Integrative phenomic analysis of psychosocial factors and their genetic associations in

major depressive disorder using a latent topic model approach

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McGill initiative in

Introduction **Results** Major depressive disorder (MDD) is 1. General analysis pipeline and 2. Inferring meaningful MDD topics ("meta-phenotypes") from 5. Gene score and pathway enrichment analysis a serious public health issue psychosocial questionnaire data latent topic model MDD meta-phenotypes are Both genetic and environmental 6 meta-phenotypes with strong positive (M6, M22, M13) or negative associated with different factors contribute to the clinical (M10, M11, M5) correlation with MDD diagnosis genes and pathways heterogeneity of MDD (Maj et al., 2020) Tonic modelin • To identify meaningful MDD-1111 related latent topics from a set of a de la psychosocial factors Topic 2 LASSO reg valuation by AU To explore the etiopathogenesis of MDD by considering both genetic ******* and psychosocial prepositions. (not at all lin PLINK Methods Zone d'Épidémiologie Psychiatrique 4. SNP-heritability 3. Classification of MDD using inferred meta-phenotypes Conclusion References du Sud-Ouest de Montréal • Higher accuracy for meta-phenotypes compared to (ZEPSOM) cohort (Caron et al., 2012) Caron, J. et al. (2012) Prevalence of raw psychosocial attributes • Topic modeling approach 1351 participants psychological distress and mental LASSO regression → "MDD probability" confers not only accurate disorders, and use of mental health \sim 8 million genetic variants (SNPs) services in the epidemiological catchment MDD risk prediction, but also Receiver operating characteristic curv area of Montreal South-West. BMC ~500 psychosocial factors (e.g., meaningful associations Psychiatry, 12(1), 183 clinical MDD diagnosis, Li, Y. et al. (2020) Inferring multimodal latent between genetic and topics from electronic health records. psychological distress, drug/alcohol

Latent topic model

abuse)

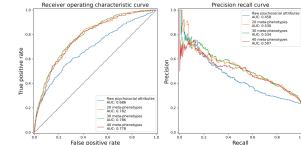
 Collapsed variational Bayesian inference of the LDA model: MixEHR (Li et al., 2020)

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Aims:

Data



- MDD probability (0.126) > diagnosis (1×10^{-6}) • Some meta-phenotypes have higher heritability
- estimates than their top-3 psychosocial attributes
- psychosocial prepositions
- Pathway analysis using MDD meta-phenotypes reveals different underlying pathways which can help interpret MDD clinical heterogeneity
- Nature Communications, 11(1), 2536.
- Mai. M. et al. (2020) The clinical characterization of the adult patient with depression aimed at personalization of management. World Psychiatry, 19, 269-293.