



McGill
Le Centre Alan-Edwards
de recherche sur la
DOULEUR



McGill
The Alan Edwards
Centre for Research on
PAIN



Canada Excellence
Research Chairs
Chaire d'excellence
en recherche du Canada

Novel Methods to Identify Pain Targets : Genomic/Genetic Approaches

Luda Diatchenko, MD, PhD

Canada Excellence Research Chair in Human Pain Genetics

Alan Edwards Centre for Research on Pain,

McGill University, Montreal, Canada

The National Academies of SCIENCE ENGINEERING MEDICINE

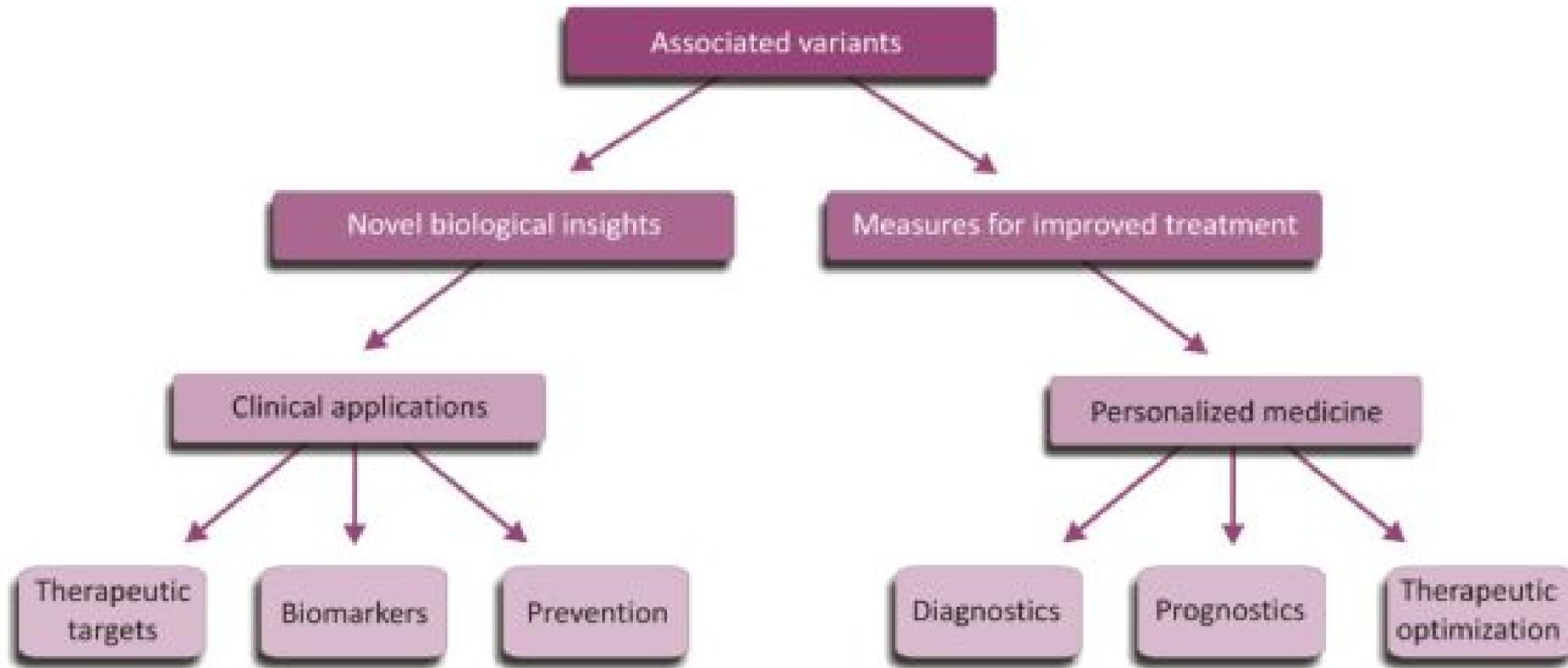
Advancing Therapeutic Development for Pain and Opioid Use Disorders

through Public-Private Partnerships: A Workshop

Washington, DC, October 11–12, 2017



Translation of Genetic Information into Clinical Practice



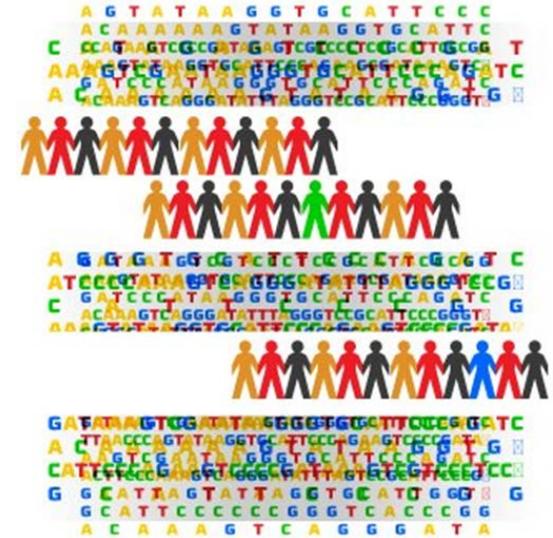
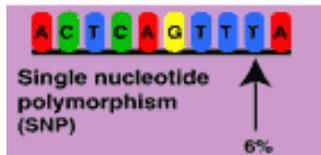
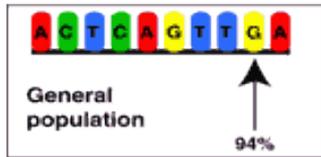
How are genomes of individuals different?

Polymorphism

"Poly" *many* "morph" *form*

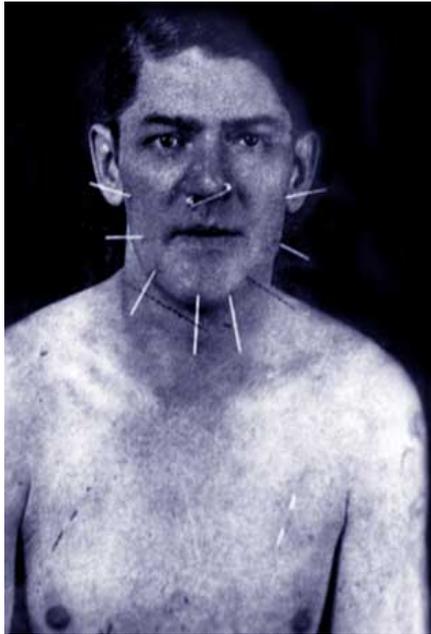


SNP



DNA contains 99.9% of identical sequence in all the individuals with only 0.1% difference. Out of this 0.1% variation, over 80% are single nucleotide polymorphisms (SNPs).

Genes Responsible for Monogenic Pain Disorders



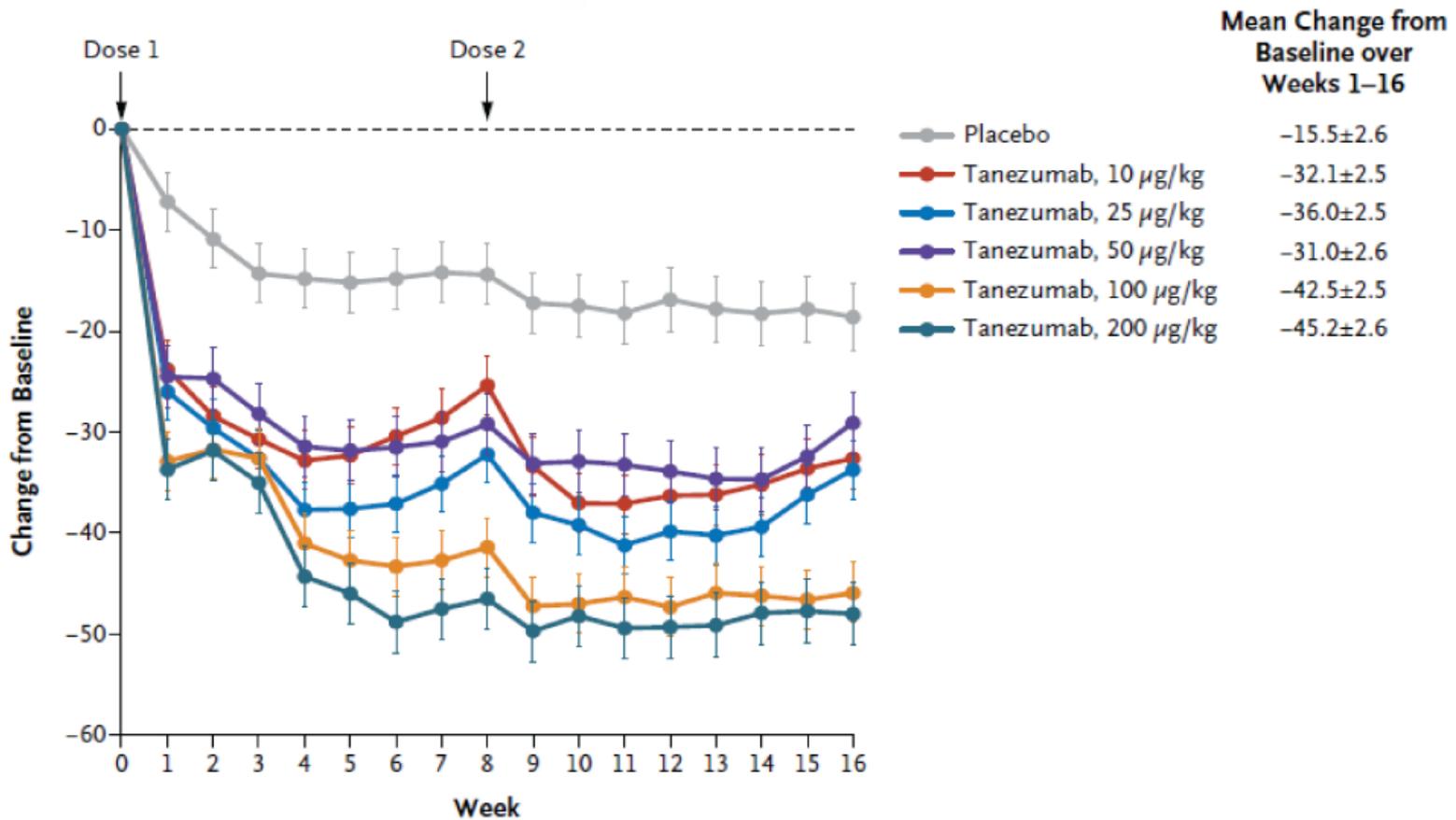
HSN Type I	<i>SPTLC1</i>	sphingolipid synthesis
HSN Type II	<i>HSN2</i>	(function unknown)
HSN Type IID	<i>SCN9A</i>	sodium (Na _v 1.7) channel
HSN Type III	<i>IKBKAP</i>	transcription factor
HSN Type IV	<i>NTRK1</i>	neurotrophin receptor
HSN Type V	<i>NGFB</i>	neurotrophin
HSN Type ?	<i>SCN11A</i>	sodium (Nav1.9) channel
FEPS	<i>TRPA1</i>	cation (TRPA1) channel
PE	<i>SCN9A</i>	sodium (Na _v 1.7) channel
PEPD	<i>SCN9A</i>	sodium (Na _v 1.7) channel
FHM Type I	<i>CACNA1A1</i>	calcium channel subunit
FHM Type II	<i>ATP1A2</i>	ion pump subunit
FHM Type III	<i>SCN1A</i>	sodium (Na _v 1.1) channel

“The Human Pincushion”
(congenital insensitivity
to pain with anhidrosis;
HSN Type IV)

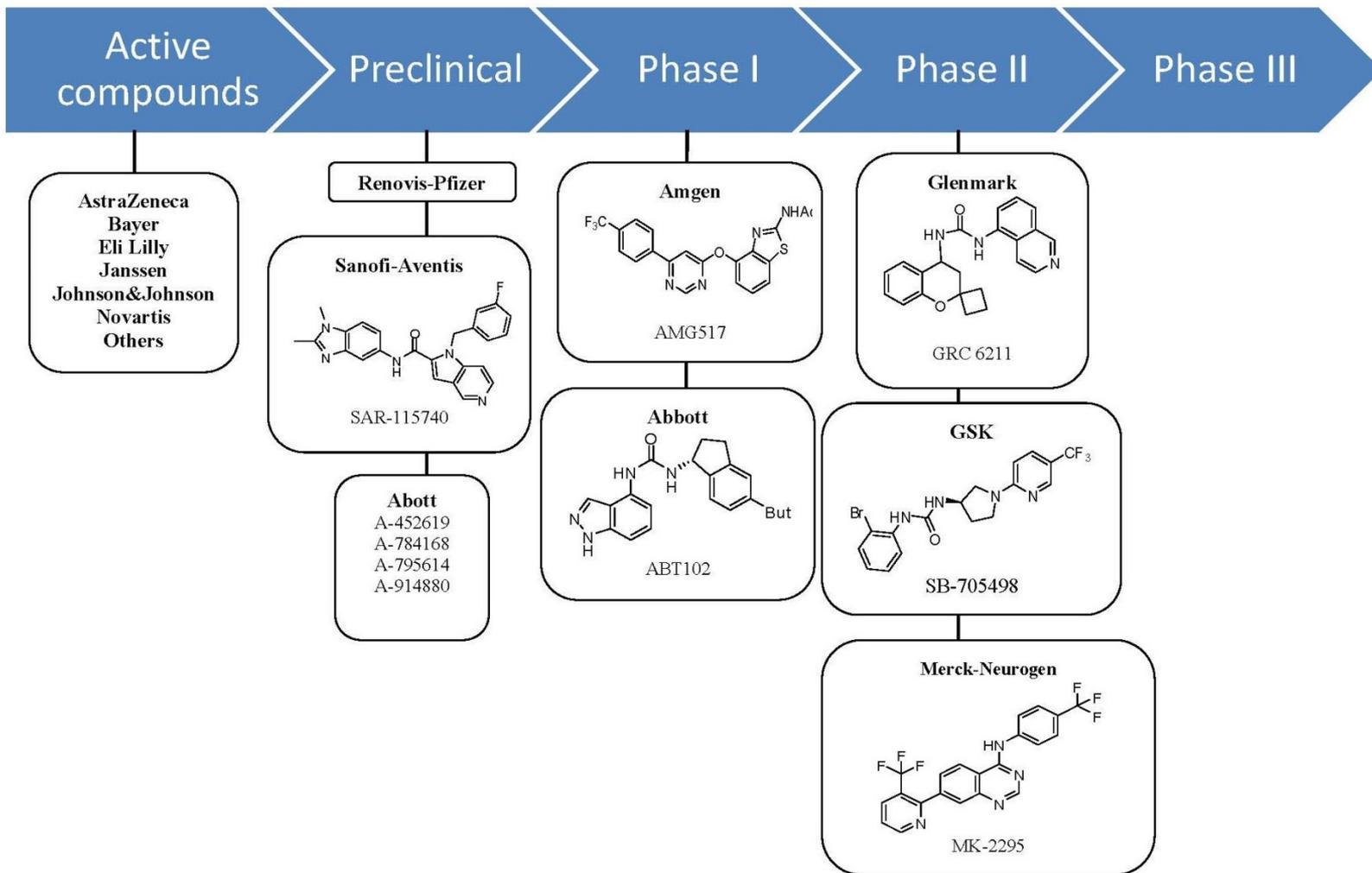
FEPS: familial episodic pain syndrome; HSN: hereditary sensory neuropathy;
PE: primary erythromelalgia; PEPD: paroxysmal extreme pain disorder; FHM: familial hereditary migraine

Drug Development Based on Monogenic Pan Disorders -

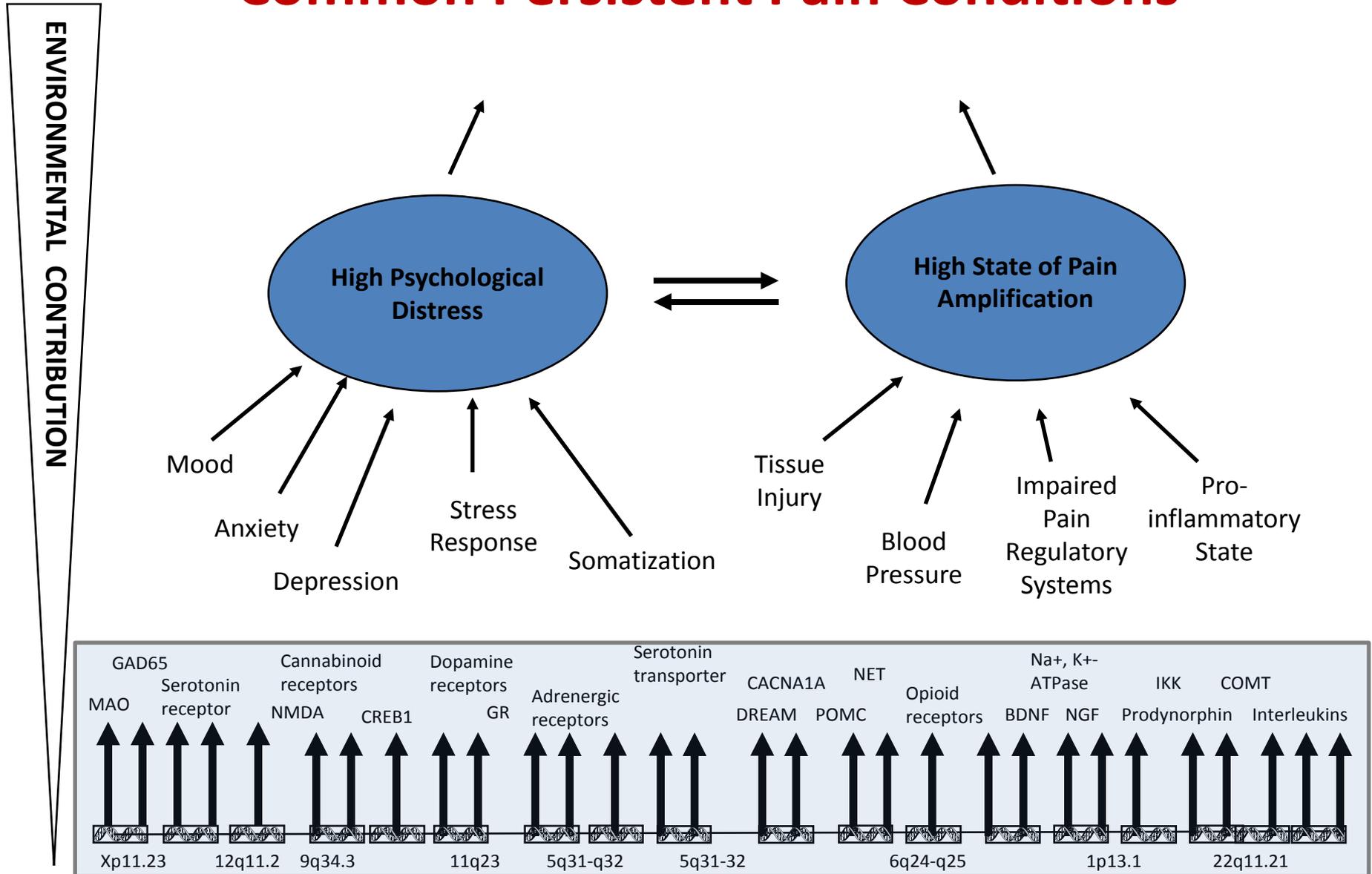
Anti-NGF attenuates knee pain while walking in osteoarthritis patients



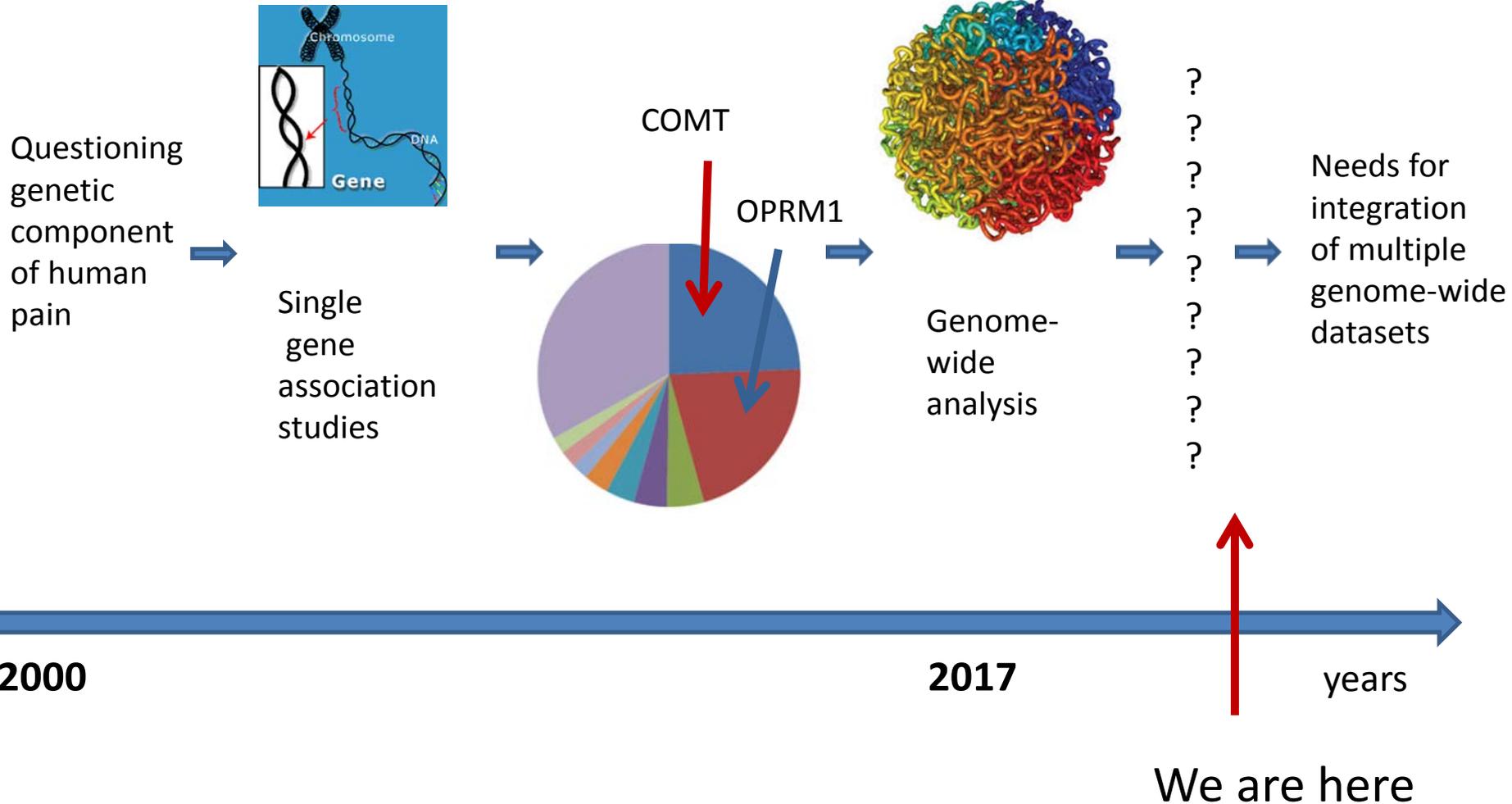
Drug Development Based on Monogenic Pan Disorders – Sodium Channel blockers



Common Persistent Pain Conditions



Development of Pain Genetics Field





To create a database that aggregates relevant and up-to-date human pain genetics data and complementary resources in one centralized location to be used as a resource for clinicians and pain researchers



Human Pain Genetics Lab Human Pain Genes Database

Home Charts Downloads Contact Us About the HPGDB

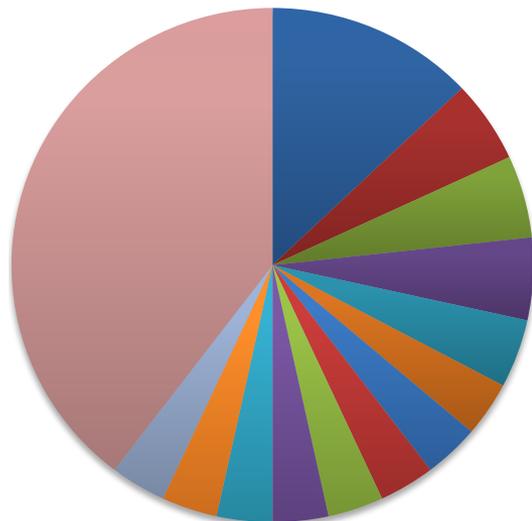
Search:

Loci	Variants	Alleles	Direction	Phenotype	Publication
ABCB1	rs1045642	C	↓	Analgesia	Campa, 2008
ABCB1	rs1045642	T/T	↓	Analgesia	Candiotti, 2013
ABCB1	rs1045642	T	↓	Analgesia	Zwisler, 2010
ABCB1	rs1045642	T/T	↑	Post-operative Pain	Sia, 2010
ABCB1	rs1045642	T/T	↑	Analgesia	Rhodin, 2013
ABCB1	rs1045642	C/C	↑	Post-operative Pain	Mamie, 2013
ABCB1	rs1045642	C	↑	Post-operative Pain	Mamie, 2013
ABCB1	rs1045642	C	↓	Analgesia	Hajj, 2015
ABCB1	rs1045642	C/C	↑	Cancer Pain	Wang, 2015

Genetic Loci Associated With

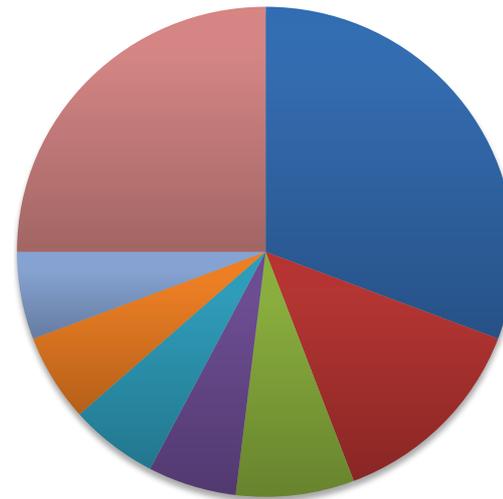
Quantified By The Number Of Genetic Association Studies

Migraine



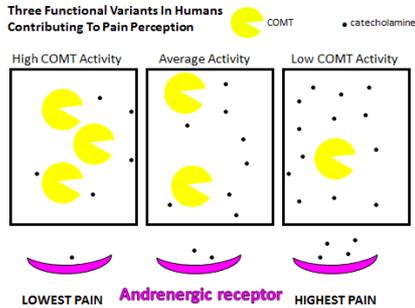
- MTHFR
- ACE
- PRDM16
- TNF
- ESR1
- AJAP1
- C7orf10
- DBH
- FHL5
- LRP1
- LTA
- MMP16
- TRPM8
- Other

Musculoskeletal Pain Disorders



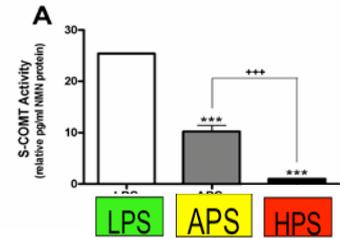
- COMT
- HTR2A
- ESR1
- ADRB2
- IL1A

The Translational Research Clock – closing the circle



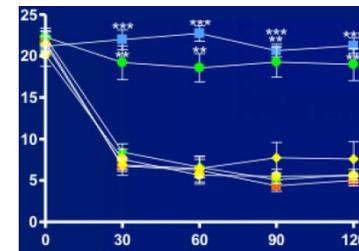
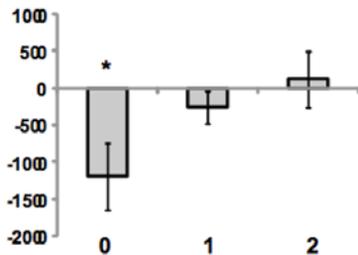
Association study
results, *HMG*,
2005
(942 citations)

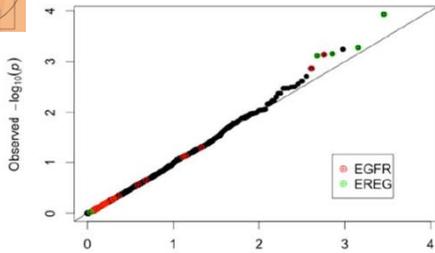
Molecular genetics
of functional
variants, *Science*,
2006
(607 citations)



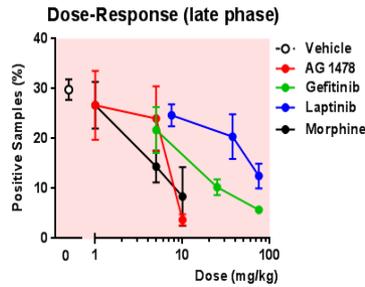
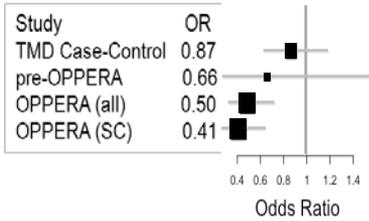
Clinical trial –
*Pharmacogenet
Genomics*, 2010
(83 citations)

The animal
behavior
study – target
identification, *Pain*
2007
(198 citations)

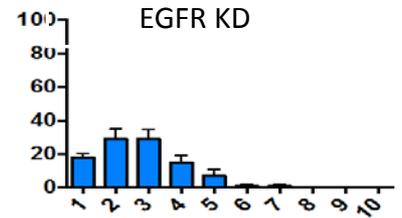
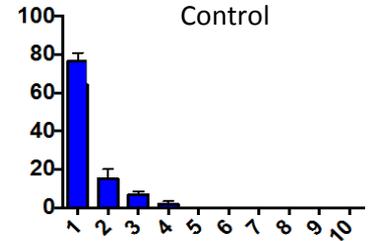
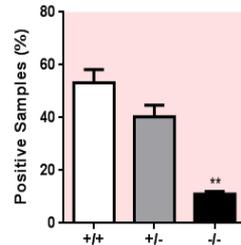




Chronic Pain Risk



Wa2 Mutant - Formalin Test (late)



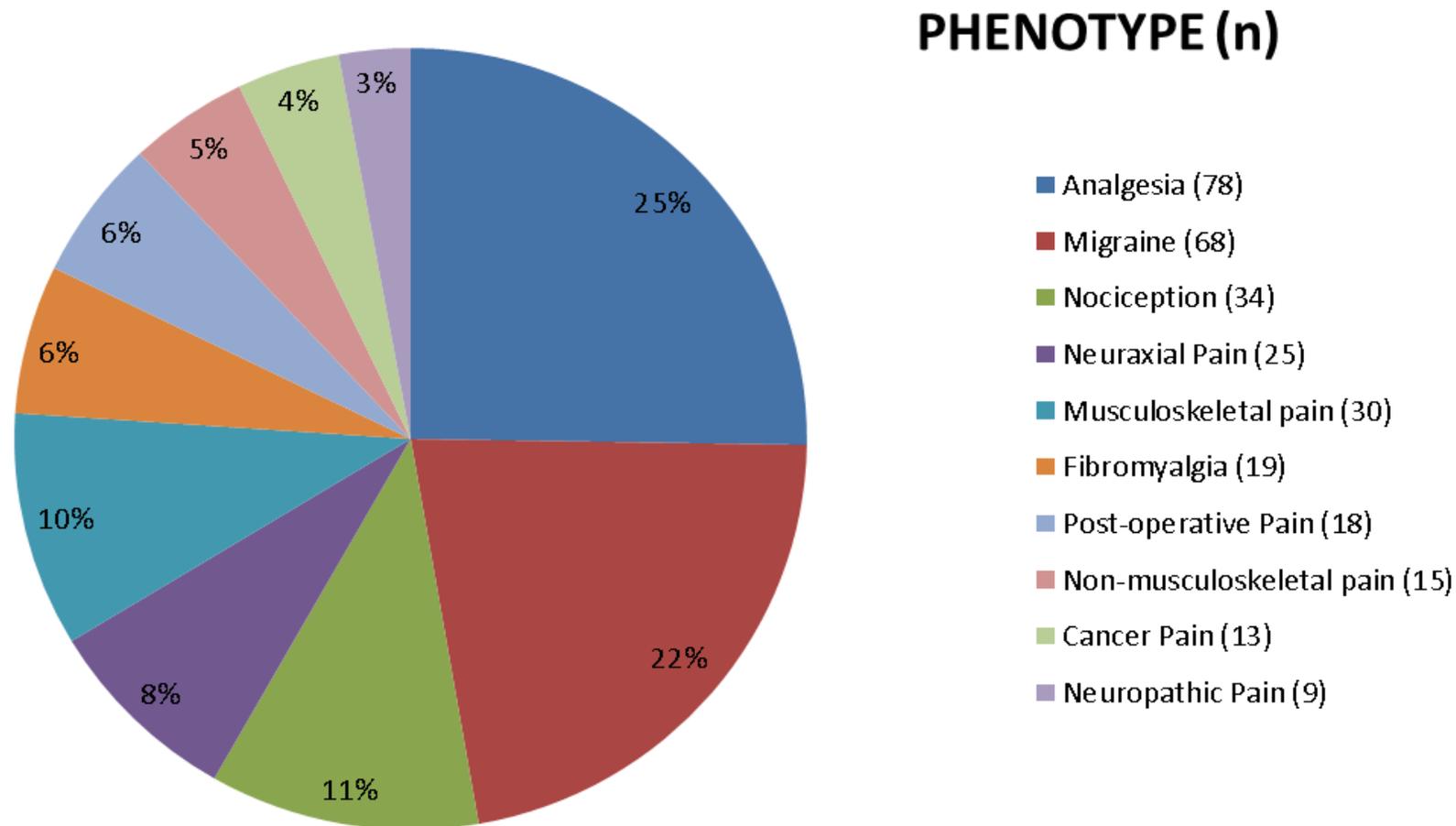
The Journal of Clinical Investigation

RESEARCH ARTICLE

Epiregulin and EGFR interactions are involved in pain processing

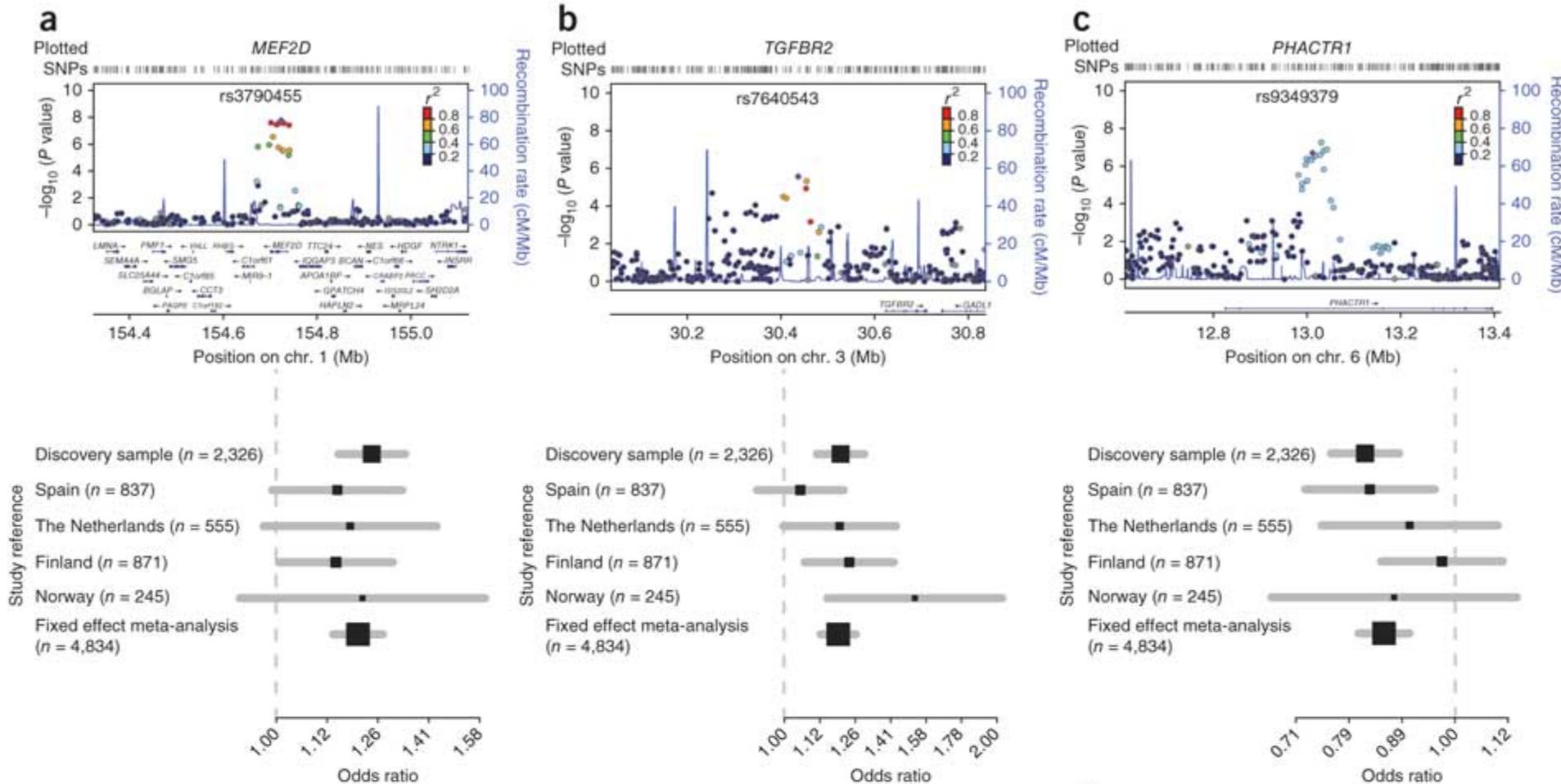
Loren J. Martin,^{1,2} Shad B. Smith,³ Arkady Khoutorsky,⁴ Claire A. Magnussen,⁵ Alexander Samoshkin,⁶ Robert E. Sorge,¹ Chulmin Cho,² Noosha Yosefpour,⁵ Sivaani Sivaselvachandran,² Sarasa Tohyama,² Tiffany Cole,⁷ Thang M. Khuong,⁷ Ellen Mir,³ Dustin G. Gibson,³ Jeffrey S. Wieskopf,¹ Susana G. Sotocinal,¹ Jean Sebastien Austin,¹ Carolina B. Meloto,⁶ Joseph H. Gitt,³ Christos Gkogkas,⁴ Nahum Sonenberg,⁴ Joel D. Greenspan,⁸ Roger B. Fillingim,⁹ Richard Ohrbach,¹⁰ Gary D. Slade,¹¹ Charles Knott,¹² Ronald Dubner,⁸ Andrea G. Nackley,³ Alfredo Ribeiro-da-Silva,⁵ G. Gregory Neely,⁷ William Maixner,³ Dmitri V. Zaykin,¹³ Jeffrey S. Mogil,¹ and Luda Diatchenko⁶

Phenotypes categories included in the HPGdb



Multiple Roads to Migraine

Freilinger et al *Nature Genetics* 2012

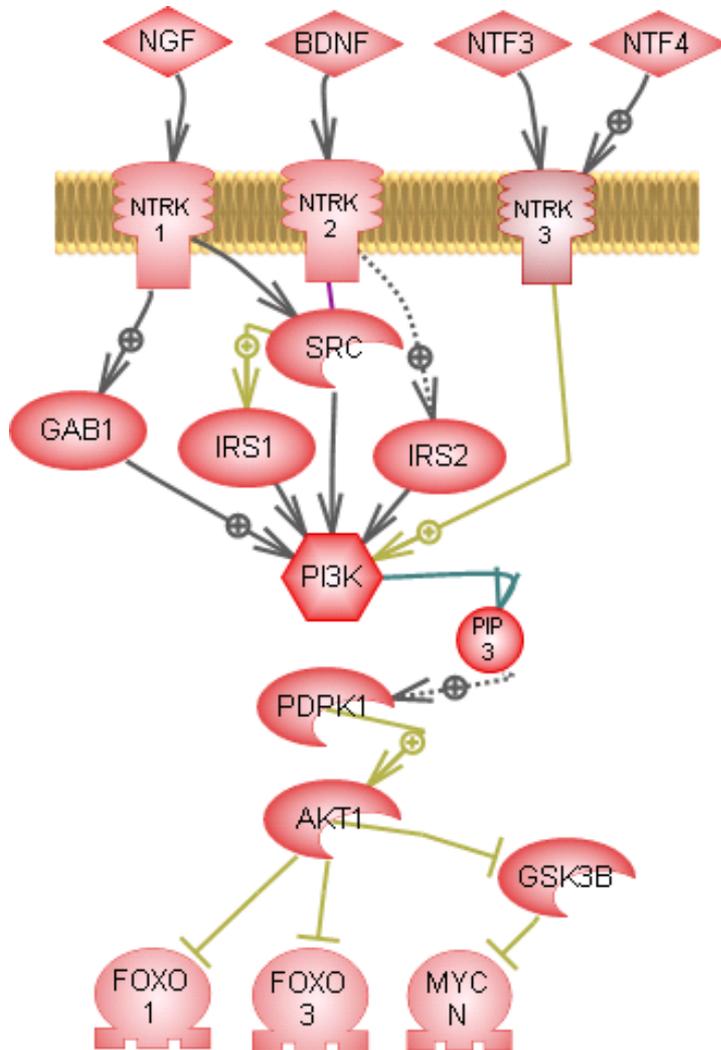


d

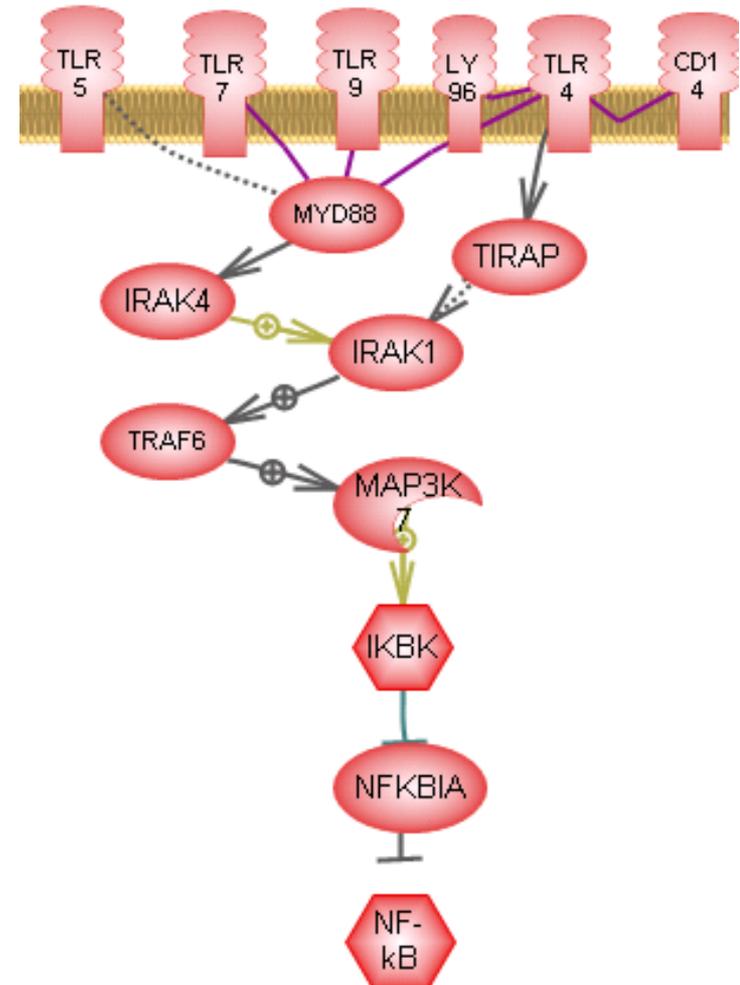
Plotted SNPs

ASTN2

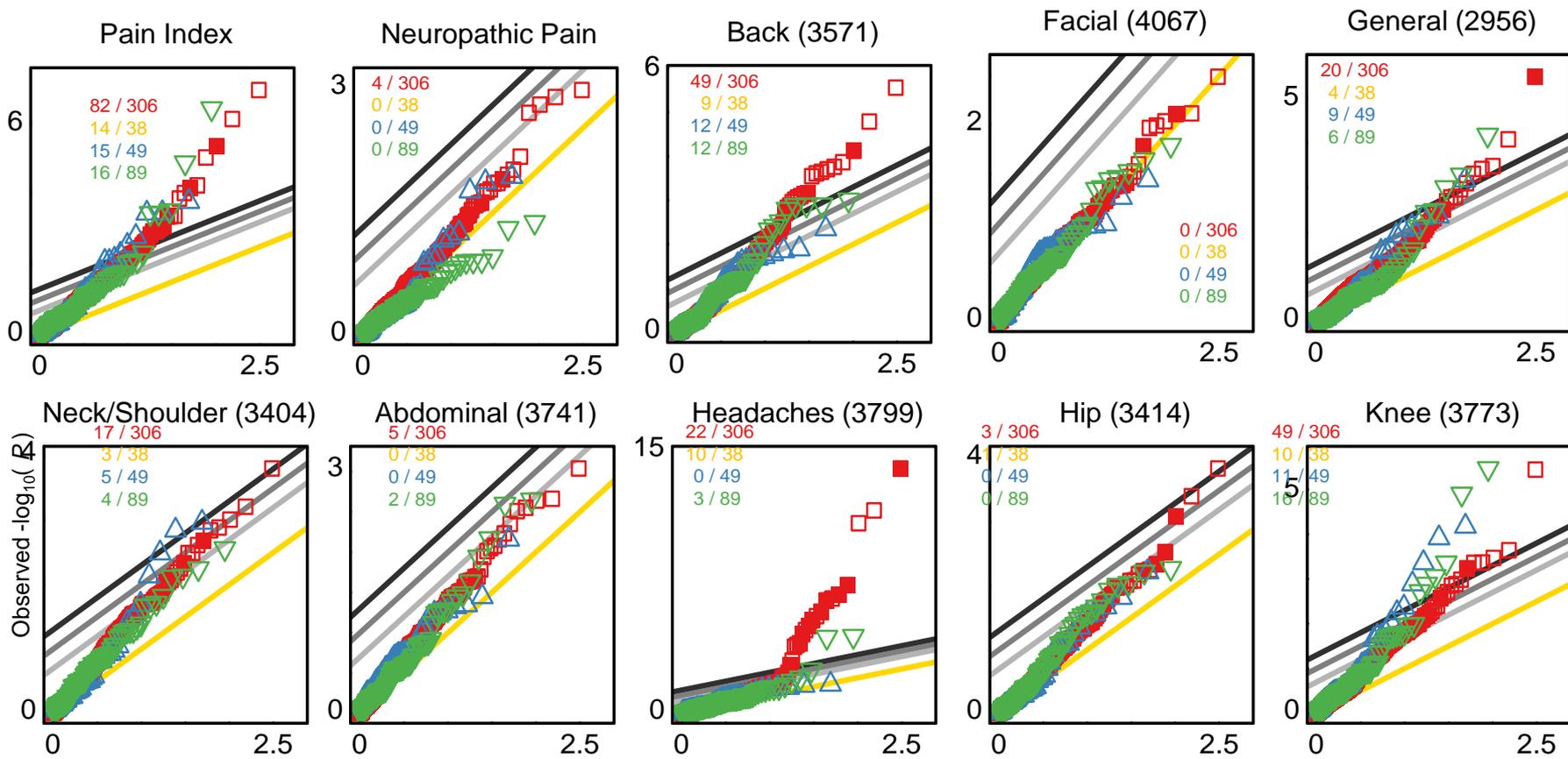
Heterogeneity of patient population



patients subgroup 1



patients subgroup 2



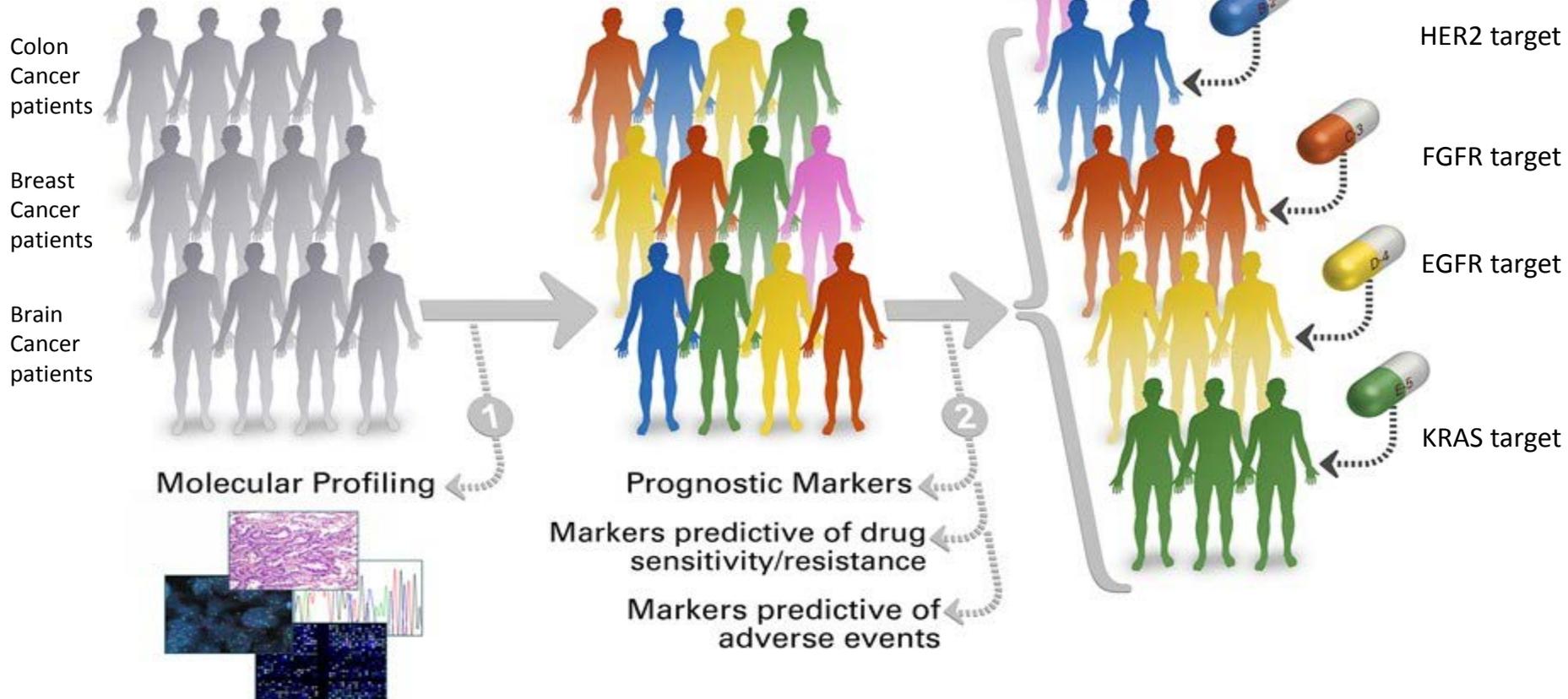
□ Clinical Pain ■ GWAS
△ Nociception
▽ Analgesia

FDR — 5%
 — 10%
 — 20%
 — null

QQ plots for SNPs replication in the UK BioBank. Ratios are number of SNPs better than 20% FDR to the total number of SNPs in respective SNP groups (SNPs from GWAS in orange).

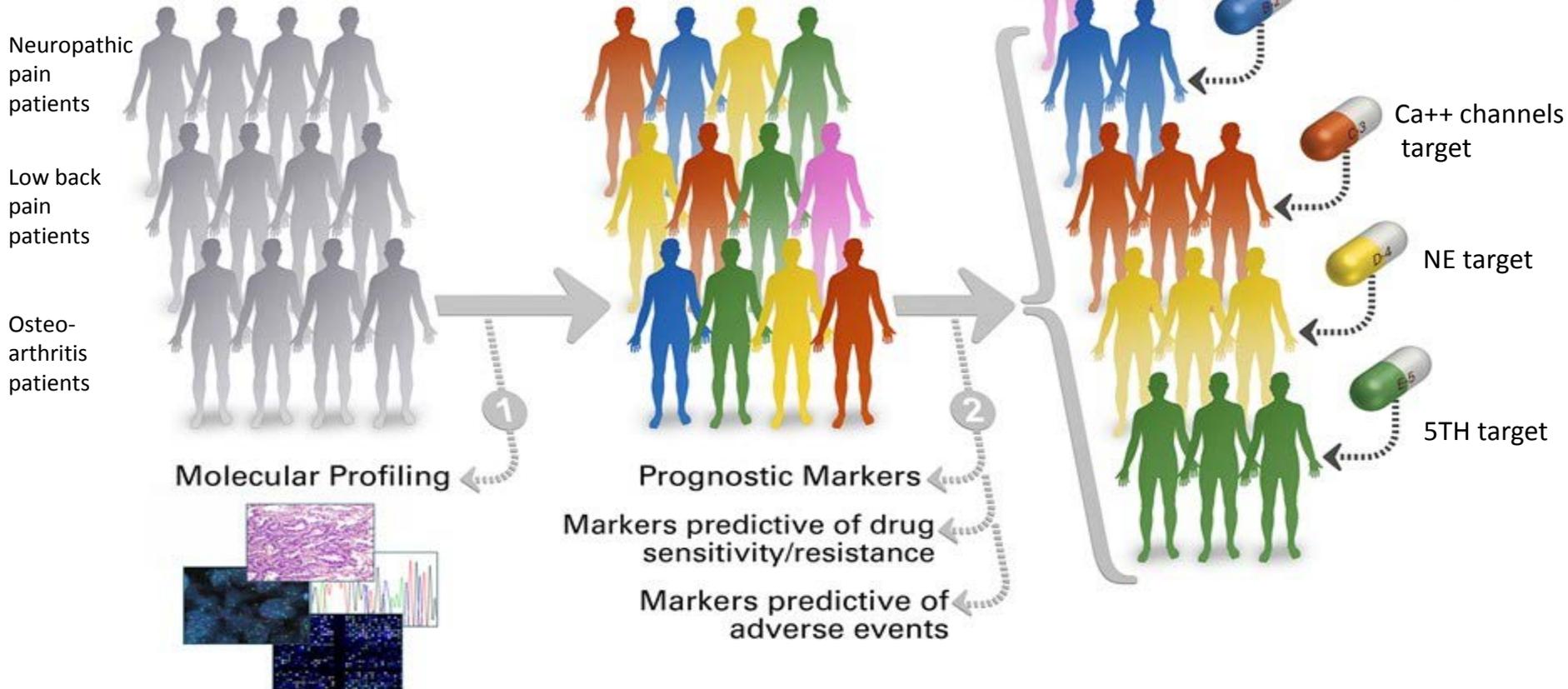
Non-organ specific approach to treatment cancer

Personalized Cancer Therapy

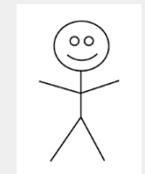
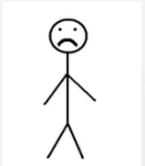
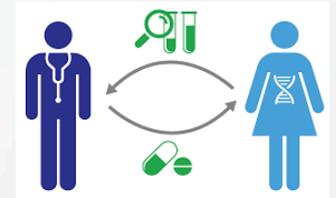
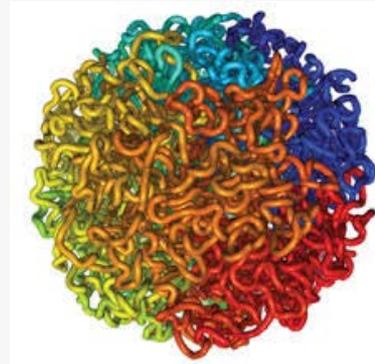
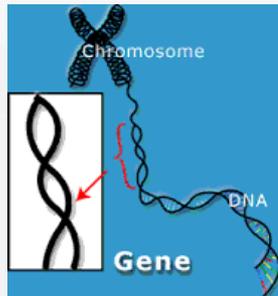


Non-organ specific approach to treatment pain

Personalized pain treatment



SUMMARY OF STEPS IN GLOBAL GENETIC PAIN TARGETS DEVELOPMENT PROGRAM



Welcome to the Human Pain Genetics Lab



Thank You

