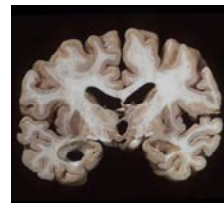
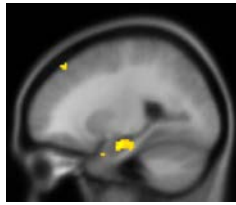


Individualisierte Medizin: Hoffnung für die Zukunft oder eher eine Illusion?



Prof. Andreas Papassotiropoulos, M.D.
Division of Molecular Neuroscience
Faculty of Psychology and University Psychiatric Clinics
Life Sciences Training Facility, Biozentrum
University of Basel, Switzerland

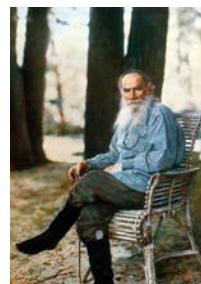
Rheinfelder Tage, Rheinfelden, 16. März 2012



Zitat zur individualisierten Medizin

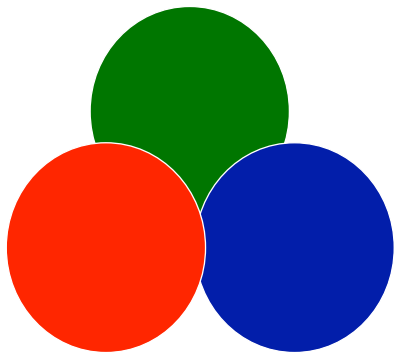
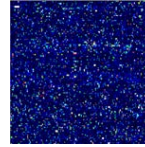
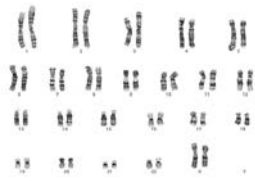
“Die Ärzte kamen sowohl einzeln als auch in Konsilien, redeten viel auf Französisch, auf Deutsch, auf Latein, missbilligten der eine den anderen, verschrieben die unterschiedlichsten Arzneien gegen alle ihnen bekannten Krankheiten; aber nicht einem von ihnen kam der einfache Gedanke in den Sinn, dass ihnen die Krankheit an der sie litt, nicht bekannt sein konnte,

Wie keine einzige Krankheit bekannt sein kann, von der ein lebendiger Mensch ergriffen ist: denn jeder lebendige Mensch hat seine Besonderheiten und immer eine besondere, ihm eigene neue, komplizierte, der Medizin unbekannte Krankheit”



Krieg und Frieden
1869
Leo Tolstoy

Individualisierte Medizin/Genetik?

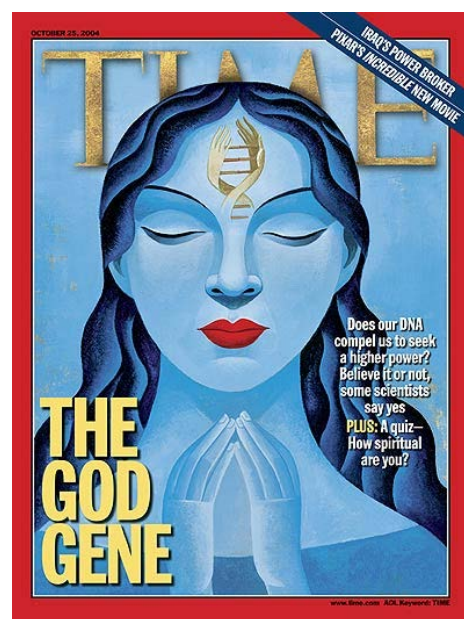
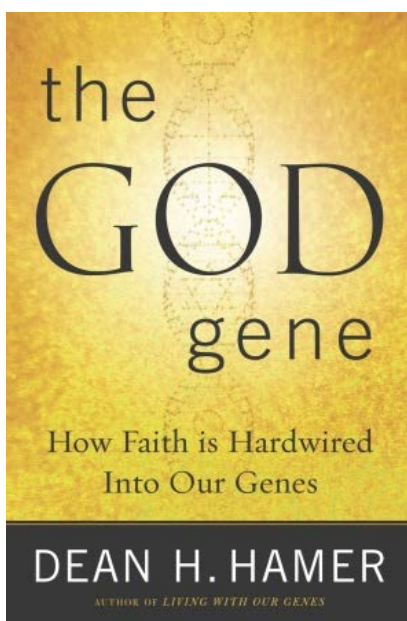


Prediction?

Diagnosis?

Pharmacogenomics?

Wichtigkeit des Phänotyps

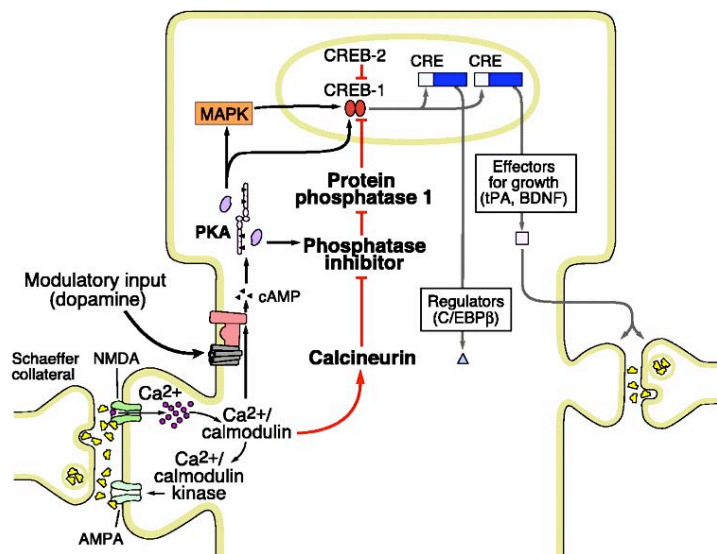
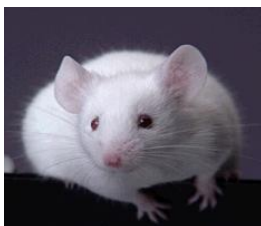


Wichtigkeit des Phänotyps

- Heritabel
- Reliabel, Valide
- Biologisches Korrelat

Memory

Molekulare Kaskade



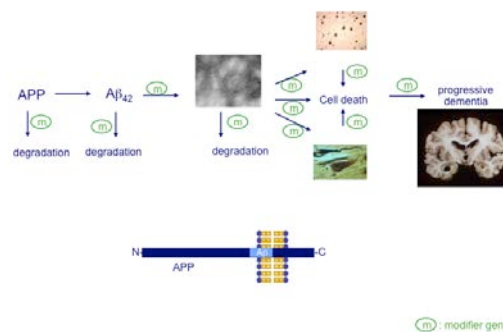
Kandel, Science, 2001

Molekulare Kaskade beim Menschen?

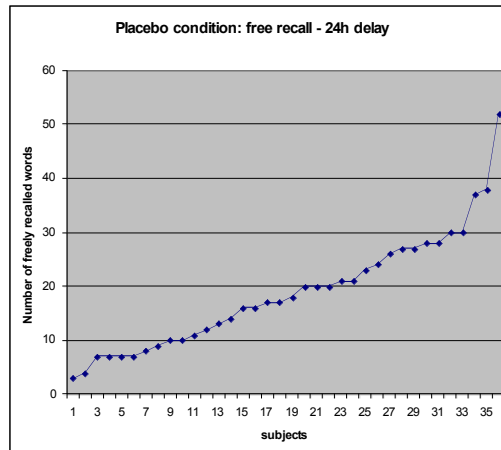


Wichtigkeit der humangenetischen Forschung

Identifizierung von gedächtnis-relevanten Molekülen



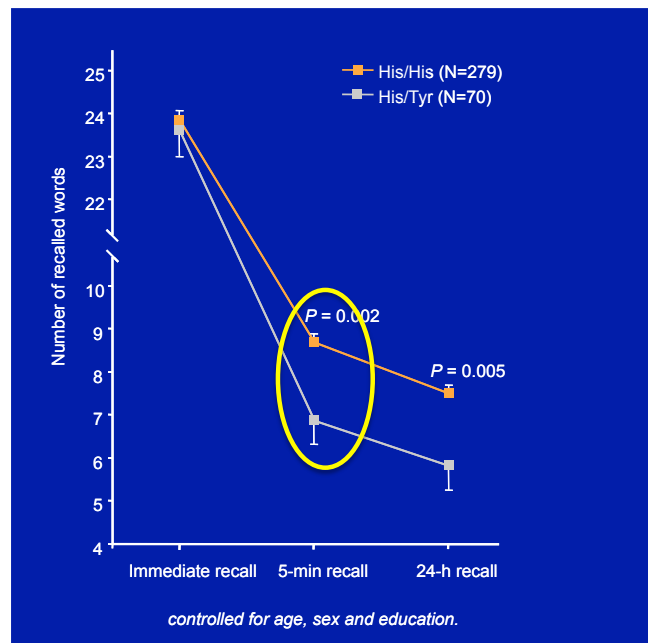
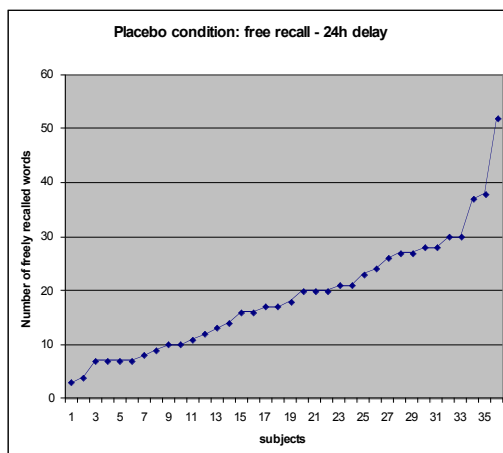
Wichtig: Physiologische Variabilität



▶ Heritabilität: 50%

▶ Gene?

5-HT2a receptor gene



de Quervain, Henke, Aerni, Coluccia, Wollmer, Hock, Nitsch, Papassotiropoulos
Nature Neuroscience (2003)

Wichtigkeit der humangenetischen Forschung

Verständnis neuronaler Mechanismen

Emotionales Gedächtnis



▶ Variabilität?

▶ Gene?

Emotionales Gedächtnis

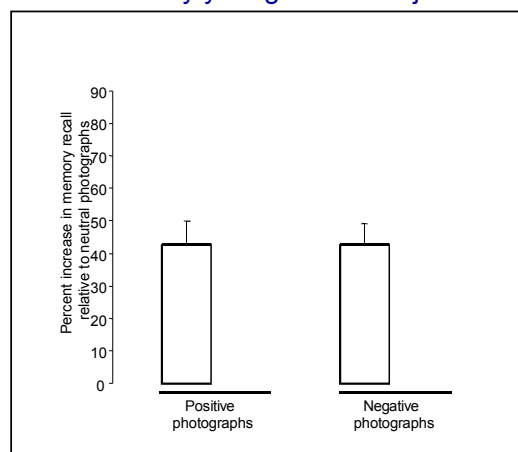
- 435 participants
- Emotional and neutral photographs



de Quervain, Kolassa, Ertl, Lamaro Onyut, Neuner, Elbert & Papassotiropoulos
Nature Neuroscience, 2007

α 2B-adrenergic receptor

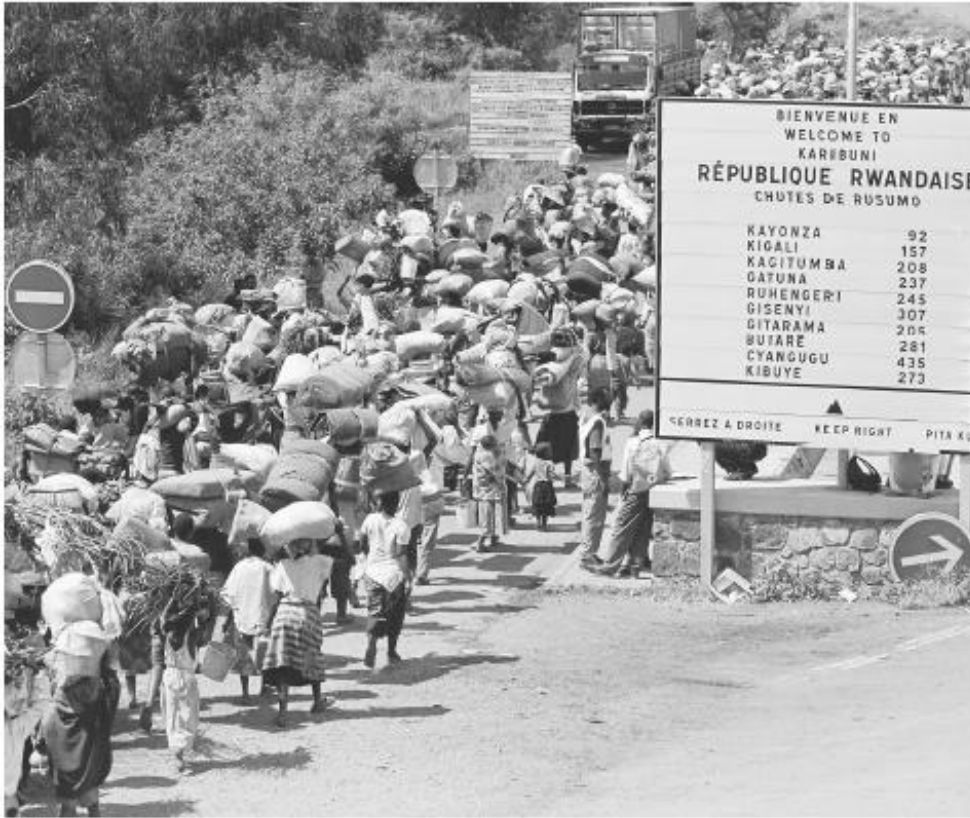
435 healthy young Swiss subjects



Traumatisches Gedächtnis?

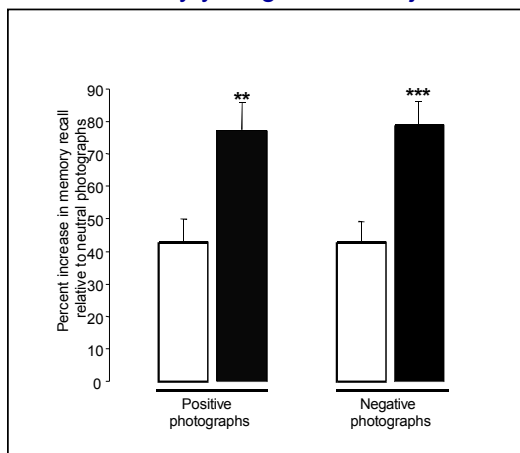
de Quervain, Kolassa, Ertl, Onyut, Neuner, Elbert, Papassotiropoulos, *Nature Neuroscience 2007*

Traumatisches Gedächtnis

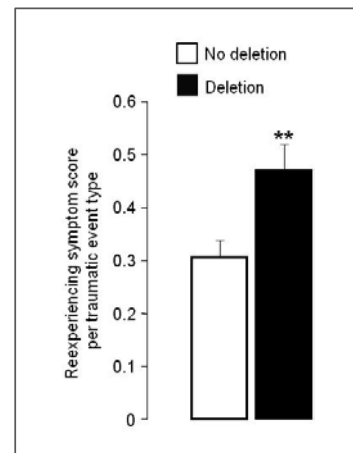


α 2B-adrenergic receptor

435 healthy young Swiss subjects

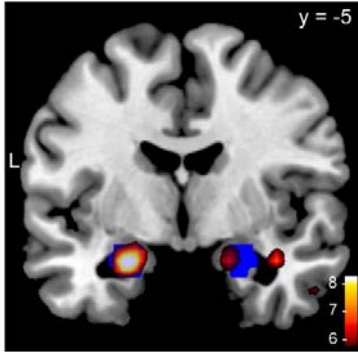


202 survivors of the Rwandan genocide

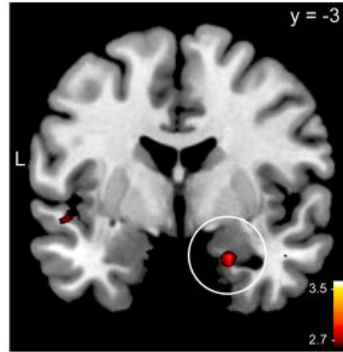


Physiologie => Pathophysiologie

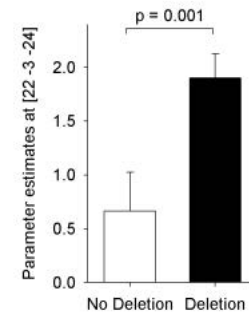
Funktionelle Bildgebung



Genotype-independent amygdala activation during encoding of negative vs. neutral pictures.



Genotype-dependent brain activity in the amygdala

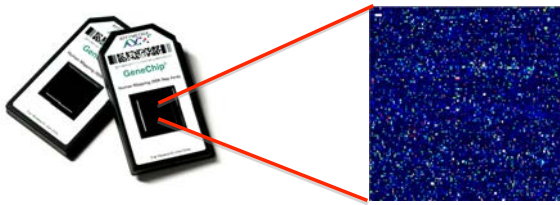


Rasch et al., *PNAS*, 2009

Wichtigkeit der humangenetischen Forschung

Identifizierung von potentiellen Medikamenten

High-throughput genome scans



> 900000 SNPs
> 900000 probes for CNVs



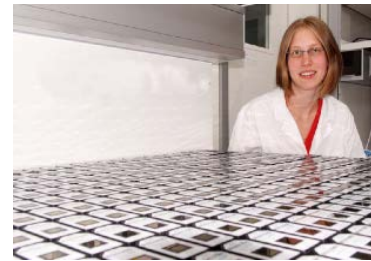
Hybridization



Washing

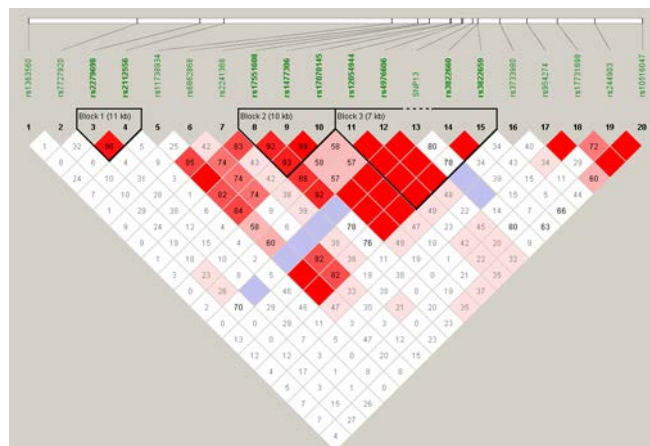
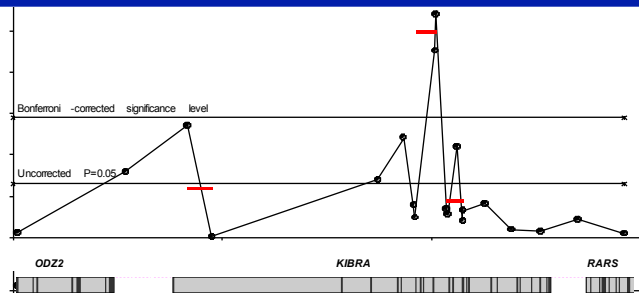
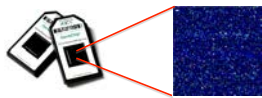


Scanner



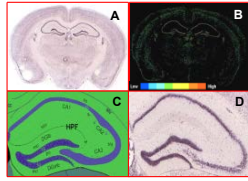
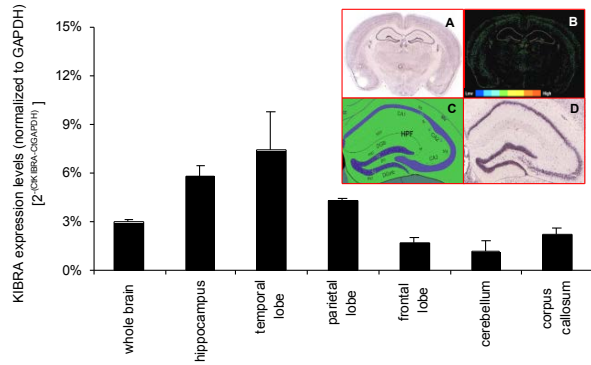
Standardized processing enables the study of large populations

KIBRA

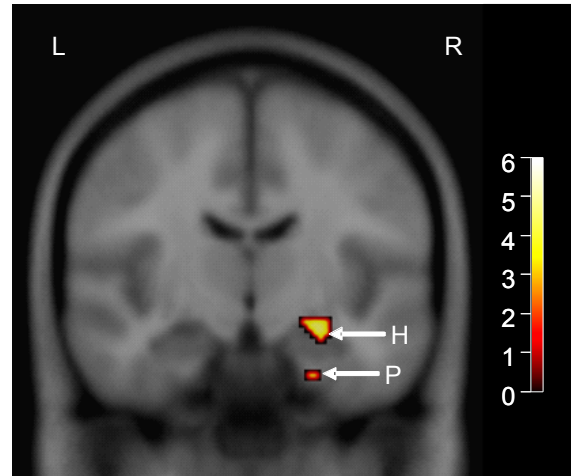


KIBRA

Gene expression



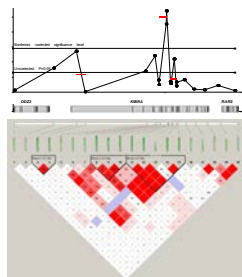
fMRI



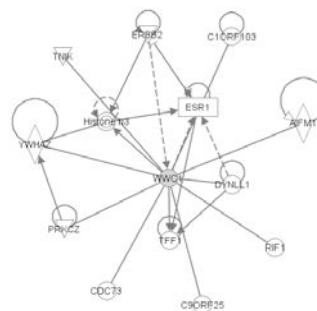
Papassotiropoulos et al. *Science*, 2006

Genetic Information => Drug targets

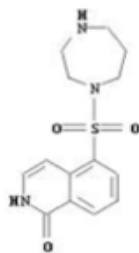
KIBRA



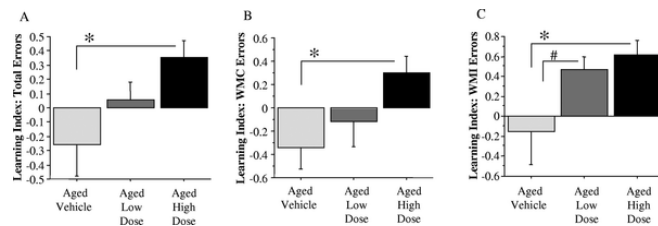
Pathway



Fasudil



Treatment



Huentelman et al., 2009

Wichtigkeit der humangenetischen Forschung

Falsche Schlüsse vermeiden!

ODER: Der Unterschied zwischen Gruppenstatistik und Individuum ist gross!
(...aber sehr schwer zu vermitteln)

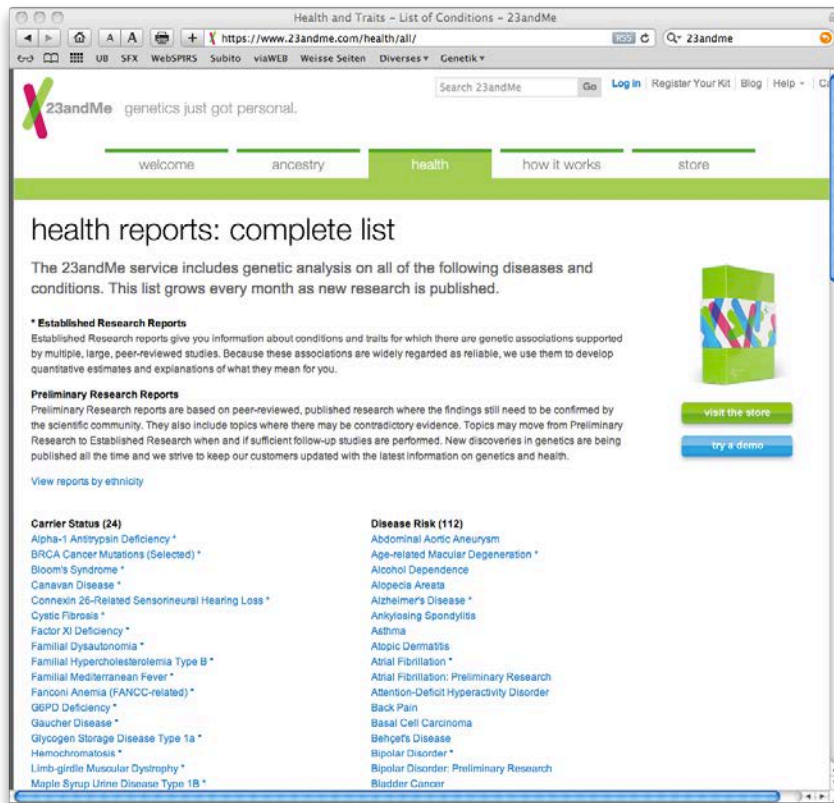
Fehler



Alcoholism-Depression Gene ID'd!

Habe ich das Gen?

Personalized medicine?



Personalized medicine?

- Colorectal Cancer
- Exfoliation Glaucoma
- Heart Attack
- Lung Cancer
- Lupus (Systemic Lupus Erythematosus)
- Multiple Sclerosis
- Obesity
- Abdominal Aortic Aneurysm
- Attention-Deficit Hyperactivity Disorder
- Alcohol Dependence
- Ankylosing Spondylitis
- Antidepressant Response
- Asthma
- Atrial Fibrillation
- Avoidance of Errors
- Back Pain
- Baldness
- Basal Cell Carcinoma
- Beta-Blocker Response
- Bipolar Disorder: Preliminary Research
- Birth Weight
- Bladder Cancer
- Blood Glucose
- Brain Aneurysm
- Breast Cancer Risk Modifiers
- Breastfeeding and IQ
- C-reactive Protein Level
- Caffeine Metabolism
- Celiac Disease: Preliminary Research
- Chronic Lymphocytic Leukemia
- Cleft Lip and Cleft Palate
- Cluster Headaches
- Creutzfeldt-Jakob Disease
- Developmental Dyslexia
- Endometriosis
- Esophageal Cancer
- Essential Tremor
- Eye Color
- Food Preference
- Freckling
- Gallstones
- Hair Color
- Hair Thickness
- HDL Cholesterol Level
- Height
- Heroin Addiction
- HIV Progression
- High Blood Pressure (Hypertension)
- Intrahepatic Cholestasis of Pregnancy
- Kidney Disease
- Larynx Cancer
- Longevity
- Lou Gehrig's Disease (ALS)
- Male Infertility
- Measures of Intelligence
- Memory
- Neuroblastoma
- Nicotine Dependence
- Obesity: Preliminary Research
- Obsessive-Compulsive Disorder
- Odor Detection
- Oral and Throat Cancer
- Osteoarthritis
- Pain Sensitivity
- Parkinson's Disease: Preliminary Research
- Peripheral Arterial Disease
- Persistent Fetal Hemoglobin
- Placental Abruption
- Preeclampsia
- Progressive Supranuclear Palsy
- Schizophrenia
- Sjögren's Syndrome
- Cutaneous Melanoma
- StatIn Response
- Stomach Cancer
- Tardive Dyskinesia
- Thyroid cancer
- Tourette's Syndrome
- Tuberculosis
- Uterine Fibroids
- Restless Legs Syndrome
- Ulcerative Colitis

Personalized medicine?

About Memory

Studies of twins suggest that genetics has about the same influence as training and other environmental factors on a person's ability to remember a limited number of items over periods of seconds to minutes. That type of recall, referred to by psychologists as short-term memory, is maintained by temporary patterns of brain activity. But long-term memory creates lasting connections between brain cells that can potentially last a lifetime.

Research Report

This Research Report includes results from studies that still need to be confirmed by the scientific community. It also includes topics where there may be contradictory evidence. The results of these studies are not conclusive.

visit the store

try a demo

Episodic memory

Journal	Science
Study Size	11
Replications	None
Contrary Studies	None
Applicable Ethnicities	European
Marker	rs17070145

Who	Genotype	What It Means
	TT	Slightly increased episodic memory.
Greg Mendel (Dad)	CT	Slightly increased episodic memory.
	CC	Typical episodic memory.

In this study, the authors asked people to read or listen to lists of words and then recall them later. People with at least one T at rs17070145 performed about 20% better than those with a C at both copies of the SNP five minutes and 24 hours after seeing or hearing the lists. The SNP lies within a gene called KIBRA that is thought to be involved with episodic memory, which involves the recall of events rather than information. (Note: the KIBRA gene is listed in the Genome Explorer as WWC1.)

Citations

Papassotiropoulos A et al. (2006). "Common Kibra alleles are associated with human memory performance." *Science* 475-8.

Personalized medicine?

- Any one have the gene for increased memory?
- I have it and was wondering how common it is?
- I thought I did...CC

	<i>n</i>	No. of words recalled		
		Immediately	After 5 min	After 24 hours
rs17070145				
CC	164	23.6 ± 0.3	7.6 ± 0.2*	6.7 ± 0.2†
CT/TT	169	24.1 ± 0.3	9.4 ± 0.2*	8.0 ± 0.2†
rs6439886				
TT	265	23.9 ± 0.2	8.4 ± 0.2‡	7.3 ± 0.2§
TC/CC	76	24.2 ± 0.4	9.8 ± 0.4‡	8.4 ± 0.4§
* <i>P</i> = 0.000004		† <i>P</i> = 0.0008	‡ <i>P</i> = 0.002	§ <i>P</i> = 0.022

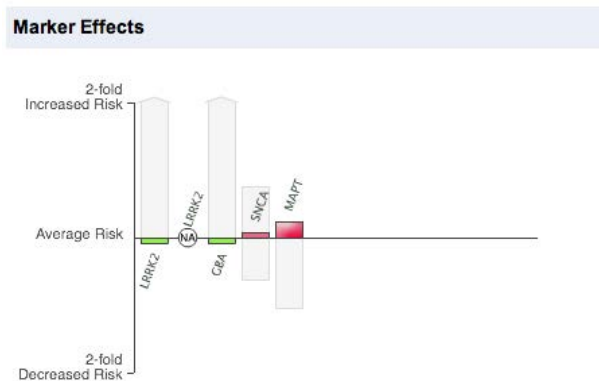
Papassotiropoulos et al. *Science*, 2006

Wichtigkeit der humangenetischen Forschung

Falsche Schlüsse vermeiden!

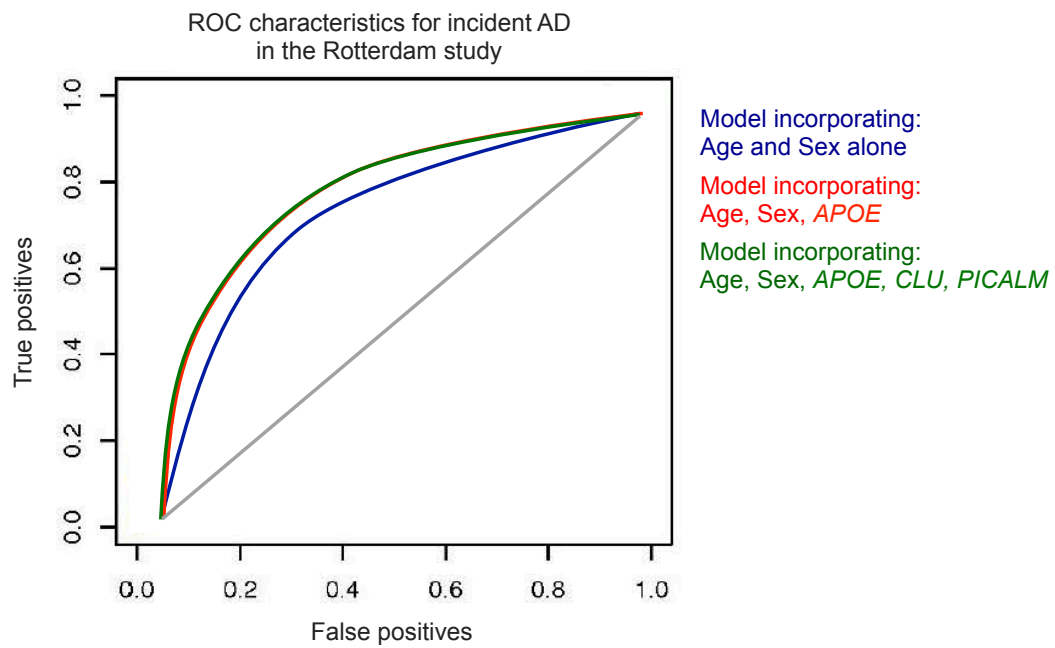
ODER: Addiere nie populations-basierte Risiken um individuelle "Vorhersagen" zu machen!
(...auch schwer zu vermitteln)

Individual risk assessment



- A simple addition implies that the effects are independent of each other
- A simple addition ignores the importance of gene-gene and gene-environment interactions

Improving prediction?



Improving prediction?

A multilocus genetic risk score for coronary heart disease: case-control and prospective cohort analyses

Samuli Ripatti, Emmi Tikkanen, Marju Orho-Melander, Aki S Havulinna, Kaisa Silander, Amitabh Sharma, Candace Guiducci, Markus Perola, Antti Jula, Juha Sinisalo, Marja-Liisa Lokki, Markku S Nieminen, Olle Melander, Veikko Salomaa, Leena Peltonen*, Sekar Kathiresan

Summary

Background Comparison of patients with coronary heart disease and controls in genome-wide association studies has revealed several single nucleotide polymorphisms (SNPs) associated with coronary heart disease. We aimed to establish the external validity of these findings and to obtain more precise risk estimates using a prospective cohort design.

Methods We tested 13 recently discovered SNPs for association with coronary heart disease in a case-control design including participants differing from those in the discovery samples (3829 participants with prevalent coronary heart disease and 48897 controls free of the disease) and a prospective cohort design including 30725 participants free of cardiovascular disease from Finland and Sweden. We modelled the 13 SNPs as a multilocus genetic risk score and used Cox proportional hazards models to estimate the association of genetic risk score with incident coronary heart disease. For case-control analyses we analysed associations between individual SNPs and quintiles of genetic risk score using logistic regression.

Findings In prospective cohort analyses, 1264 participants had a first coronary heart disease event during a median 10.7 years' follow-up (IQR 6.7–13.6). Genetic risk score was associated with a first coronary heart disease event. When compared with the bottom quintile of genetic risk score, participants in the top quintile were at 1.66-times increased risk of coronary heart disease in a model adjusting for traditional risk factors (95% CI 1.35–2.04, p value for linear trend = 7.3×10^{-10}). Adjustment for family history did not change these estimates. Genetic risk score did not improve C index over traditional risk factors and family history ($p=0.19$), nor did it have a significant effect on net reclassification improvement (2.2%, $p=0.18$); however, it did have a small effect on integrated discrimination index (0.004, $p=0.0006$). Results of the case-control analyses were similar to those of the prospective cohort analyses.

The Lancet, 2010

Personalized medicine?

Psynomics – Genomics For The New Psychiatry

http://www.psynomics.com/

UB SFX WebSPIRS Subito viaWEB Weisse Seiten Diverses Genetik

Psynomics™
genomics for the *new* psychiatry

Welcome to Psynomics

Psynomics is the first and only company in the world to offer DNA-based diagnostic and therapeutic tests to help millions of people suffering from mental illness.

Our first two products:

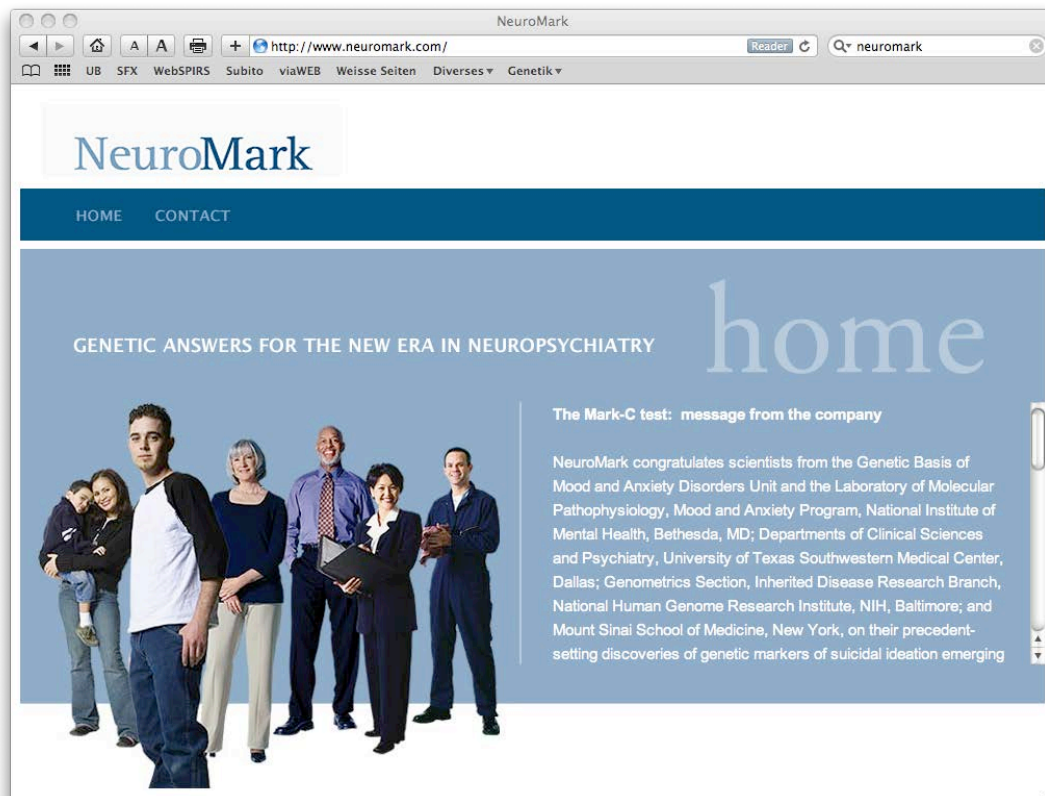
Psynome™ – tests for two mutations of a gene that are associated with bipolar disorder.

Psynome2™ – tests for gene mutations in the Promoter L allele gene that predicts patient response to serotonin-based drugs, the most commonly prescribed drug therapies in psychiatry today. These tests are useful to your doctor in making a timely and accurate diagnosis of your condition and prescribing the right medication. The tests can be ordered individually or combined.

LEARN
more about our company

[Privacy Policy](#) | [Consent Form](#) | [Mailer Instructions](#) | [Model Report](#)

Personalized medicine?



Personalized medicine?

Article

Genetic Markers of Suicidal Ideation Emerging During Citalopram Treatment of Major Depression

Gonzalo Laje, M.D.

Silvia Paddock, Ph.D.

Husseini Manji, M.D.

A. John Rush, M.D.

Alexander F. Wilson, Ph.D.

Dennis Charney, M.D.

Francis J. McMahon, M.D.

Objective: Suicidal ideation is an uncommon symptom than can emerge during antidepressant treatment. The biological basis of treatment-emergent suicidal ideation is unknown. Genetic markers may shed light on the causes of treatment-emergent suicidal ideation and help identify individuals at high risk who may benefit from closer monitoring, alternative treatments, or specialty care.

Method: A clinically representative cohort of outpatients with major depressive disorder who enrolled in the Sequenced Treatment Alternatives to Relieve Depression (STAR*D) trial were treated with citalopram under a standard protocol for up to 14 weeks. DNA samples from 1,915 participants were genotyped for 768 single-nucleotide polymorphisms in 68 candidate genes. Allele and genotype frequencies were compared between the 120 participants who developed treat-

ment-emergent suicidal ideation and those who did not.

Results: Two markers were significantly associated with treatment-emergent suicidal ideation in this sample (marker rs4825476, $p=0.0000784$, odds ratio=1.94; permutation $p=0.01$; marker rs2518224, $p=0.0000243$, odds ratio=8.23; permutation $p=0.003$). These markers reside within the genes GRIA3 and GRIK2, respectively, both of which encode ionotropic glutamate receptors.

Conclusions: Markers within GRIK2 and GRIA3 were associated with treatment-emergent suicidal ideation during citalopram therapy. If replicated, these findings may shed light on the biological basis of this potentially dangerous adverse event and help identify patients at increased risk.

Personalized medicine


Companies address important issues, BUT

Many implications not supported by current data!!

Personalized medicine

naturenews Full text access provided to **University of Basel** by Acquisitions

[nature news home](#) [news archive](#) [specials](#) [opinion](#) [features](#) [news blog](#) [events blog](#) [nature journal](#)

 [comments on this story](#) Published online 23 November 2009 | Nature | doi:10.1038/news.2009.1102


News

Diagnosing the future of genomics

Eric Green discusses his priorities as newly appointed director of the US National Human Genome Research Institute.

[Erika Check Hayden](#)

On 17 November, US National Institutes of Health (NIH) director Francis Collins named Eric Green as head of the National Human Genome Research Institute in Bethesda, Maryland — the post held by Collins before he became NIH director. Green talked to *Nature* about his plans for the institute, which has a budget of almost US\$500



Eric Green is the new director of the US National Human Genome Research Institute. *M. Bartlett, NHGRI*

most recent **commented**

- [Single-celled life does a lot with very little](#)
26 November 2009
- [Medical Research Council chief to step down](#)
26 November 2009
- [Japanese scientists rally against government cuts](#)
26 November 2009
- [Plans for cutting emissions could also benefit health](#)
25 November 2009
- [Spin success for silicon](#)
25 November 2009

Related stories

- [The new head of the NIH](#)
01 October 2009
- [Collins sets out his vision for the NIH](#)
18 August 2009


Stories by subject

- [Genetics](#)
- [Policy](#)

Stories by keywords

- [Eric Green](#)
- [National Institutes of Health](#)
- [National Human Genome Research Institute](#)
- [genomics](#)
- [personal genomics](#)

This article elsewhere

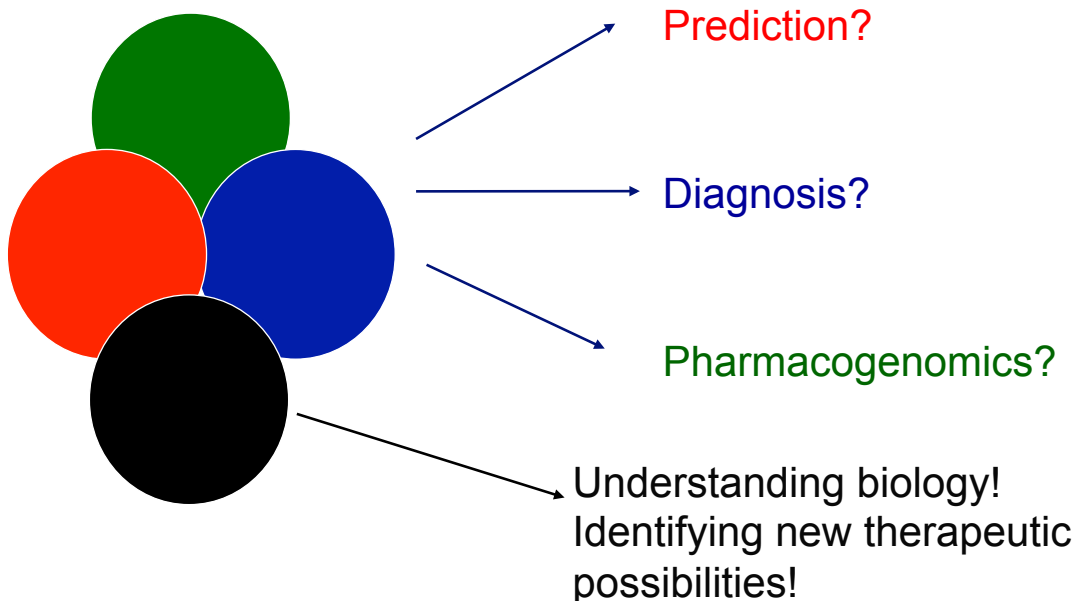
 [Blogs linking to this article](#)

Personalized medicine

What advice would you have for people who are considering buying personal genomics services from a company to find out their genetic risk for common diseases?

I haven't yet gone to get that information, because I think that the amount of information available at this time wouldn't really change anything that I am doing. A lot of what I know about my own health is based on family history — I think that understanding family history, and making sure your physician knows that, is incredibly valuable, and that's where I would put my priority at the moment. But it is a changing landscape, so I don't think any advice I would give today would be the same a year from now.

Personalized medicine?

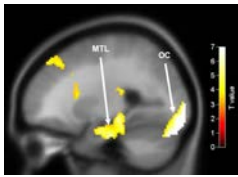




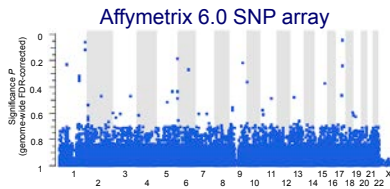
Current status



memory tasks in health and disease
N > 3000



structural and functional brain imaging
Current N = 600

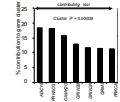



Clinical trials

single loci



gene/loci groups



computational modeling



biological validation

