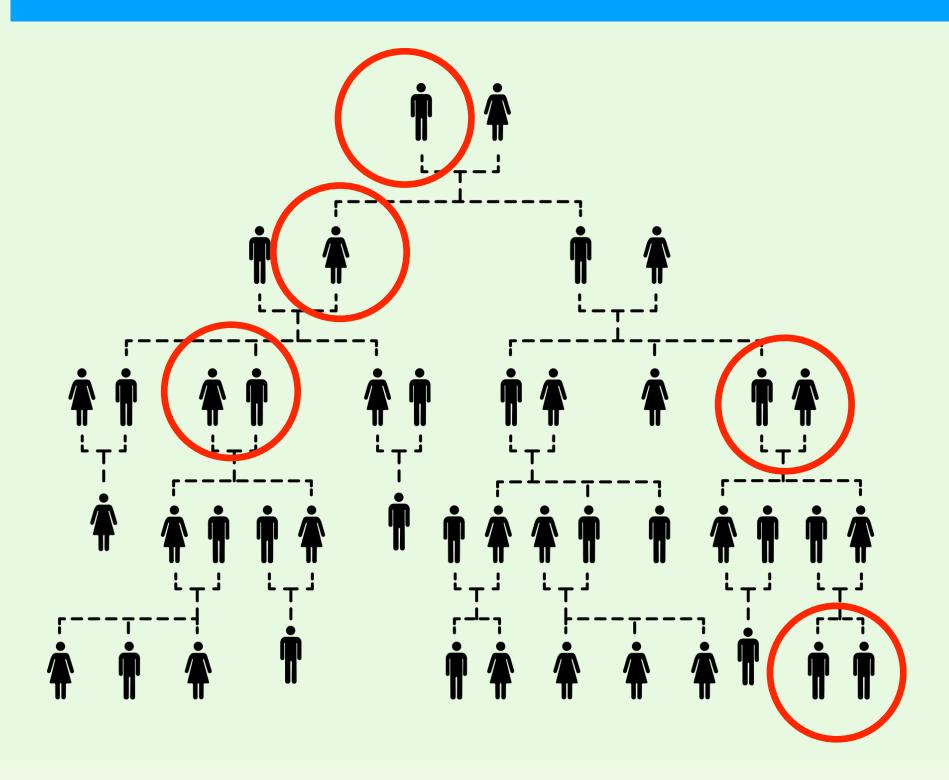


Mikko Kallela - disclosures

Mikko Kallela has served on Advisory Boards for MSD, Allergan, TEVA, Lilly and Lundbeck; has received funding for travel and/or speaker honoraria from MSD, Allergan, TEVA, Novartis, Genzyme and Lundbeck; has received compensation for producing educational material from TEVA and Allergan; has received research support from Helsinki University Central Hospital; and holds stock/stock options and/or has received Board of Directors compensation from Helsinki Headache Center.

Linkage - analysis



Affected and non-affected family members

Rare causal genes such as CACNA1A (calcium channel), ATP1A2 (ion pump), SCN1A (sodium channel)

Genomwide association studies (GWAS)



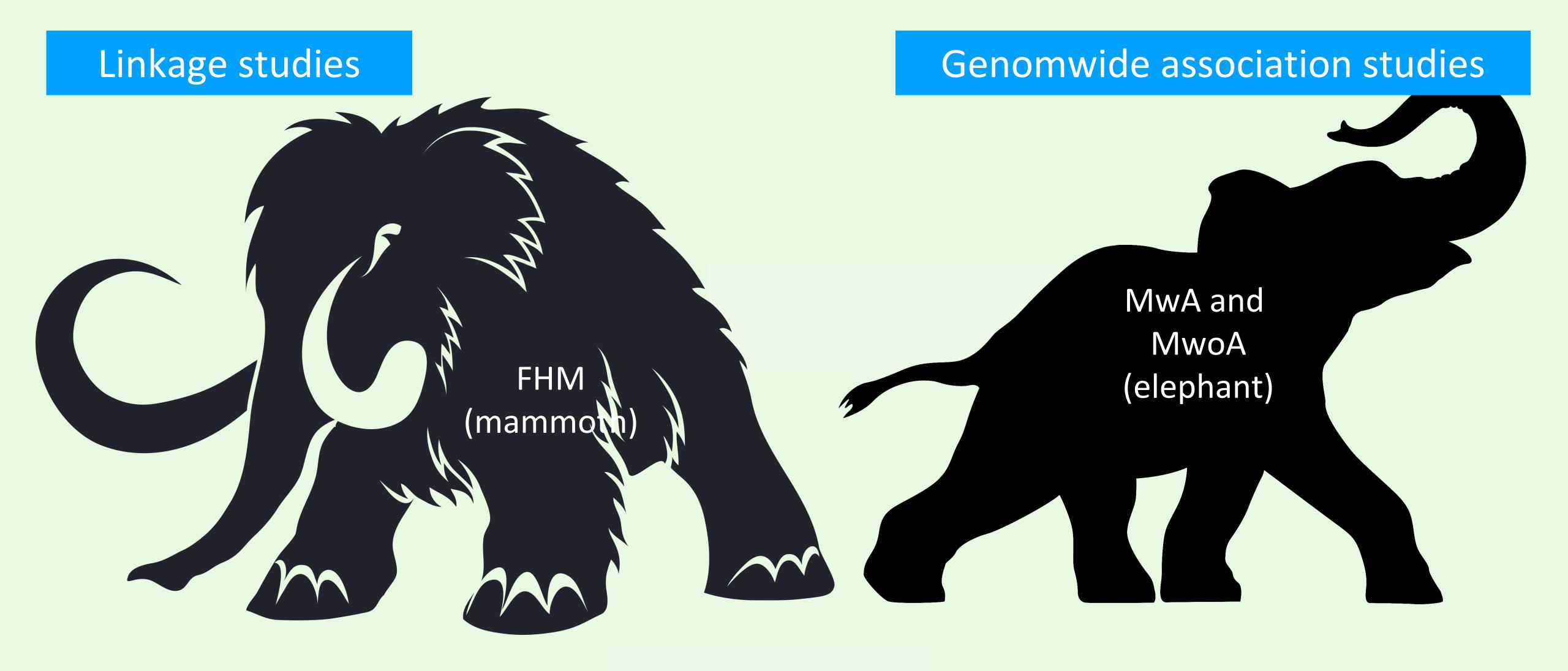
Cases and controls



Common variants - increase the risk to have the disorder, such as TPRM8

Stam AH, van den Maagdenberg AM, Haan J, Terwindt GM, Ferrari MD. Genetics of migraine: an update with special attention to genetic comorbidity. Curr Opin Neurol. 2008;21(3):288-93.

Gormley P, Anttila V, Winsvold BS, Palta P, Esko T, Pers TH, et al. Meta-analysis of 375,000 individuals identifies 38 susceptibility loci for migraine. Nat Genet. 2016;48(8):856-66.



Rare phenotypes like FHM

MwA = migraine with aura MwoA = migraine without aura FHM = familial hemiplegic migraine

Common phenotypes like MwoA and MwoA



Molecular genetic diagnosis



Strange symptoms, no diagnosis, no treatment, no hope, live with it

1.2.3.1.1 CACNA1A (calcium channel)

1.2.3.1.2 ATP1A2 (ion pump)

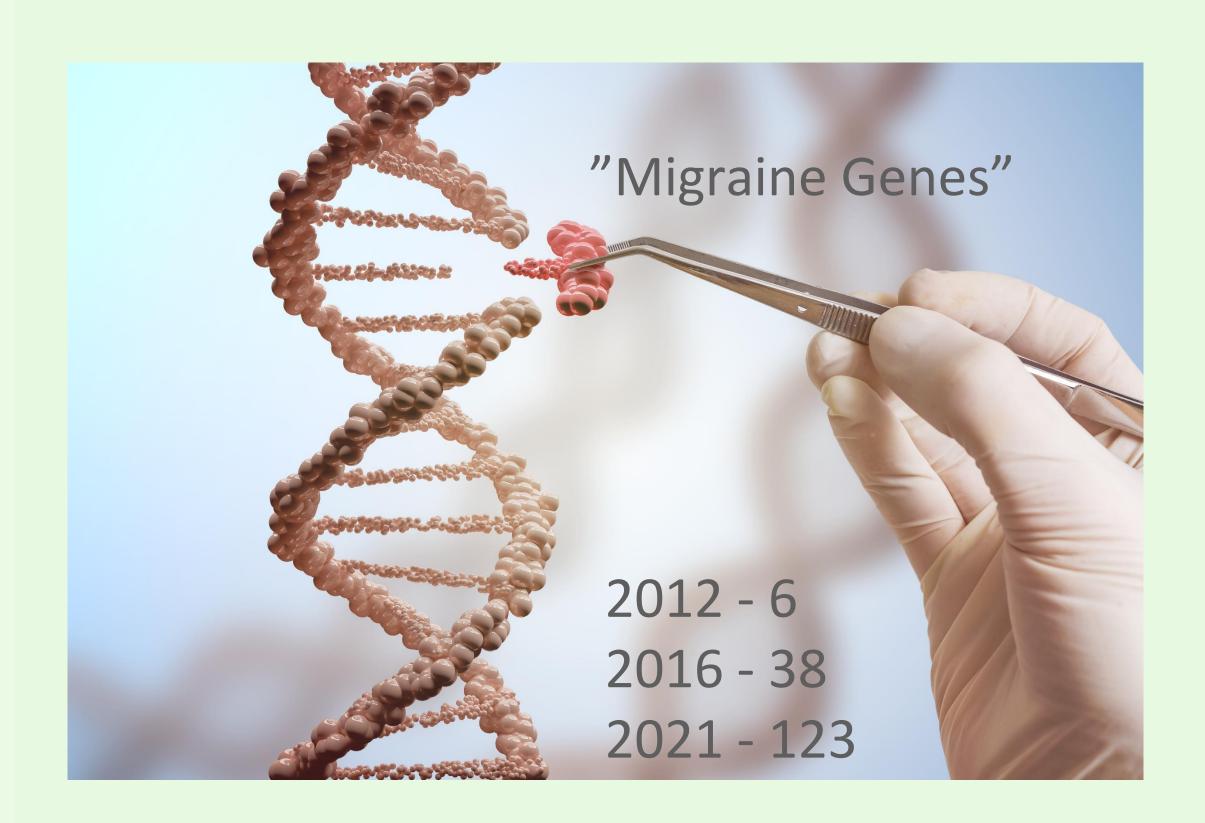
1.2.3.1.3 SCN1A (sodium channel)

Strange symptoms, proper diagnosis, treatment and prognosis

Part of migraine is a chanellopathy

Carrera P, Stenirri S, Ferrari M, Battistini S. Familial hemiplegic migraine: a ion channel disorder. Brain Res Bull. 2001;56(3-4):239-41.

Genomwide association studies (GWAS)



Common variants - increase the risk to have the disorder, such as MEF2D, TGFBR2, PHACTR1, ASTN2, LRP1, TPRM8



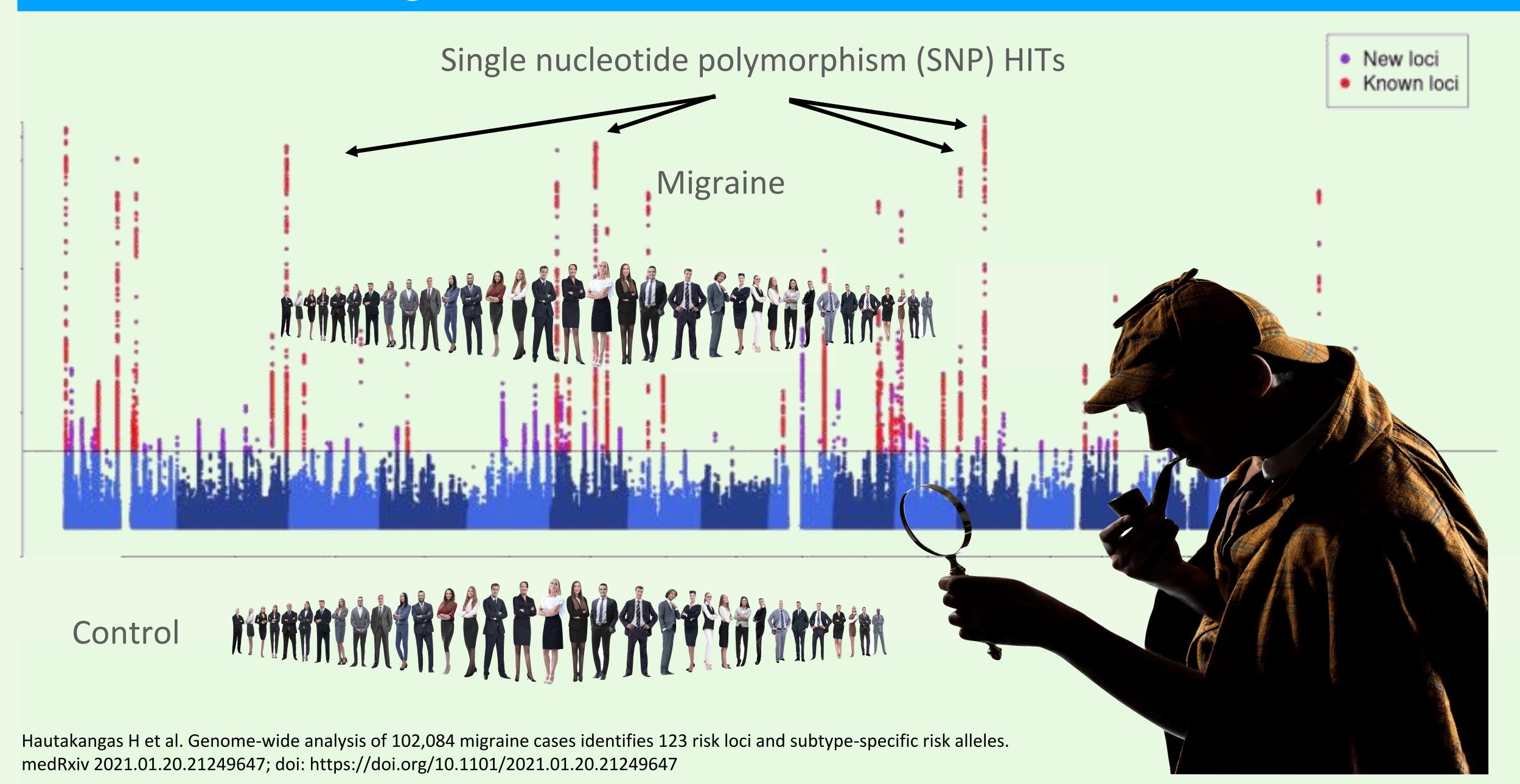
Cases and controls

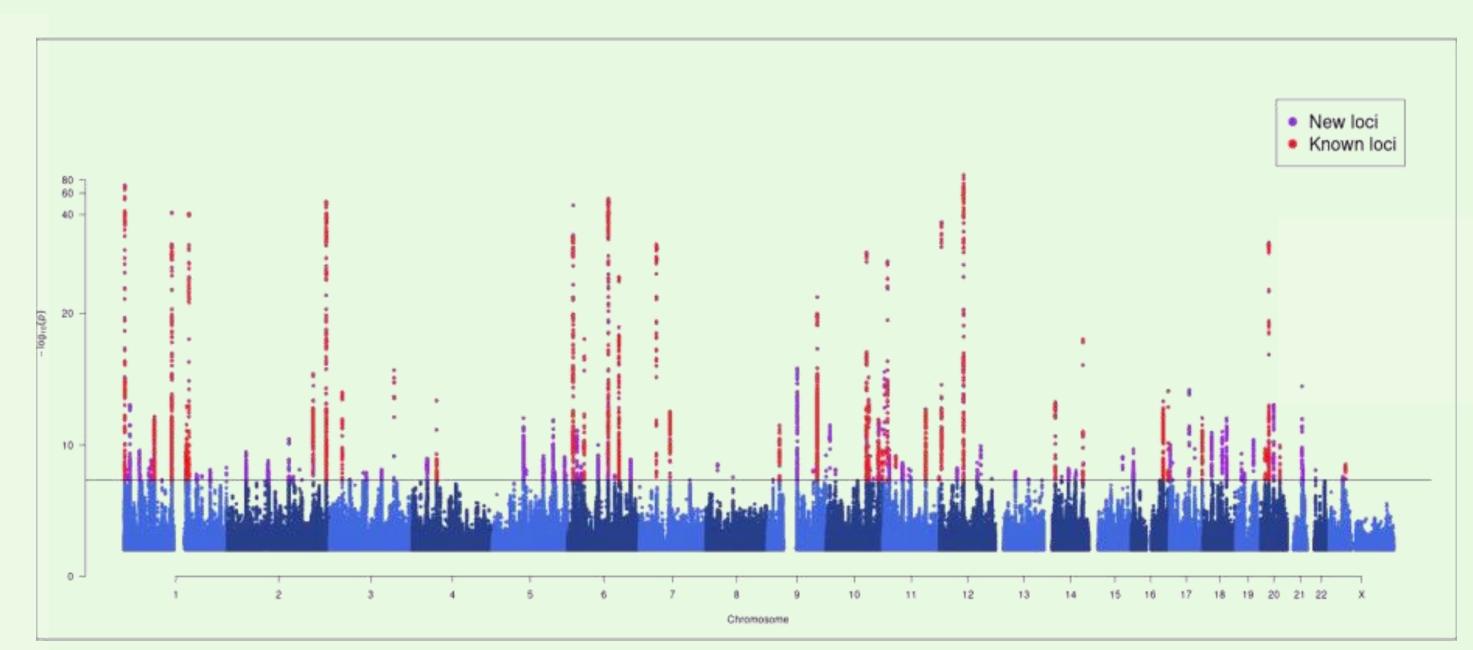


Rare causal genes such as CACNA1A (calcium channel), ATP1A2 (ion pump), SCN1A (sodium channel) can (and will be) be missed

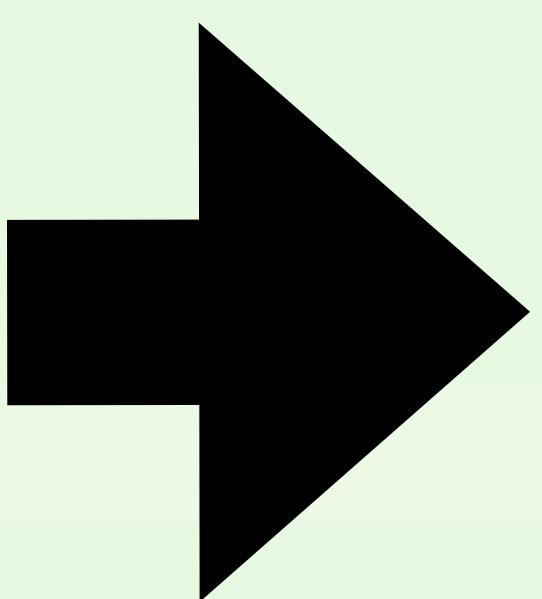
Freilinger T, Anttila V, de Vries B, Malik R, Kallela M, Terwindt GM, et al. Genome-wide association analysis identifies susceptibility loci for migraine without aura. Nat Genet. 2012;44(7):777-82; Gormley P, Anttila V, Winsvold BS, Palta P, Esko T, Pers TH, et al. Meta-analysis of 375,000 individuals identifies 38 susceptibility loci for migraine. Nat Genet. 2016;48(8):856-66; Hautakangas et al. Genome-wide analysis of 102,084 migraine cases identifies 123 risk loci and subtype-specific risk alleles 2021.

100 000 migraine cases - 500 000 controls - 123 SNP hits





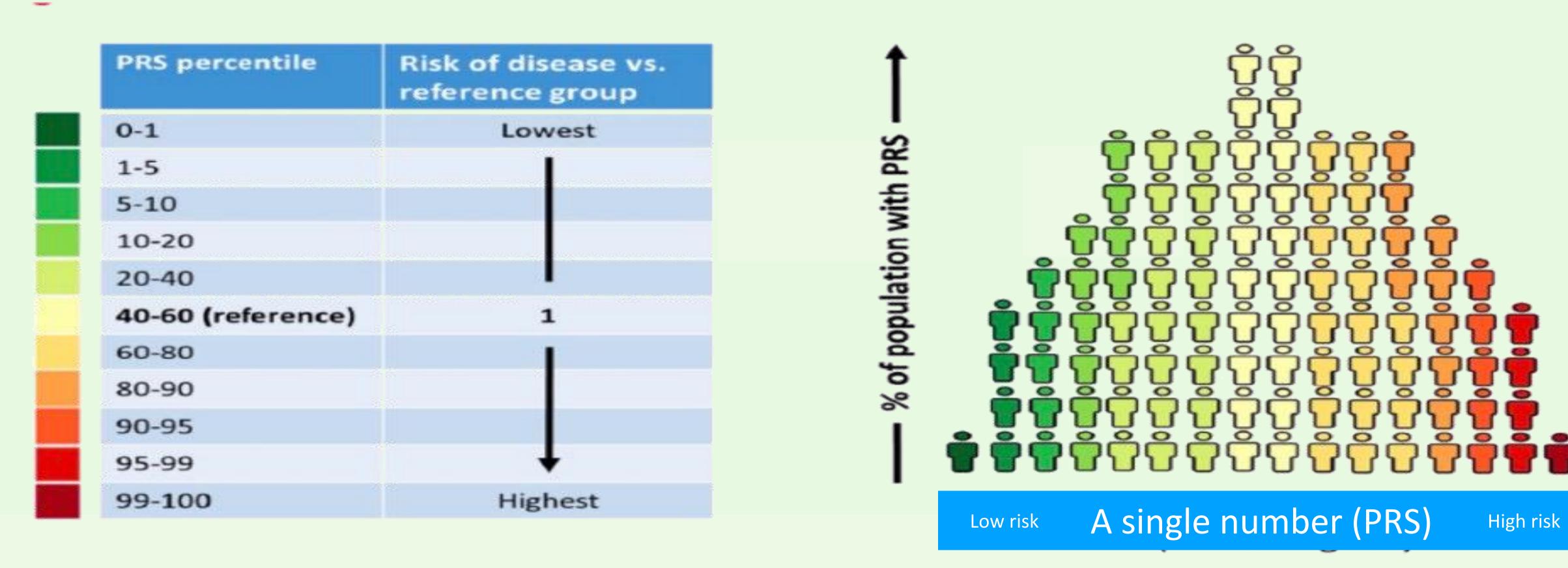
How to use the GWAS - data?





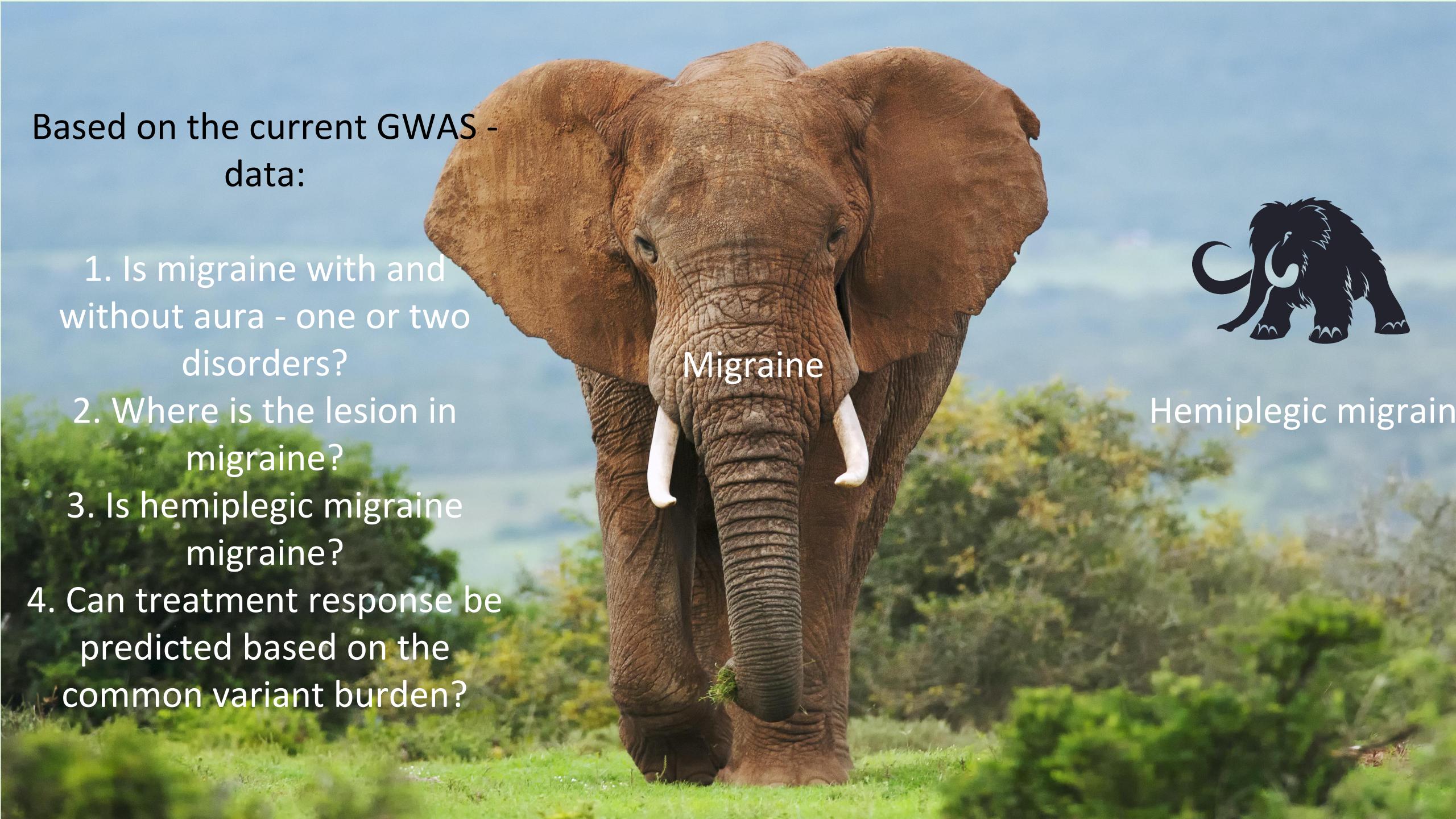
Hautakangas H et al. Genome-wide analysis of 102,084 migraine cases identifies 123 risk loci and subtype-specific risk alleles. medRxiv 2021.01.20.21249647; doi: https://doi.org/10.1101/2021.01.20.21249647

Common variants contribution to a disease in a single number = polygenic risk score = PRS



Source: RGA

Polygenic Risk Scores = Statistical geneticists have developed the 'polygenic risk score' (PRS), identifying hundreds, thousands and even millions of SNPs (variants) that can be included in a single score that measures the individual's genetic predisposition to specific diseases or traits. Source: RGA = Reinsurance Group of America



1. Is migraine with and without aura - one or two disorders?

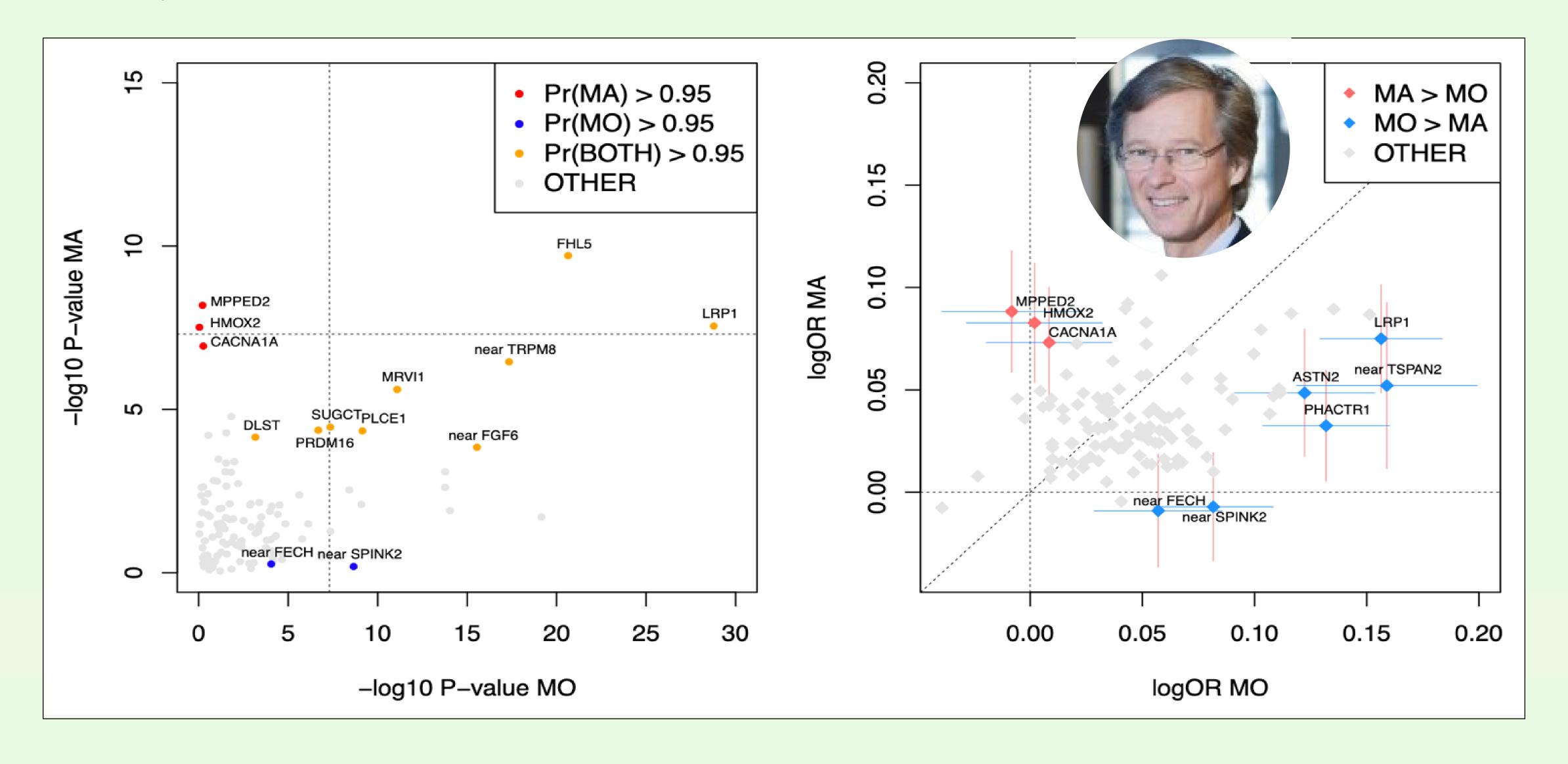
MwA + MwoA MwoA

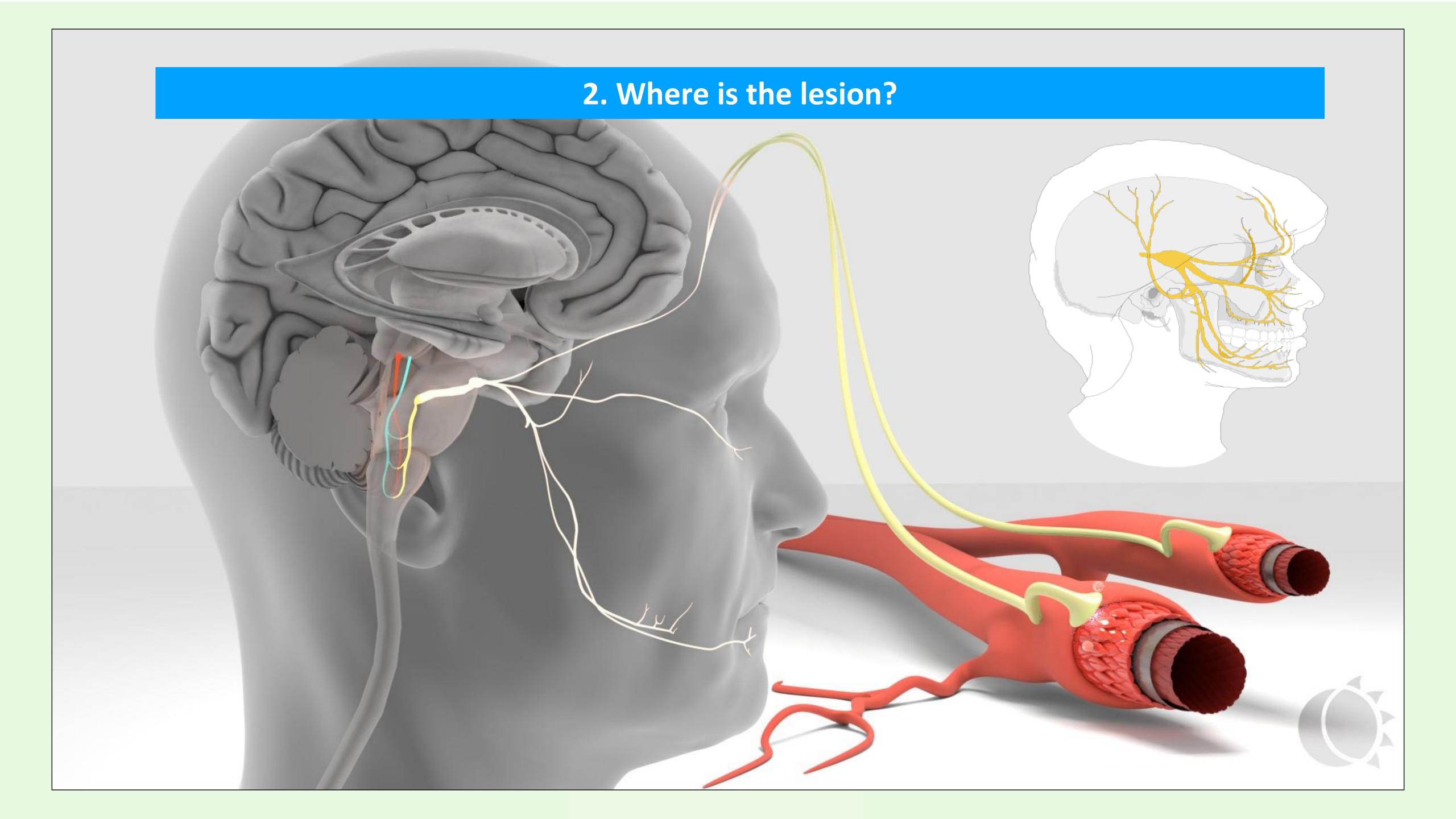


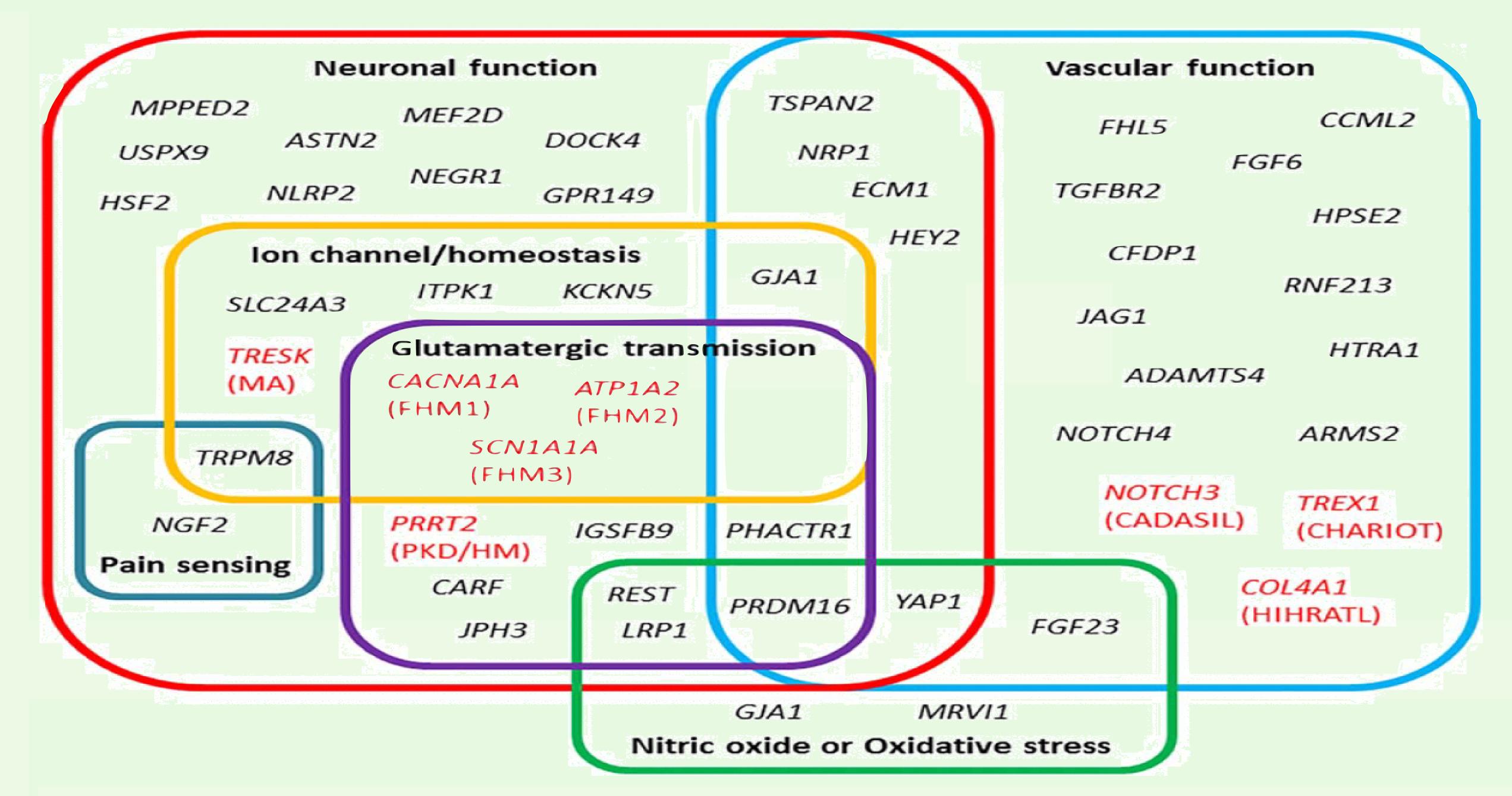


MwA = migraine with aura, MwoA = migraine without aura.

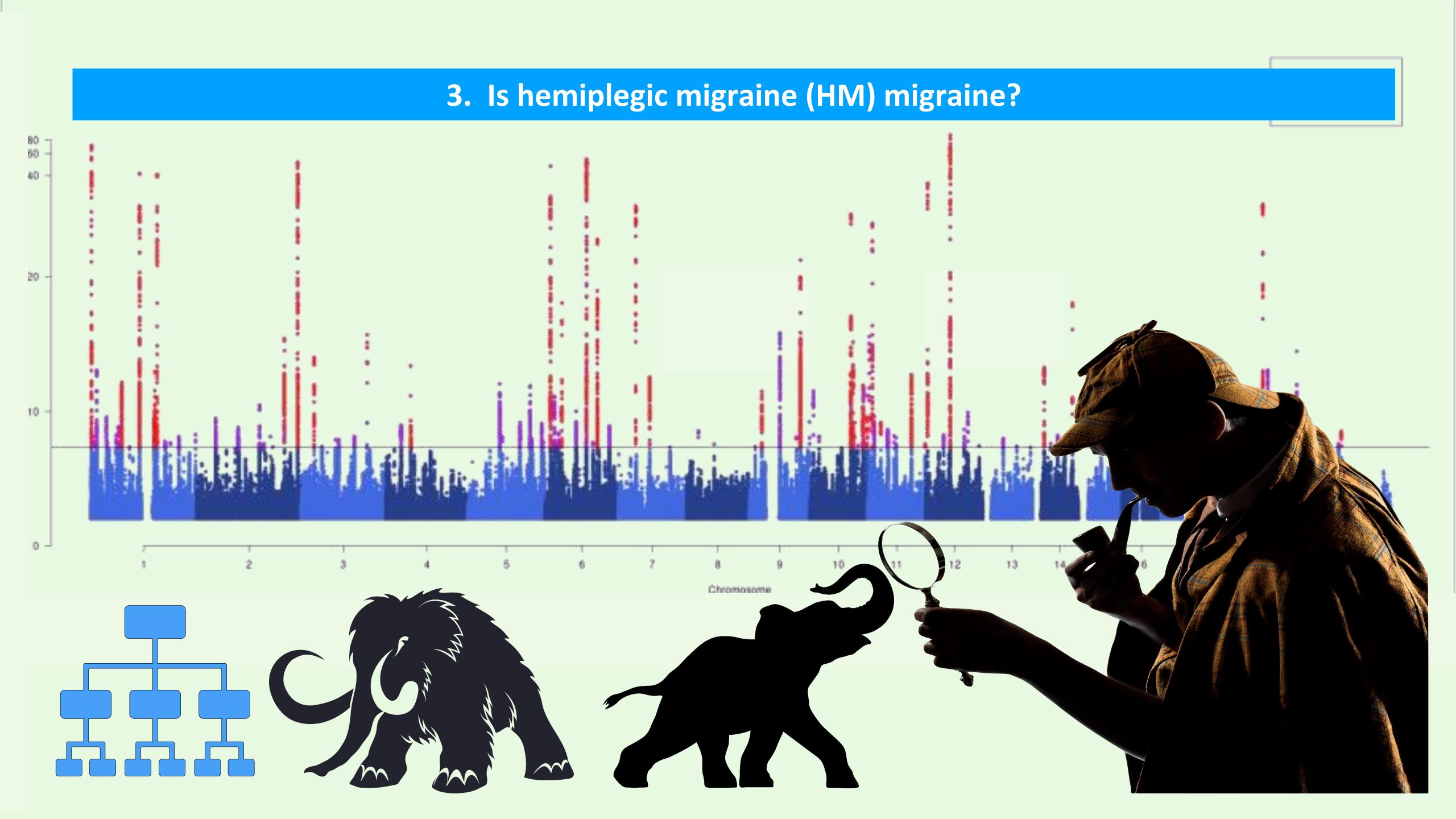
Maybe three - there are aura variants, headache variants, aura and headache variants

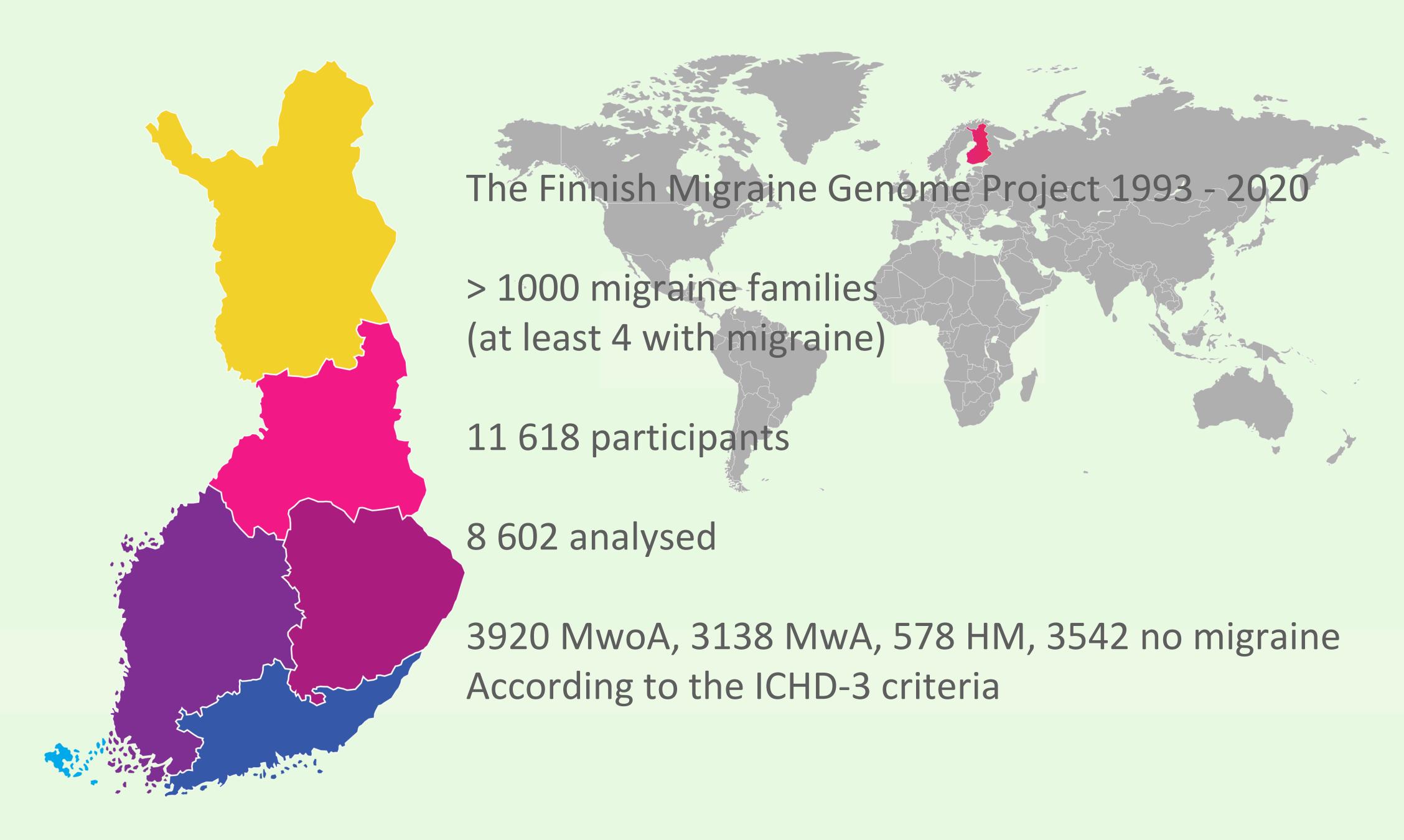




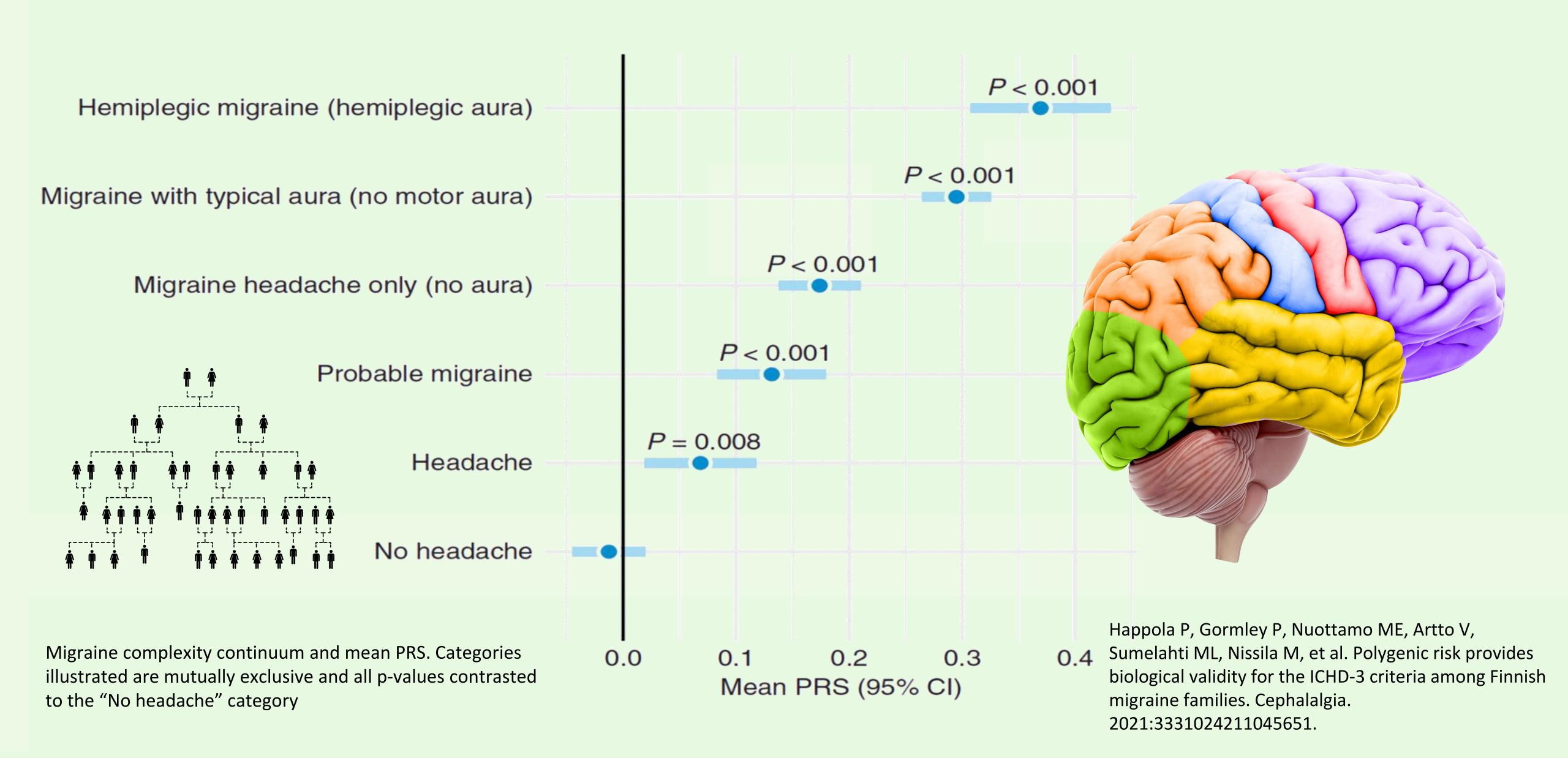


Sutherland HG, Griffiths LR. Genetics of Migraine: Insights into the Molecular Basis of Migraine Disorders. Headache. 2017;57(4):537-69.

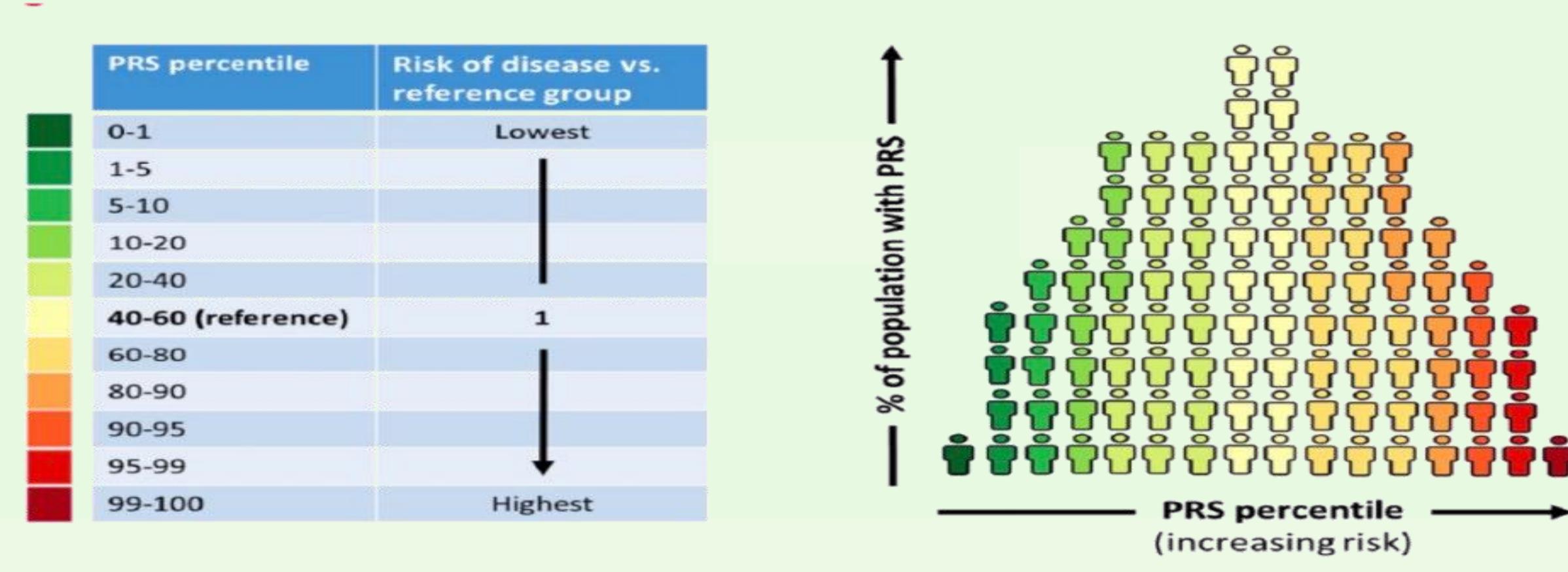




Hemiplegic migraine belongs to the migraine spectrum



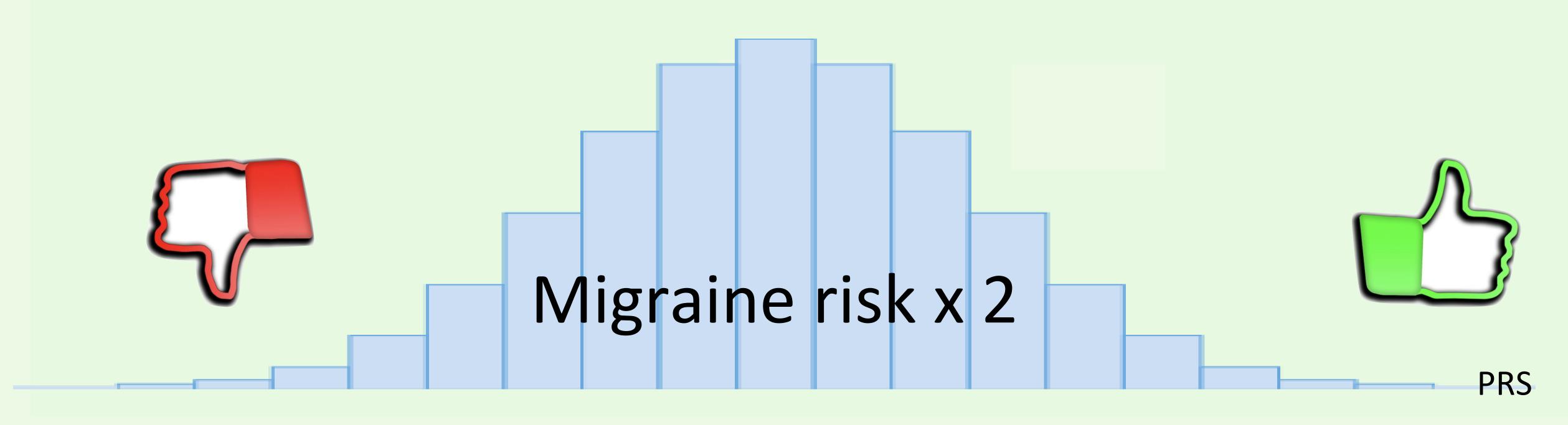
4. Can treatment response predicted based on the common variant burden?



Source: RGA

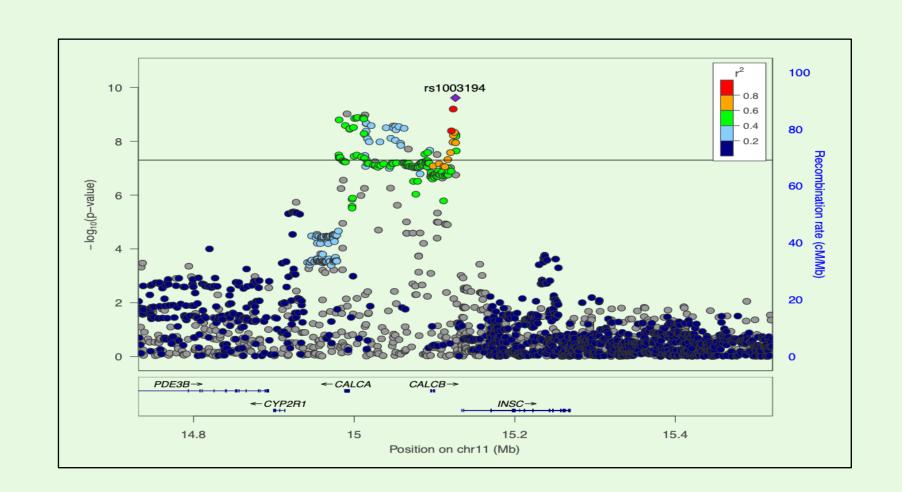
Polygenic Risk Scores = Statistical geneticists have developed the 'polygenic risk score' (PRS), identifying hundreds, thousands and even millions of SNPs (variants) that can be included in a single score that measures the individual's genetic predisposition to specific diseases or traits. Source: RGA = Reinsurance Group of America

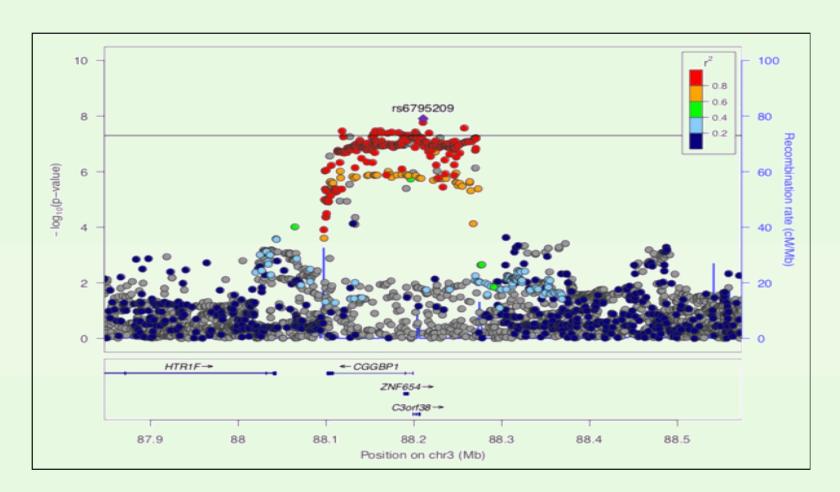
Response to triptans and genetics



A twofold increase in migraine risk associates with positive response to migraine-specific acute treatment (odds ratio [OR] = 1.25 [95% con!dence interval (CI) = 1.05-1.49).

Hot Topic - genetics and migraine medications

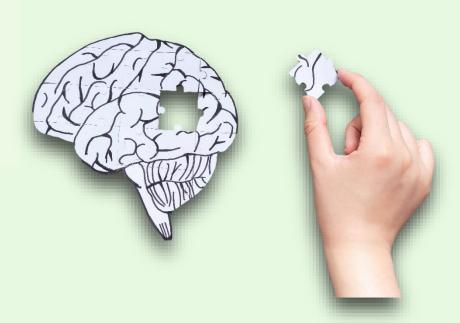




mAbs and gepants = CGRP-antagonists

CALCA CALCB

Chromosome 11



5-HT-1F Ditans (lasmiditan = 5-HT-1F agonist)

Chromosome 3

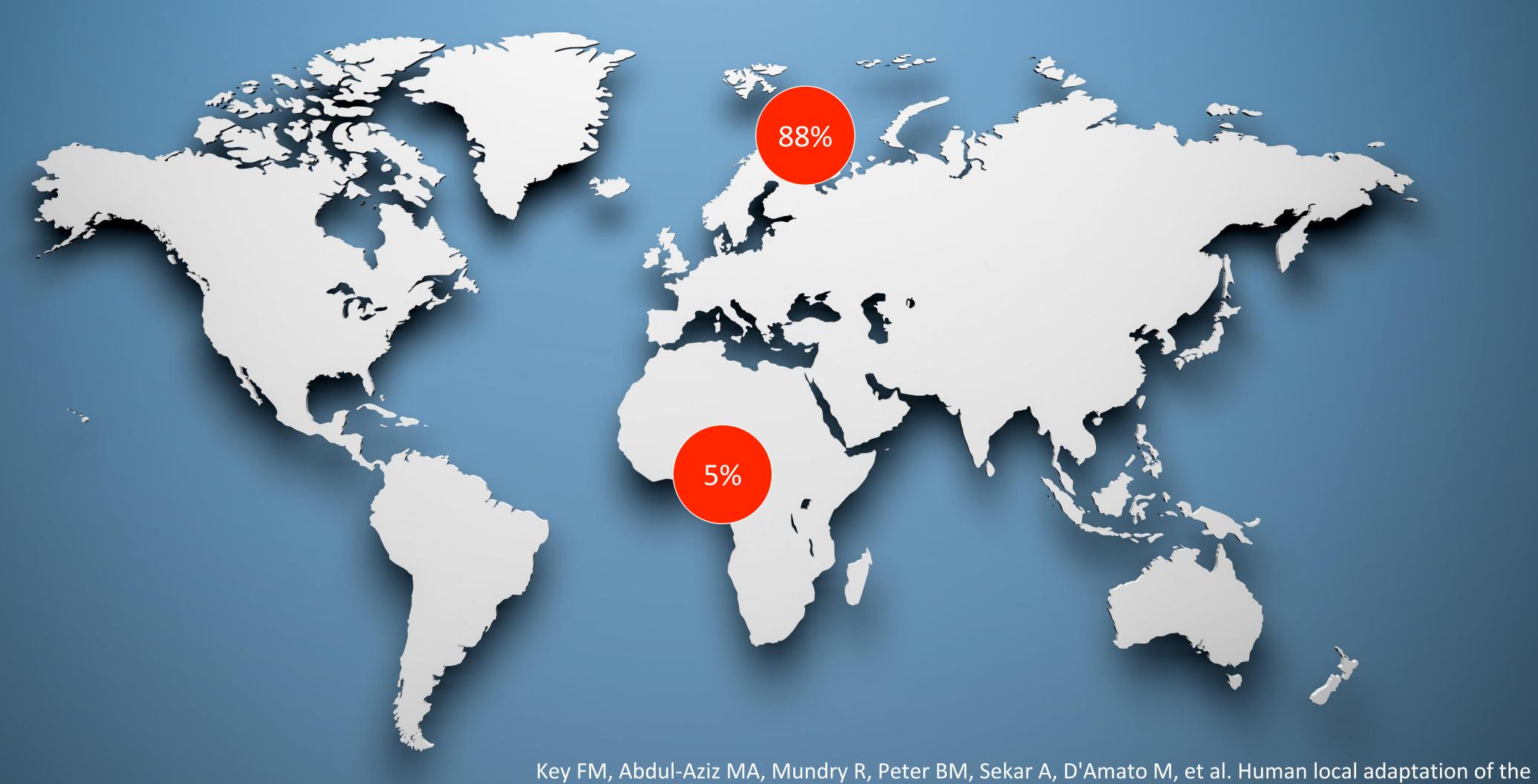
Chan C, Goadsby PJ. Recent Advances in Pharmacotherapy for Episodic Migraine. CNS Drugs. 2019

Hautakangas H et al. Genome-wide analysis of 102,084 migraine cases identifies 123 risk loci and subtype-specific risk alleles. medRxiv 2021.01.20.21249647; doi: https://doi.org/10.1101/2021.01.20.21249647

Some local molecular genetic observations



1. TPRM8 The upstream variant rs10166942 shows extreme population differentiation, with frequencies that range from 5% in Nigeria to 88% in Finland



TRPM8 cold receptor along a latitudinal cline. PLoS Genet. 2018;14(5):e1007298.

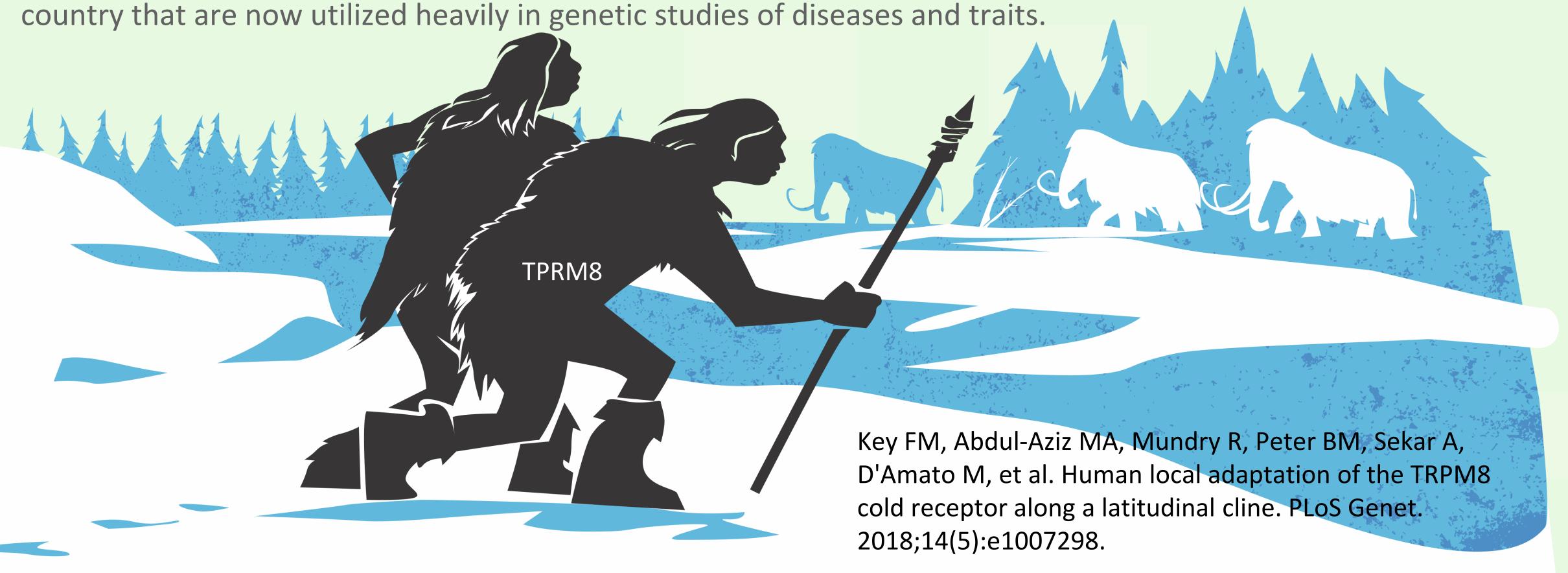
TPRM8 = Transient receptor potential cation channel subfamily M member 8

TPRM8 is a receptor for cold and menthol and also one of the migraine variants Key FM, Abdul-Aziz MA, Mundry R, Peter BM, Sekar A, D'Amato M, et al. Human local adaptation of the TRPM8 cold

receptor along a latitudinal cline. PLoS Genet. 2018;14(5):e1007298.

Two Finns going to the market anno 2021

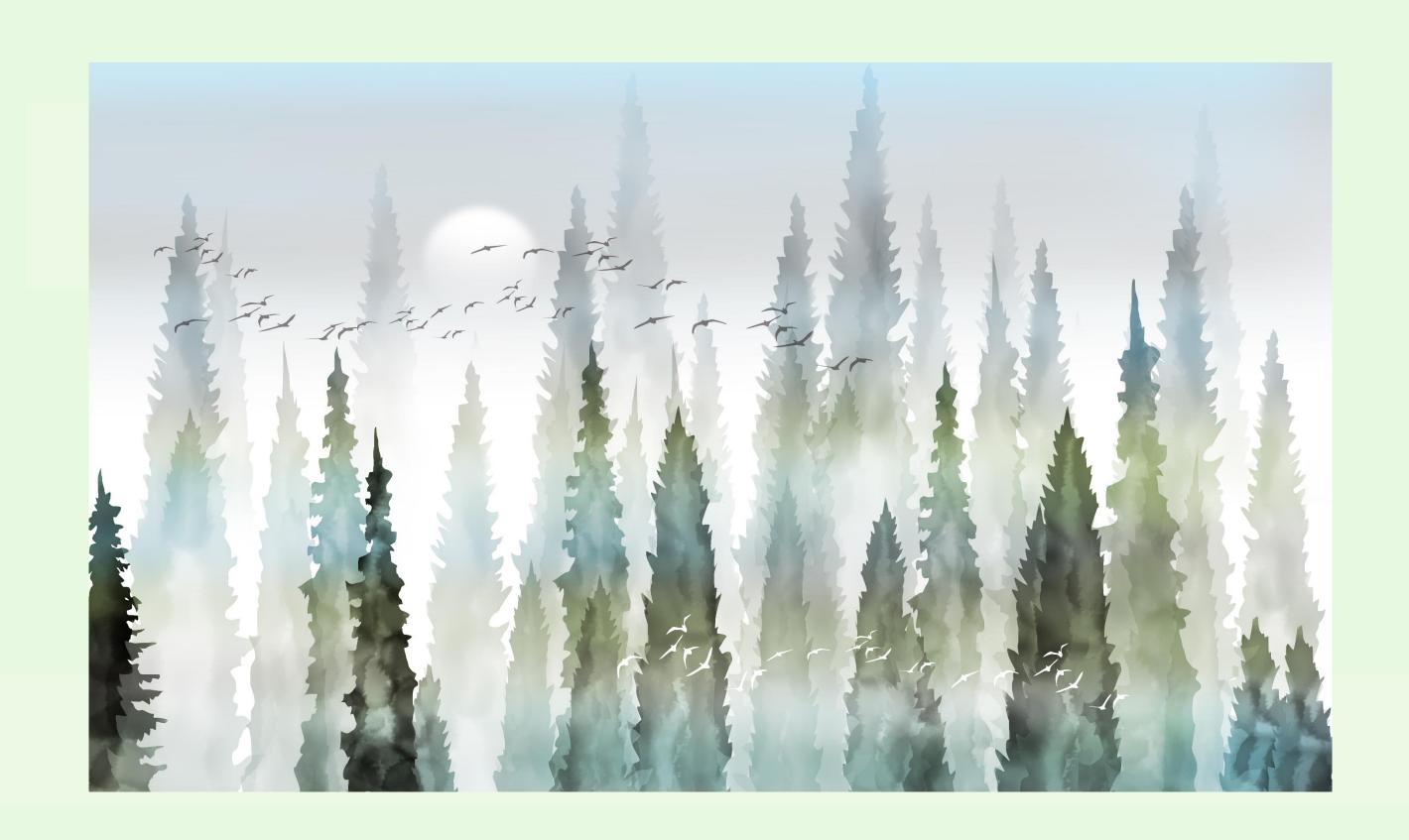
The Finnish population is one of the most genetically studied in the world. A relatively small number of founder individuals and strong genetic isolation over centuries have shaped the unique genetic makeup across the

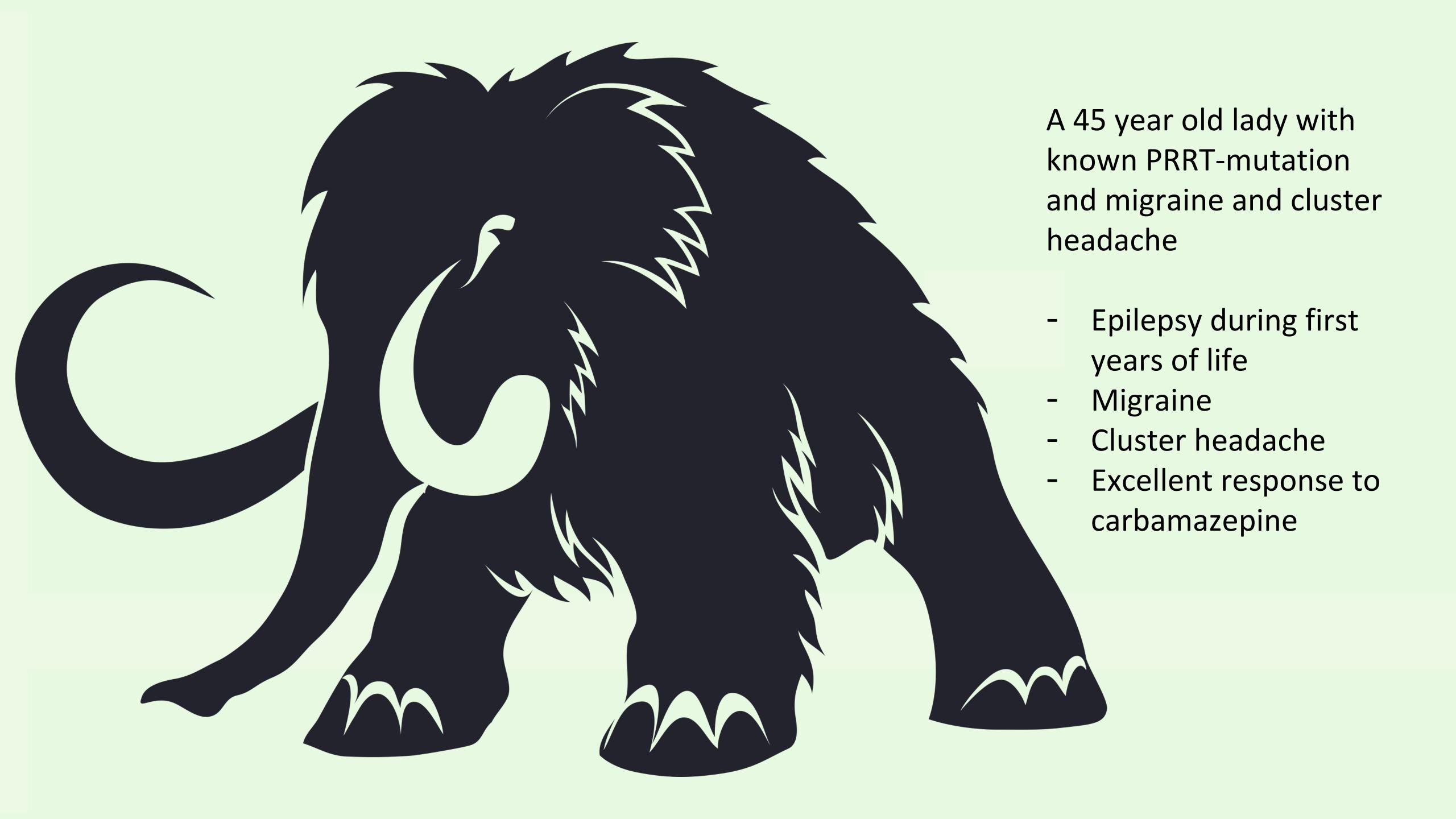


TPRM8 = Transient receptor potential cation channel subfamily M member 8

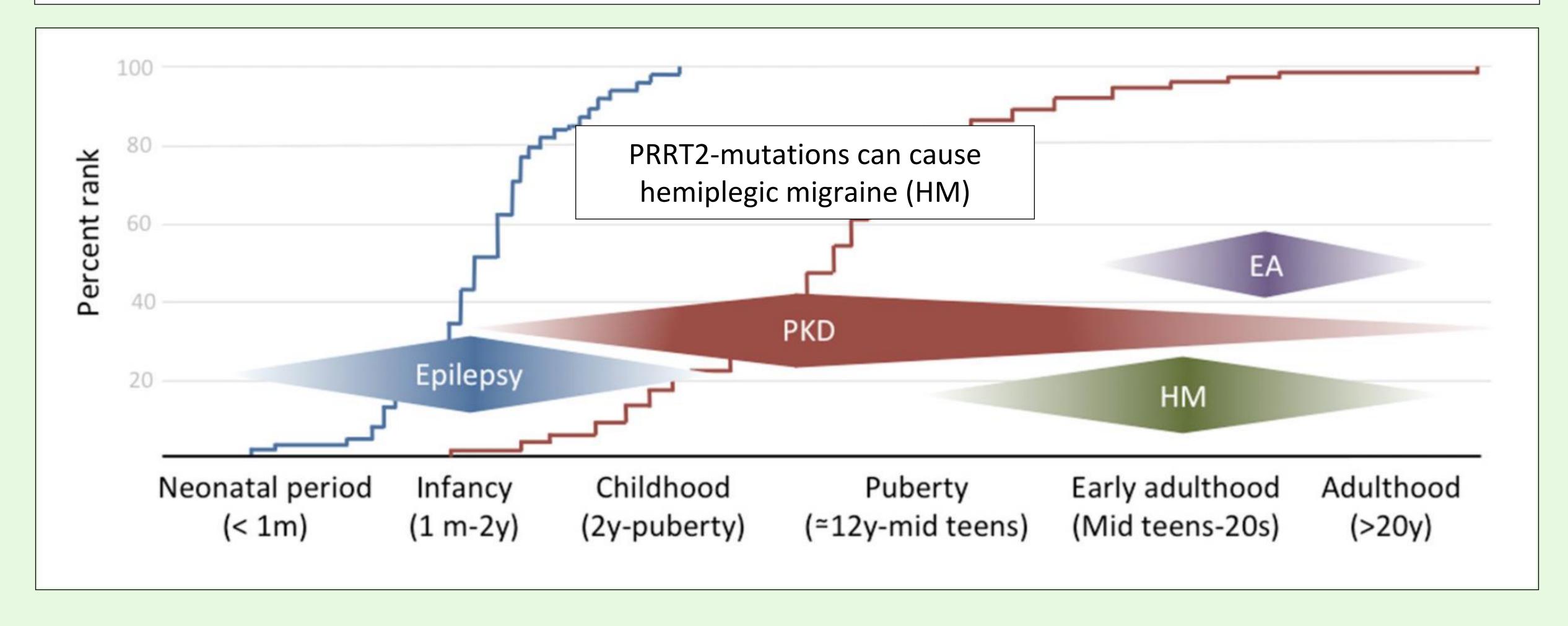
The 2021 Nobel Prize in Physiology or Medicine was awarded to David Julius and Ardem Patapoutian for their discoveries of thermal and mechanical transducers including TRPV1 (receptor for capsaicin) and TPRM8 (receptor for cold and menthol)

https://www.nobelprize.org/prizes/medicine/2021/s ummary/





Hemiplegic migraine with excellent response to Na-channel blocker carbamazepine



Hemiplegic migraine (HM), epilepsy, paroxysmal kinesigenic dyskinesia (PKD), episodic ataxia (EA), excellent response to Na-channel blocker carbamazepine

PRRT2 = proline-rich transmembrane protein 2 (PRRT2)

Landolfi A, Barone P, Erro R. The Spectrum of PRRT2-Associated Disorders: Update on Clinical Features and Pathophysiology. Front Neurol. 2021;12:629747.

What does future look like in migraine genetics?



Aarno Palotie - to make individual decisions it takes larger data sets Jogi Berra - future is not what its used to be

Migraine - multiple mechanisms - multiple chances for progress



Detailed clinical and genetic phenotyping - better migraine spesific tretments



Clinical problem

Randomised placebo controlled trial (RCT)

Treatment versus placebo



Clinical diagnostic criteria

Treatment superior to placebo in Phase 3 study



Patients with hemiplegic and brainstem aura excluded

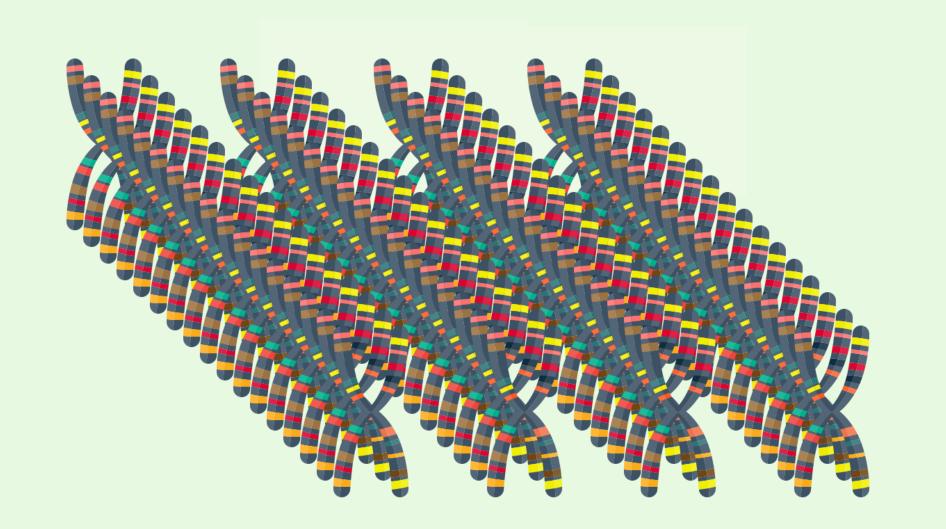
Clinical problem

Randomised placebo controlled trial (RCT) including PRS - data

Clinical trial - data and PRS) of the patient



Clinical diagnostic criteria

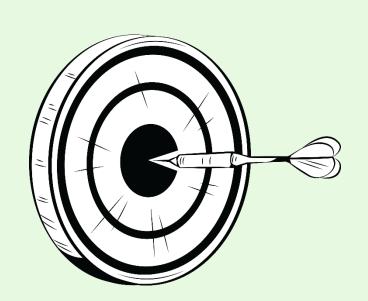


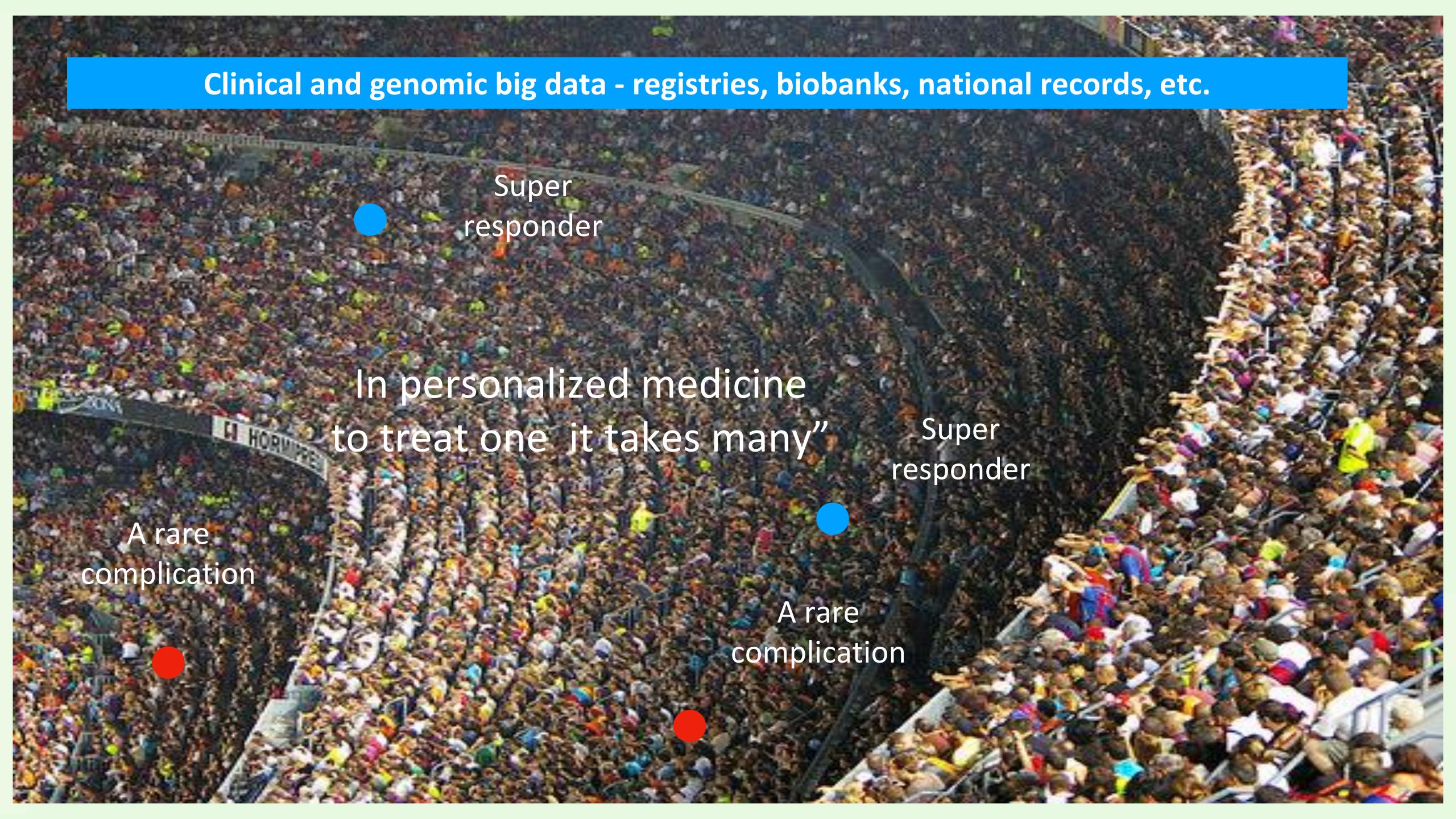
Treatment based on clinical trials and the PRS of the patient



PRS based on >> 100 000 genotypes







Migraine is common and rare

