(Pharmaco)genetics in schizophrenia

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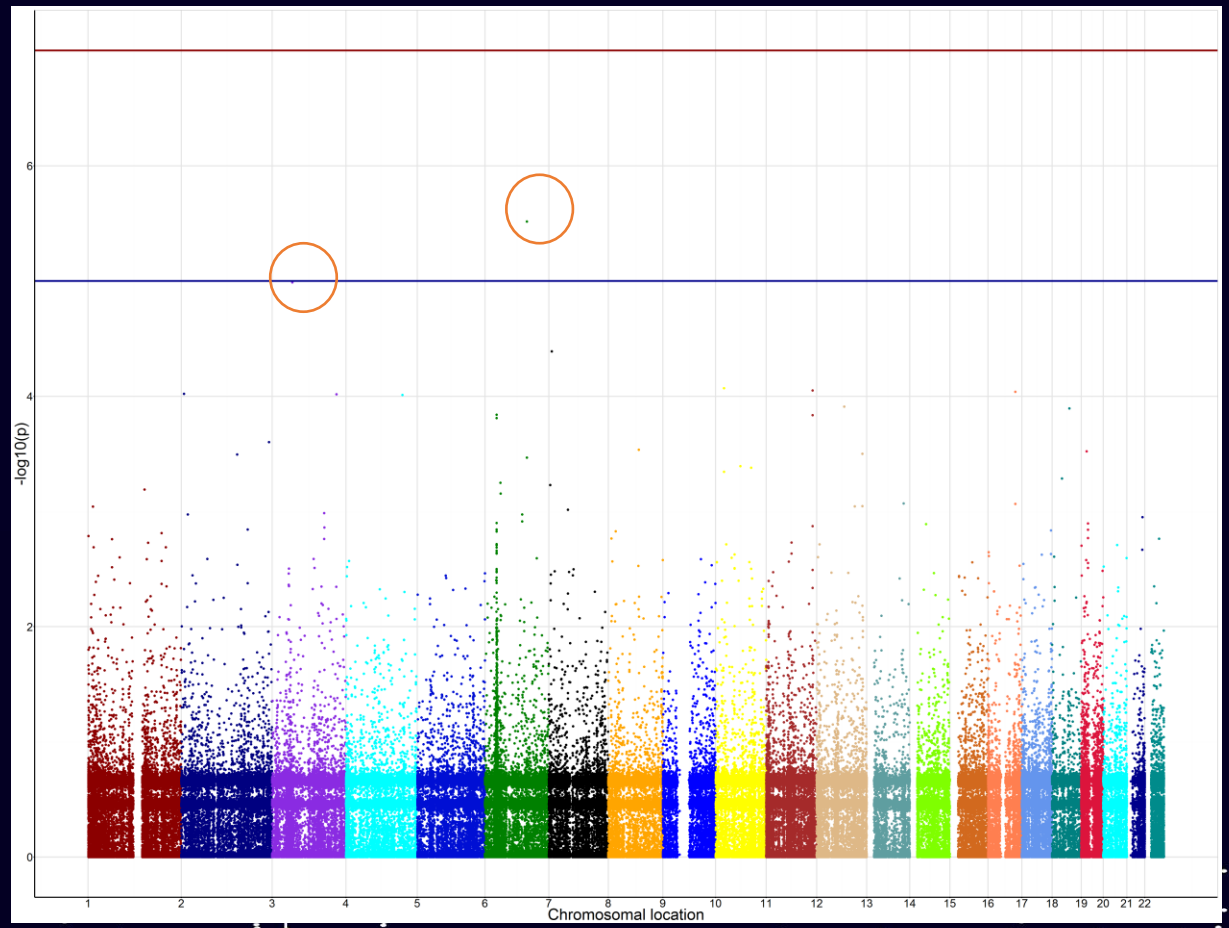
Case-control study of exonic variation

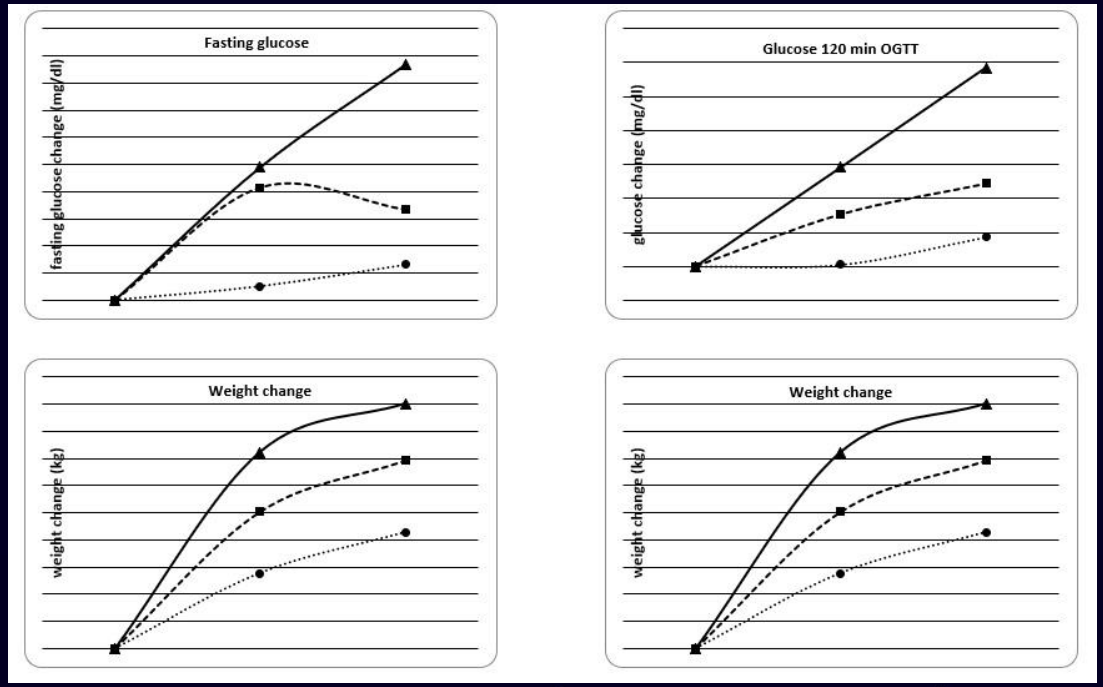
- 493 patients with schizophrenia or schizoaffective D
- 484 healthy controls

- No significant findings
  - Best 2 genes
    - WISP3 (6q21 region):
      - rs1230345 ($p=3.05*10^{-6}$)
    - CACNA2D3
      - rs9311525 ($p=1.03*10^{-6}$) and rs1558557 ($p=3.85*10^{-5}$)



Genetic influence on metabolic syndrome

- Genetic association studies:
  - MTHFR
    - Prospective design
    - C allele of A1298C had the worst metabolic outcomes
  - Circadian clock and involvement in metabolic syndrome
    - Cross-sectional study
    - NR3C1 rs6196 A allele carriers had worse metabolic parameters



IGF variation and DNA methylation

- Cross-sectional study of genetic and epigenetic variantion
  - DNA variation:
    - 27 SNPs in Insulin Growth Factors
    - 7 differentially methylated regions (Methylation as epigenetic marker)
  - No significant findings related to metabolic outcomes
  - But: clear association between genetic variation and methylation
- Note: Current literature on (hypothesis driven) genetic association studies