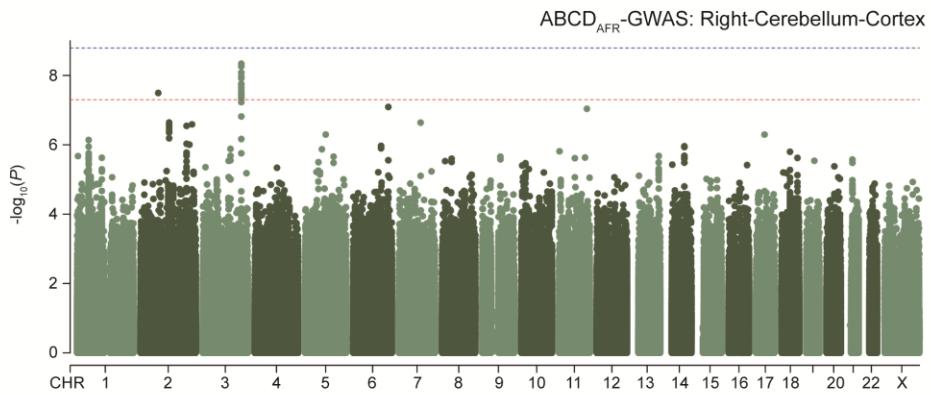


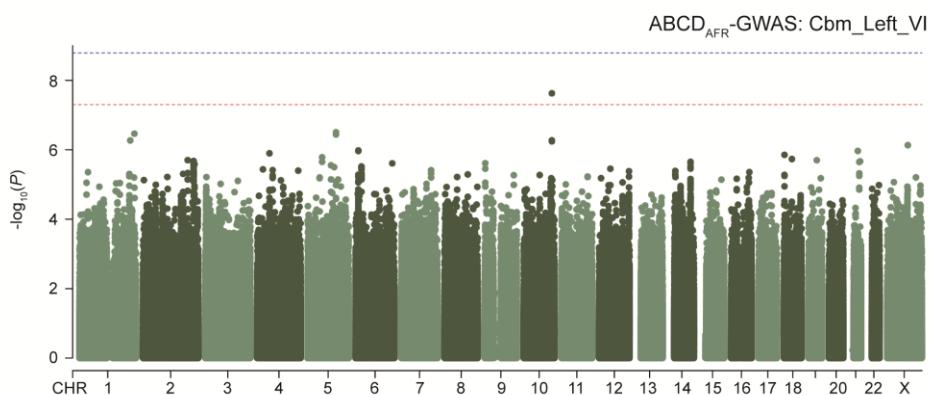
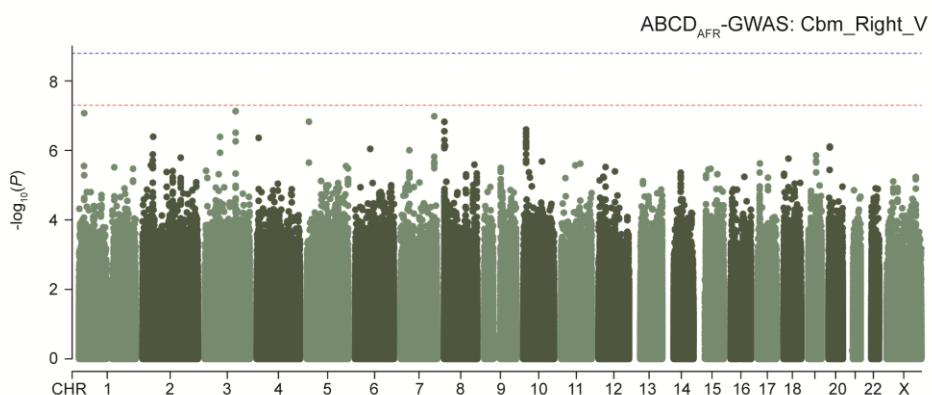
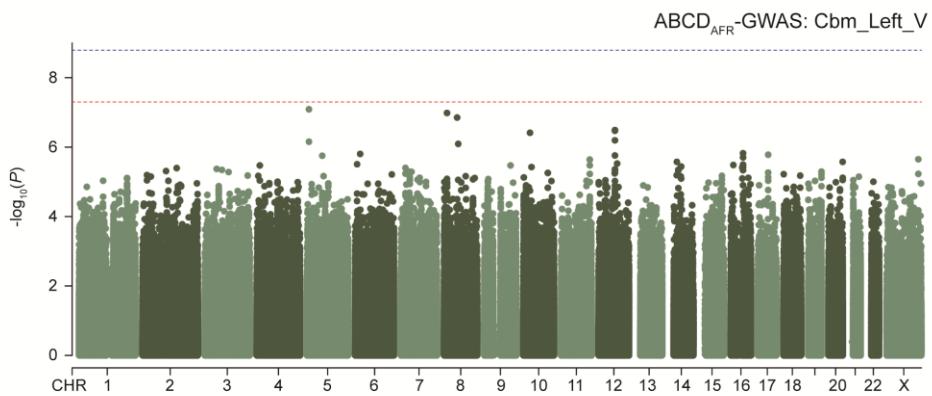
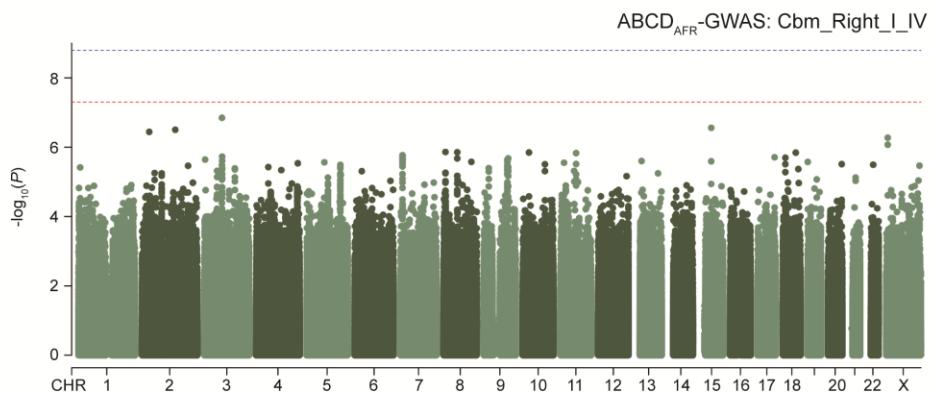


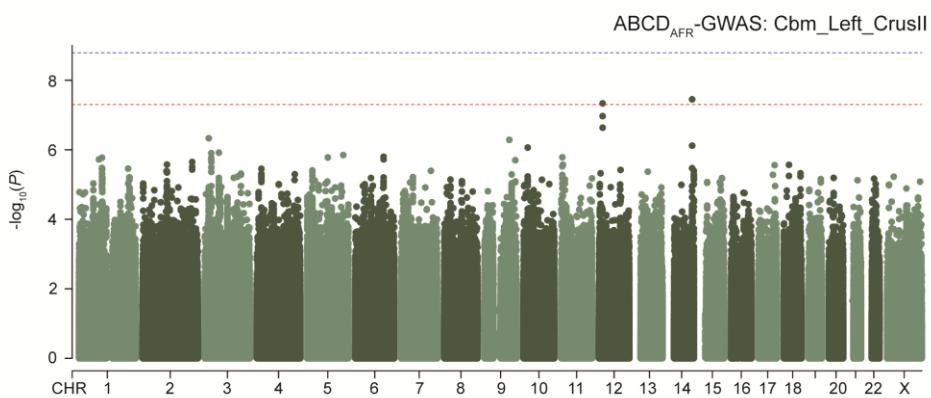
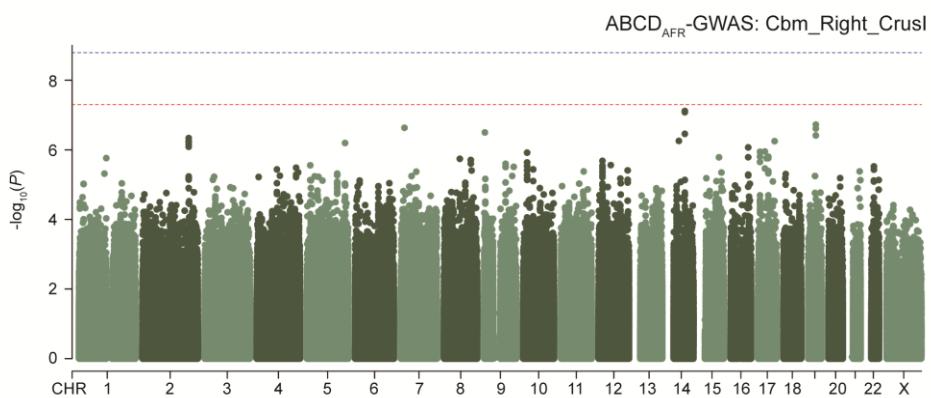
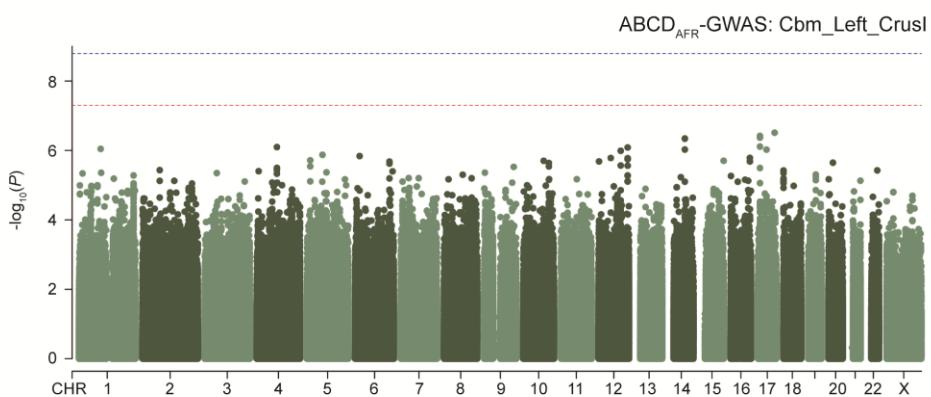
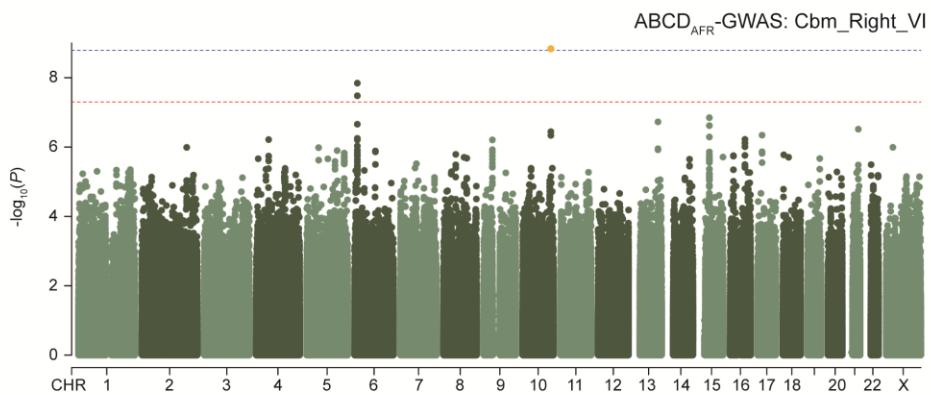


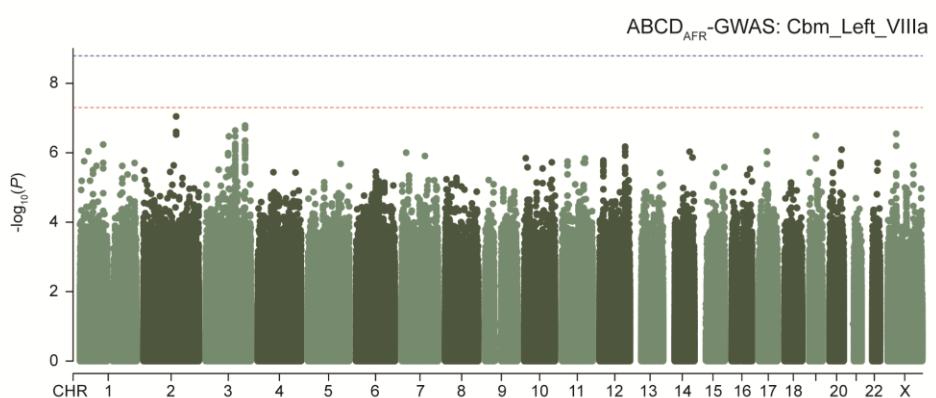
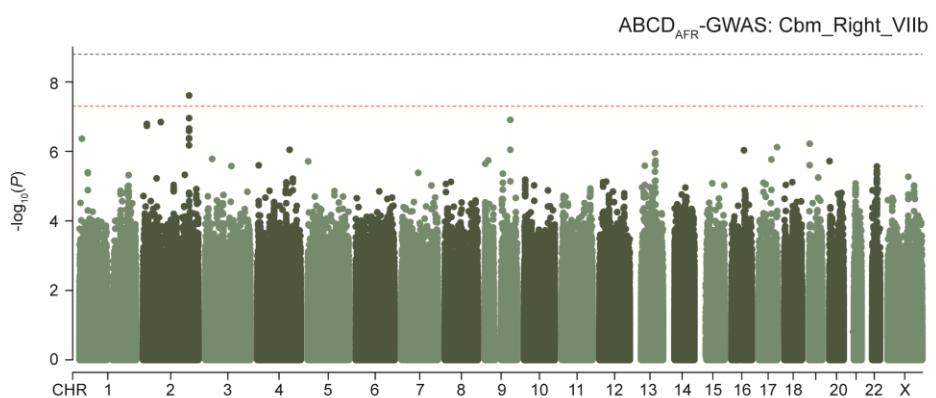
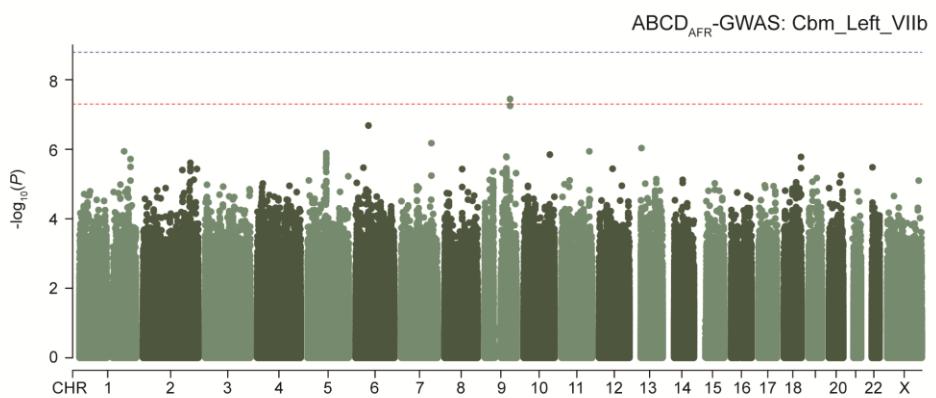
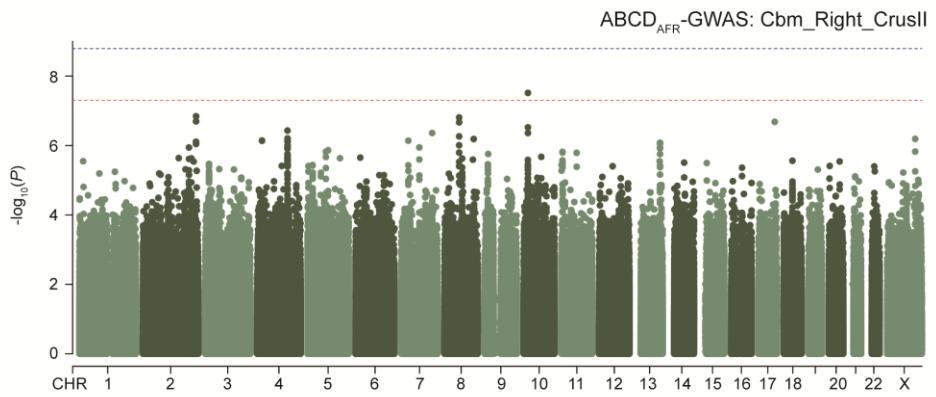


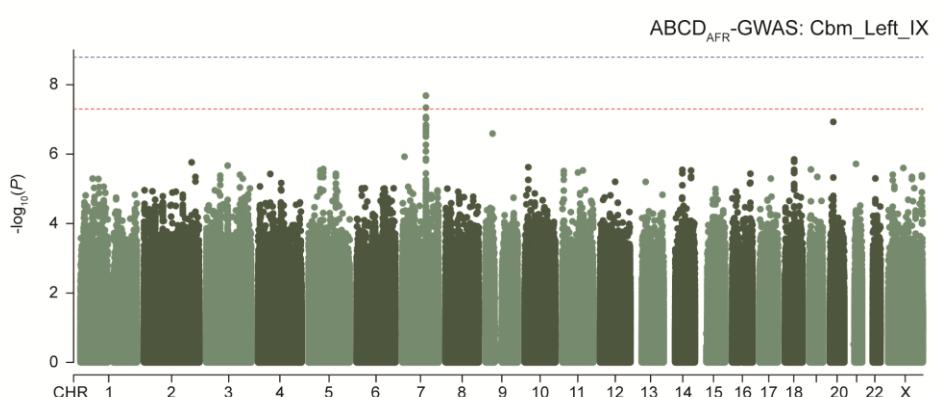
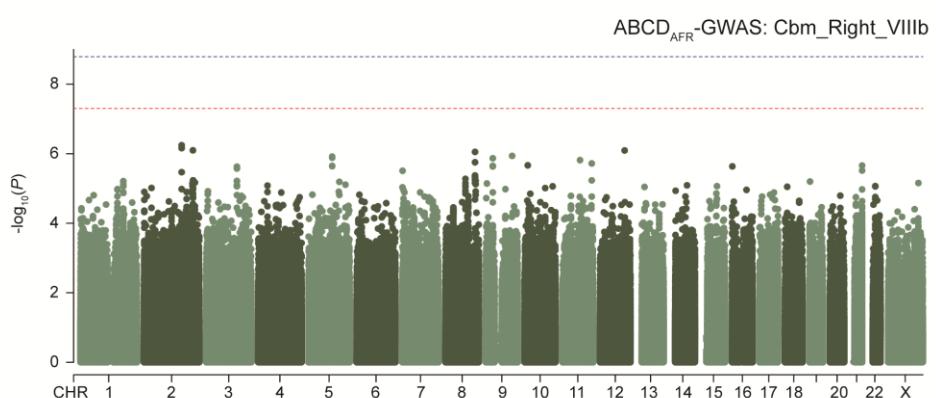
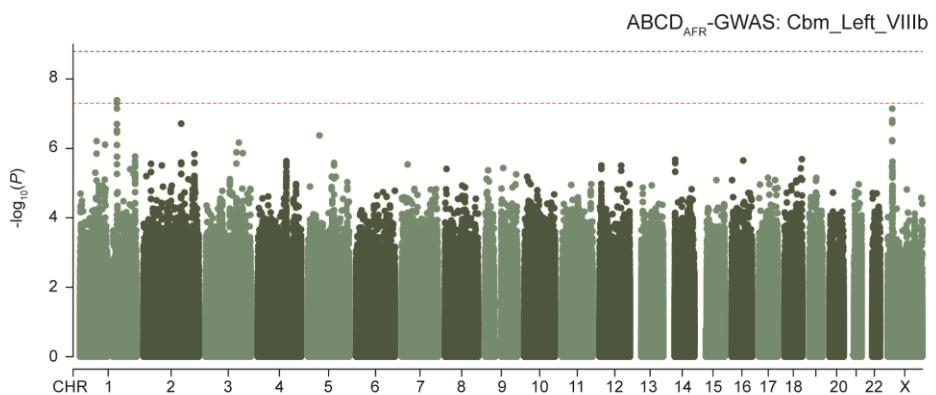
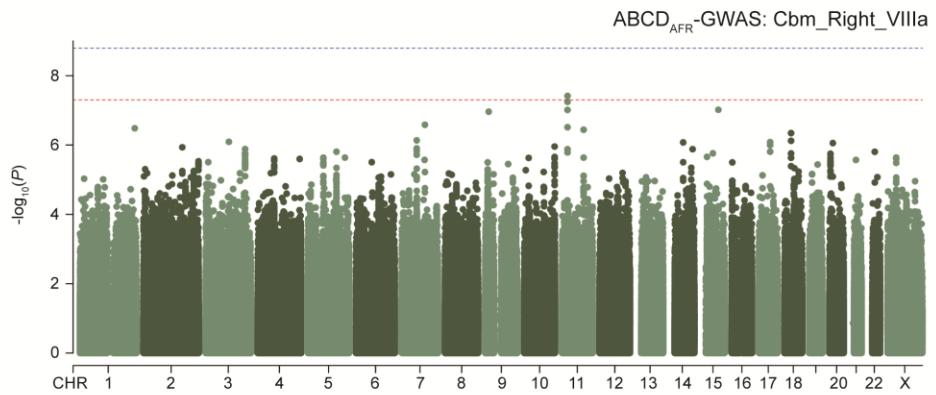


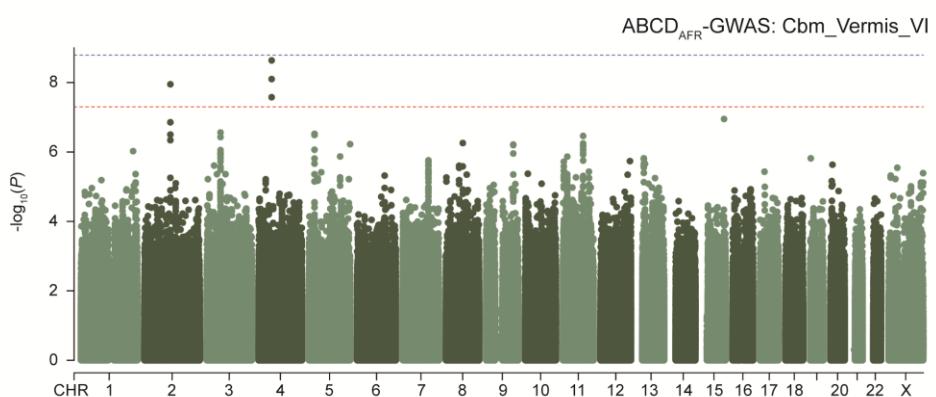
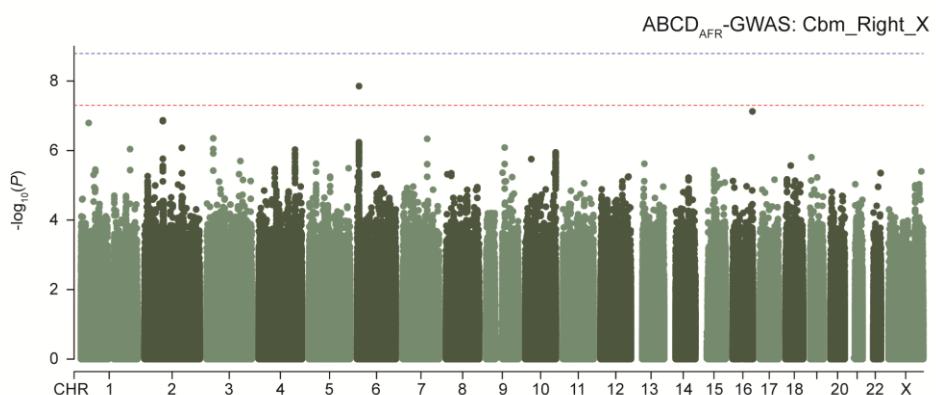
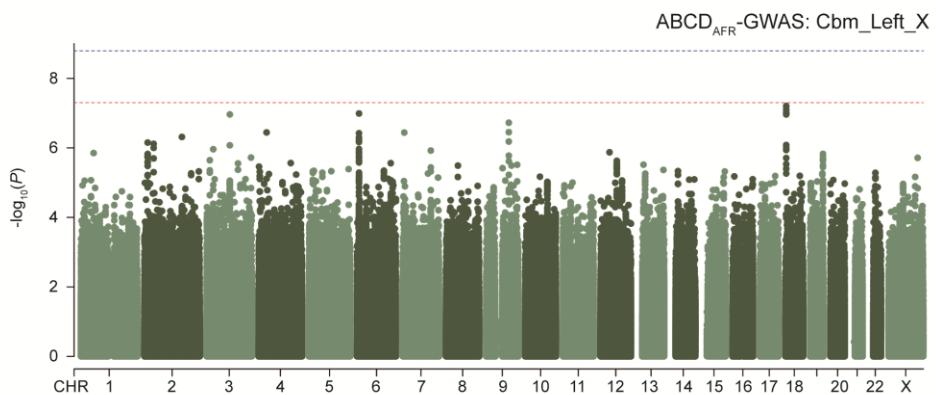
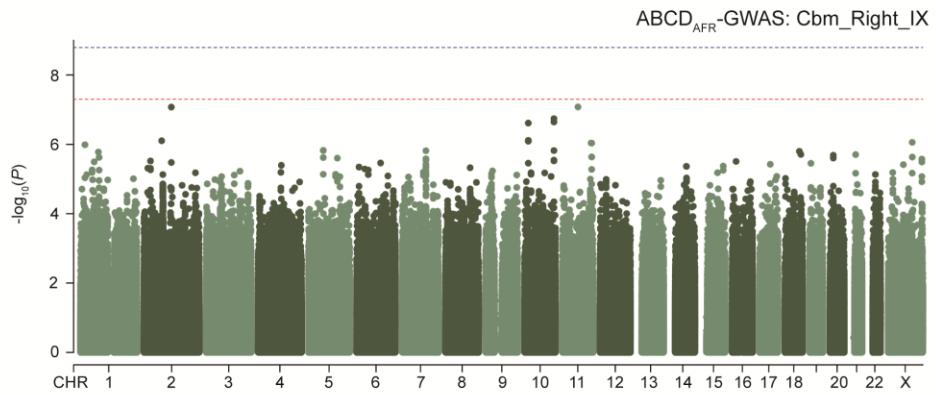


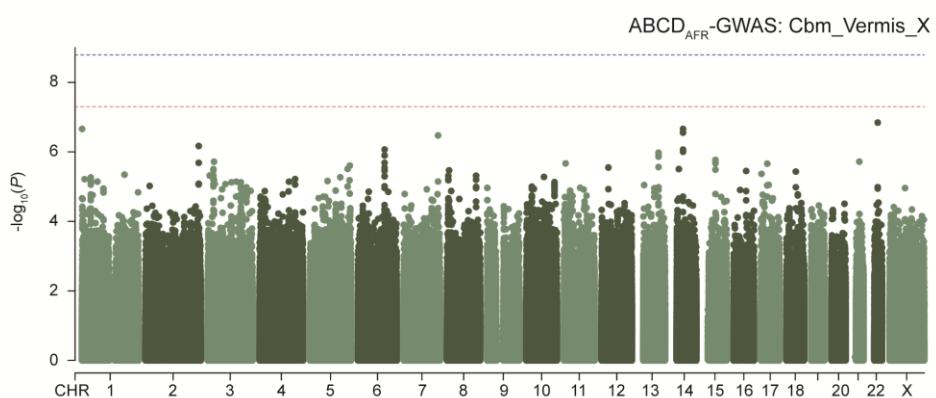
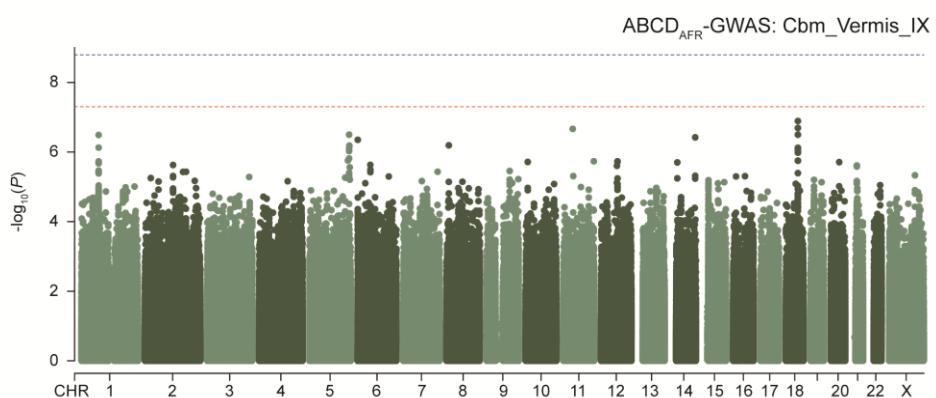
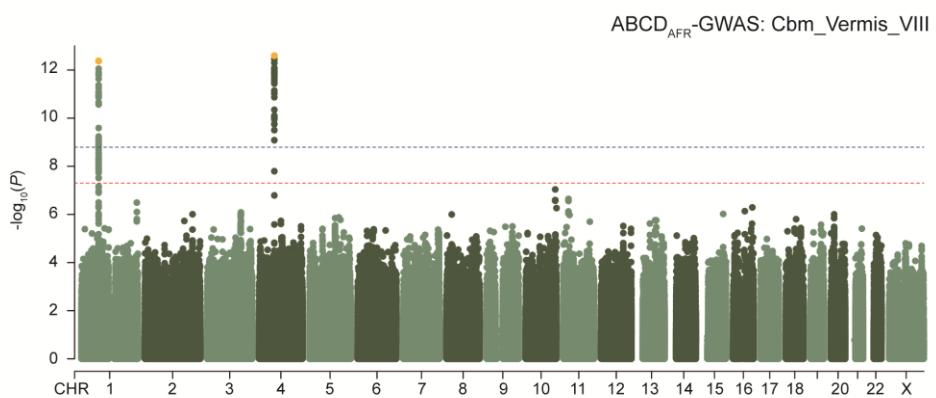
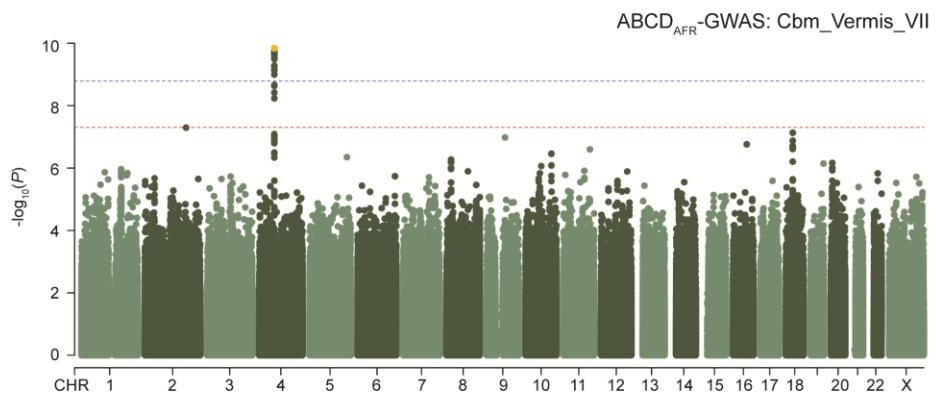


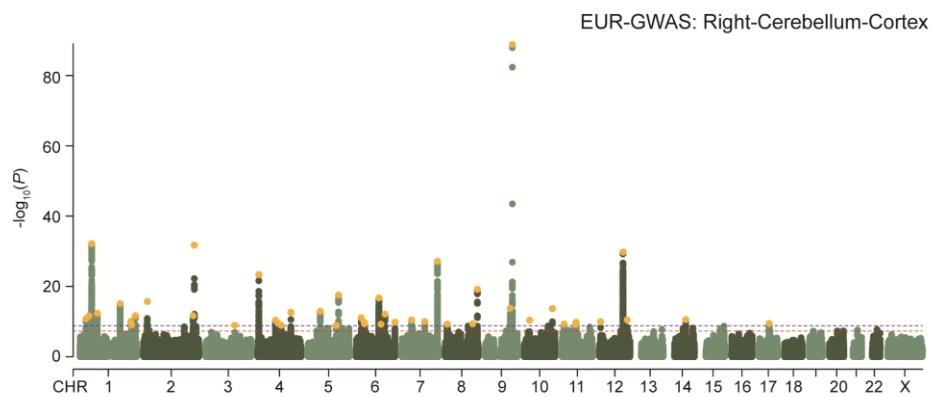
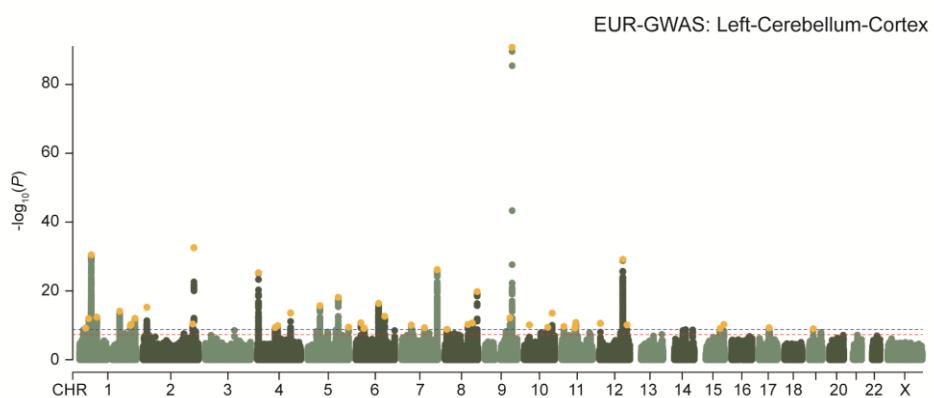
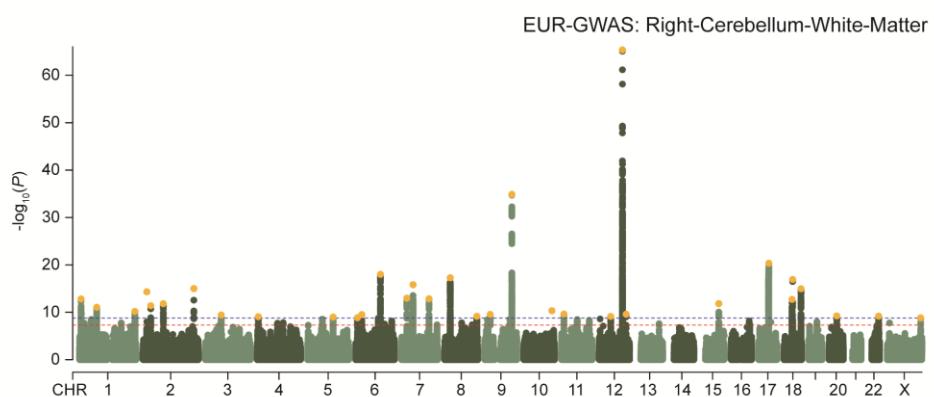
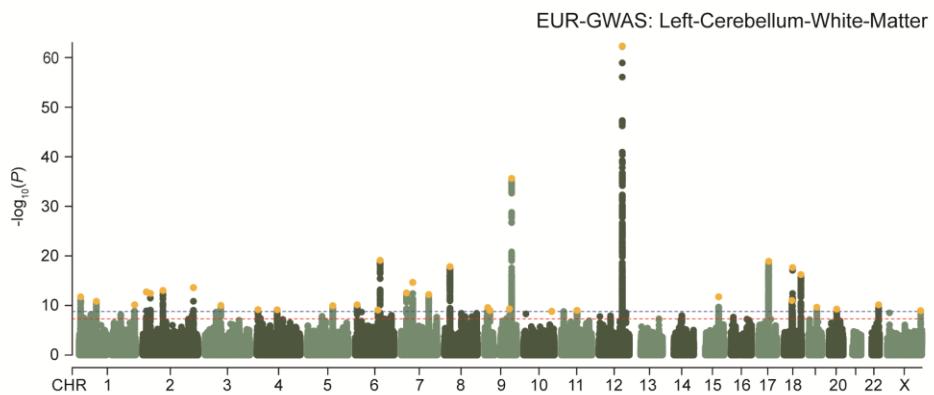




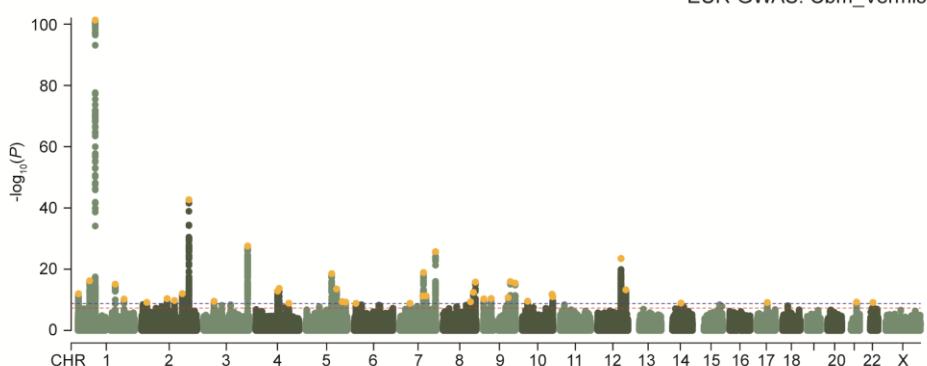




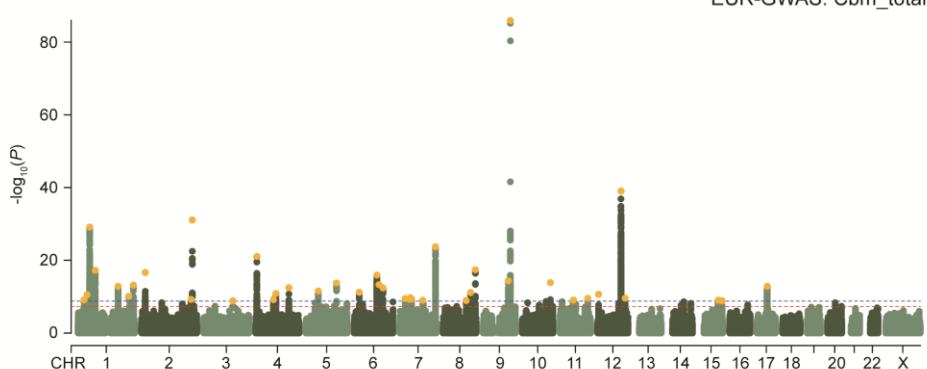




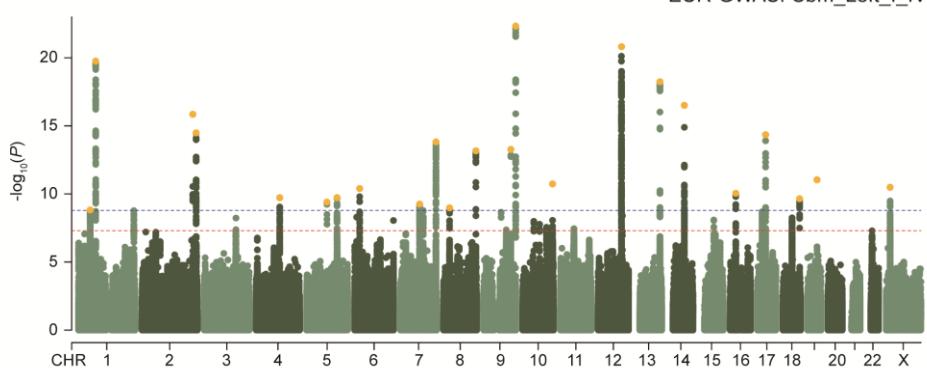
EUR-GWAS: Cbm_Vermis



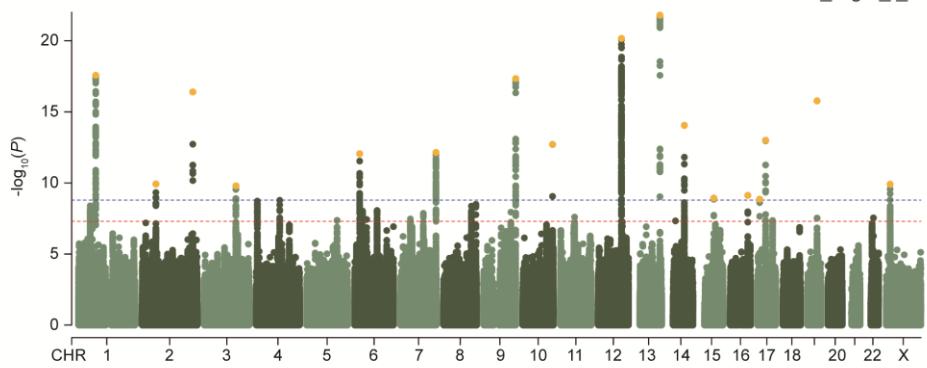
EUR-GWAS: Cbm_total

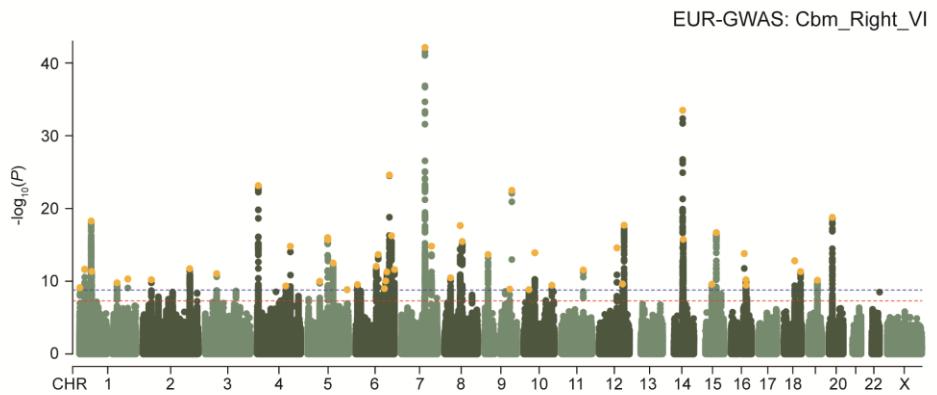
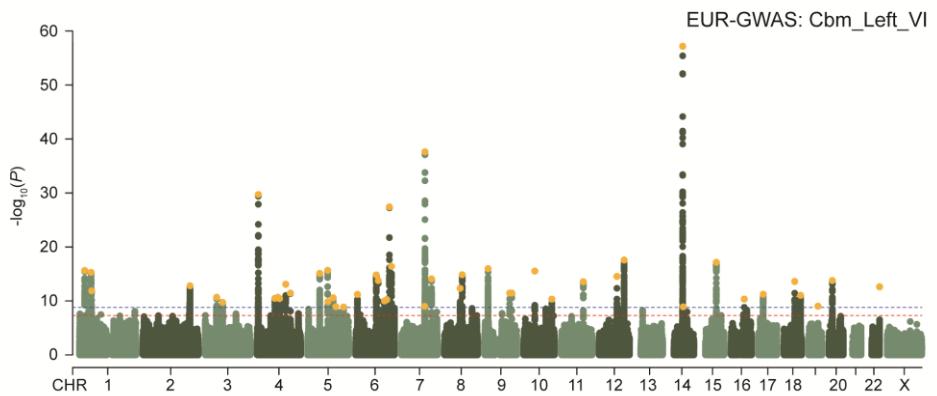
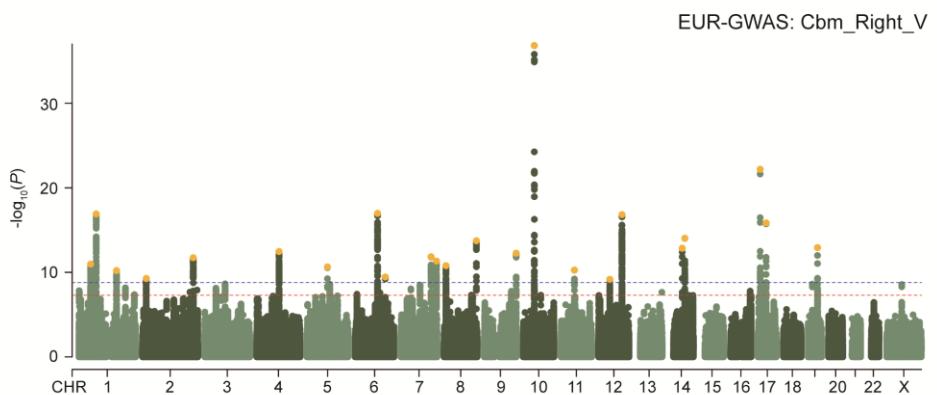
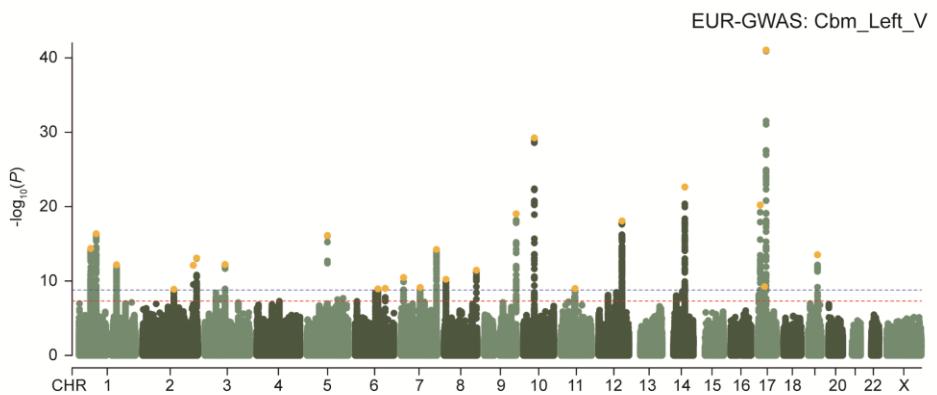


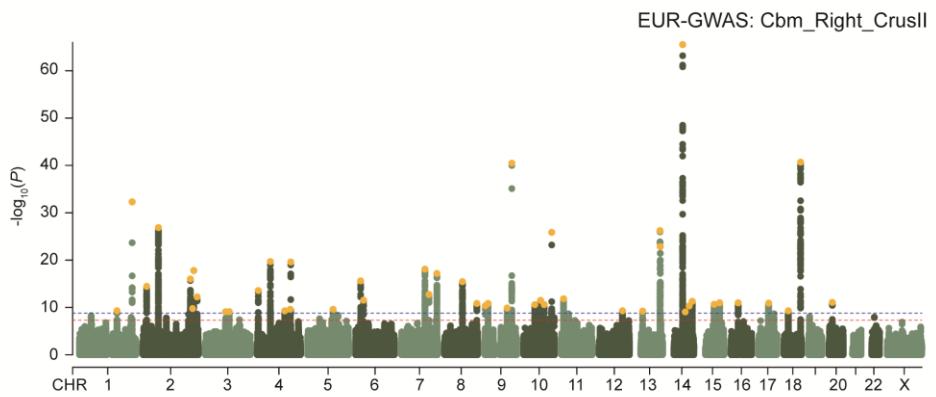
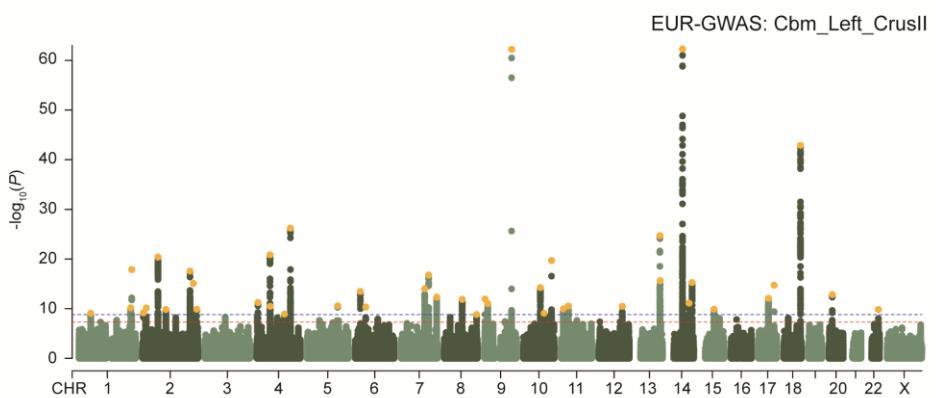
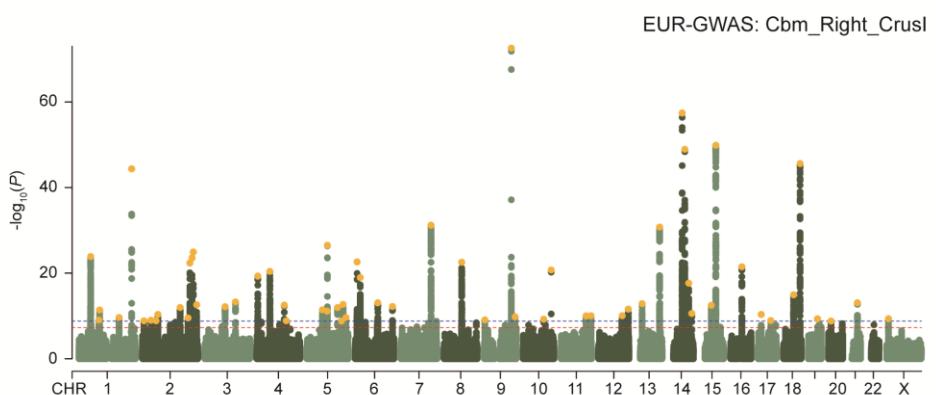
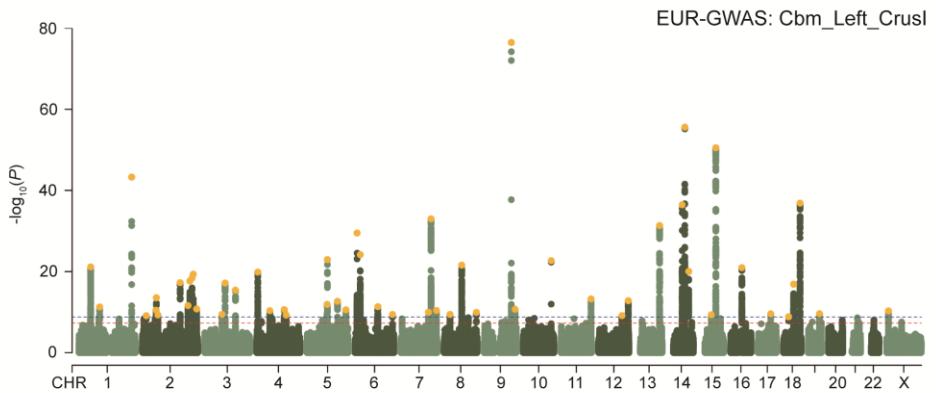
EUR-GWAS: Cbm_Left_I_IV

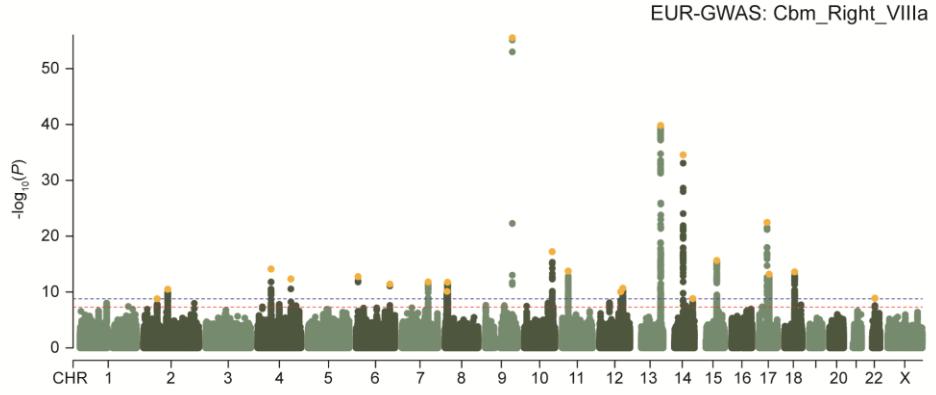
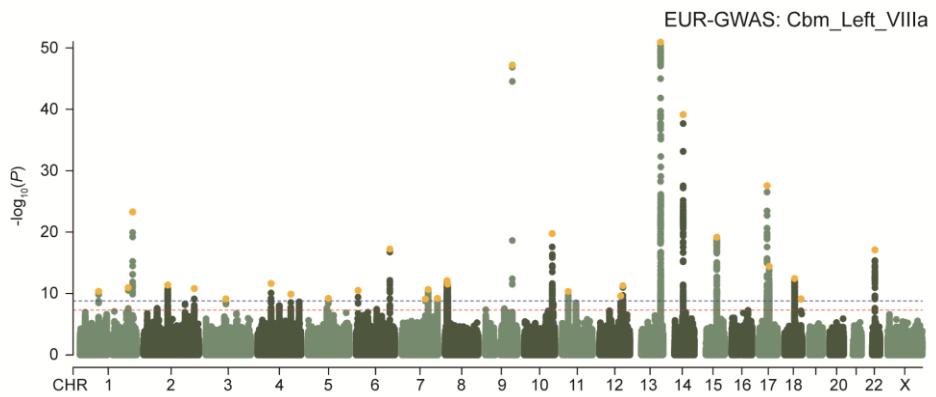
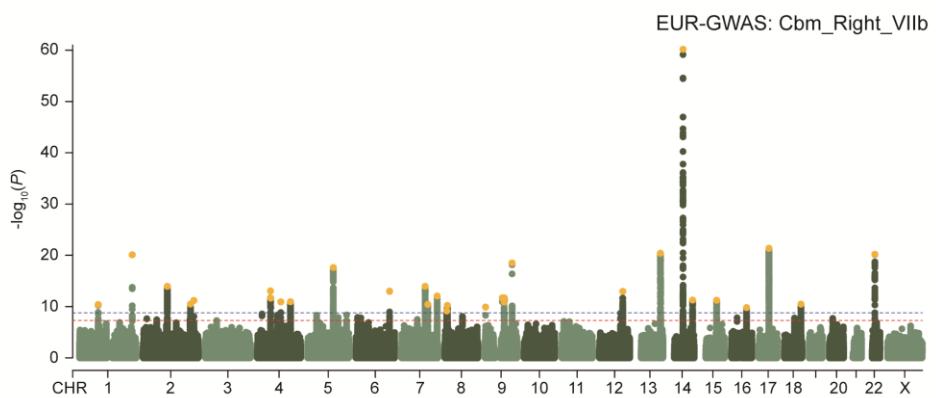
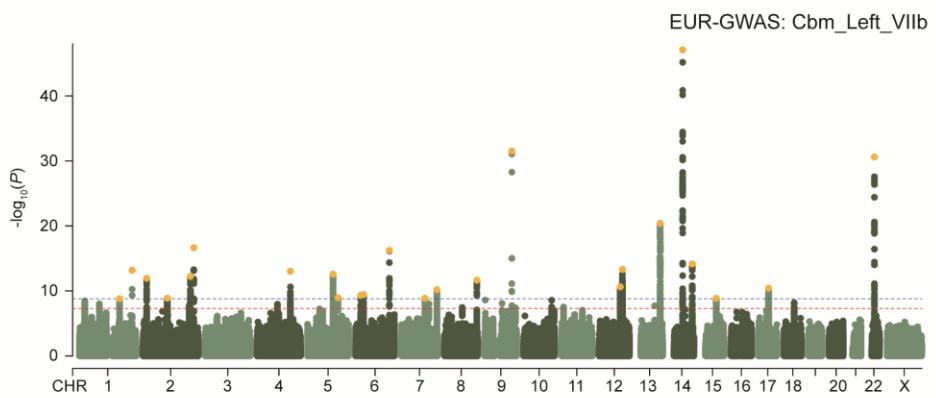


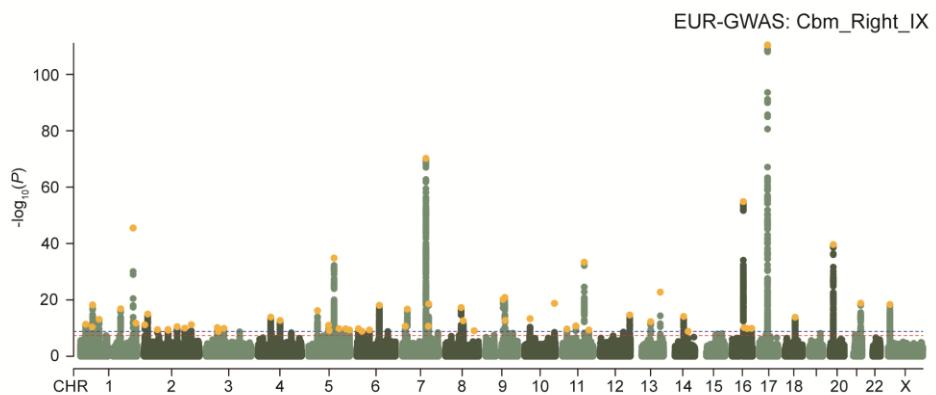
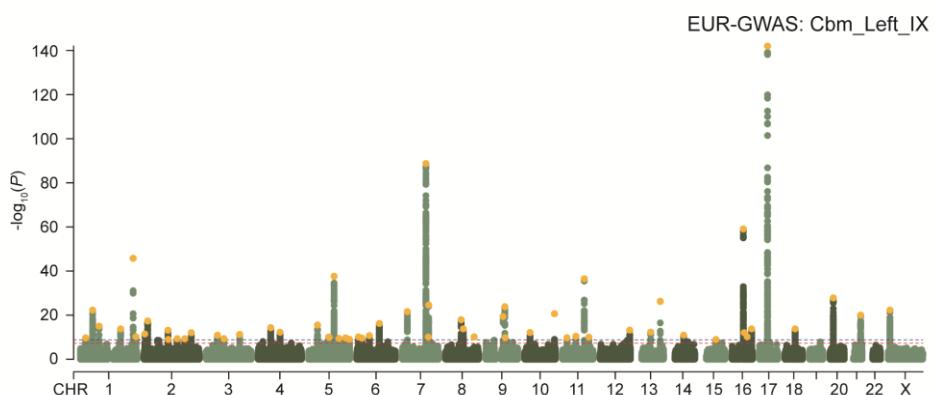
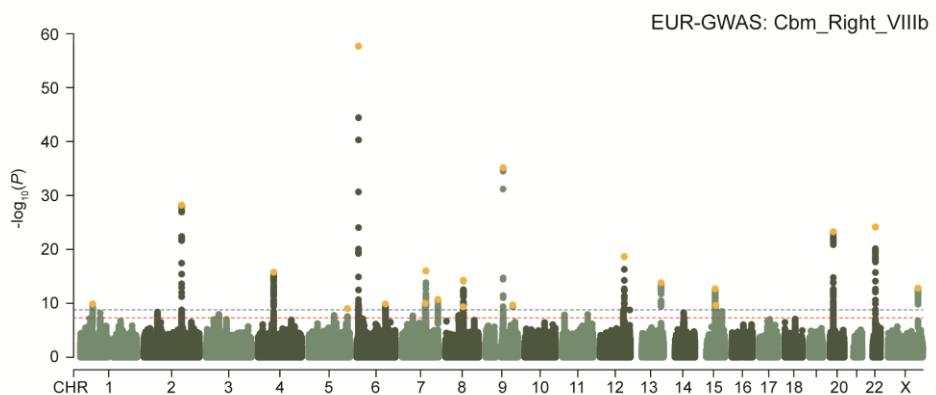
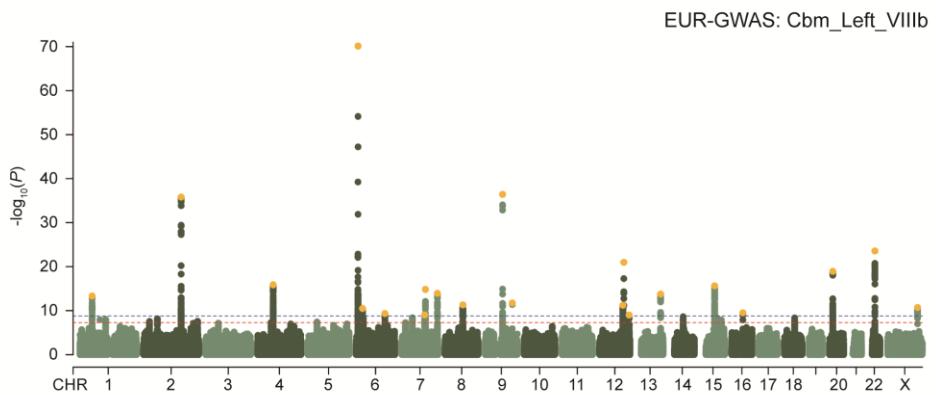
EUR-GWAS: Cbm_Right_I_IV

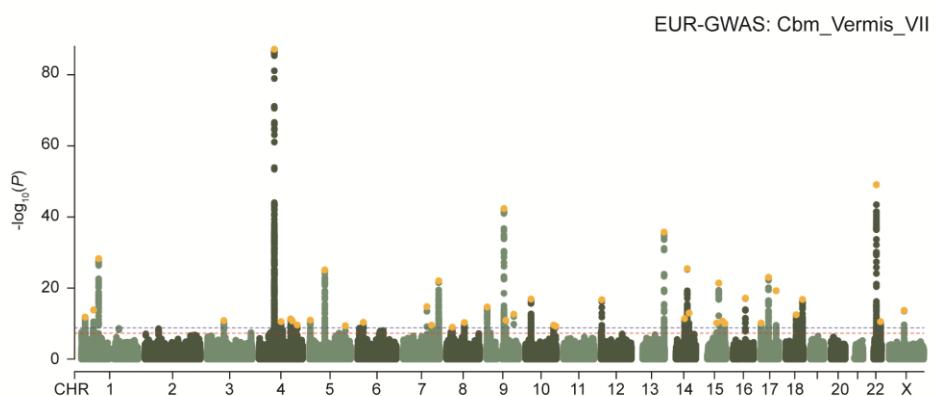
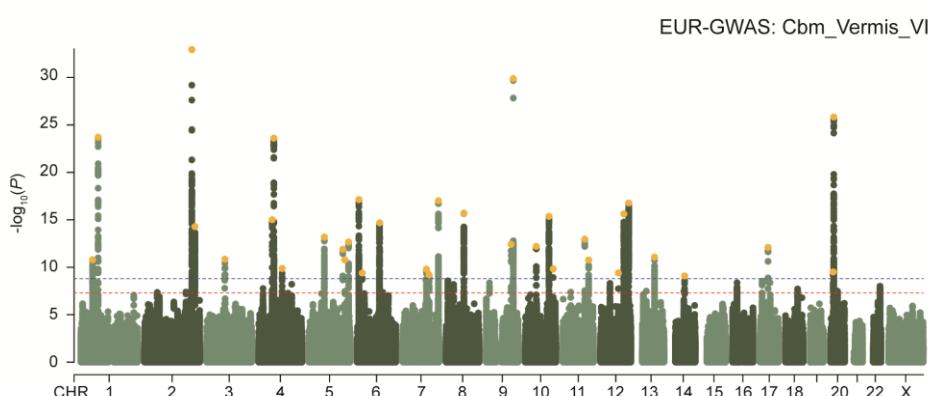
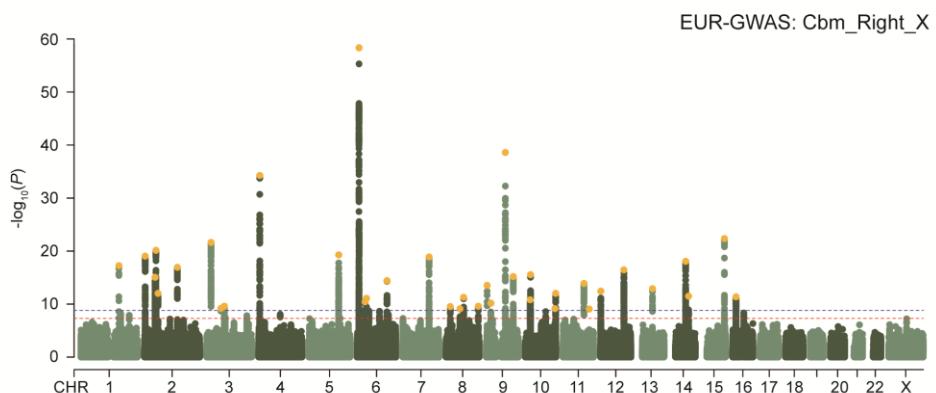
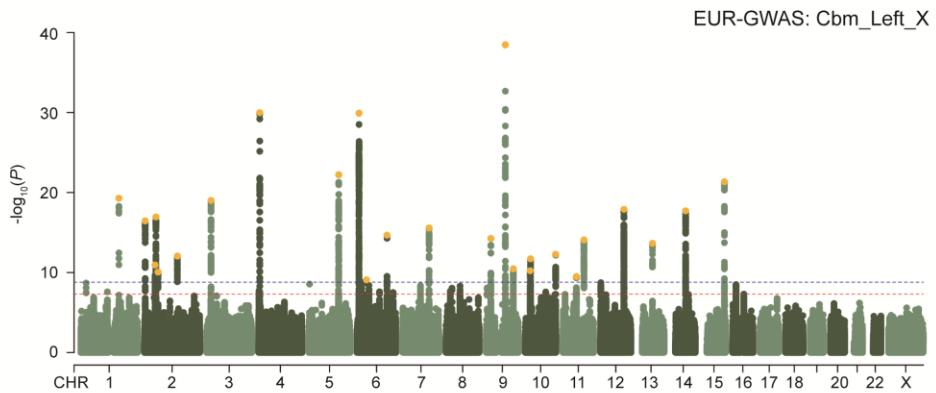




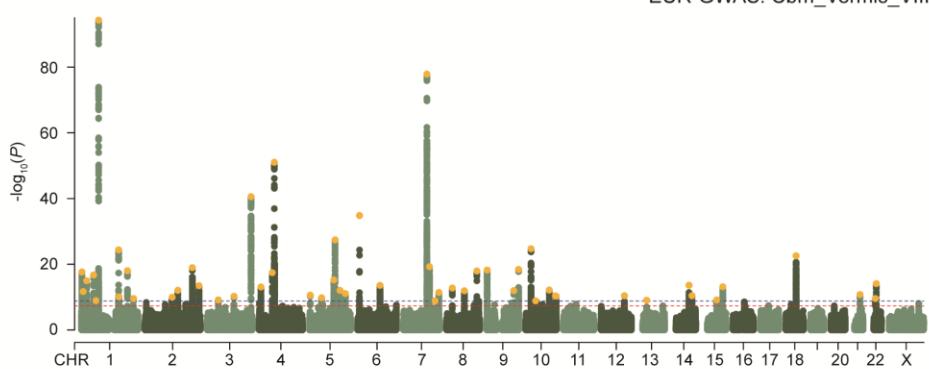




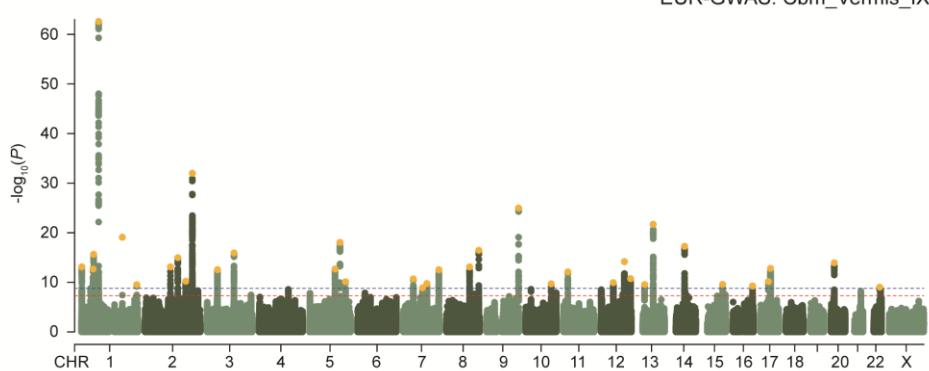




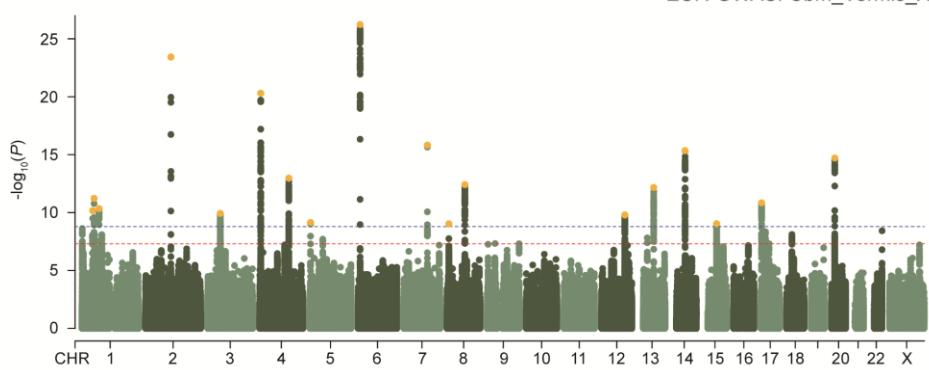
EUR-GWAS: Cbm_Vermis_VIII



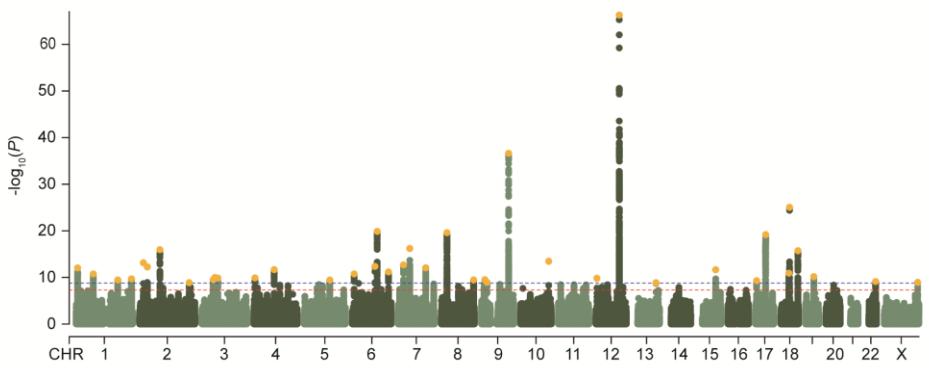
EUR-GWAS: Cbm_Vermis_IX

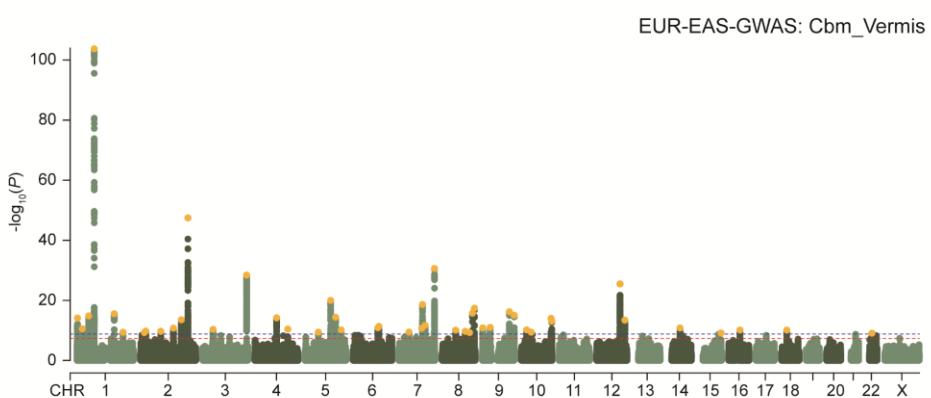
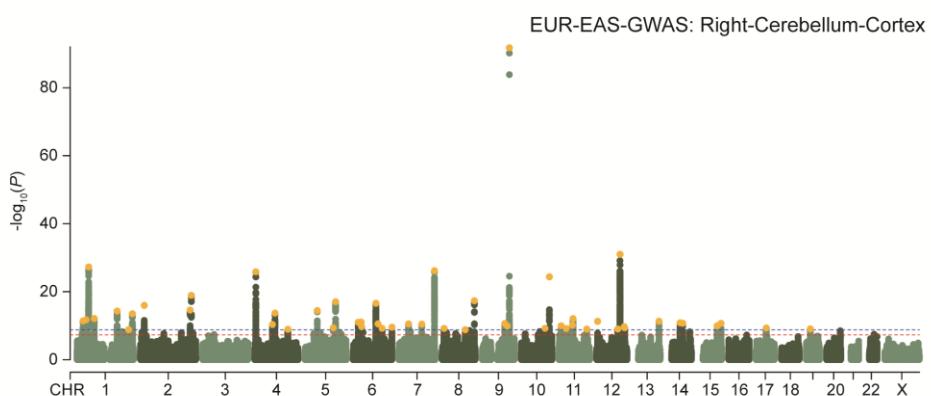
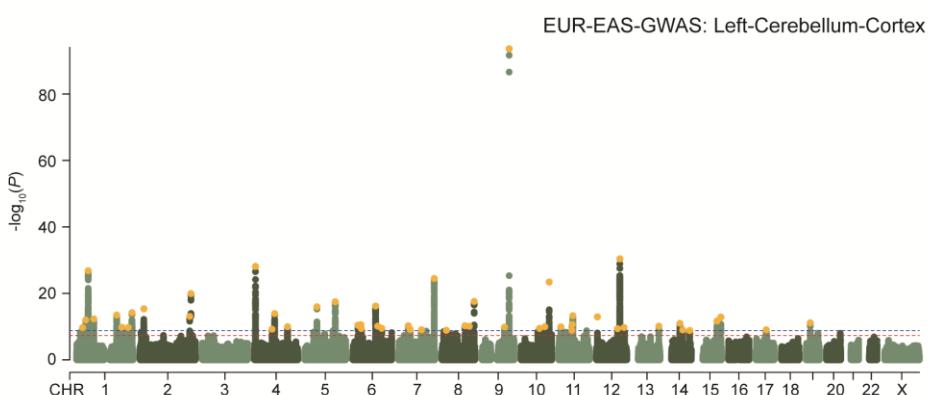
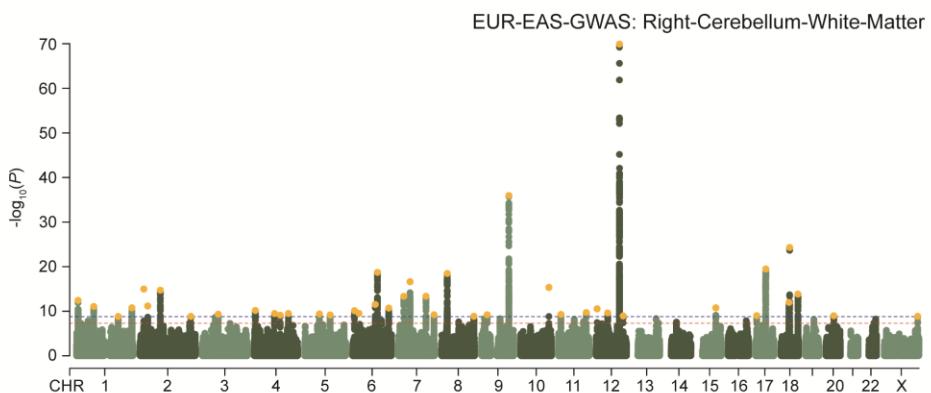


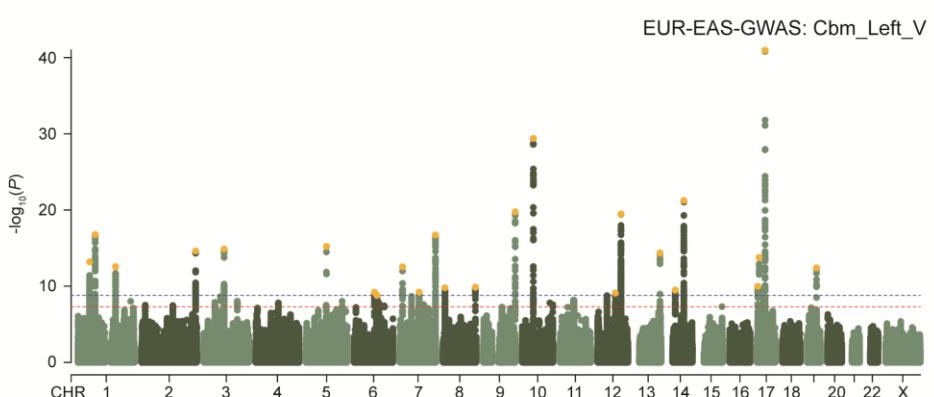
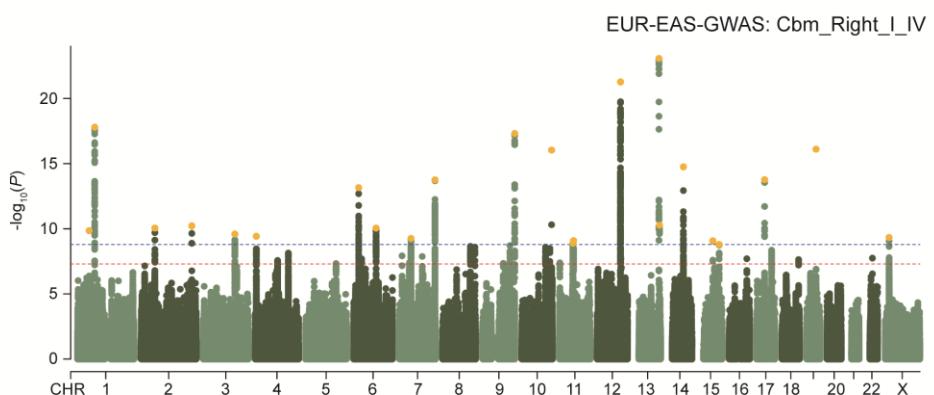
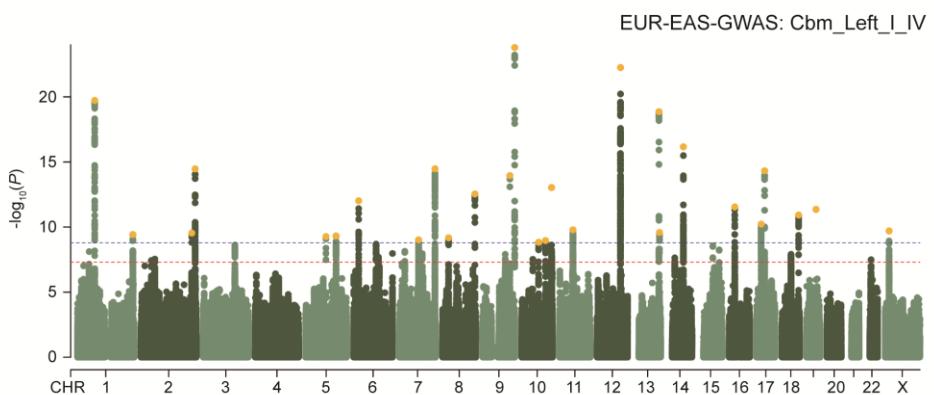
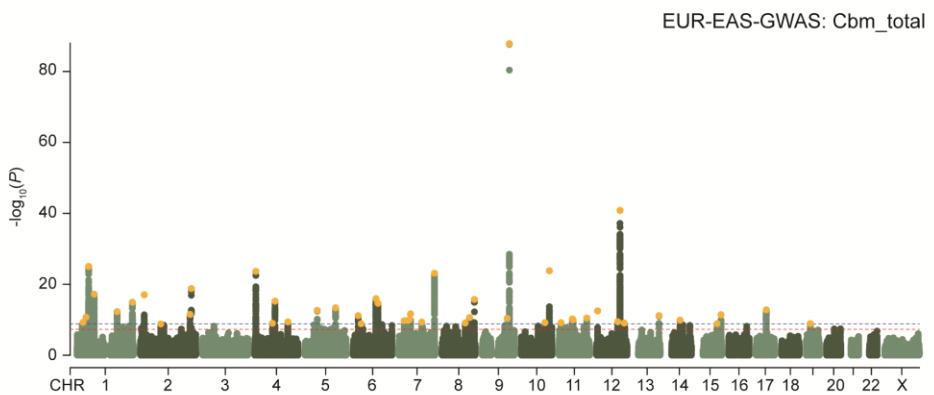
EUR-GWAS: Cbm_Vermis_X

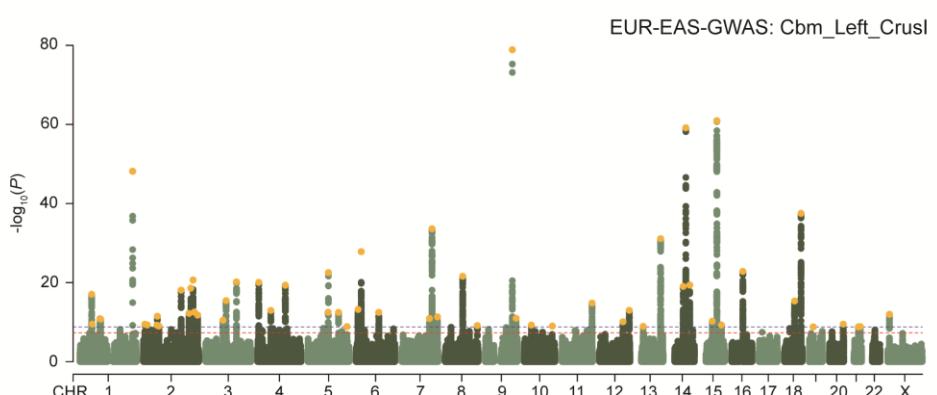
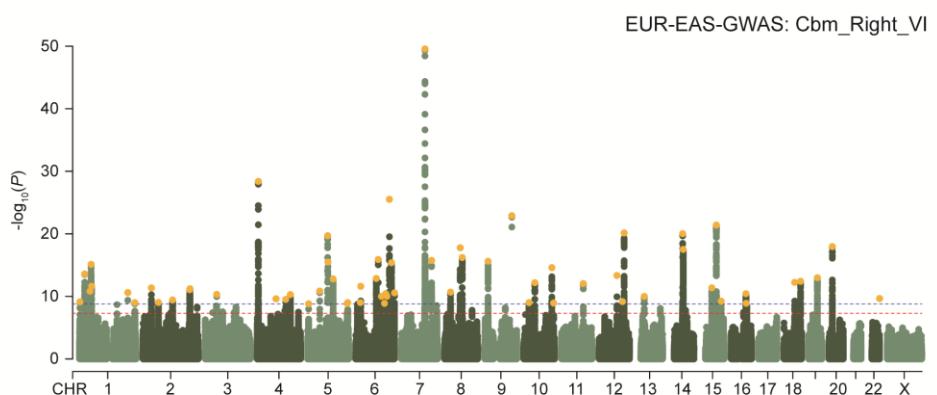
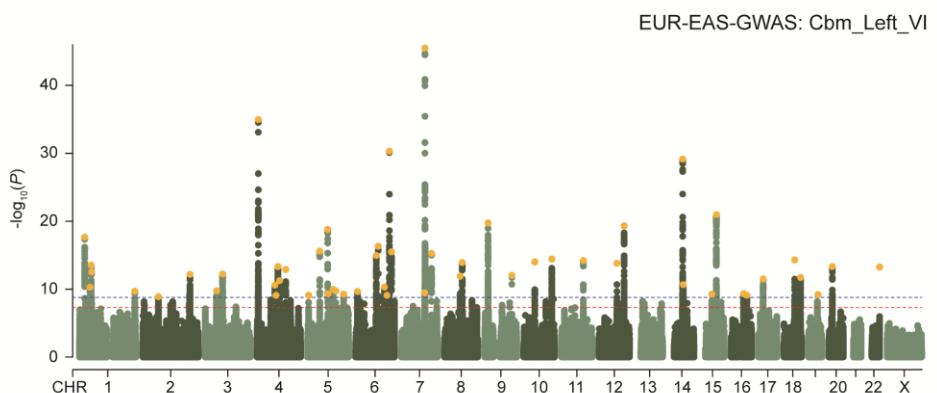
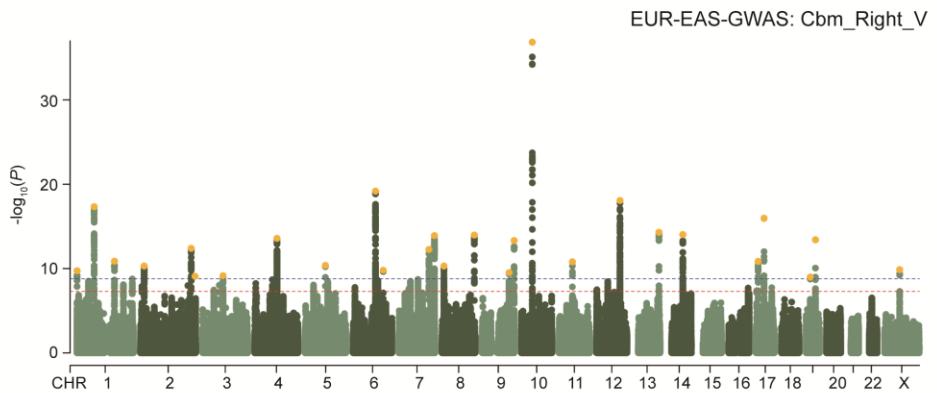


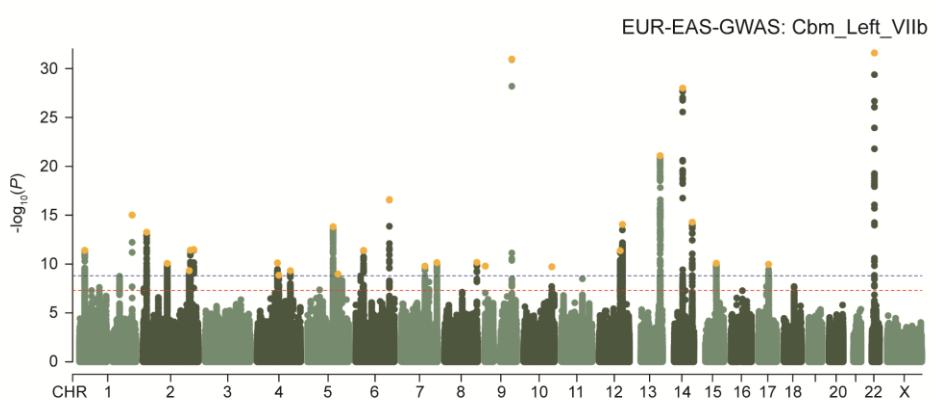
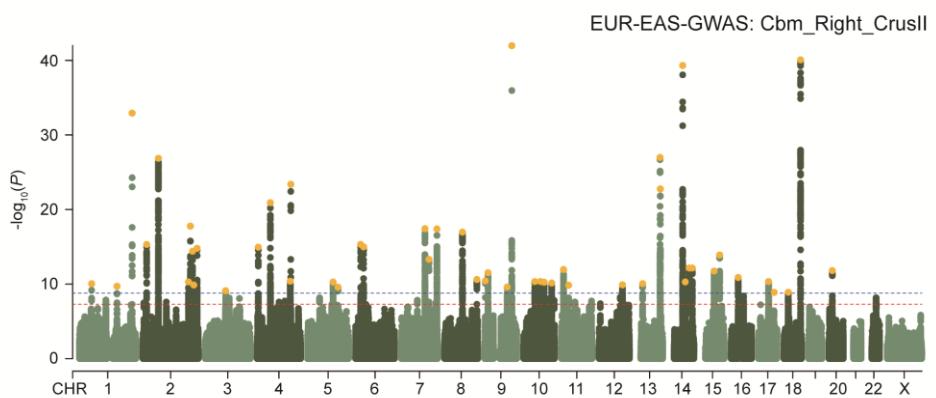
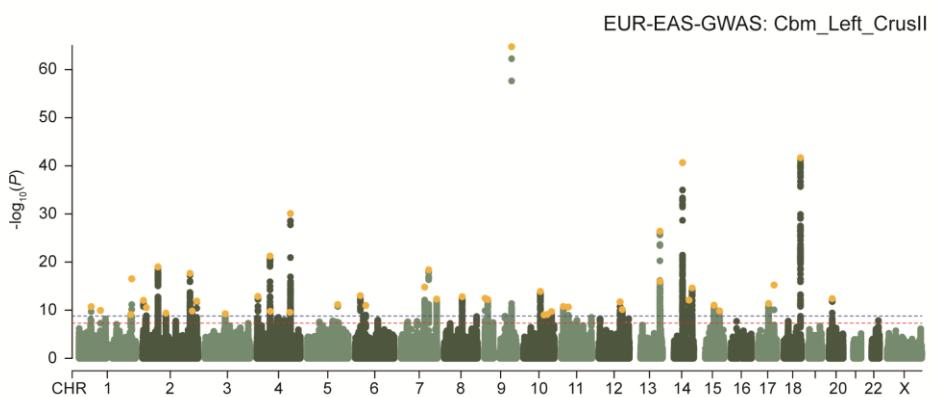
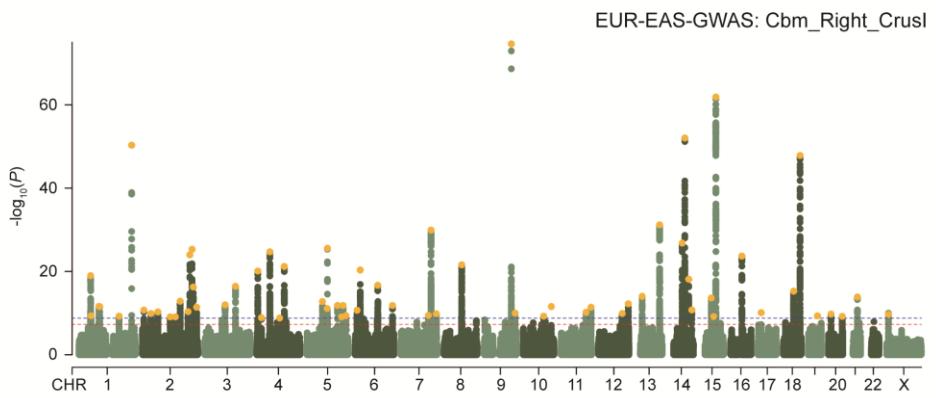
EUR-EAS-GWAS: Left-Cerebellum-White-Matter

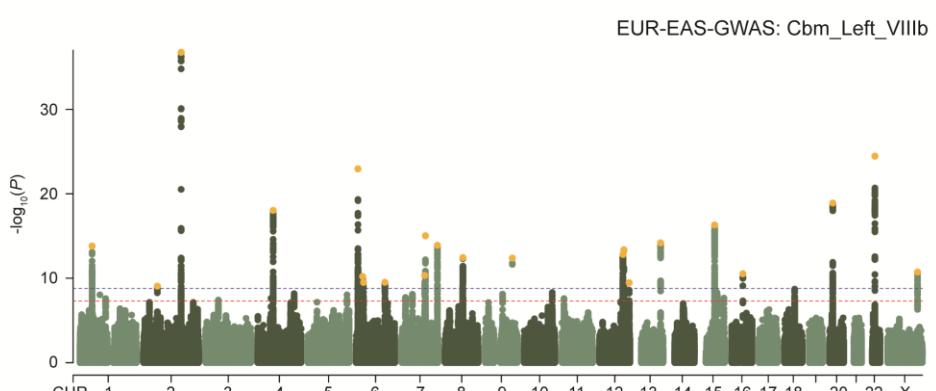
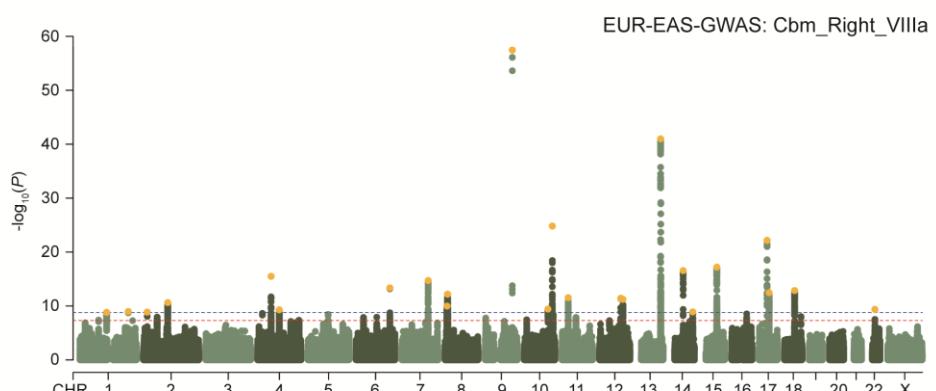
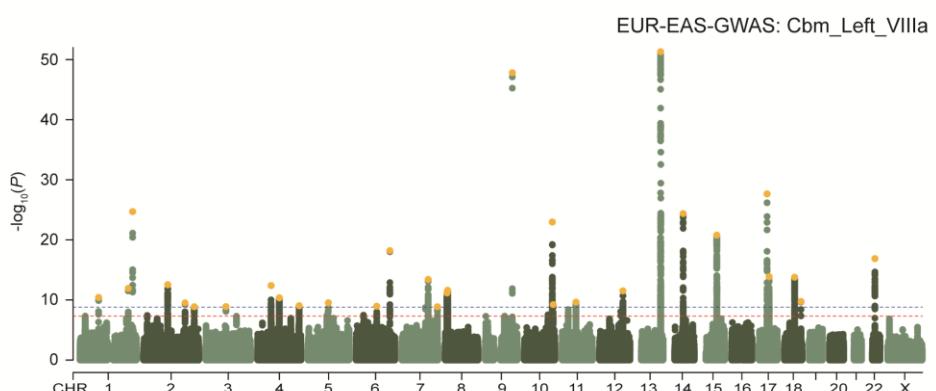
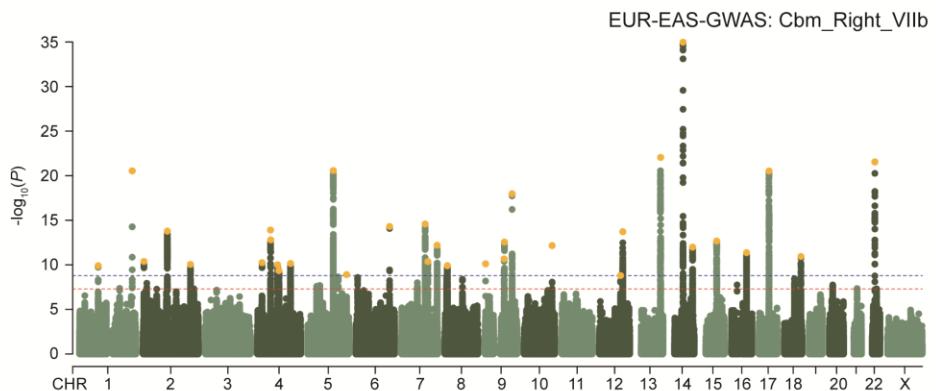


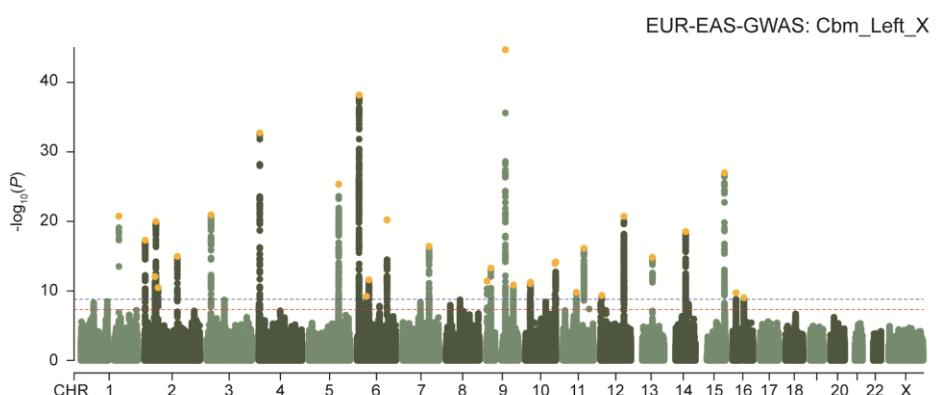
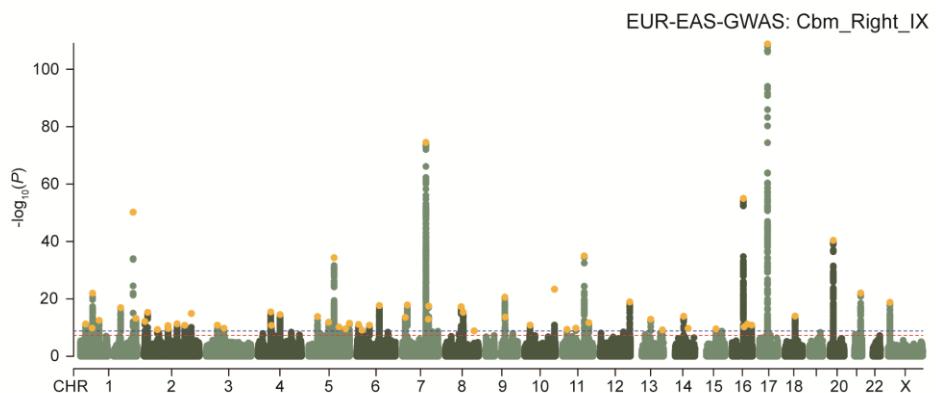
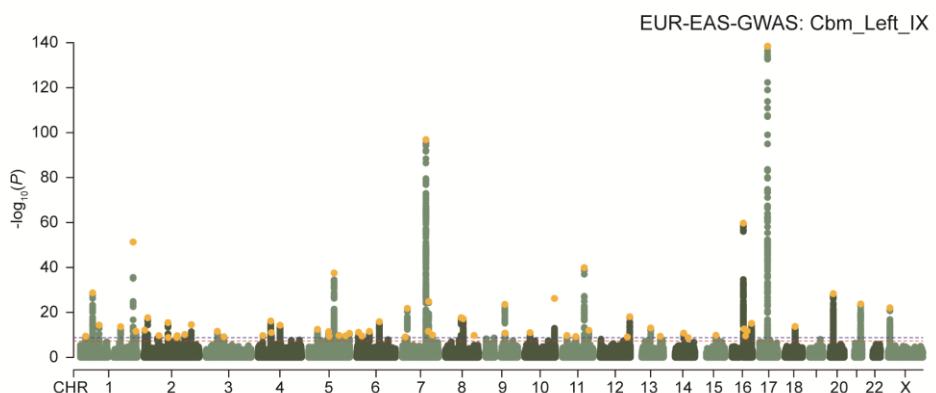
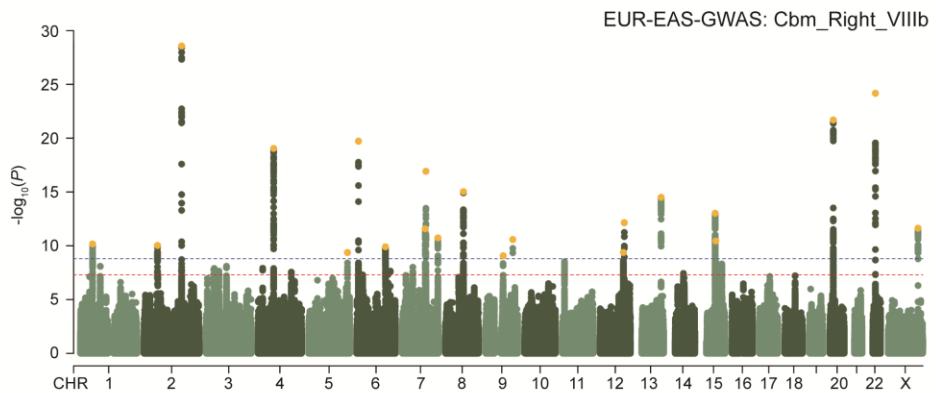




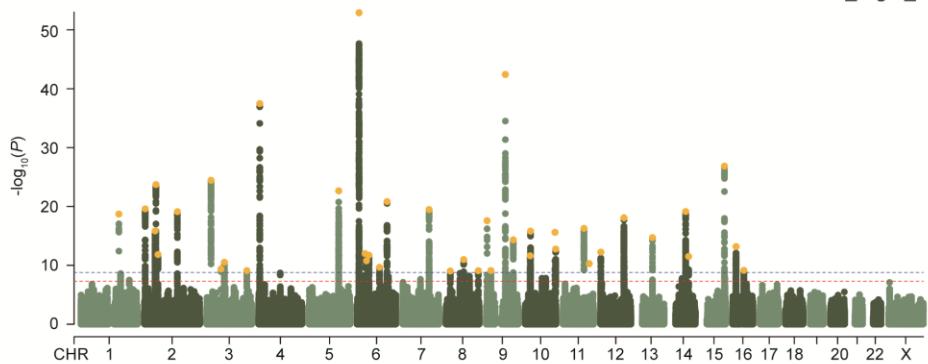




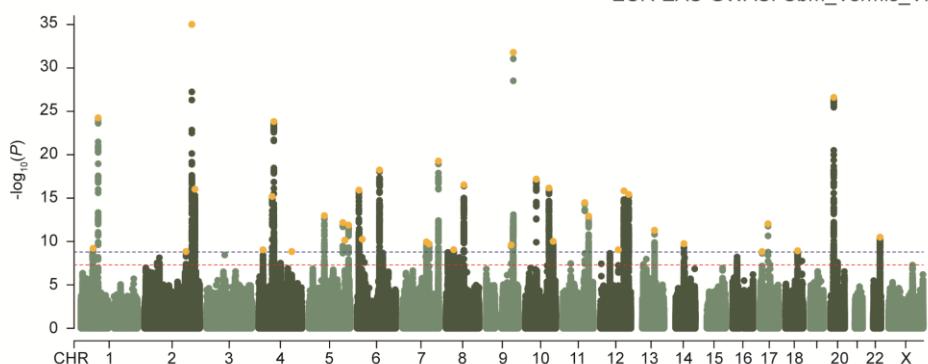




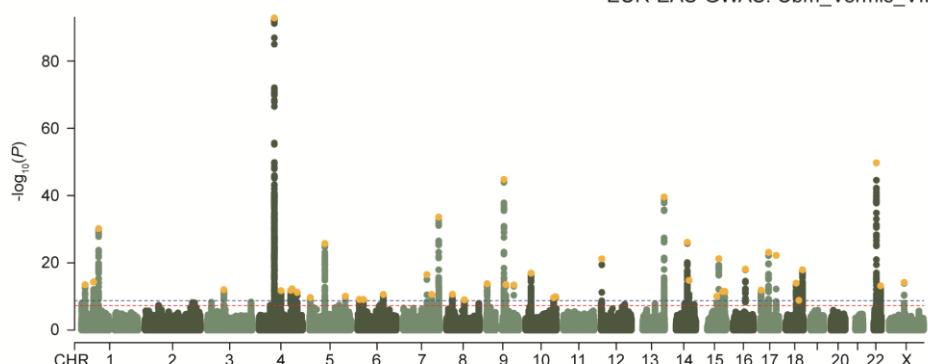
EUR-EAS-GWAS: Cbm_Right_X



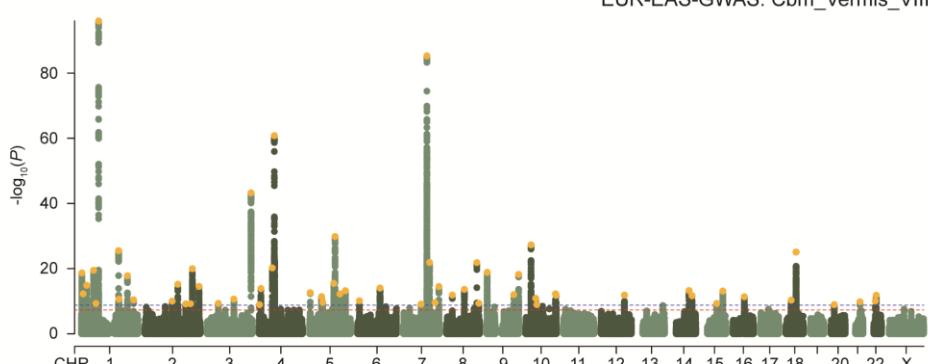
EUR-EAS-GWAS: Cbm_Vermis_VI



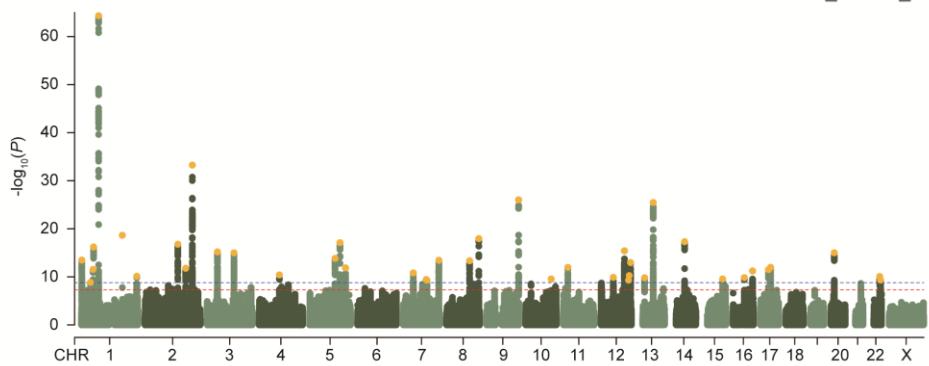
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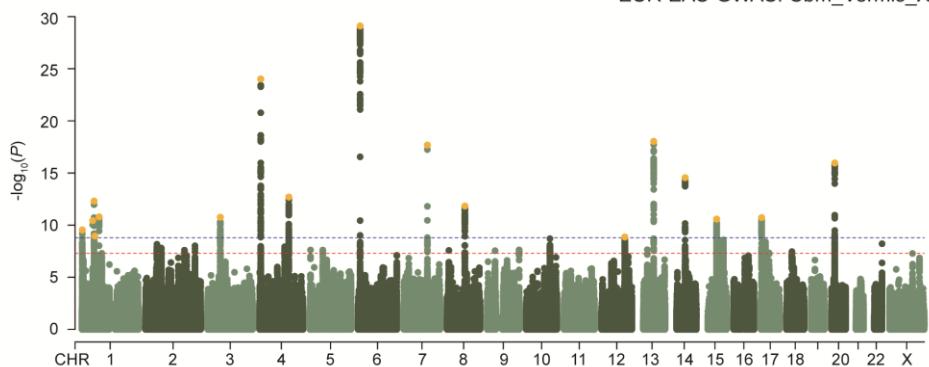
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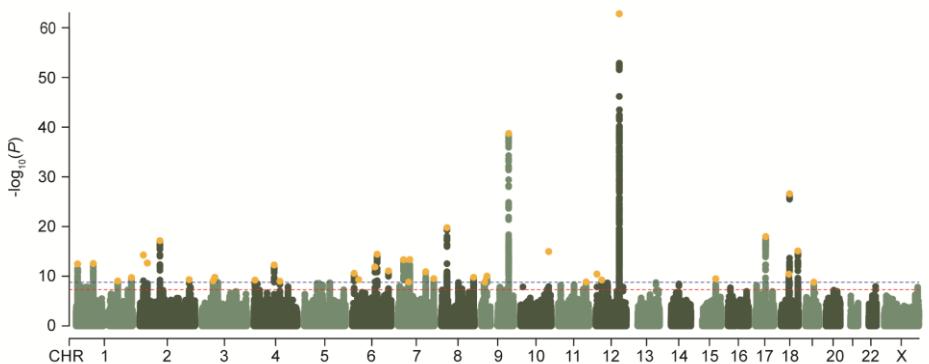
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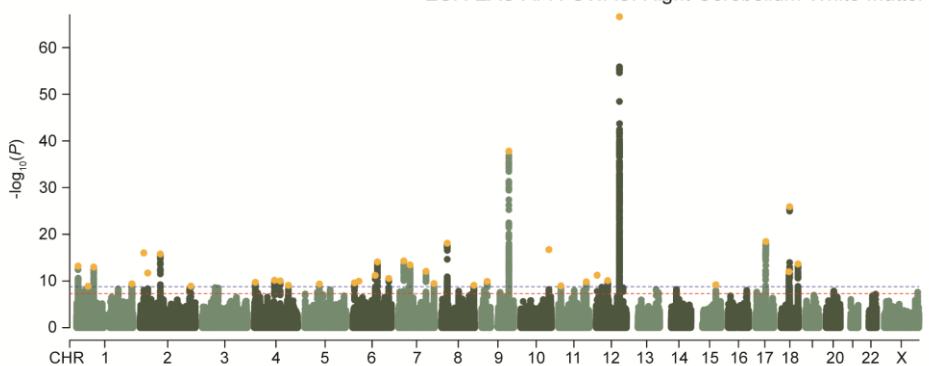
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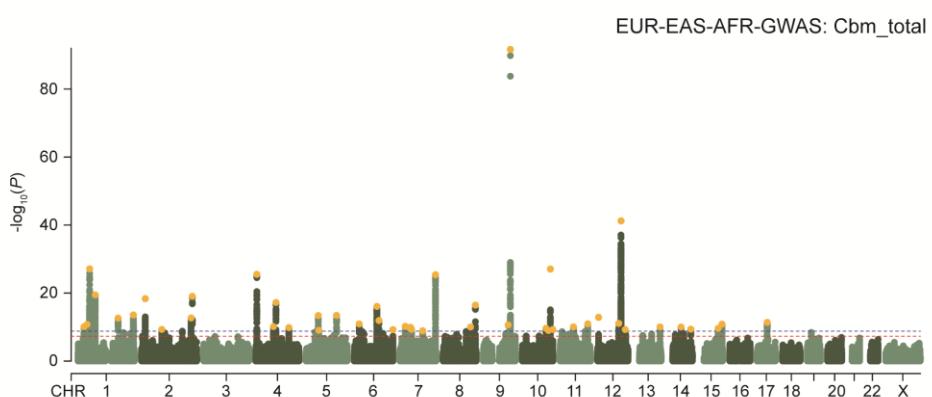
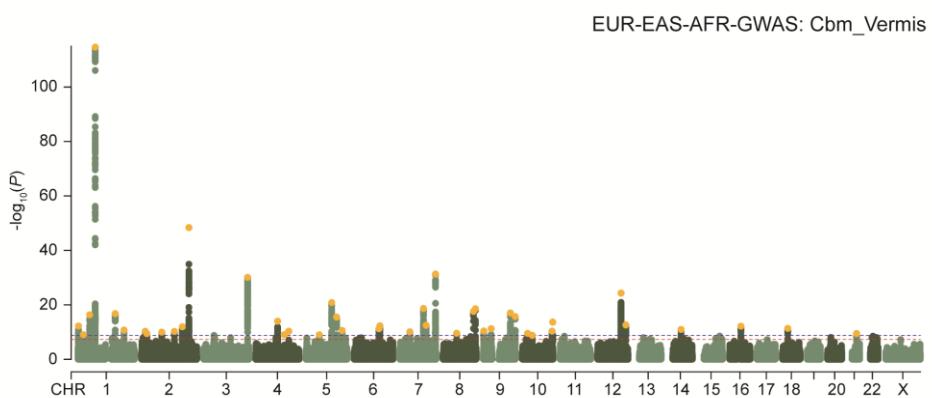
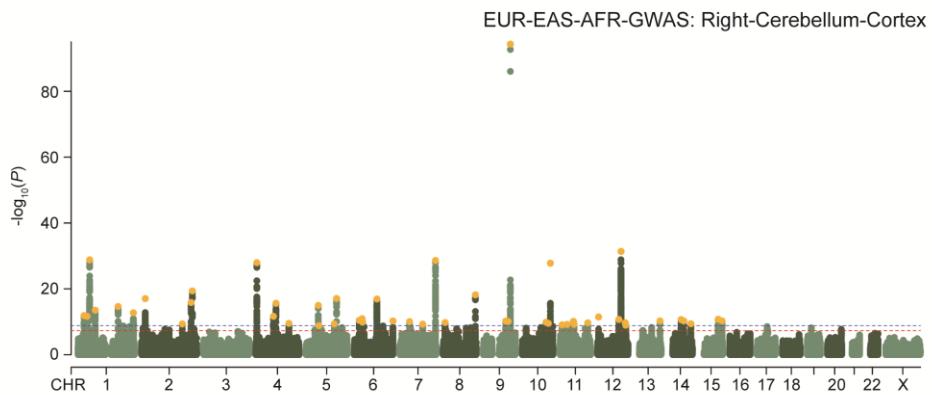
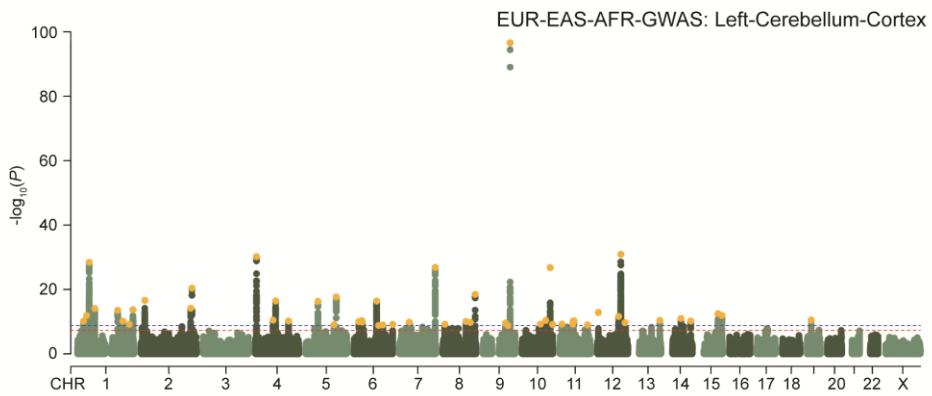


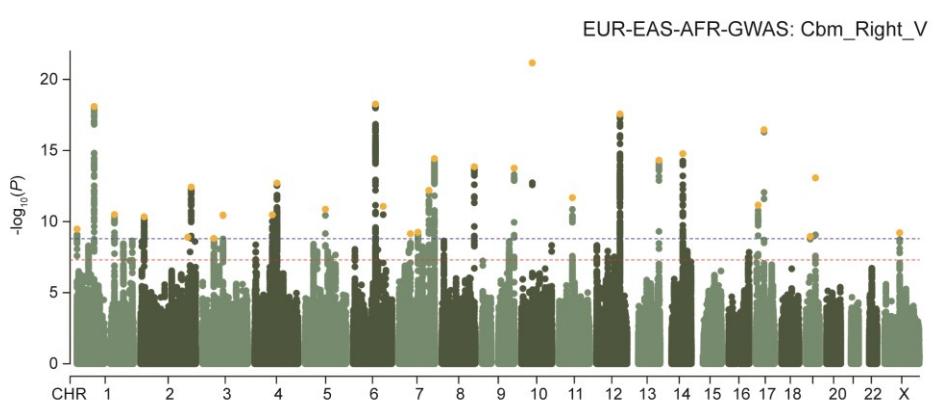
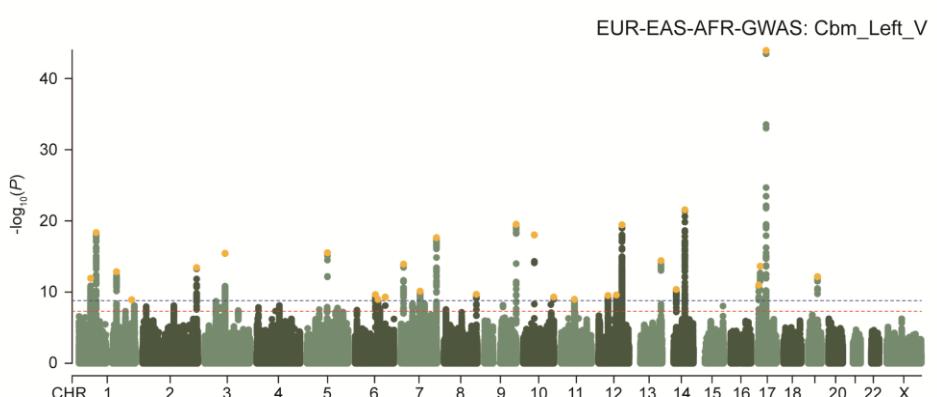
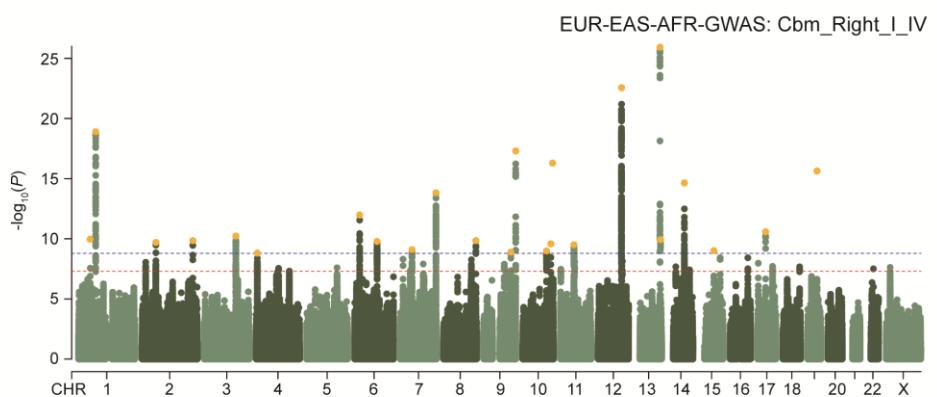
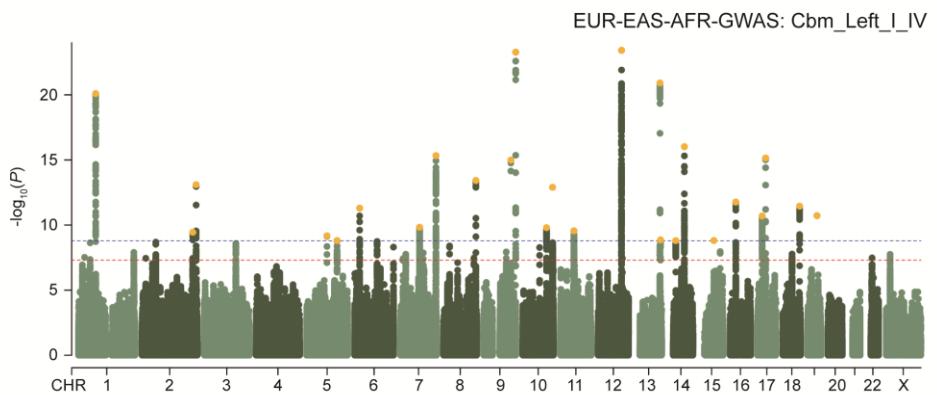
EUR-EAS-AFR-GWAS: Left-Cerebellum-White-Matter

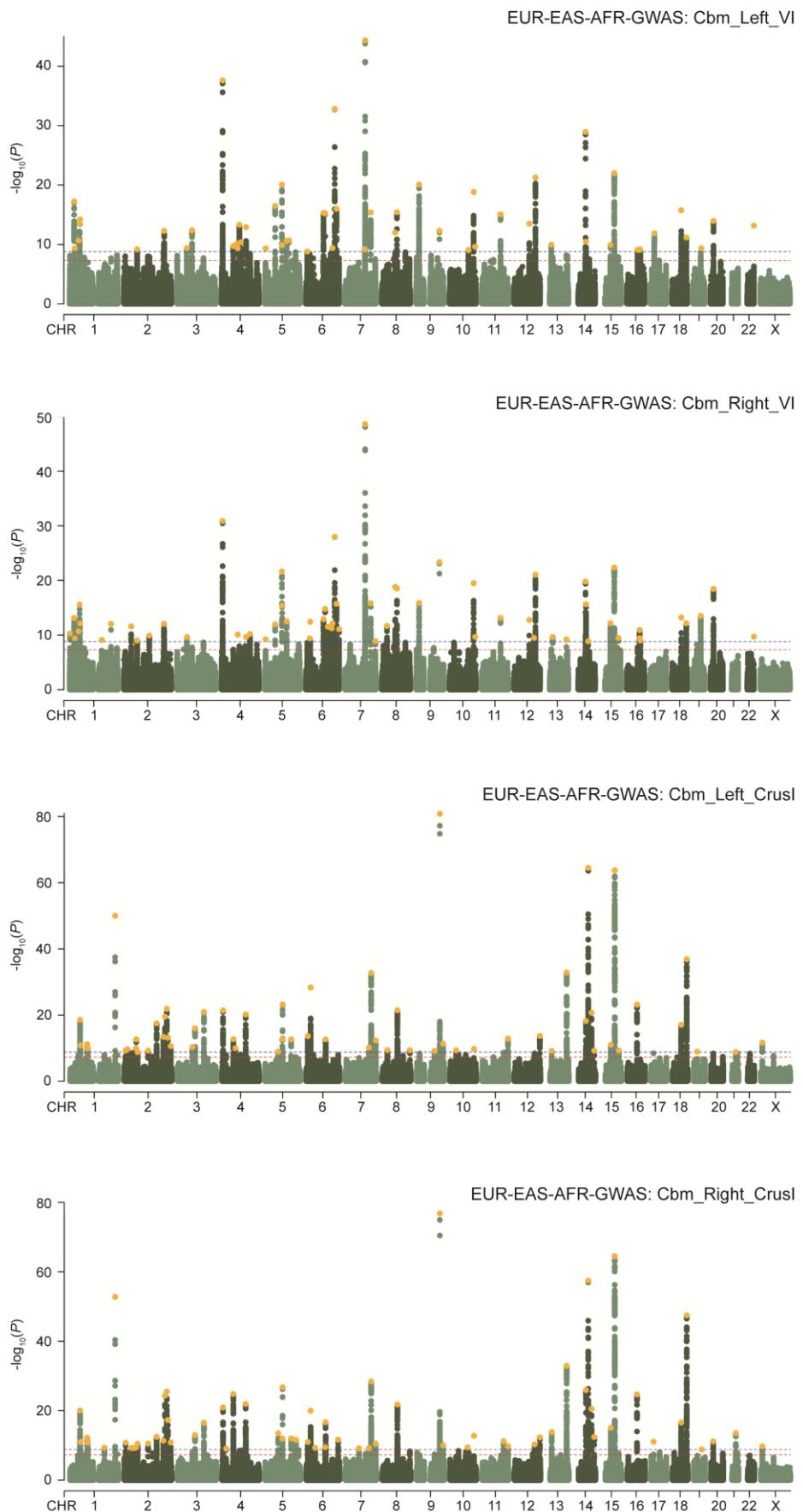


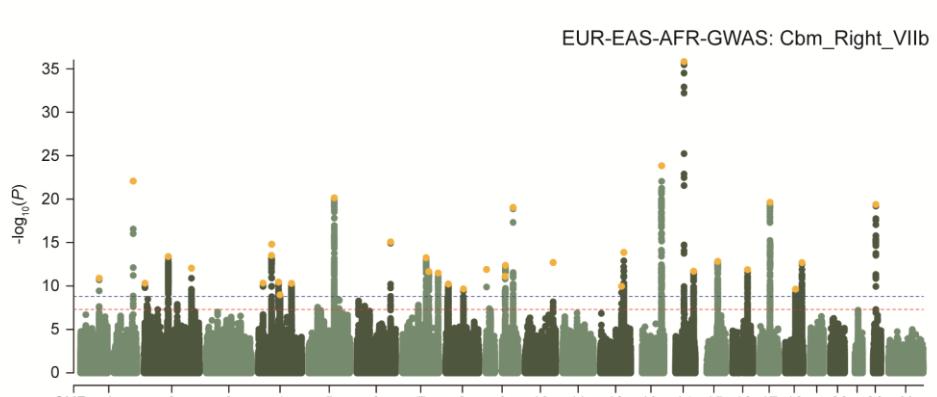
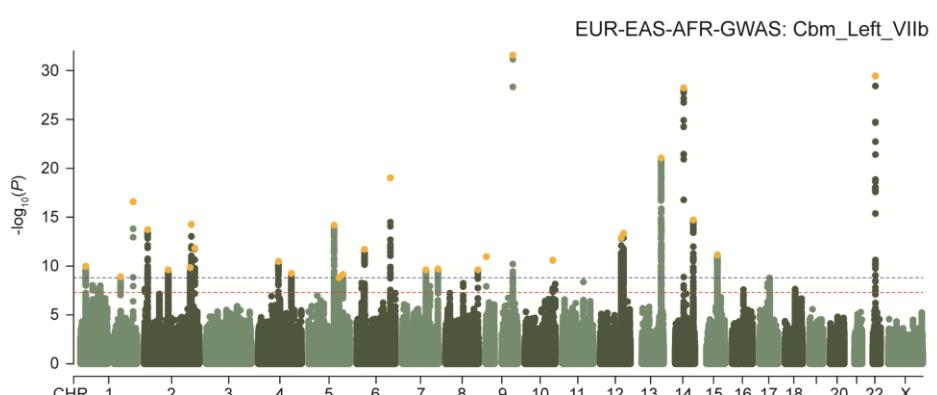
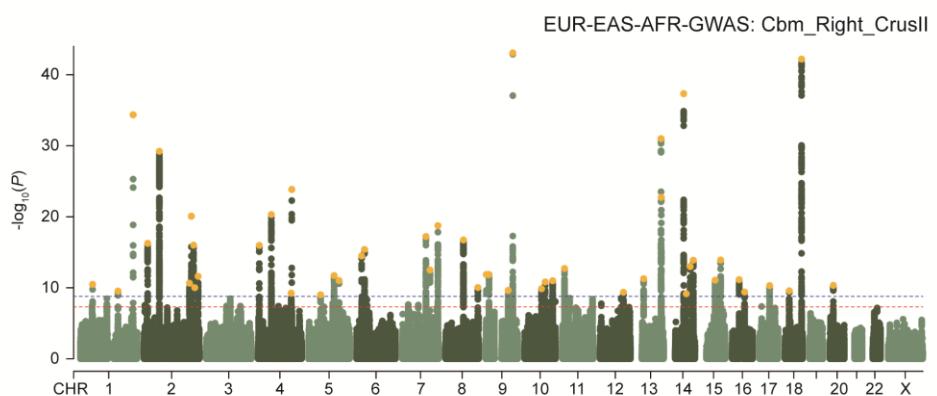
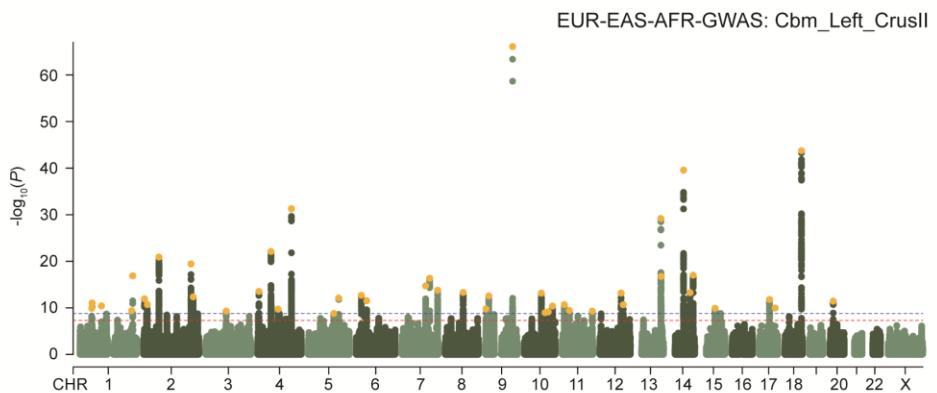
EUR-EAS-AFR-GWAS: Right-Cerebellum-White-Matter

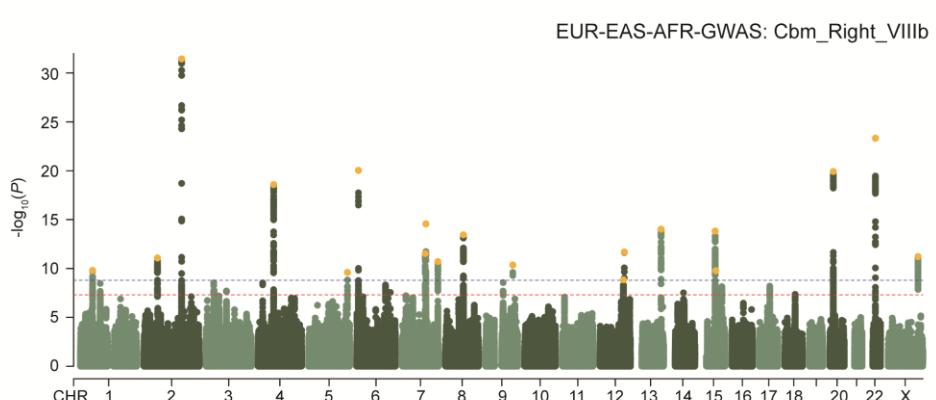
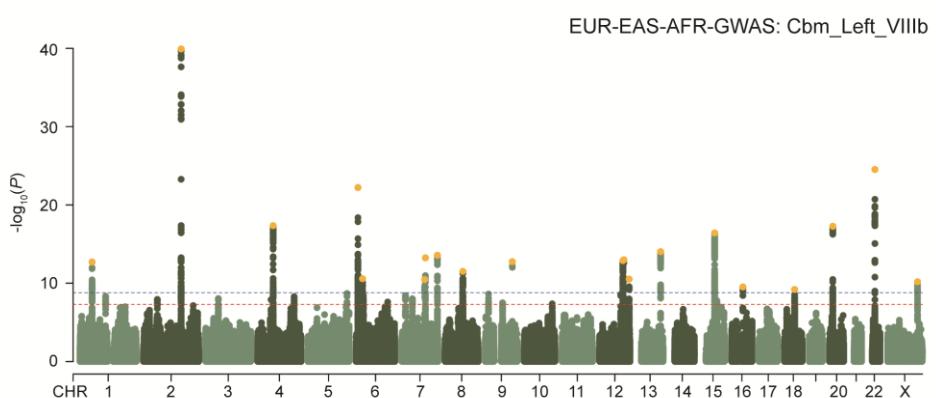
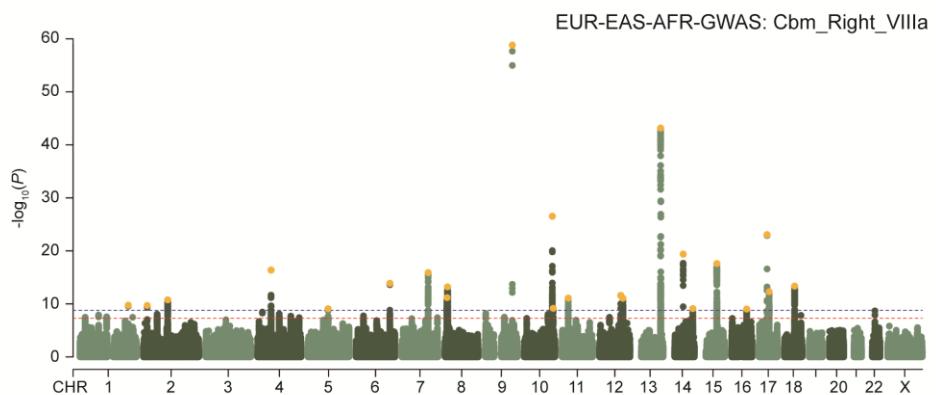
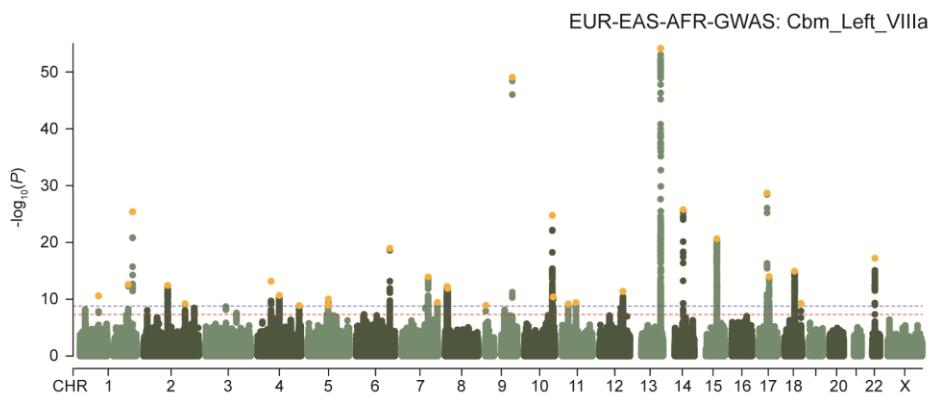


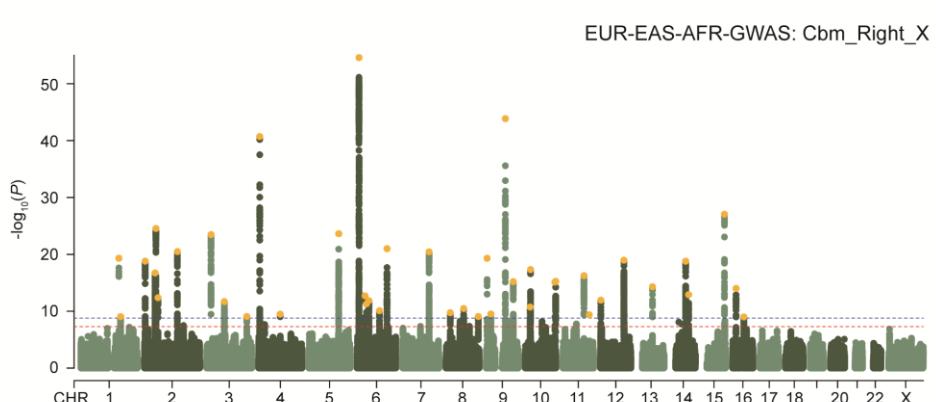
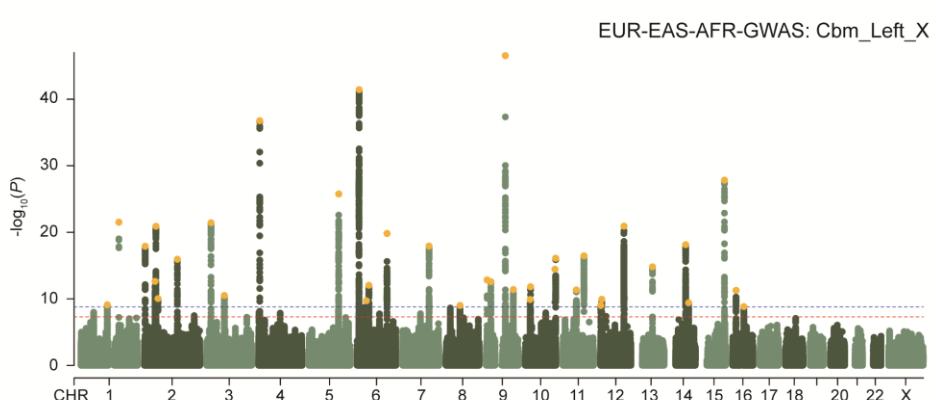
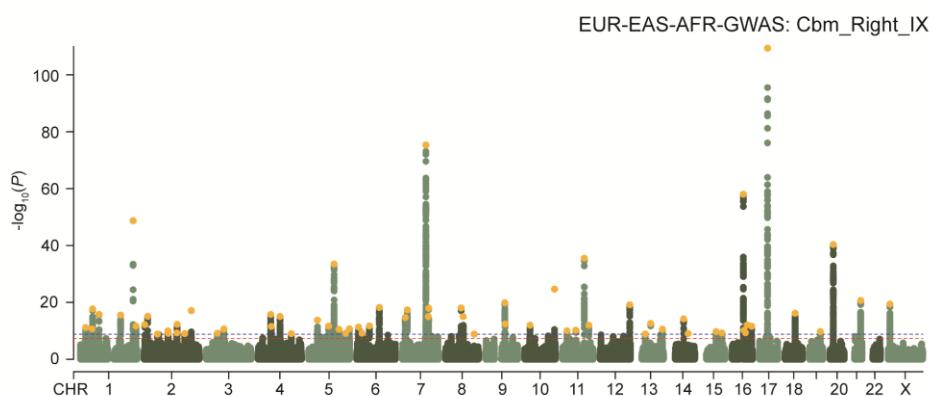
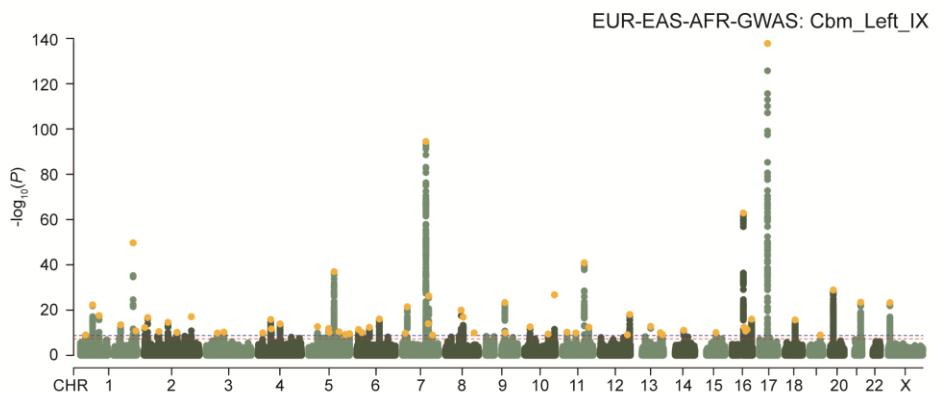


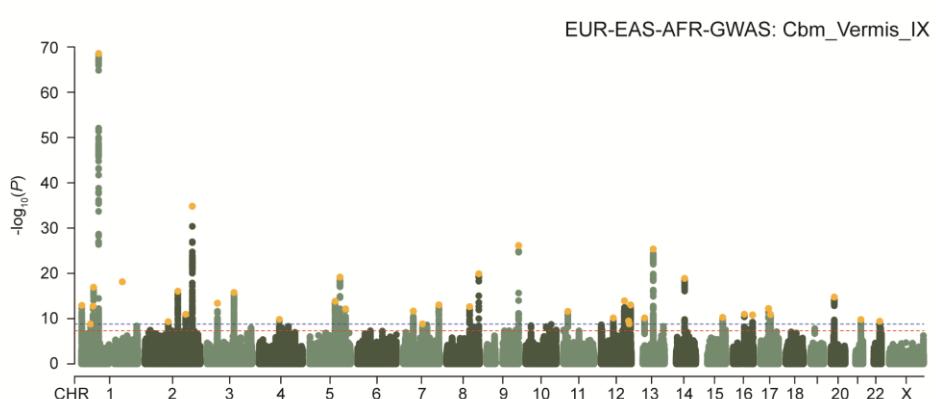
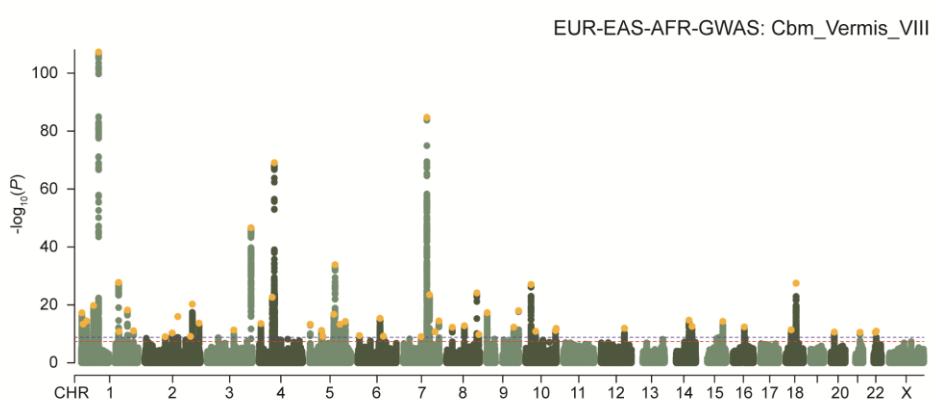
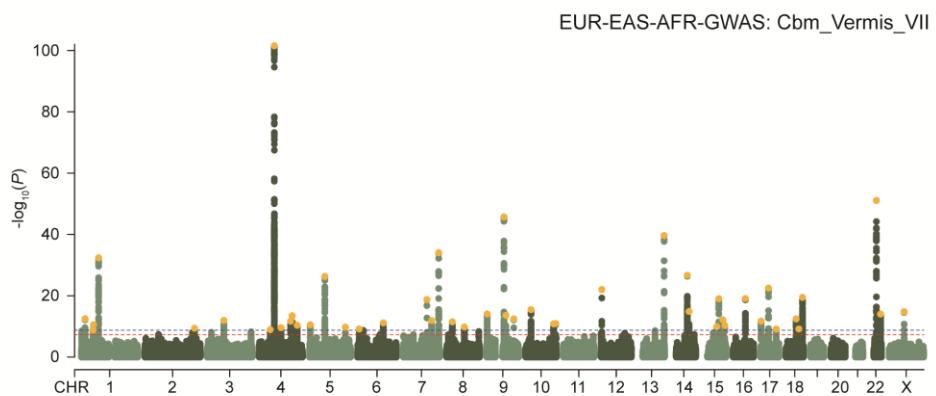
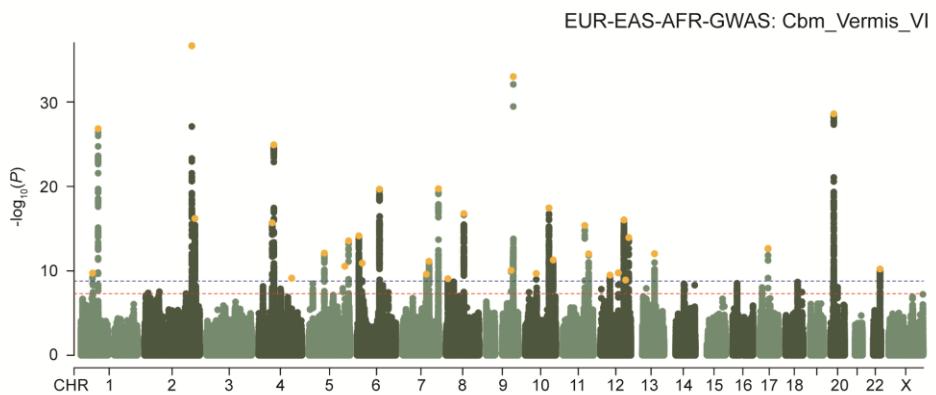












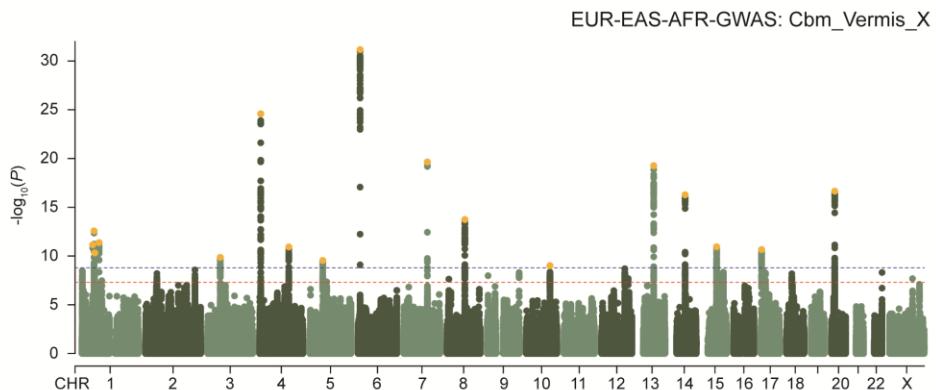
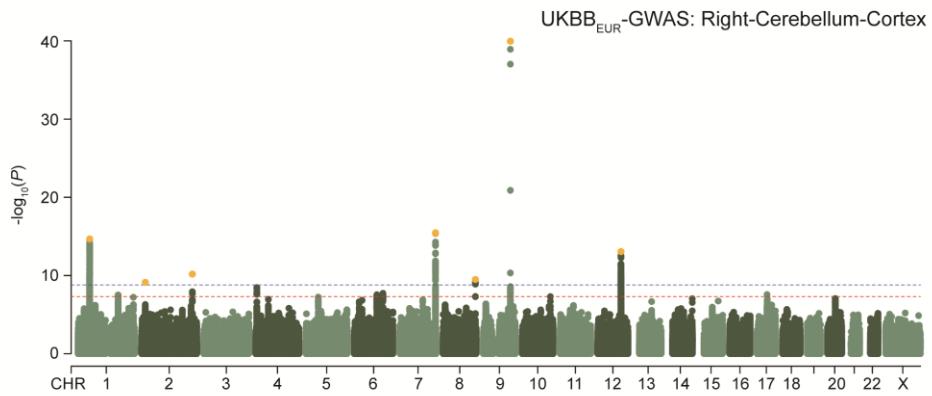
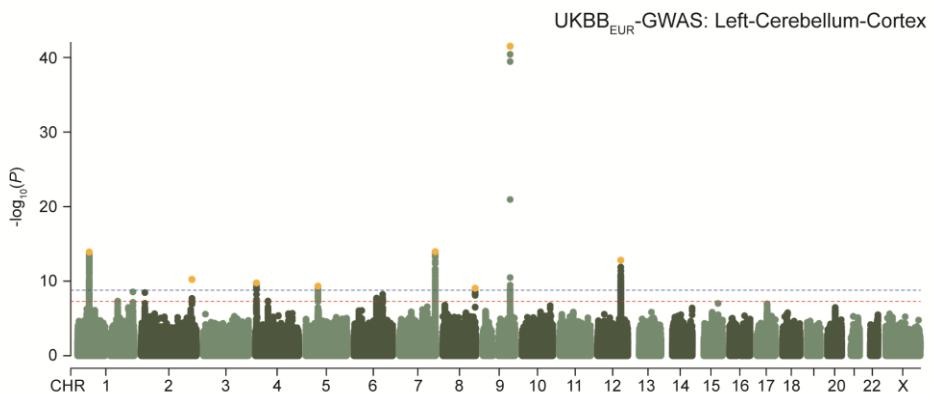
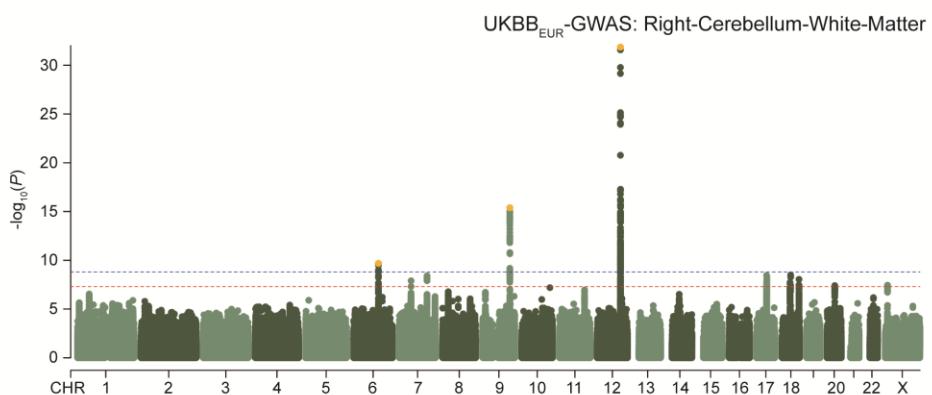
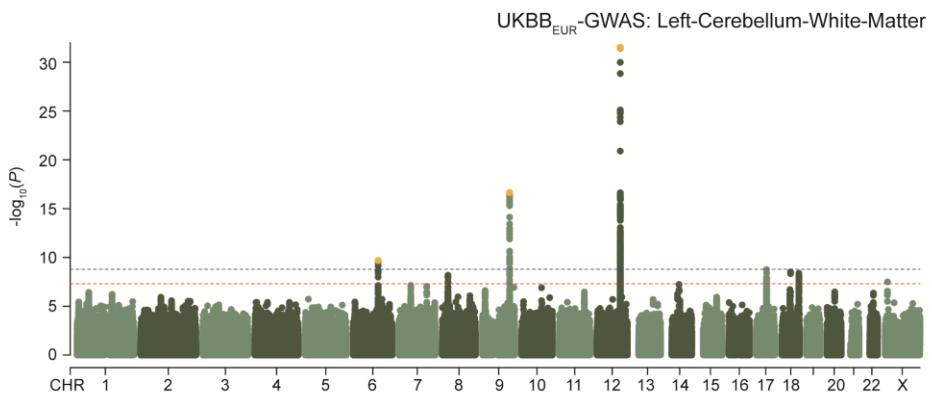
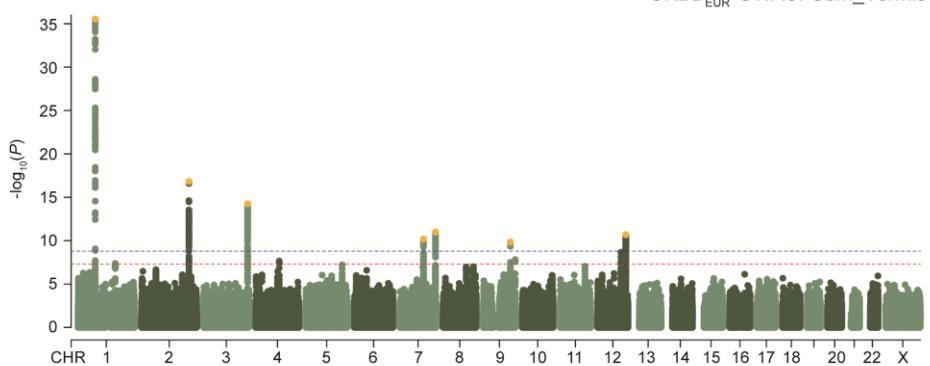


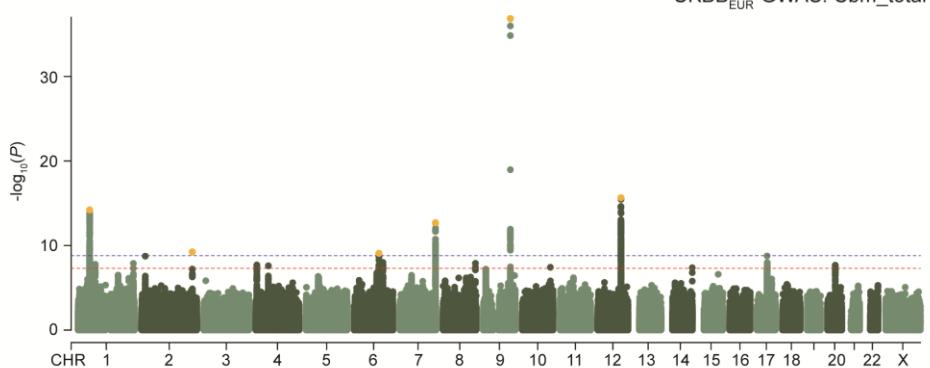
Fig. S3 | Genetic discovery of seven sets of sex-combined univariate GWAS. The Manhattan plots show the results from seven sex-combined univariate GWASs (UKBB_{EUR}-GWAS, ABCD_{EUR}-GWAS, CHIMGEN_{EAS}-GWAS, ABCD_{AFR}-GWAS, EUR-GWAS, EUR-EAS-GWAS, and EUR-EAS-AFR-GWAS) on the 31 cerebellar volumetric traits. Each plot shows genome-wide (red dashed line: $P = 5 \times 10^{-8}$) and study-wide (blue dashed line: $P = 1.61 \times 10^{-9}$) significant associations with 31 cerebellar volumetric traits. Each point represents a single genetic variant plotted according to its genomic position (x axis) and its $-\log_{10}(P)$ value for the association (y axis). Each yellow point represents an independent locus with study-wide significance ($P = 1.61 \times 10^{-9}$). Abbreviations: ABCD, the Adolescent Brain Cognitive Development study; AFR, African ancestry; CHIMGEN, the Chinese Imaging Genetics study; EAS, East Asian ancestry; EUR, European ancestry; GWAS, genome-wide association study; UKBB, the UK Biobank study.



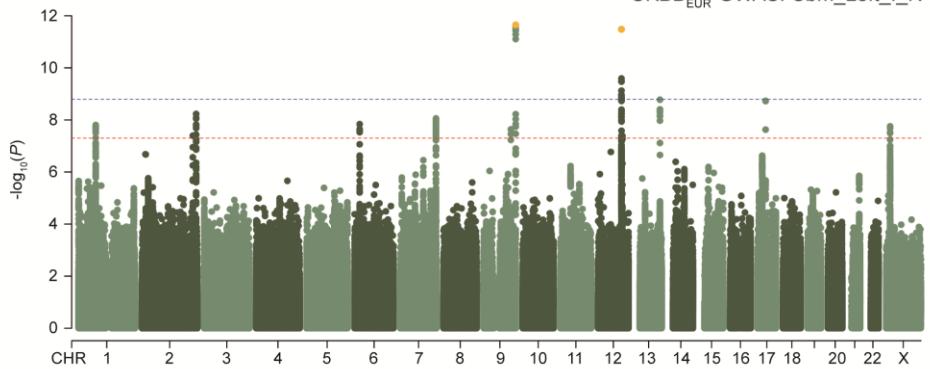
UKBB_{EUR}-GWAS: Cbm_Vermis



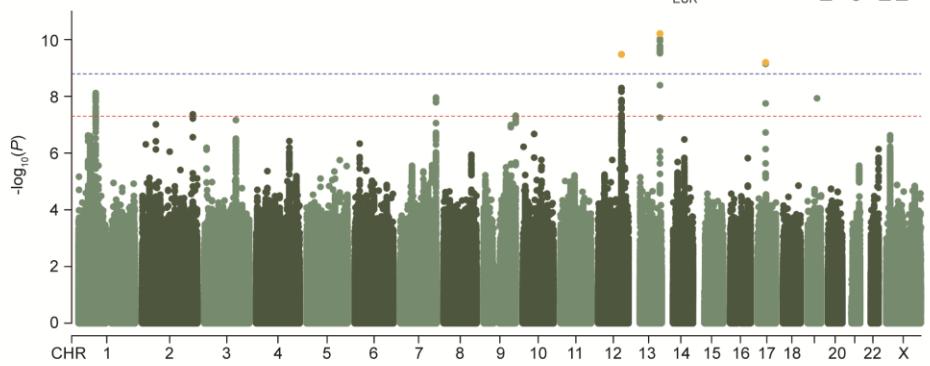
UKBB_{EUR}-GWAS: Cbm_total

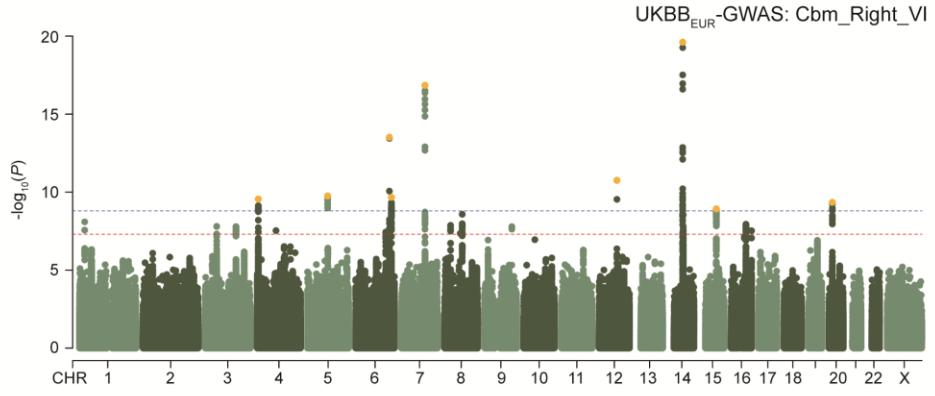
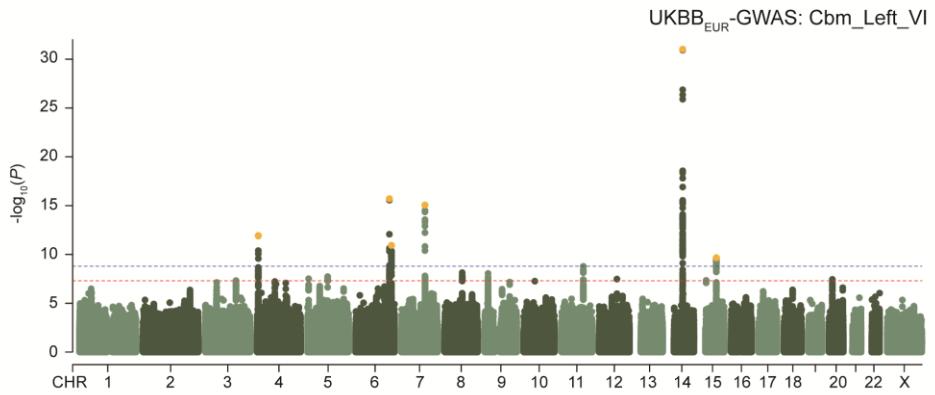
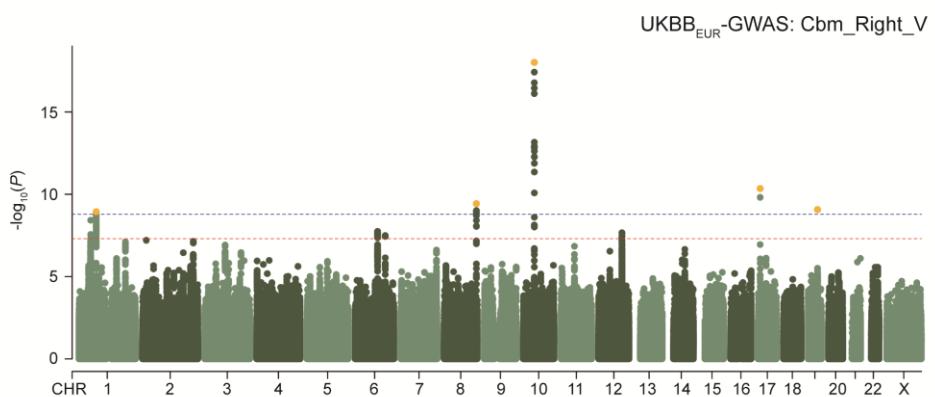
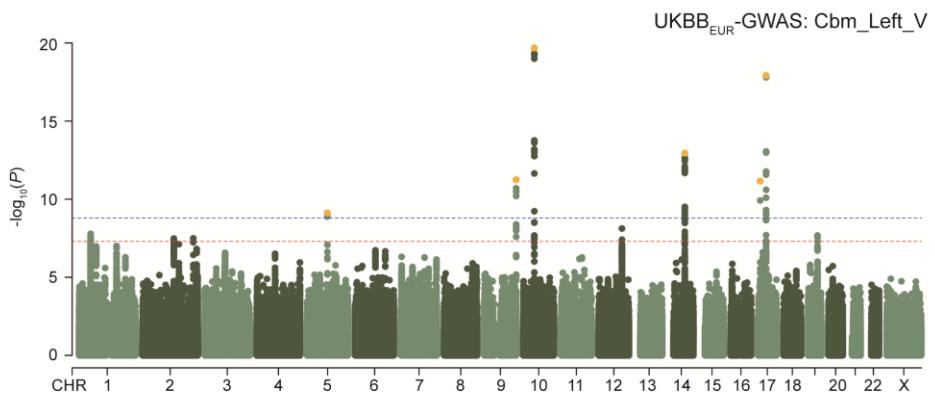


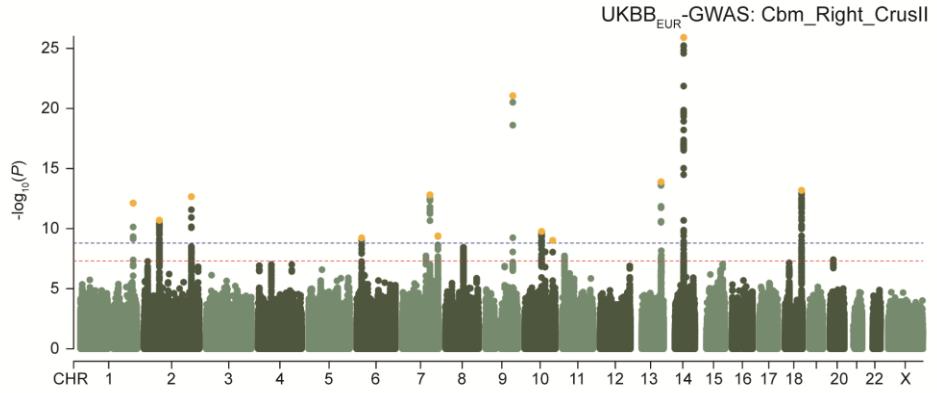
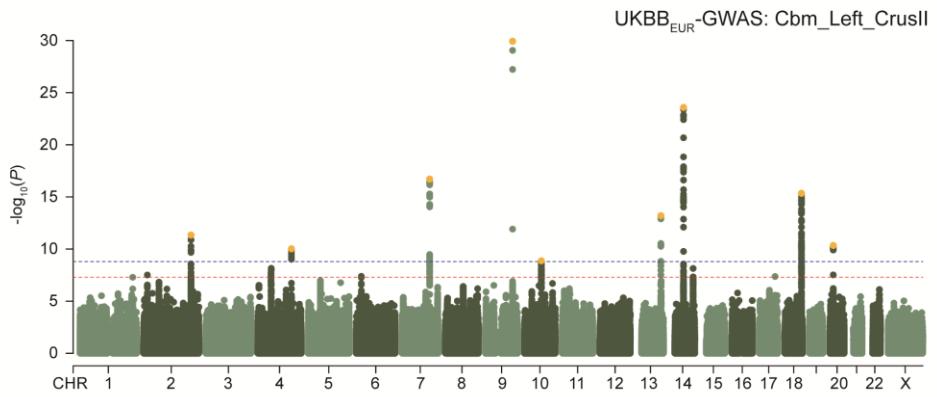
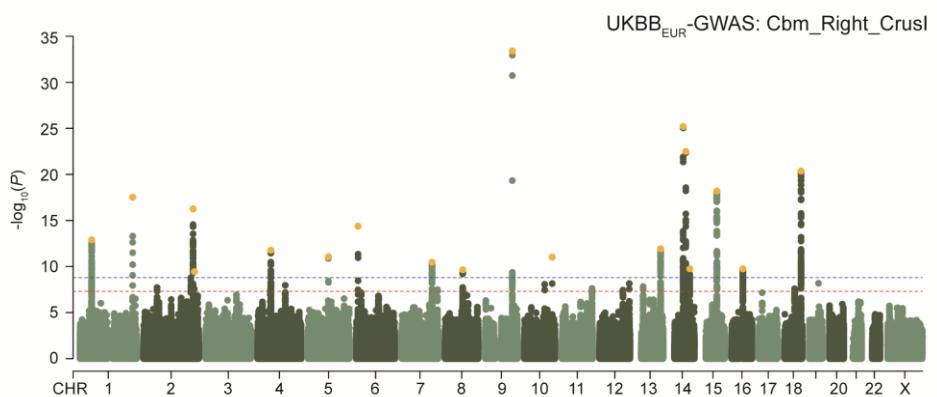
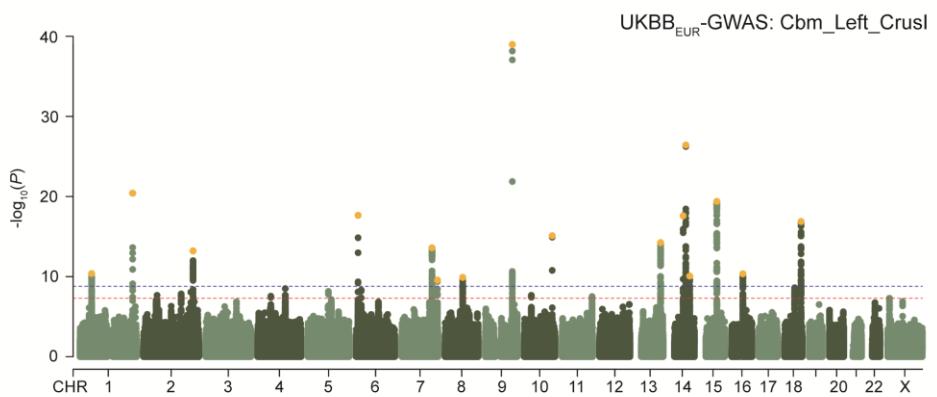
UKBB_{EUR}-GWAS: Cbm_Left_I_IV

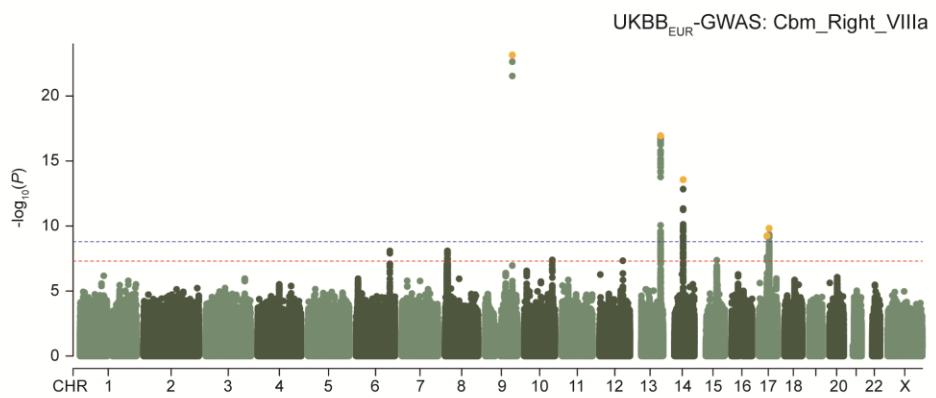
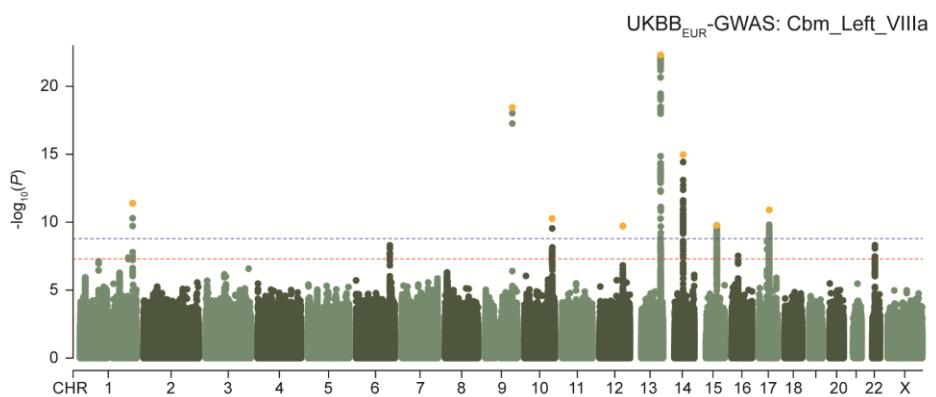
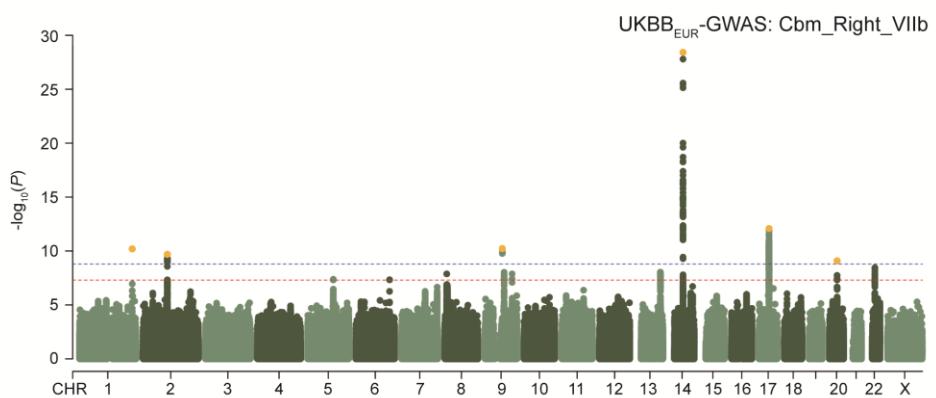
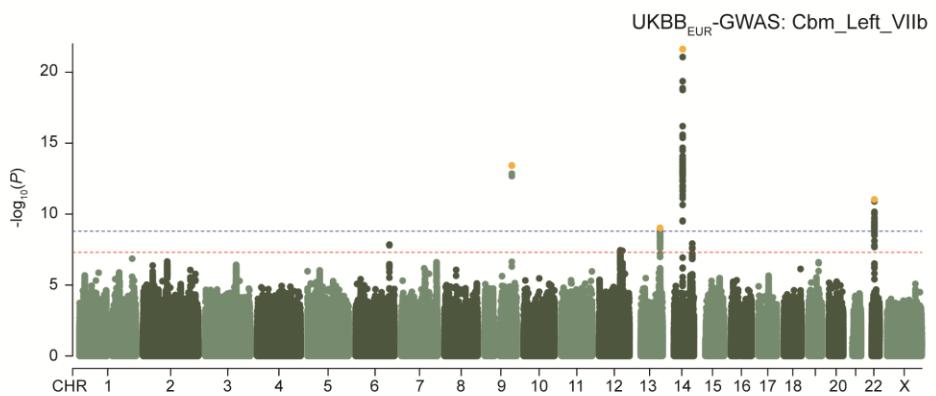


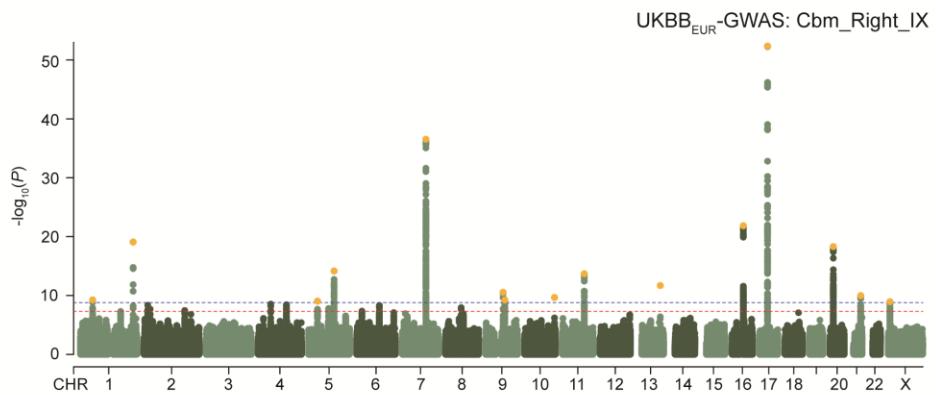
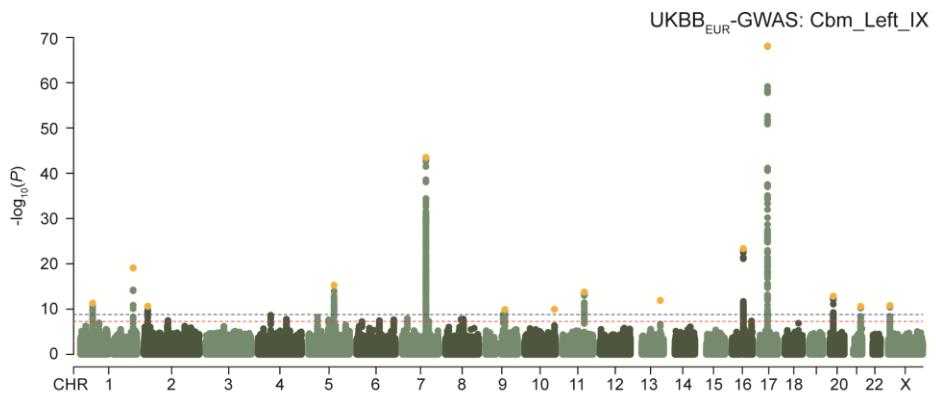
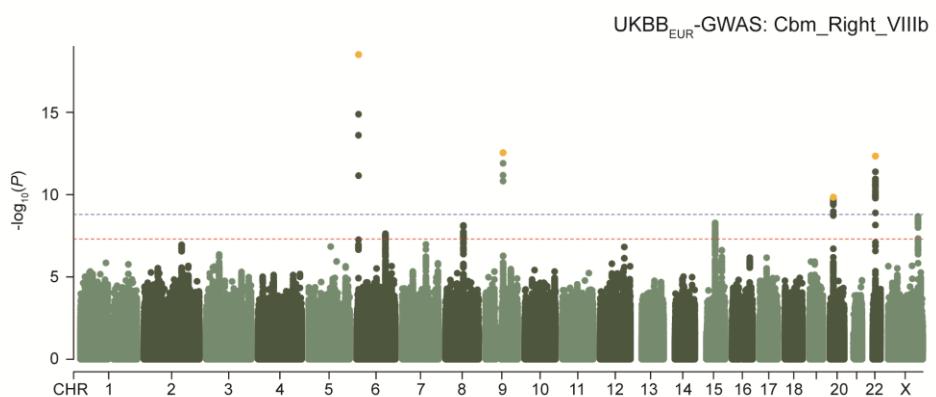
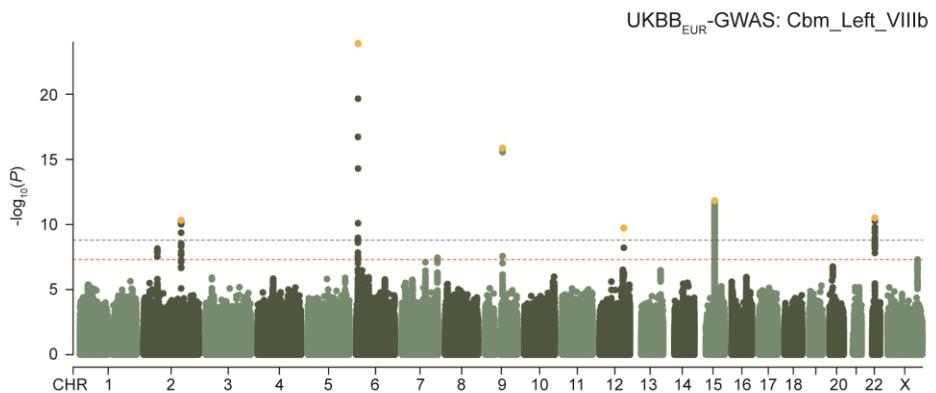
UKBB_{EUR}-GWAS: Cbm_Right_I_IV

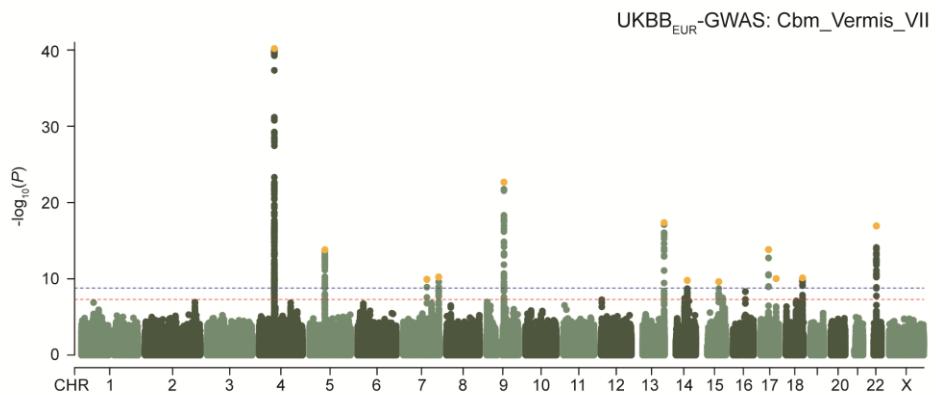
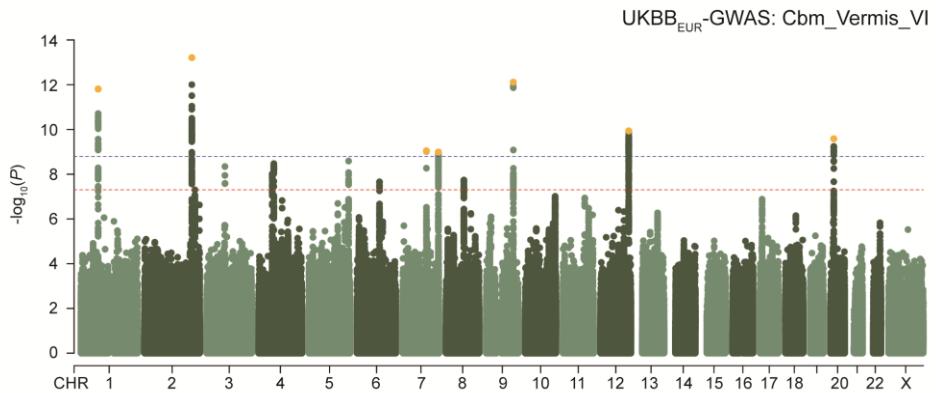
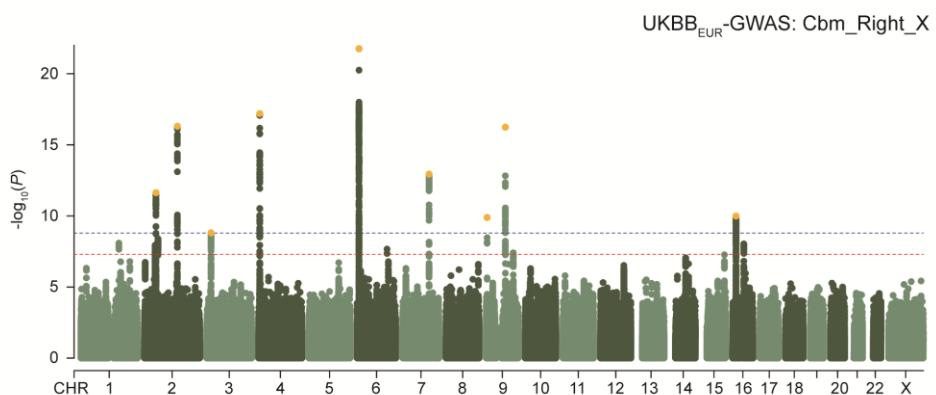
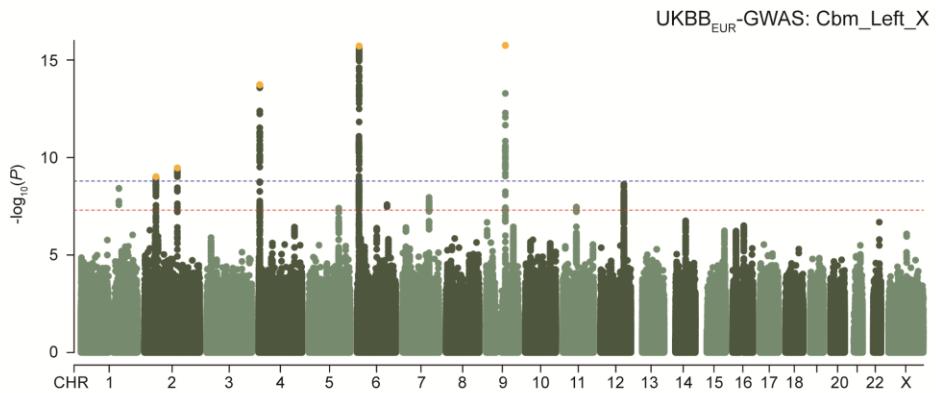


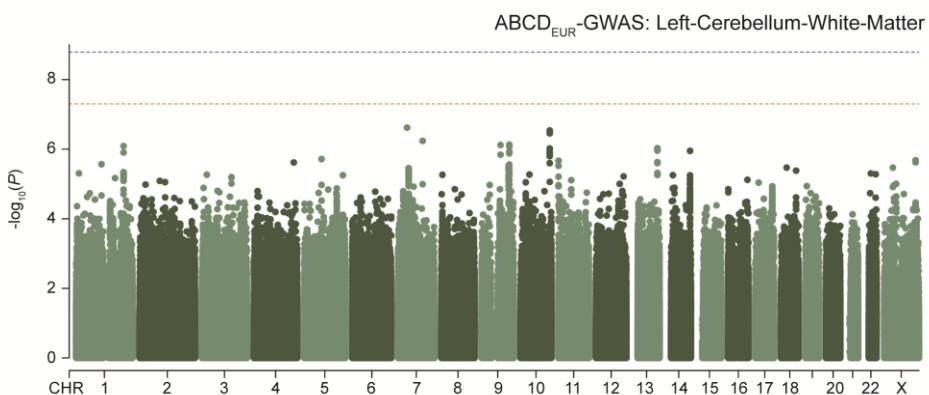
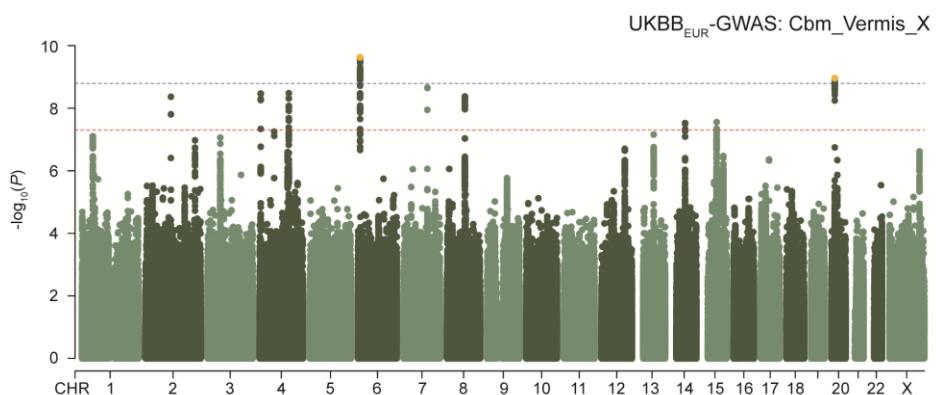
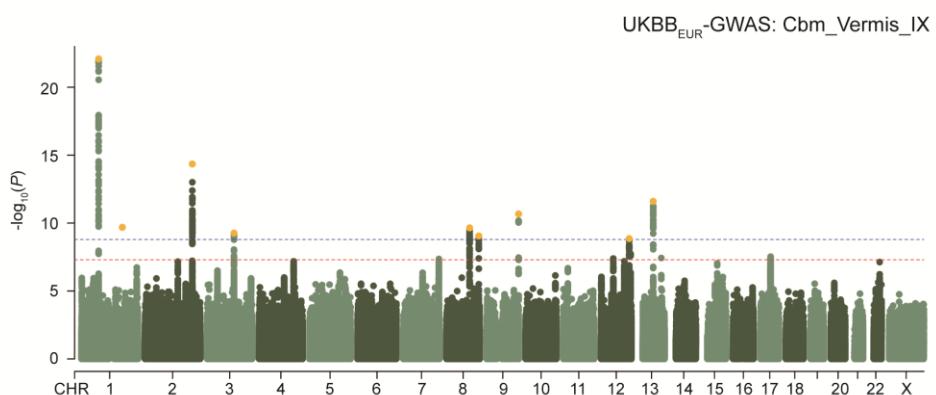
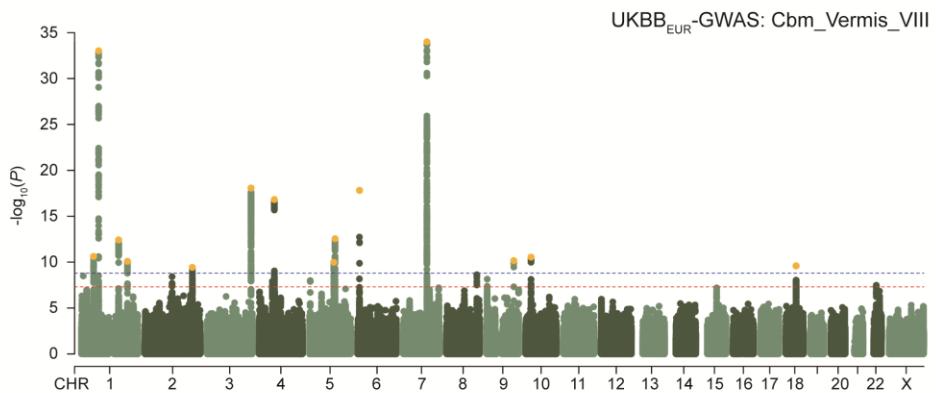




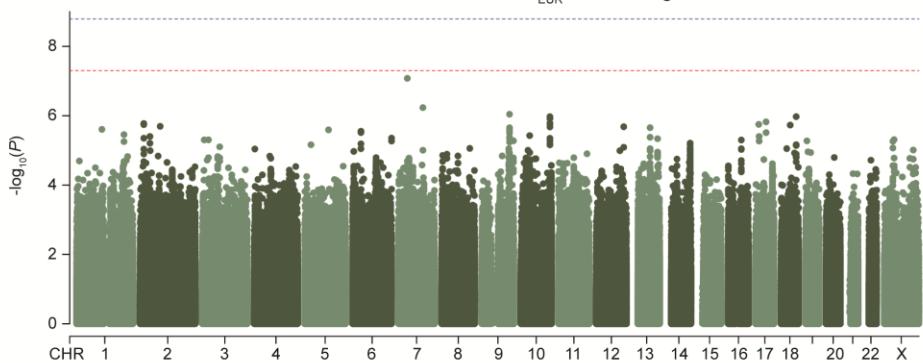




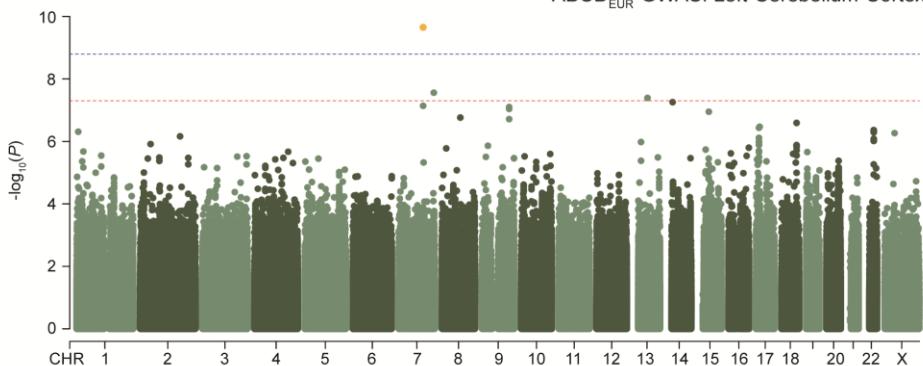




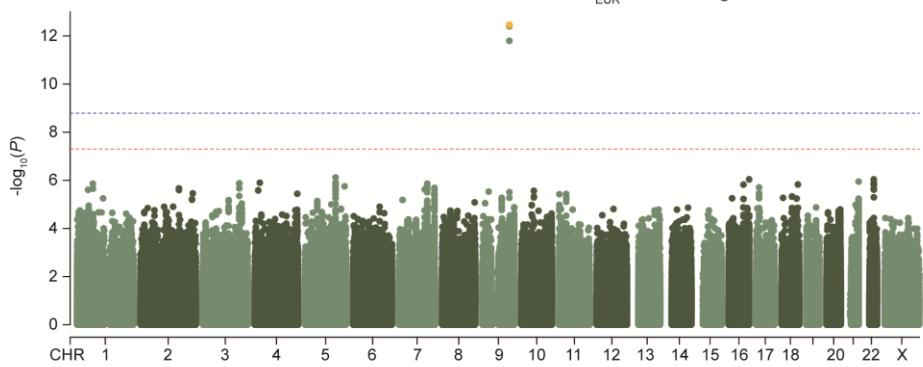
ABCD_{EUR}-GWAS: Right-Cerebellum-White-Matter



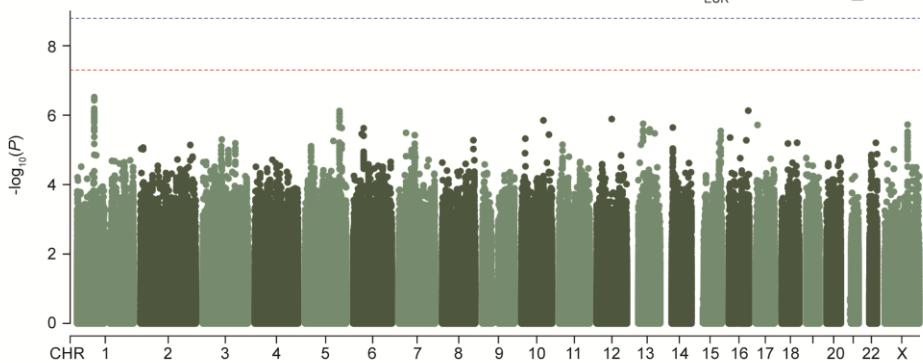
ABCD_{EUR}-GWAS: Left-Cerebellum-Cortex

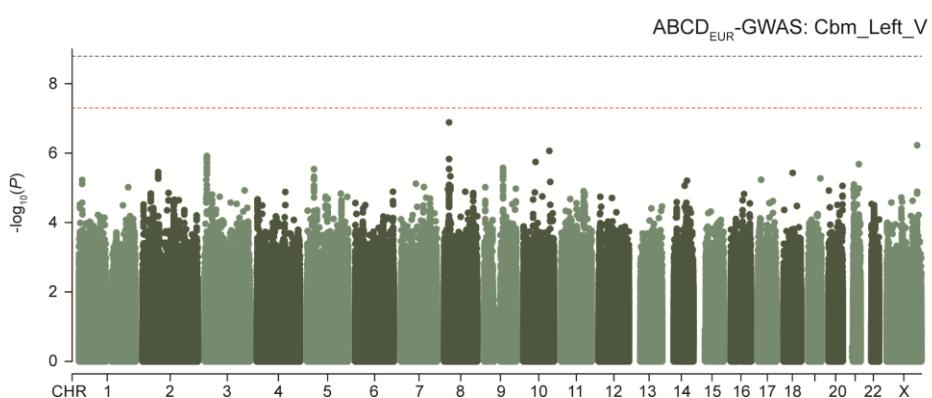
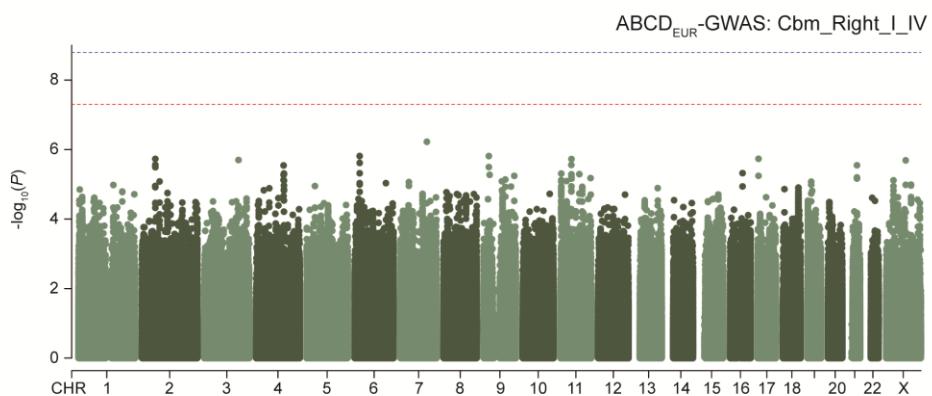
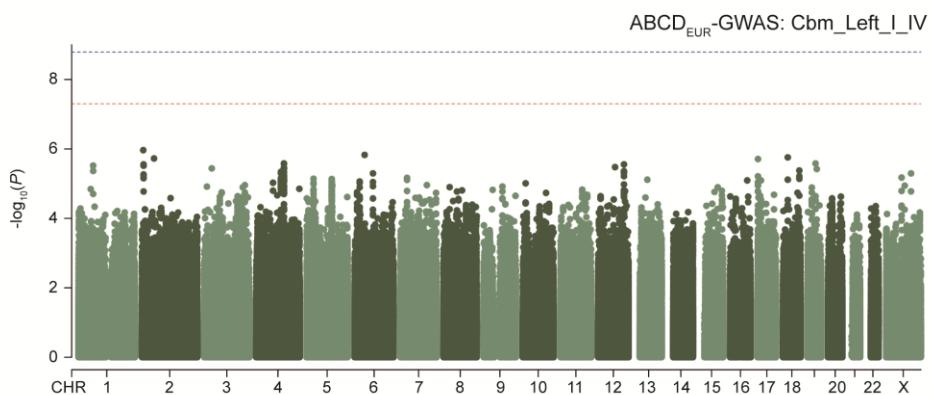
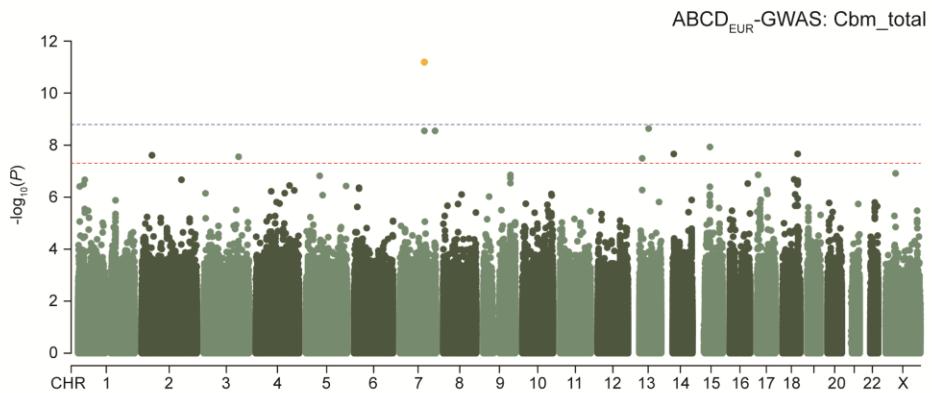


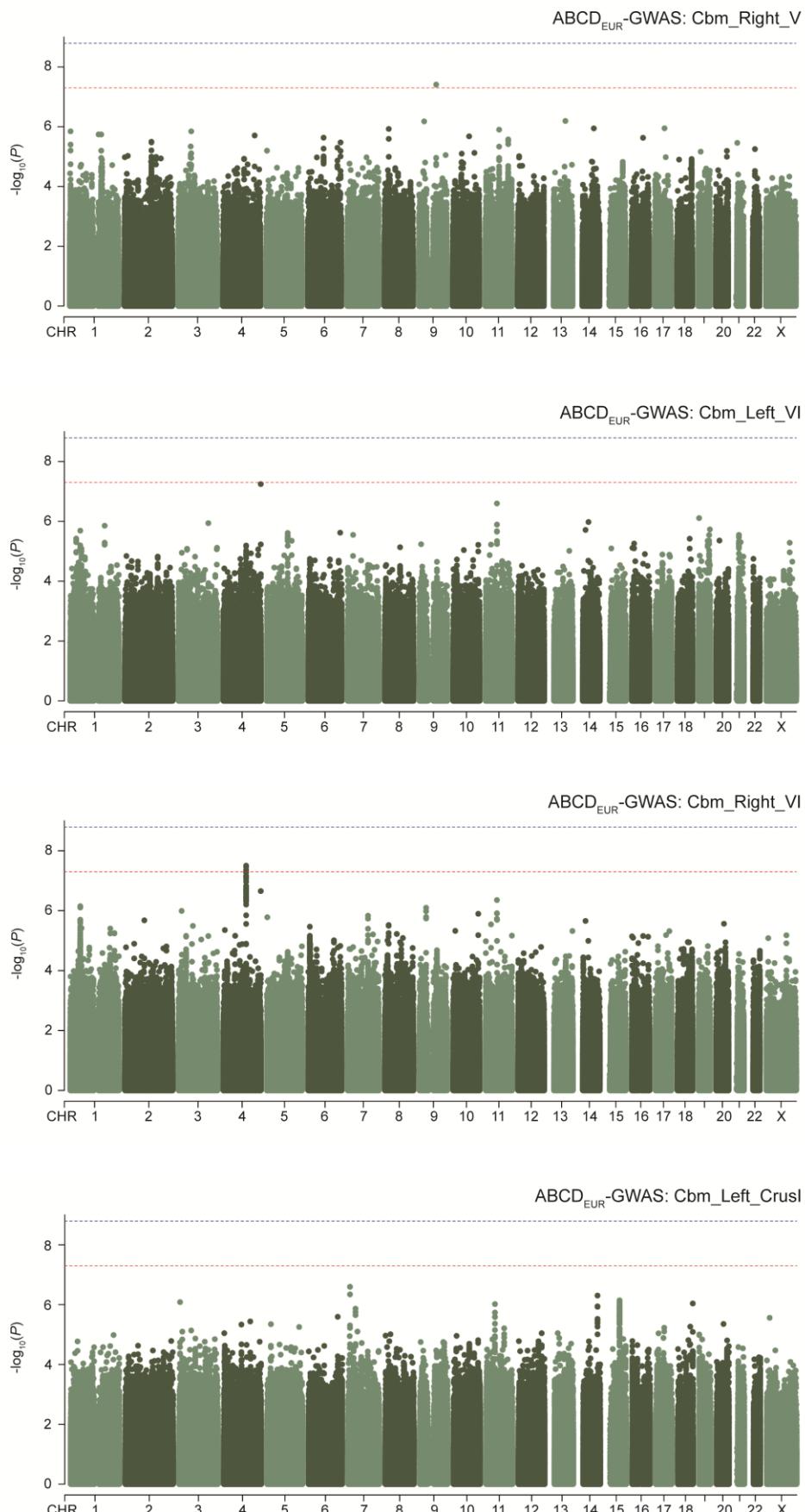
ABCD_{EUR}-GWAS: Right-Cerebellum-Cortex

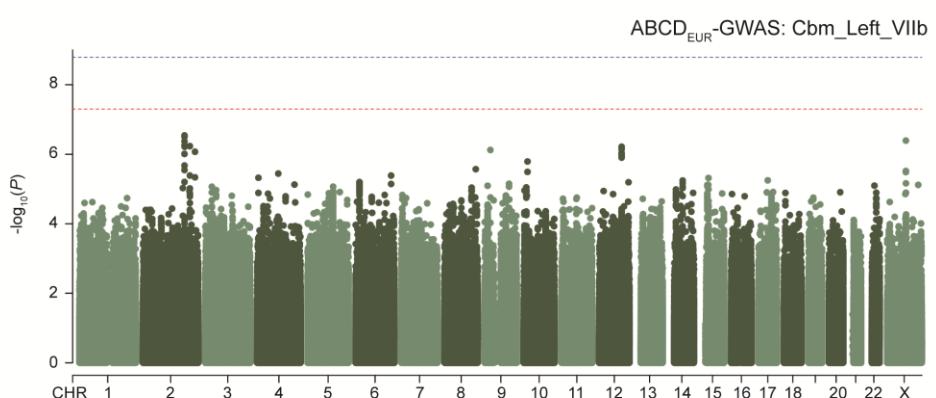
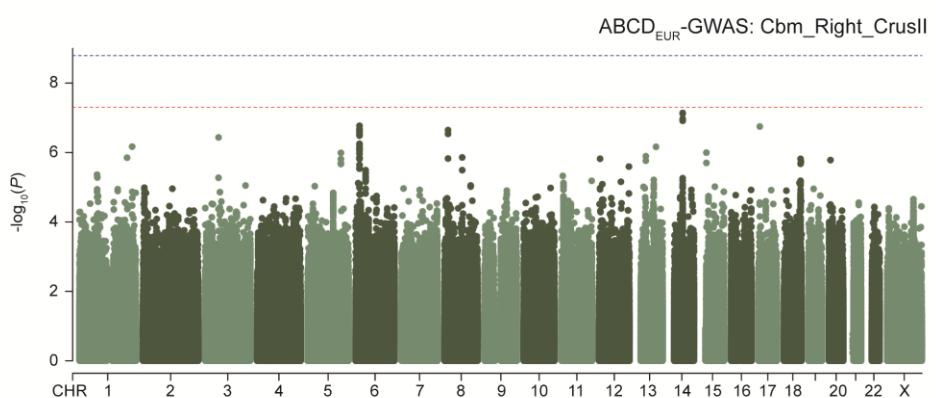
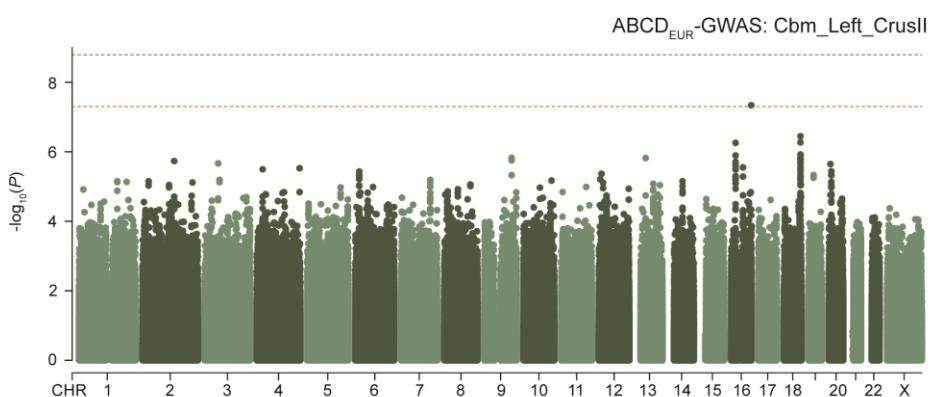
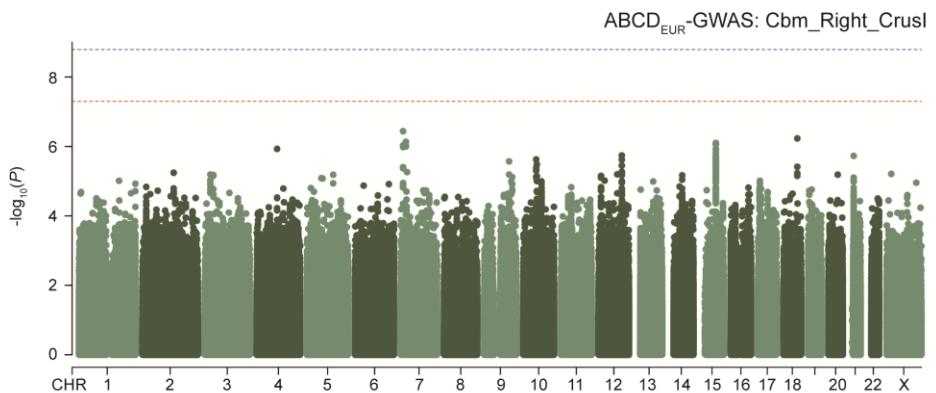


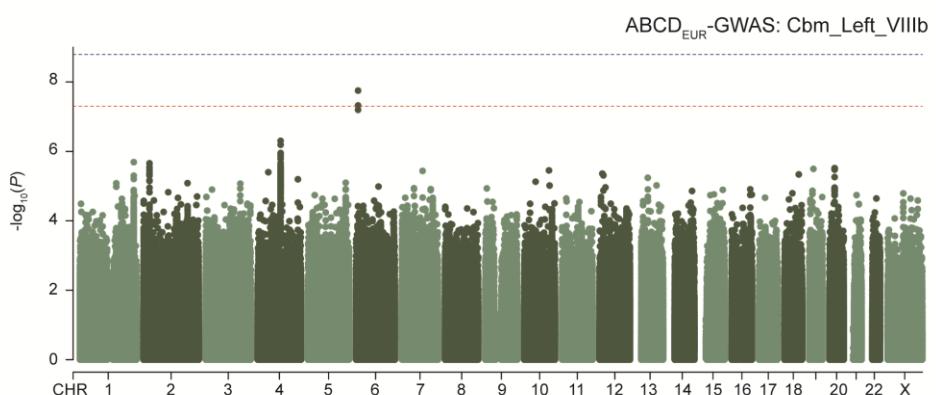
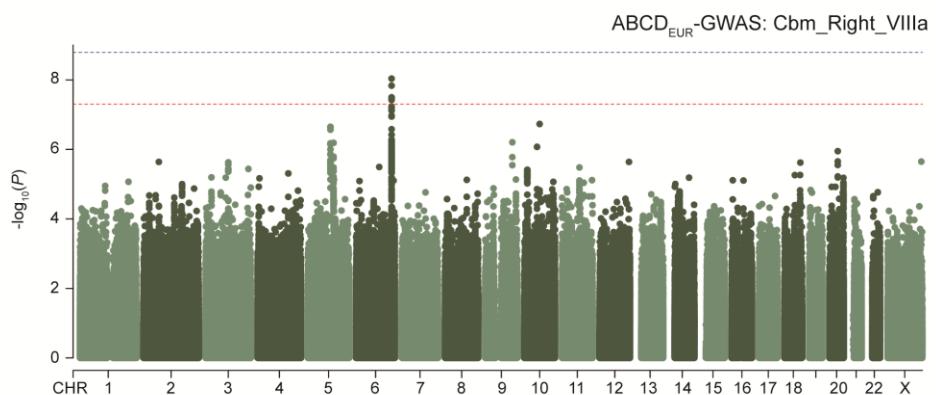
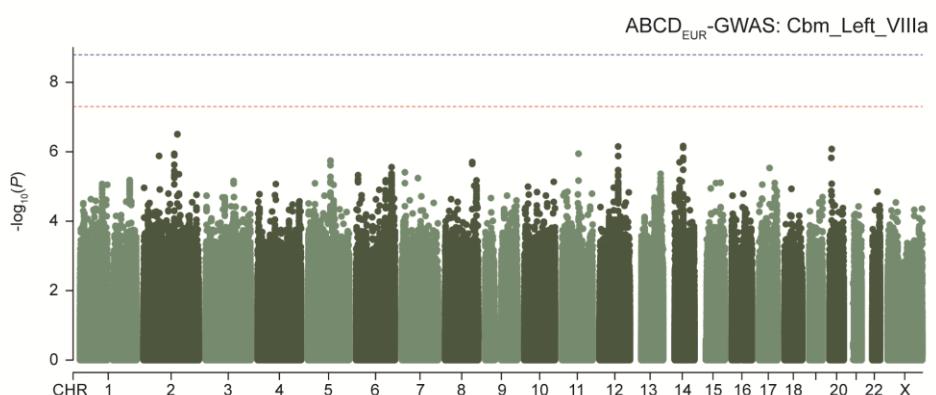
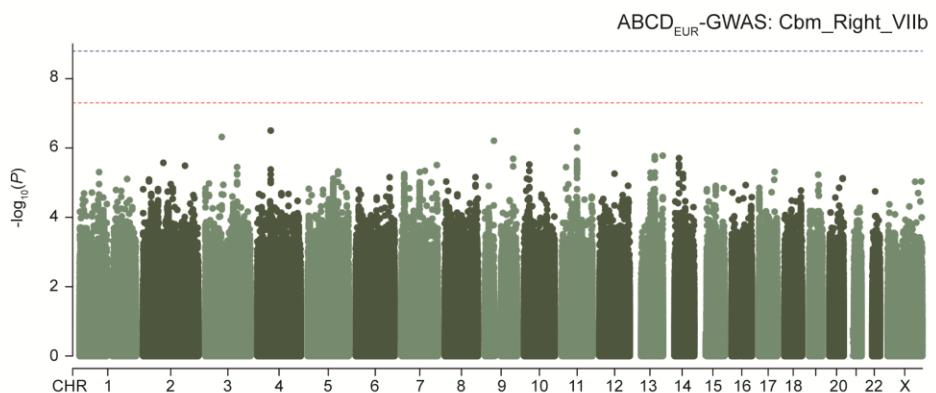
ABCD_{EUR}-GWAS: Cbm_Vermis

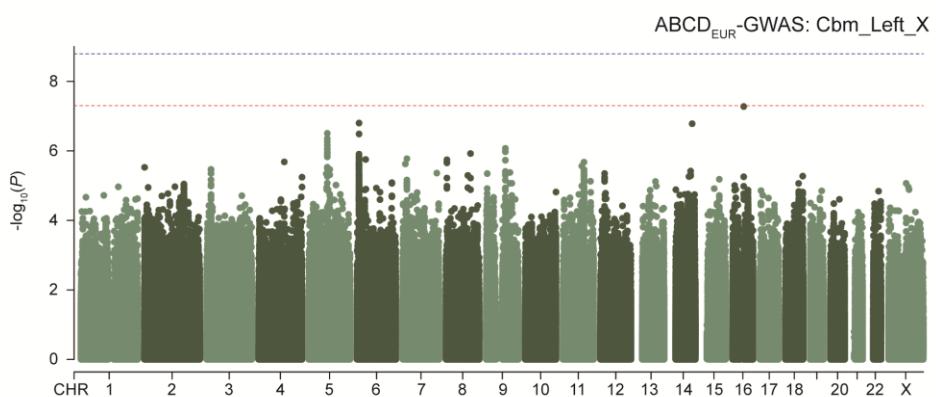
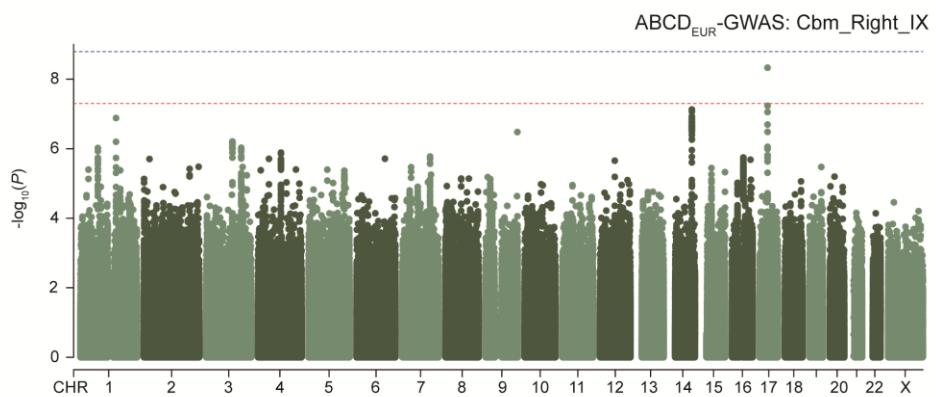
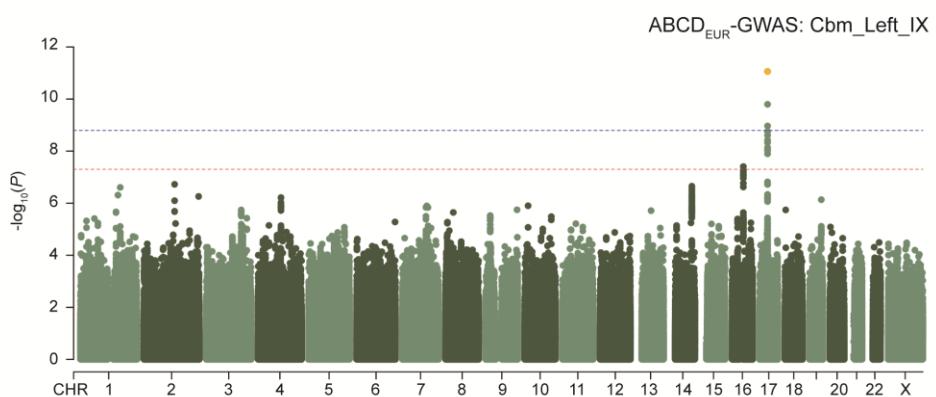
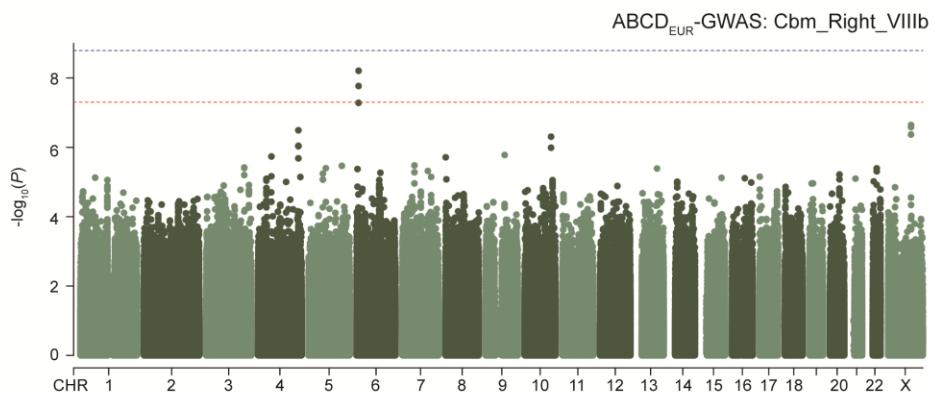


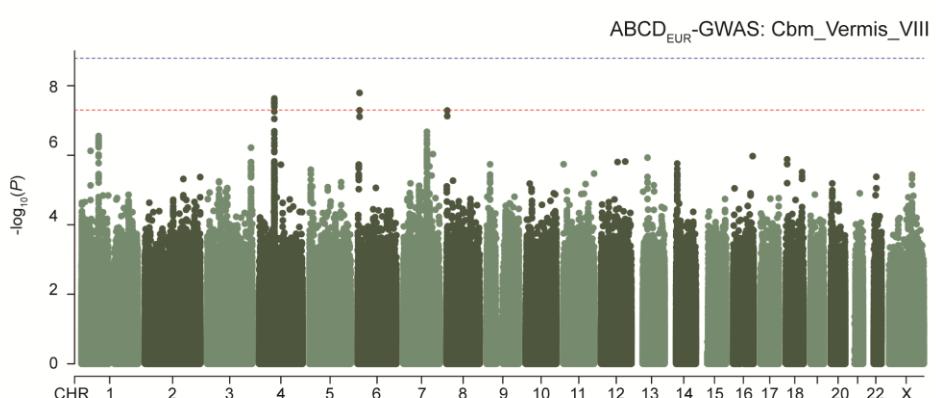
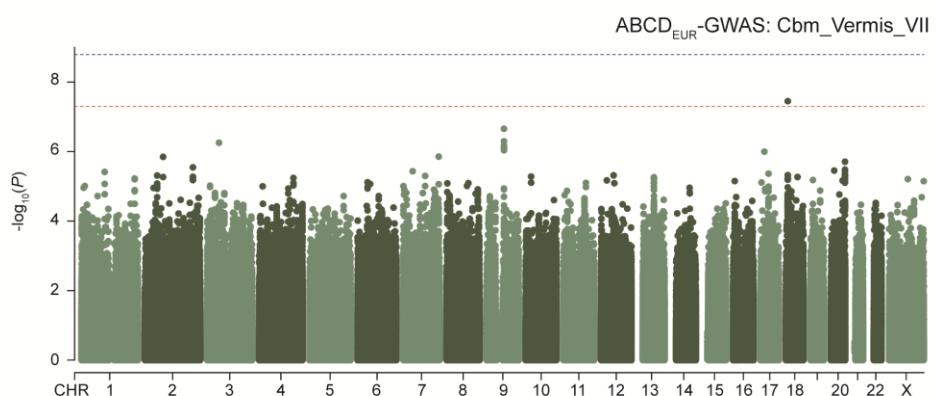
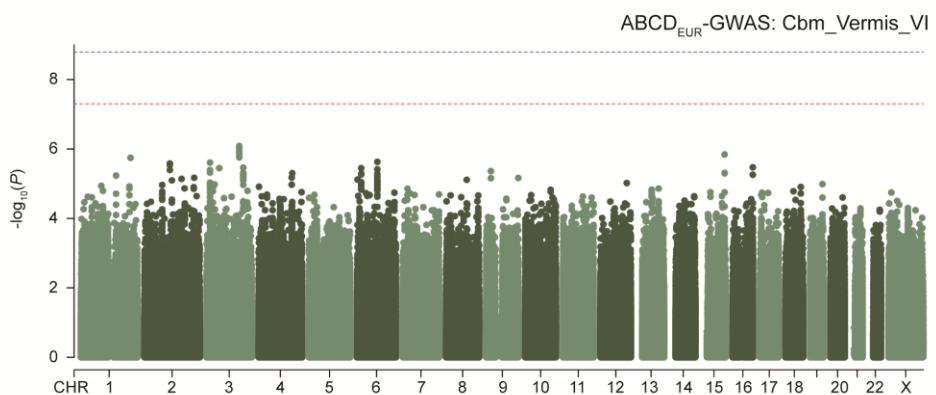
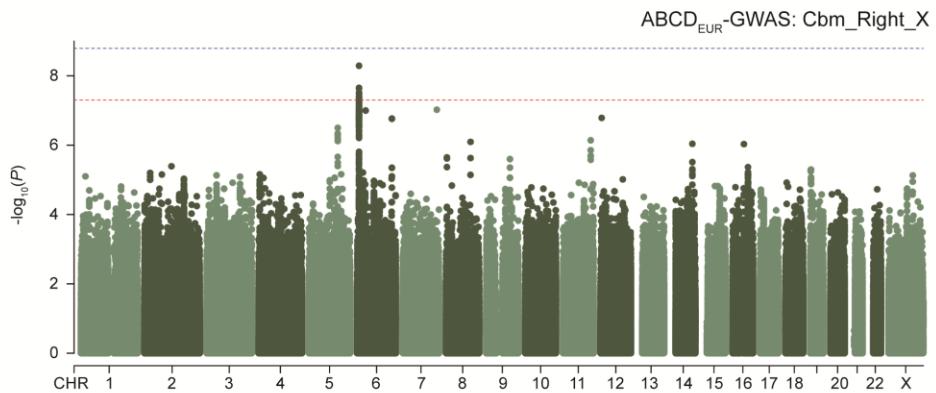


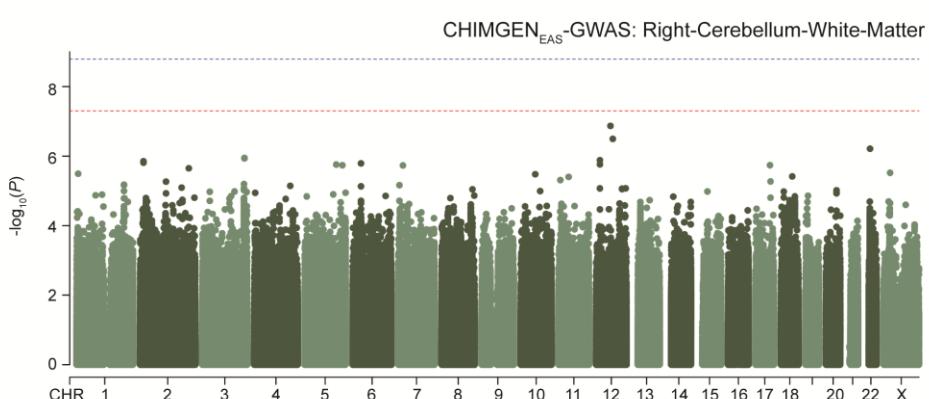
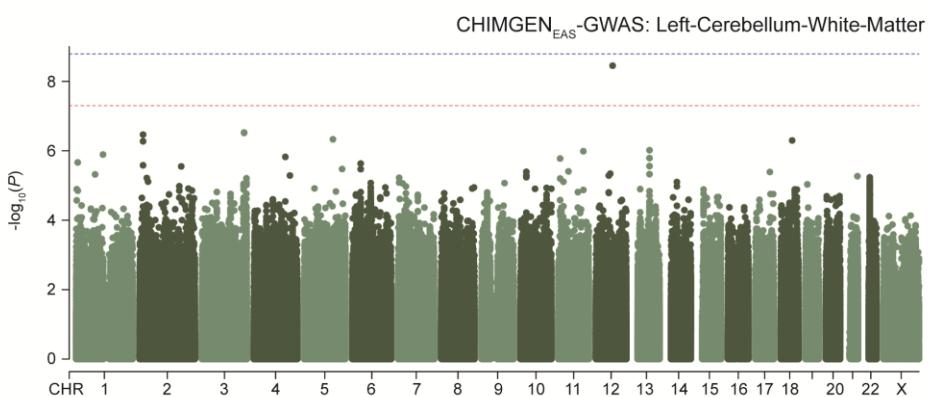
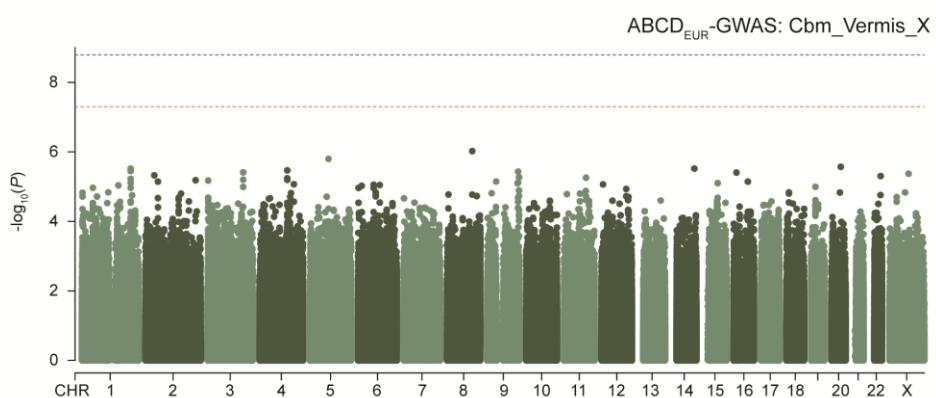
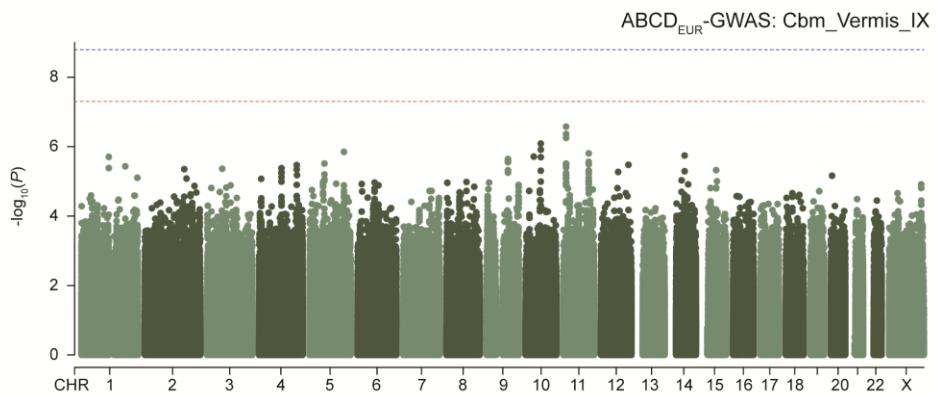


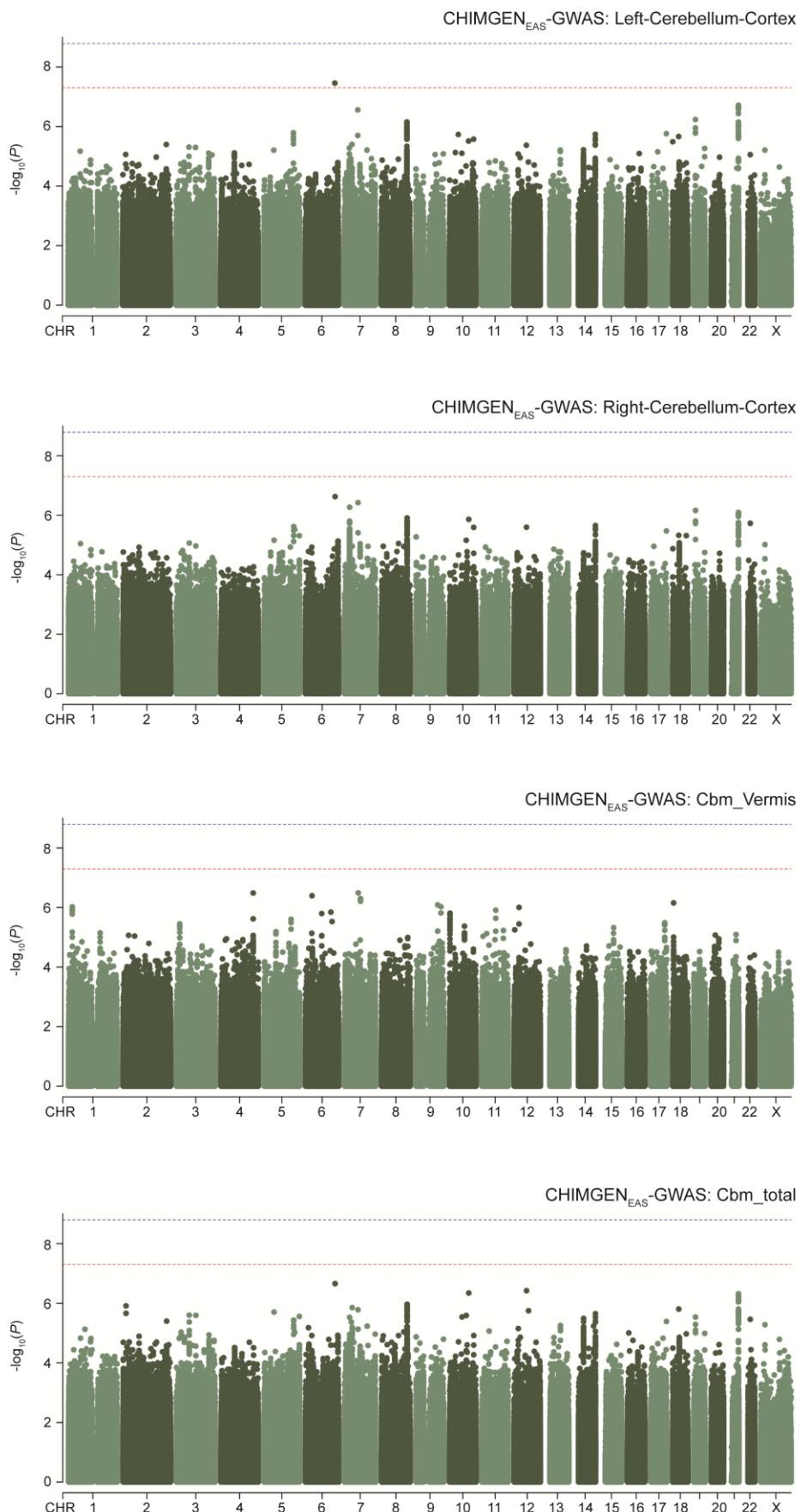


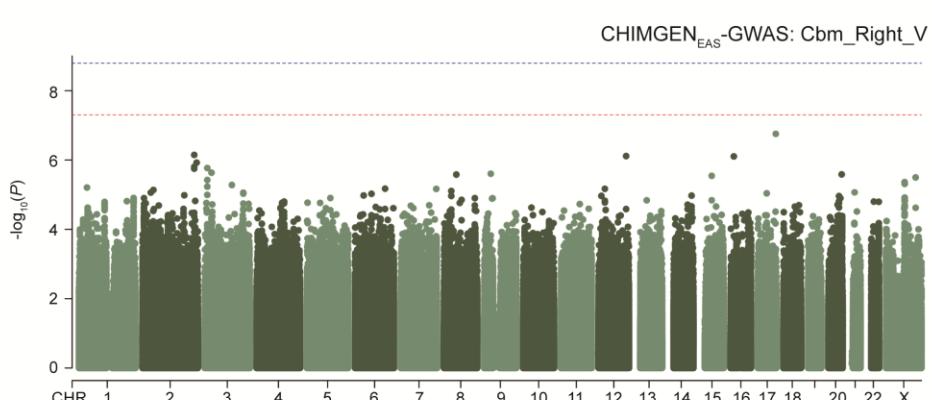
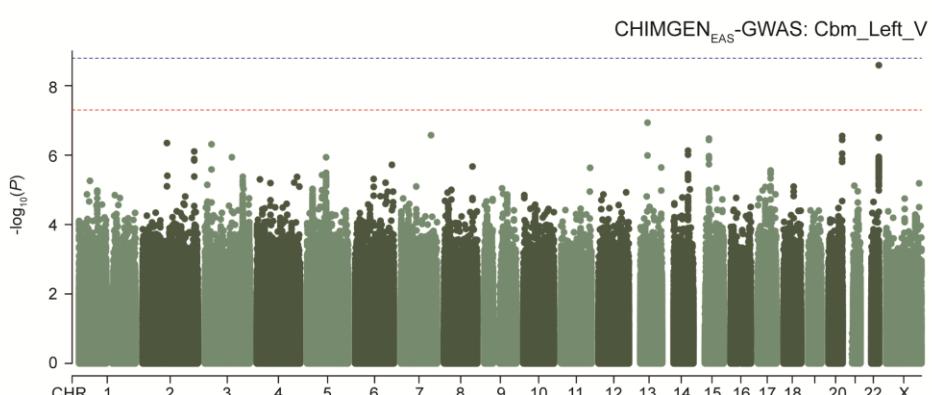
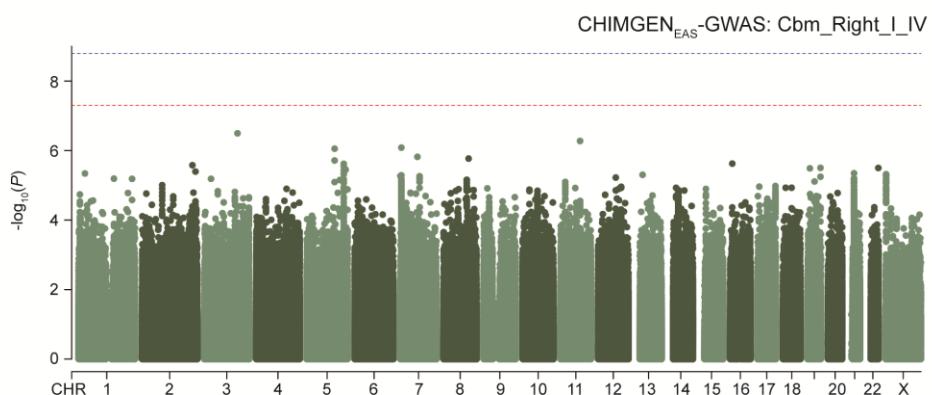
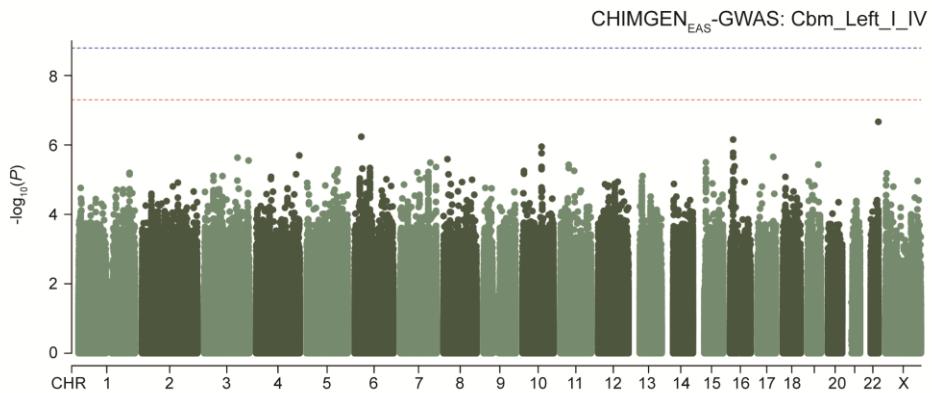


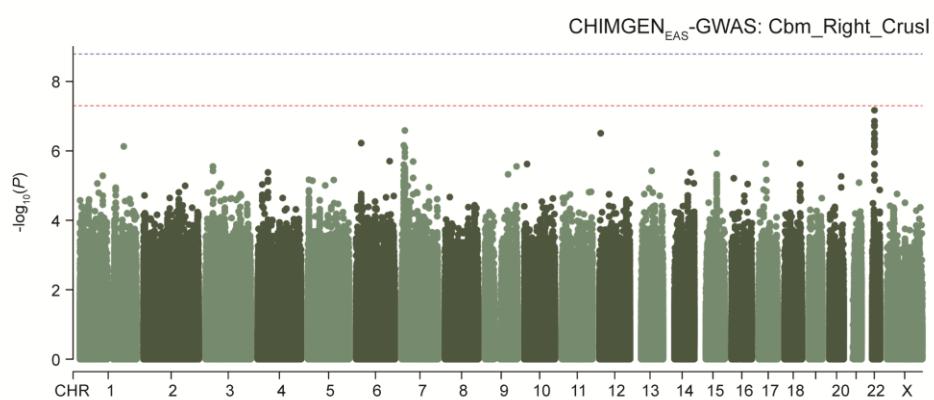
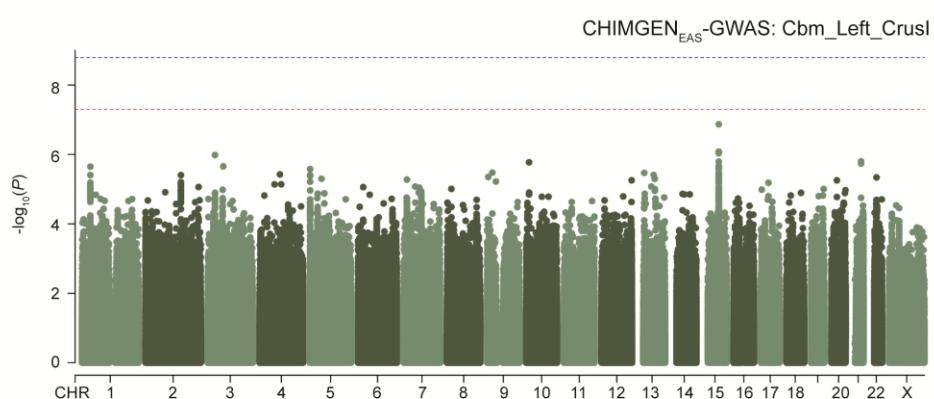
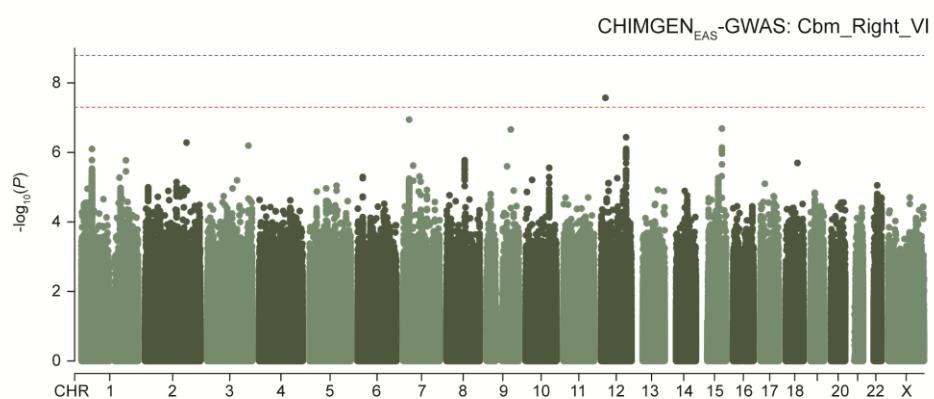
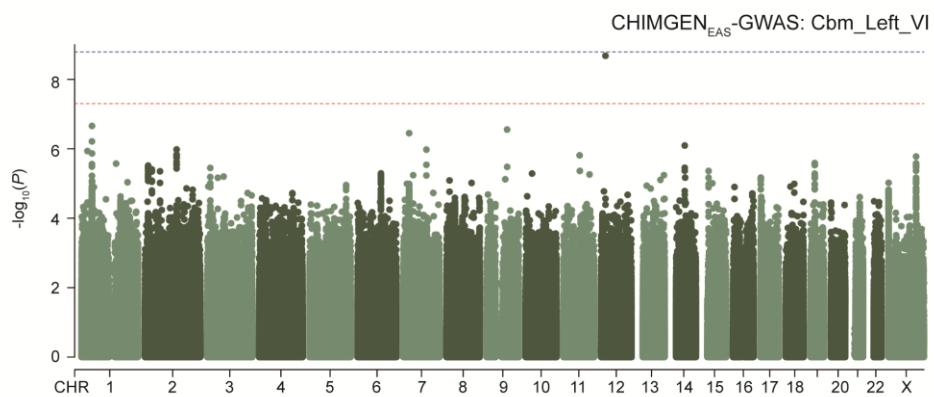


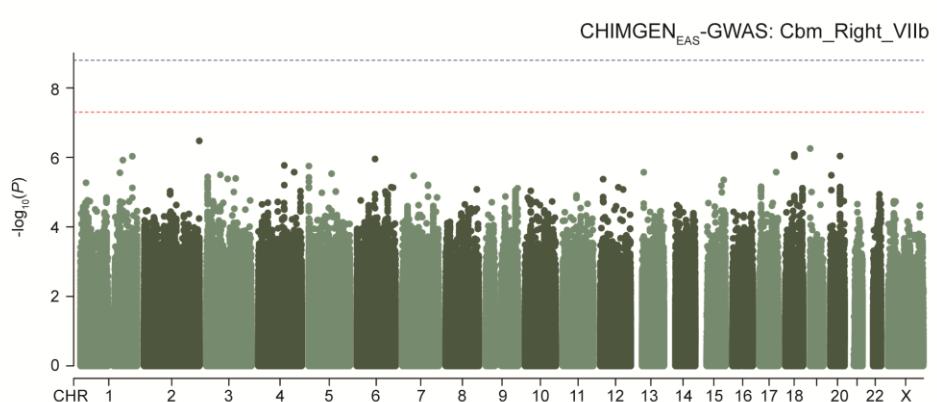
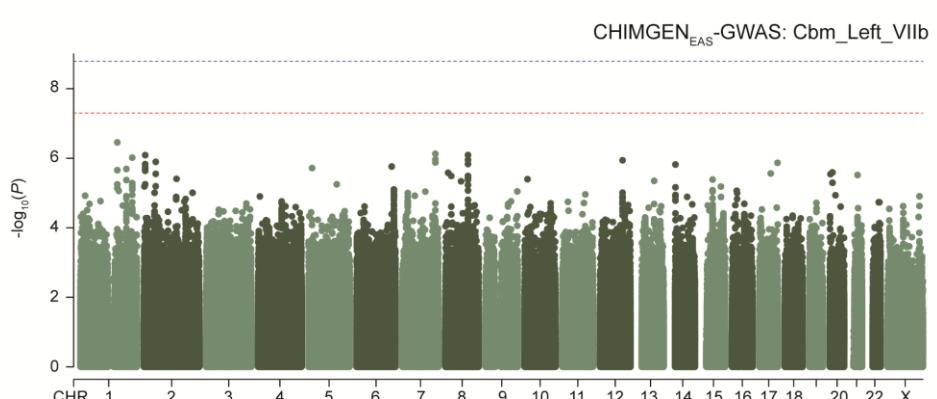
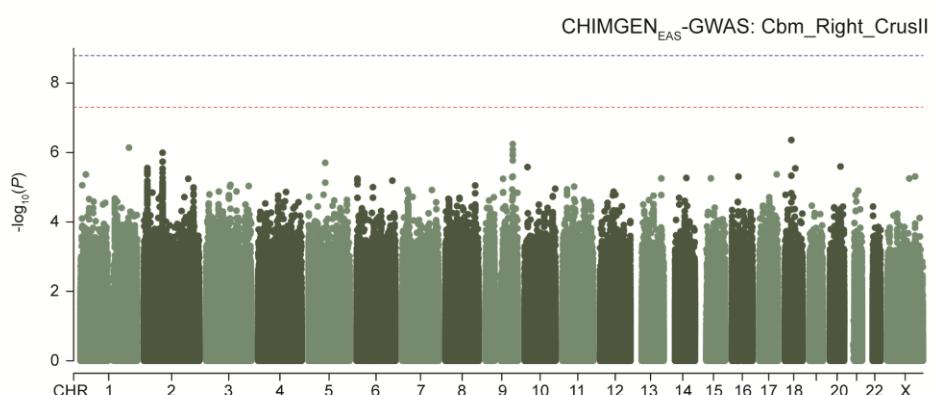
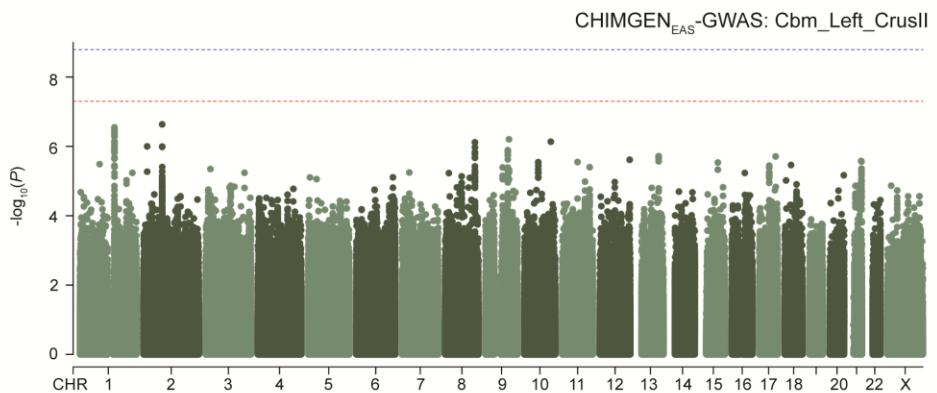


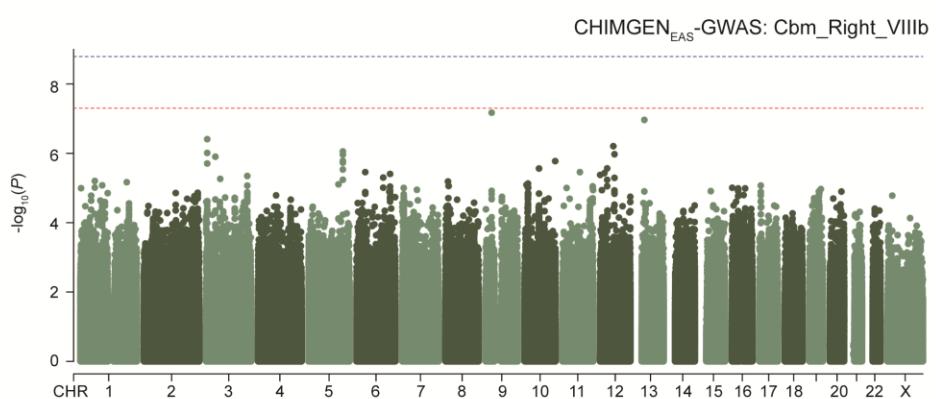
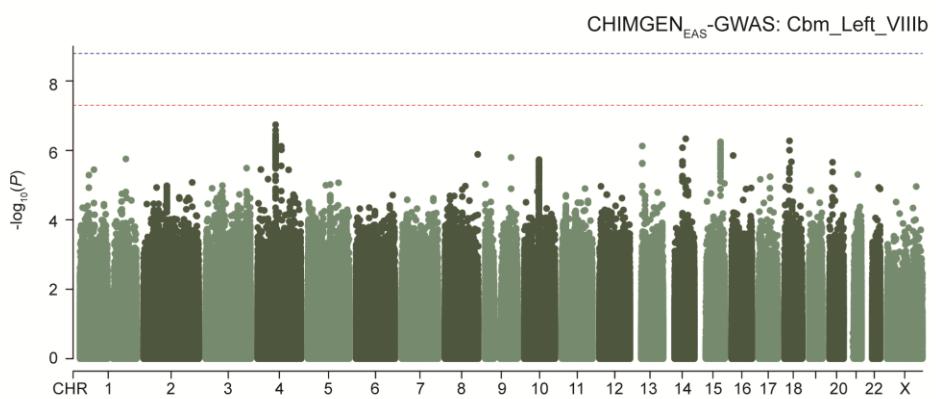
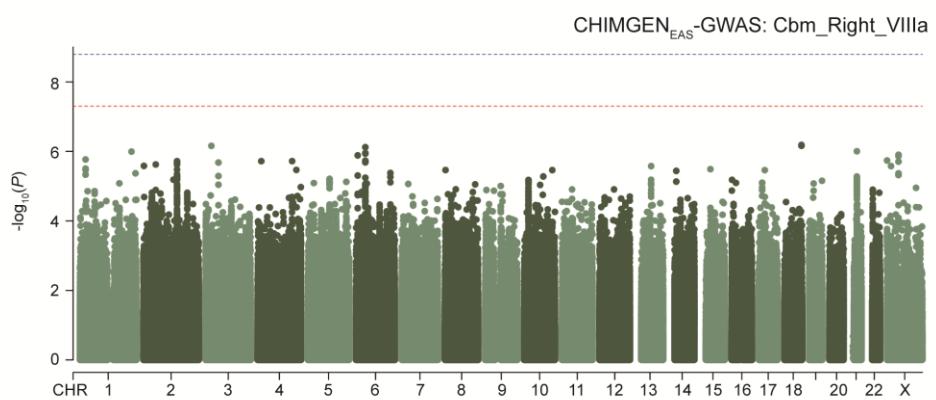
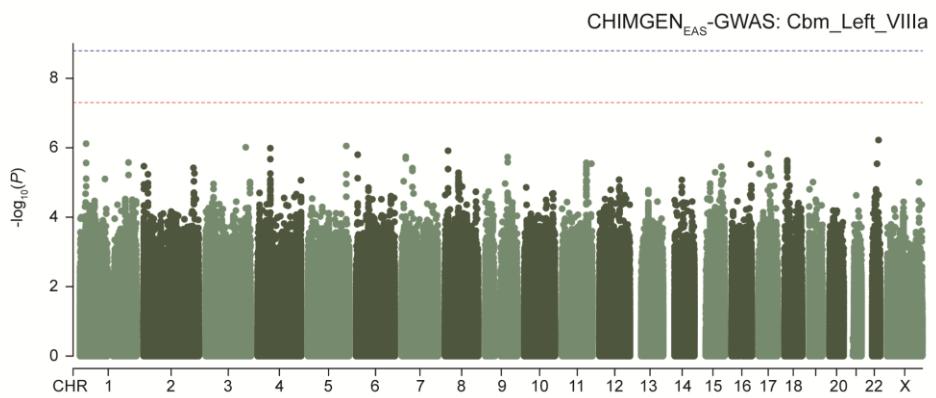


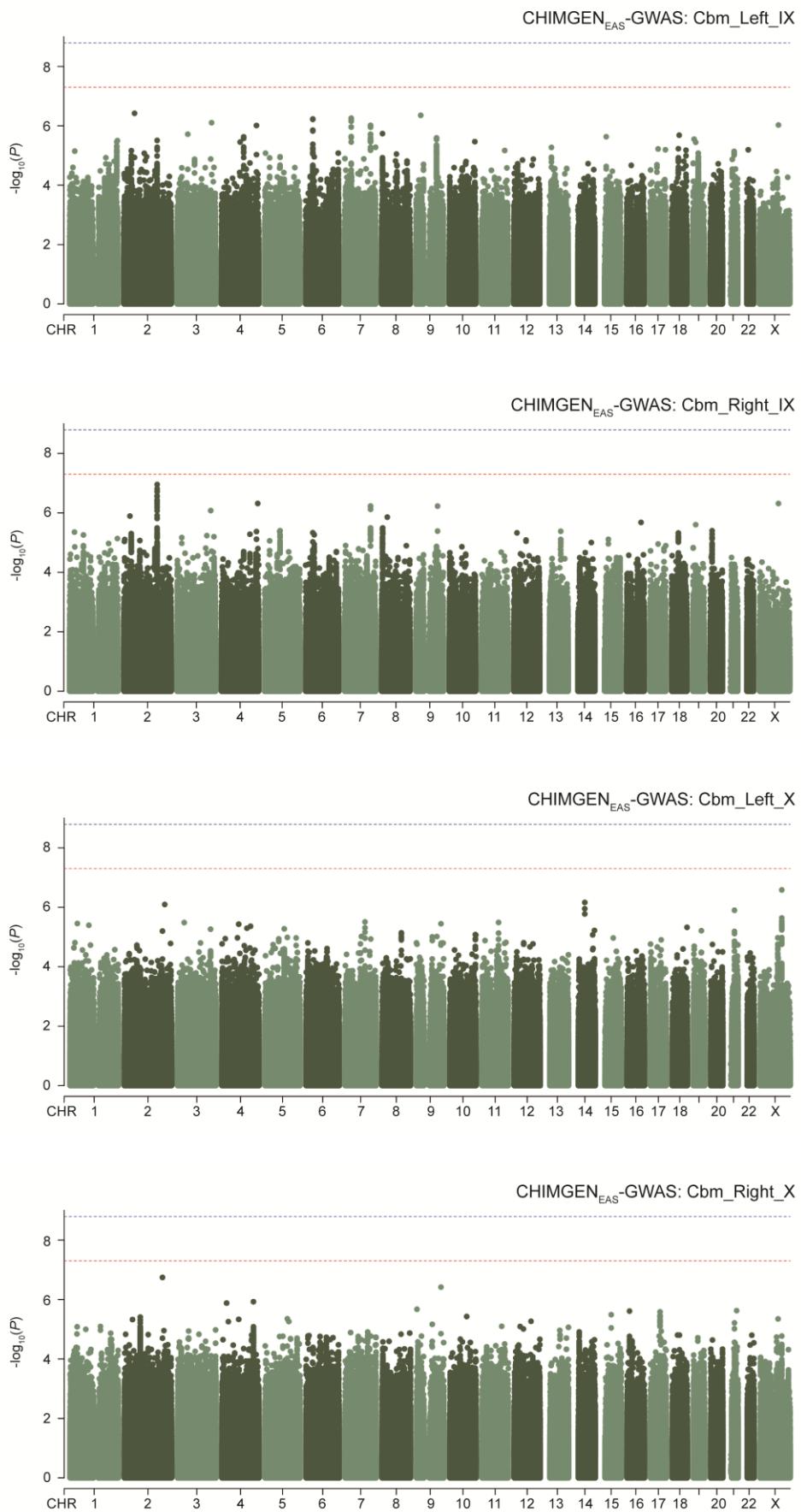


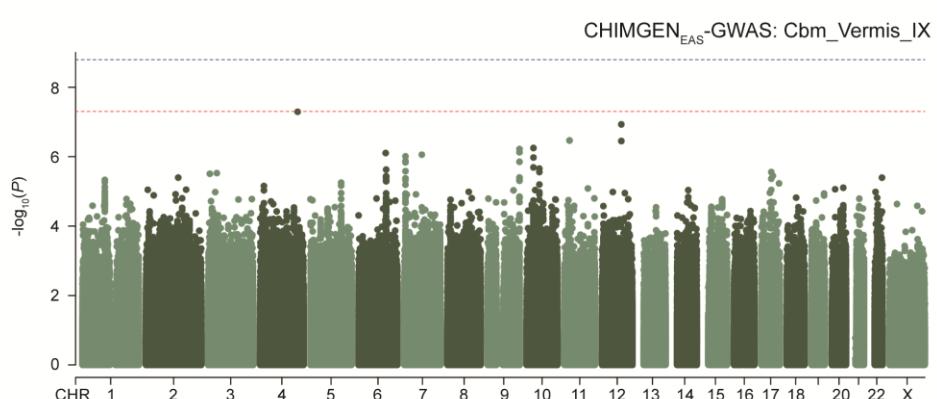
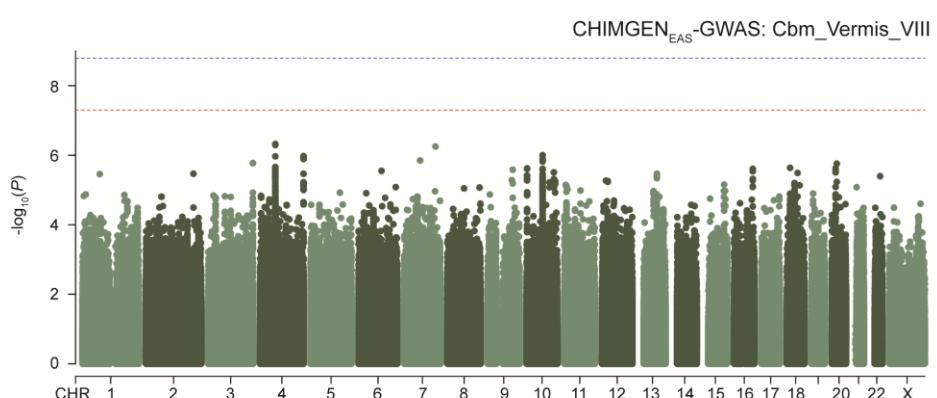
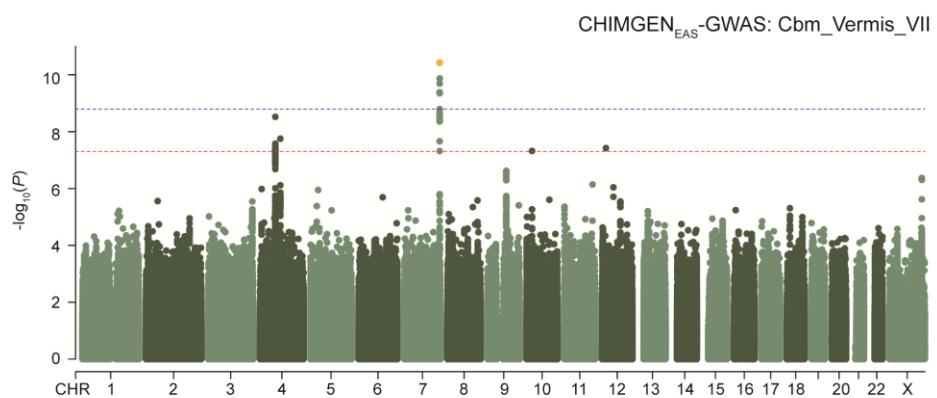
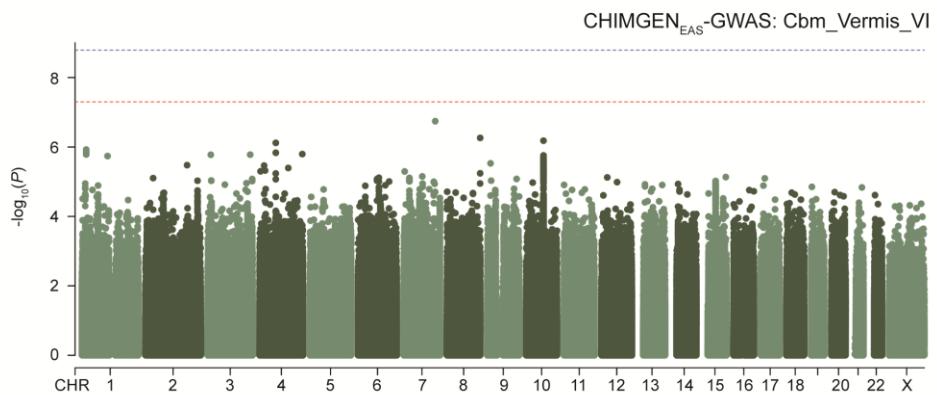


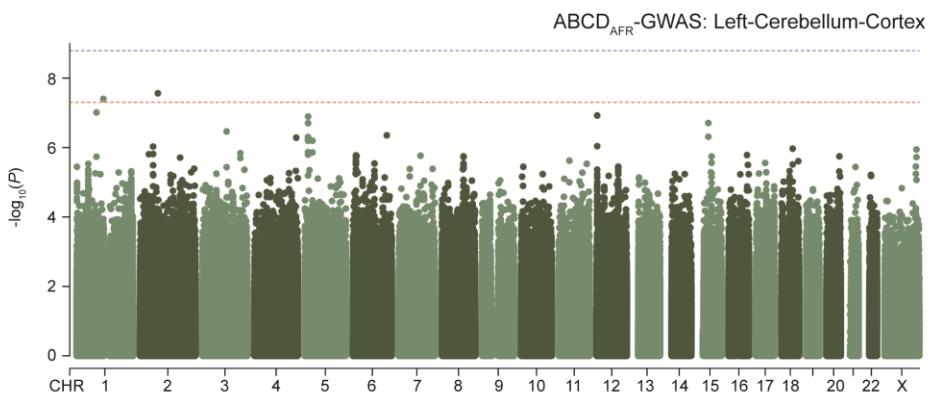
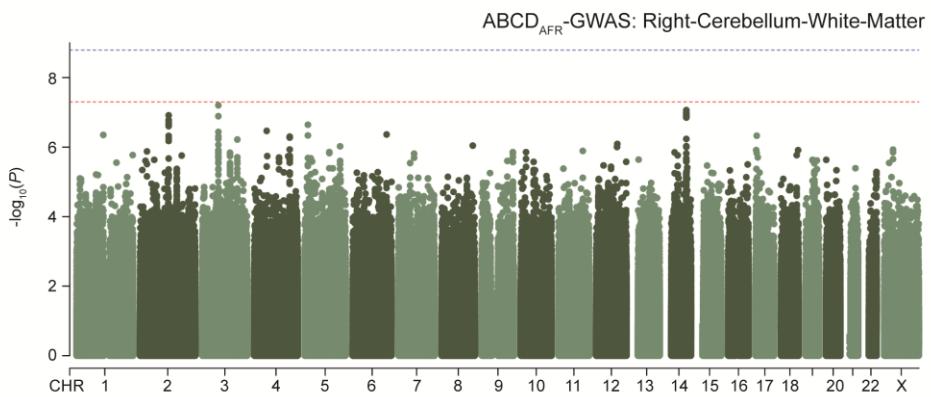
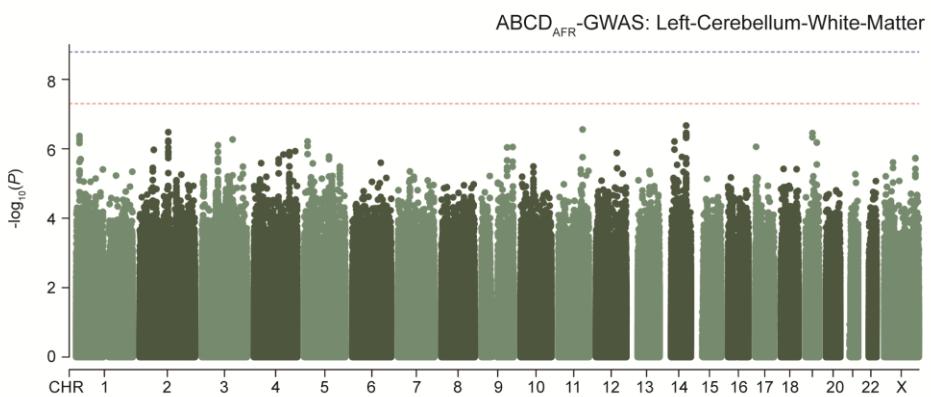
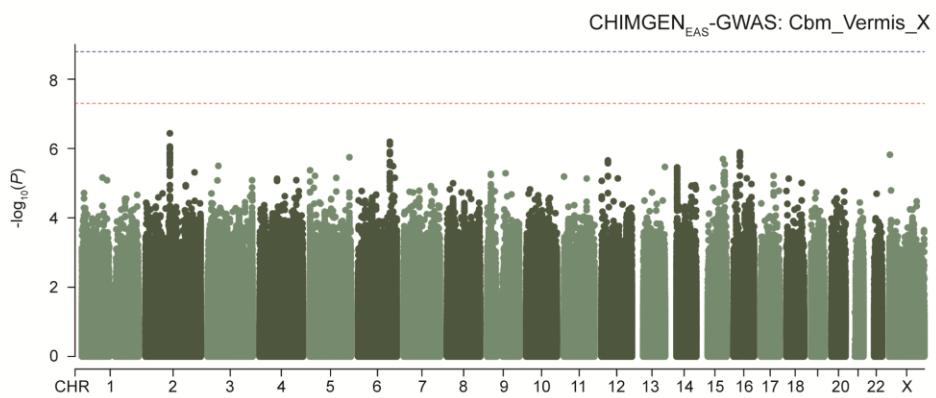




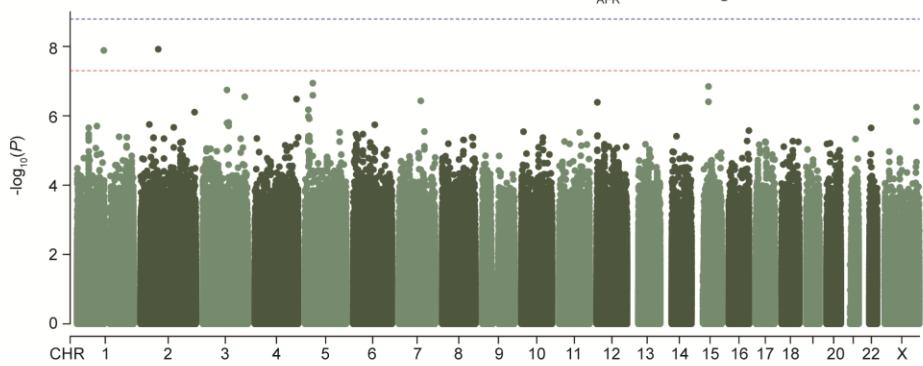




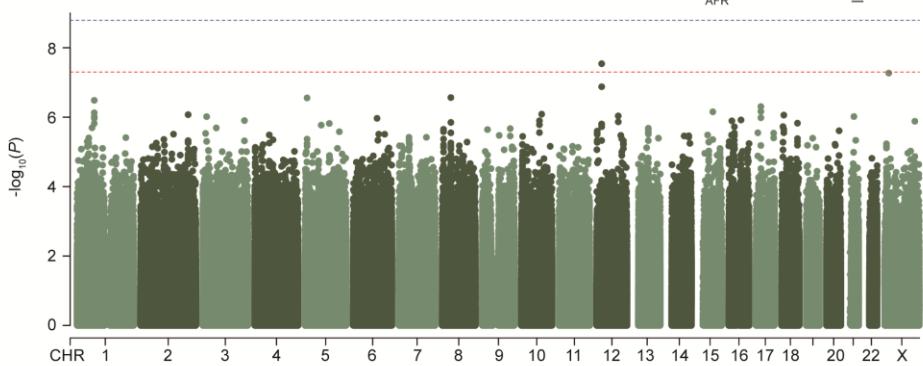




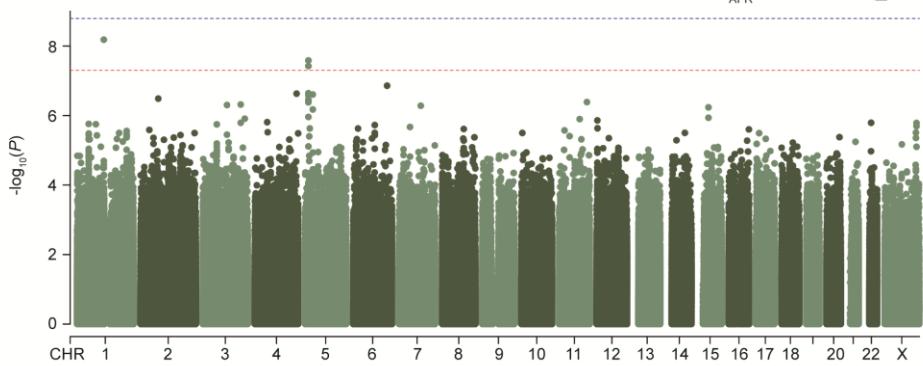
ABCD_{AFR}-GWAS: Right-Cerebellum-Cortex



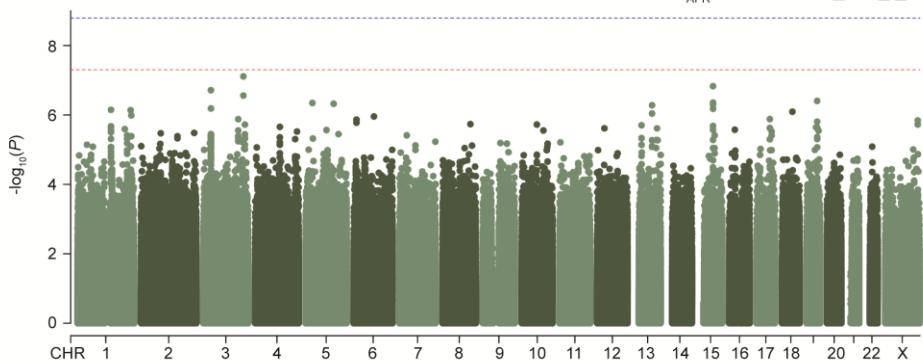
ABCD_{AFR}-GWAS: Cbm_Vermis

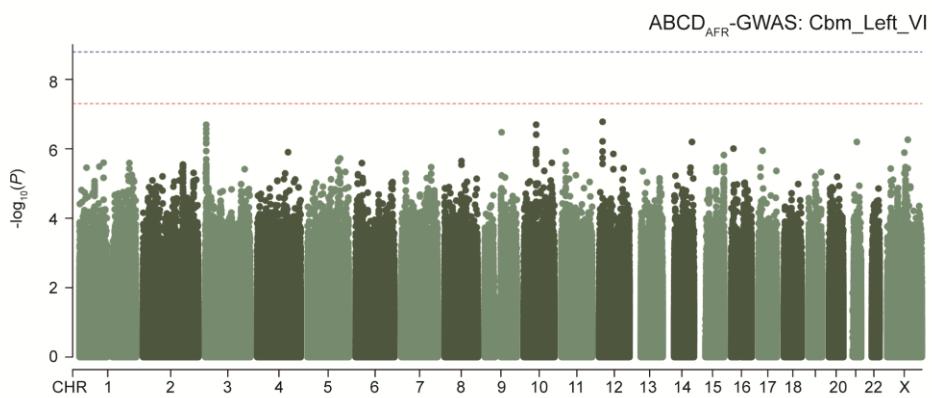
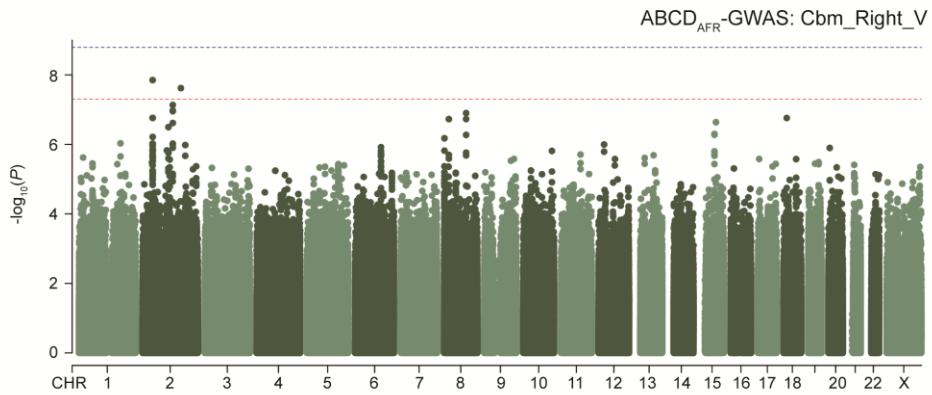
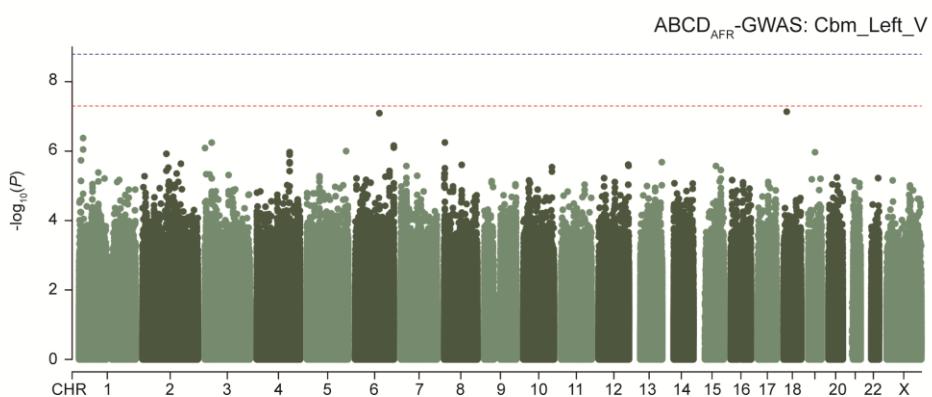
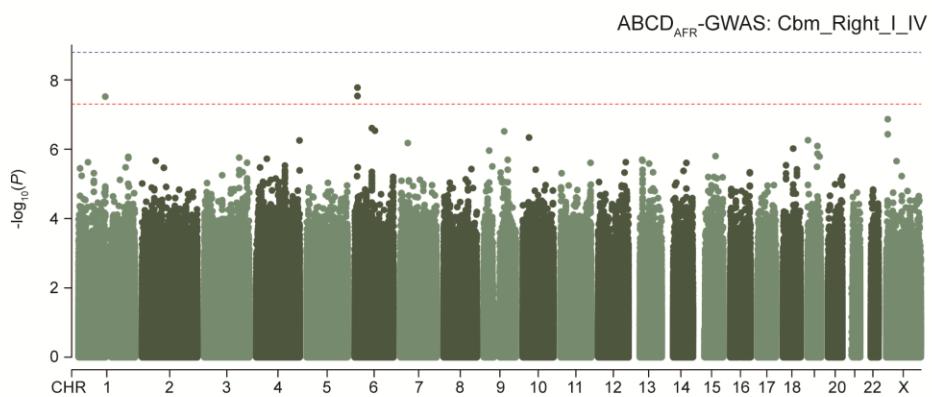


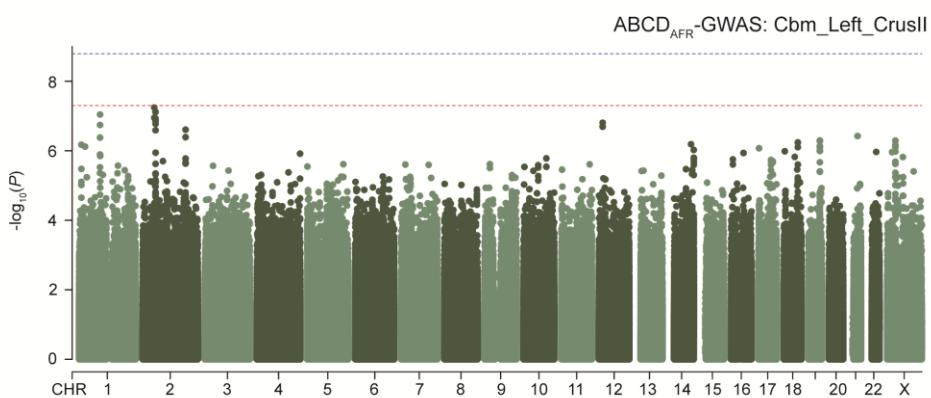
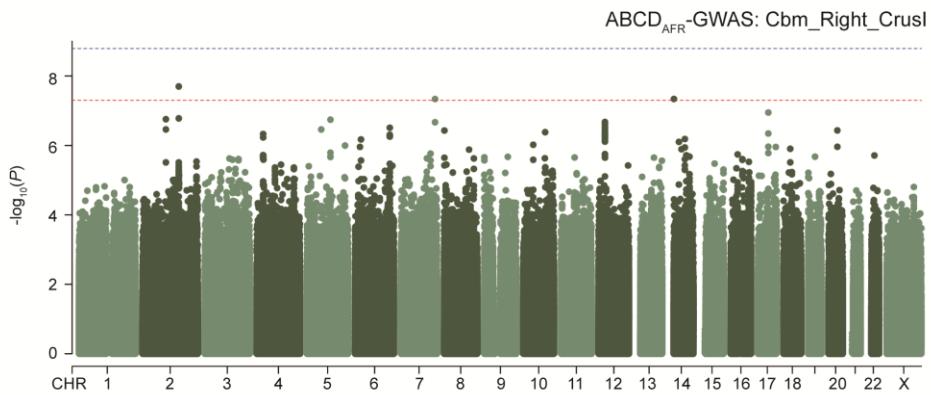
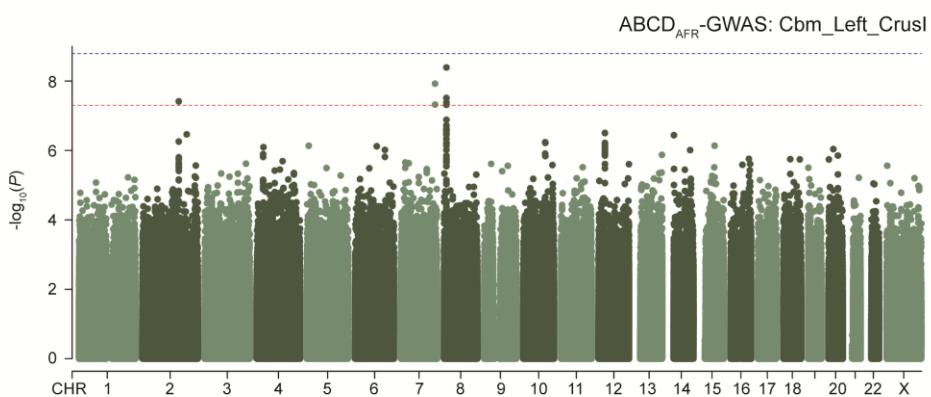
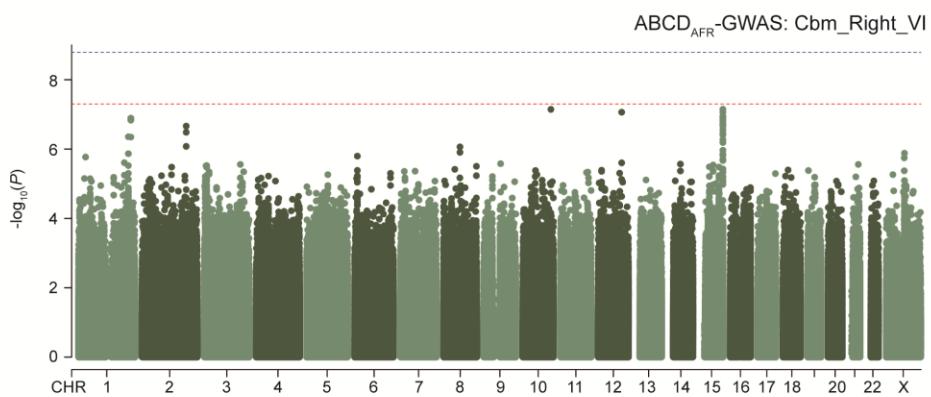
ABCD_{AFR}-GWAS: Cbm_total

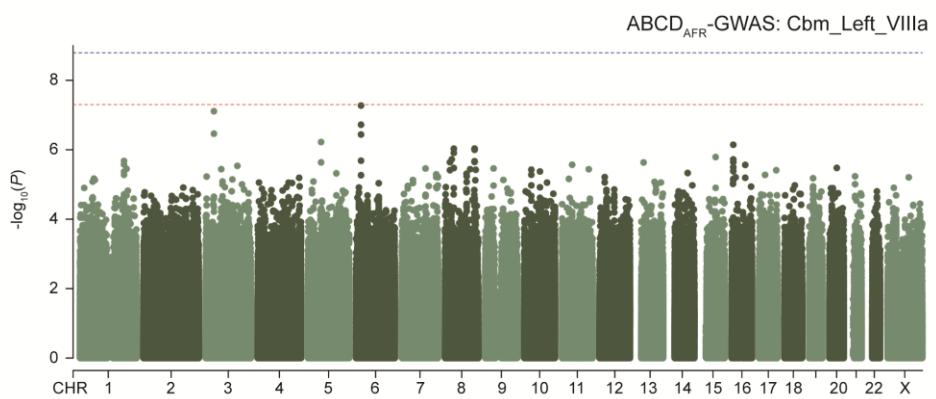
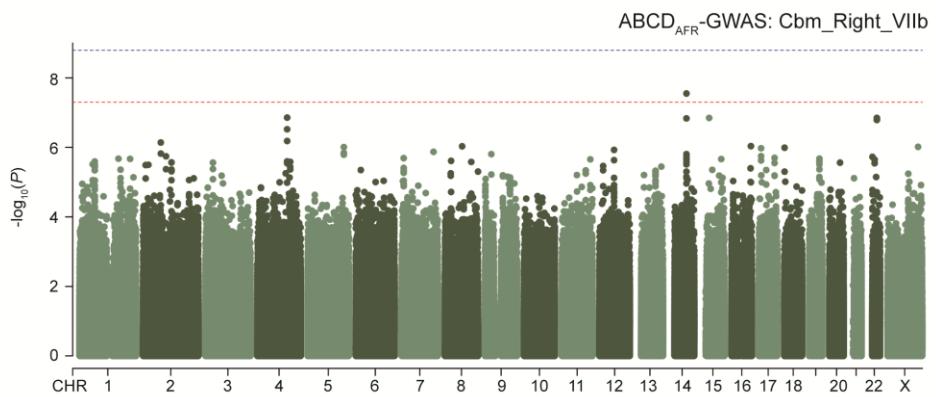
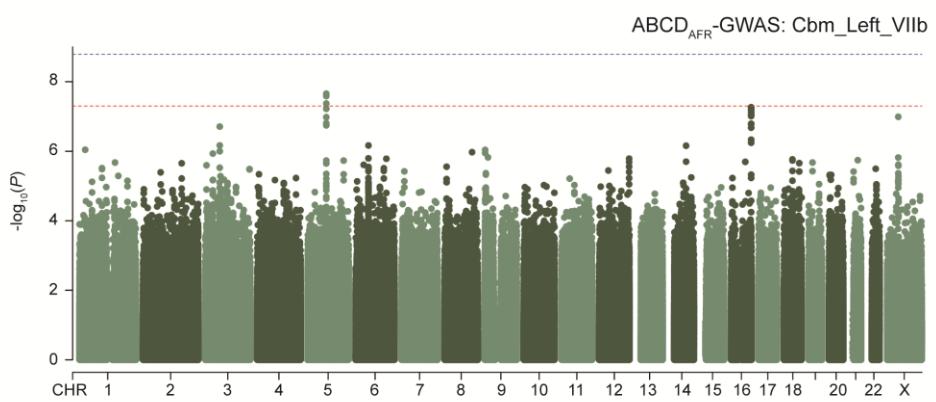
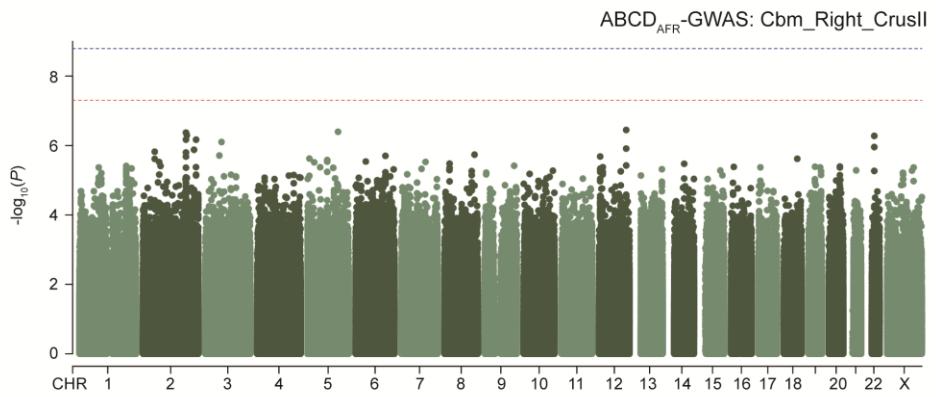


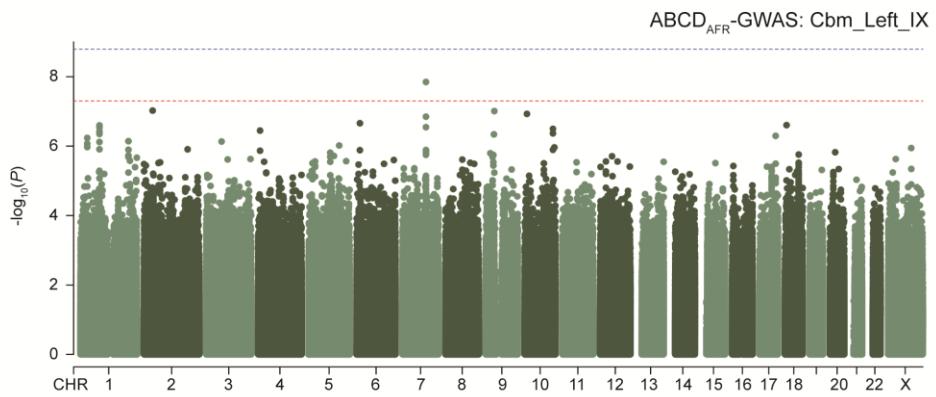
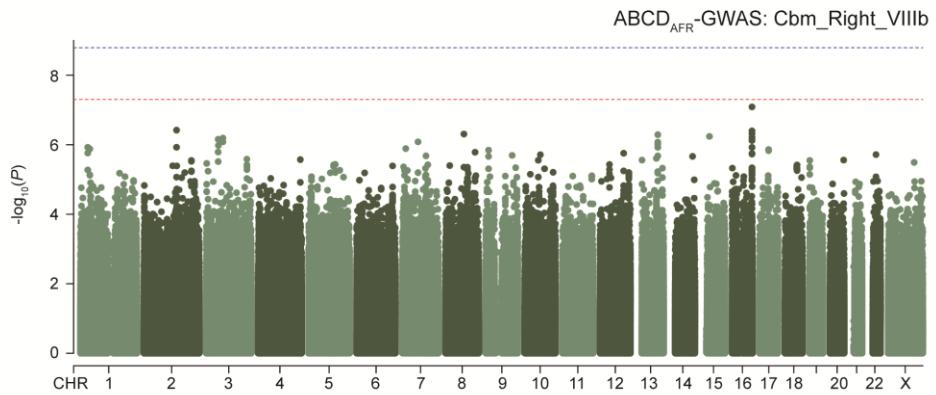
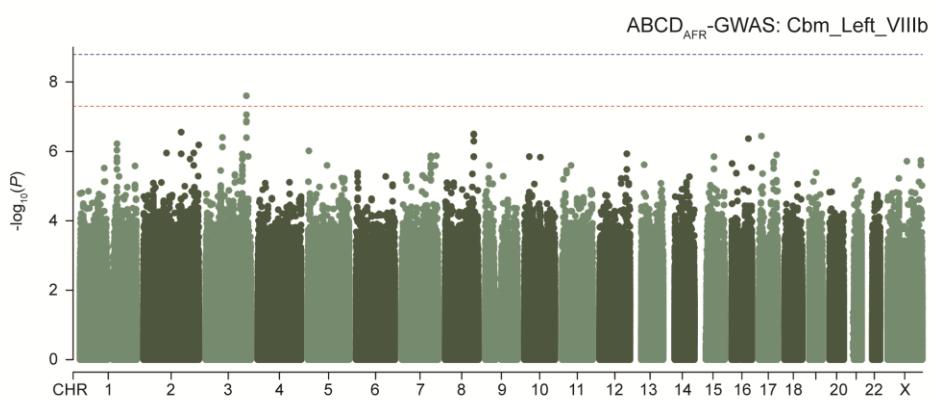
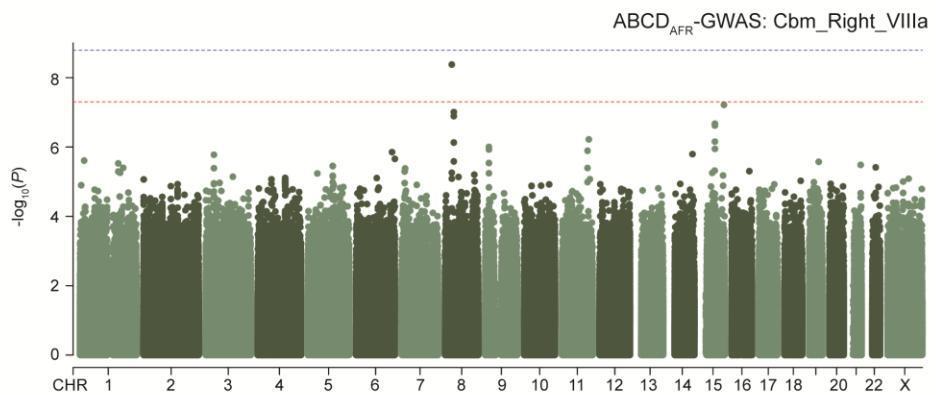
ABCD_{AFR}-GWAS: Cbm_Left_I_IV

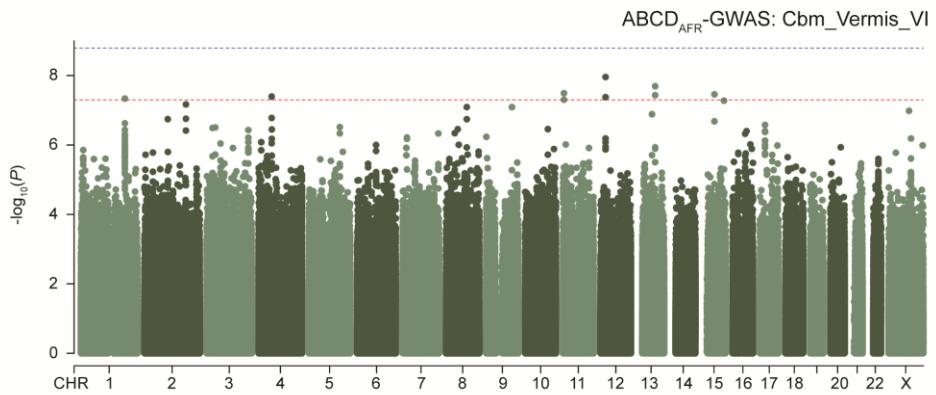
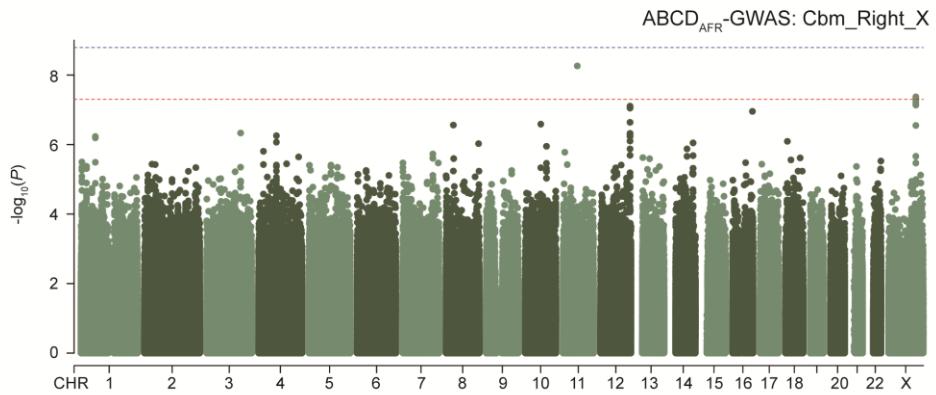
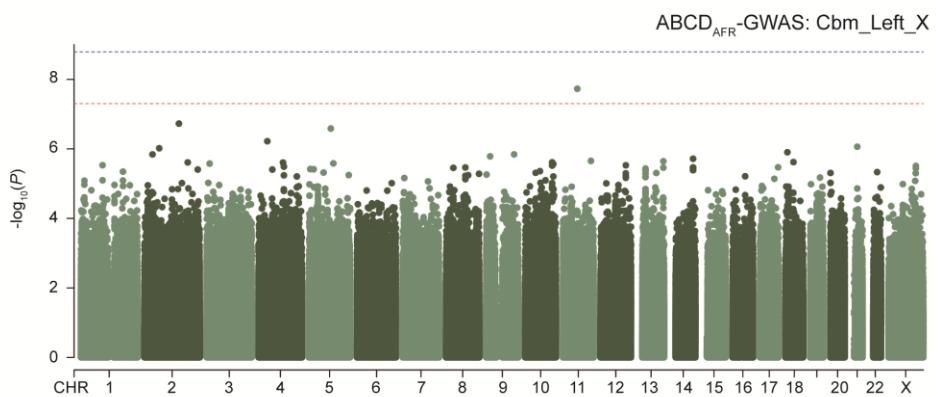
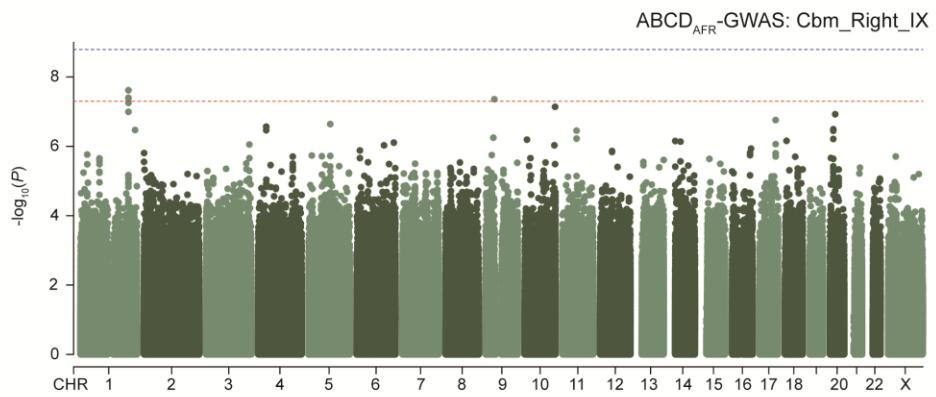




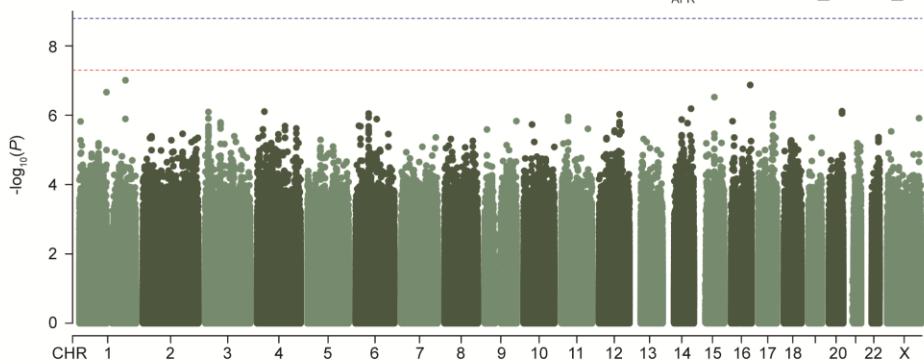




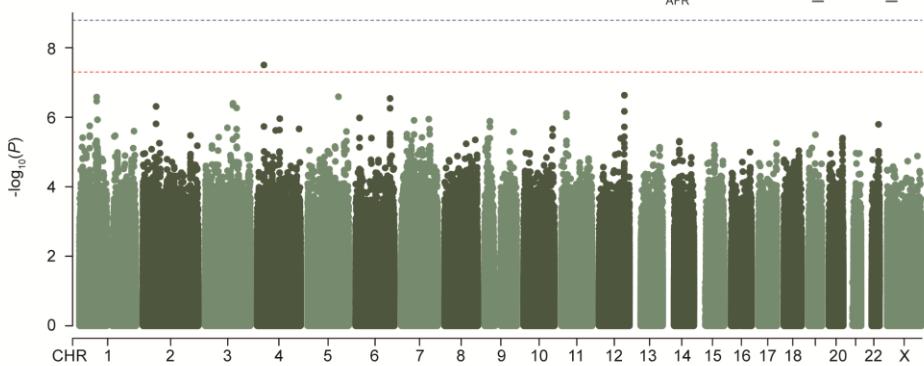




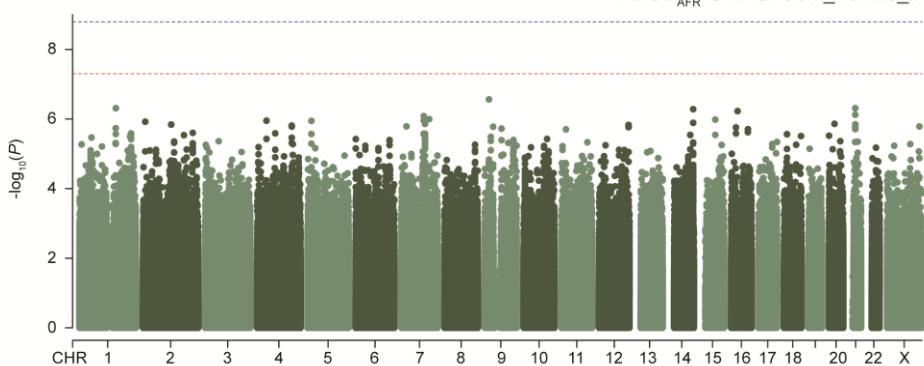
ABCD_{AFR}-GWAS: Cbm_Vermis_VII



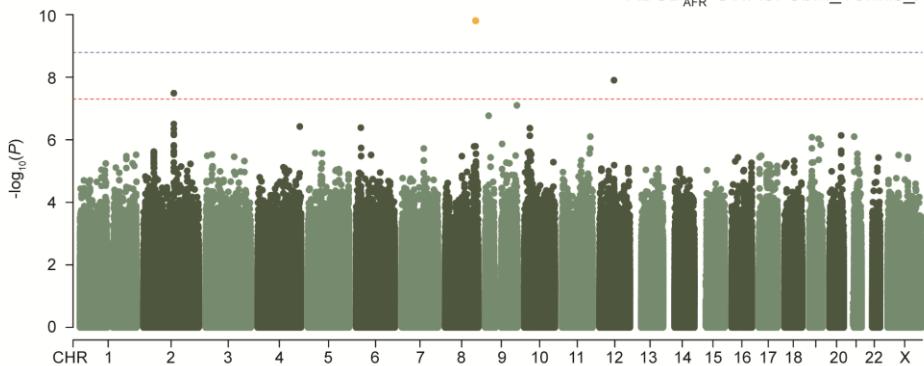
ABCD_{AFR}-GWAS: Cbm_Vermis_VIII



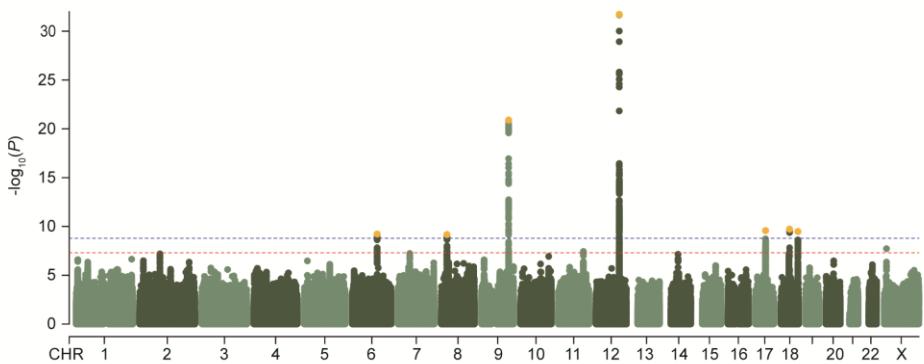
ABCD_{AFR}-GWAS: Cbm_Vermis_IX



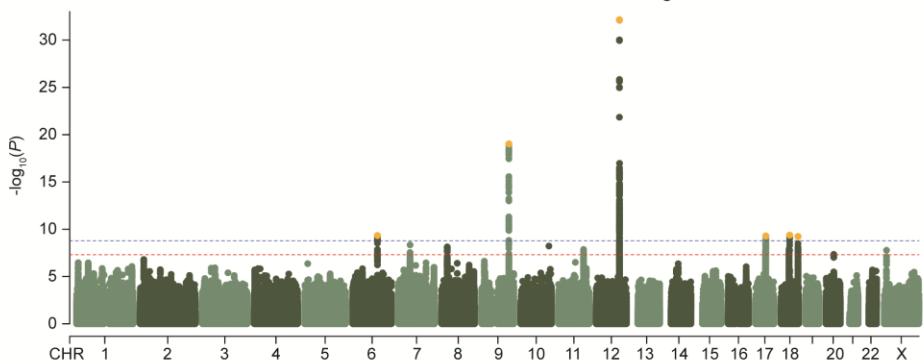
ABCD_{AFR}-GWAS: Cbm_Vermis_X



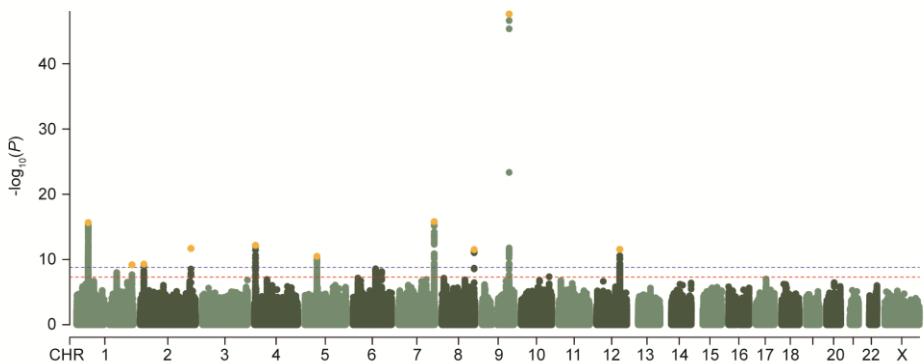
EUR-GWAS: Left-Cerebellum-White-Matter



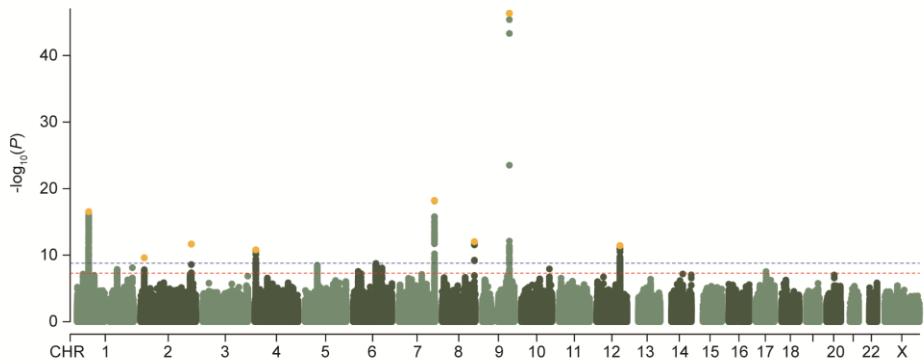
EUR-GWAS: Right-Cerebellum-White-Matter



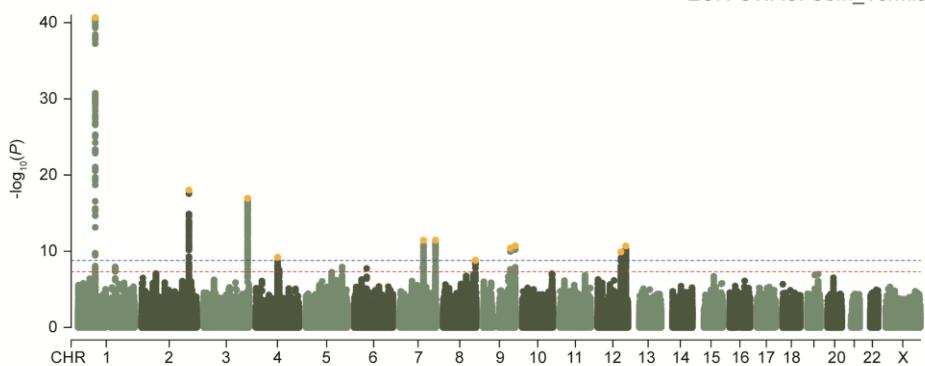
EUR-GWAS: Left-Cerebellum-Cortex



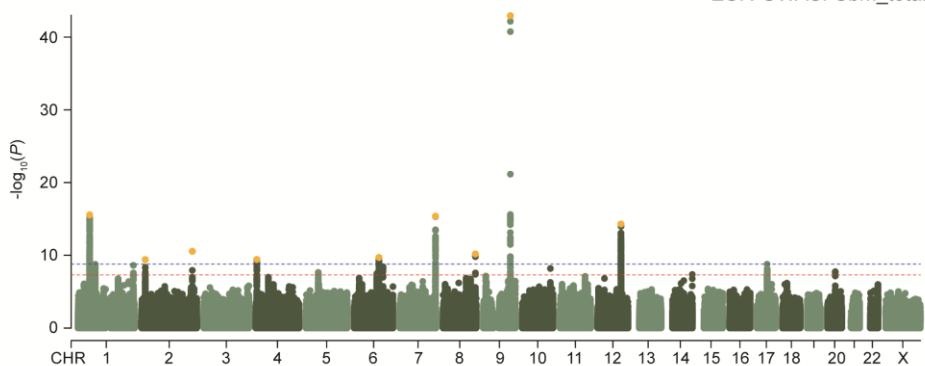
EUR-GWAS: Right-Cerebellum-Cortex



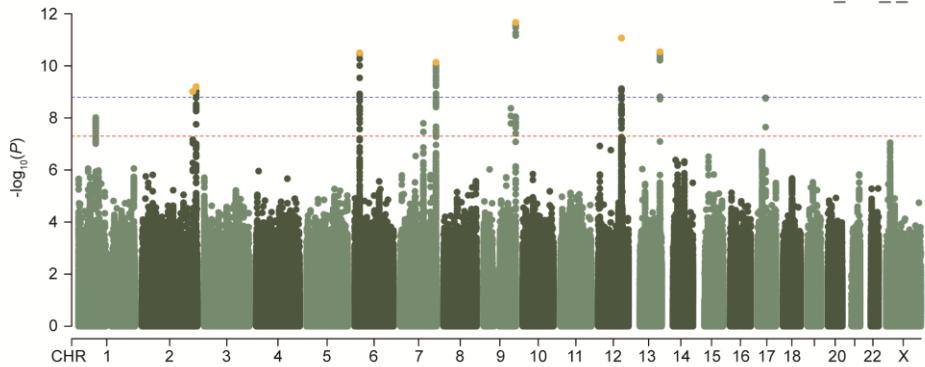
EUR-GWAS: Cbm_Vermis



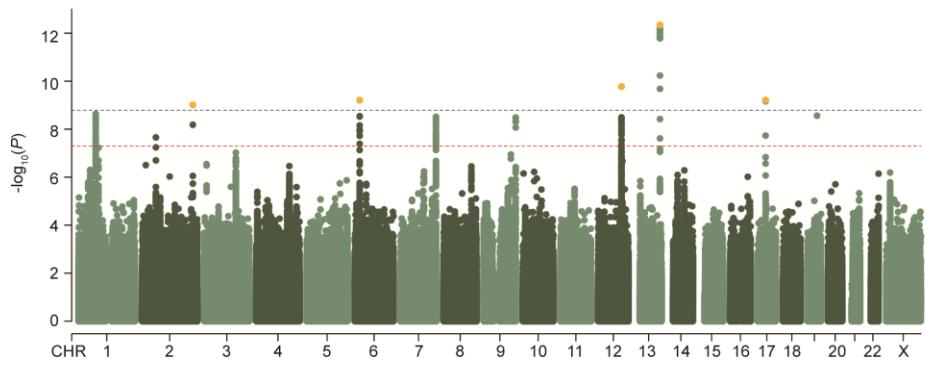
EUR-GWAS: Cbm_total

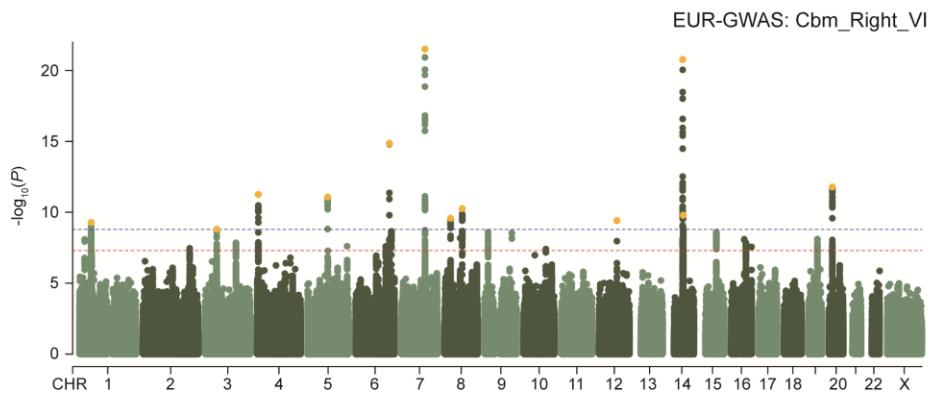
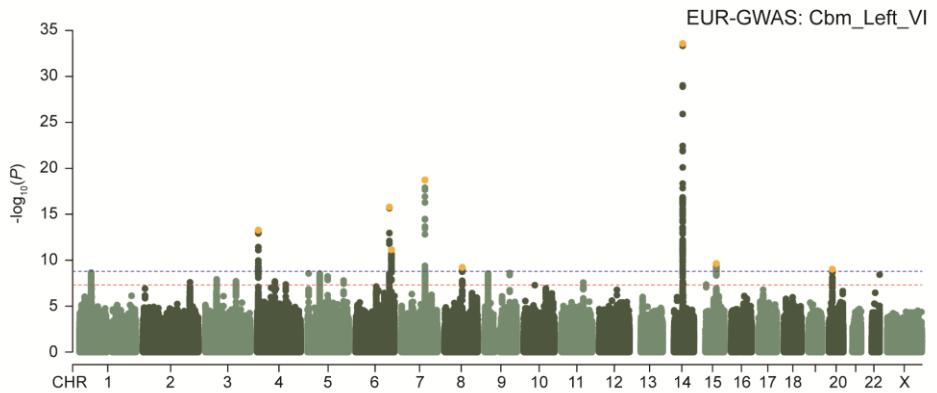
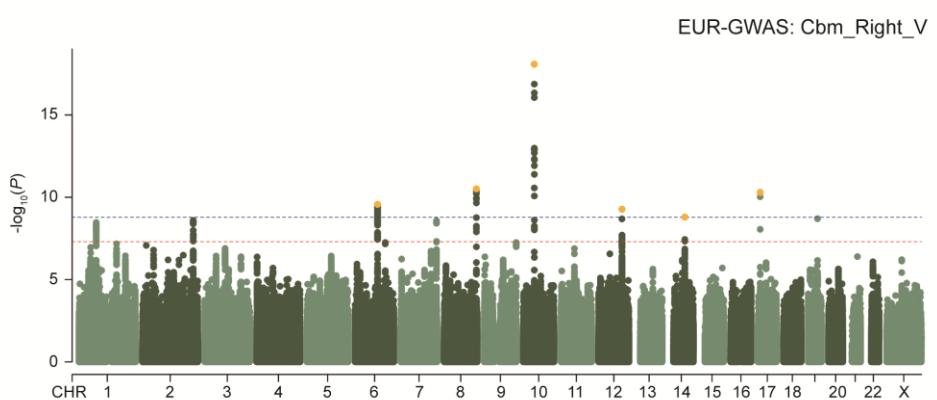
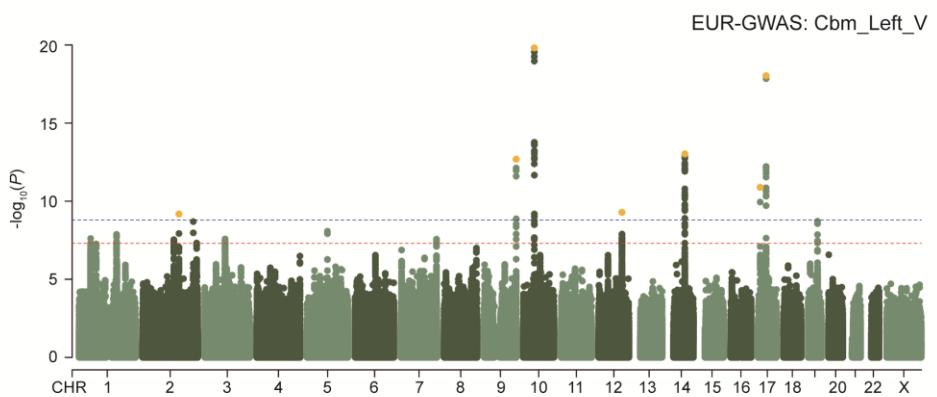


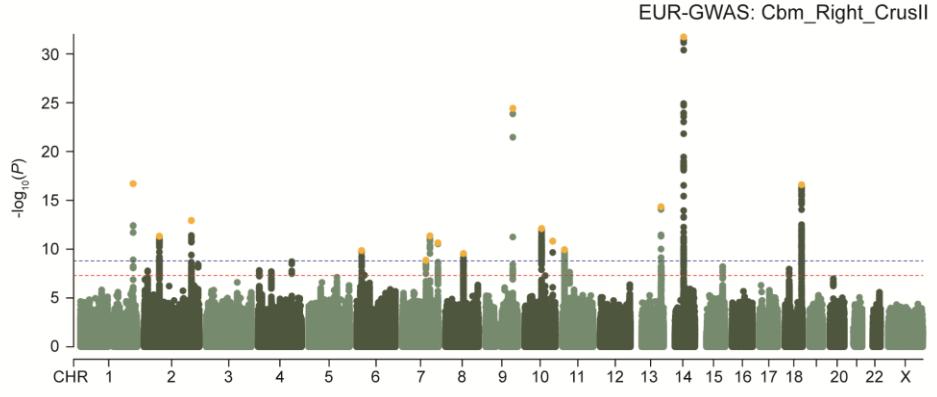
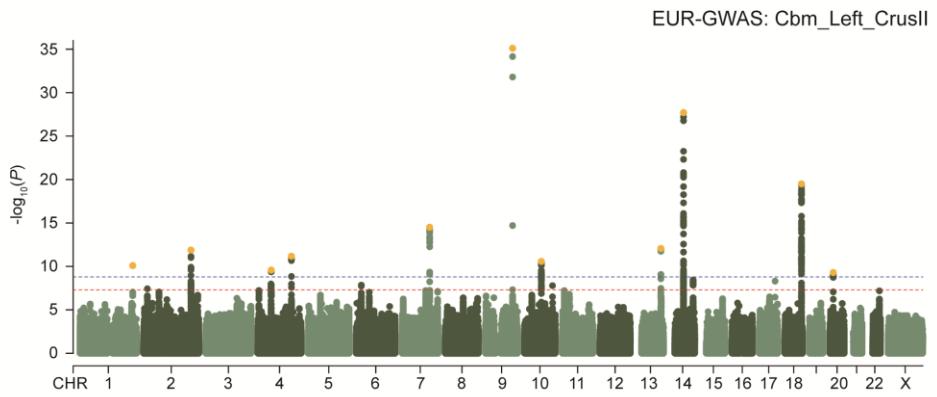
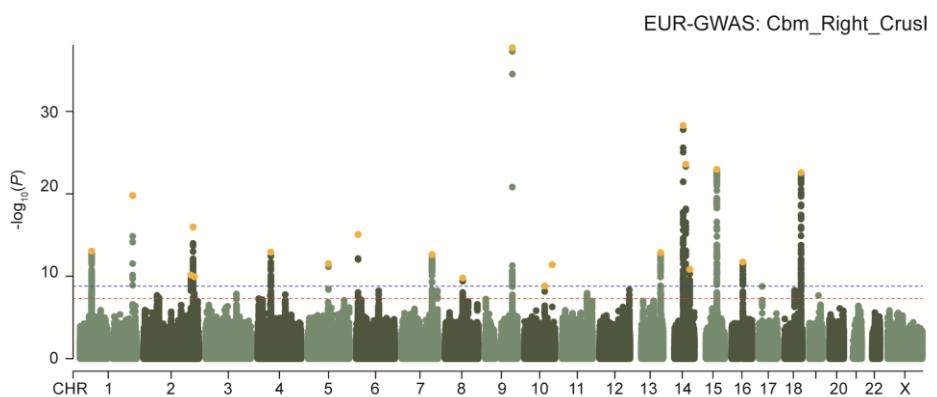
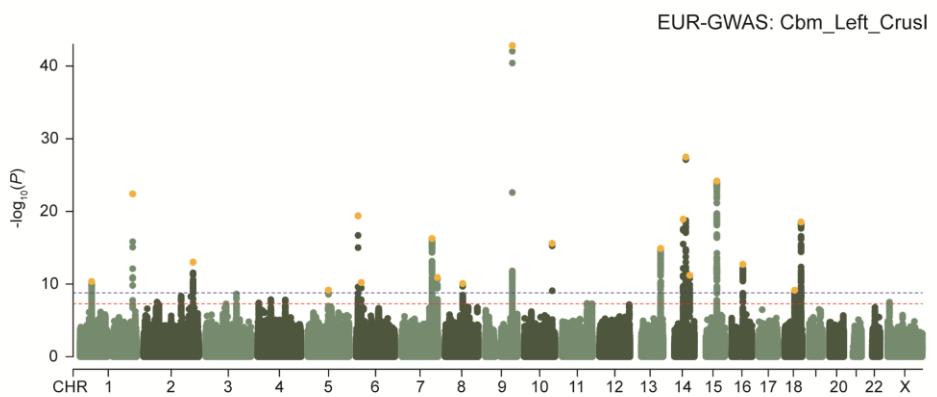
EUR-GWAS: Cbm_Left_I_IV

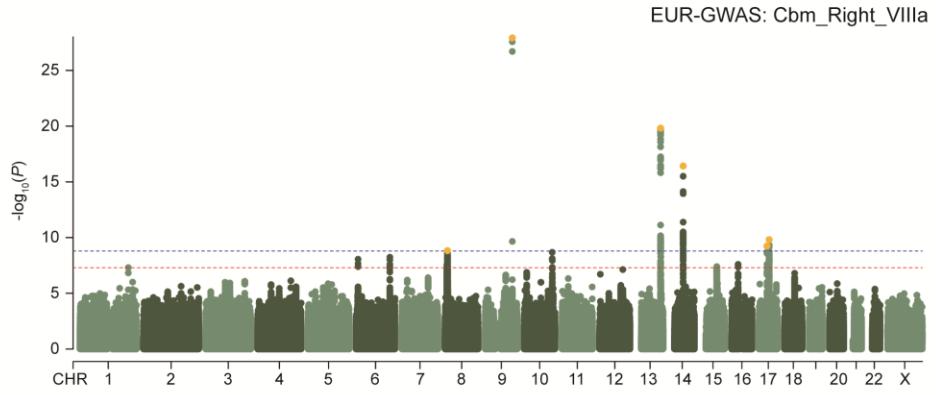
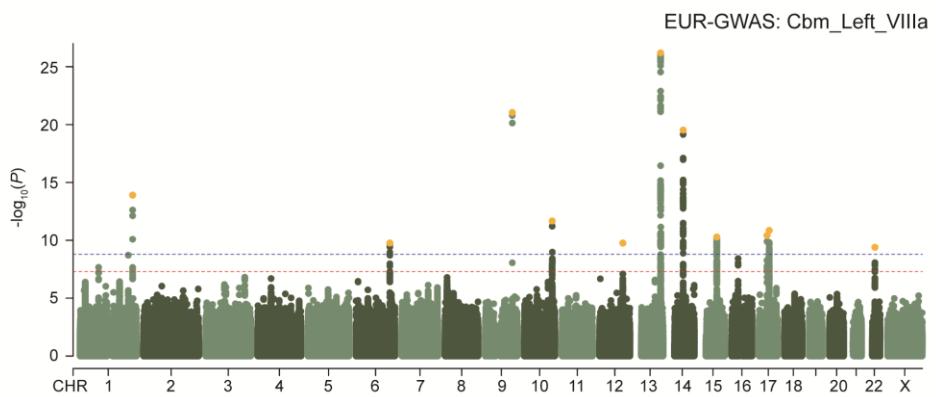
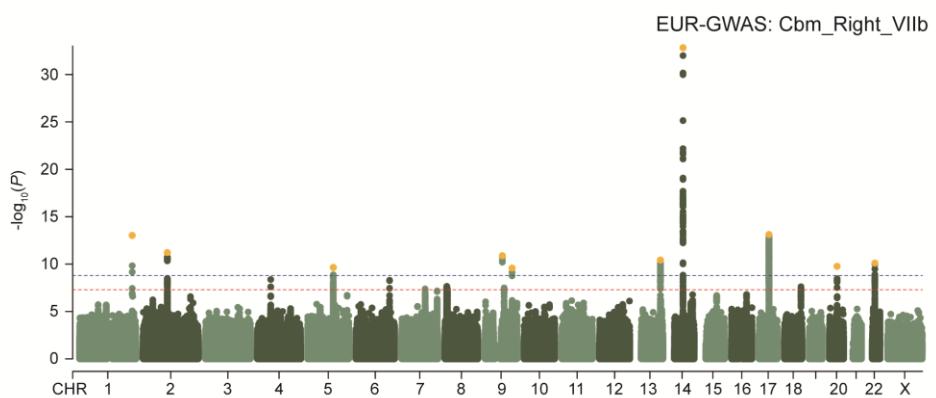
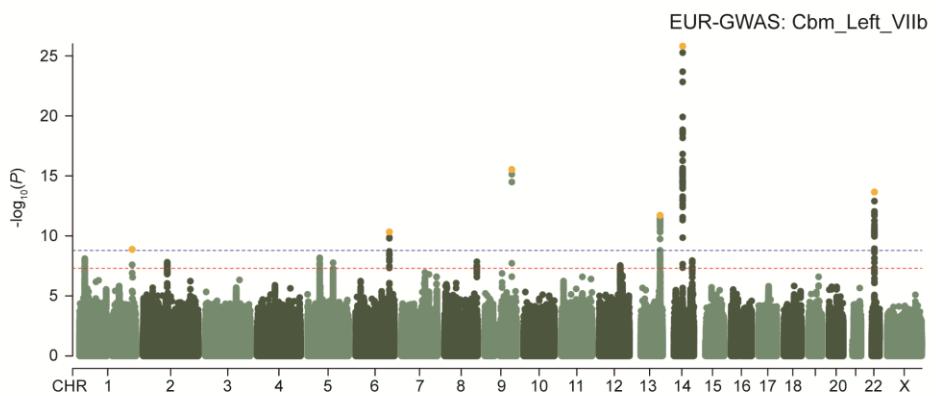


EUR-GWAS: Cbm_Right_I_IV

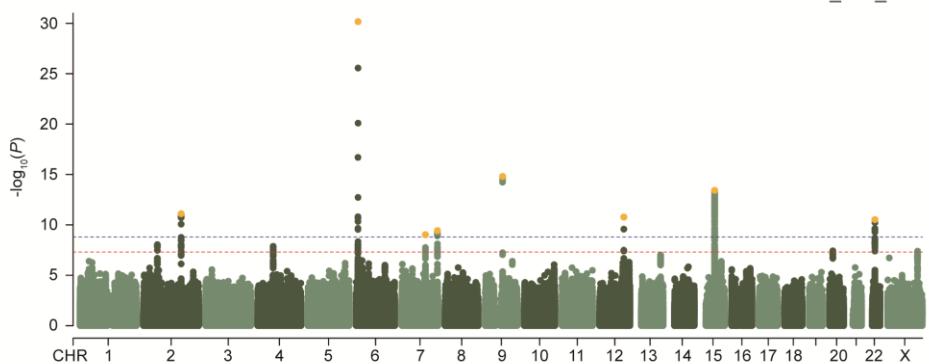




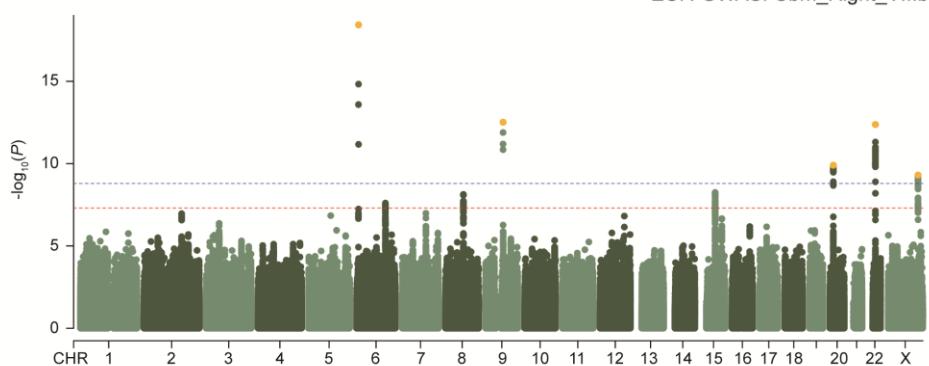




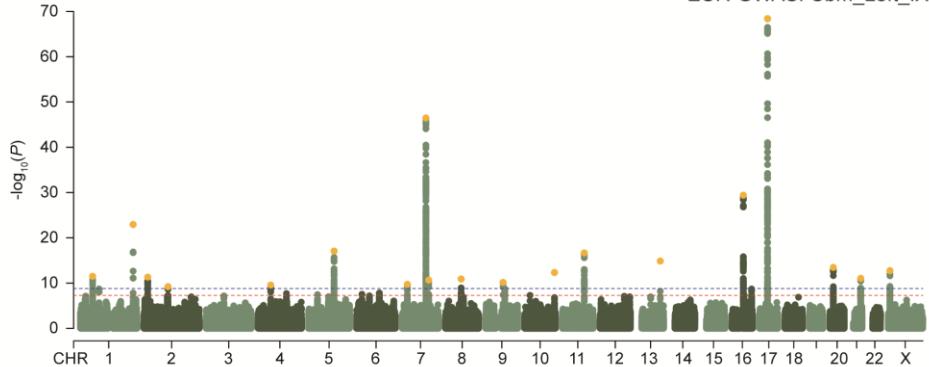
EUR-GWAS: Cbm_Left_VIIIb



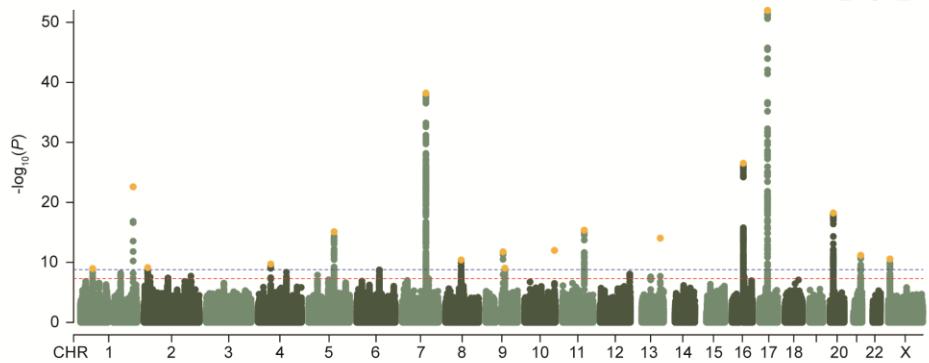
EUR-GWAS: Cbm_Right_VIIIb

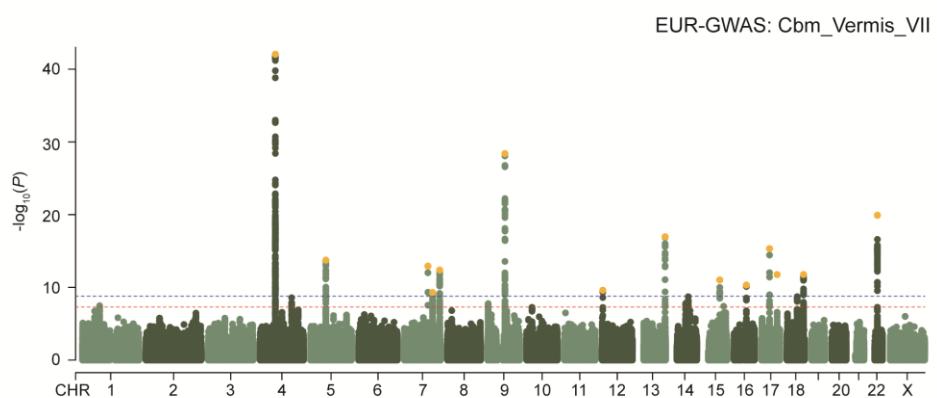
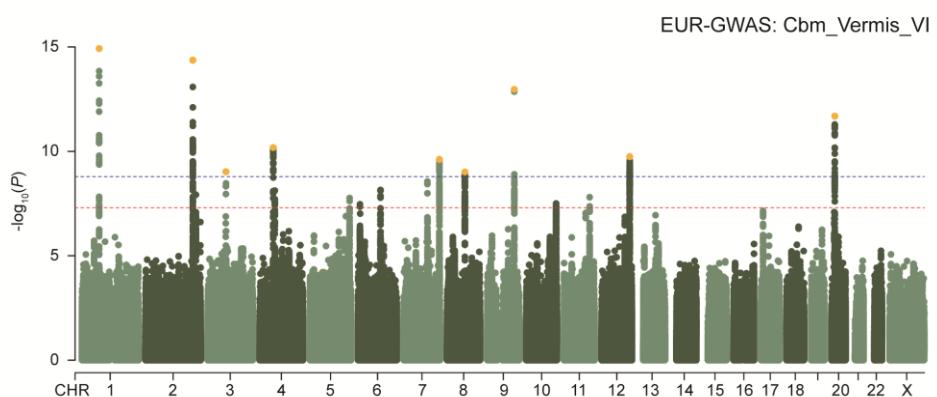
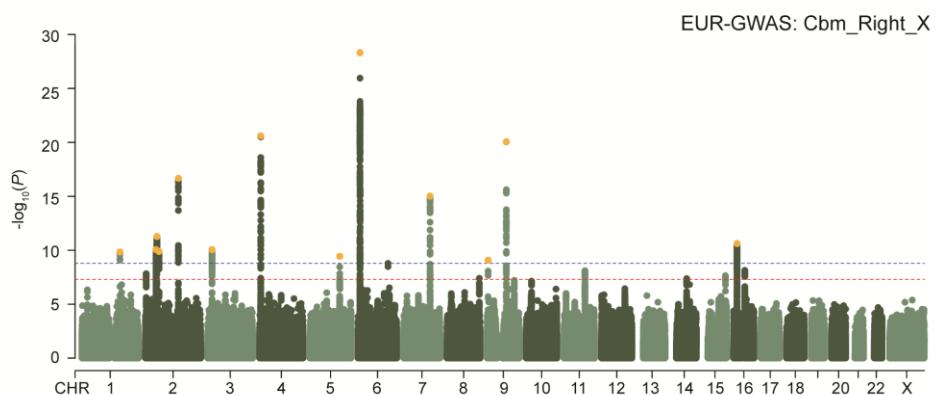
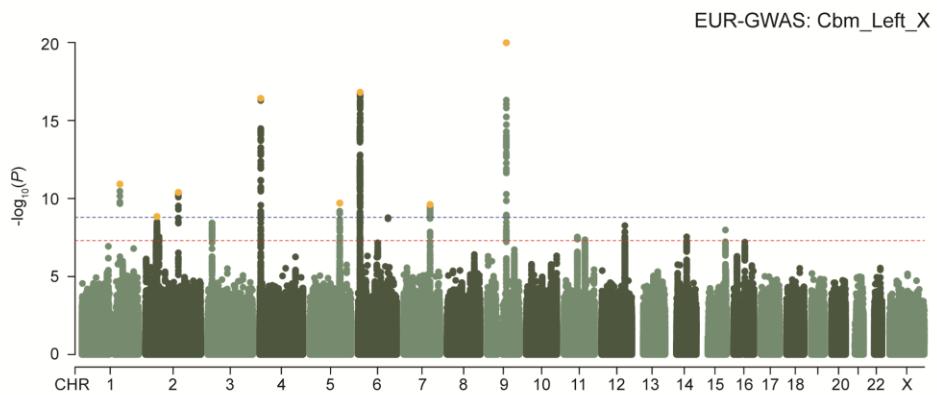


EUR-GWAS: Cbm_Left_IX

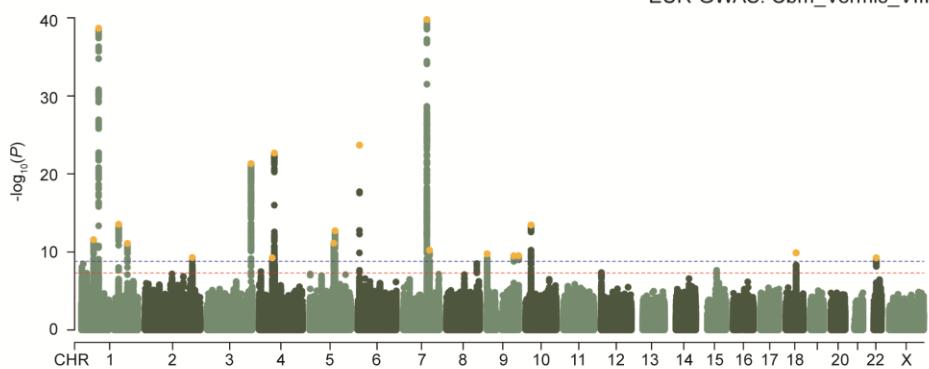


EUR-GWAS: Cbm_Right_IX

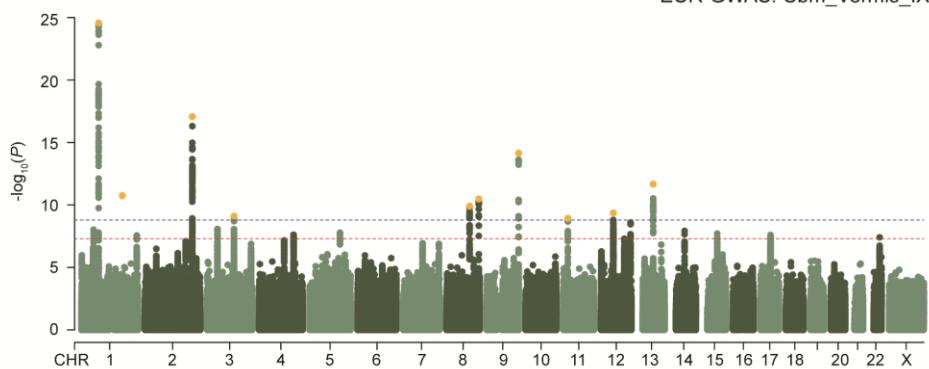




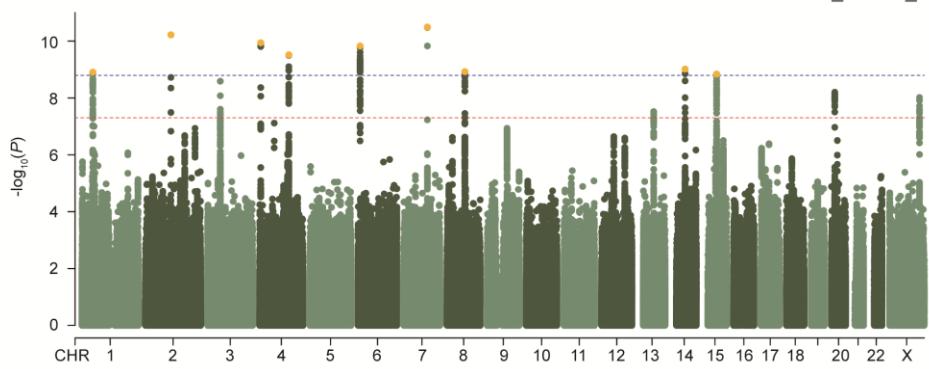
EUR-GWAS: Cbm_Vermis_VIII



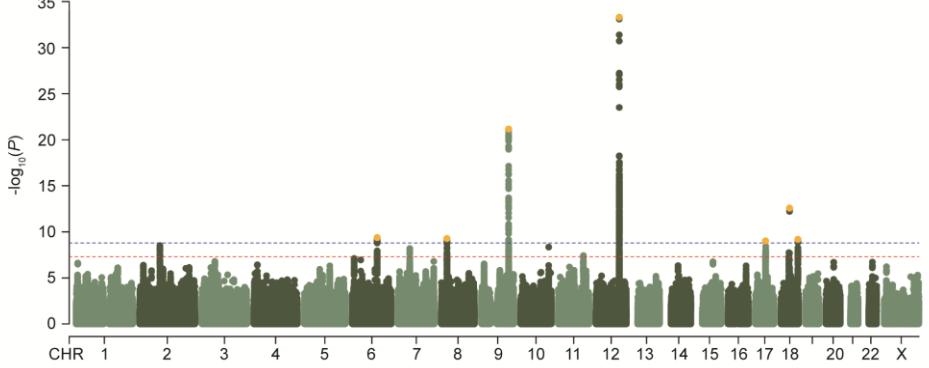
EUR-GWAS: Cbm_Vermis_IX

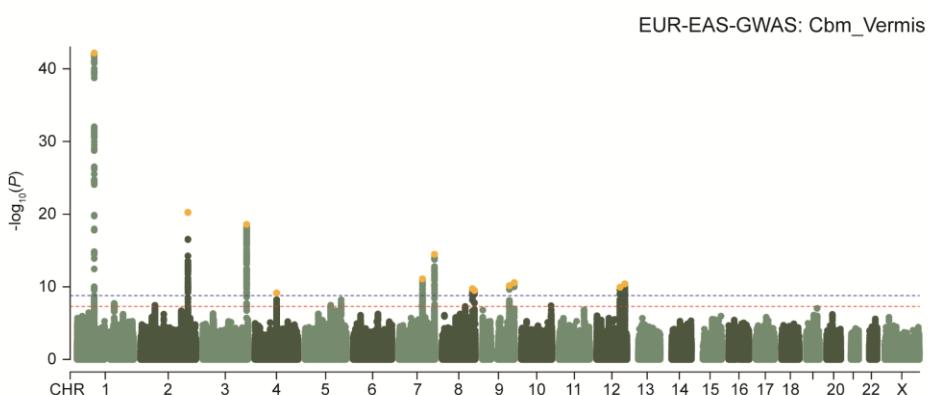
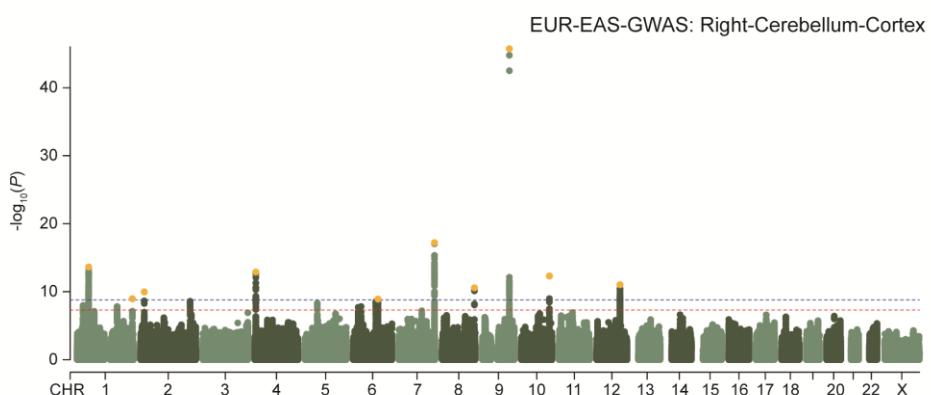
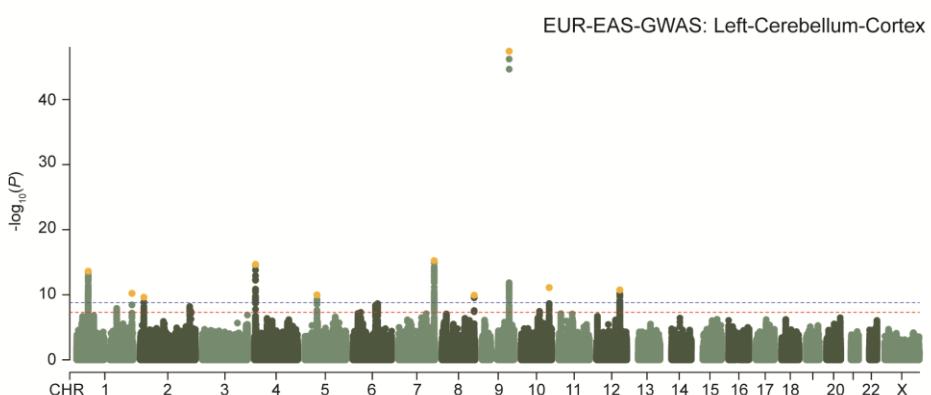
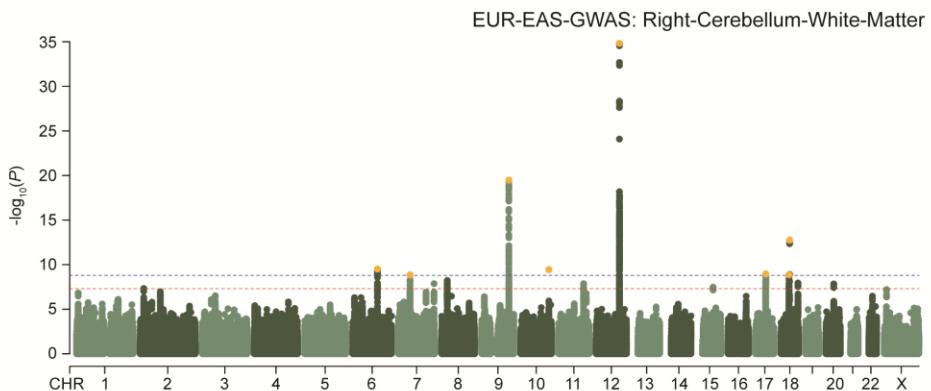


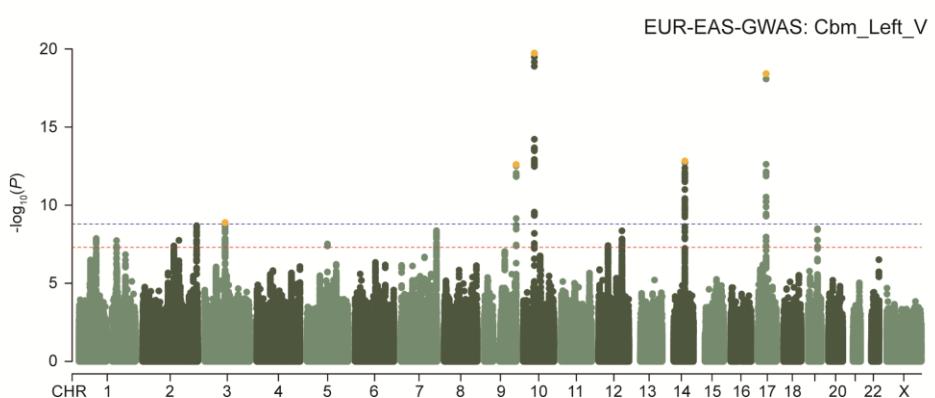
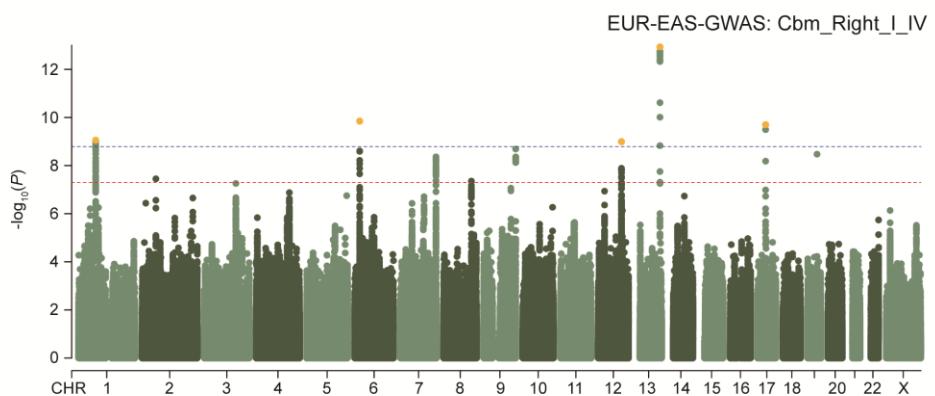
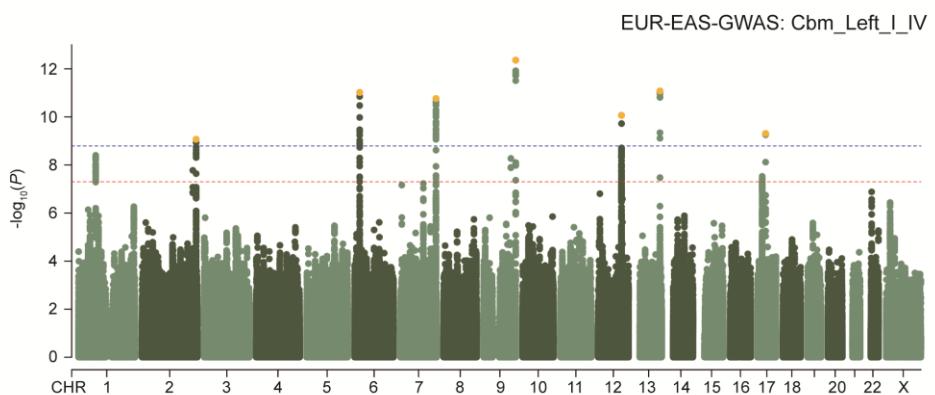
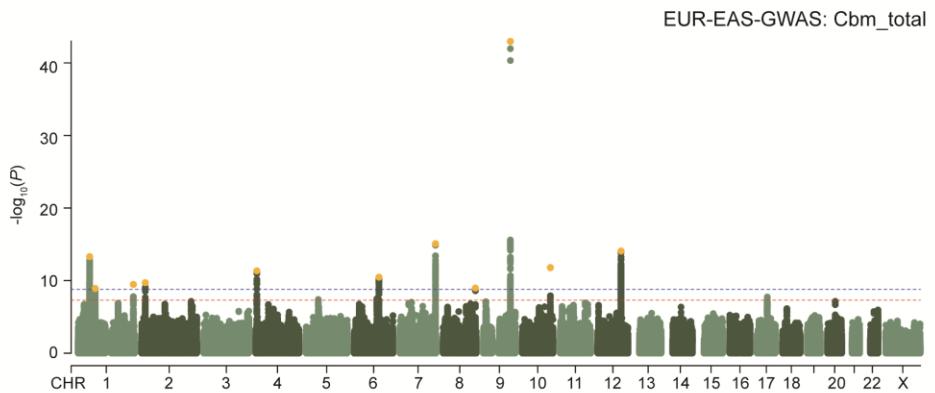
EUR-GWAS: Cbm_Vermis_X

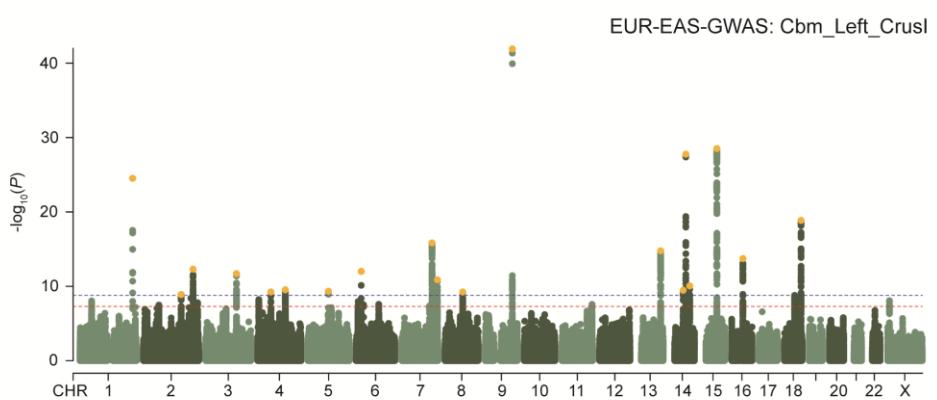
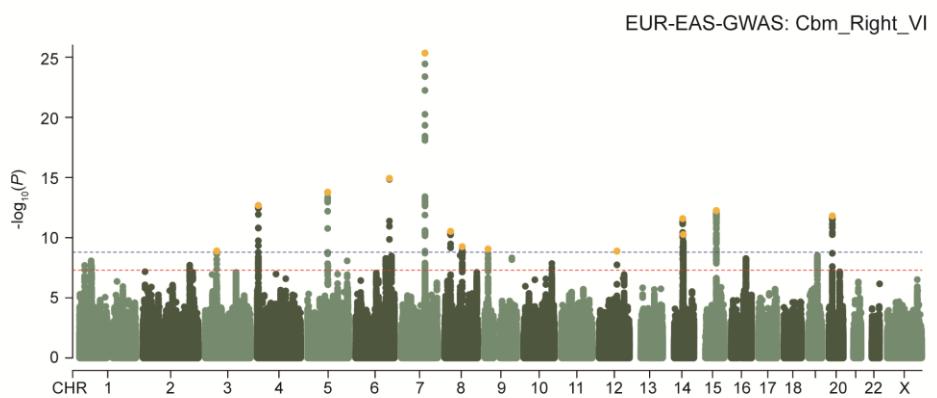
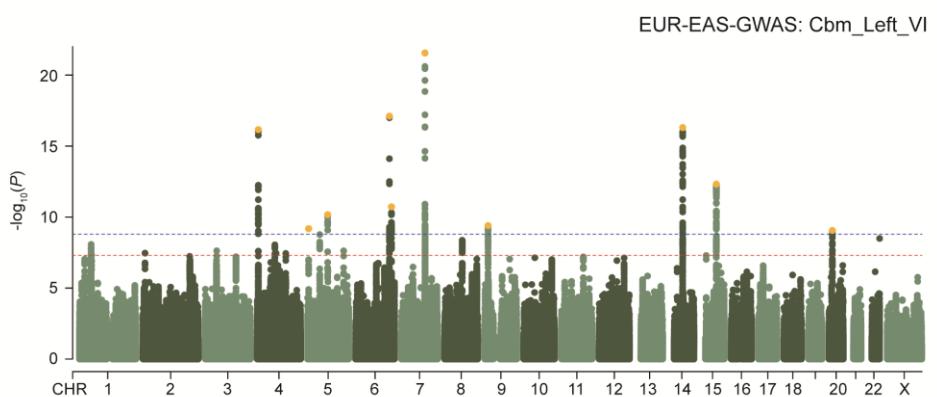
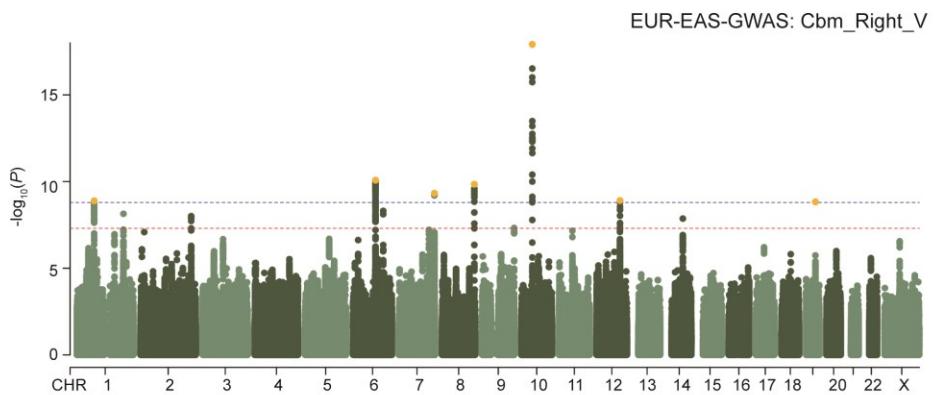


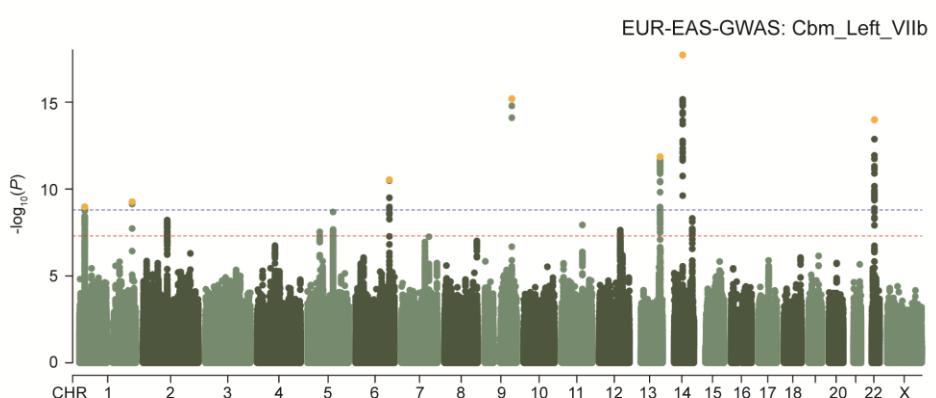
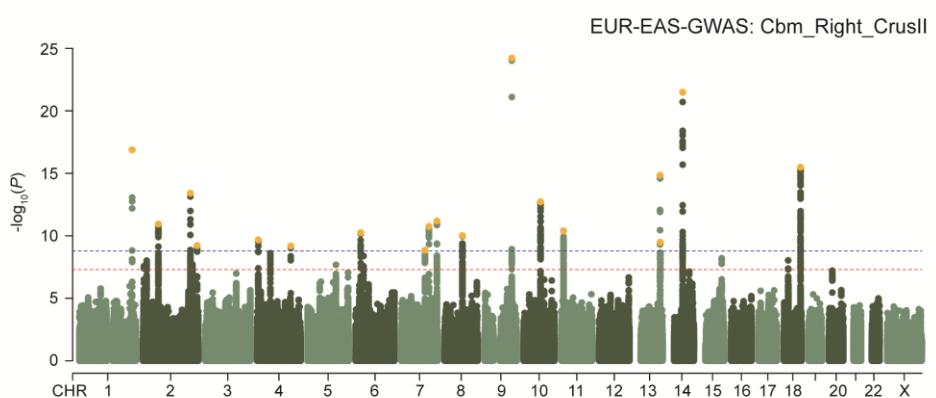
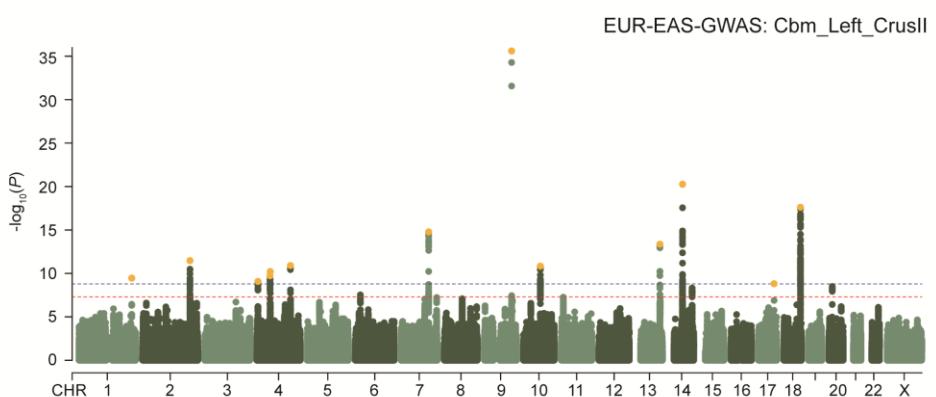
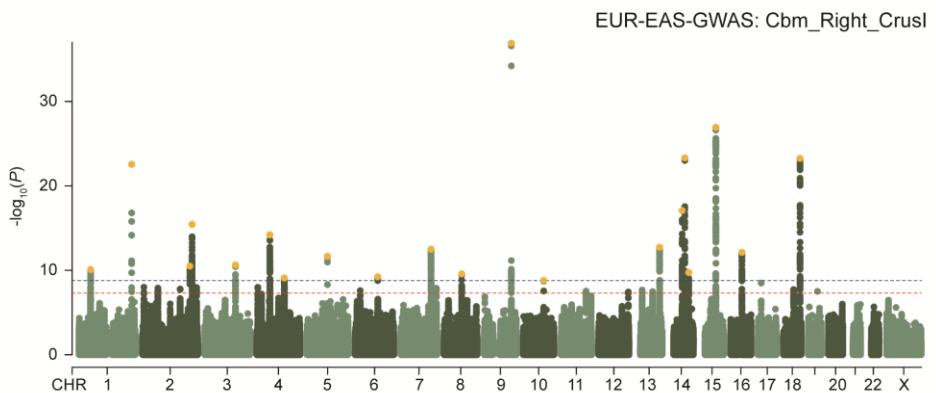
EUR-EAS-GWAS: Left-Cerebellum-White-Matter

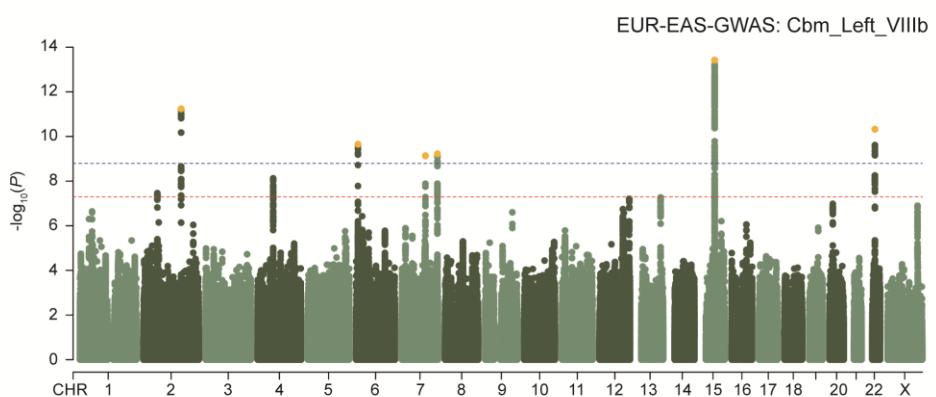
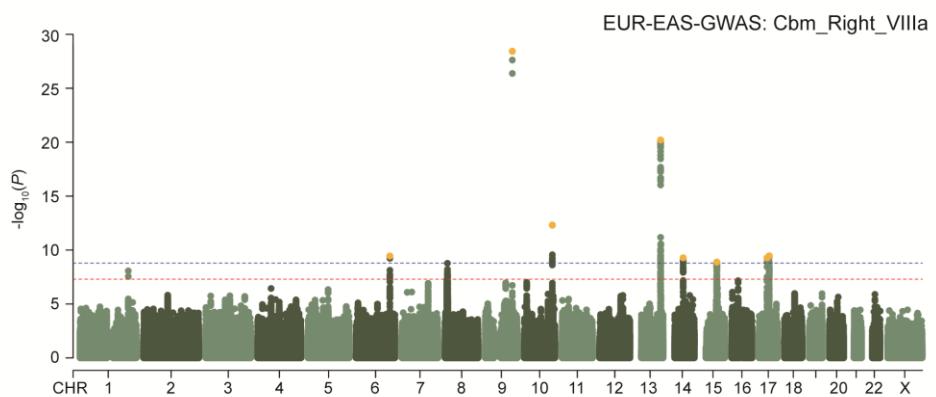
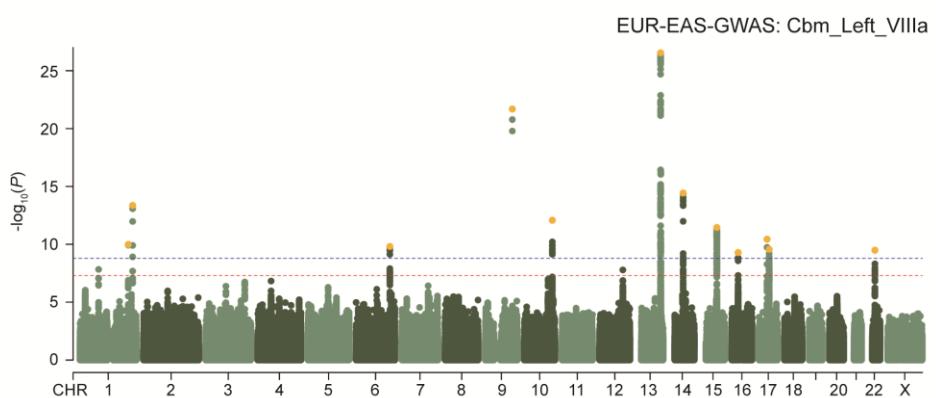
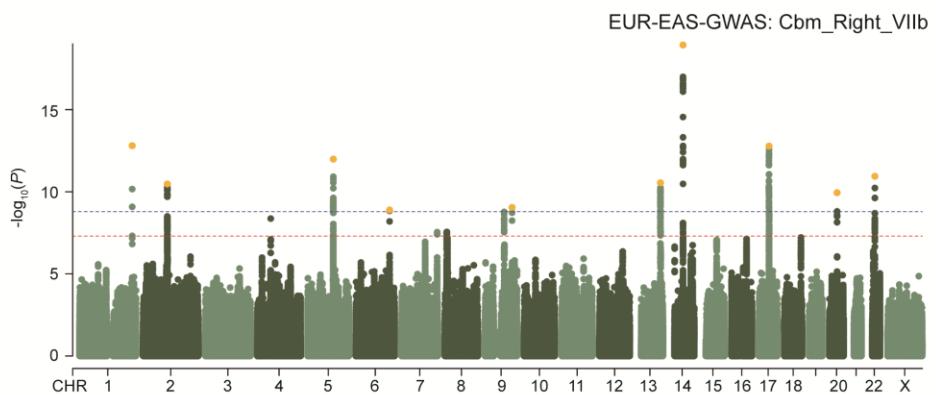




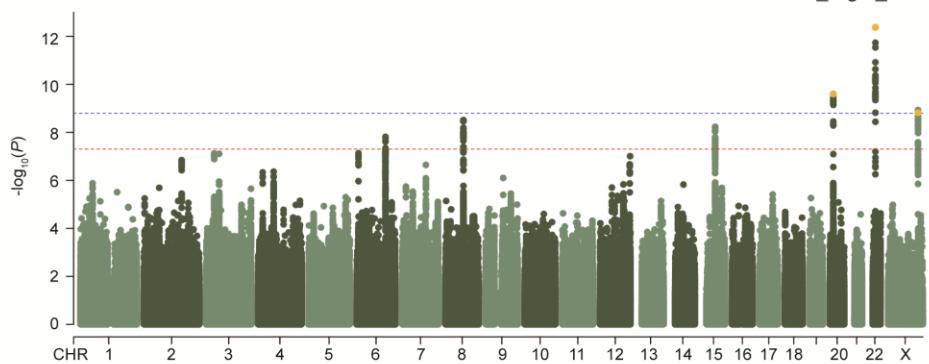




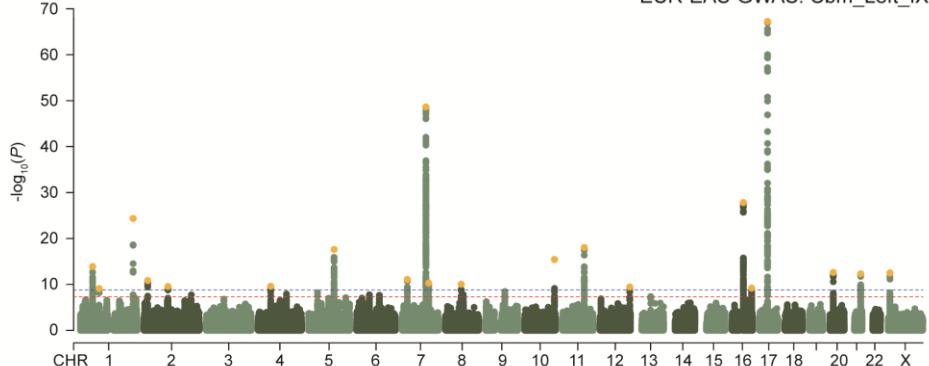




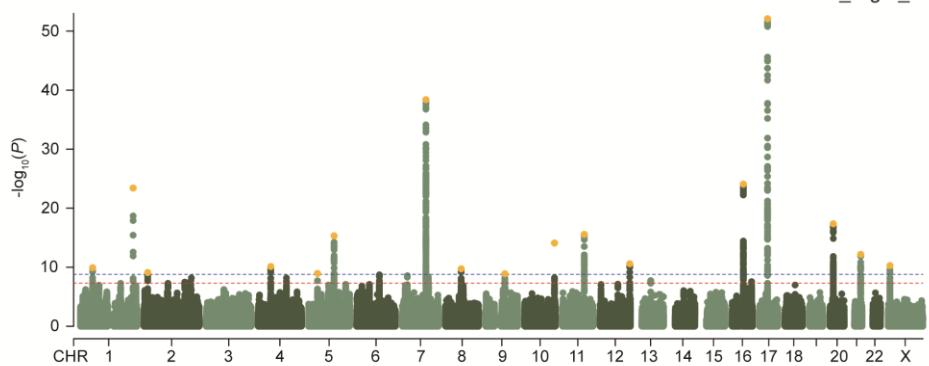
EUR-EAS-GWAS: Cbm_Right_VIIIb



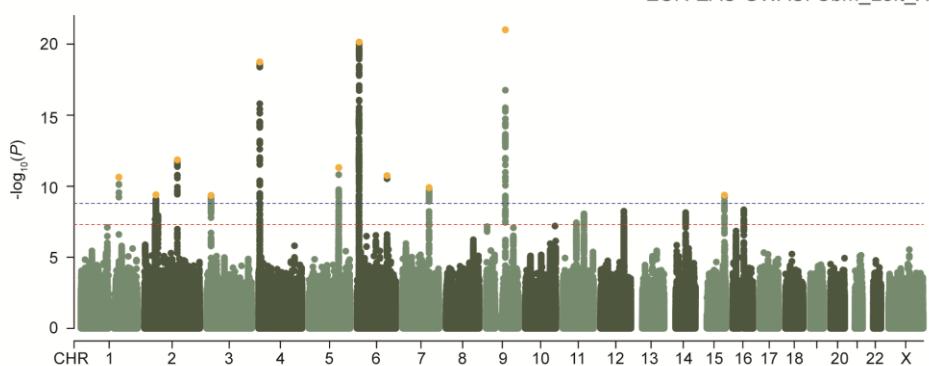
EUR-EAS-GWAS: Cbm_Left_IX



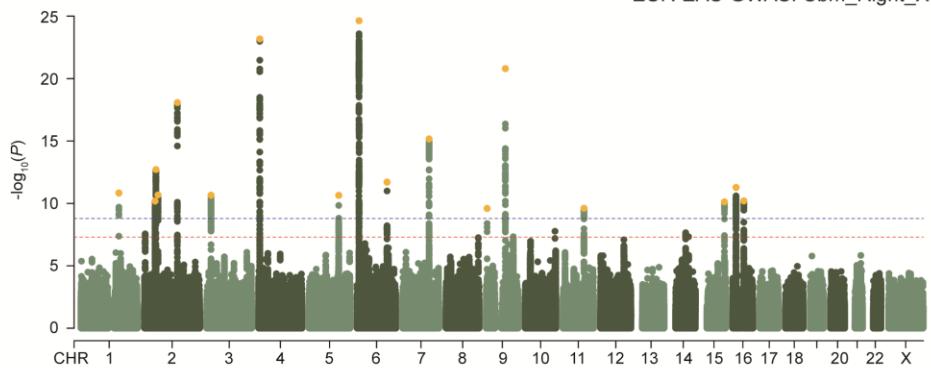
EUR-EAS-GWAS: Cbm_Right_IX



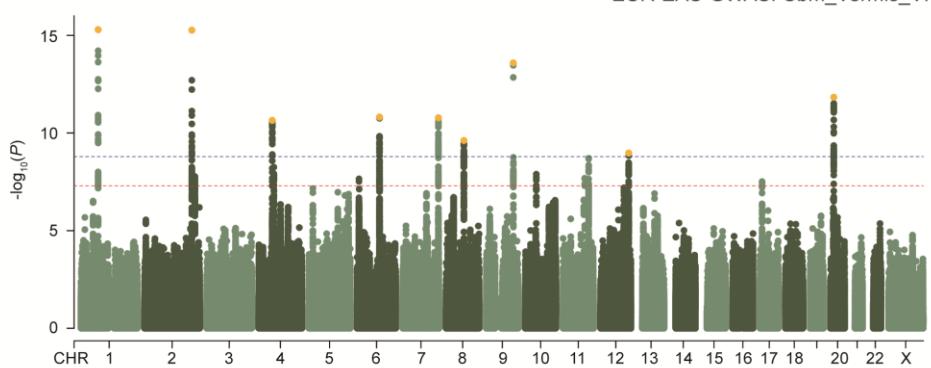
EUR-EAS-GWAS: Cbm_Left_X



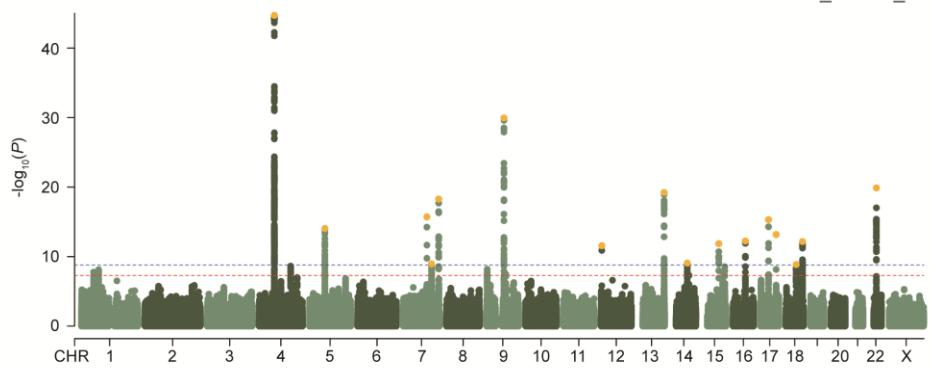
EUR-EAS-GWAS: Cbm_Right_X



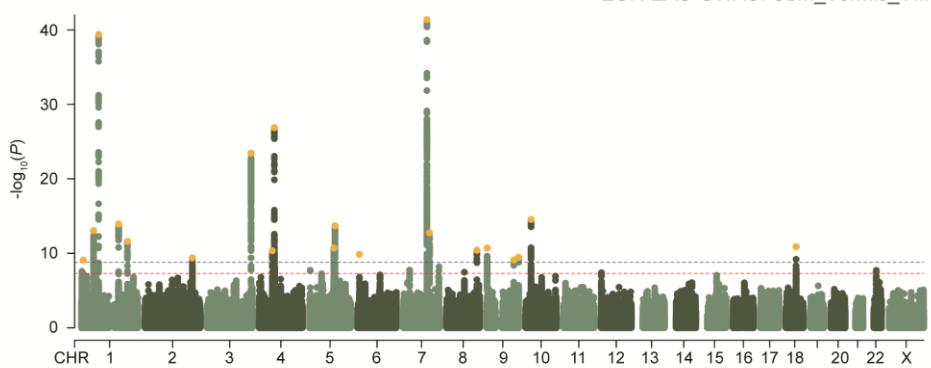
EUR-EAS-GWAS: Cbm_Vermis_VI



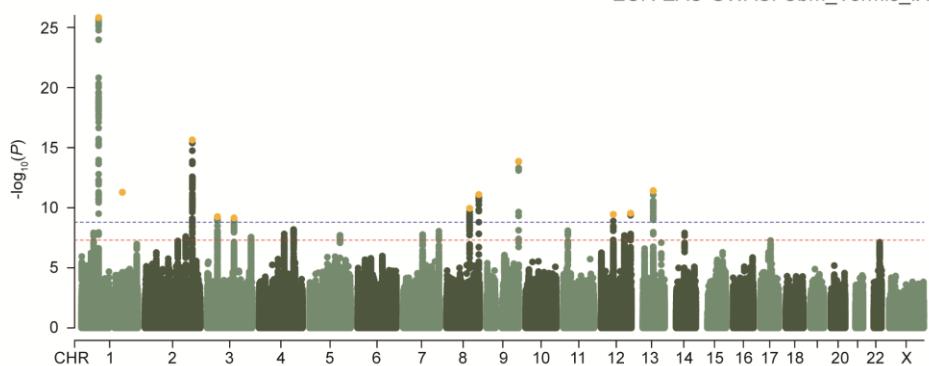
EUR-EAS-GWAS: Cbm_Vermis_VII



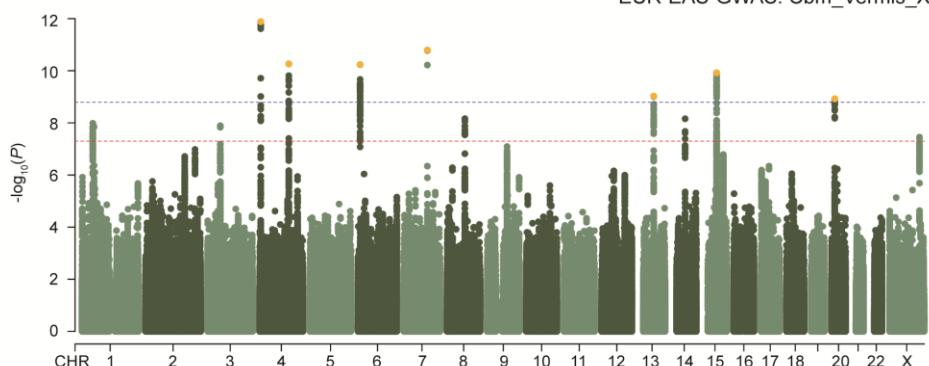
EUR-EAS-GWAS: Cbm_Vermis_VIII



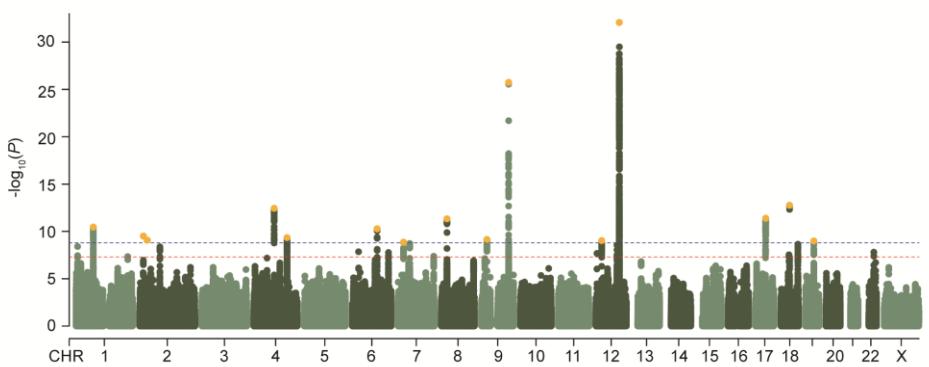
EUR-EAS-GWAS: Cbm_Vermis_IX



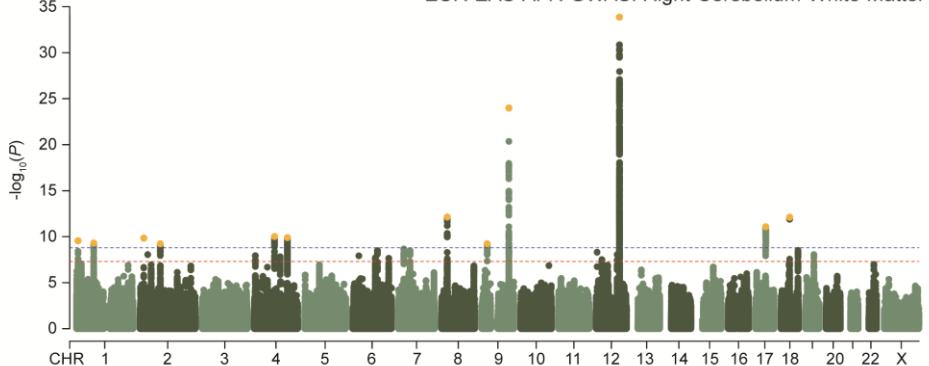
EUR-EAS-GWAS: Cbm_Vermis_X



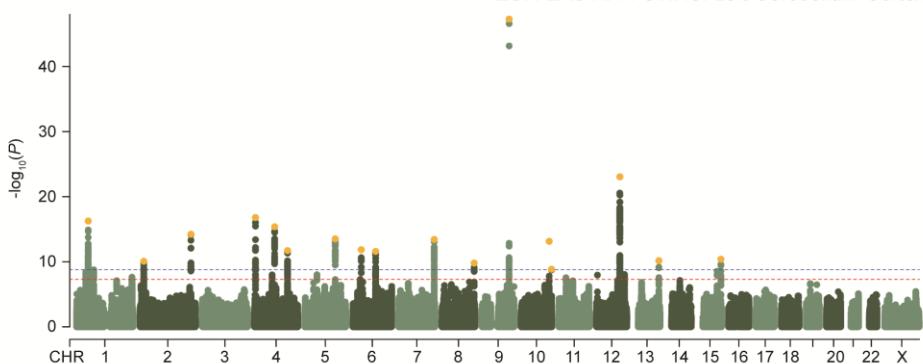
EUR-EAS-AFR-GWAS: Left-Cerebellum-White-Matter



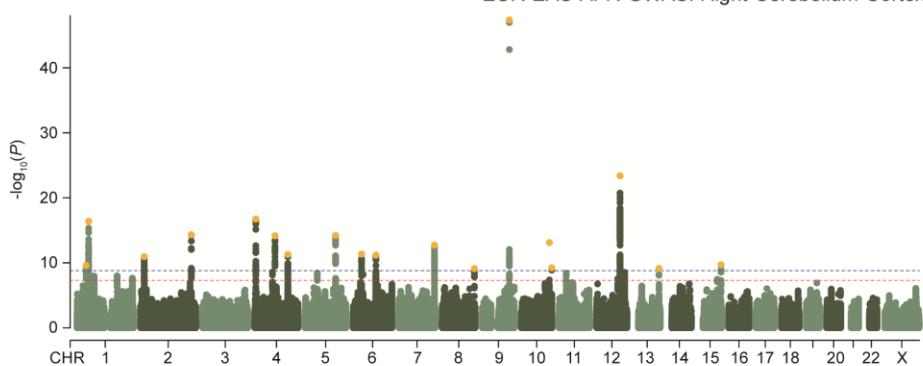
EUR-EAS-AFR-GWAS: Right-Cerebellum-White-Matter



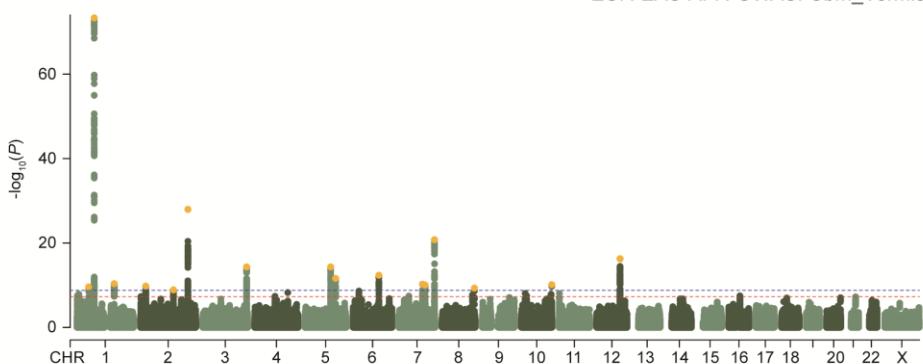
EUR-EAS-AFR-GWAS: Left-Cerebellum-Cortex



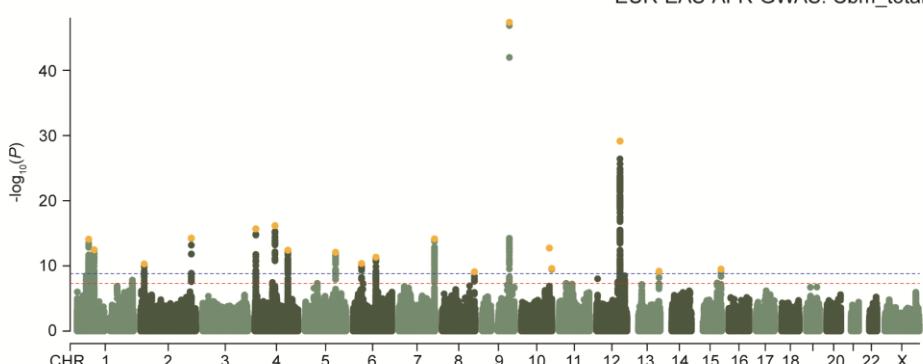
EUR-EAS-AFR-GWAS: Right-Cerebellum-Cortex

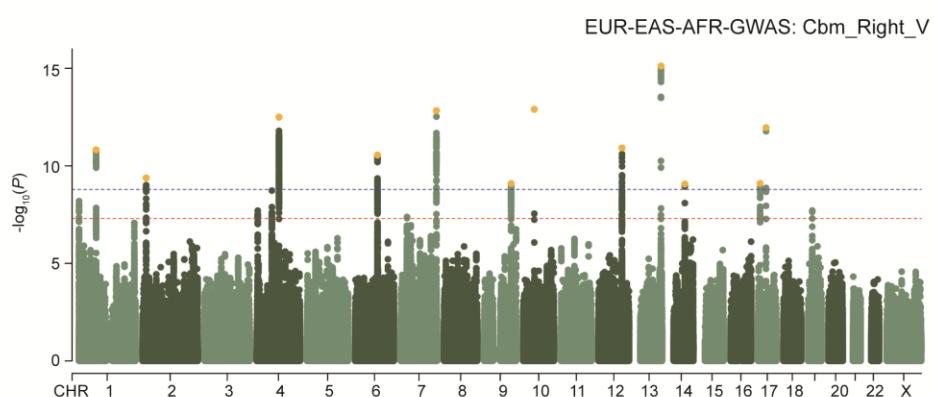
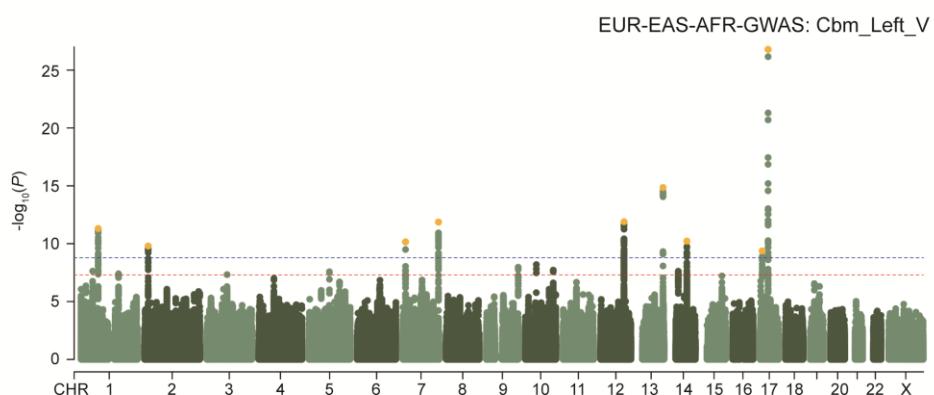
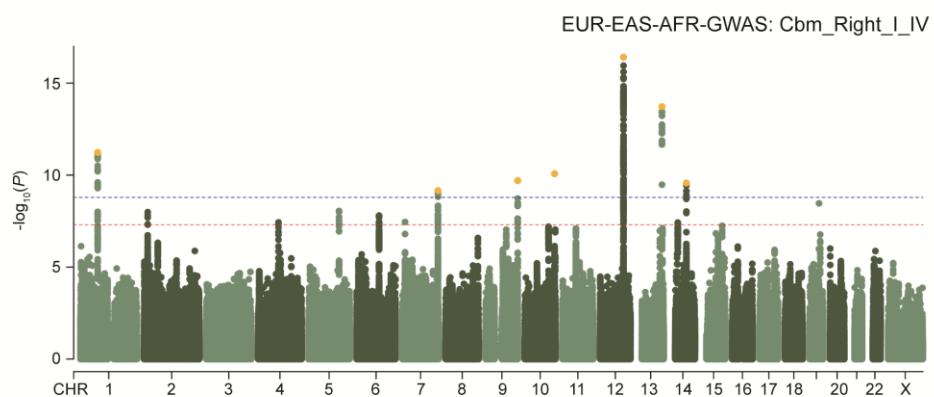
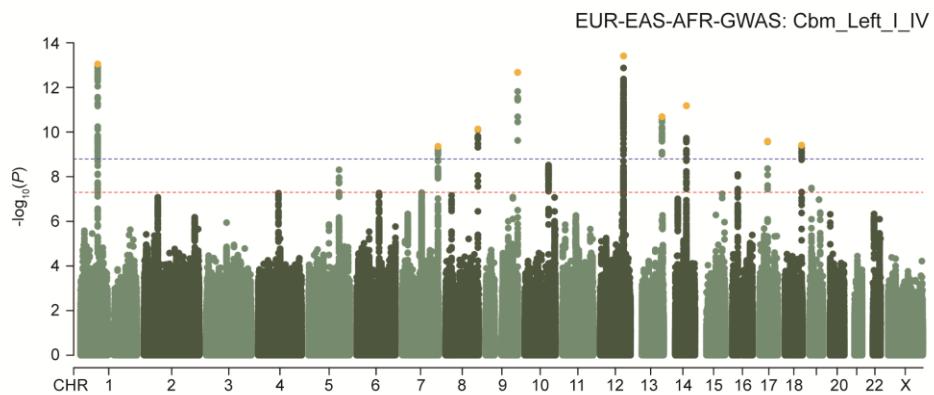


EUR-EAS-AFR-GWAS: Cbm_Vermis

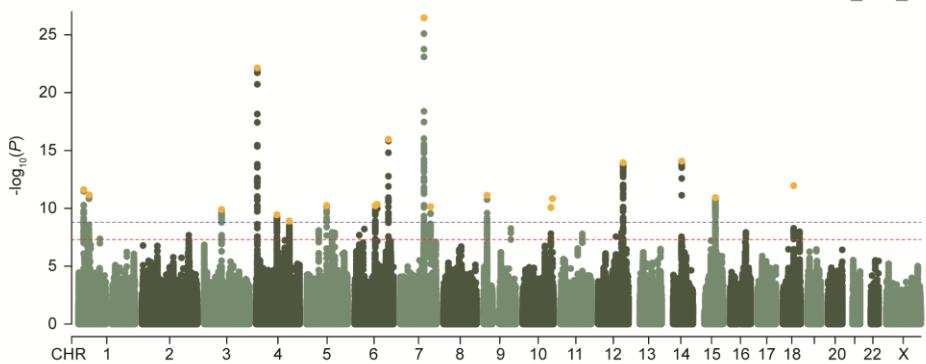


EUR-EAS-AFR-GWAS: Cbm_total

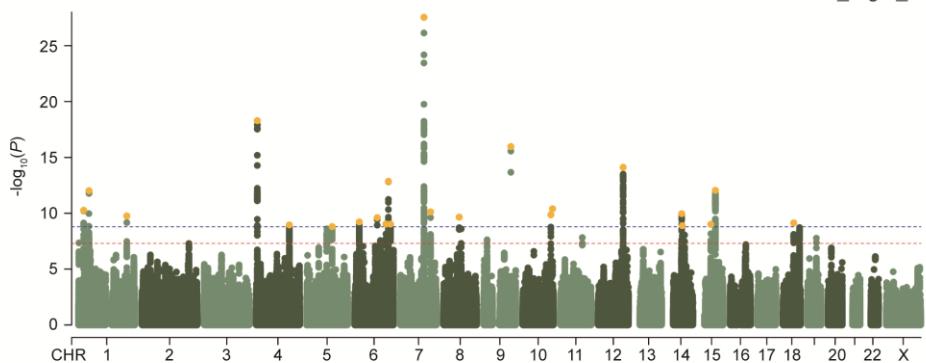




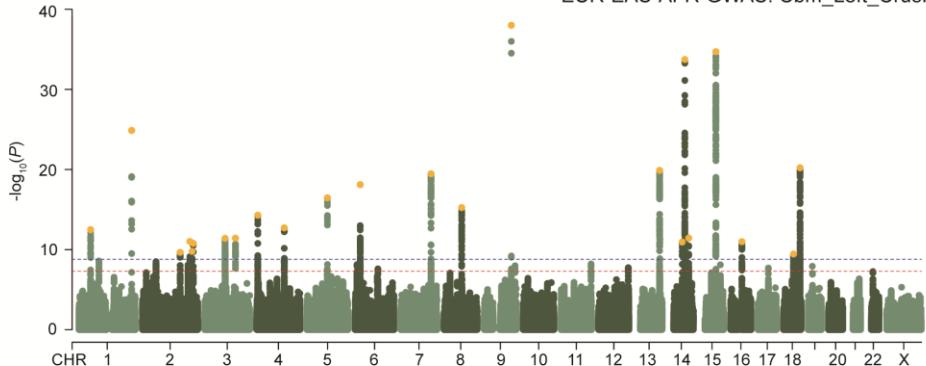
EUR-EAS-AFR-GWAS: Cbm_Left_VI



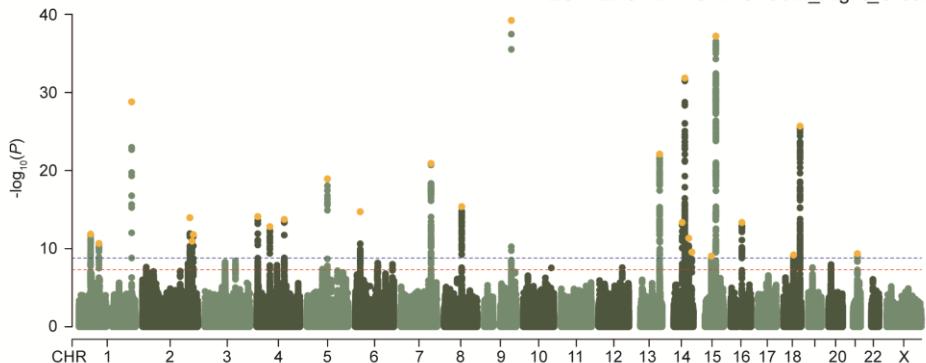
EUR-EAS-AFR-GWAS: Cbm_Right_VI

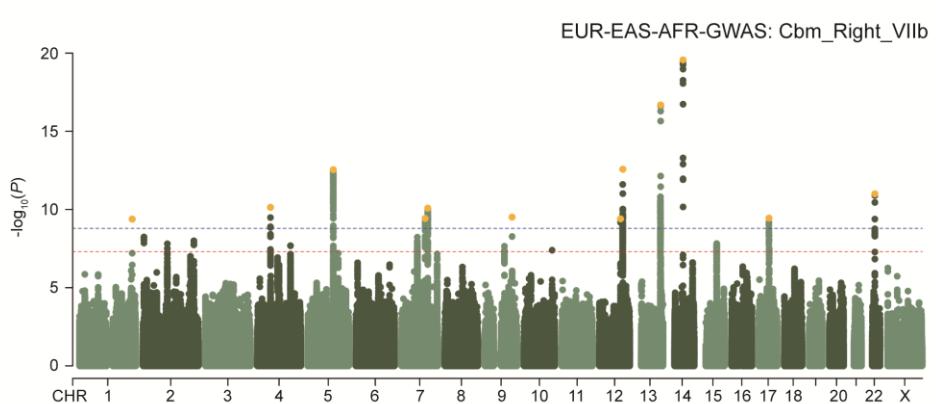
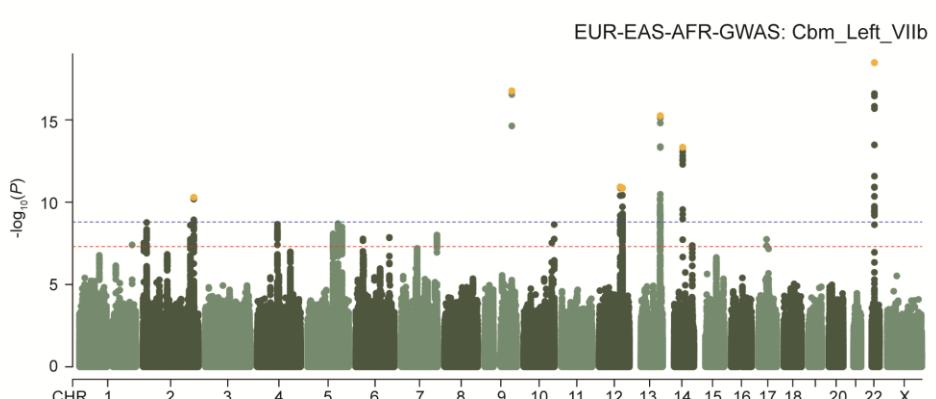
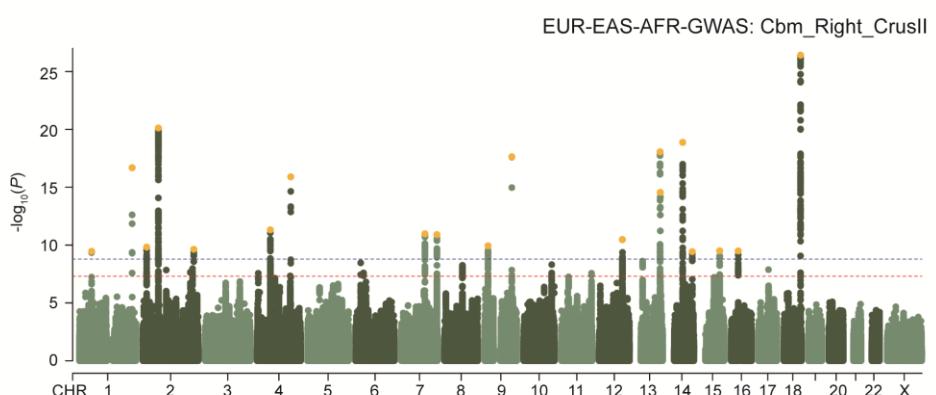
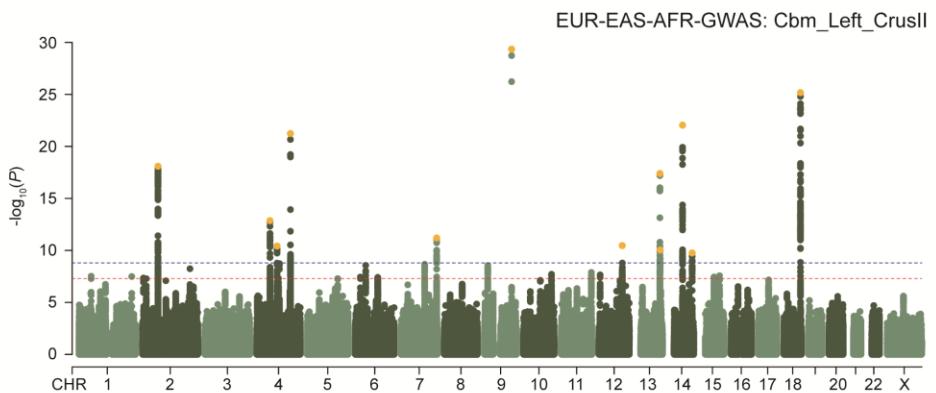


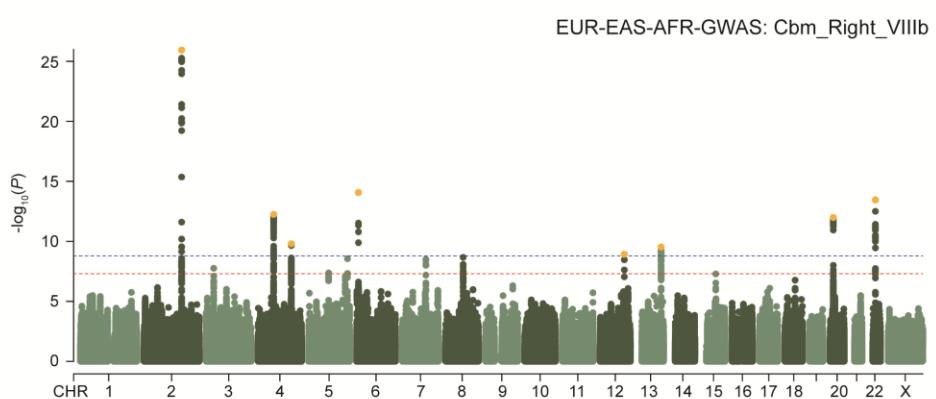
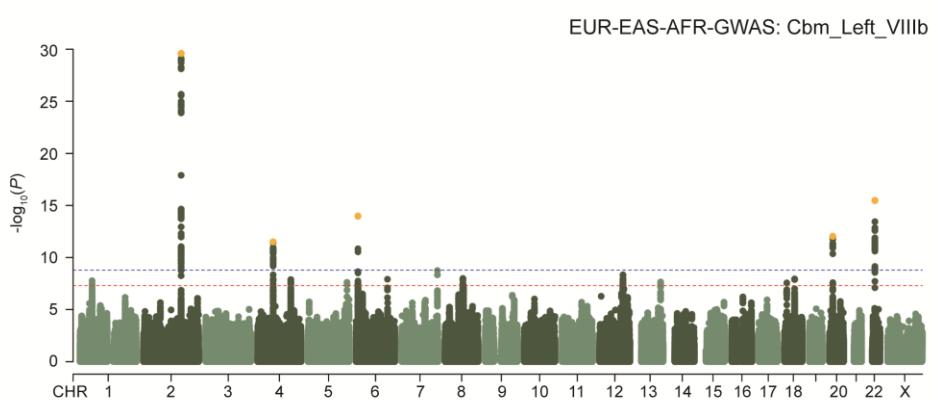
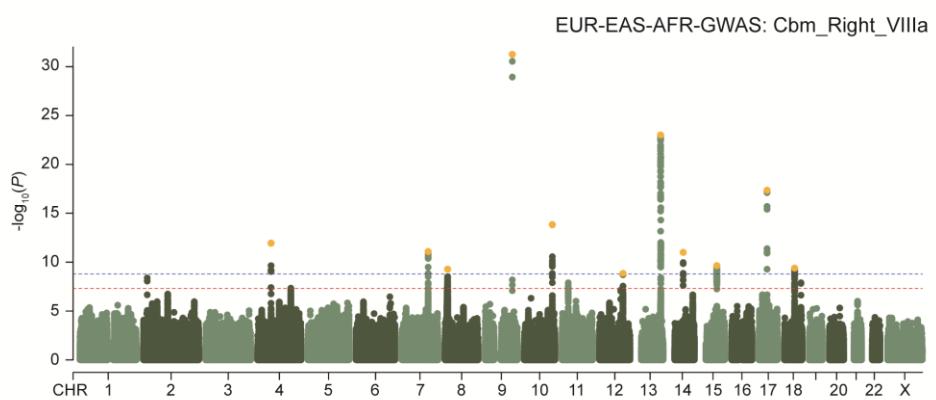
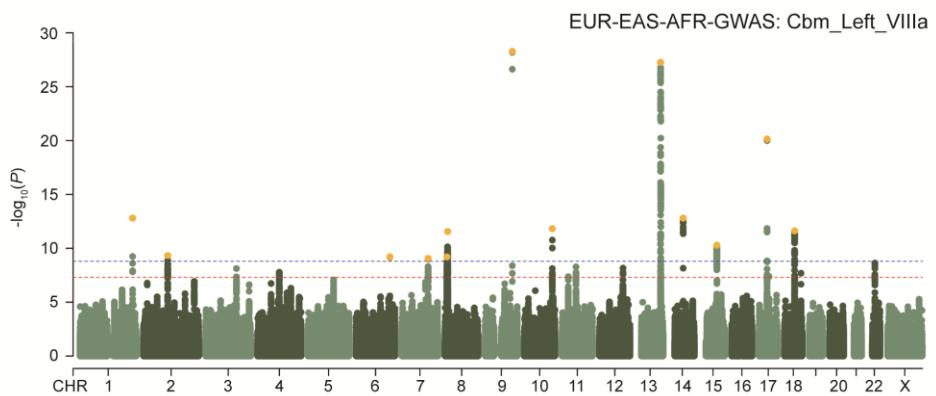
EUR-EAS-AFR-GWAS: Cbm_Left_CrusI

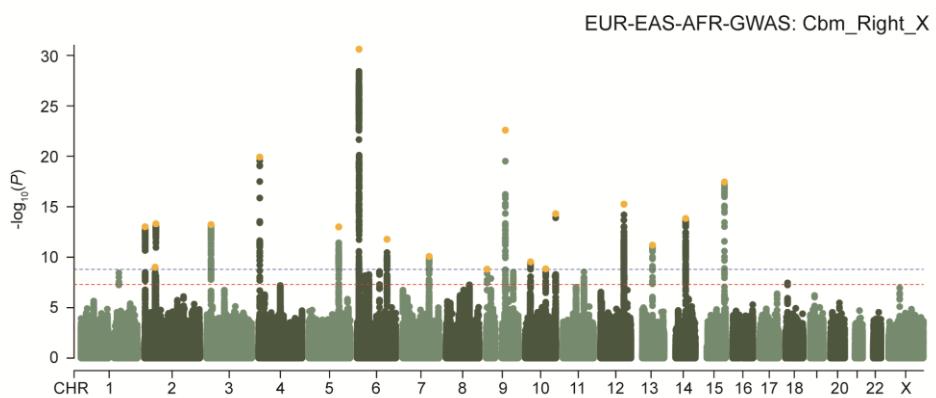
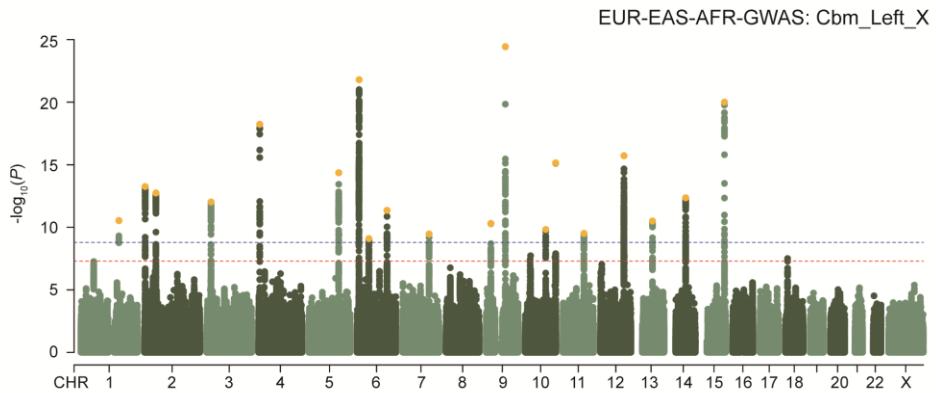
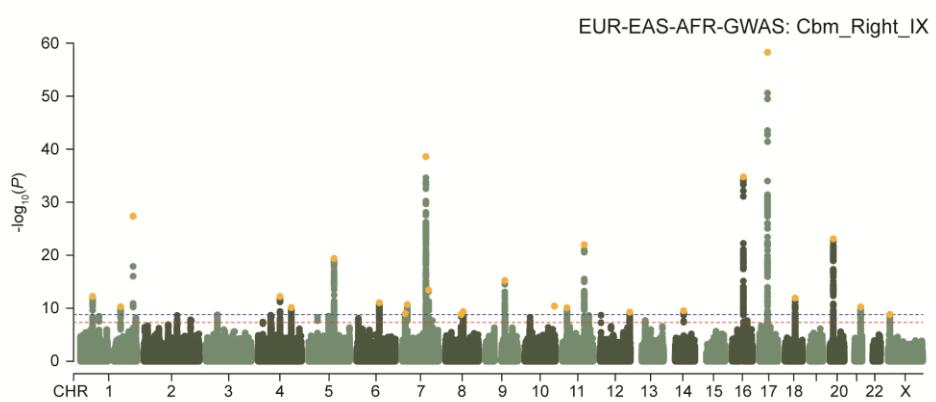
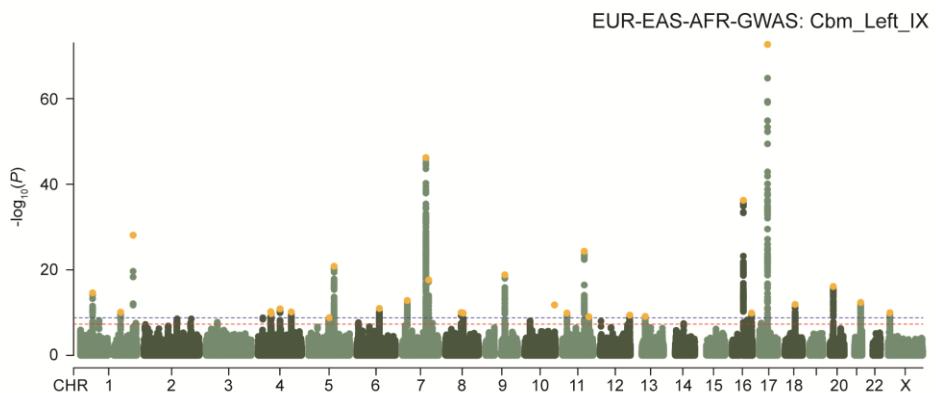


EUR-EAS-AFR-GWAS: Cbm_Right_CrusI

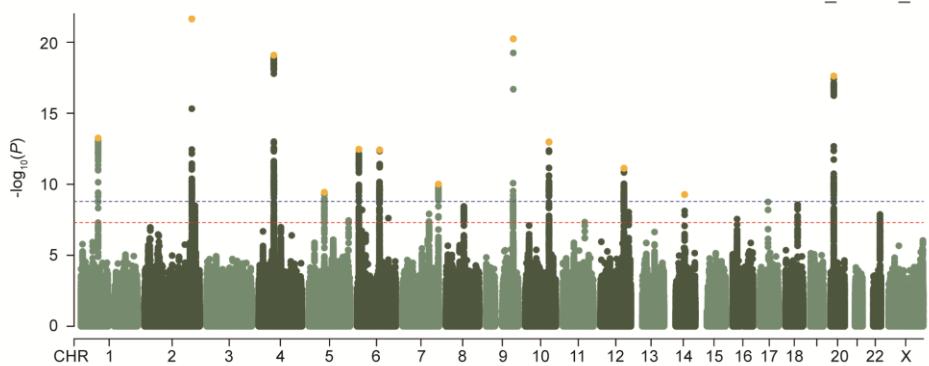




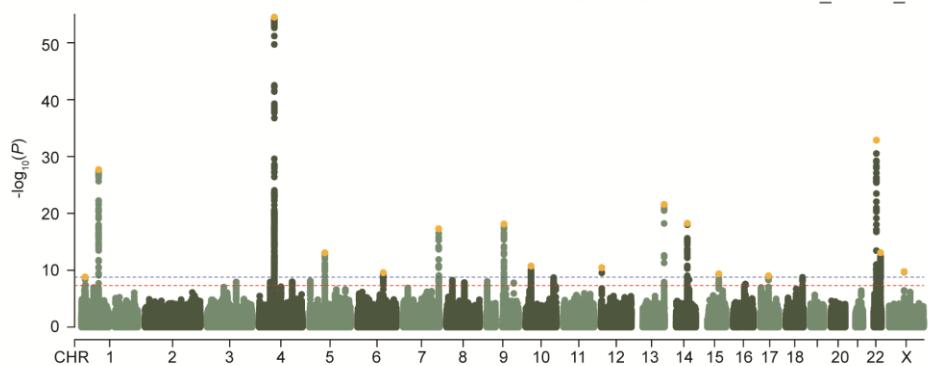




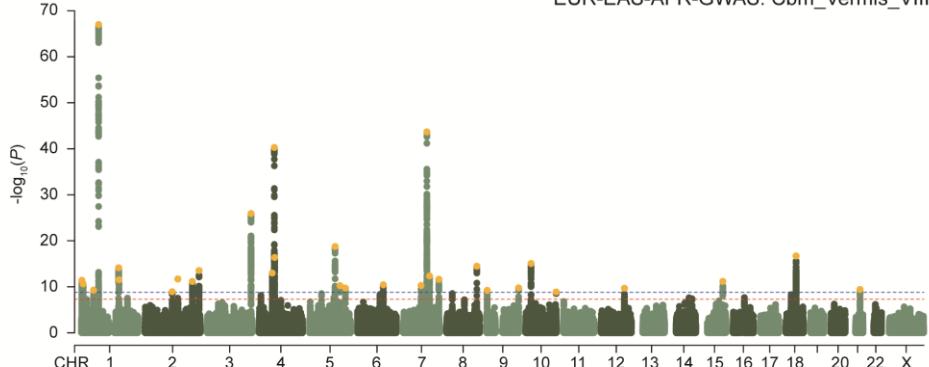
EUR-EAS-AFR-GWAS: Cbm_Vermis_VI



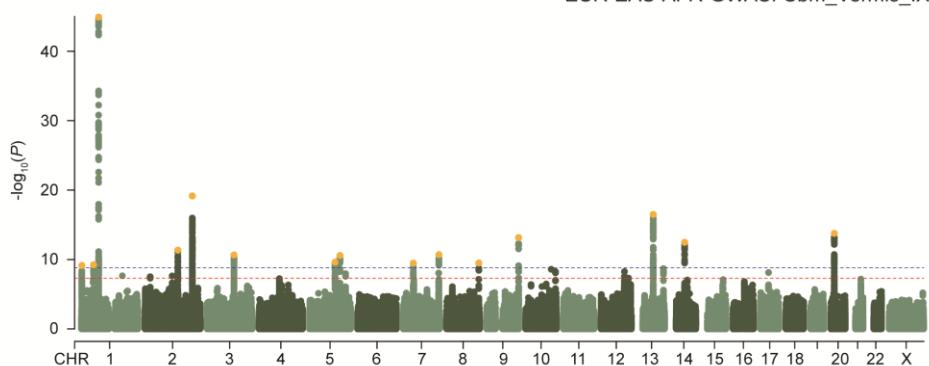
EUR-EAS-AFR-GWAS: Cbm_Vermis_VII



EUR-EAS-AFR-GWAS: Cbm_Vermis_VIII



EUR-EAS-AFR-GWAS: Cbm_Vermis_IX



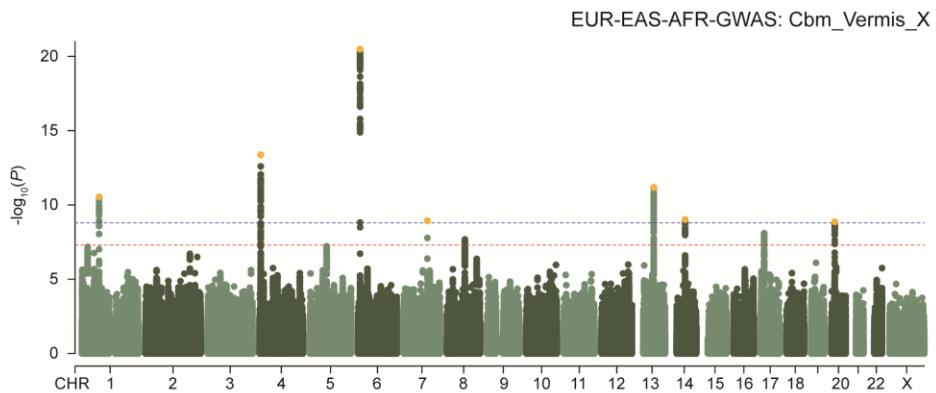
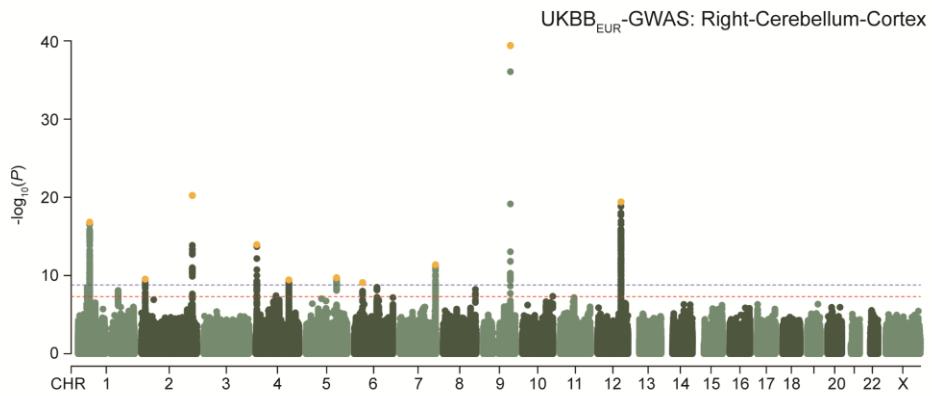
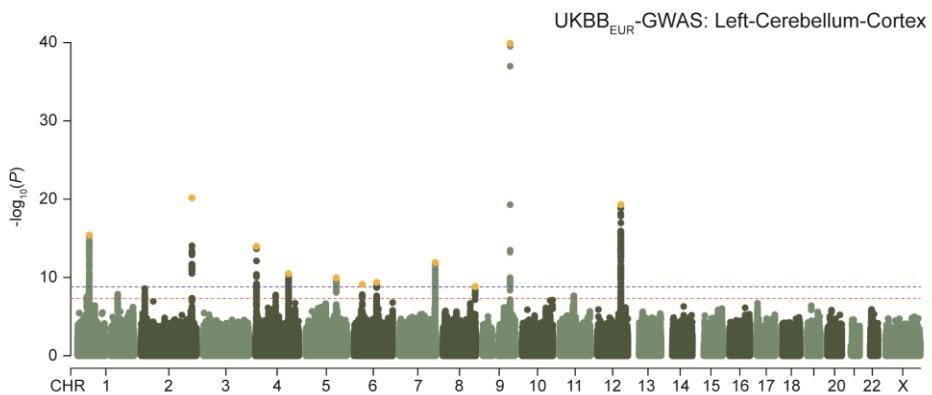
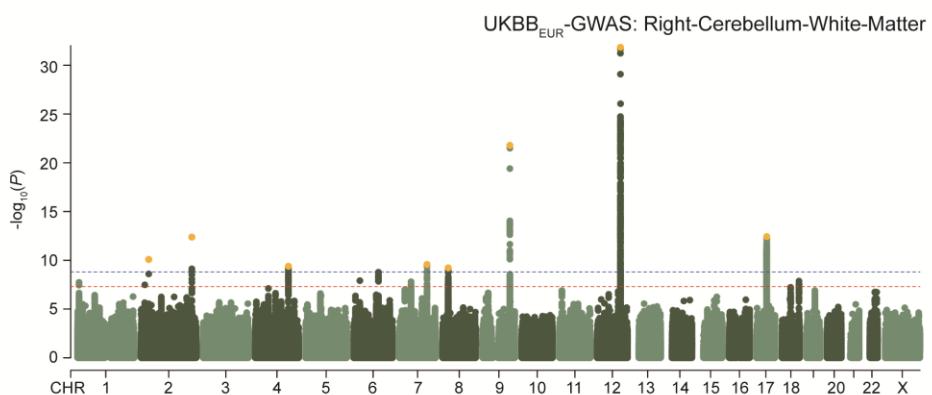
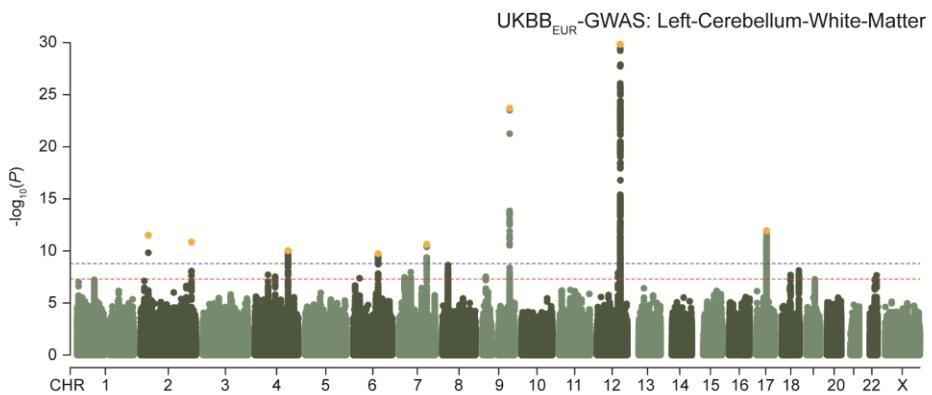
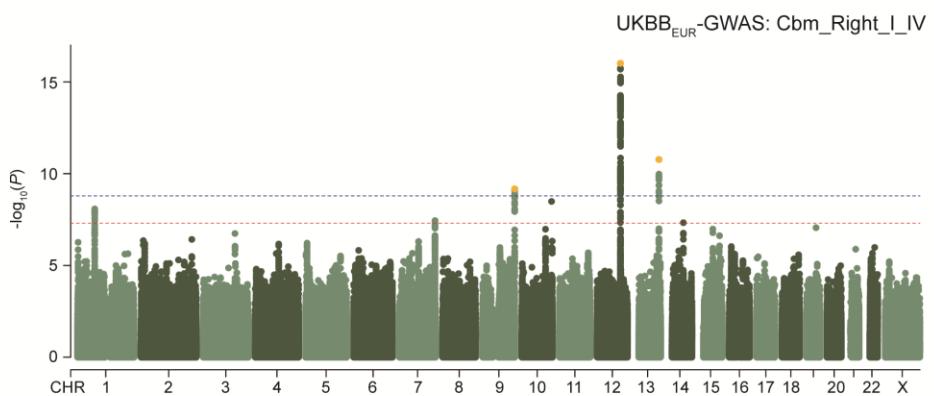
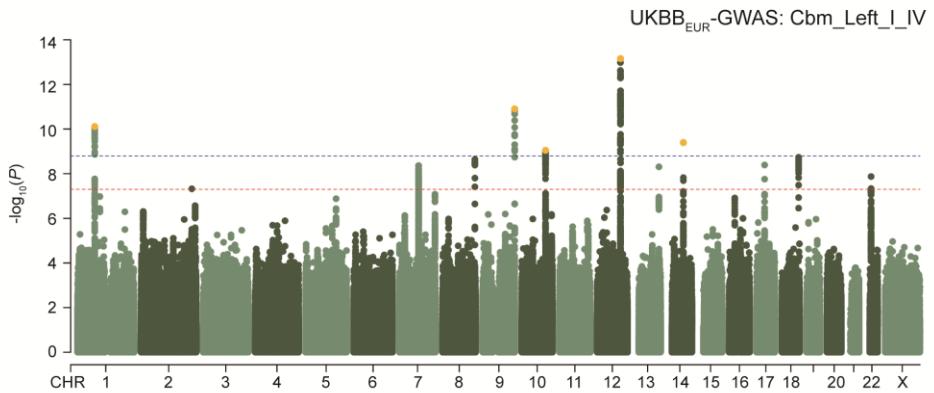
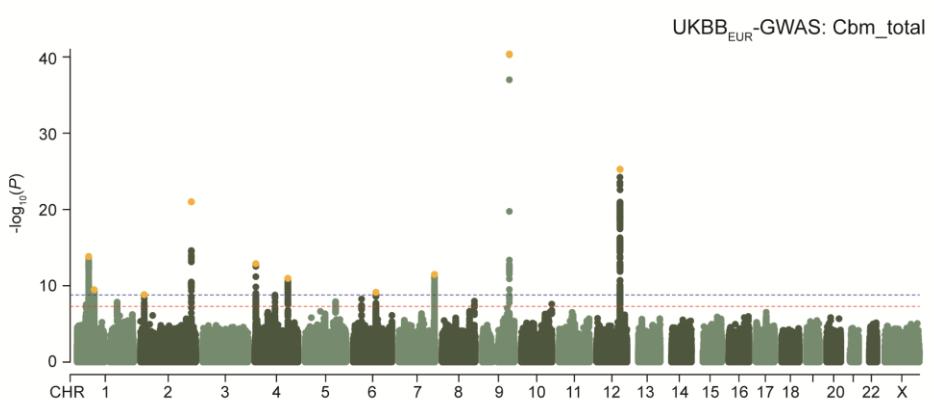
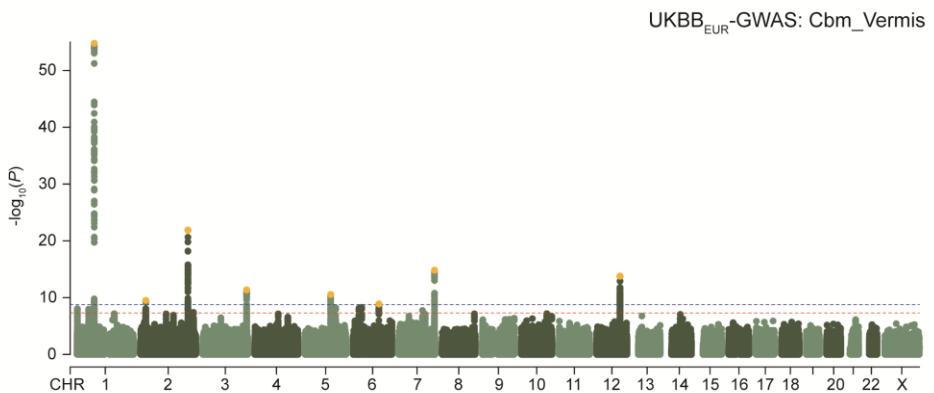
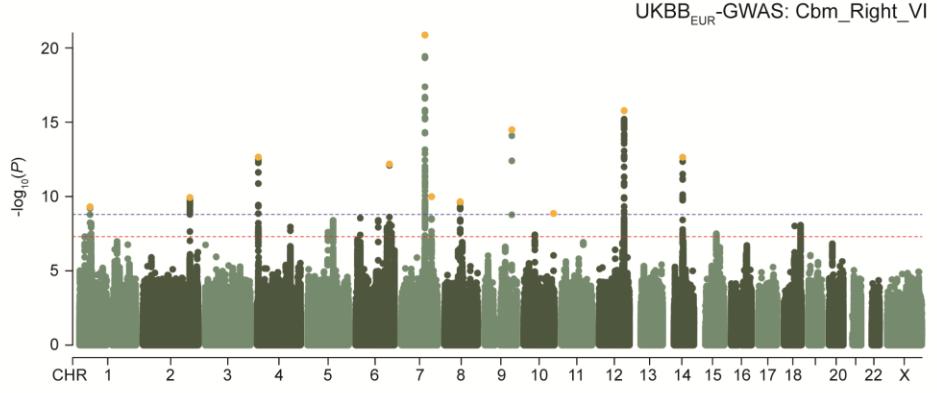
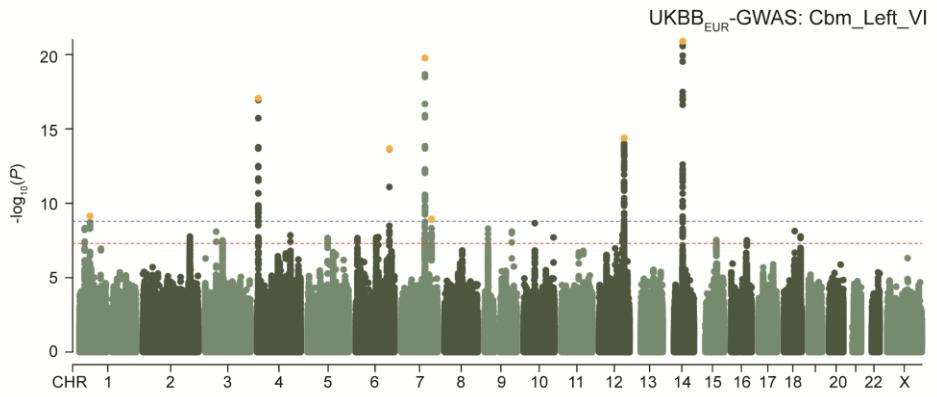
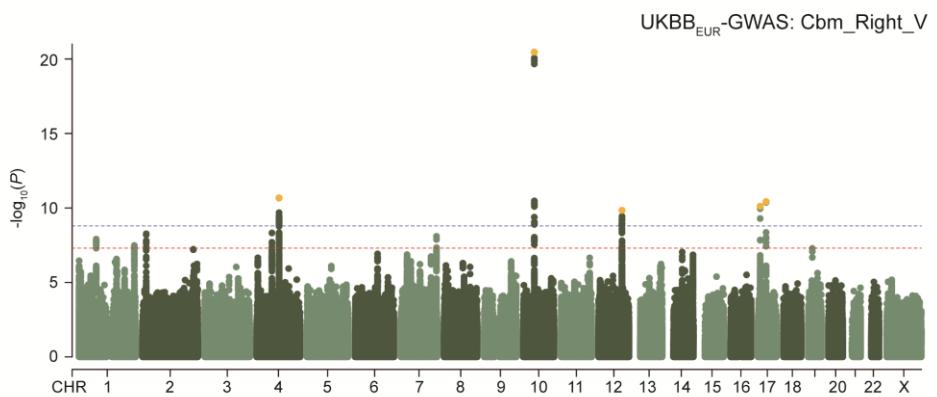
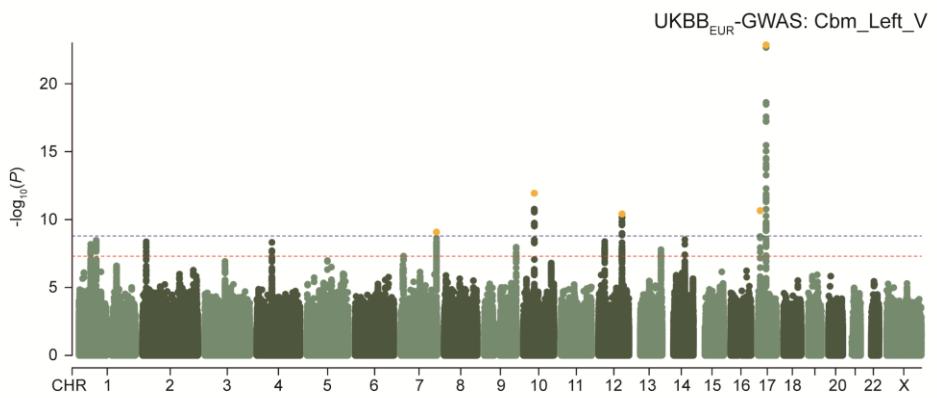
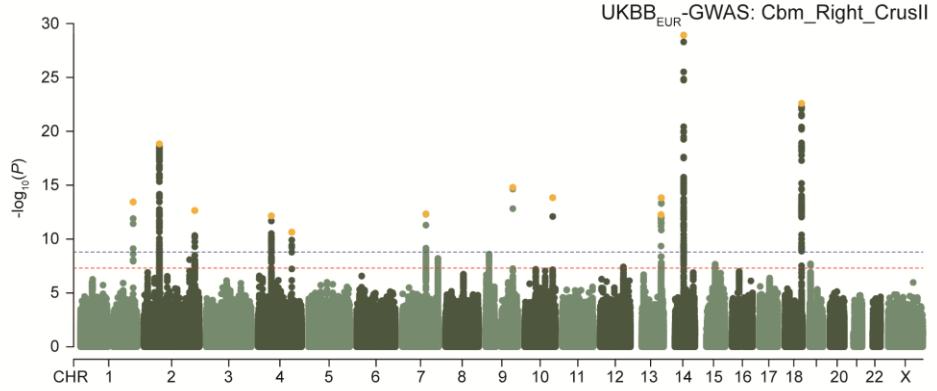
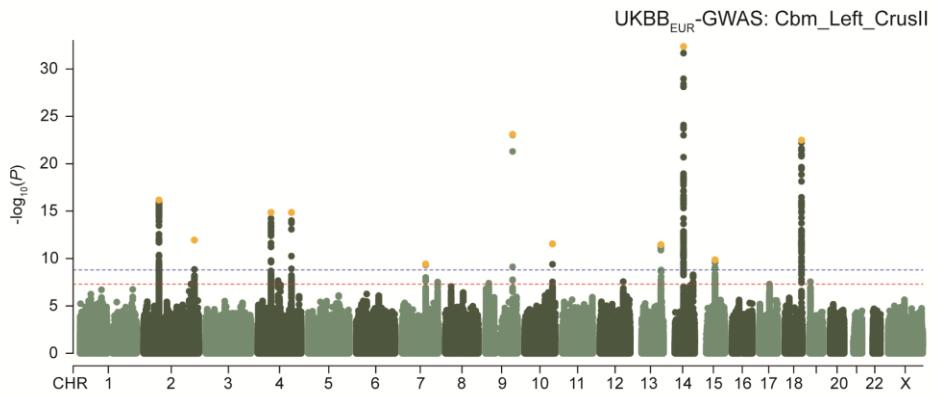
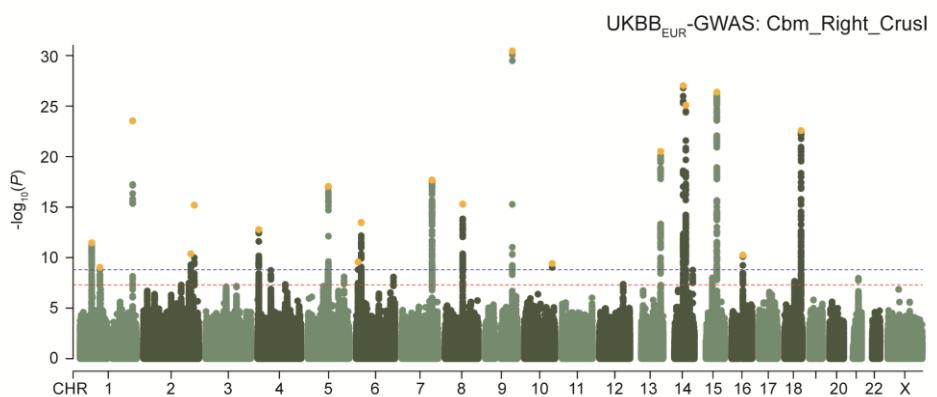
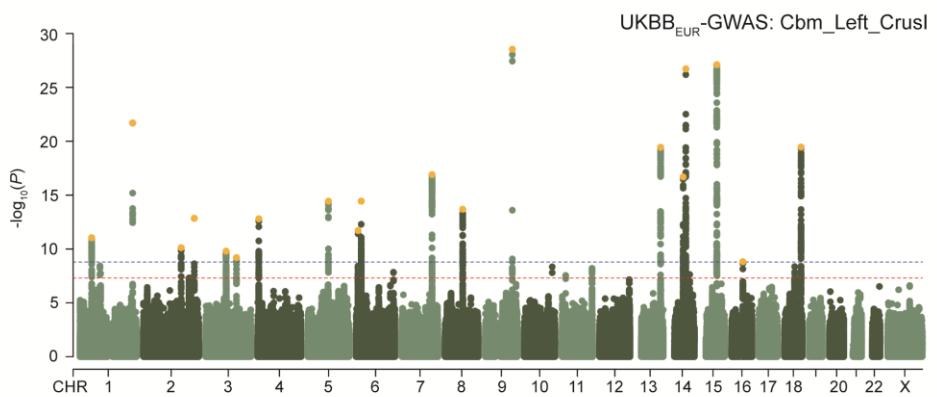


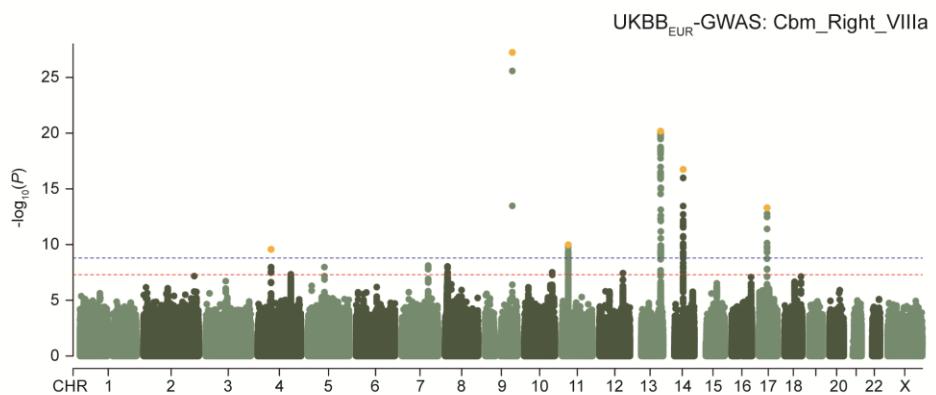
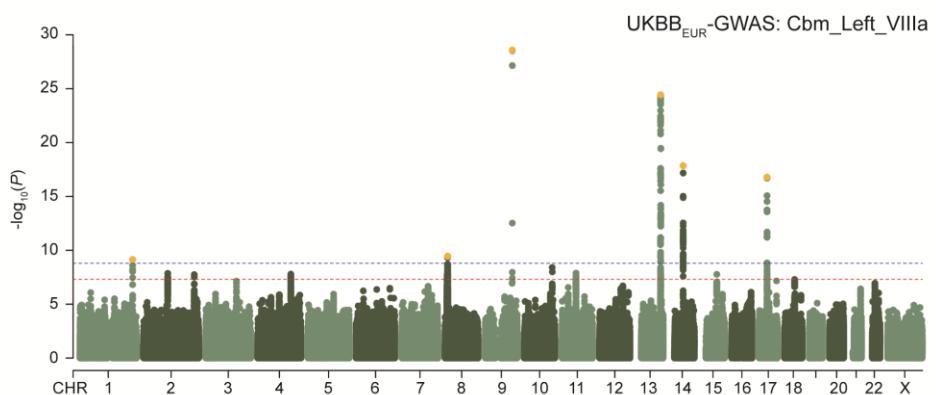
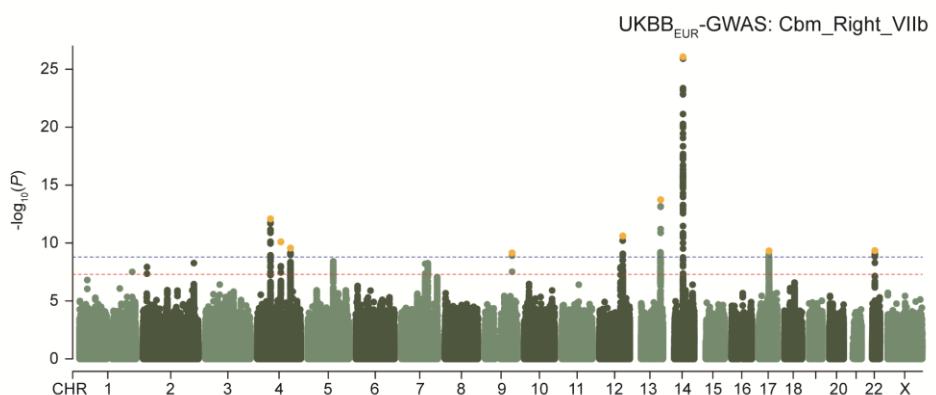
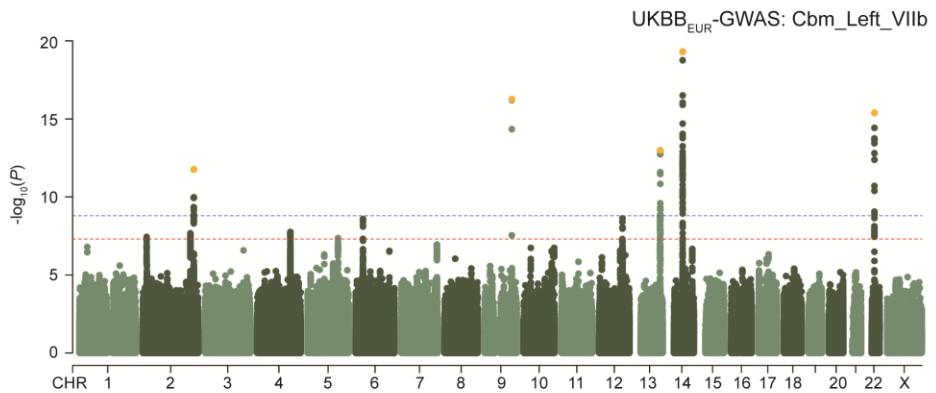
Fig. S4 | Genetic discovery of male-specific univariate GWAS. The Manhattan plots show the results from male-specific univariate GWASs (UKBB_{EUR}-GWAS, ABCD_{EUR}-GWAS, CHIMGEN_{EAS}-GWAS, ABCD_{AFR}-GWAS, EUR-GWAS, EUR-EAS-GWAS, and EUR-EAS-AFR-GWAS) on the 31 cerebellar volumetric traits. Each plot shows genome-wide (red dashed line: $P = 5 \times 10^{-8}$) and study-wide (blue dashed line: $P = 1.61 \times 10^{-9}$) significant associations. Each point represents a single genetic variant plotted according to its genomic position (x axis) and its $-\log_{10}(P)$ value for the association (y axis). Each yellow point represents an independent locus with study-wide significance ($P = 1.61 \times 10^{-9}$). Abbreviations: ABCD, the Adolescent Brain Cognitive Development study; AFR, African ancestry; CHIMGEN, the Chinese Imaging Genetics study; EAS, East Asian ancestry; EUR, European ancestry; GWAS, genome-wide association study; UKBB, the UK Biobank study.

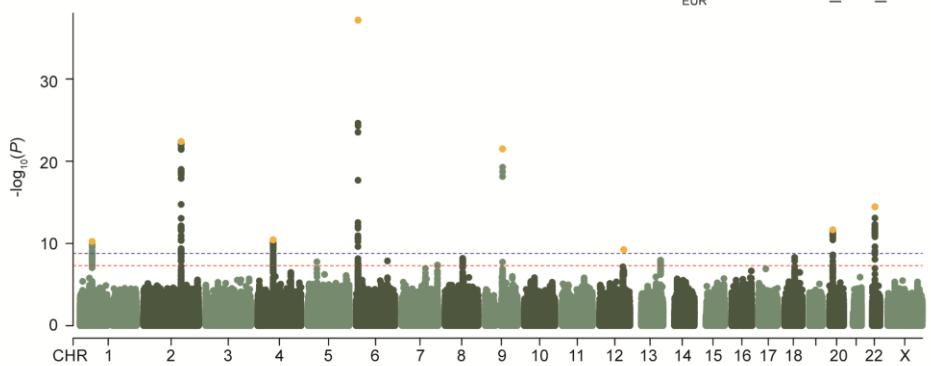
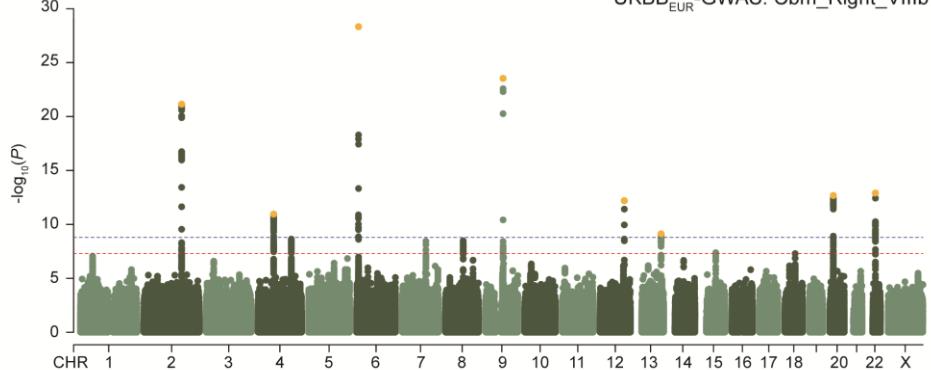
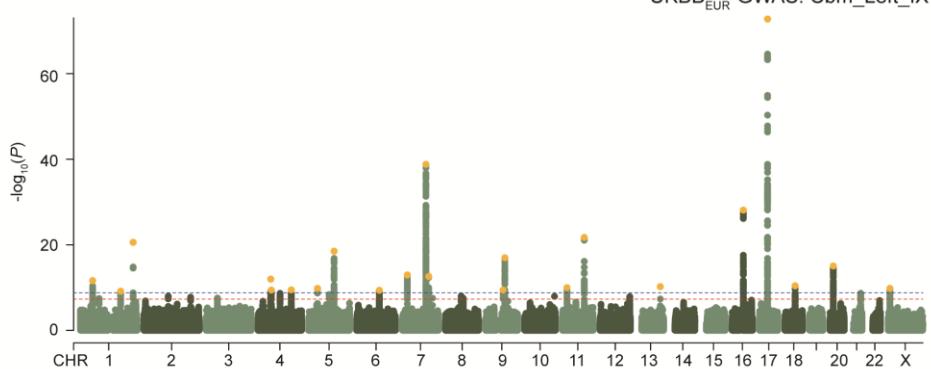
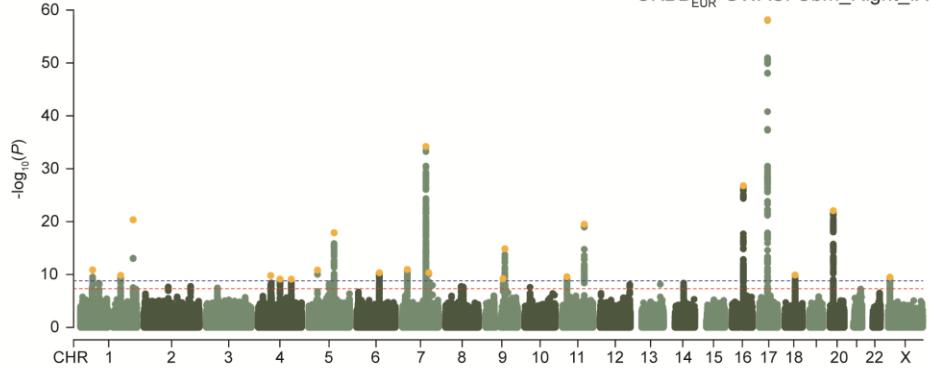


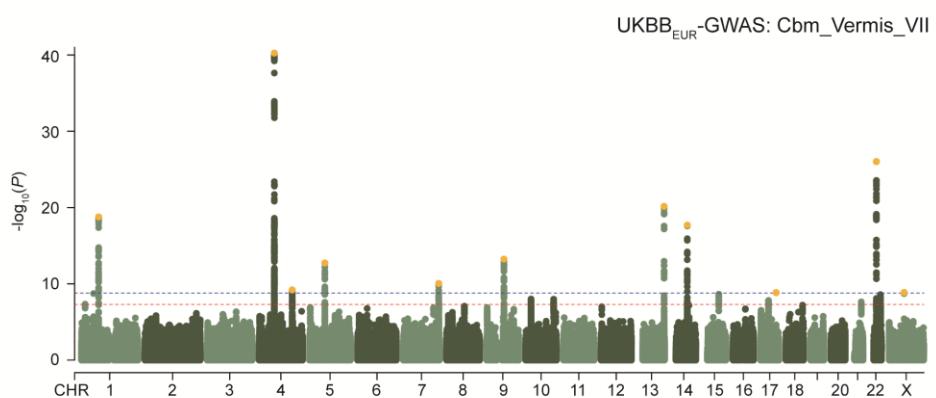
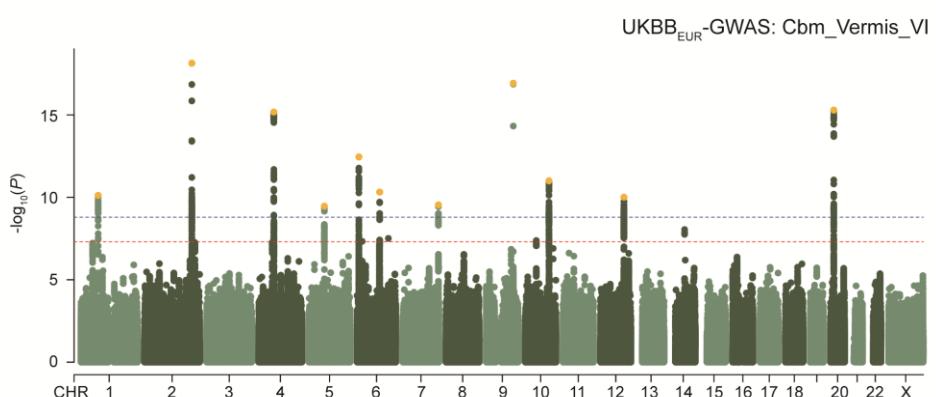
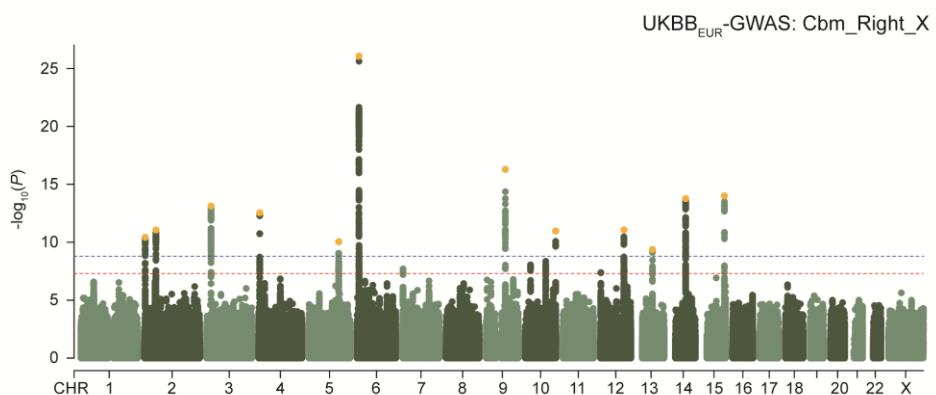
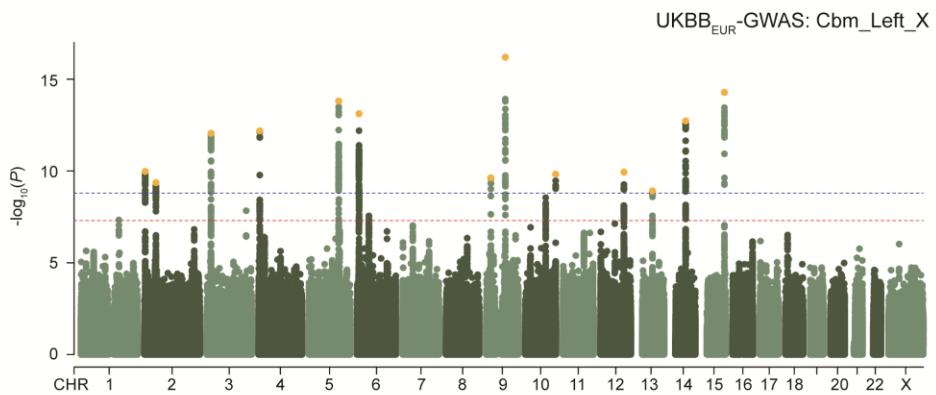




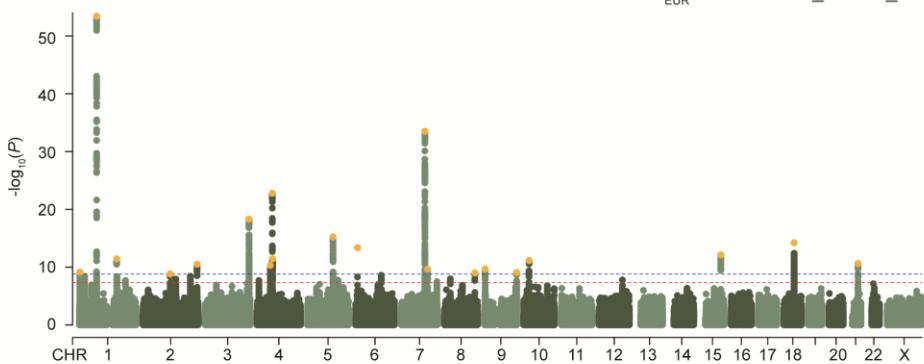




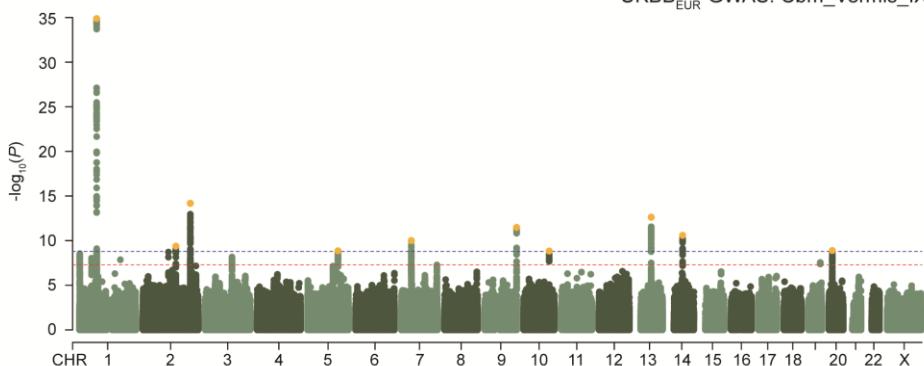
UKBB_{EUR}-GWAS: Cbm_Left_VIIIbUKBB_{EUR}-GWAS: Cbm_Right_VIIIbUKBB_{EUR}-GWAS: Cbm_Left_IXUKBB_{EUR}-GWAS: Cbm_Right_IX



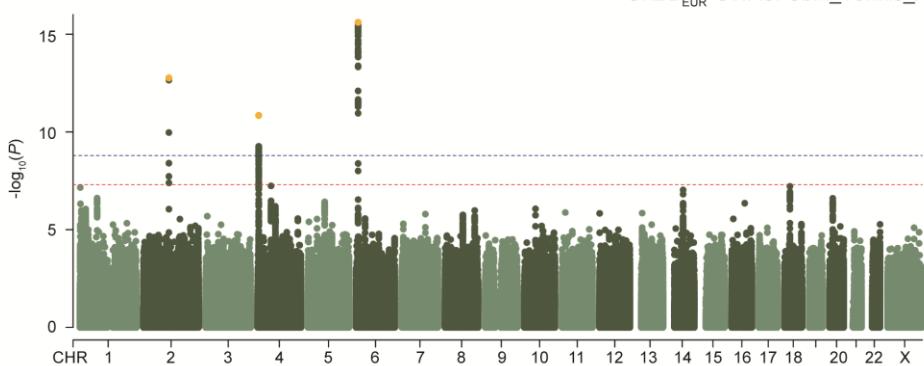
UKBB_{EUR}-GWAS: Cbm_Vermis_VIII



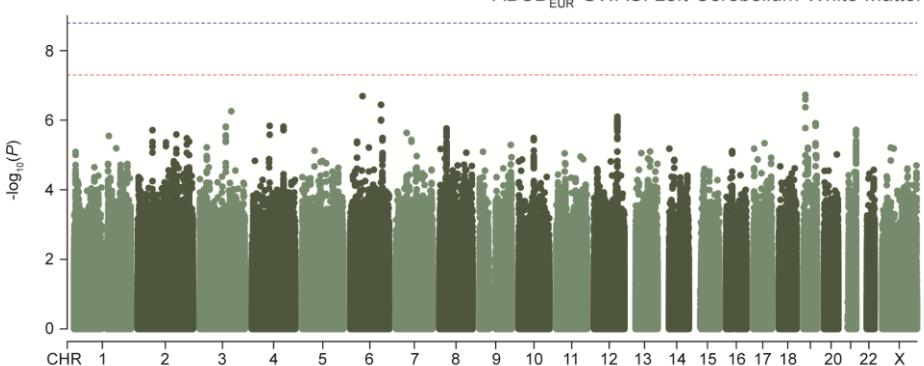
UKBB_{EUR}-GWAS: Cbm_Vermis_IX

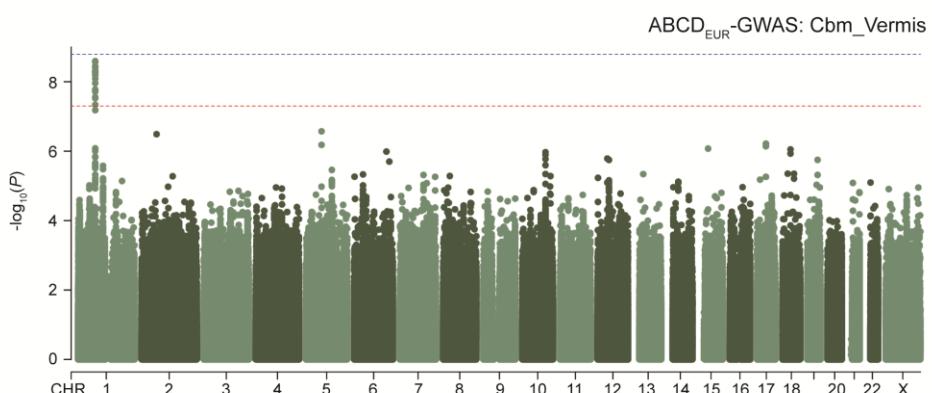
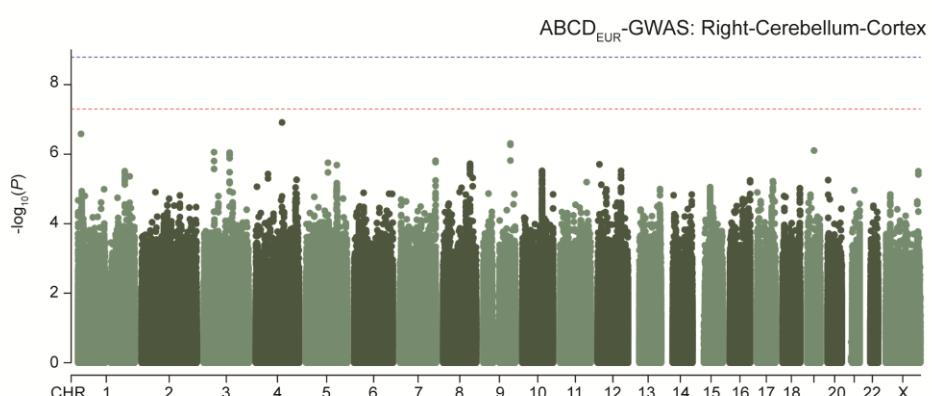
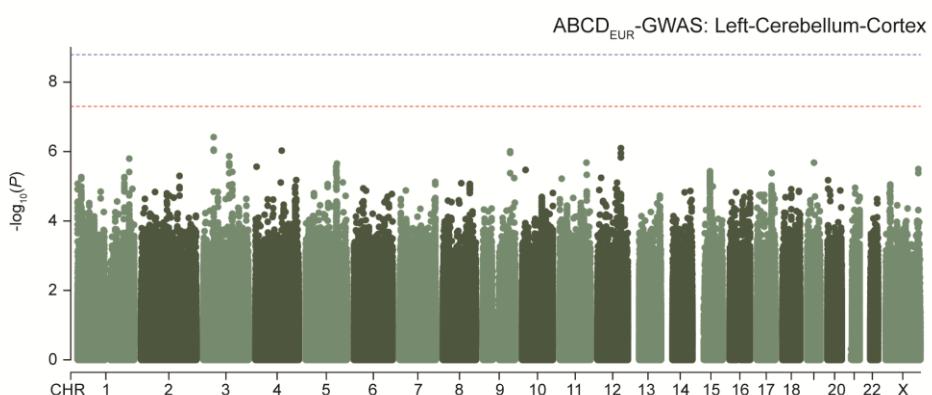
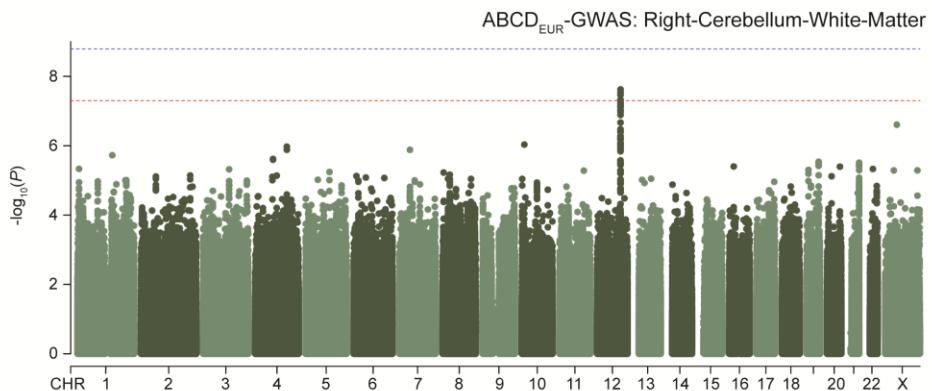


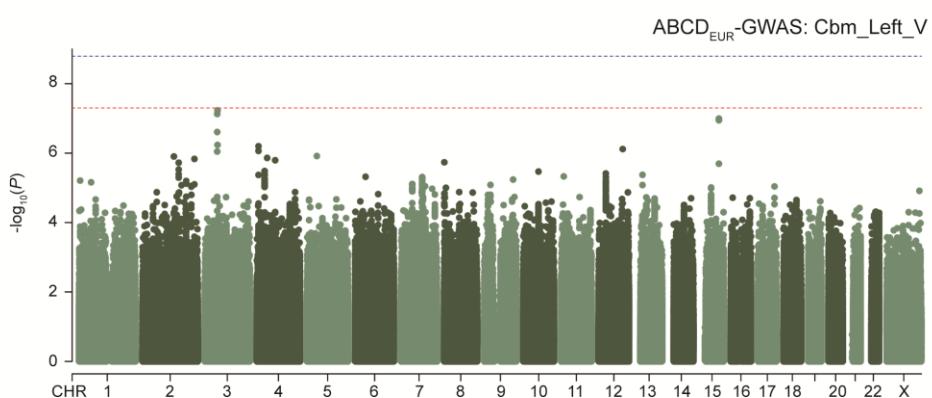
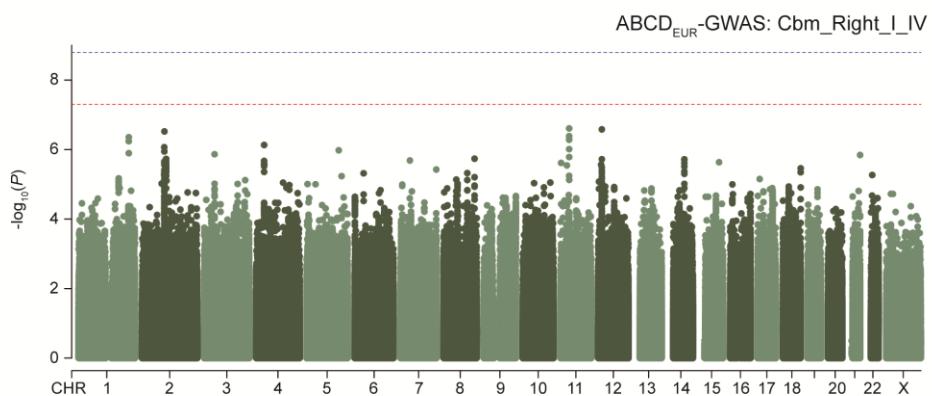
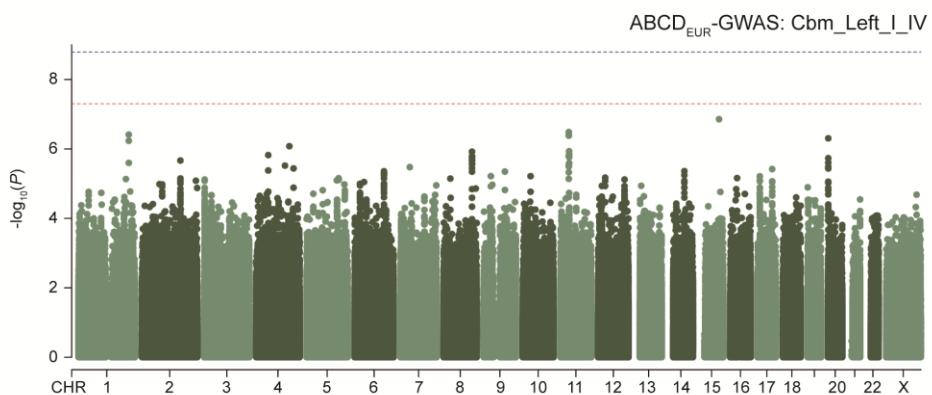
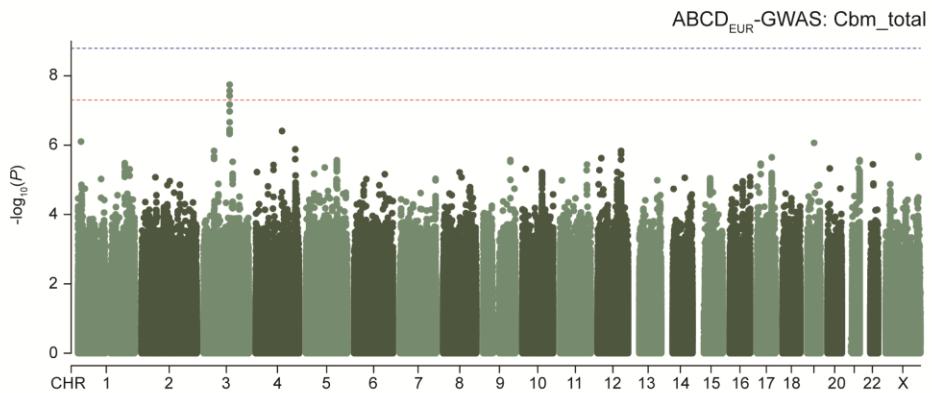
UKBB_{EUR}-GWAS: Cbm_Vermis_X

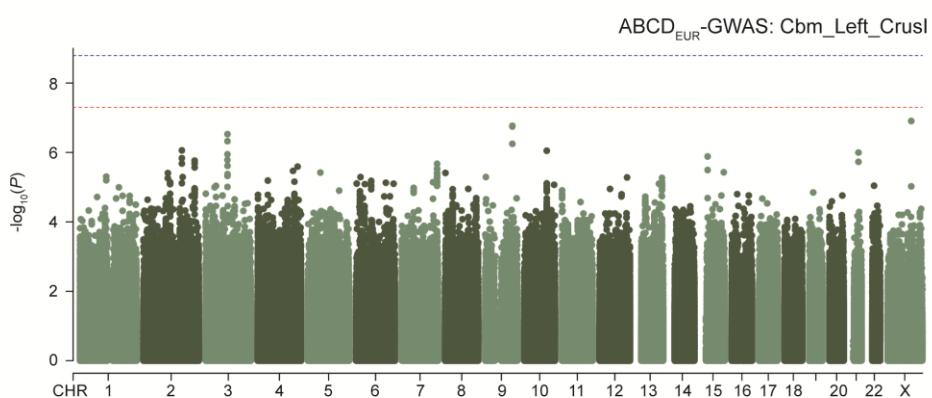
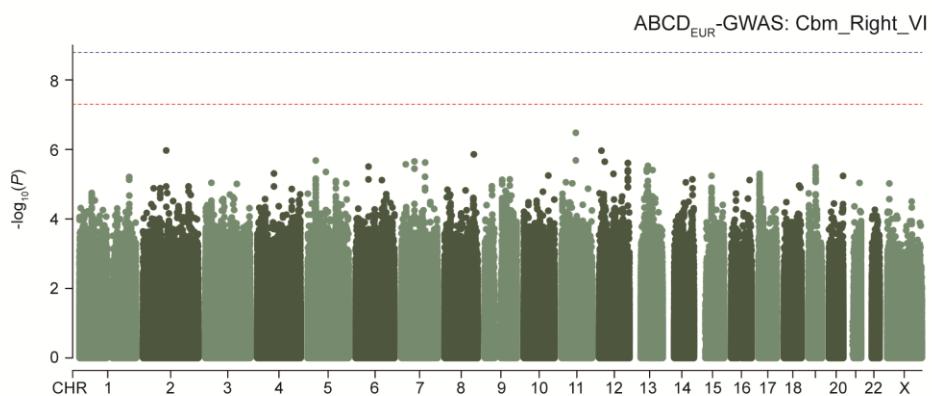
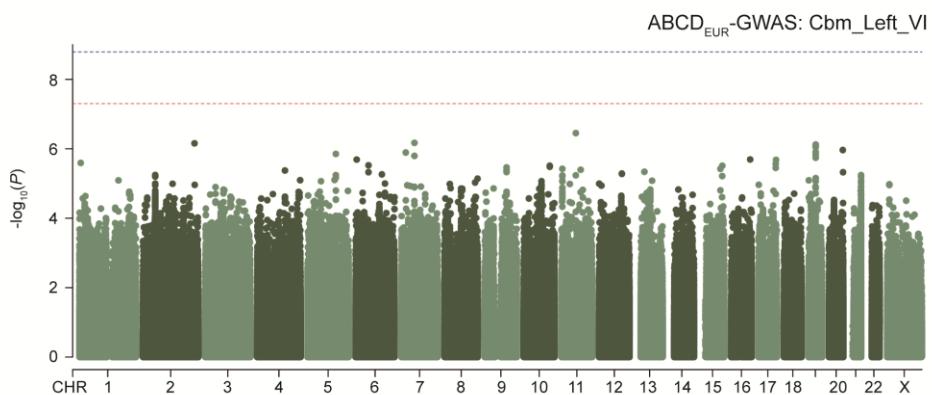
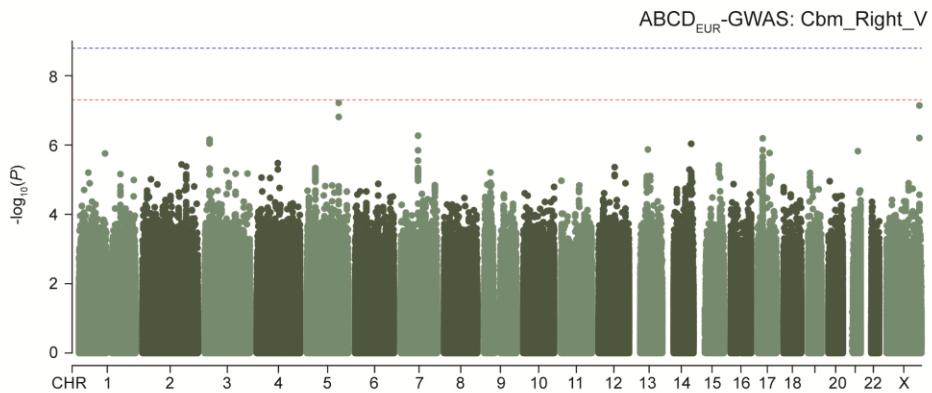


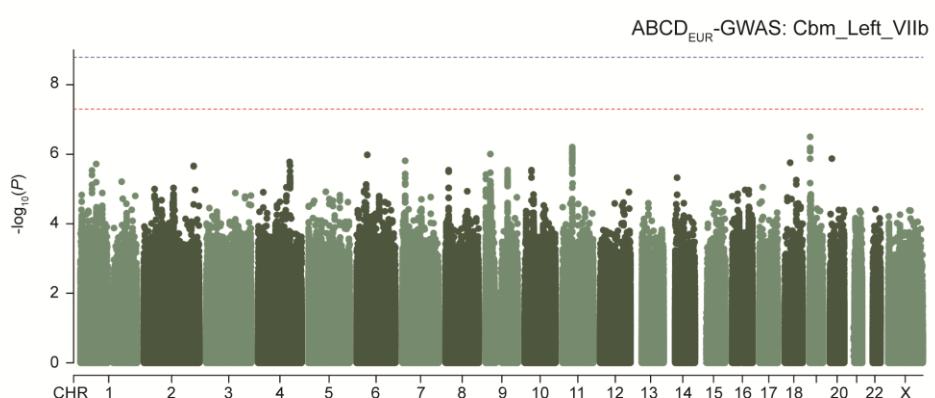
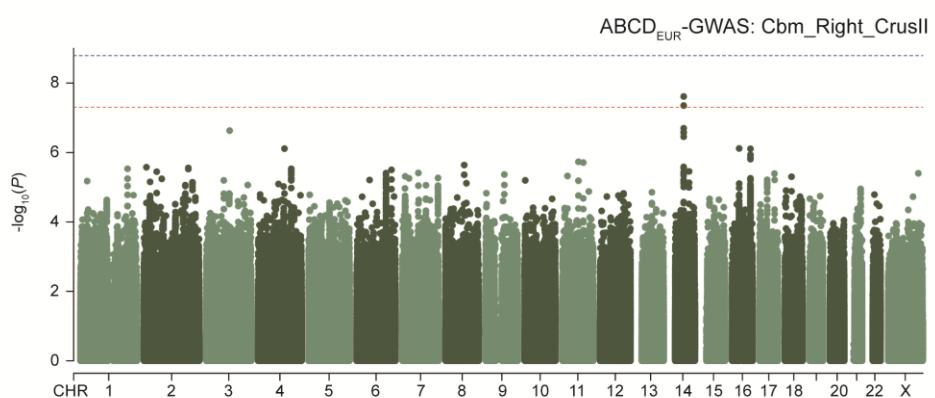
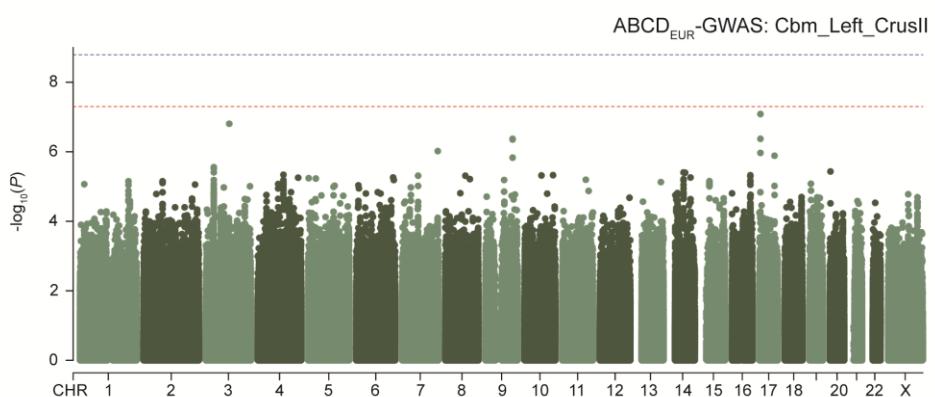
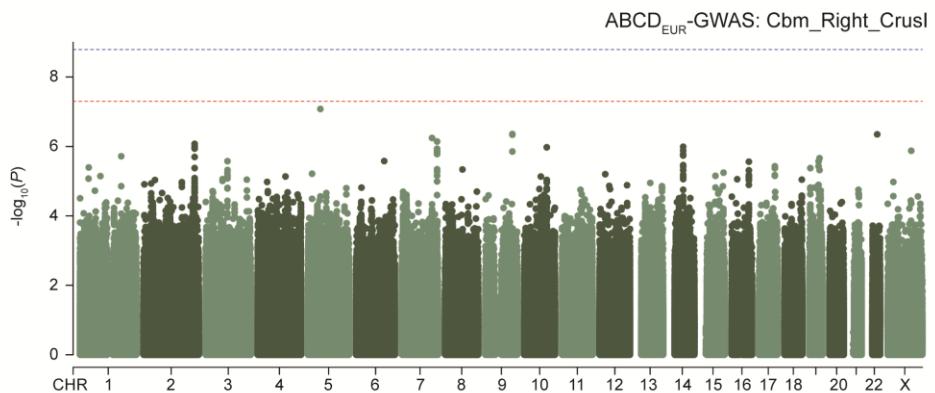
ABCD_{EUR}-GWAS: Left-Cerebellum-White-Matter

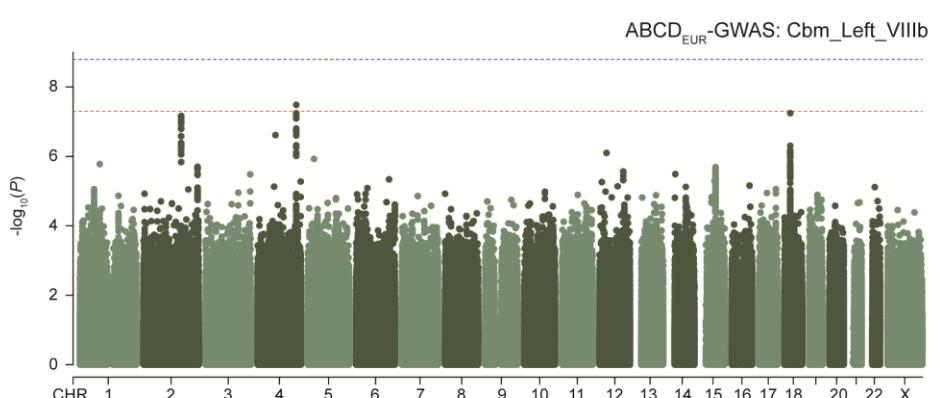
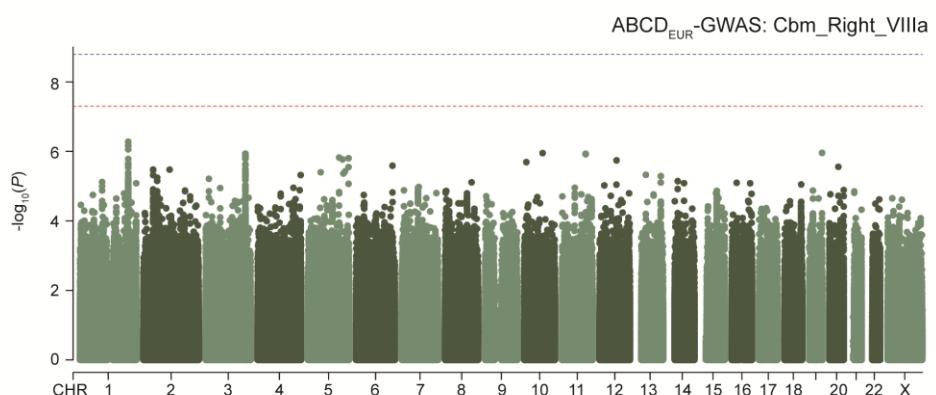
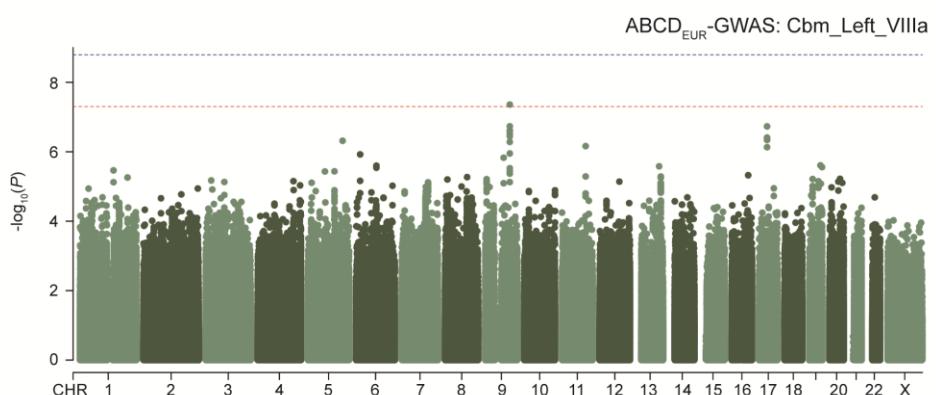
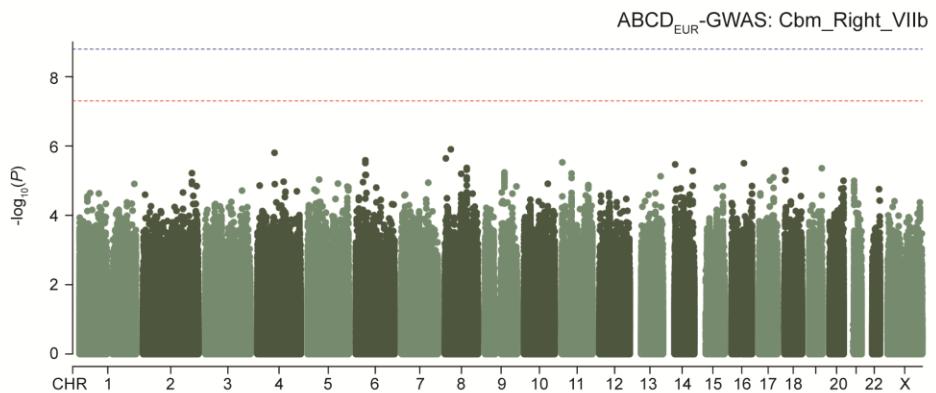


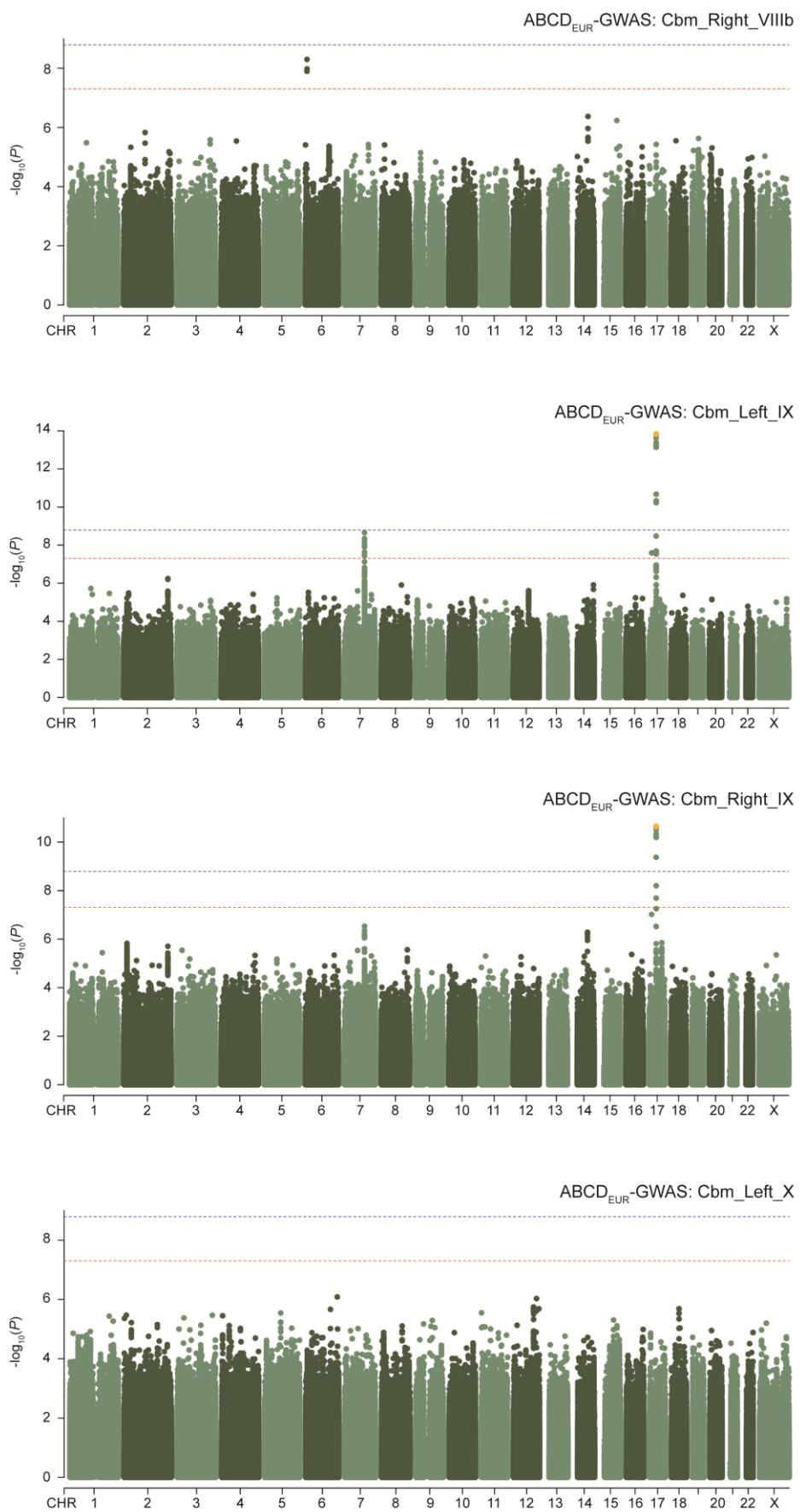


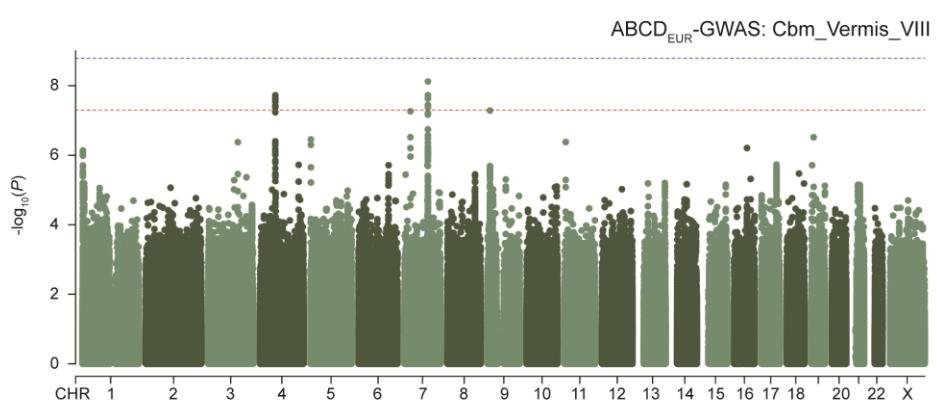
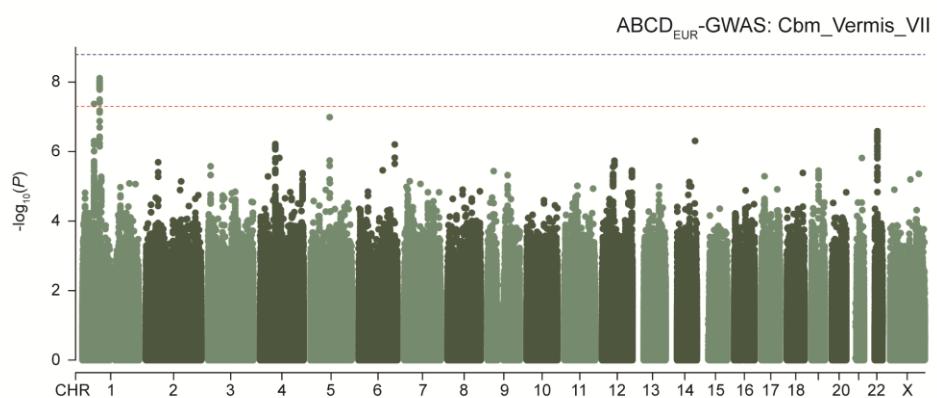
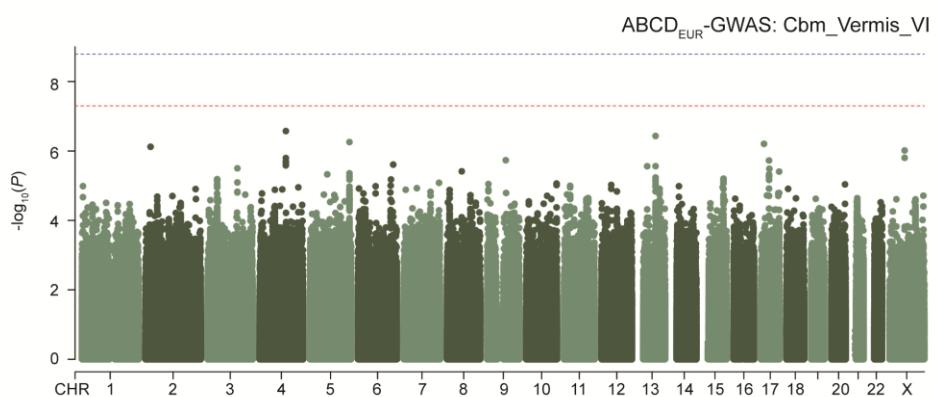
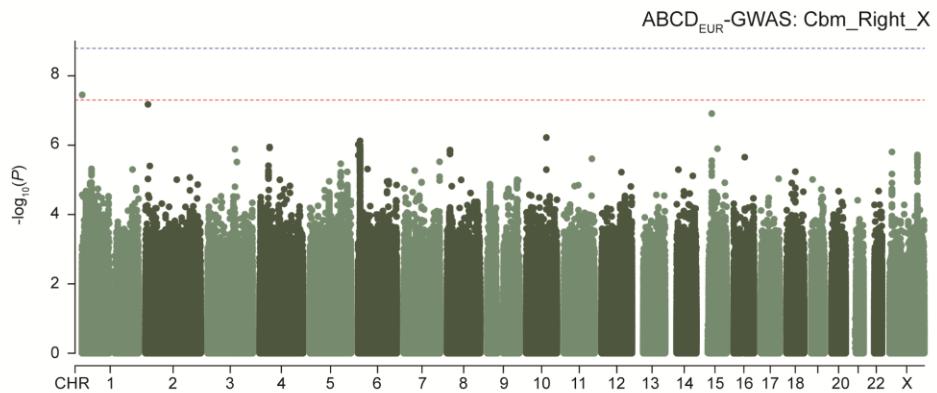


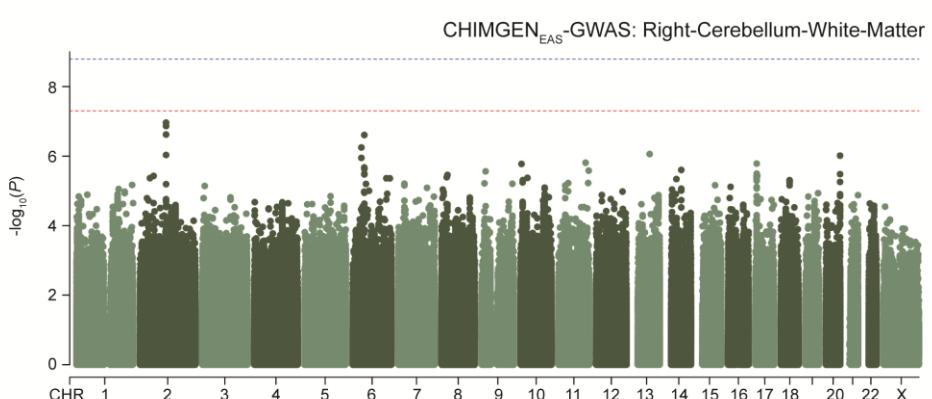
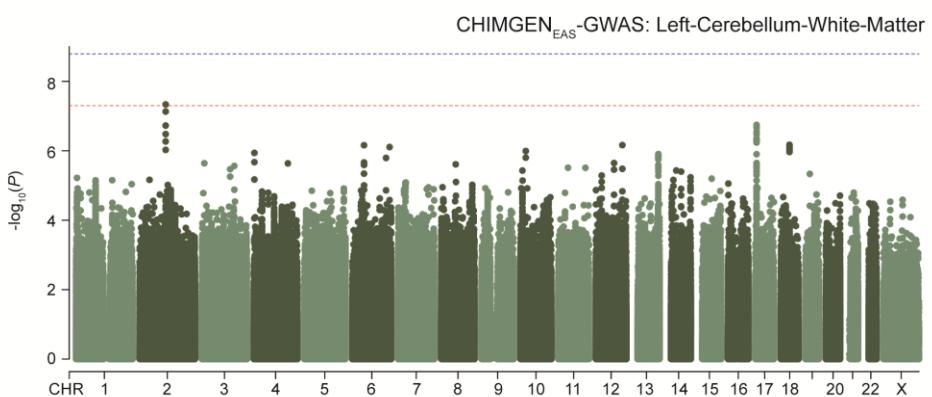
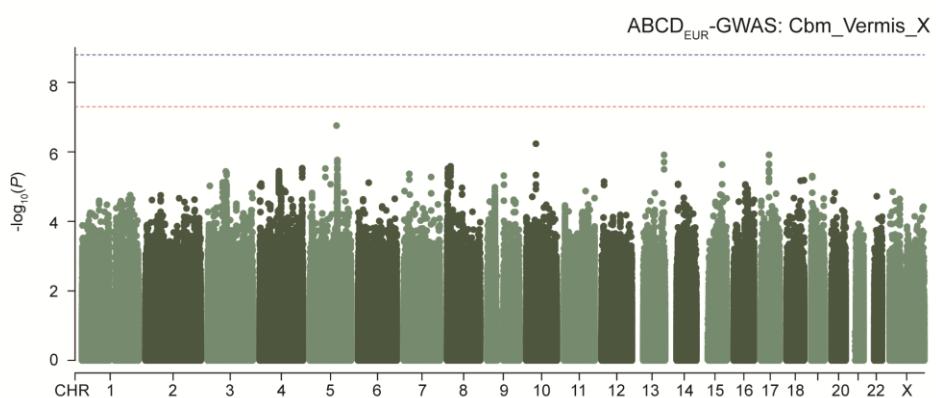
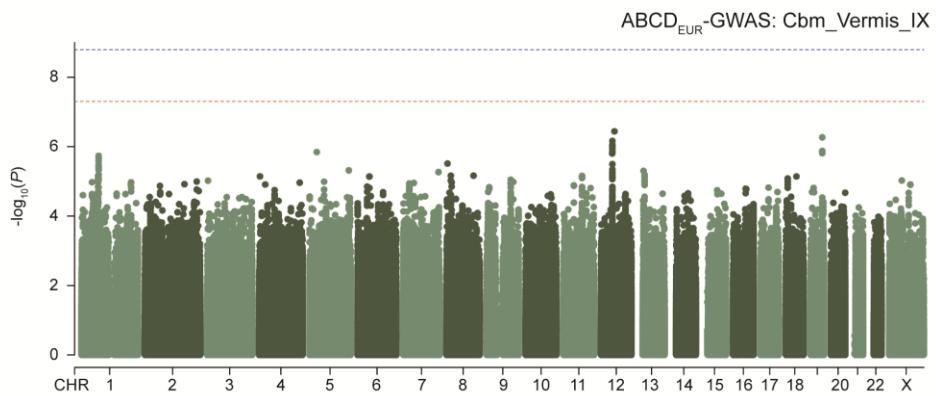


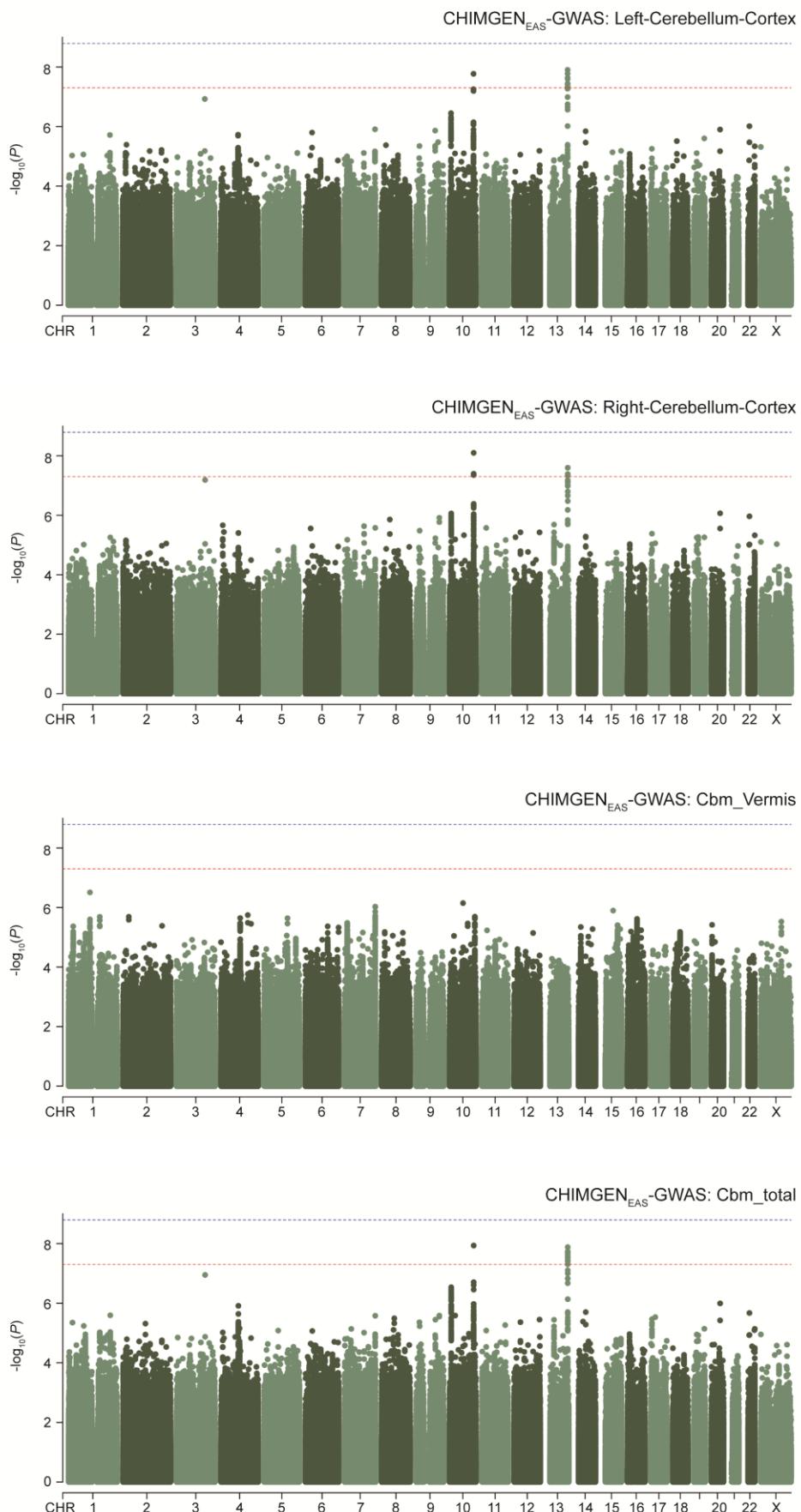


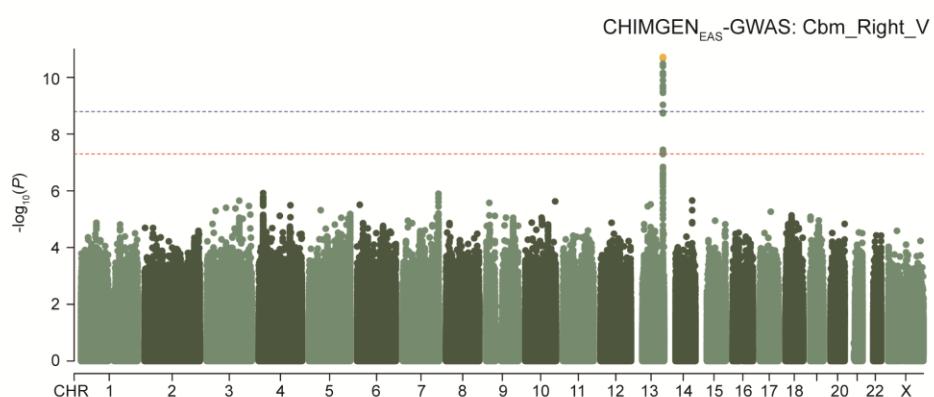
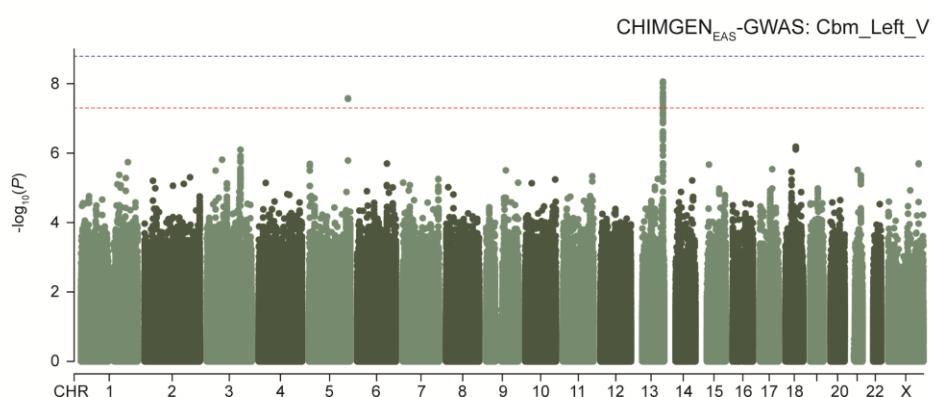
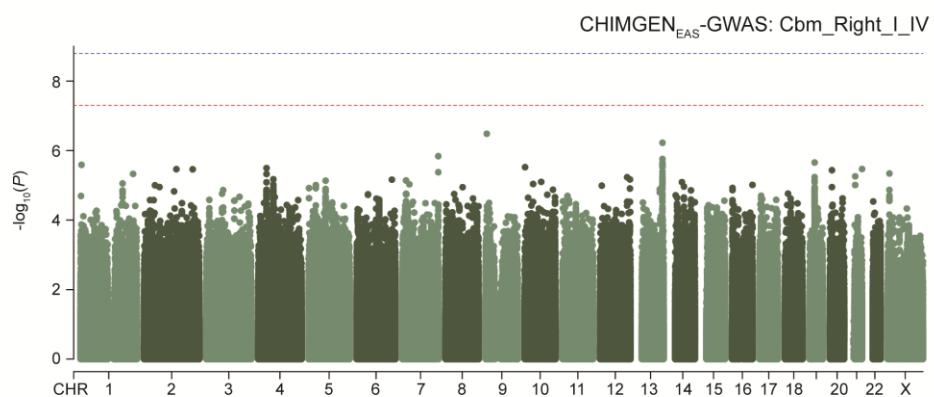
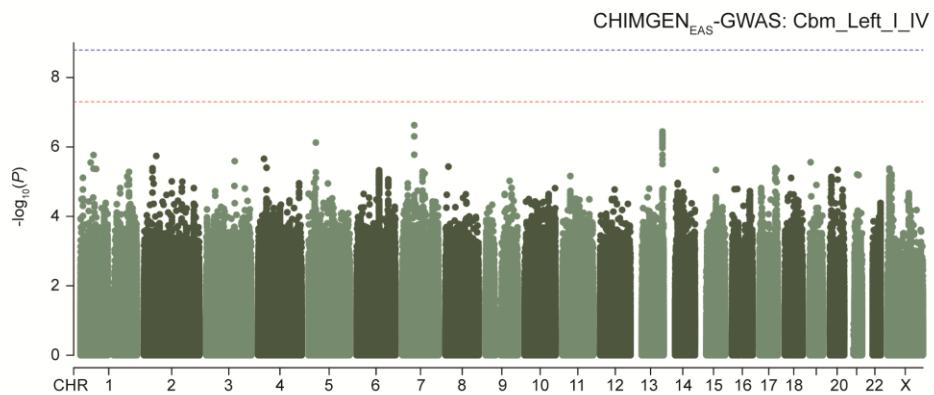


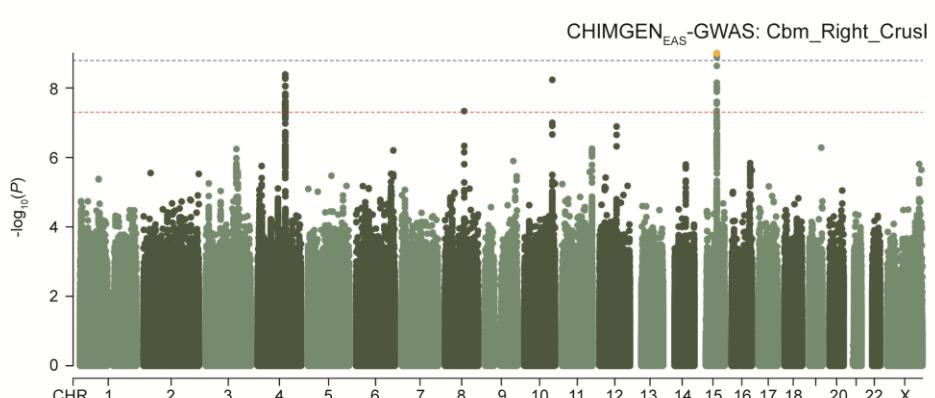
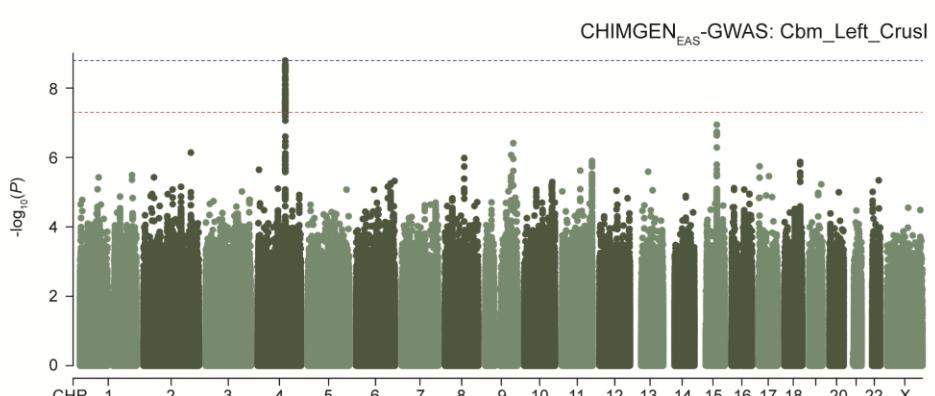
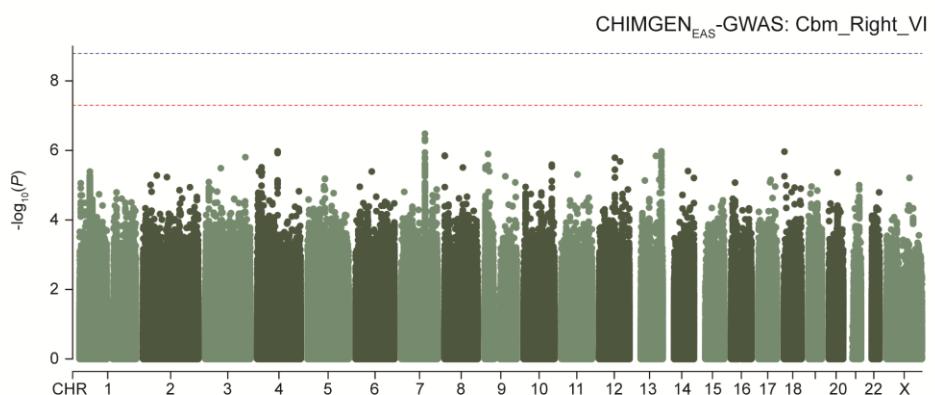
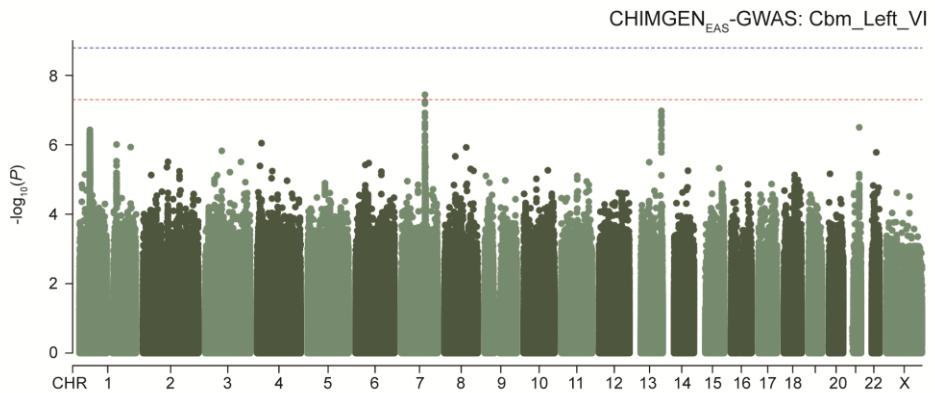


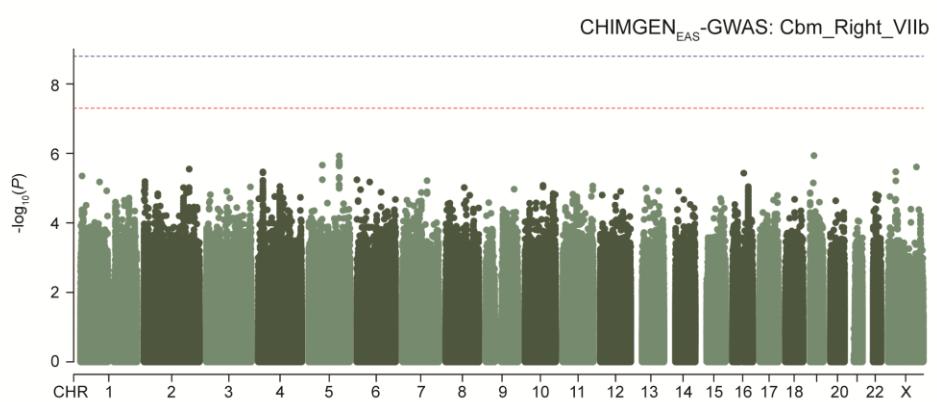
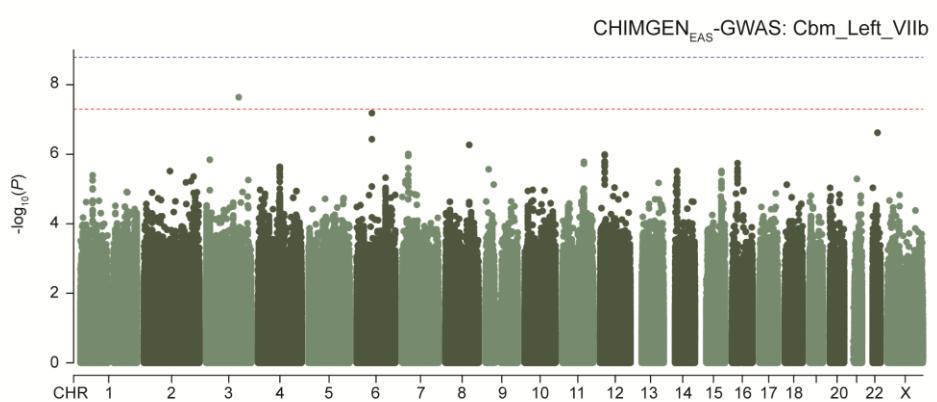
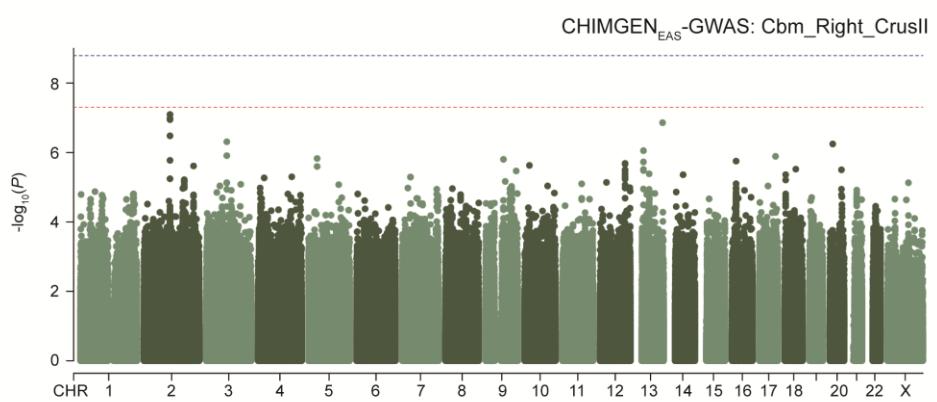
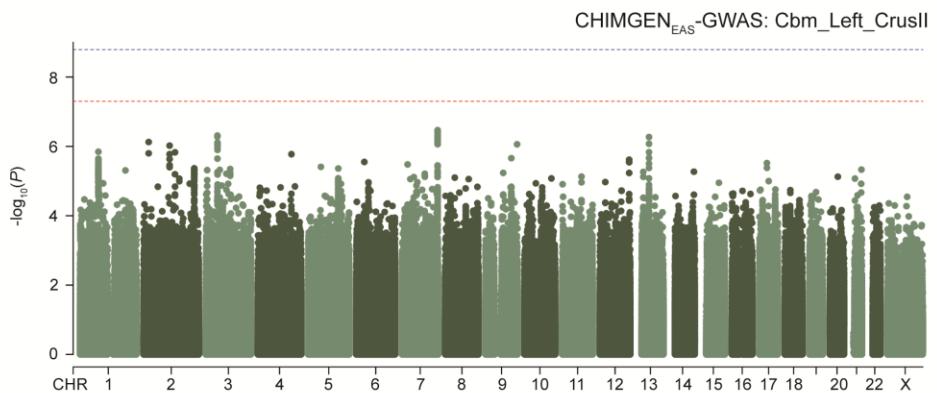


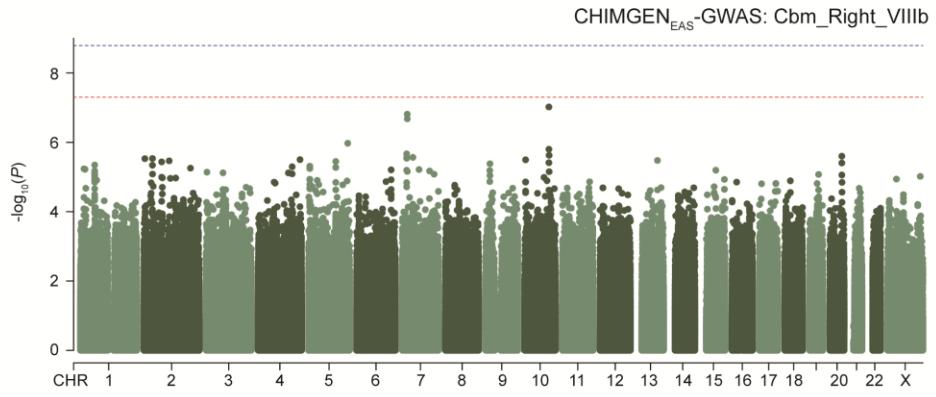
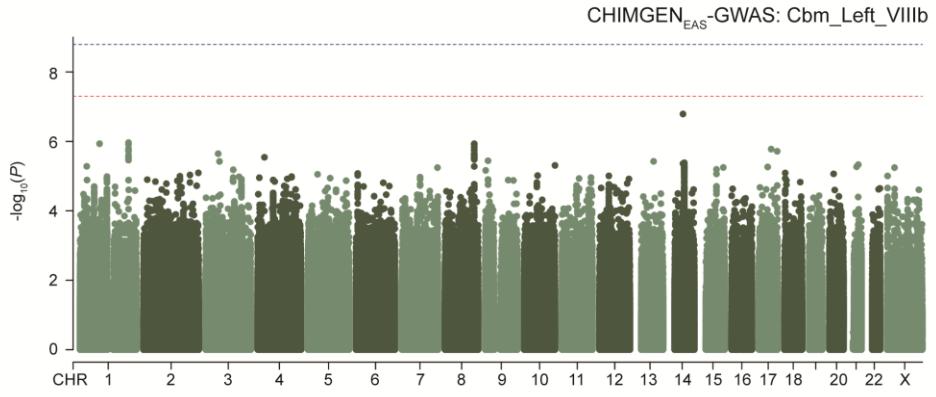
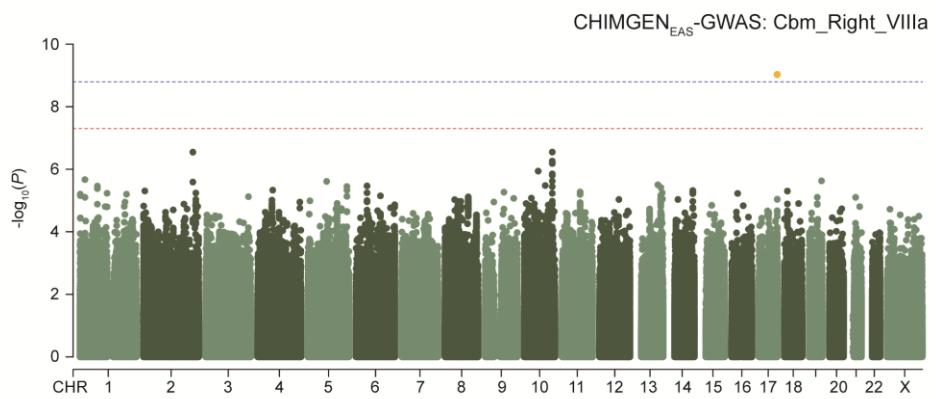
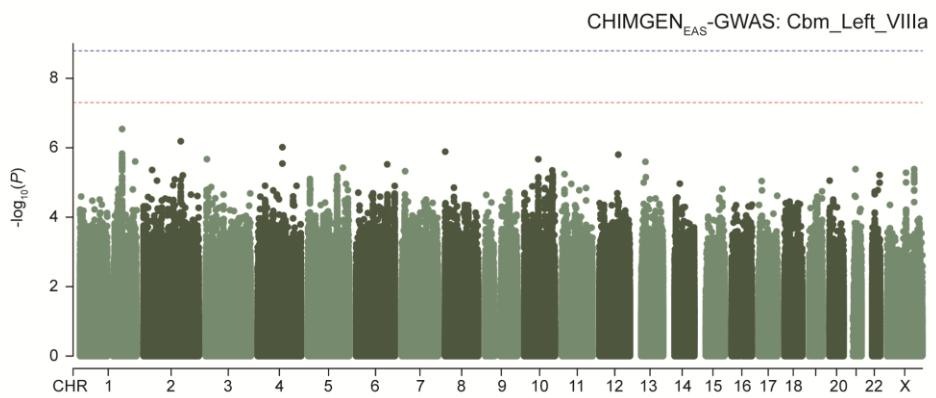


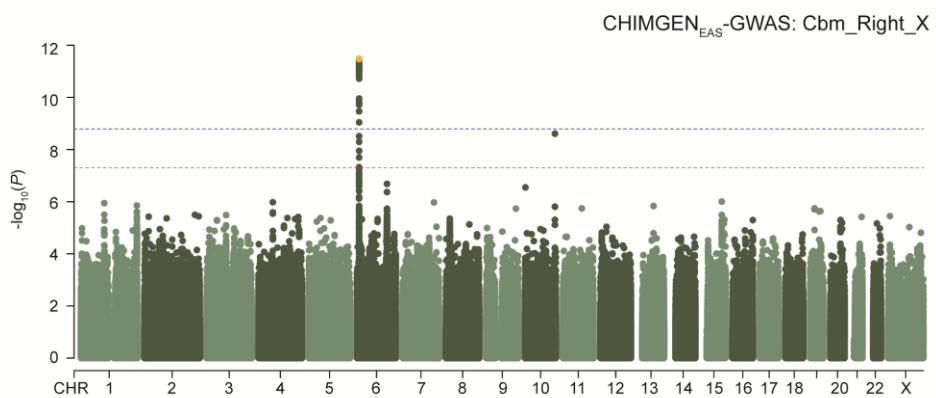
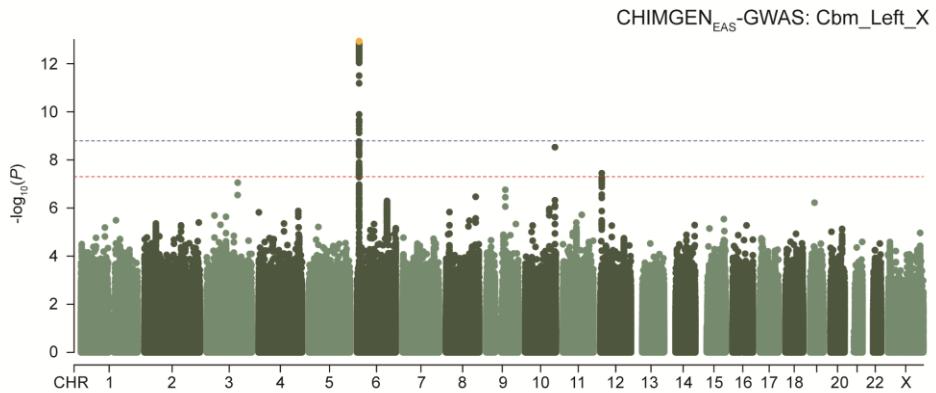
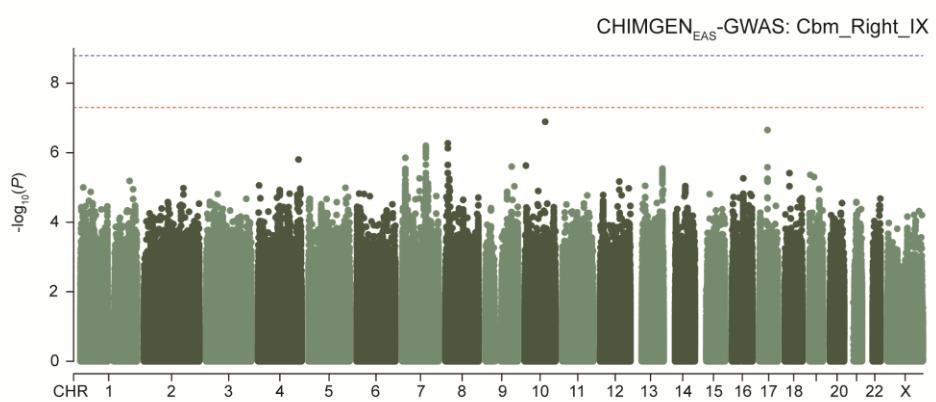
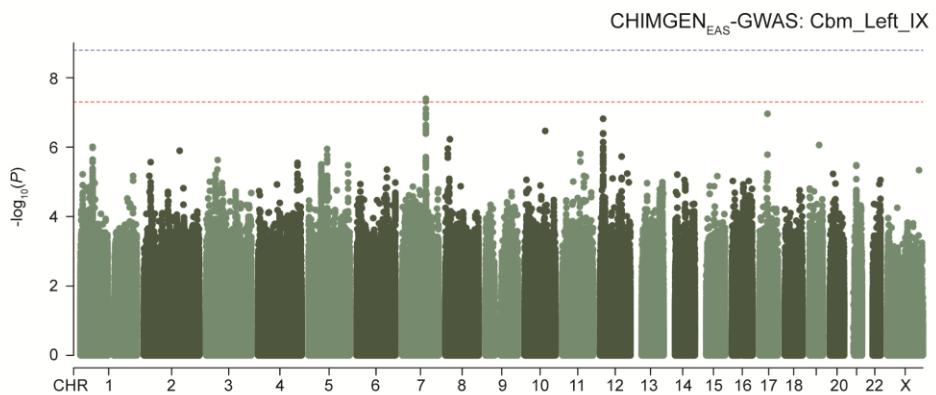


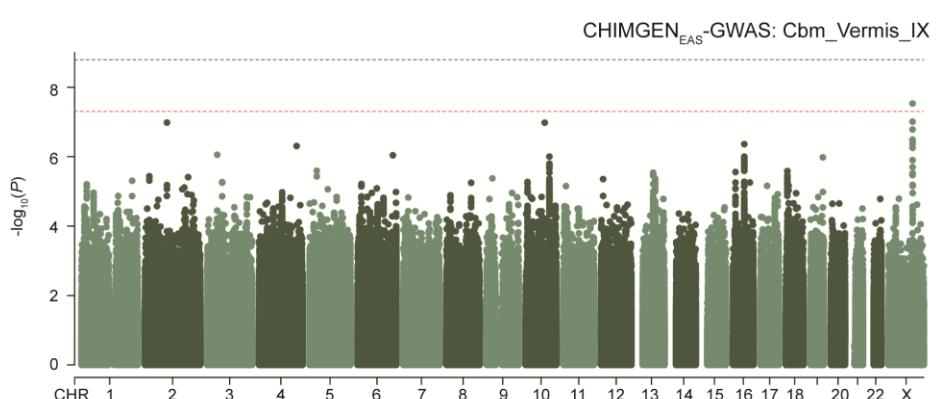
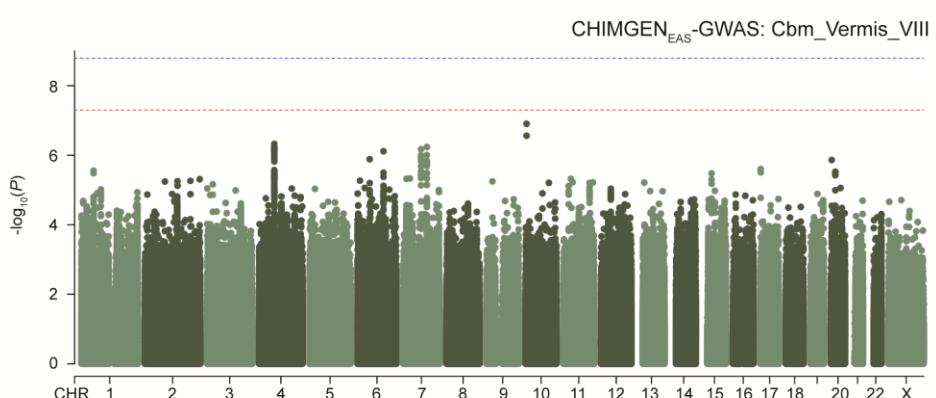
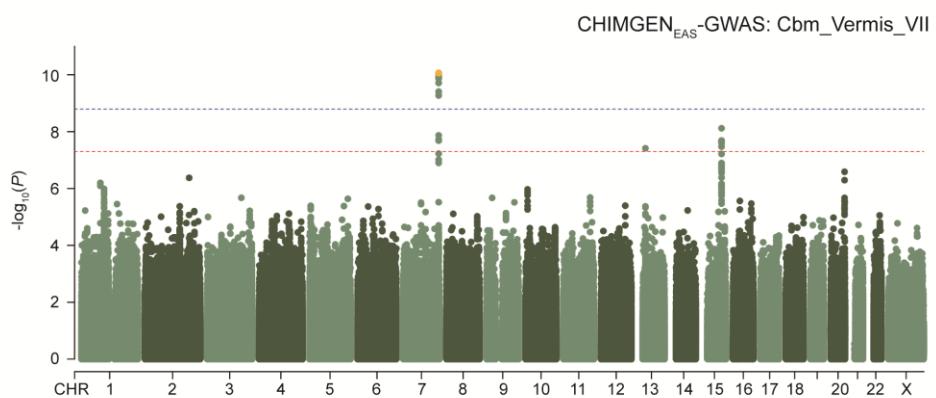
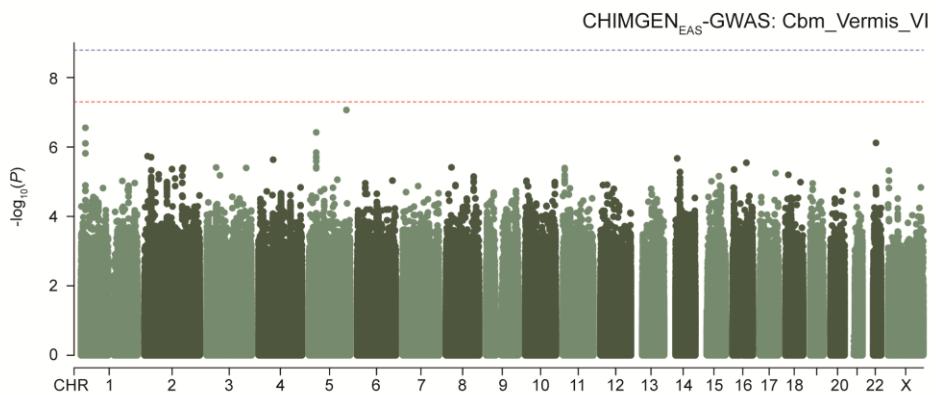


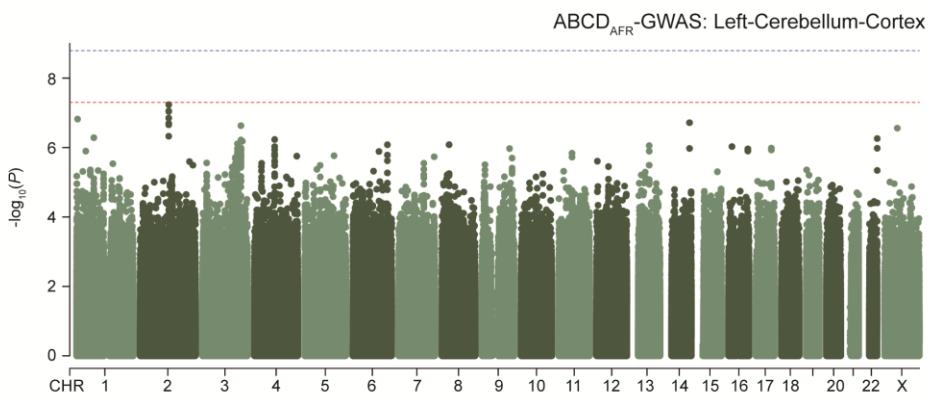
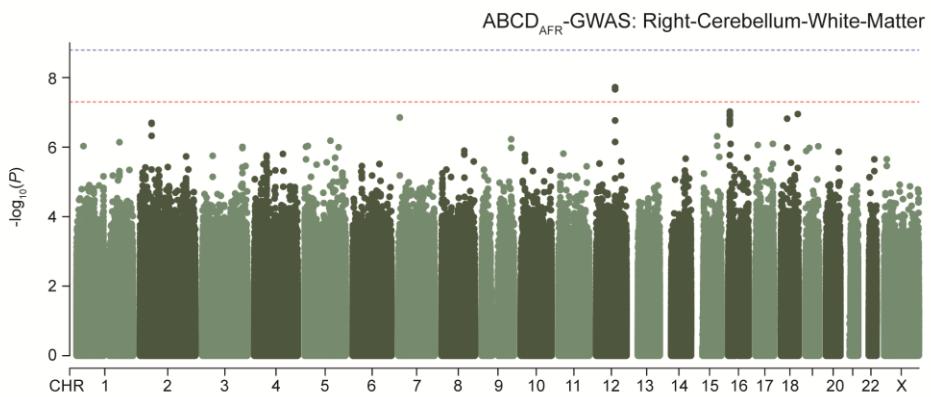
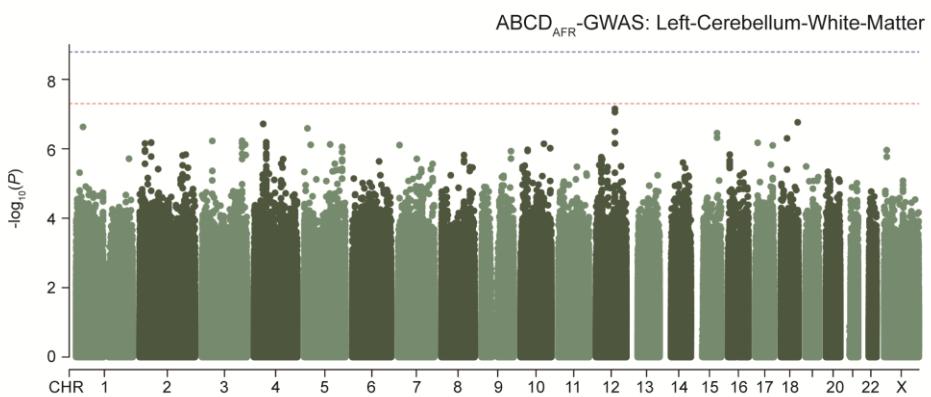
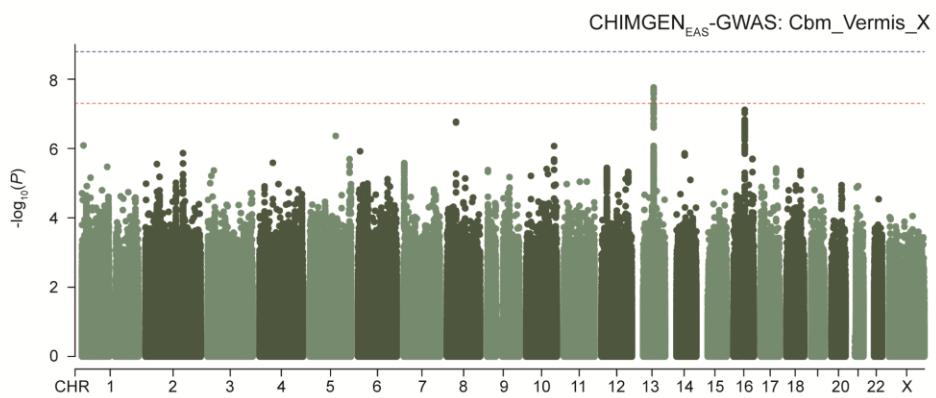




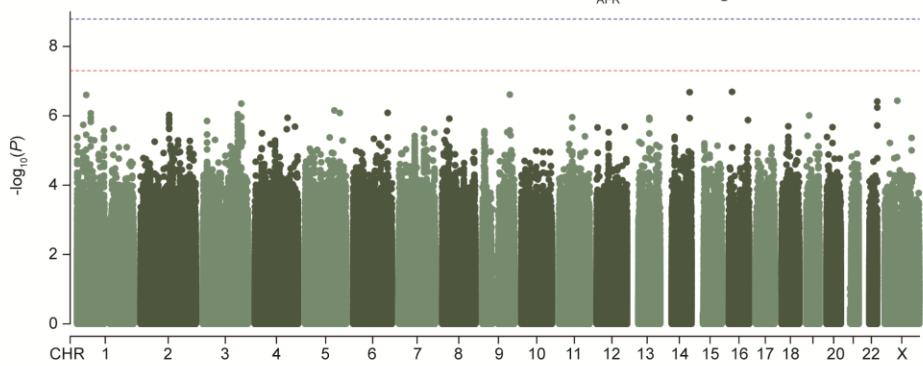




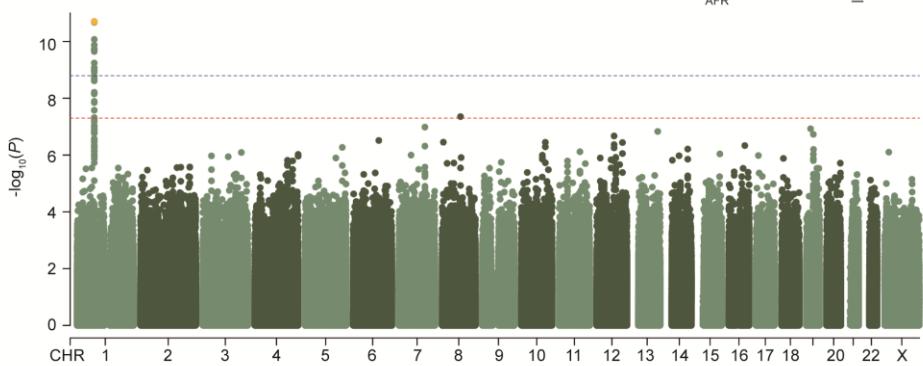




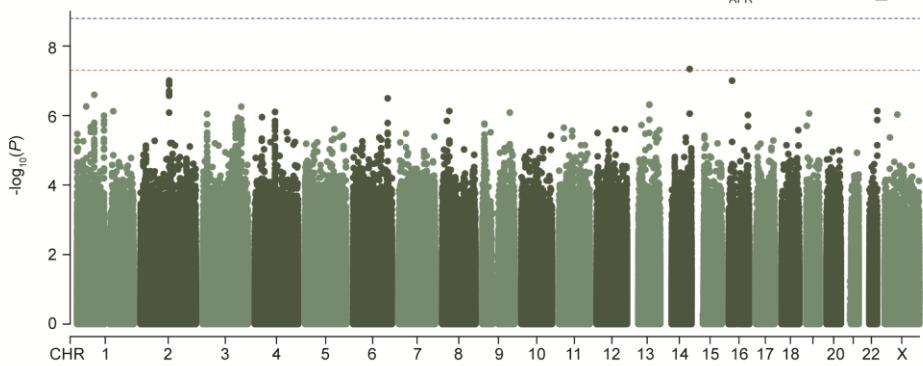
ABCD_{AFR}-GWAS: Right-Cerebellum-Cortex



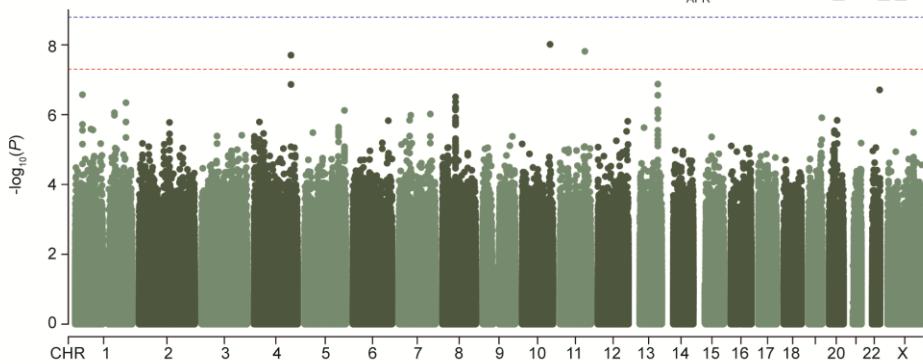
ABCD_{AFR}-GWAS: Cbm_Vermis

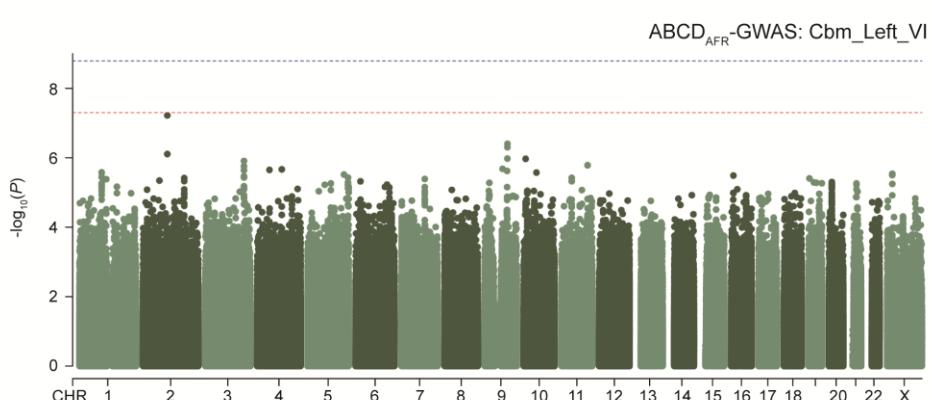
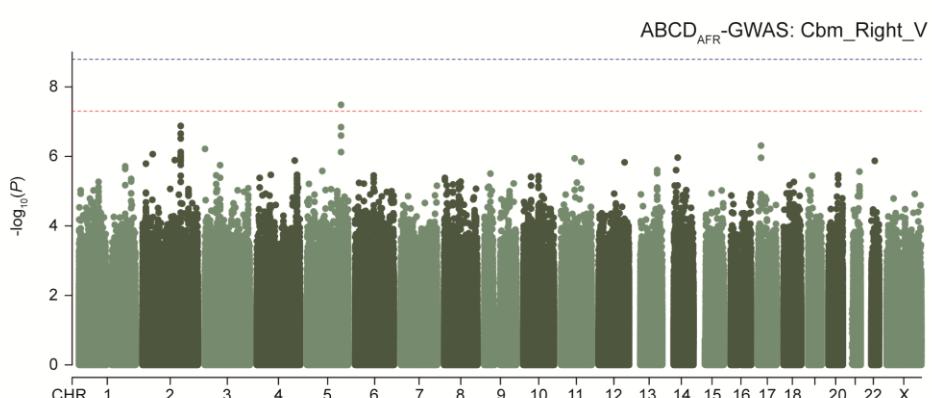
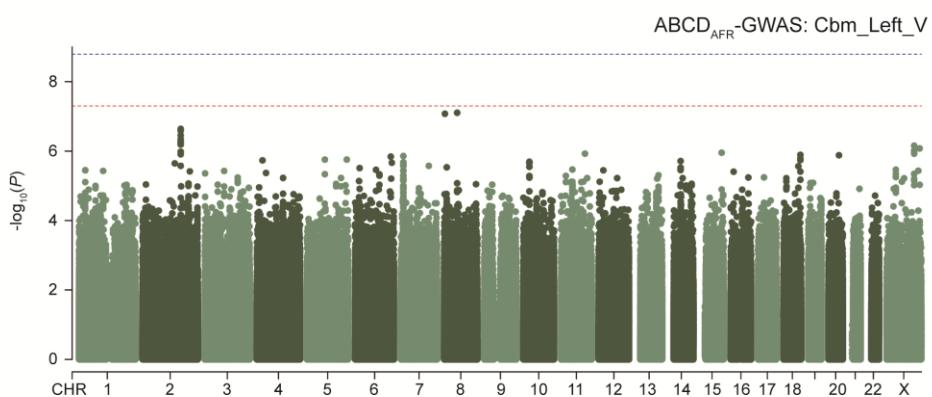
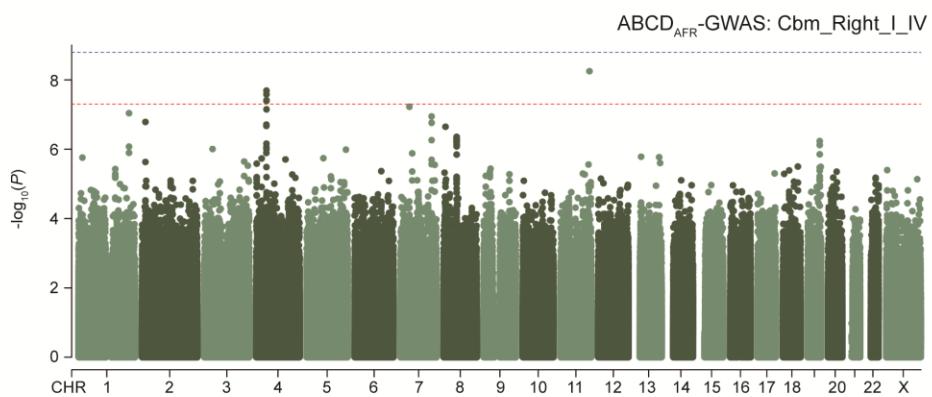


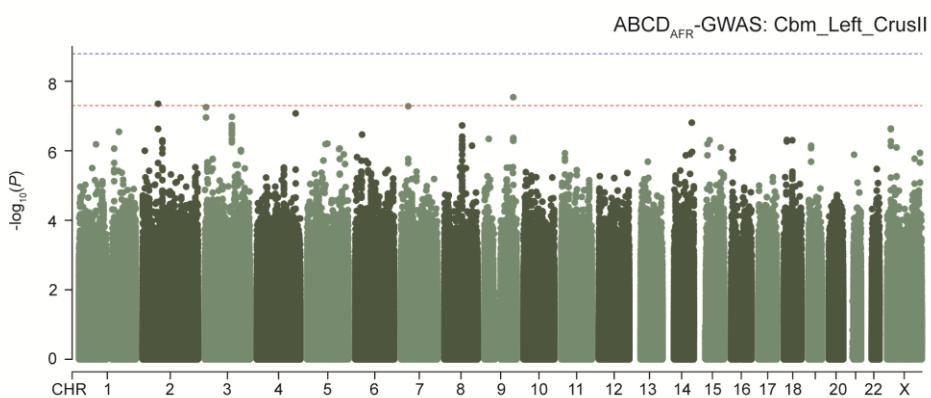
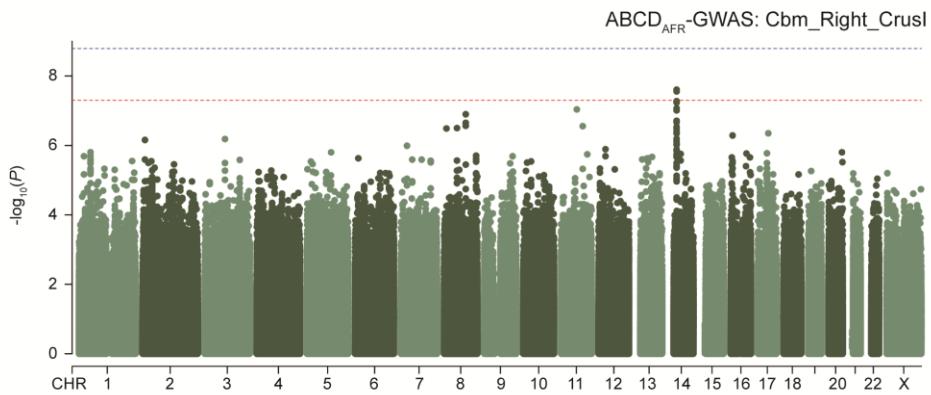
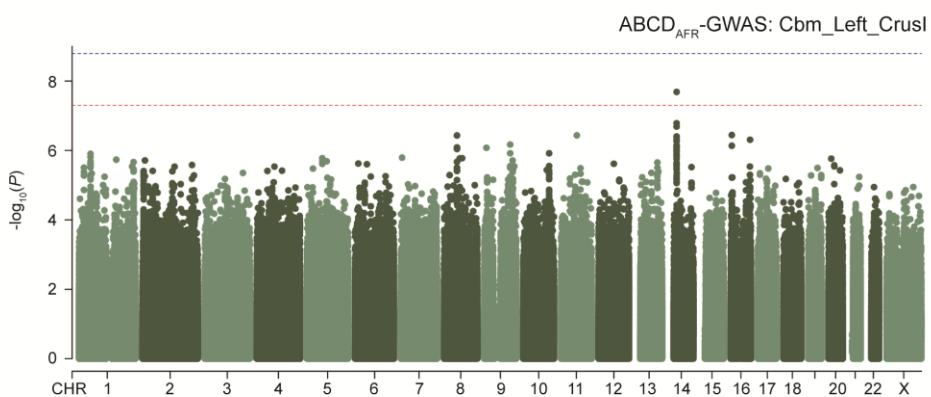
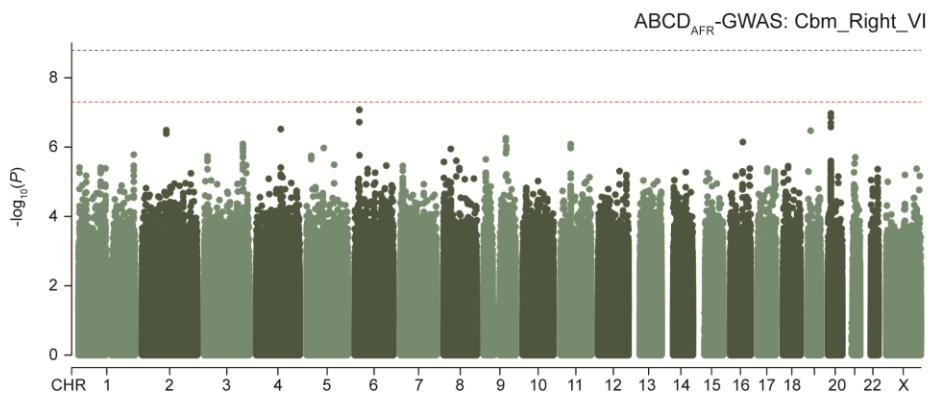
ABCD_{AFR}-GWAS: Cbm_total

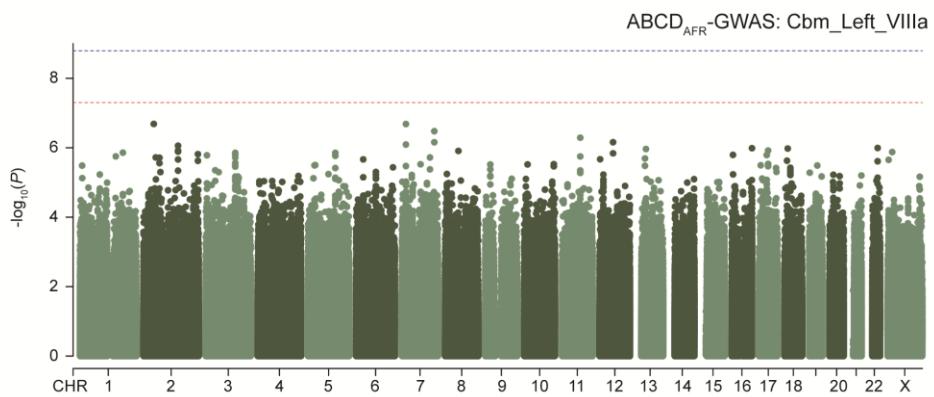
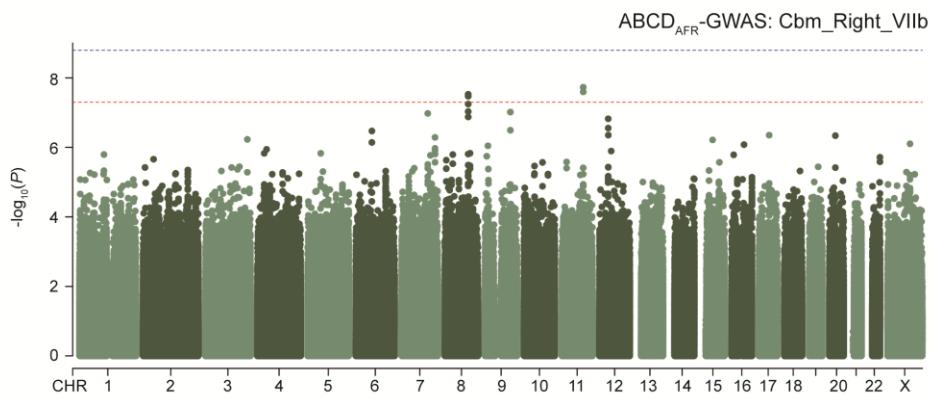
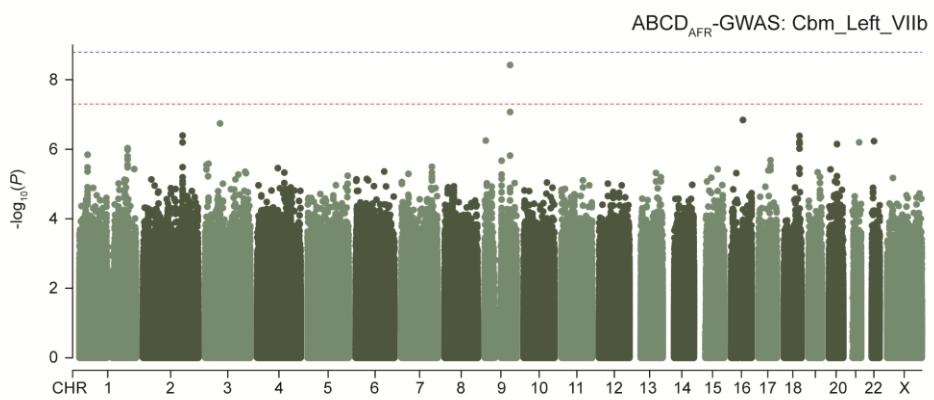
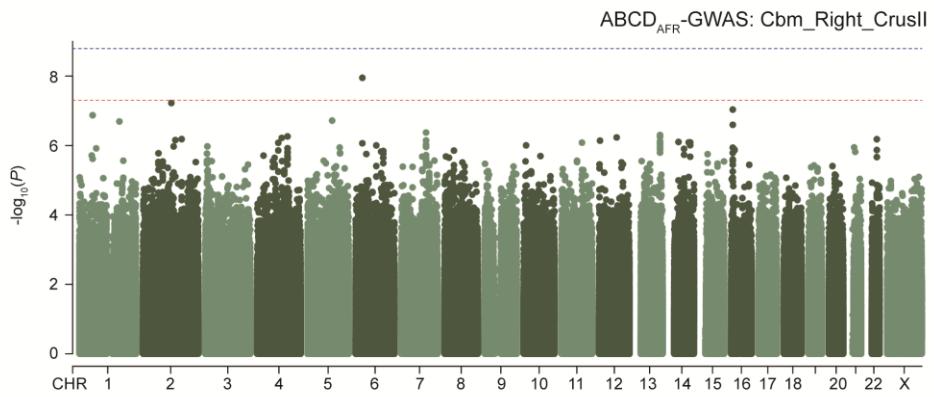


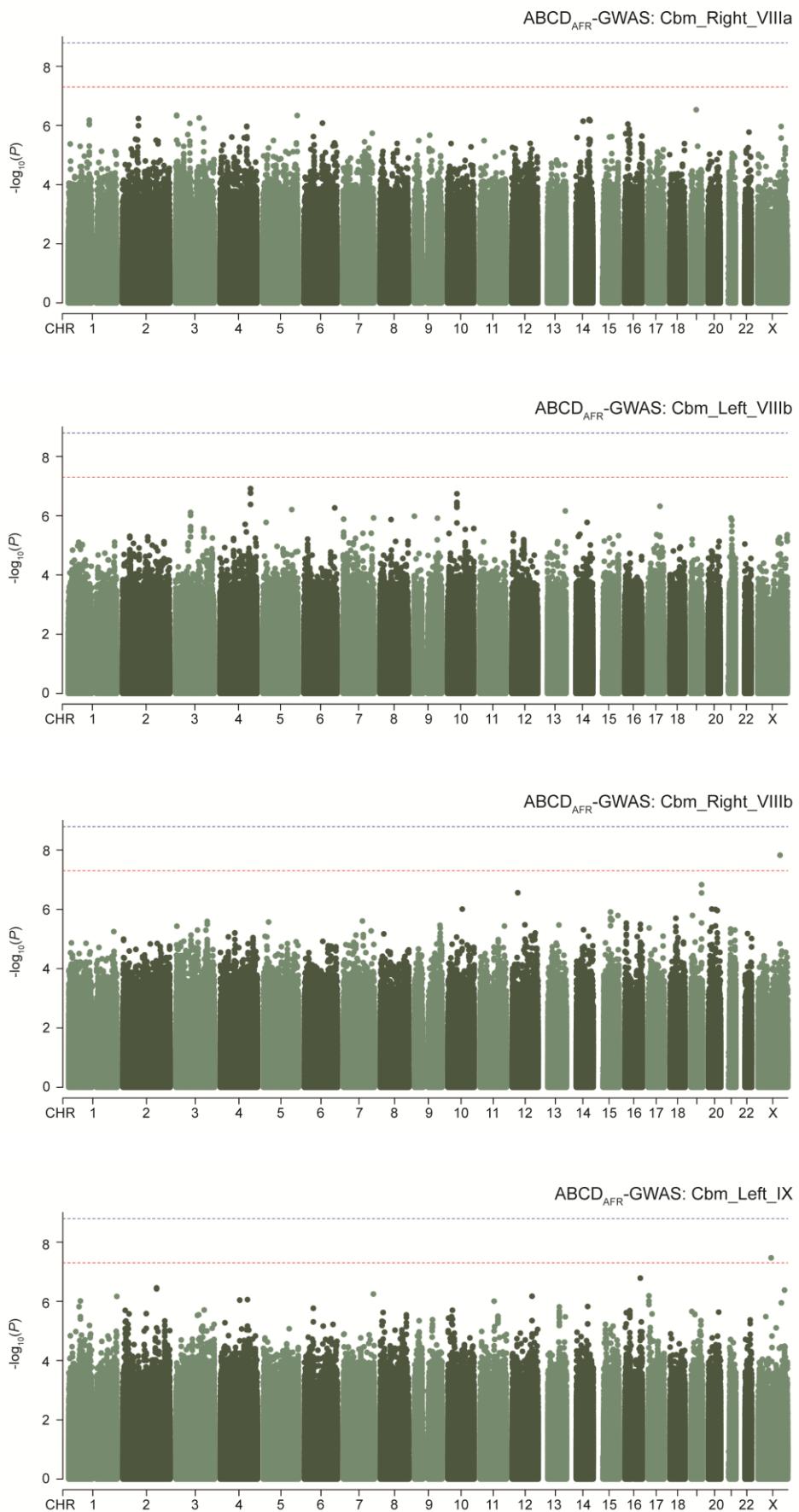
ABCD_{AFR}-GWAS: Cbm_Left_I_IV

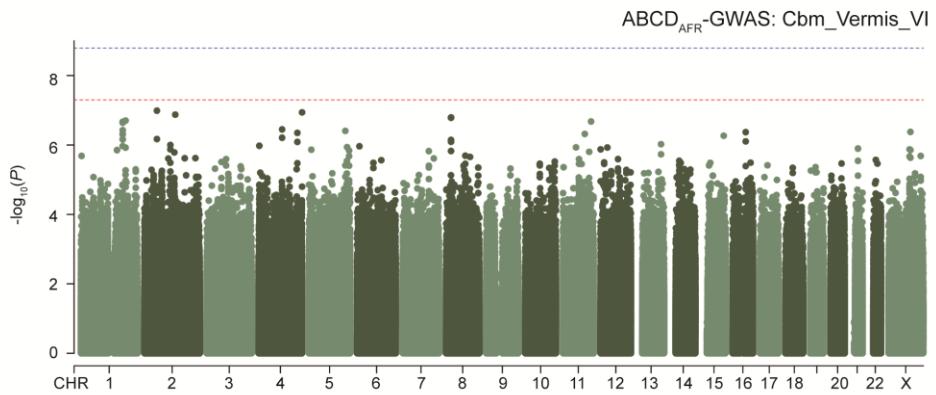
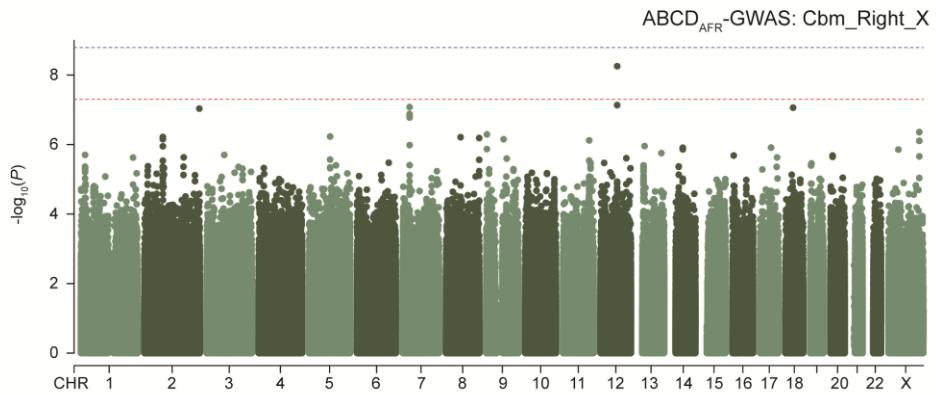
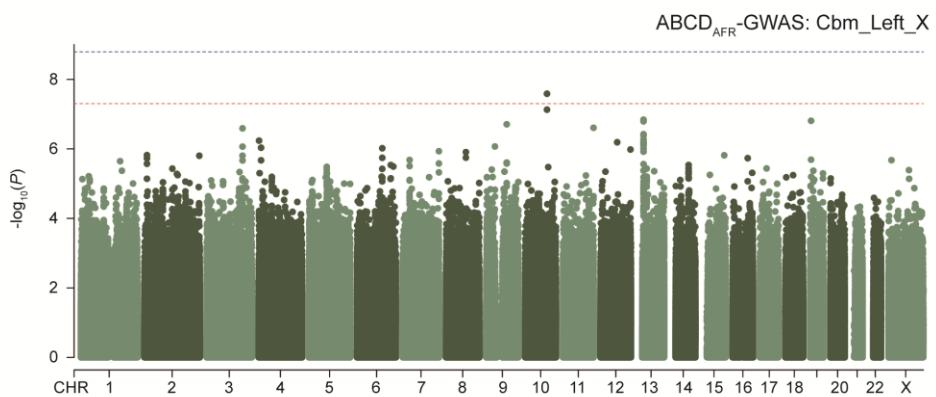
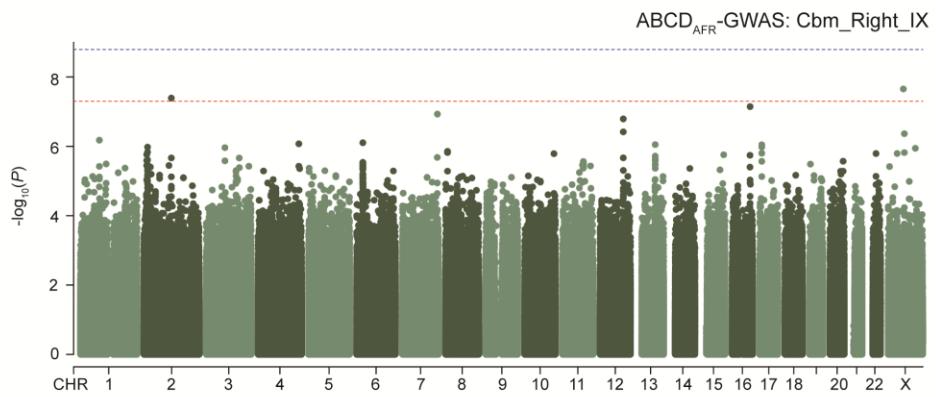




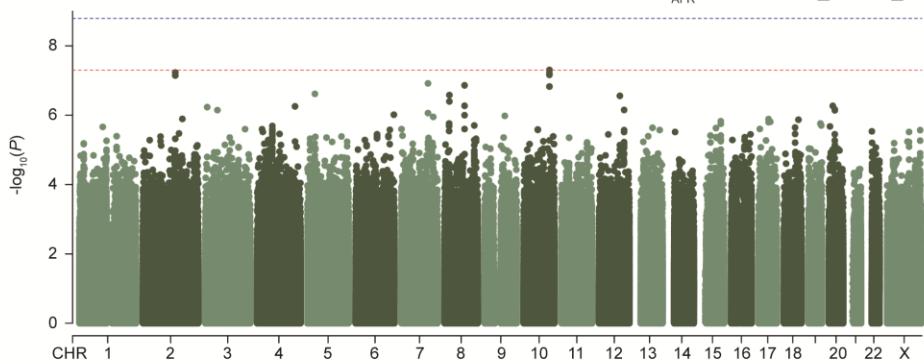




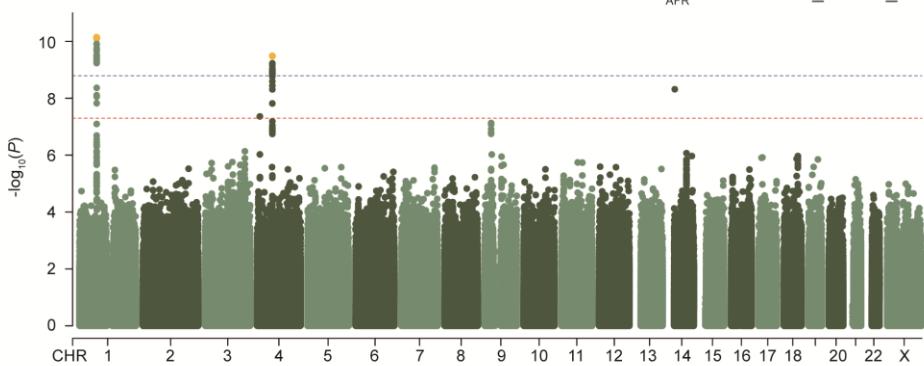




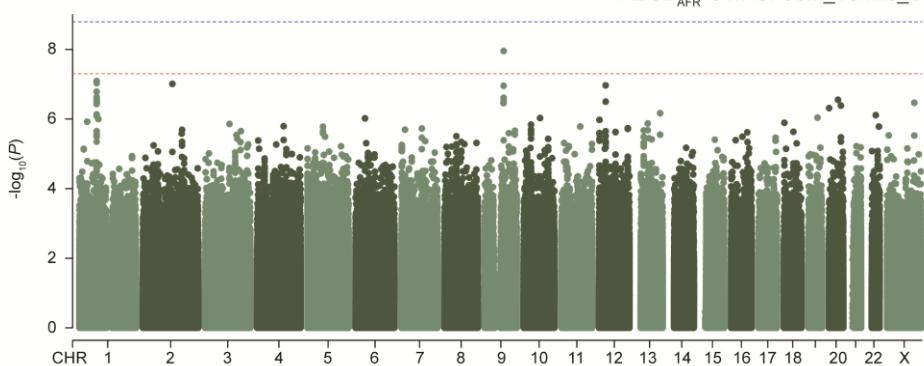
ABCD_{AFR}-GWAS: Cbm_Vermis_VII



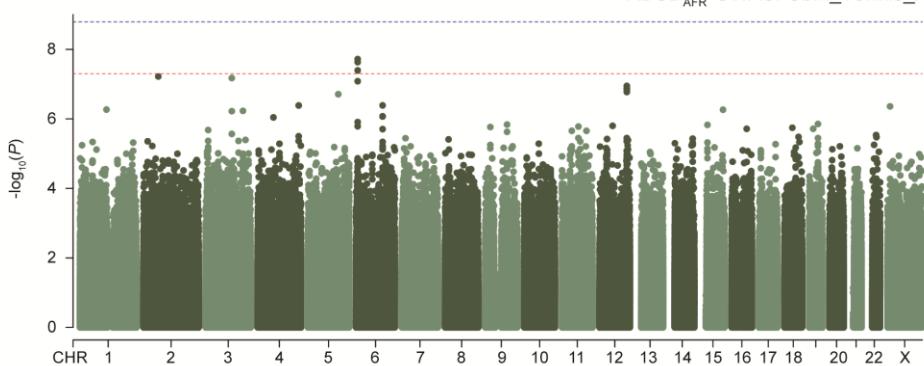
ABCD_{AFR}-GWAS: Cbm_Vermis_VIII



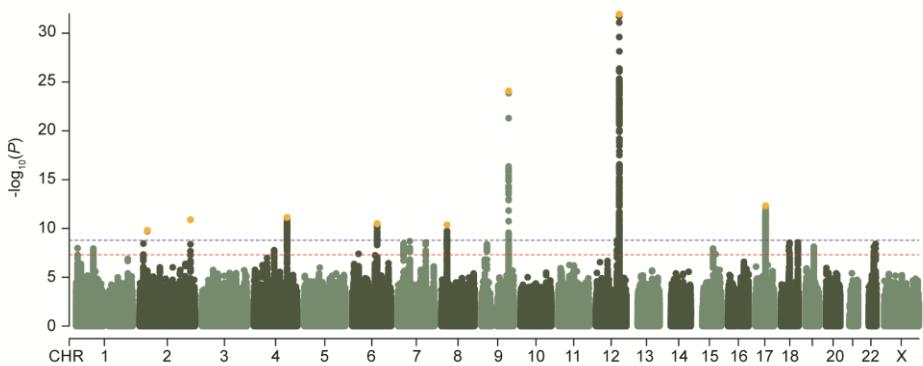
ABCD_{AFR}-GWAS: Cbm_Vermis_IX



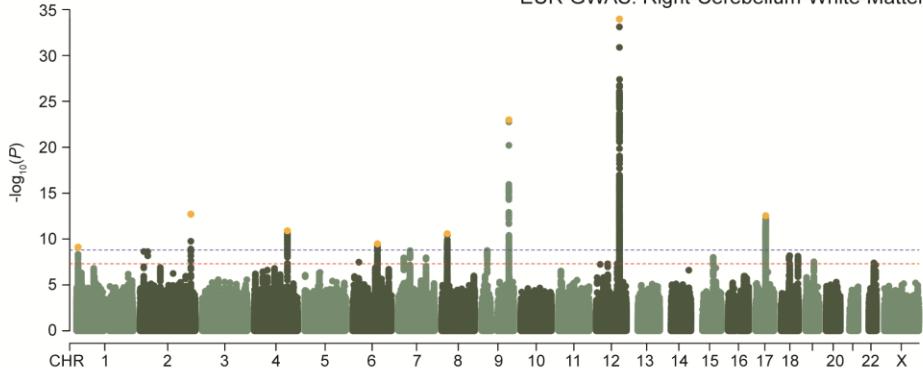
ABCD_{AFR}-GWAS: Cbm_Vermis_X



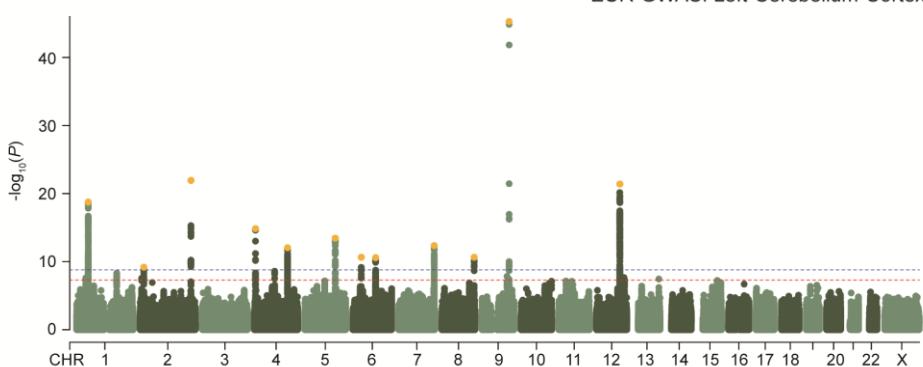
EUR-GWAS: Left-Cerebellum-White-Matter



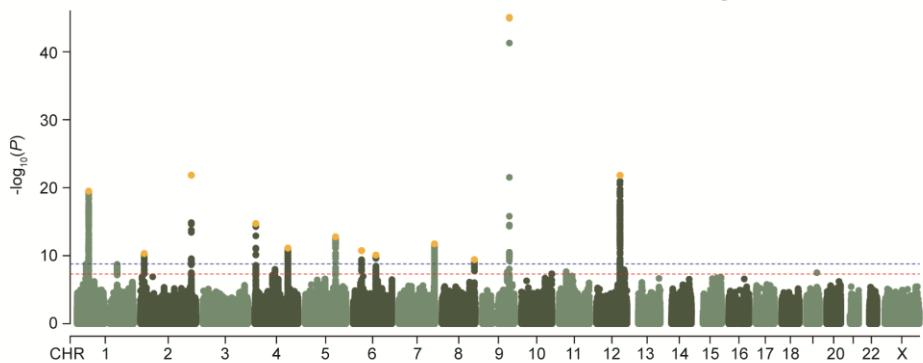
EUR-GWAS: Right-Cerebellum-White-Matter



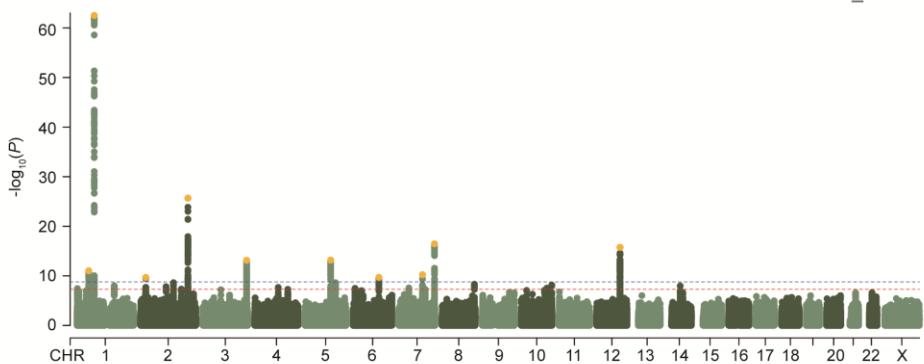
EUR-GWAS: Left-Cerebellum-Cortex



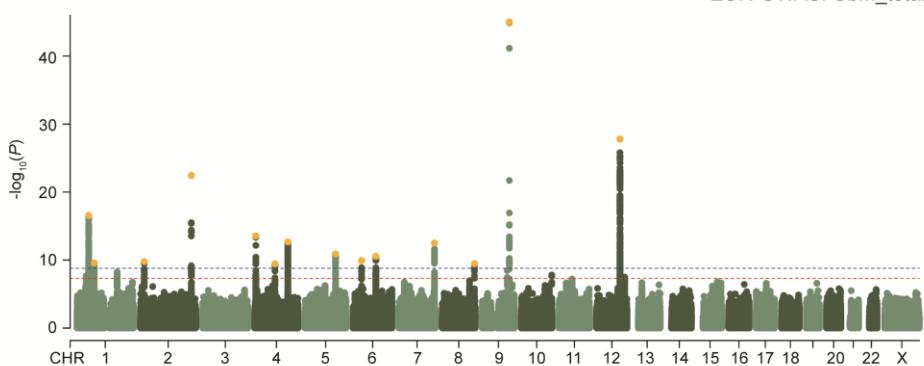
EUR-GWAS: Right-Cerebellum-Cortex



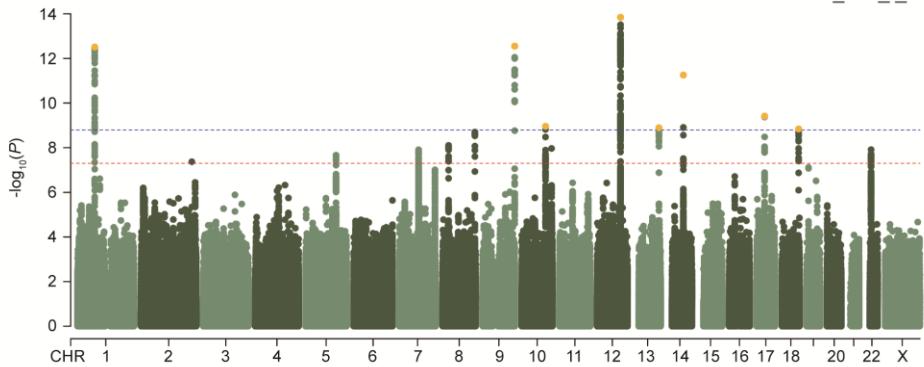
EUR-GWAS: Cbm_Vermis



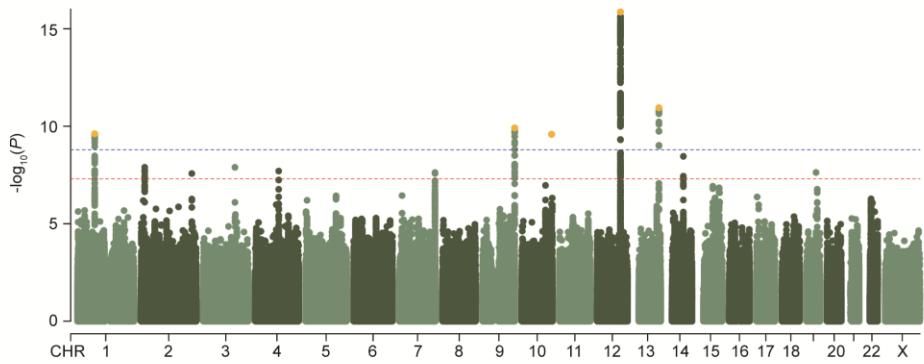
EUR-GWAS: Cbm_total

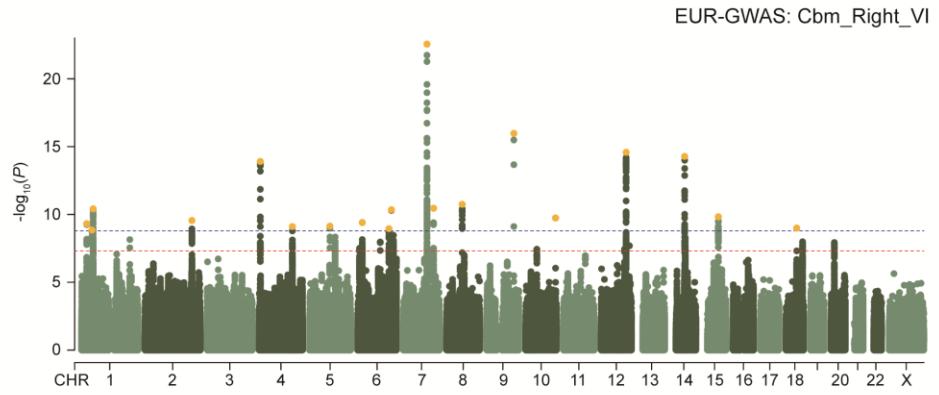
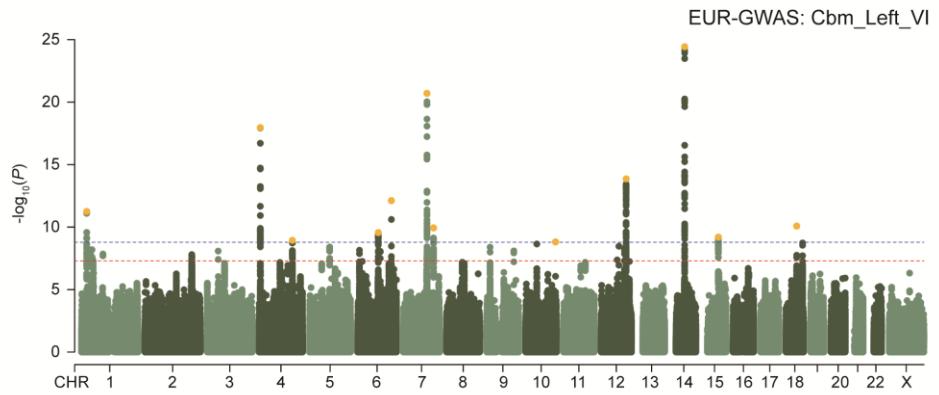
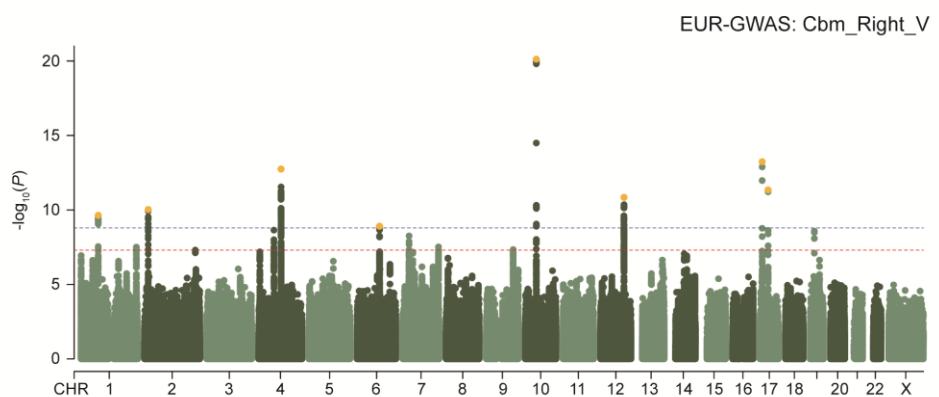
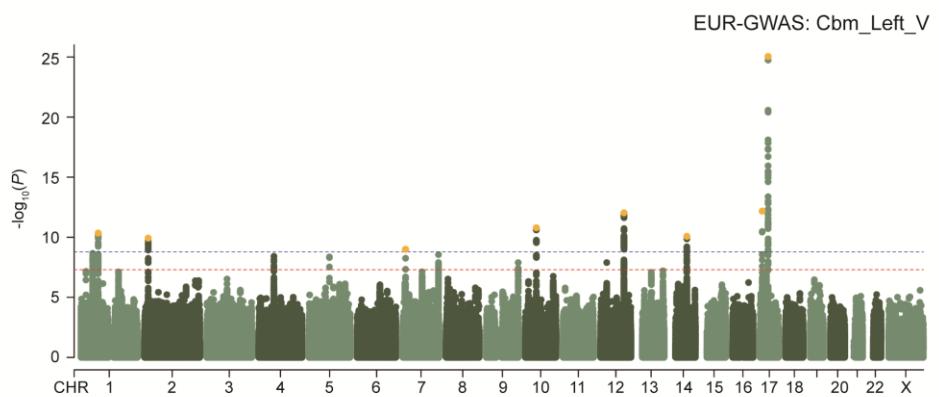


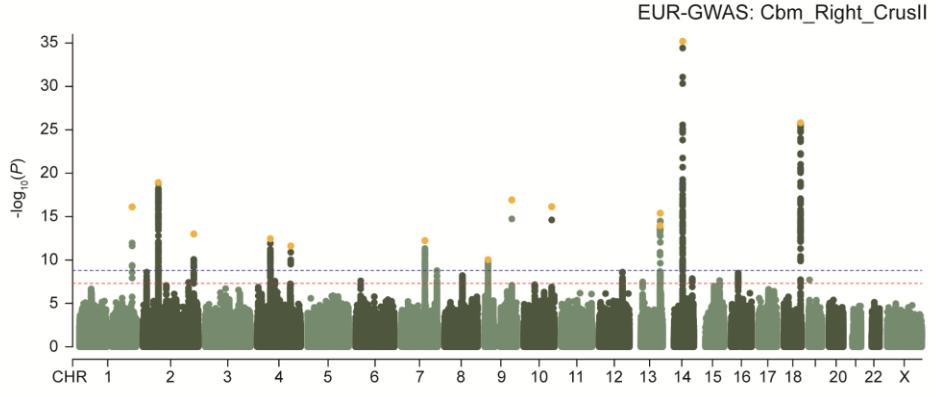
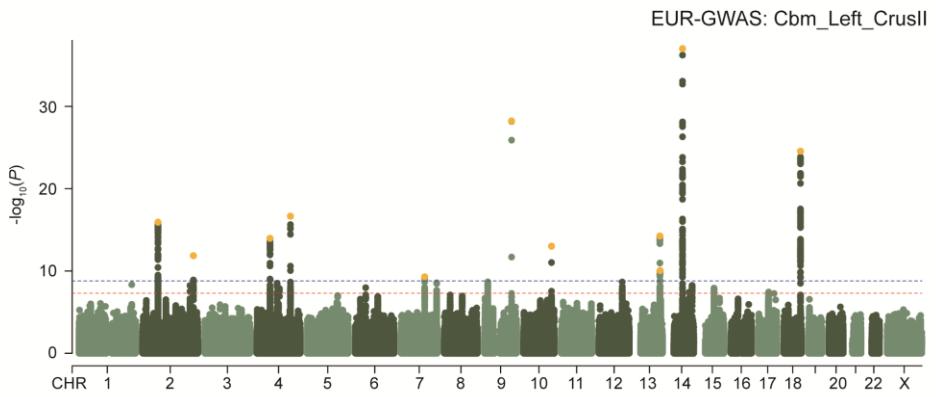
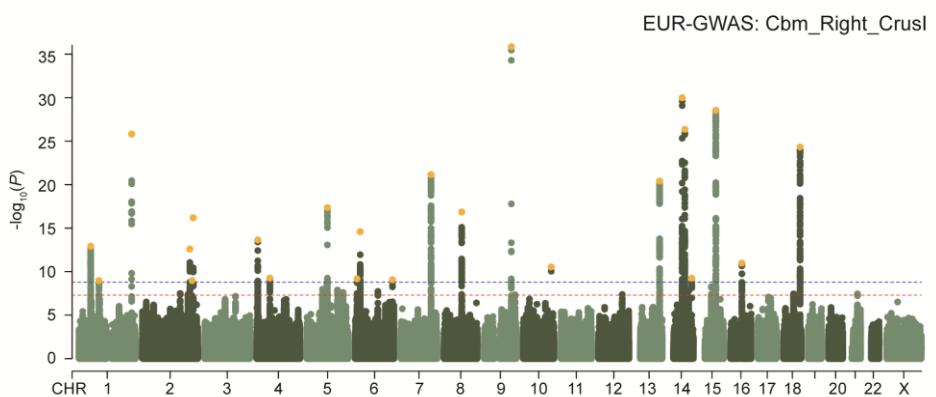
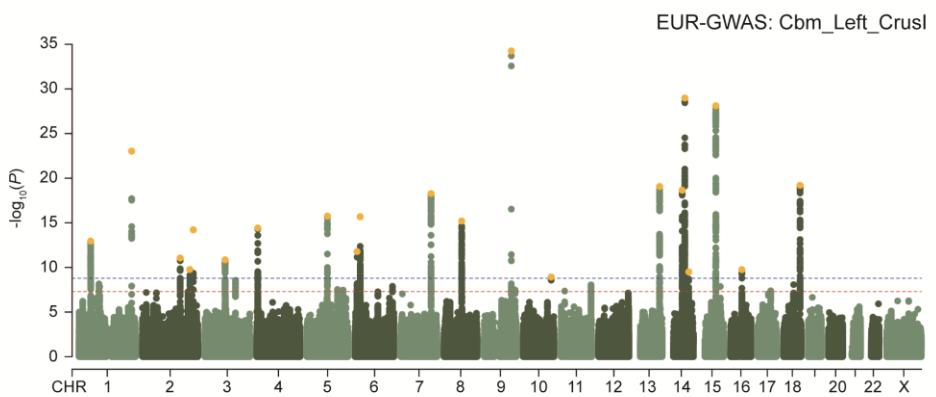
EUR-GWAS: Cbm_Left_I_IV

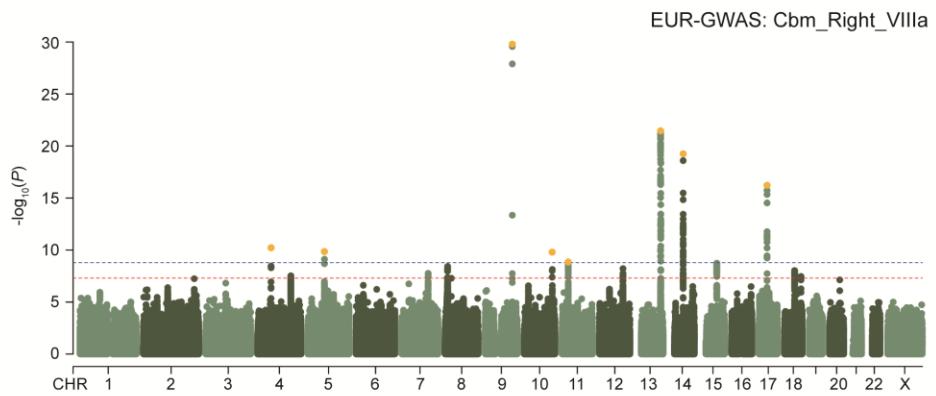
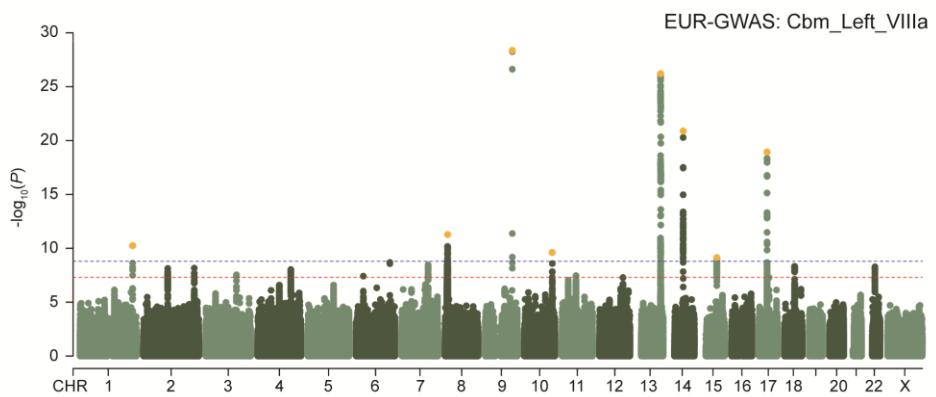
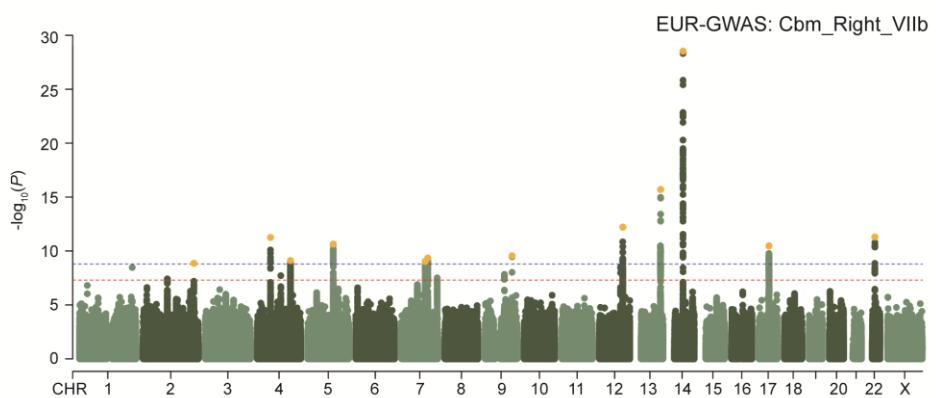
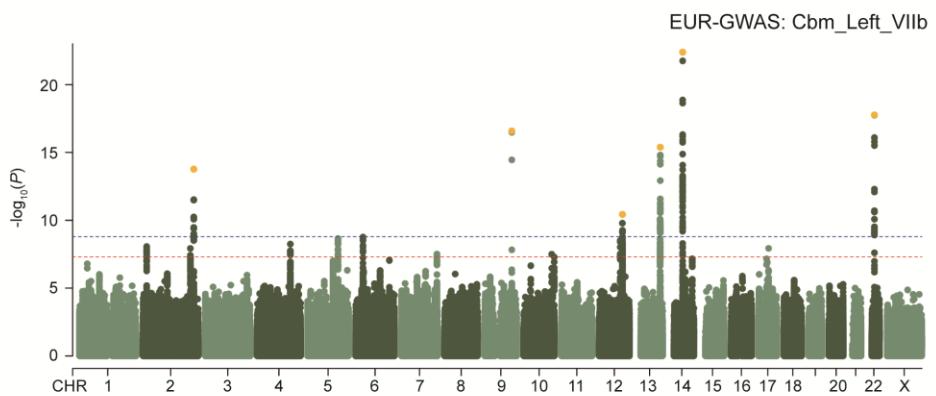


EUR-GWAS: Cbm_Right_I_IV

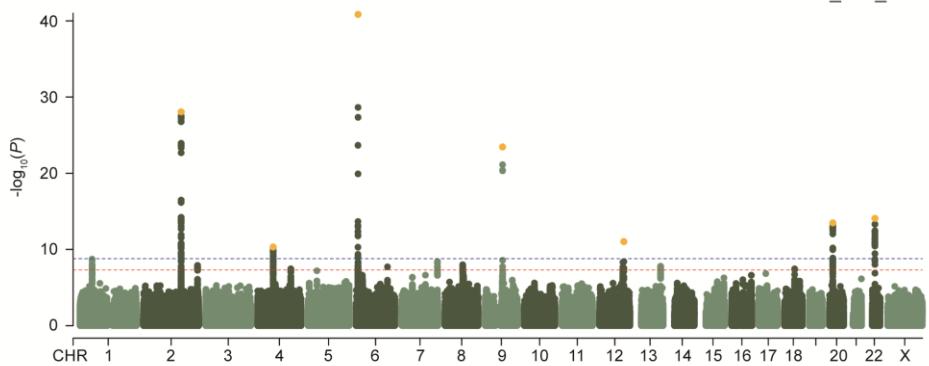




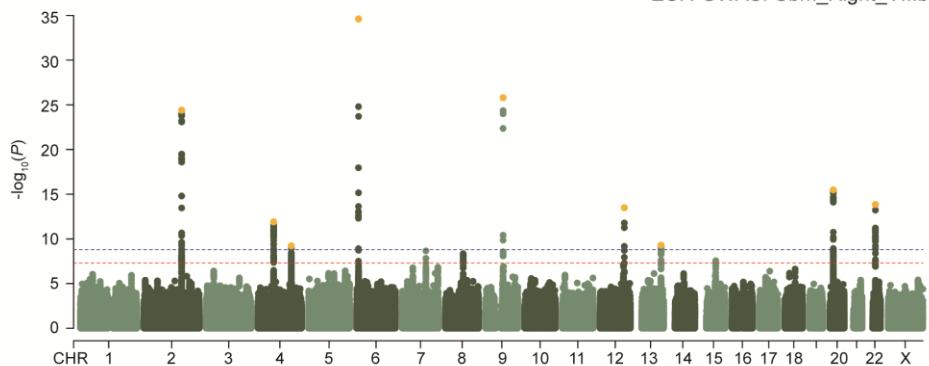




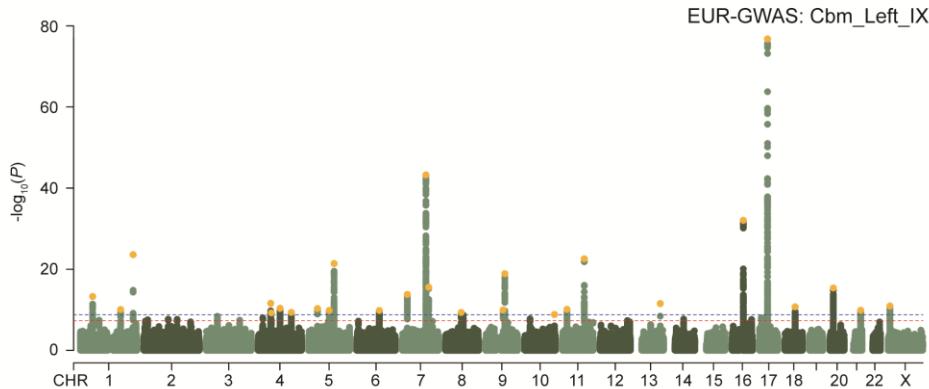
EUR-GWAS: Cbm_Left_VIIIb



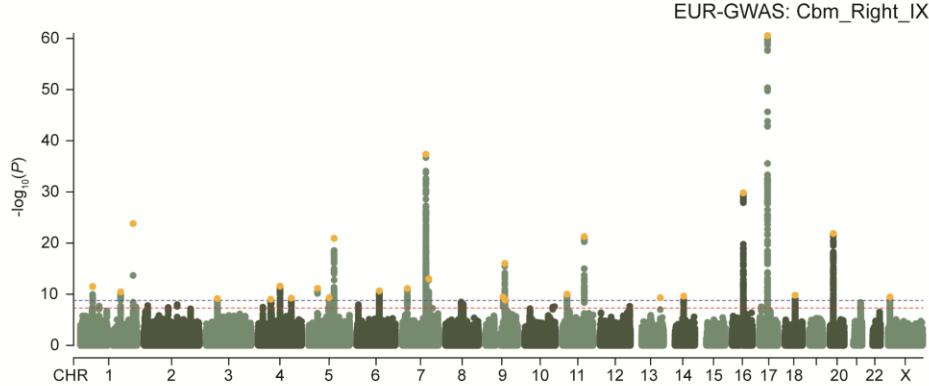
EUR-GWAS: Cbm_Right_VIIIb

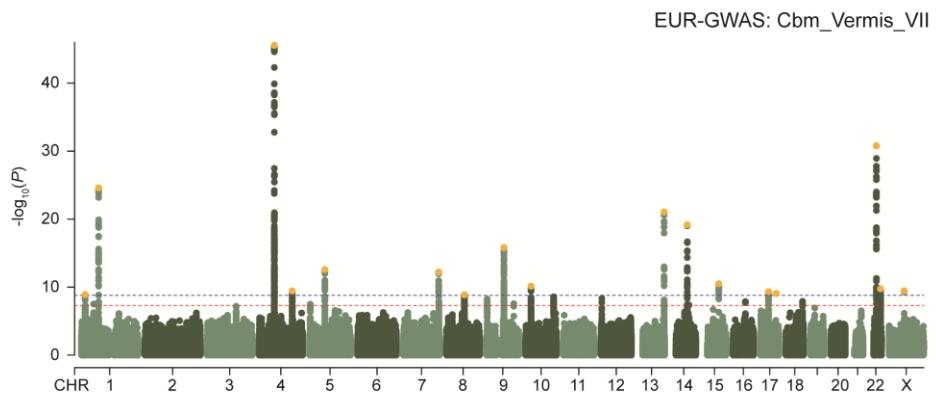
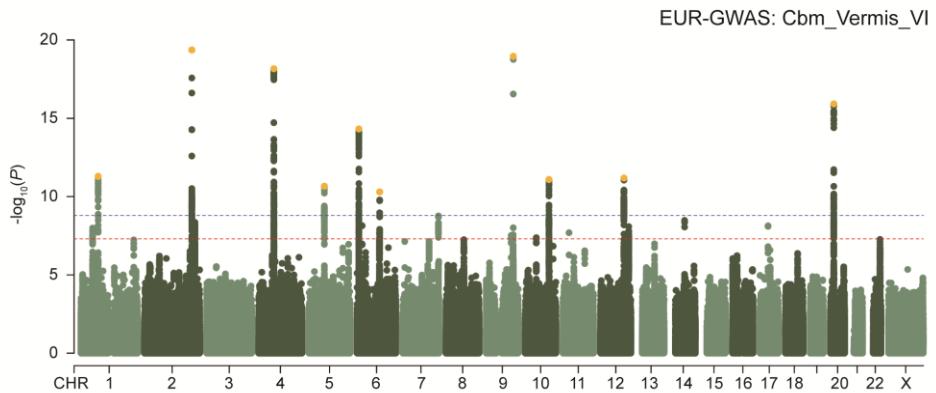
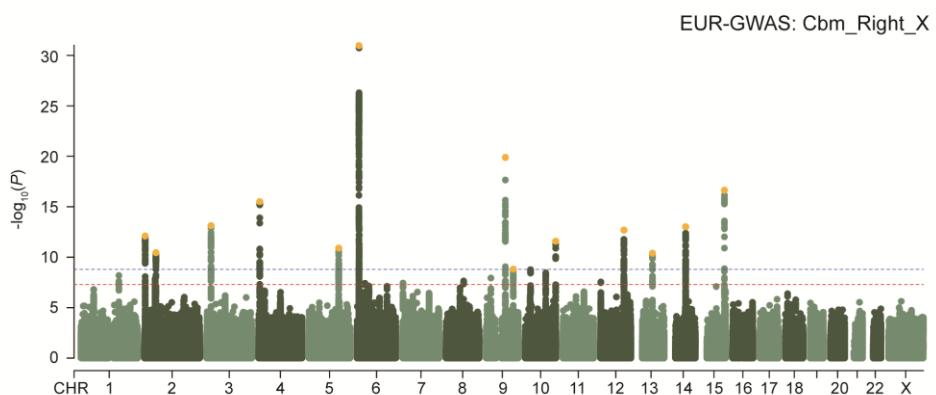
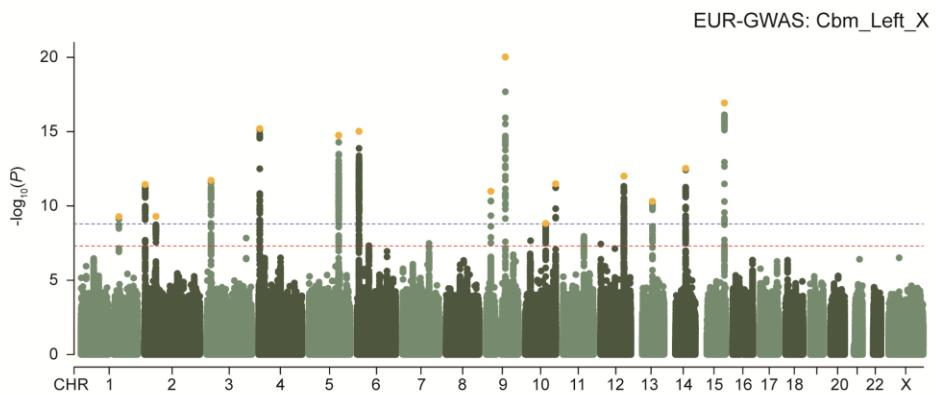


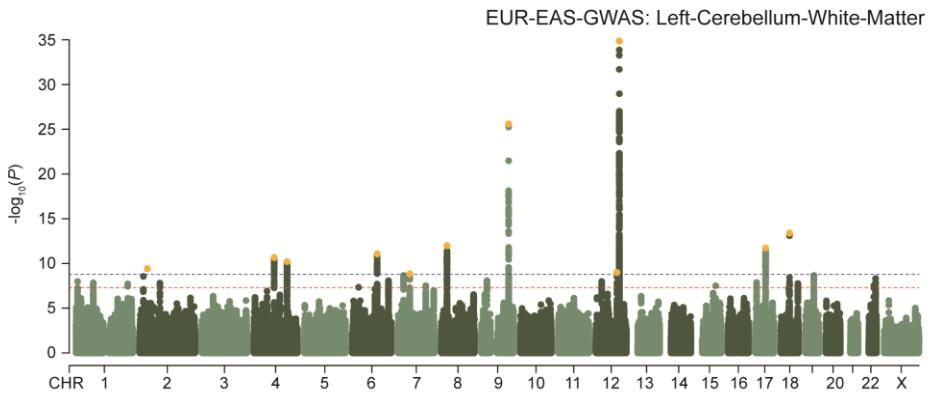
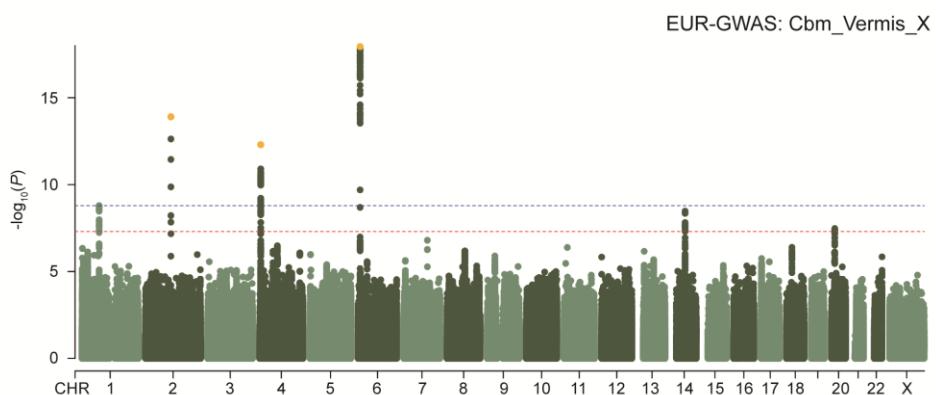
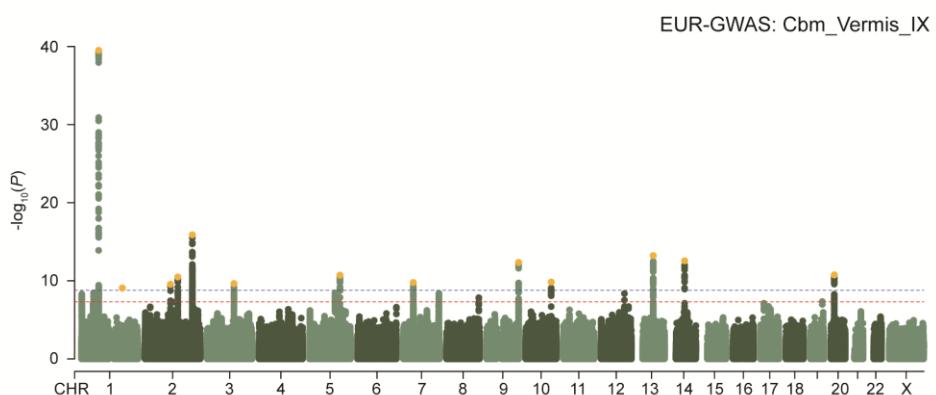
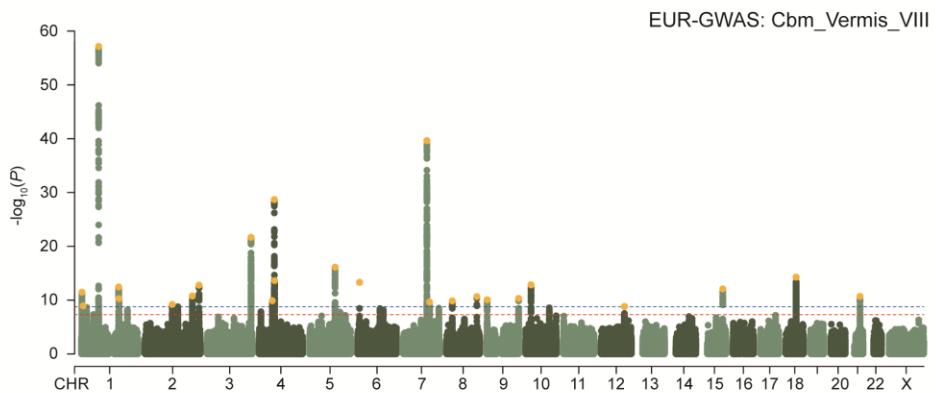
EUR-GWAS: Cbm_Left_IX



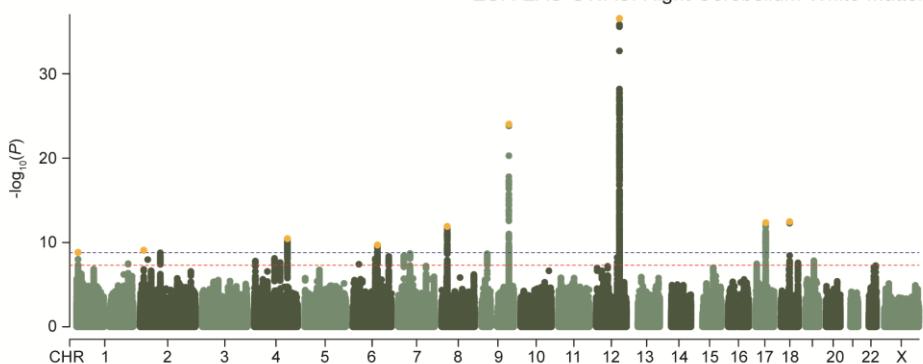
EUR-GWAS: Cbm_Right_IX



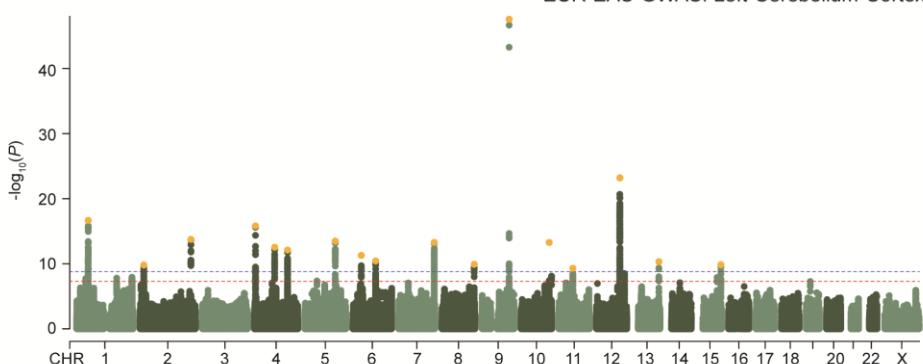




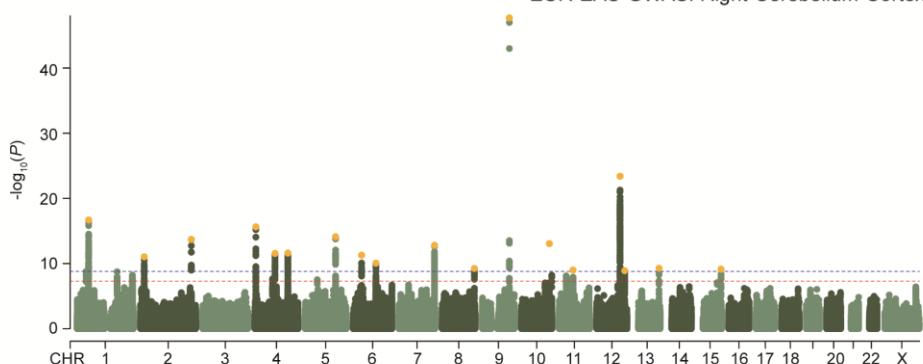
EUR-EAS-GWAS: Right-Cerebellum-White-Matter



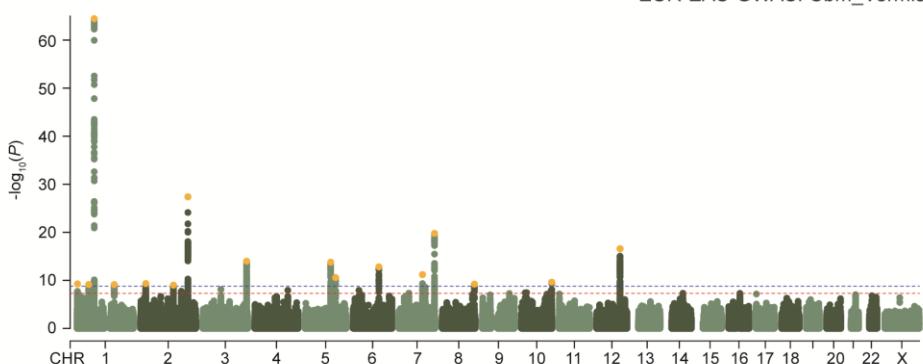
EUR-EAS-GWAS: Left-Cerebellum-Cortex

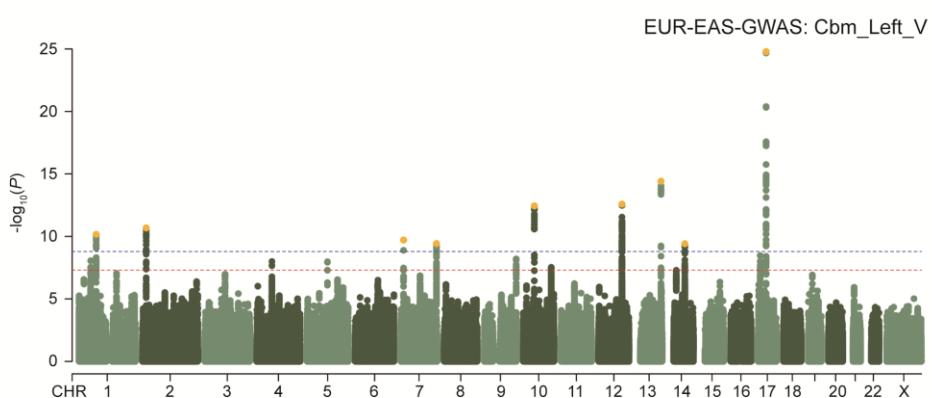
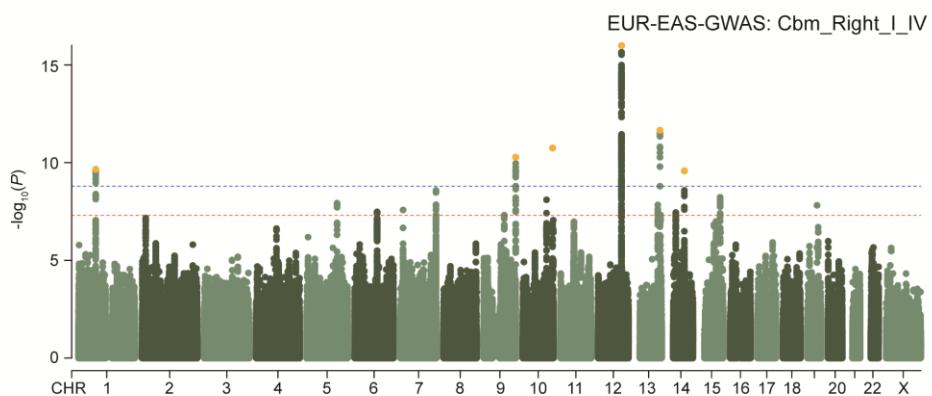
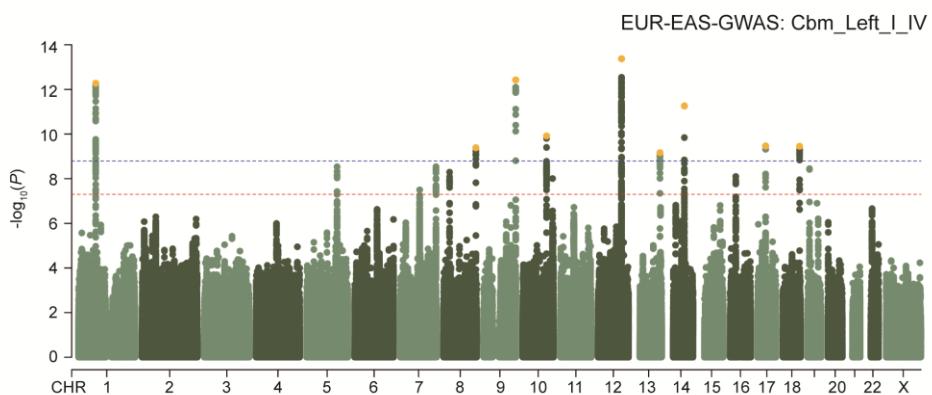
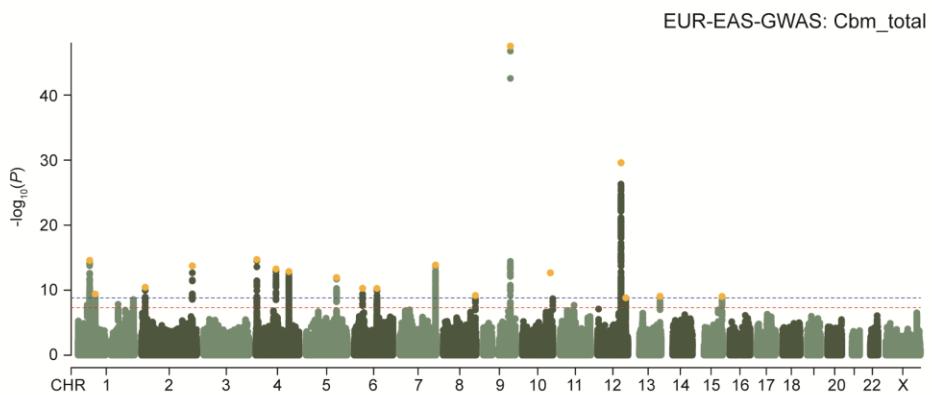


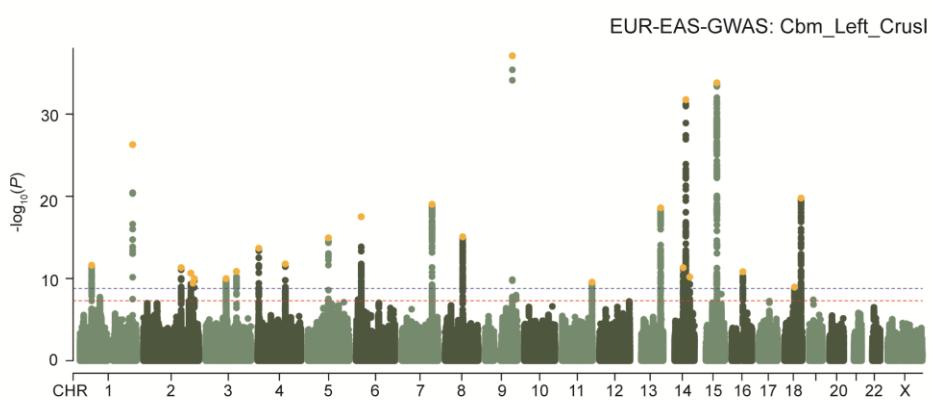
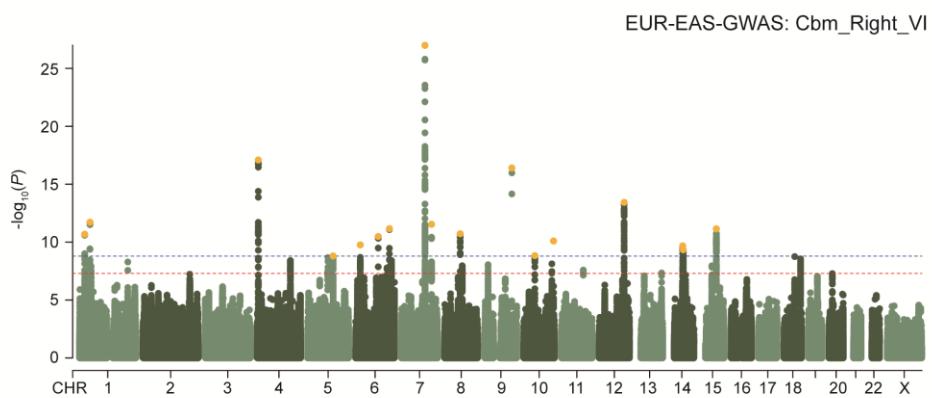
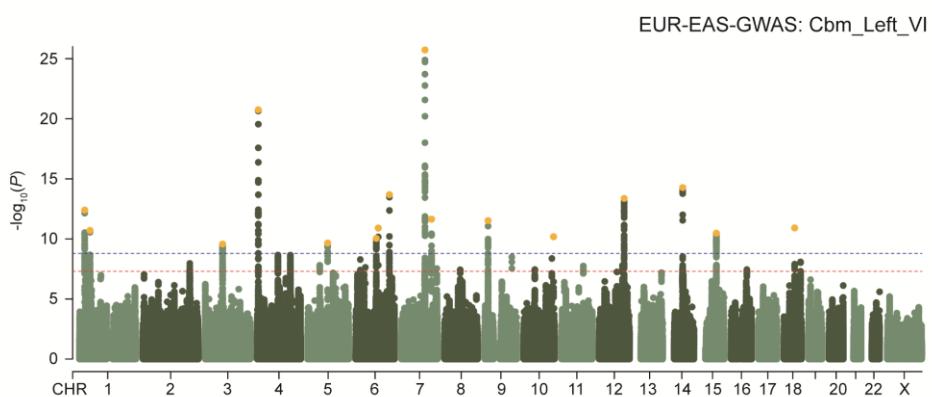
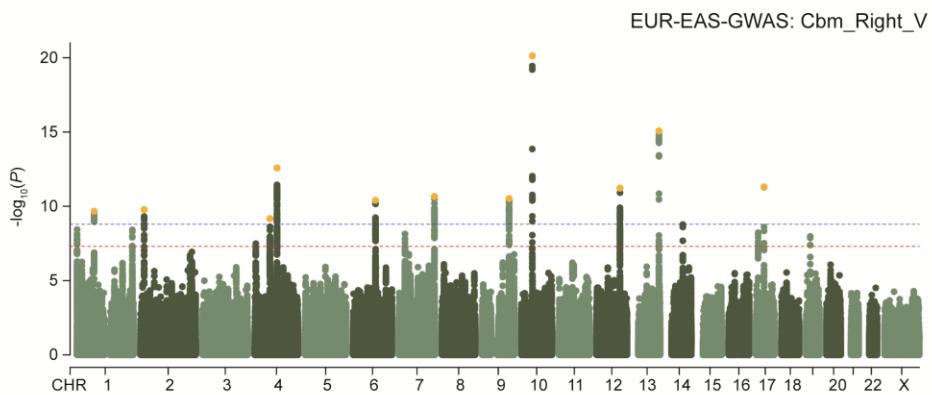
EUR-EAS-GWAS: Right-Cerebellum-Cortex

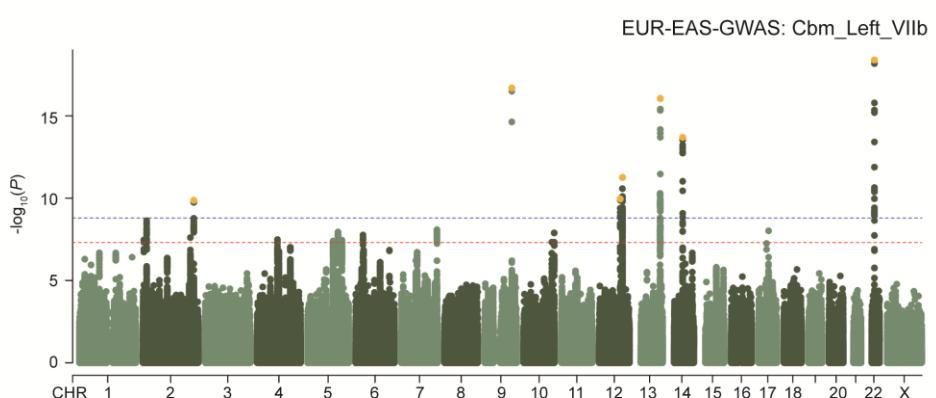
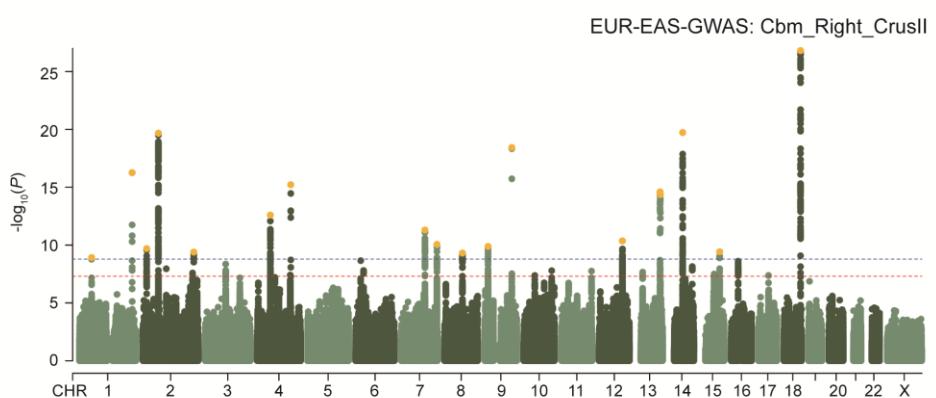
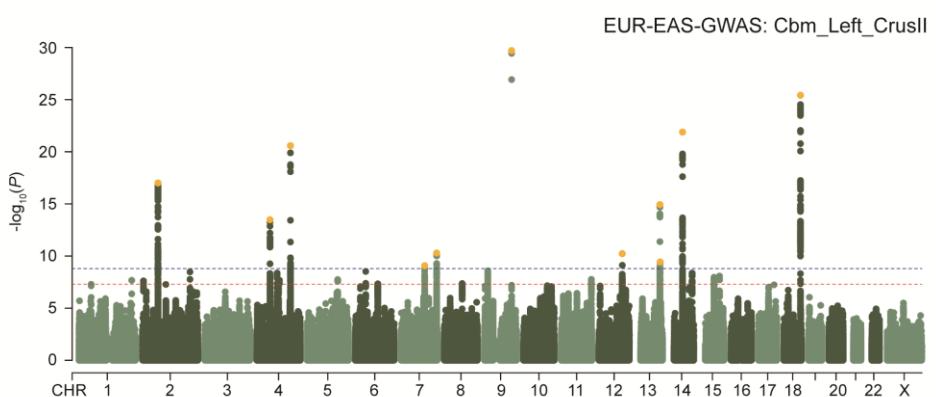
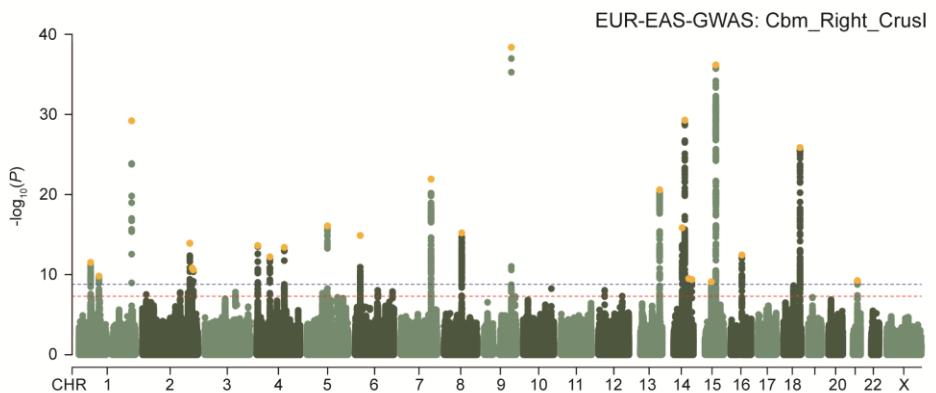


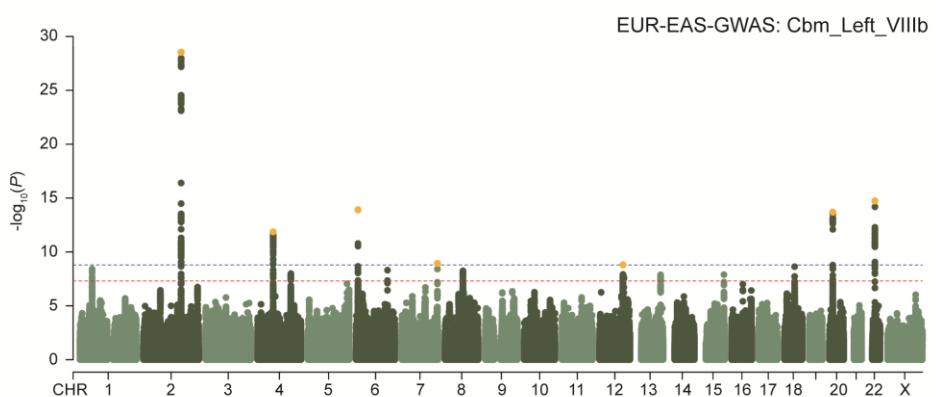
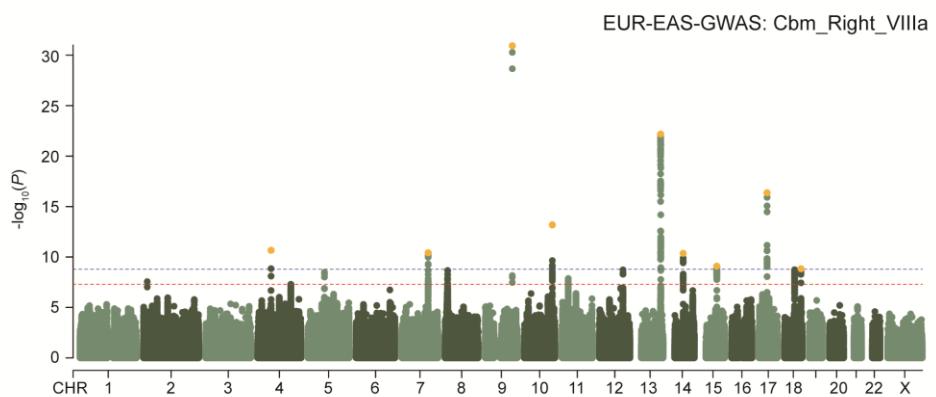
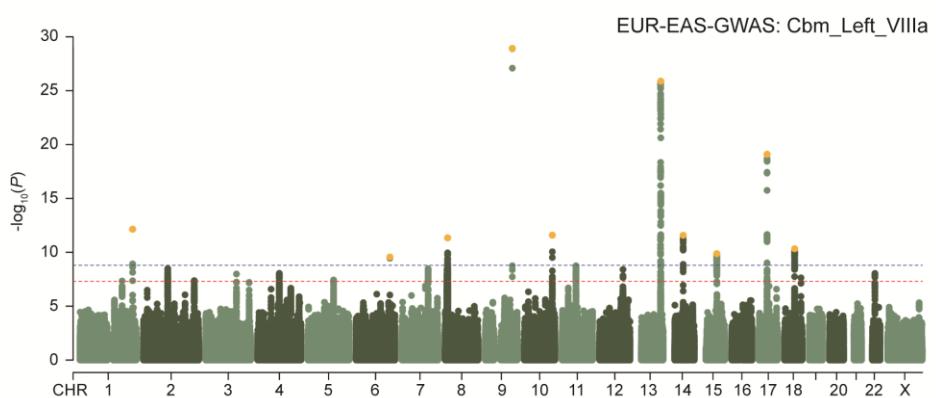
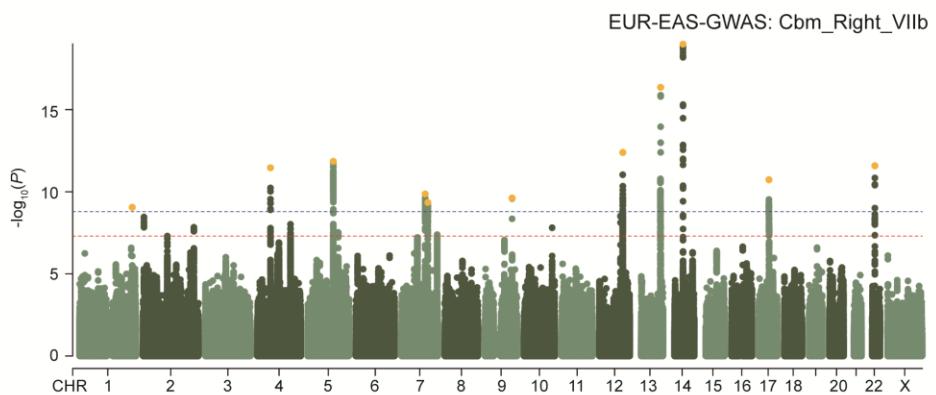
EUR-EAS-GWAS: Cbm_Vermis

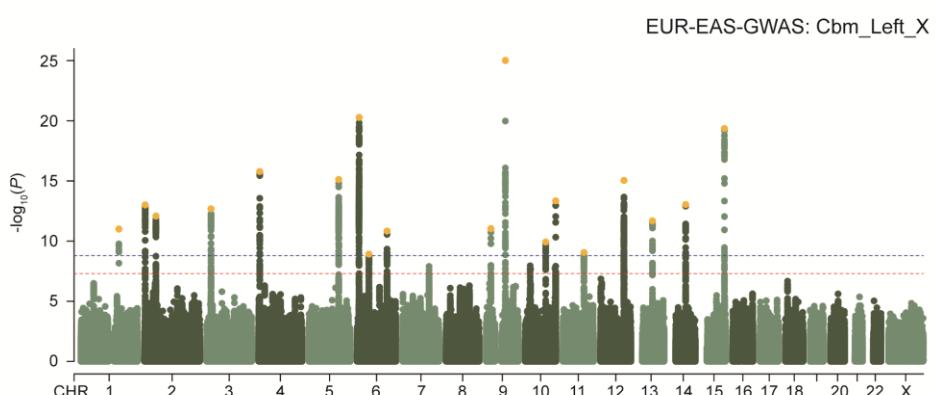
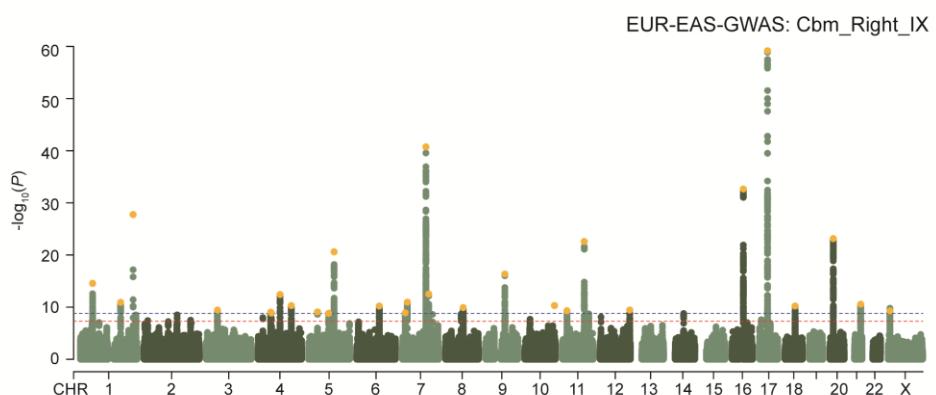
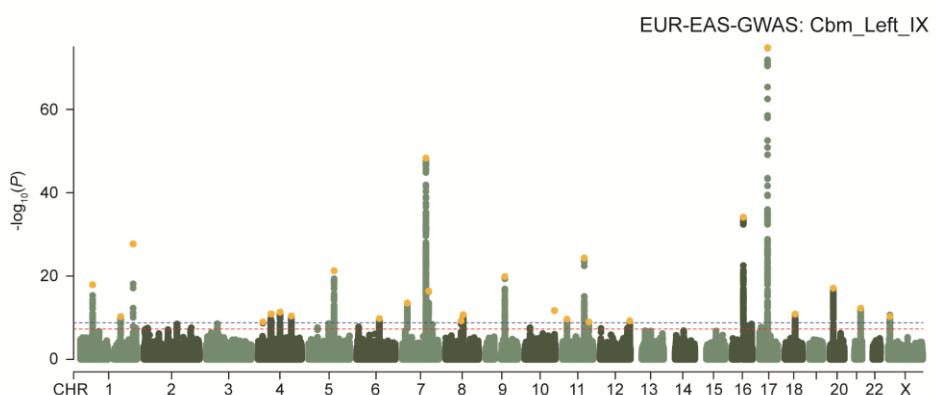
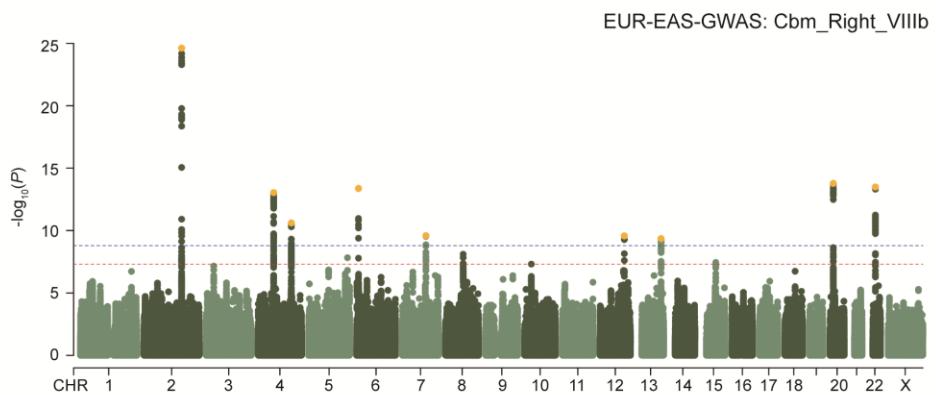




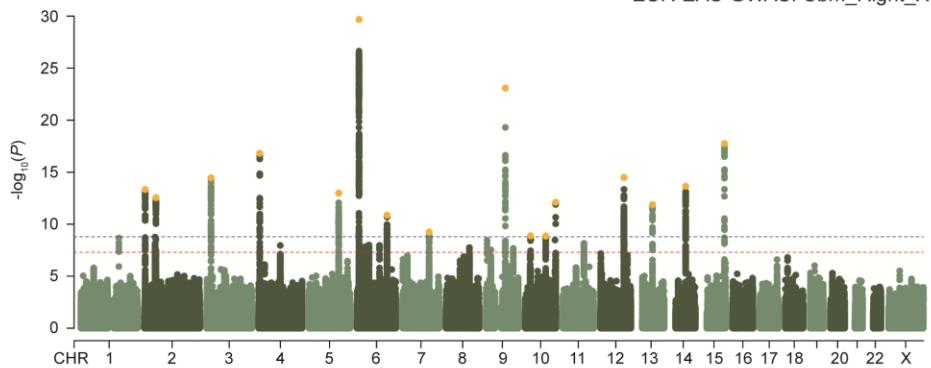




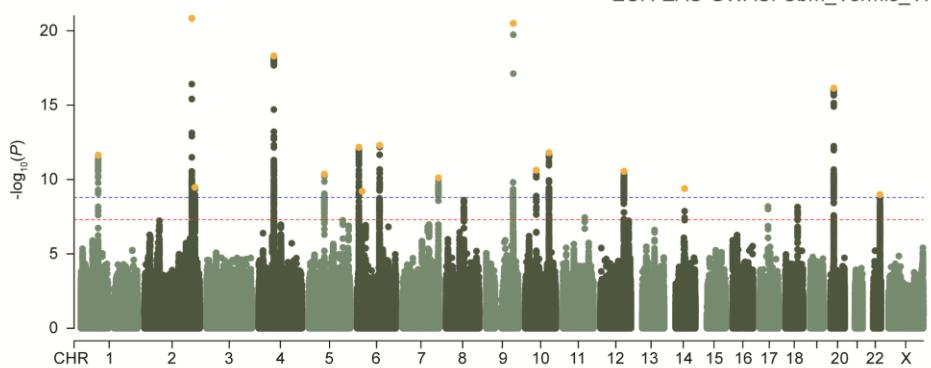




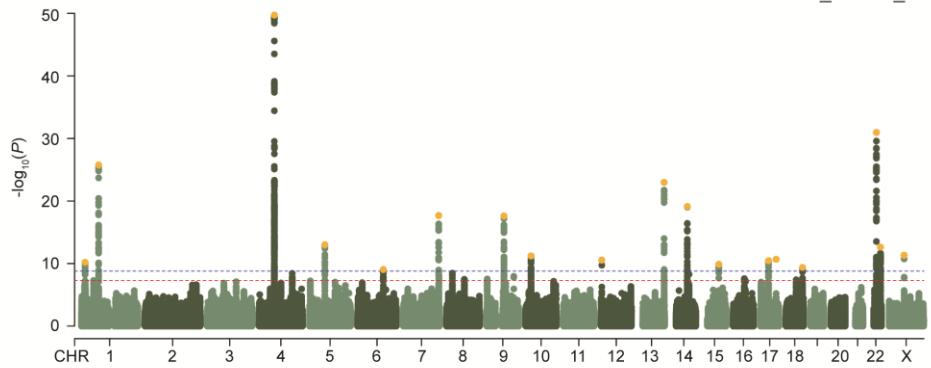
EUR-EAS-GWAS: Cbm_Right_X



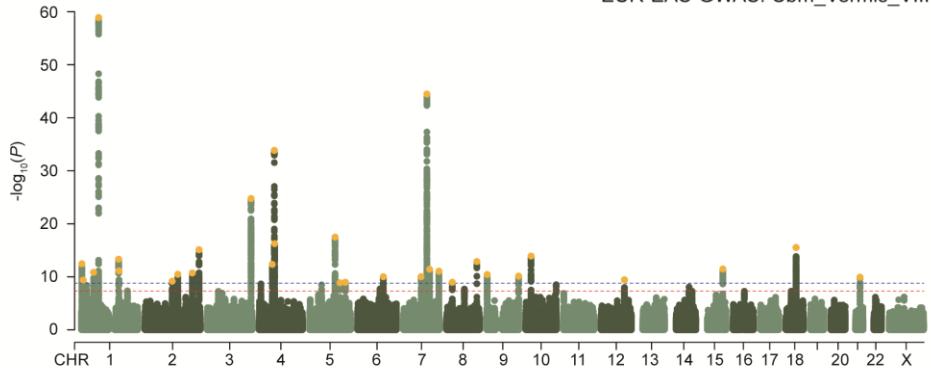
EUR-EAS-GWAS: Cbm_Vermis_VI



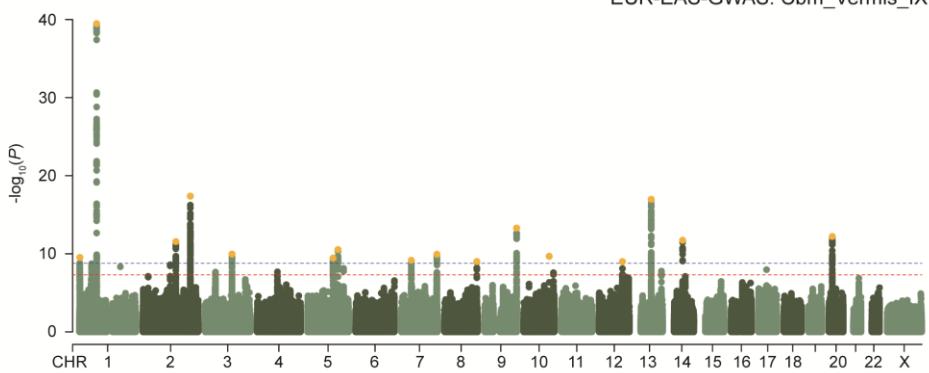
EUR-EAS-GWAS: Cbm_Vermis_VII



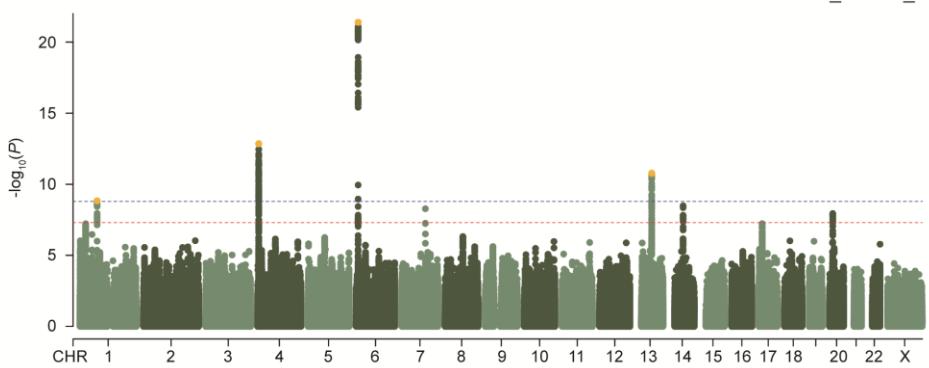
EUR-EAS-GWAS: Cbm_Vermis_VIII



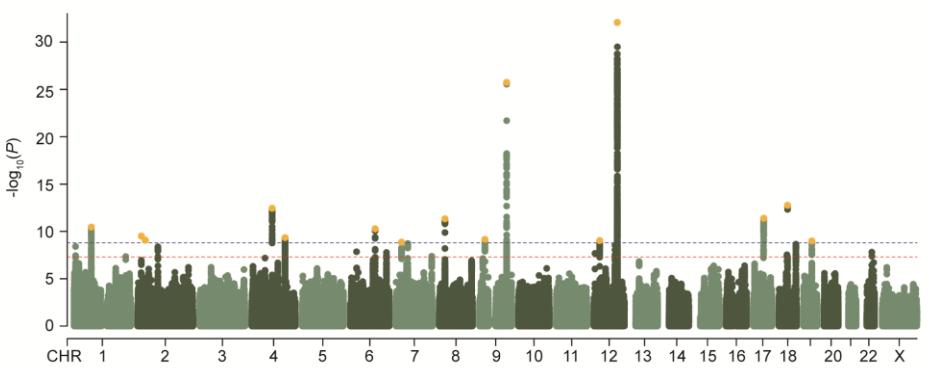
EUR-EAS-GWAS: Cbm_Vermis_IX



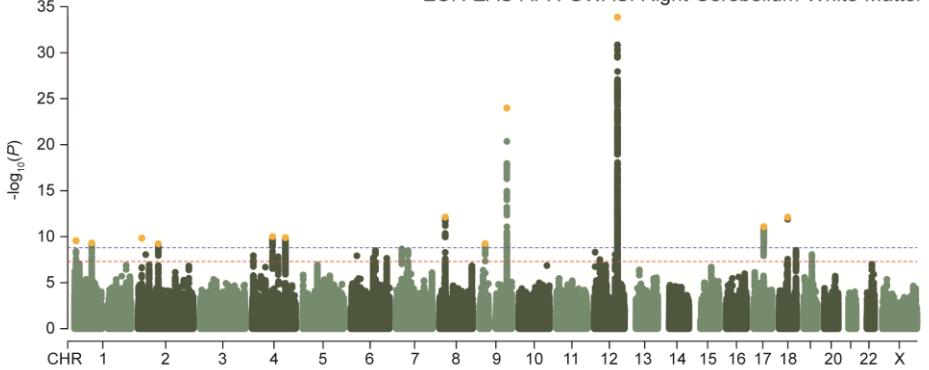
EUR-EAS-GWAS: Cbm_Vermis_X



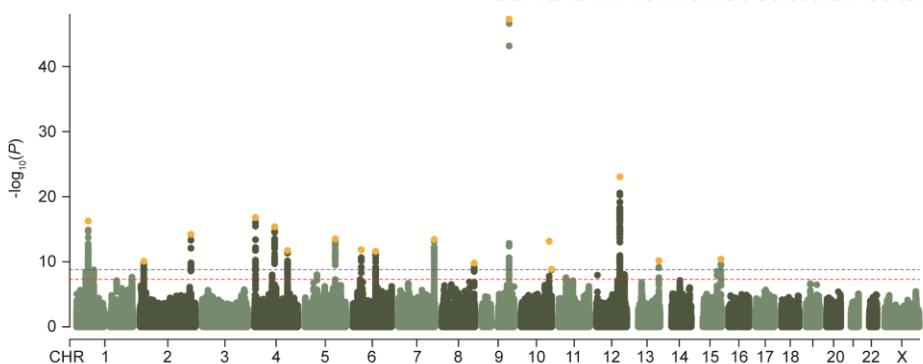
EUR-EAS-AFR-GWAS: Left-Cerebellum-White-Matter



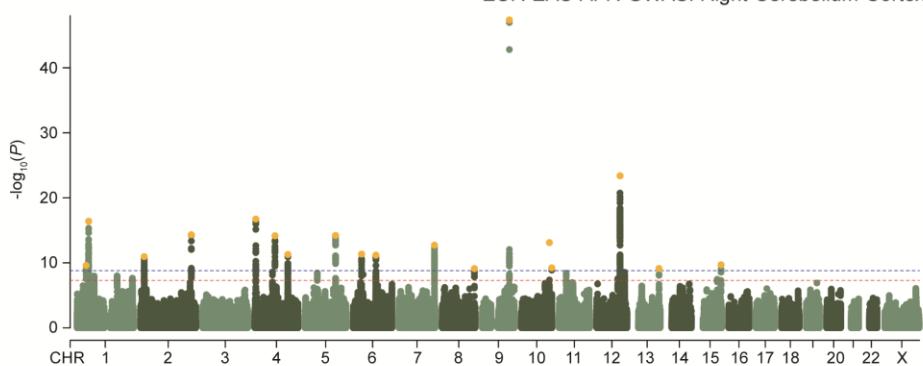
EUR-EAS-AFR-GWAS: Right-Cerebellum-White-Matter



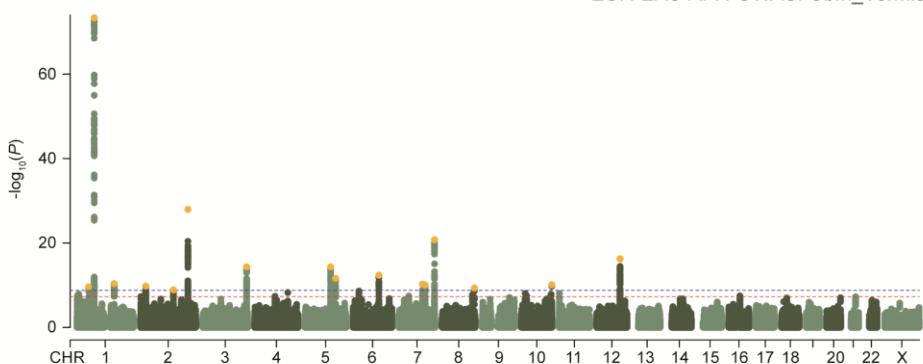
EUR-EAS-AFR-GWAS: Left-Cerebellum-Cortex



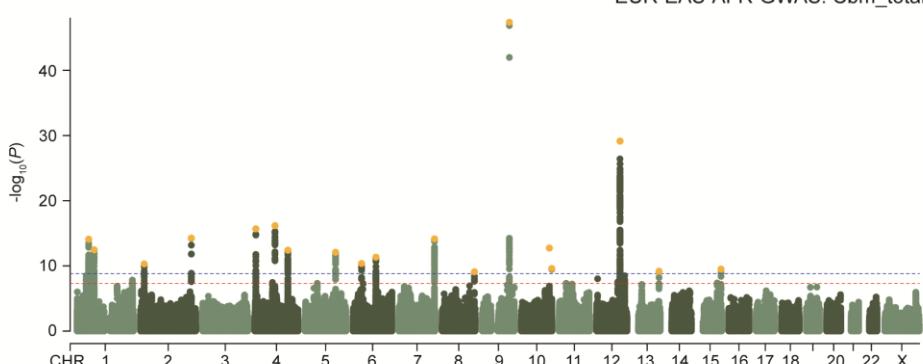
EUR-EAS-AFR-GWAS: Right-Cerebellum-Cortex

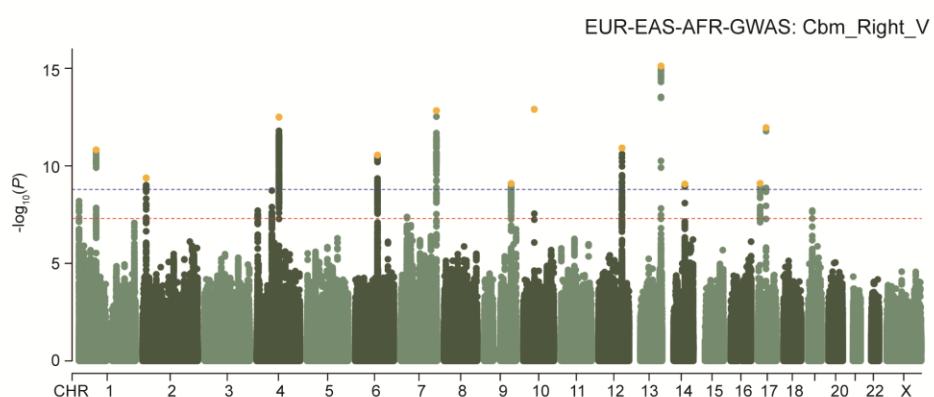
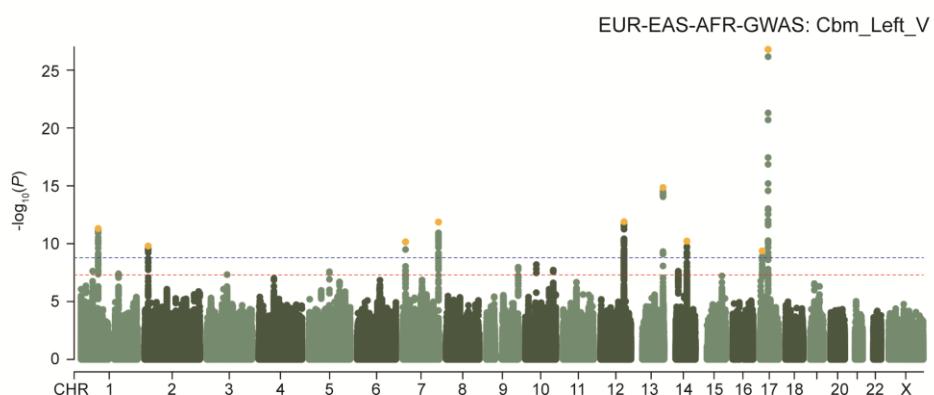
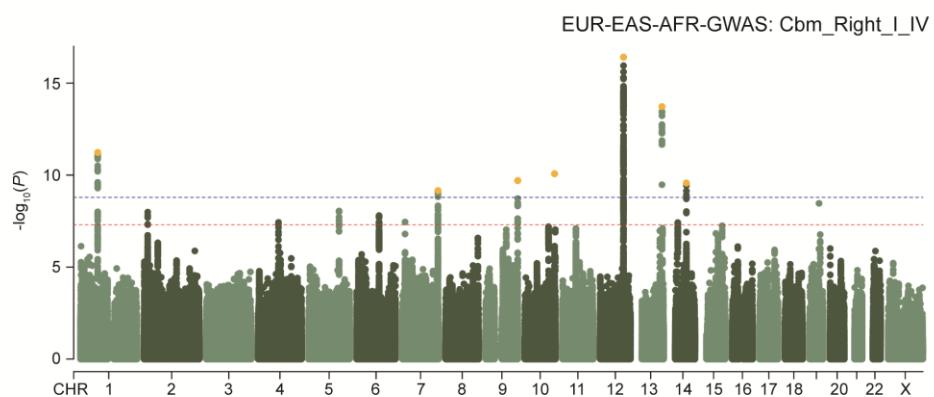
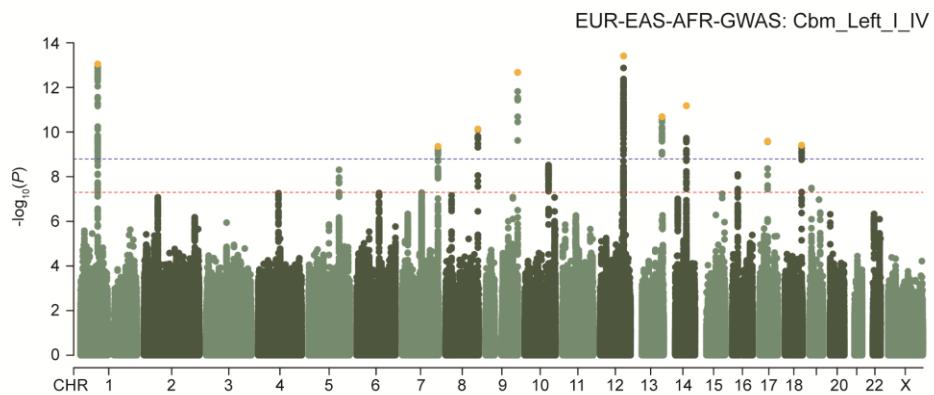


EUR-EAS-AFR-GWAS: Cbm_Vermis

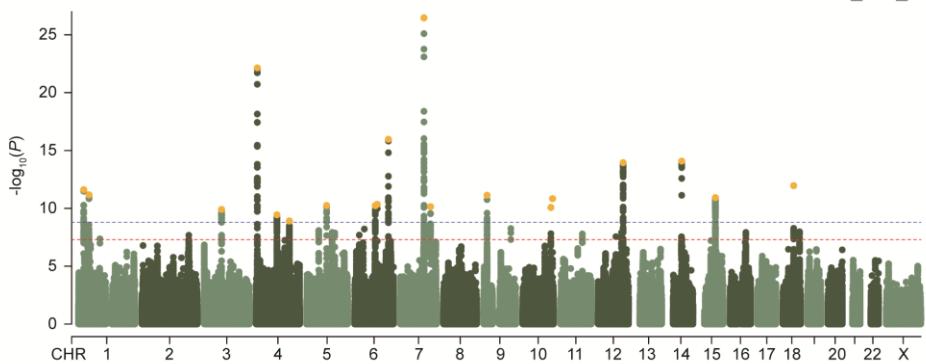


EUR-EAS-AFR-GWAS: Cbm_total

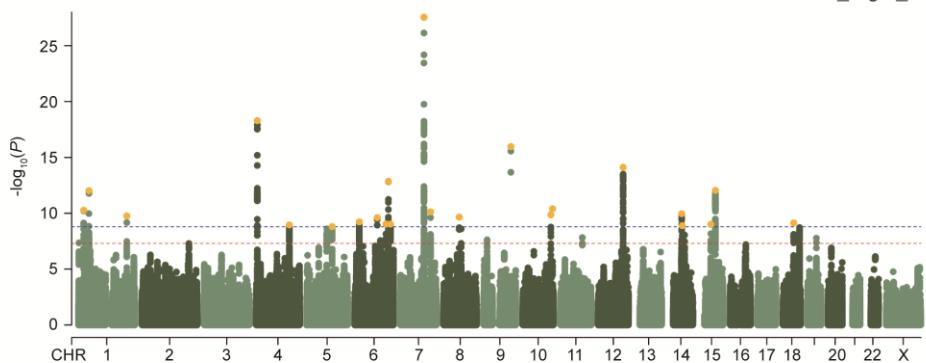




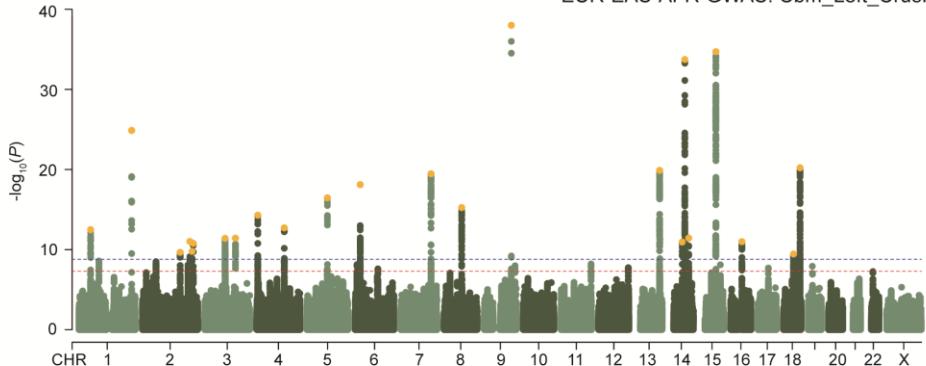
EUR-EAS-AFR-GWAS: Cbm_Left_VI



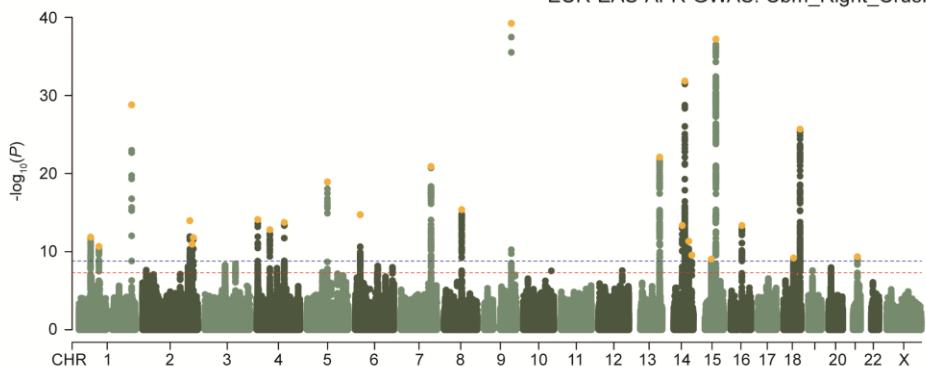
EUR-EAS-AFR-GWAS: Cbm_Right_VI

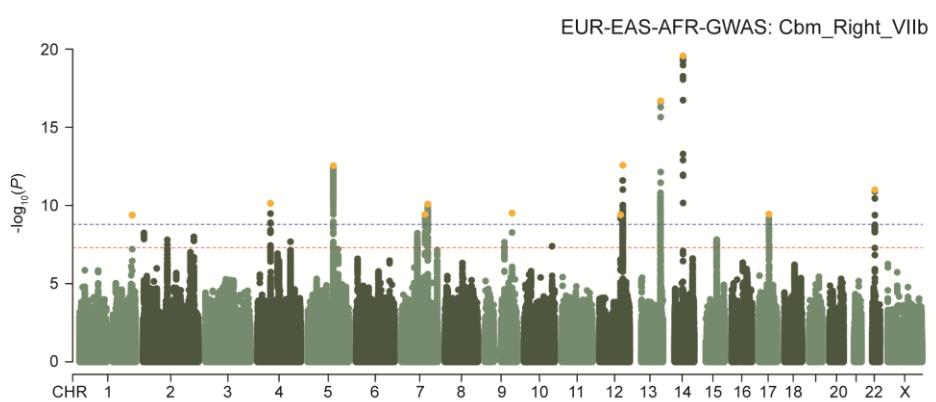
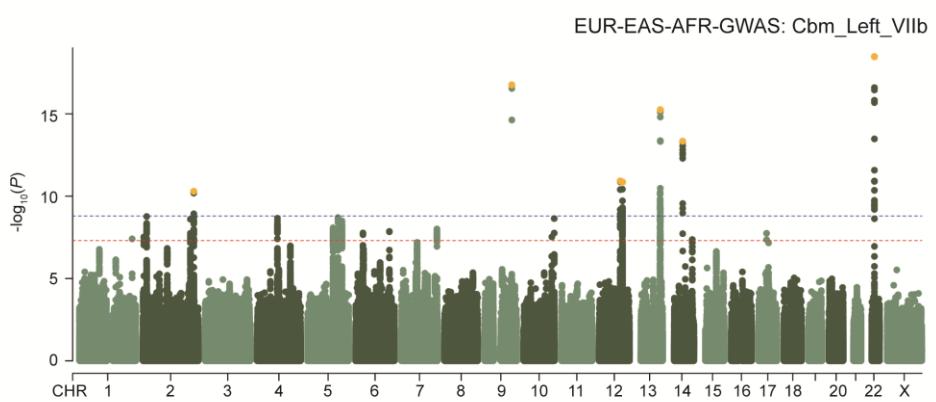
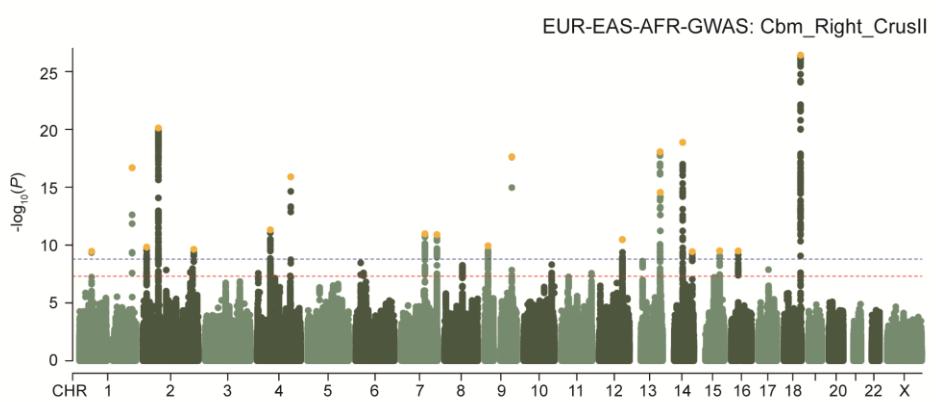
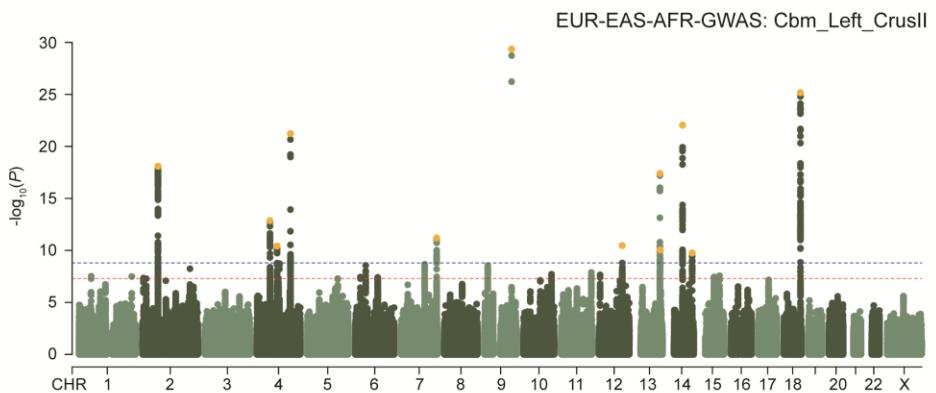


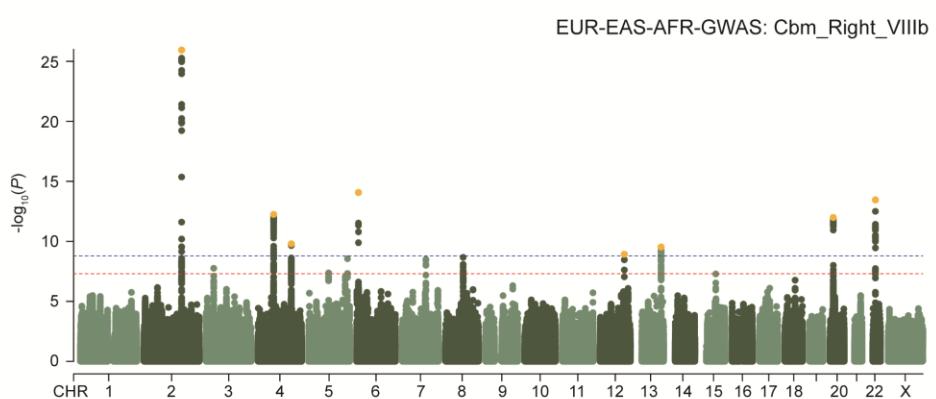
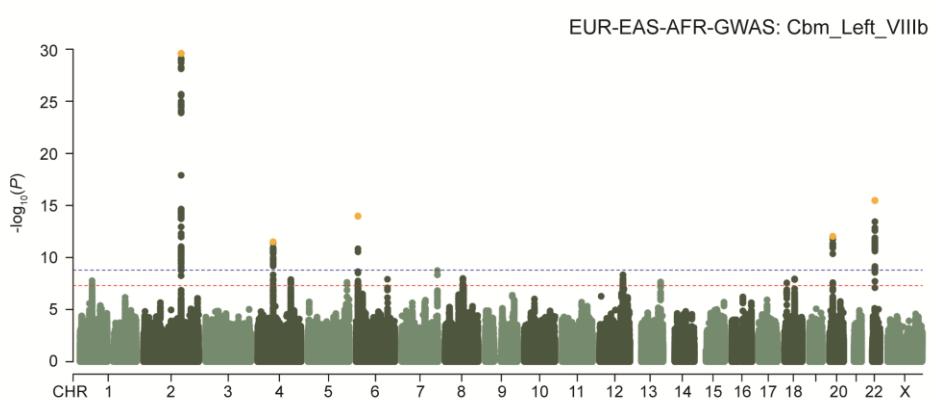
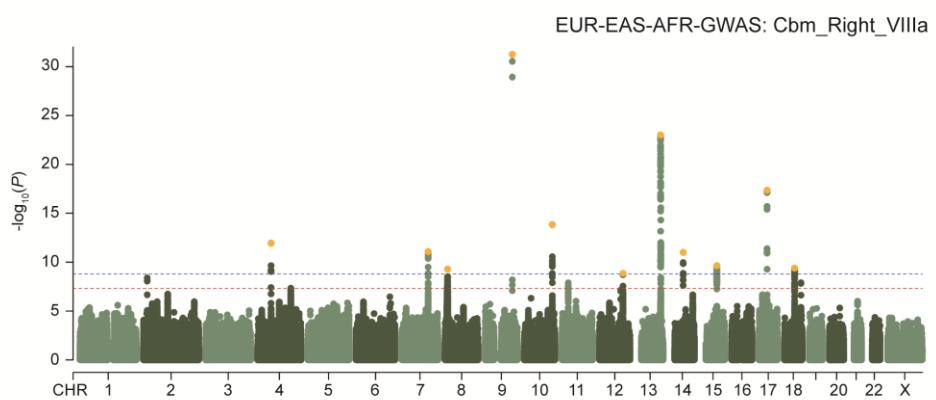
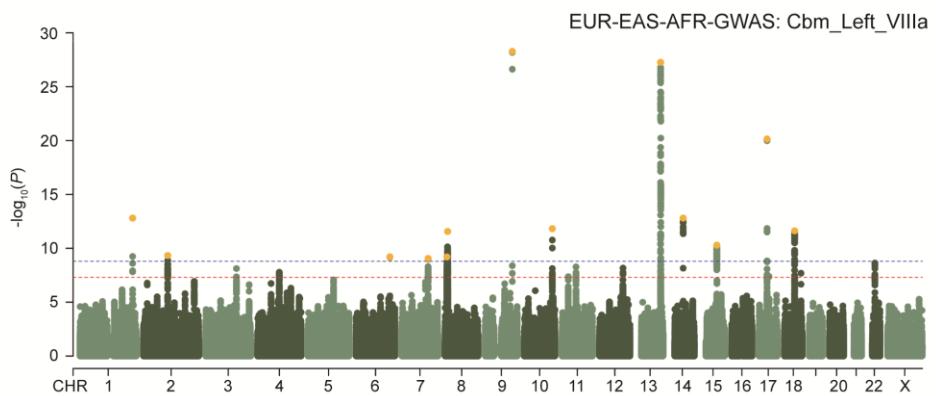
EUR-EAS-AFR-GWAS: Cbm_Left_CrusI

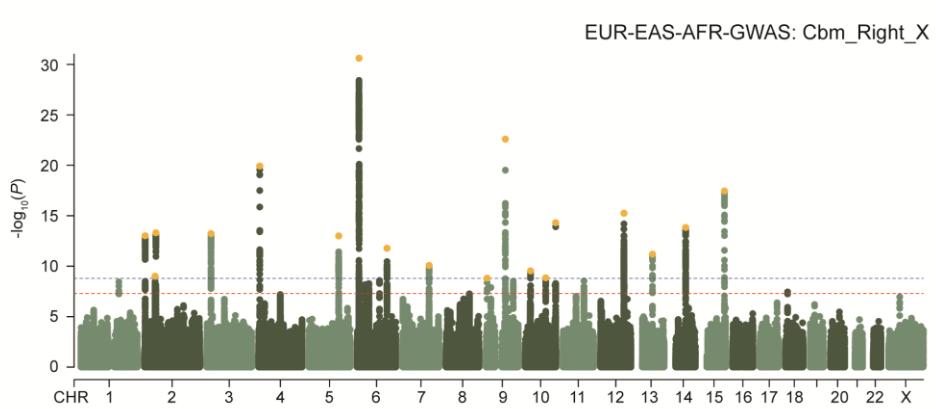
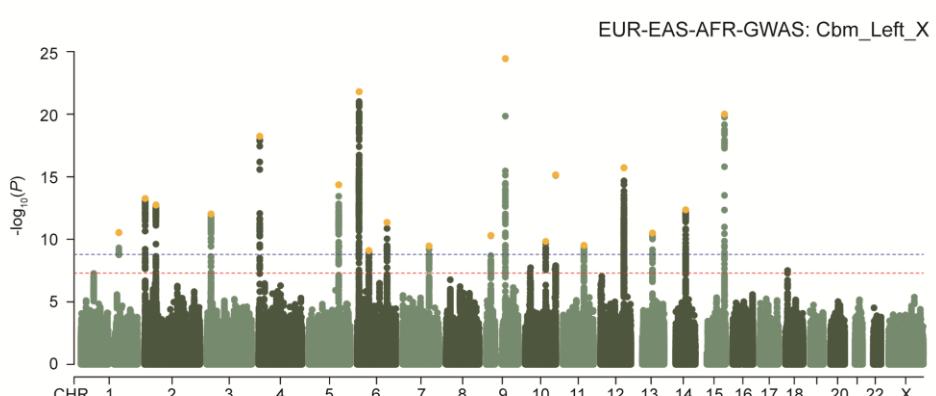
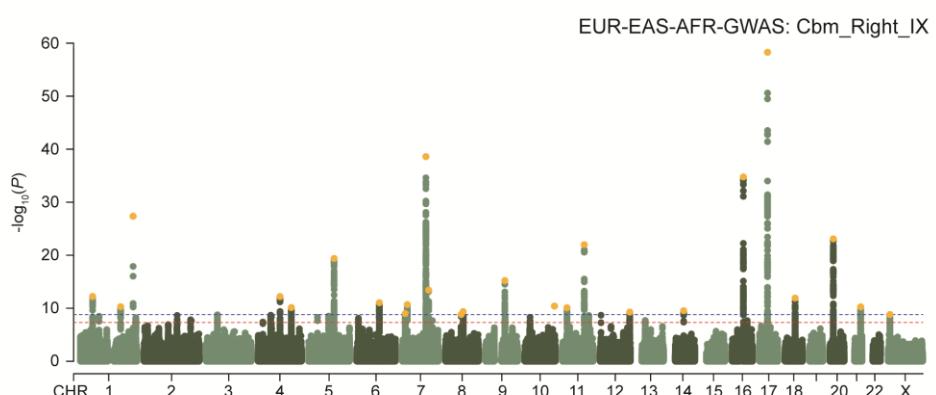
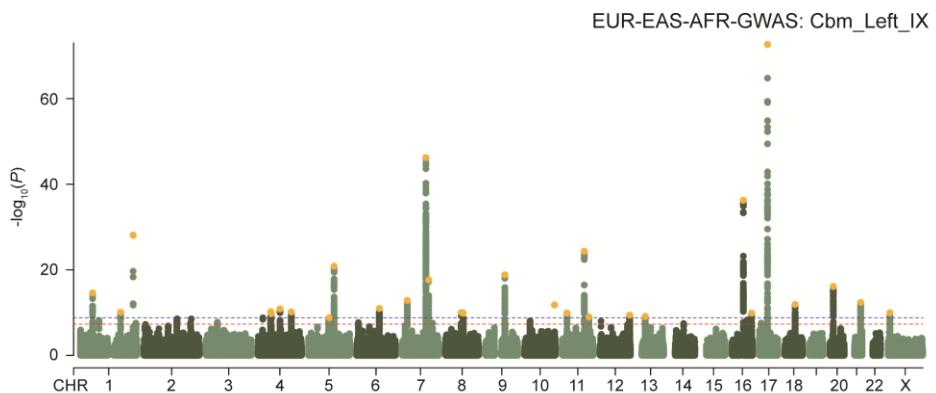


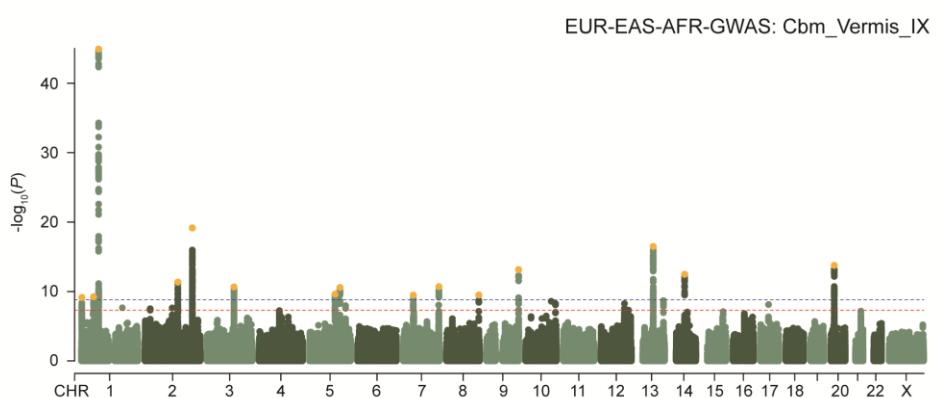
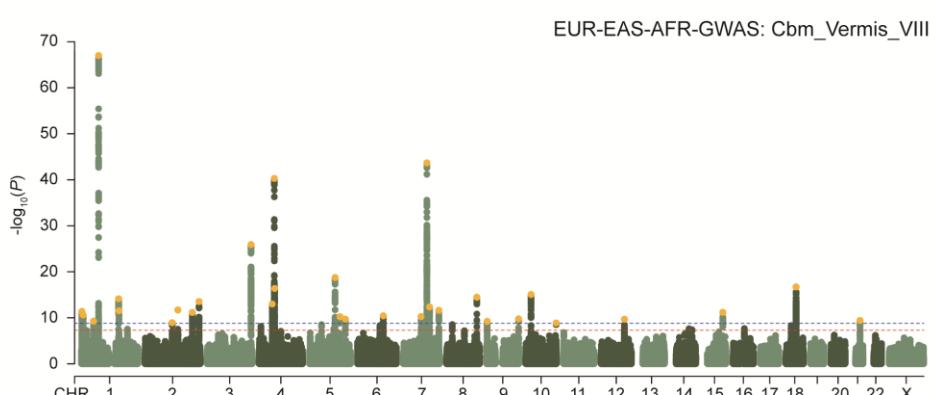
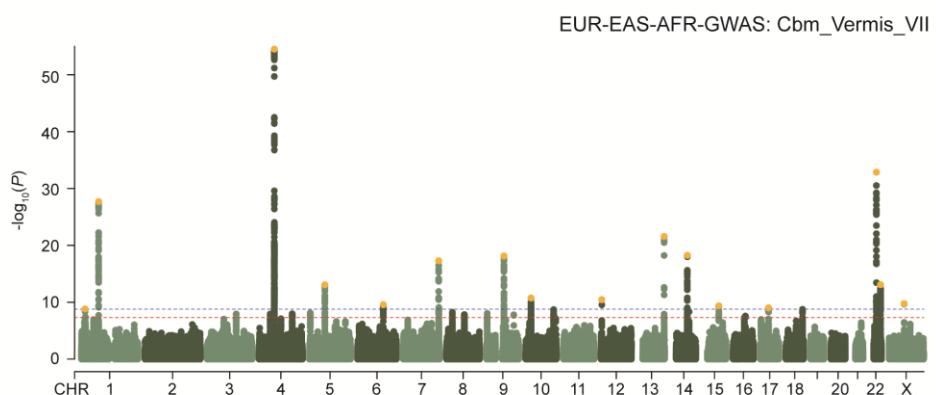
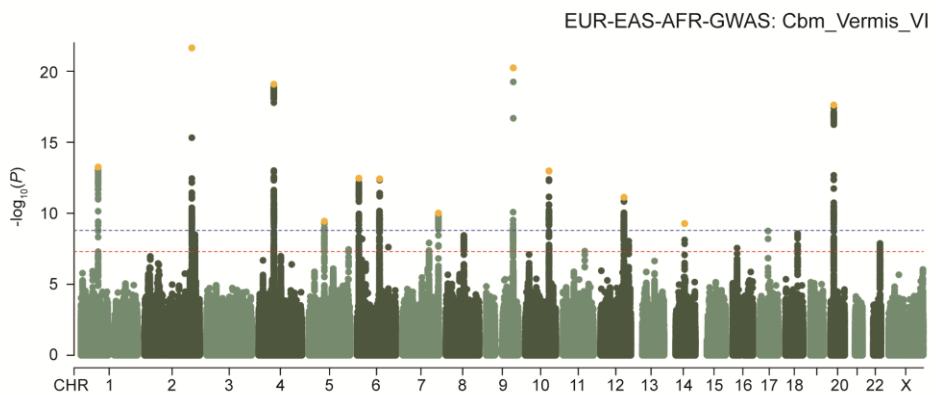
EUR-EAS-AFR-GWAS: Cbm_Right_CrusI











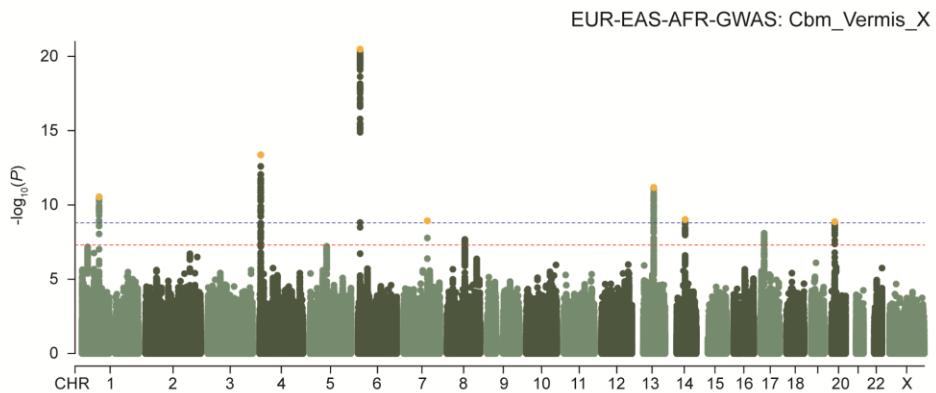
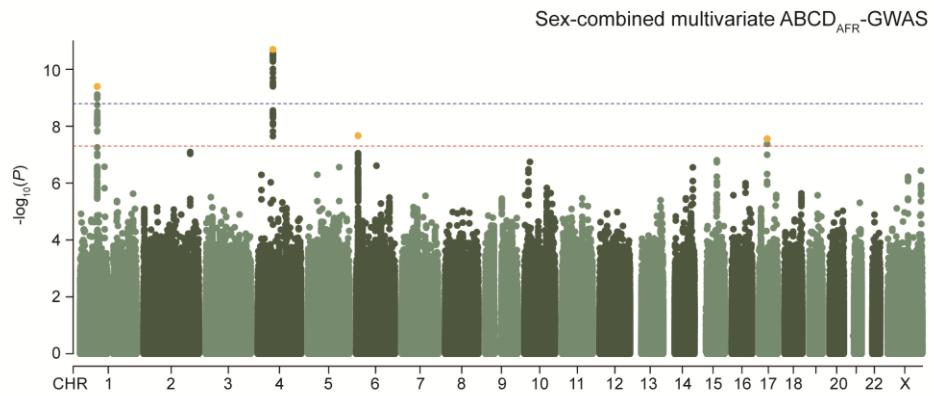
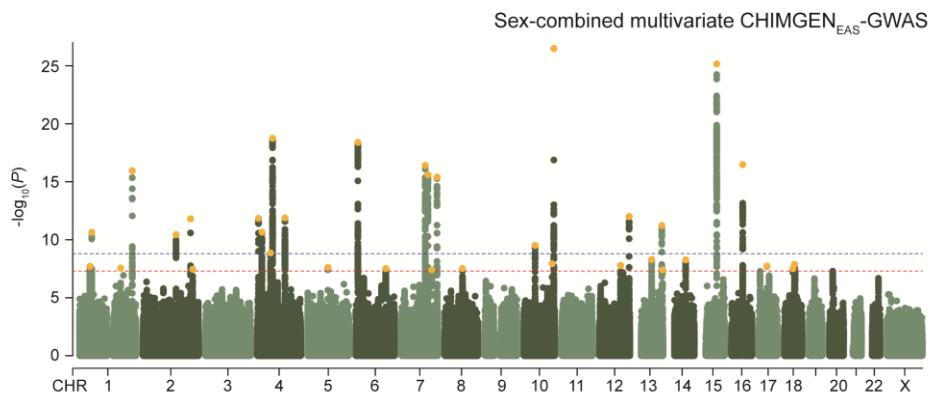
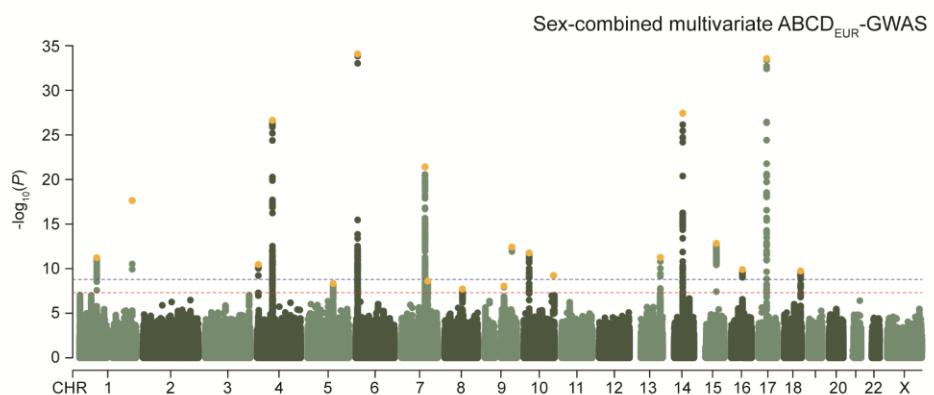
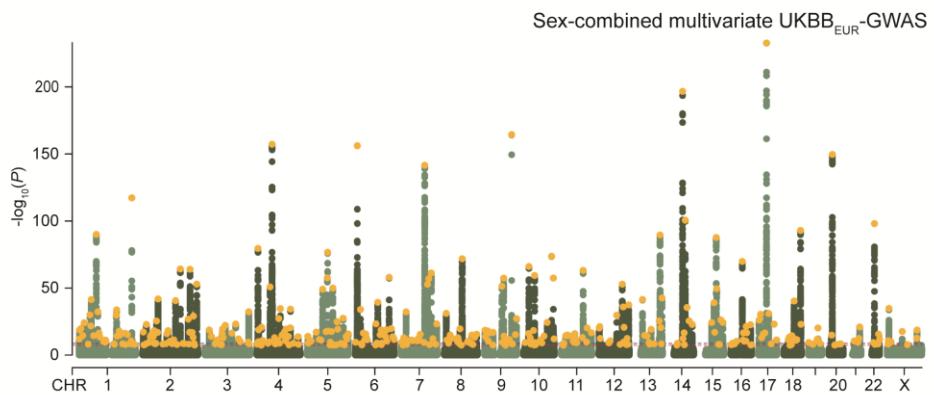
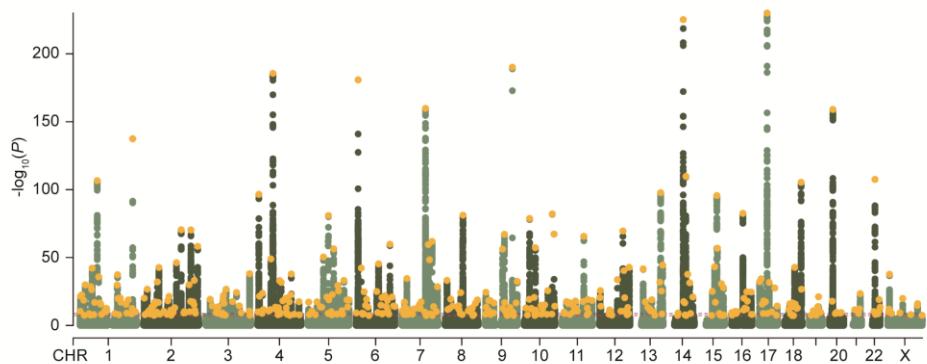


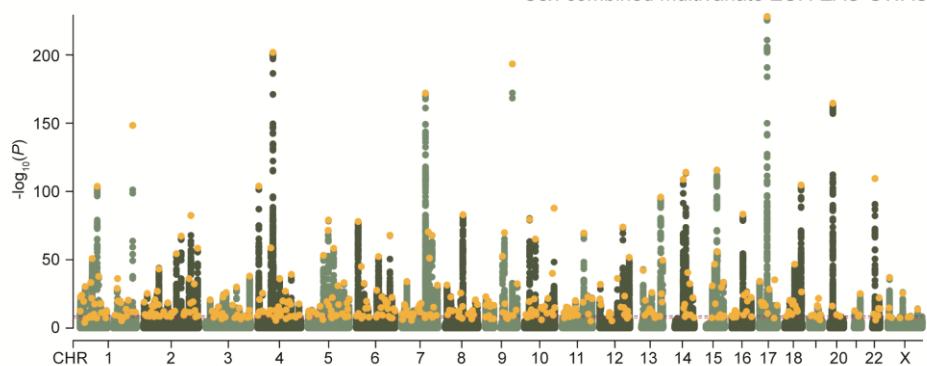
Fig. S5 | Genetic discovery of female-specific univariate GWAS. The Manhattan plots illustrate the results from female-specific univariate GWASs (UKBB_{EUR}-GWAS, ABCD_{EUR}-GWAS, CHIMGEN_{EAS}-GWAS, ABCD_{AFR}-GWAS, EUR-GWAS, EUR-EAS-GWAS, and EUR-EAS-AFR-GWAS) on the 31 cerebellar volumetric traits. Each plot shows genome-wide (red dashed line: $P = 5 \times 10^{-8}$) and study-wide (blue dashed line: $P = 1.61 \times 10^{-9}$) significant associations. Each point represents a single genetic variant plotted according to its genomic position (x axis) and its $-\log_{10}(P)$ value for the association (y axis). Each yellow point represents an independent locus with study-wide significance ($P = 1.61 \times 10^{-9}$). Abbreviations: ABCD, the Adolescent Brain Cognitive Development study; AFR, African ancestry; CHIMGEN, the Chinese Imaging Genetics study; EAS, East Asian ancestry; EUR, European ancestry; GWAS, genome-wide association study; UKBB, the UK Biobank study.



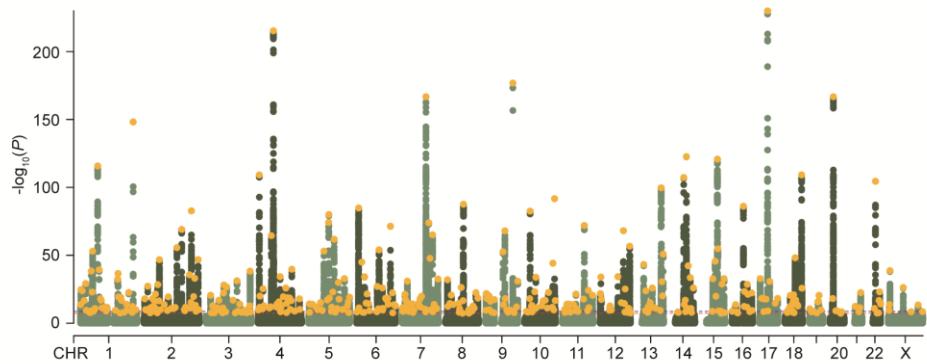
Sex-combined multivariate EUR-GWAS



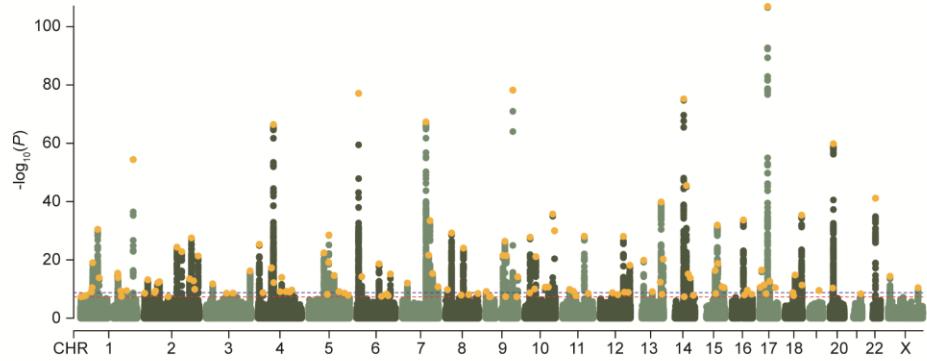
Sex-combined multivariate EUR-EAS-GWAS



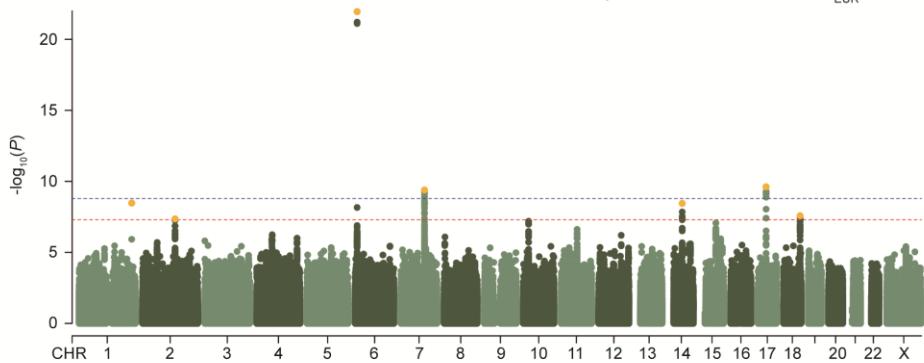
Sex-combined multivariate EUR-EAS-AFR-GWAS



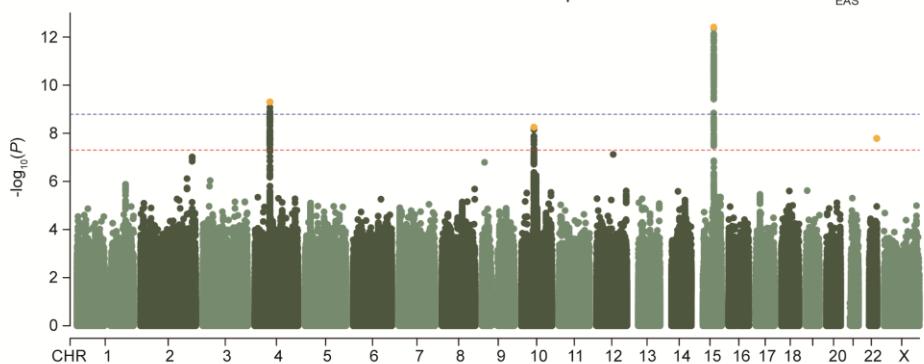
Male-specific multivariate UKBB_{EUR}-GWAS



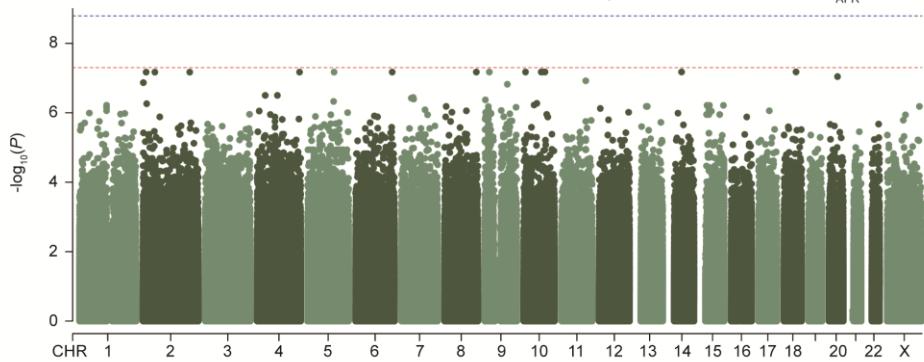
Male-specific multivariate ABCD_{EUR}-GWAS



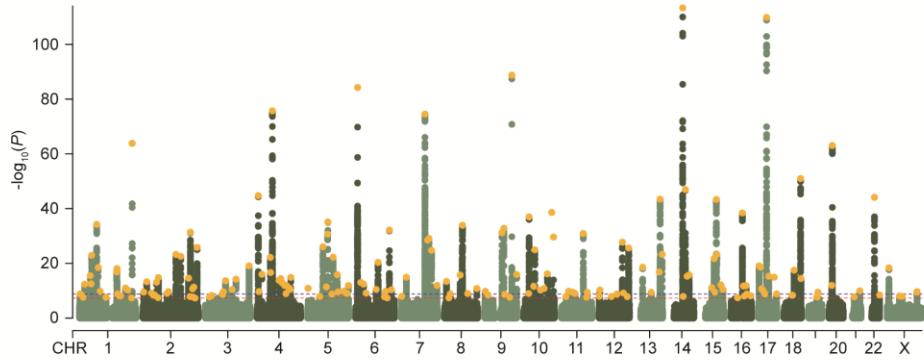
Male-specific multivariate CHIMGEN_{EAS}-GWAS



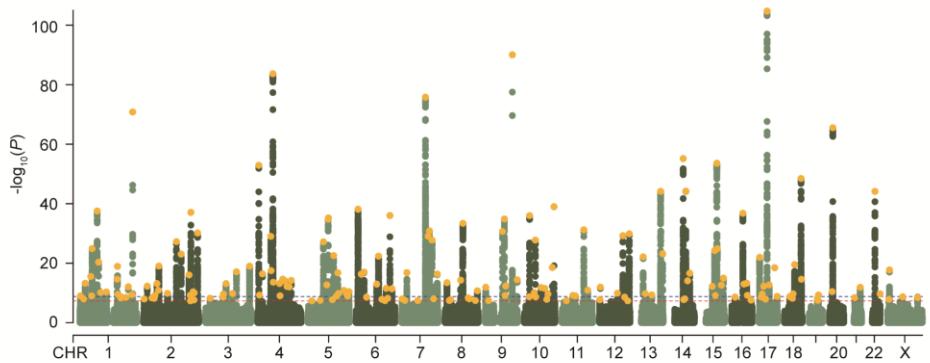
Male-specific multivariate ABCD_{AFR}-GWAS



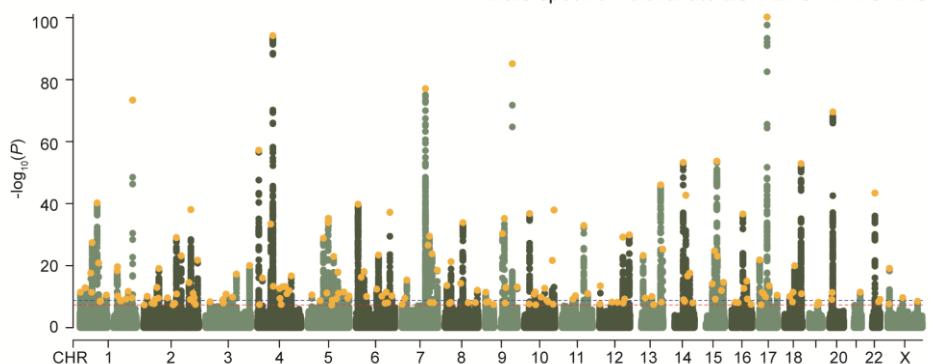
Male-specific multivariate EUR-GWAS



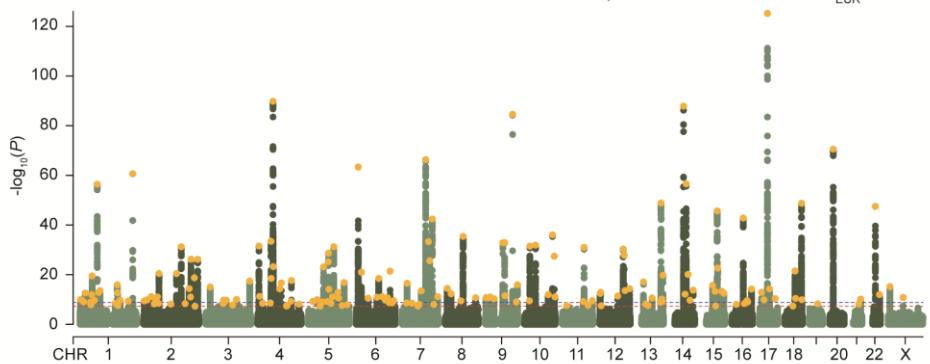
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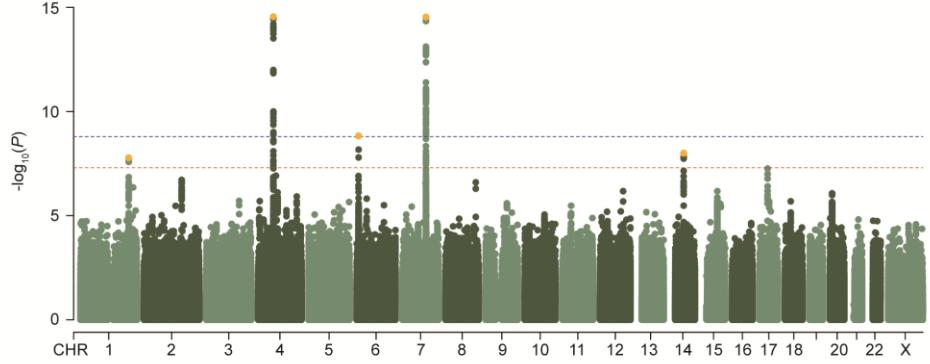
Male-specific multivariate EUR-EAS-AFR-GWAS



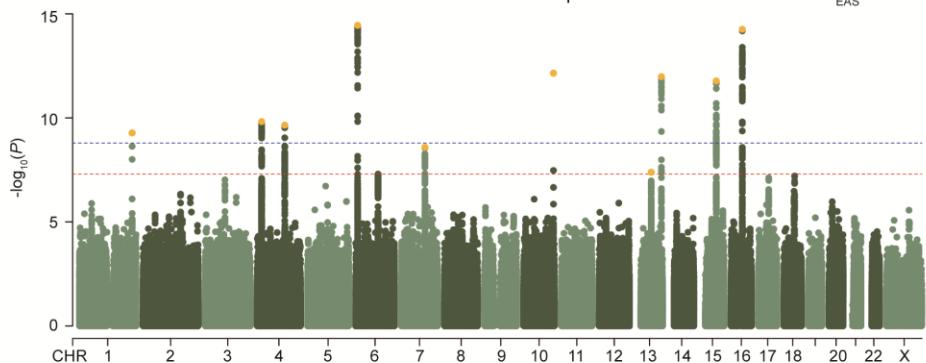
Female-specific multivariate UKBB_{EUR}-GWAS



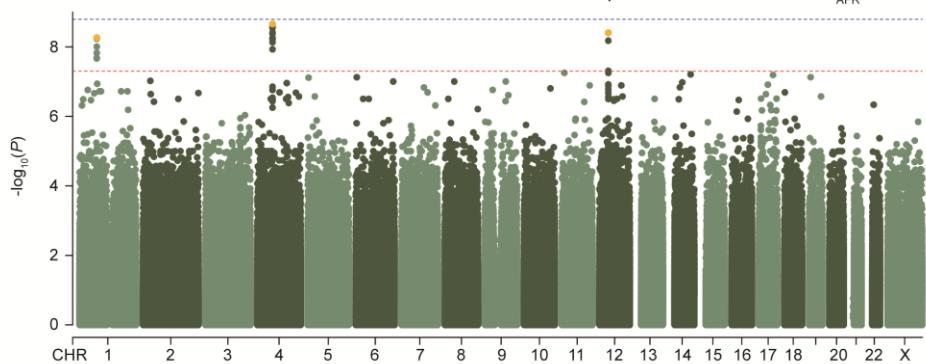
Female-specific multivariate ABCD_{EUR}-GWAS



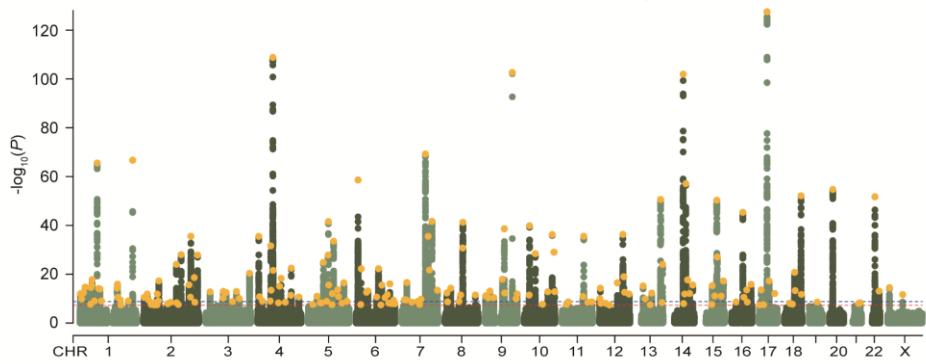
Female-specific multivariate CHIMGEN_{EAS}-GWAS



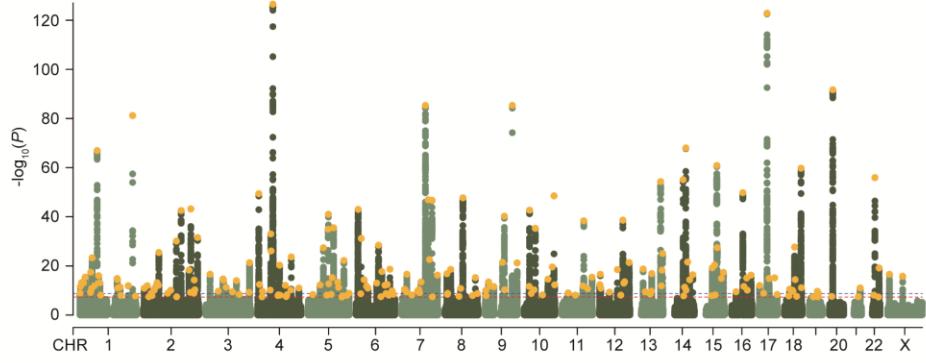
Female-specific multivariate ABCD_{AFR}-GWAS



Female-specific multivariate EUR-GWAS



Female-specific multivariate EUR-EAS-GWAS



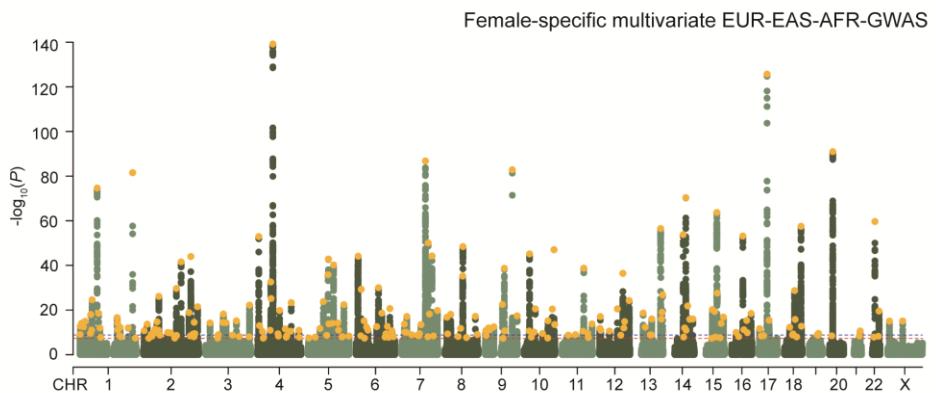


Fig. S6 | Genetic discovery of multivariate GWASs. The Manhattan plots show the results of multivariate GWASs on 27 non-overlapping cerebellar volumetric traits based on sex-combined, male-specific, and female-specific univariate GWASs. Each plot shows both genome-wide (red dashed line: $P = 5 \times 10^{-8}$) and study-wide (blue dashed line: $P = 1.61 \times 10^{-9}$) significance thresholds. Each point represents a single genetic variant plotted according to its genomic position (x axis) and its $-\log_{10}(P)$ value for the association (y axis). Each yellow point represents an independent locus with genome-wide significant association ($P = 5 \times 10^{-8}$). Abbreviations: ABCD, the Adolescent Brain Cognitive Development study; AFR, African ancestry; CHIMGEN, the Chinese Imaging Genetics study; EAS, East Asian ancestry; EUR, European ancestry; GWAS, genome-wide association study; UKBB, the UK Biobank study.

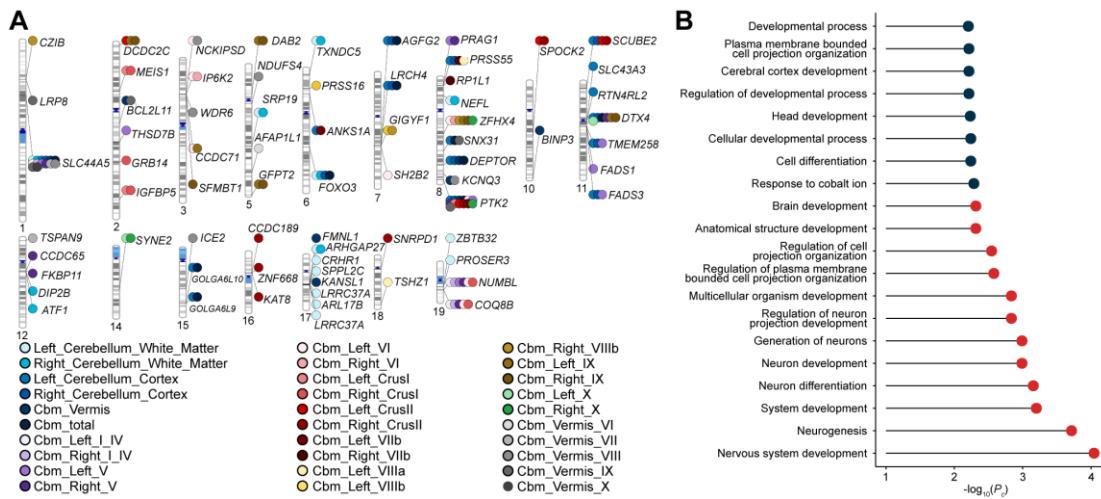


Fig. S7 | Prioritized causal genes and their enriched pathways. **(A)** The ideogram shows the prioritized causal genes identified by the genetic colocalization (PP.H4 > 0.8) analysis between locus-trait associations from EUR-GWASs and eQTLs in cerebellar tissue. **(B)** The Lollipop chart shows the top 20 pathways enriched by the 71 prioritized causal genes. The significant pathways are red colored ($P_c < 0.05$, Benjamini-Hochberg FDR corrected).

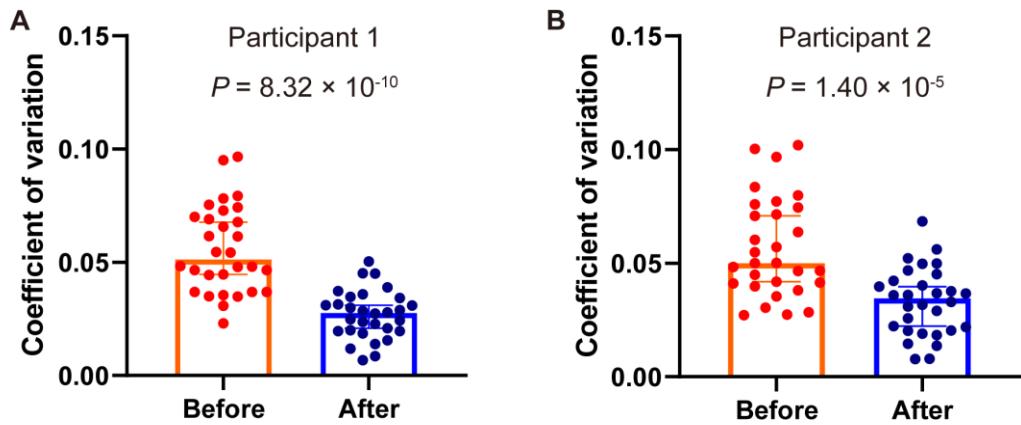


Fig. S8 | Inter-scanner consistency of 30 cerebellar substructure volumes calculated from CerebNet from the same participant before and after Combat harmonization. Two CHIMGEN participants travel to different centers and are scanned with 28 MRI scanners. In each subject, we calculate 30 cerebellar substructure volumes by CerebNet based on the structural MRI data acquired from each scanner and computed CV of each trait across scanners. After applying Combat harmonization to cerebellar substructure volumes, we re-calculate the CV. The Wilcoxon sum rank test is used to investigate the differences in CVs before (red dots) and after (blue dots) harmonization and we found that CVs of all traits before harmonization are reduced after harmonization in both two participants (Wilcoxon rank-sum test, Participant 1: $P = 8.32 \times 10^{-10}$; Participant 2: $P = 1.40 \times 10^{-5}$). The lines indicate the interquartile range of CVs and the box indicate the median of CVs. Abbreviations: CHIMGEN, Chinese imaging genetics; CV, coefficient of variation; MRI, magnetic resonance imaging.