# Genes Associated with Cardiovascular Risk in Anxiety/Stress Disorders

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September 25, 2019



## Relationship to genetics

- Brain-heart interaction in psychological disorders not studied thoroughly
- ▶ GWAS studies not focused on brain-heart interactions
- ► Could be used to identify at-risk individuals

#### Methods

- NCBI Phenotype-Genotype integration system (merges NHGRI, dbGaP, OMIM, eQTL, and dbSNP)
- MeSH term search for phenotype identification: arrhythmia, CAD, depression, anxiety, psychological stress, PTSD
- ▶ significance threshold of p < 1e-5 for gene selection
- script to find overlapping phenotypes and genotypes
- manual annotation of results

### Results

#### Overview

- ▶ 1095 genes initially identified
- data included gene location, chromosome, location, and relevant studies
- ▶ identified 41 genes of appropriate phenotype-genotype overlap
- further annotated results by significance of relationship/overlap and primary literature

#### BIN1

- ▶ BIN1 (chromosome 2, rs10207628),
- bridging integrator for synaptic vesicle endocytosis
- associated with SCD and depression, expressed in brain
- regulates calcium ion transport, cardiac AP, neuronal differentiation
- locates to cardiac T-tubules, releases microparticles<sup>1,2</sup>
- ▶ assocated with tau protein in dementia<sup>3</sup>

#### PHARCTR1

- ► PHARCTR1 (chromosome 6, rs4615376)
- phosphatase and actin regulator of endothelial cell survival
- associated with CAD and depression and expressed in the brain and heart
- migraine pathogenesis<sup>4</sup> and susceptibility for CAD in type 2 diabetes mellitus<sup>5</sup>

#### CNNM2

- CNNM2 (chromosome 10, rs12413409),
- divalent metal cation transport mediator
- associated with CAD and depression and expressed in the kidneys
- magnesium ion homeostasis and transmembrane transport
- common among major psychiatric disorders<sup>6</sup>
- ▶ susceptibility locus in CAD<sup>7</sup> and increases the risk of hypertension<sup>8</sup>

#### PRTFDC1

- ► PRTFDC1 (chromosome 10, rs11014306)
- phosphoribosyl transferase domain protein
- associated with SCD and anxiety/stress disorders and is expressed in the brain and adrenal glands
- functions as part of purine salvage pathways
- cardiometabolic profile in heart failure<sup>9</sup>
- predictor of combat stress vulnerability in the development of PTSD<sup>10</sup>

#### CDH13

- CDH13 (chromosome 16, rs8055236)
- cadherin protein
- associated with CAD and depression and is expressed in the brain and heart
- functions in protein signal transduction, endothelial cell migration, and is protective against apoptosis
- provides resistance to atherosclerosis, and is part of neural differentiation
- ▶ role in the cardioprotective effects of sleep and in incident coronary artery disease<sup>11</sup>
- identified in hyperactivity, impulsivity, violent behavior, and extraverted personality traits<sup>6,12,13</sup>

#### BMP2

- ► BMP2 (chromosome 20, rs6117734)
- bone morphogenetic transforming growth factor-beta protein
- associated with CAD and depression and is expressed broadly
- functions as part of cardiac epithelial transition and cardiomyocyte differentiation
- associated with depressive traits and stressful life events<sup>14</sup>
- ▶ also in cardiac progenitor cell differentiation<sup>15</sup> and atherosclerosis in type 2 diabetes mellitus<sup>16</sup>

# Additional genes identified I

- ► RORA (chromosome 15, rs12912233)
- ► GRIN2A (chromosome 16, rs8058295)
- ► FAM155A (chromosome 13, rs1509091)
- ENOX1 (chromosome 13, rs17538444)
- QKI (chromosome 6, rs7756185)
- ► EGFLAM (chromosome 5, rs2561805)
- ► SNX7 (chromosome 1, rs11581859)
- ► ACVR1 (chromosome 2, rs35806662)
- ► MYL10 (chromosome 7, rs1722229)
- KSR2 (chromosome 12, rs7973260)
- ► PARVA (chromosome 11, rs7120489)
- SNCA (chromosome 4, rs356228)
- SORCS3 (chromosome 10, rs7074335)
- MAML3 (chromosome 4, rs1877075)
- DCLK2 (chromosome 4, rs150175932)
- LDDDE (abromacomo 1 val 200461)
- ► LPPR5 (chromosome 1, rs1329461)
- Pseudogenes: RNA5SP87, RPL26P5, RNA5SP404, HSPE1P20, MTCO3P1, RPL6P18, and MTCL1P1

## References

#### References I

- 1. Xu B, Fu Y, Liu Y, et al. The ESCRT-III pathway facilitates cardiomyocyte release of cBIN1-containing microparticles. Lo C, ed. *PLoS Biology.* 2017;15(8):e2002354. doi:10.1371/journal.pbio.2002354
- 2. Hong TT, Smyth JW, Gao D, et al. BIN1 localizes the L-type calcium channel to cardiac T-tubules. Chien KR, ed. *PLoS Biology*. 2010;8(2):e1000312. doi:10.1371/journal.pbio.1000312
- 3. Zhou Y, Hayashi I, Wong J, Tugusheva K, Renger JJ, Zerbinatti C. Intracellular clusterin interacts with brain isoforms of the bridging integrator 1 and with the microtubule-associated protein Tau in Alzheimer's Disease. Götz J, ed. *PLoS ONE*. 2014;9(7):e103187. doi:10.1371/journal.pone.0103187
- 4. Anttila V, Winsvold BS, Gormley P, et al. Genome-wide meta-analysis identifies new susceptibility loci for migraine. *Nature Genetics*. 2013;45(8):912-917. doi:10.1038/ng.2676

#### References II

- 5. Qi L, Parast L, Cai T, et al. Genetic susceptibility to coronary heart disease in type 2 diabetes: 3 independent studies. *Journal of the American College of Cardiology.* 2011;58(25):2675-2682. doi:10.1016/j.jacc.2011.08.054
- 6. Smoller JW, Kendler K, Craddock N, et al. Identification of risk loci with shared effects on five major psychiatric disorders: a genome-wide analysis. *The Lancet*. 2013;381(9875):1371-1379. doi:10.1016/S0140-6736(12)62129-1
- 7. Schunkert H, König IR, Kathiresan S, et al. Large-scale association analysis identifies 13 new susceptibility loci for coronary artery disease. *Nature Genetics*. 2011;43(4):333-340. doi:10.1038/ng.784
- 8. Liu X, Chen L, Zhang Y, et al. Associations between polymorphisms of the CXCL12 and CNNM2 gene and hypertension risk: A case-control study. *Gene.* 2018;675:185-190. doi:10.1016/j.gene.2018.06.107

#### References III

- 9. Yu B, Zheng Y, Alexander D, et al. Genome-Wide association study of a heart failure related metabolomic profile among african americans in the atherosclerosis risk in communities (ARIC) study. *Genetic Epidemiology*. 2013;37(8):840-845. doi:10.1002/gepi.21752
- 10. Nievergelt CM, Maihofer AX, Mustapic M, et al. Genomic predictors of combat stress vulnerability and resilience in U.S. Marines: A genome-wide association study across multiple ancestries implicates PRTFDC1 as a potential PTSD gene. *Psychoneuroendocrinology*. 2015;51:459-471. doi:10.1016/j.psyneuen.2014.10.017
- 11. Li G, Feng D, Wang Y, et al. Loss of Cardio-Protective Effects at the CDH13 Locus Due to Gene-Sleep Interaction: The BCAMS Study. *EBioMedicine*. 2018;32:164-171. doi:10.1016/j.ebiom.2018.05.033
- 12. Tiihonen J, Rautiainen MR, Ollila HM, et al. Genetic background of extreme violent behavior. *Molecular Psychiatry*. 2015;20(6):786-792. doi:10.1038/mp.2014.130

#### References IV

- 13. Terracciano A, Tanaka T, Sutin AR, et al. Genome-wide association scan of trait depression. *Biological Psychiatry*. 2010;68(9):811-817. doi:10.1016/j.biopsych.2010.06.030
- 14. Ikeda M, Shimasaki A, Takahashi A, et al. Genome-wide environment interaction between depressive state and stressful life events. 2016;77:e29-e30. doi:10.4088/JCP.15l10127
- 15. Bylund JB, Trinh LT, Awgulewitsch CP, et al. Coordinated Proliferation and Differentiation of Human-Induced Pluripotent Stem Cell-Derived Cardiac Progenitor Cells Depend on Bone Morphogenetic Protein Signaling Regulation by GREMLIN 2. *Stem Cells and Development*. 2017;26(9):678-693. doi:10.1089/scd.2016.0226
- 16. Zhang L, Guo T, Xi B, et al. Automatic recognition of cardiac arrhythmias based on the geometric patterns of poincaré plots. *Physiological Measurement*. 2015;36(2):283-301. doi:10.1088/0967-3334/36/2/283