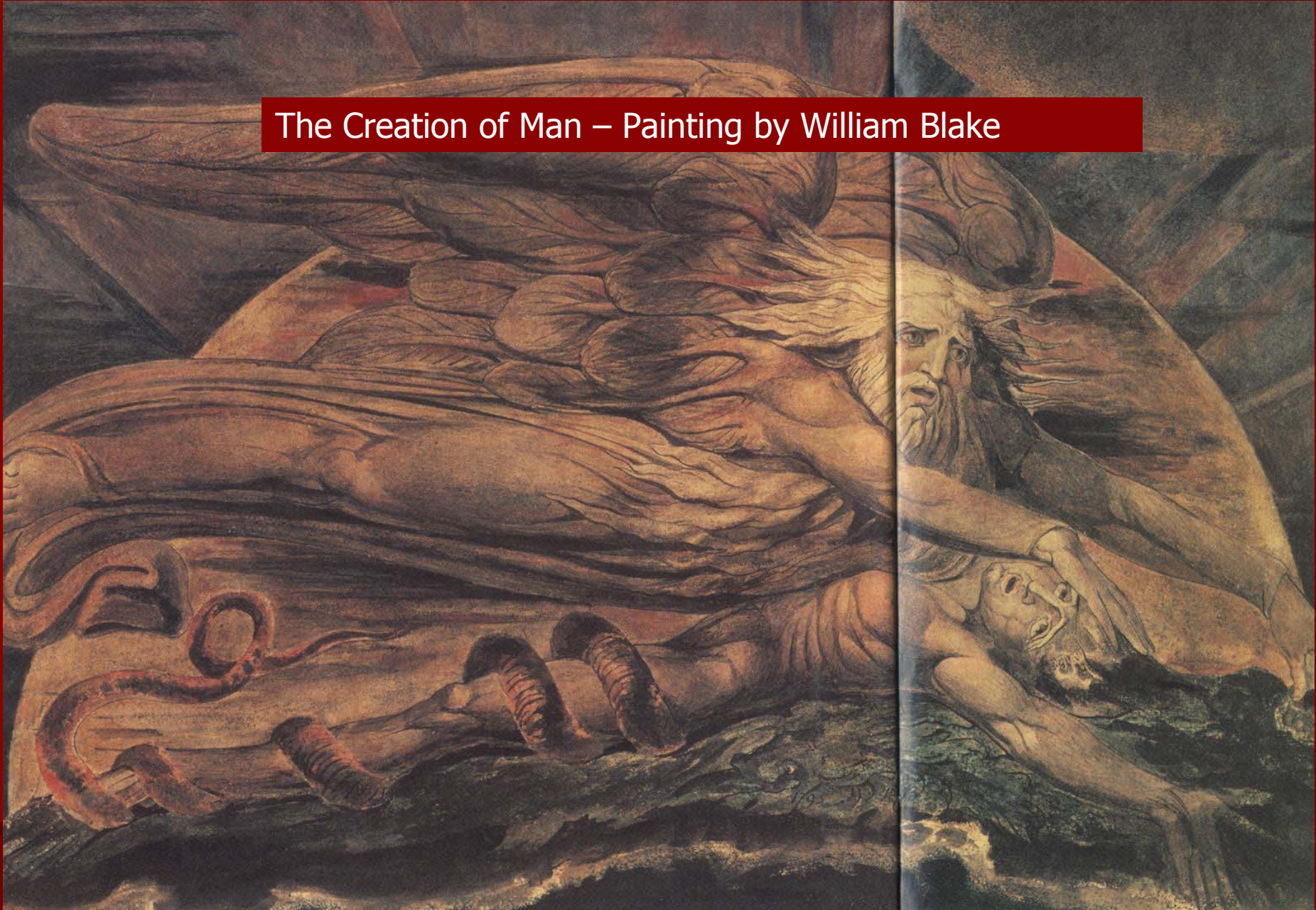


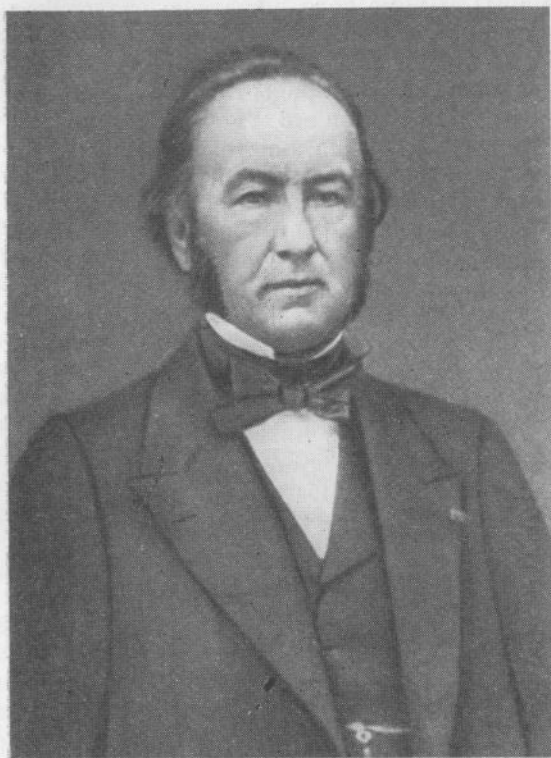


А.Л.Маркель

ГЕНЕТИЧЕСКИЕ
МОДЕЛИ
ПАТОЛОГИИ
ЧЕЛОВЕКА:
АРТРИАЛЬНАЯ ГИПЕРТОНИЯ И ДР.

The Creation of Man – Painting by William Blake





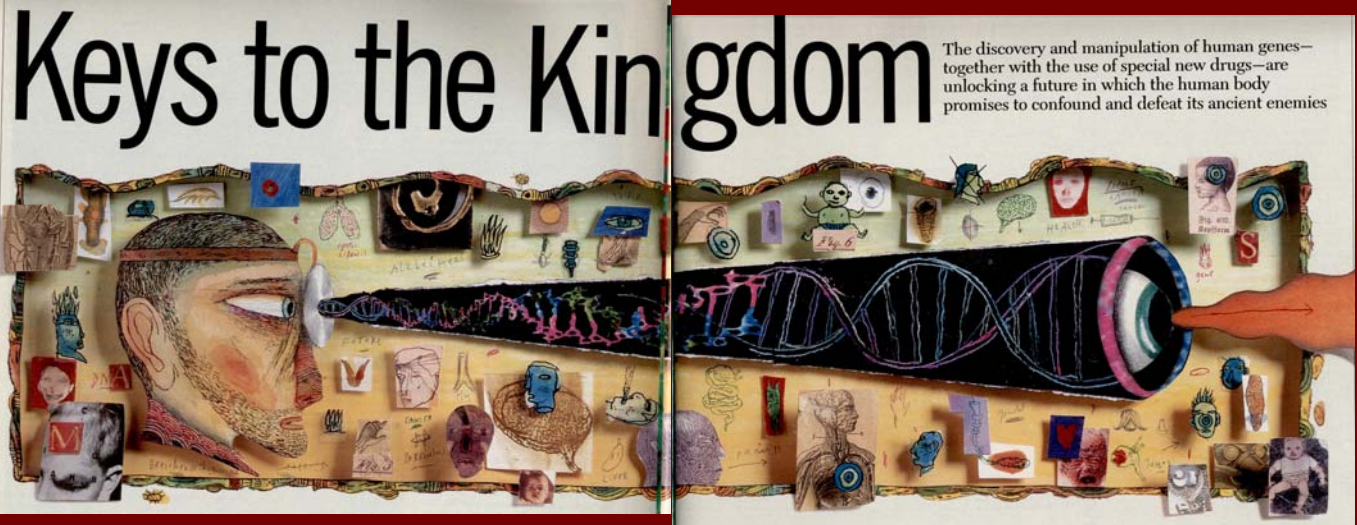
Клод Бернар (1813-1878)

Введение к изучению опытной медицины (1865)

«Единственной нашей целью всегда было и есть – содействовать тому, чтобы всем известные начала опытного метода проникли в медицинские науки»

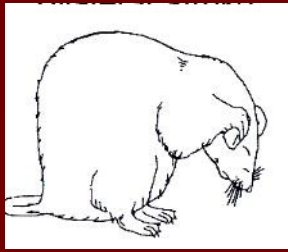
«Всепроникающее око экспериментальной биологии и медицины»

TIME, v148, N14





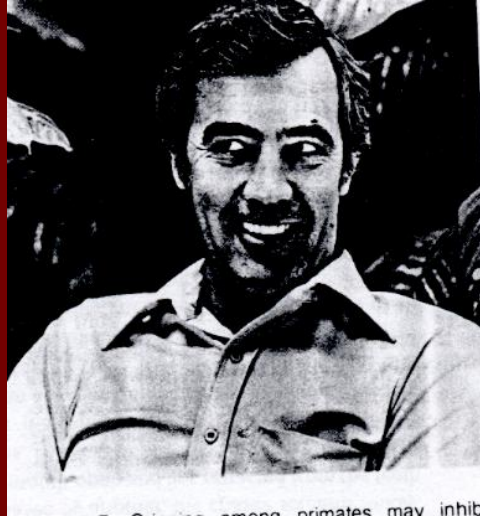
Displacement activity



«умывание»



«Песни любви»



Some species of primates may inhibit

"The Morbid Anatomy of the Human Genome" (MacKusick)

Cystic fibrosis (Mucoviscidosis)

Severe progressive disease of the bronchial system and gastrointestinal tract

Disturbed function of a chloride ion channel by mutations of the *CFTR* gene

Autosomal recessive

Gene locus 7q31.3

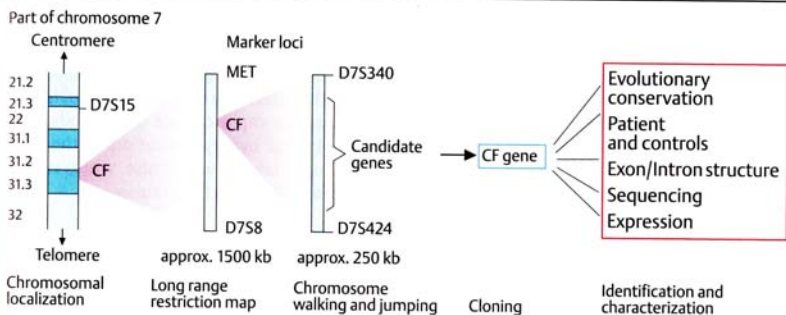
Disease incidence approx. 1:2500

Heterozygote frequency approx. 1:25

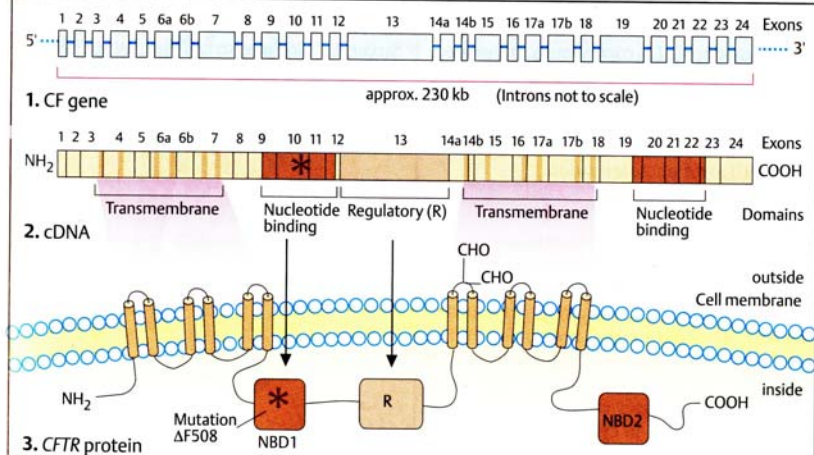
Mutation $\Delta F508$ in approx. 70%



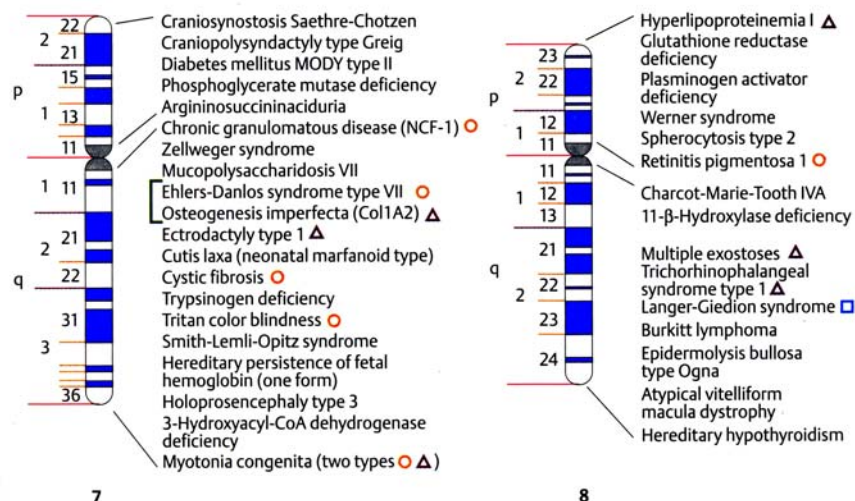
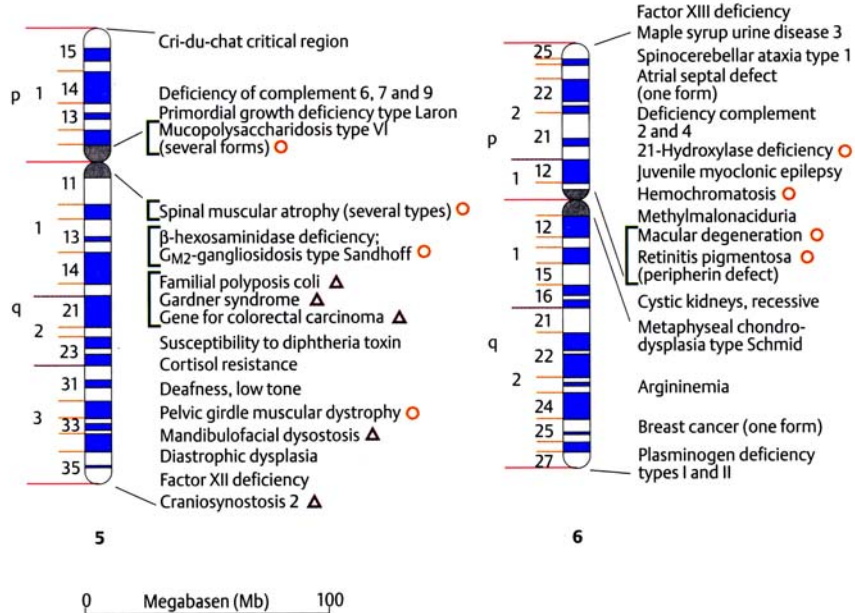
A. Cystic fibrosis, a very frequent recessive disease in Europe and North America



B. Positional cloning of the gene for cystic fibrosis (CF)



C. CF gene and CFTR protein



Prolonged QT interval in the electrocardiogram



Prolonged QT

Syncope

Sudden death

Autosomal dominant

Six genes involved (LQT1 - LQT6)



Romano Ward syndrome (Long-QT syndrome)

| Type | Locus | Gene |
|------|---------|-------------------------|
| LQT1 | 11p15.5 | KCNQ1 (<i>KVLQT1</i>) |
| LQT2 | 7q35-36 | <i>HERG</i> |
| LQT3 | 3p21-24 | <i>SCNA5</i> |
| LQT4 | 4q25-27 | unknown |
| LQT5 | 21q22.1 | <i>KCNE1</i> |
| LQT6 | 21q21.1 | <i>KCNE2</i> |

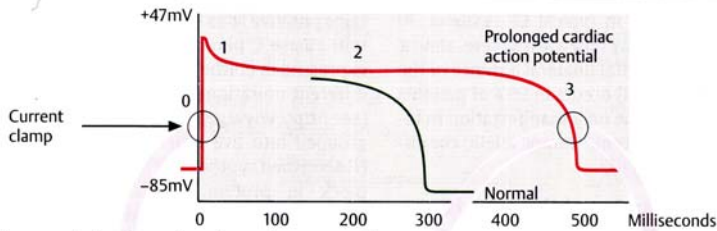
Long-QT and Deafness (Jervell and Lange-Nielsen) due to allelic mutations at LQT1 and LQT5 (autosomal recessive)

1. Main features

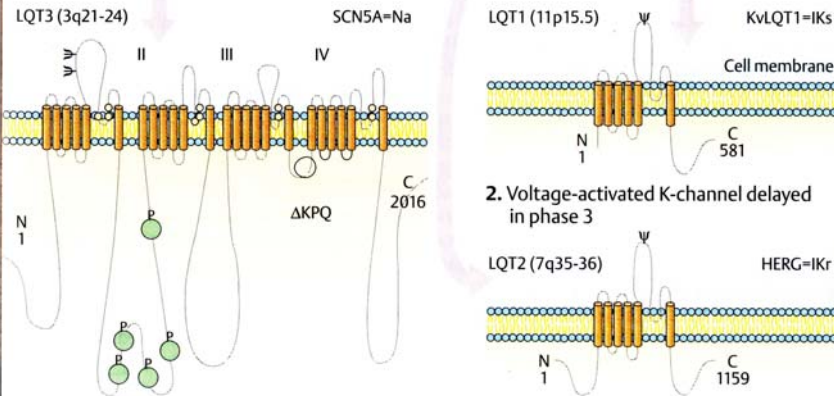
2. Electrocardiogram

3. Genetics

A. Long-QT syndrome, a genetic cardiac arrhythmia

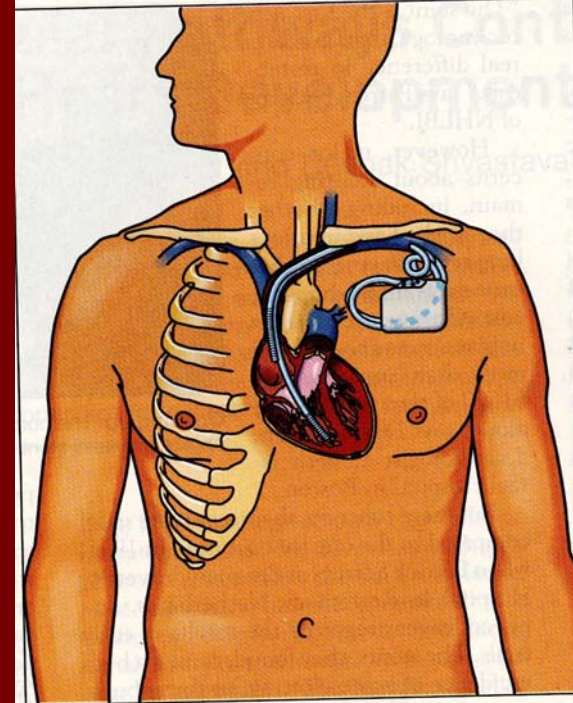


1. Increased duration of cardiac action potential



4. Na-channel fails to inactivate completely during phase 0

B. Different molecular types of long-QT syndrome

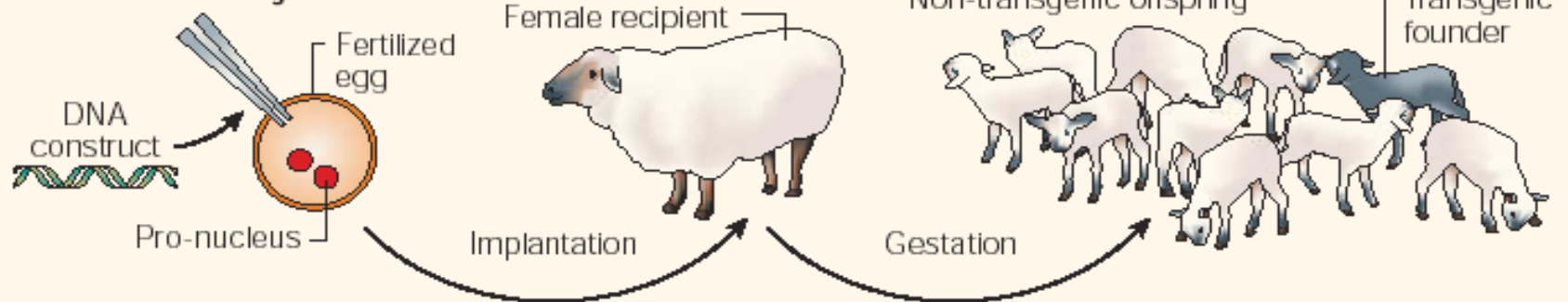


A shock in time. Implantable defibrillators, such as the one shown here, are already saving lives.

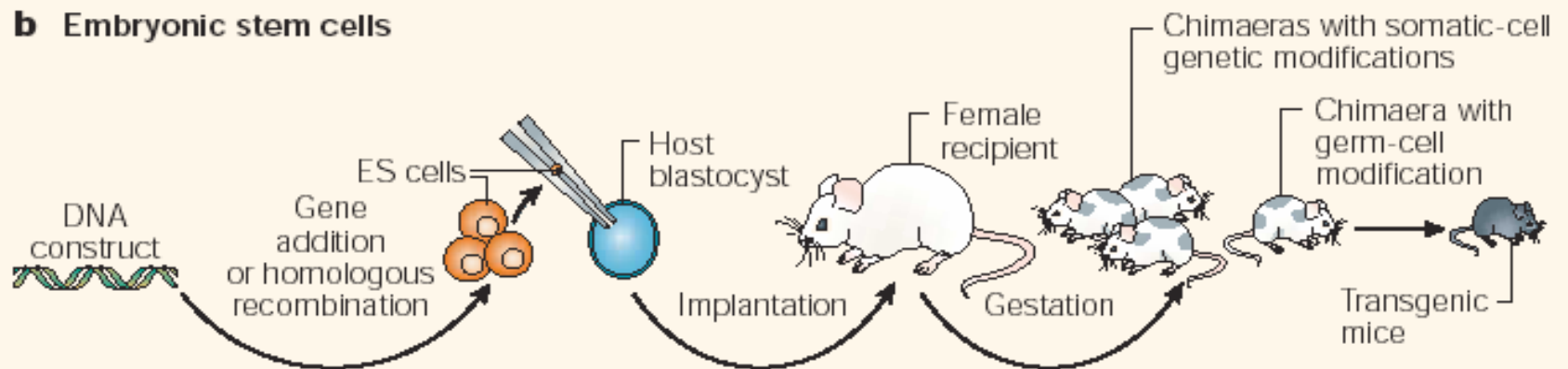
“My whole body jumped, and I was hollering pretty loud. But at least those shocks saved my life” –

remembered 71-year-old patient Salvatore Viviani, who received seven shocks soon after his defibrillator was implanted.

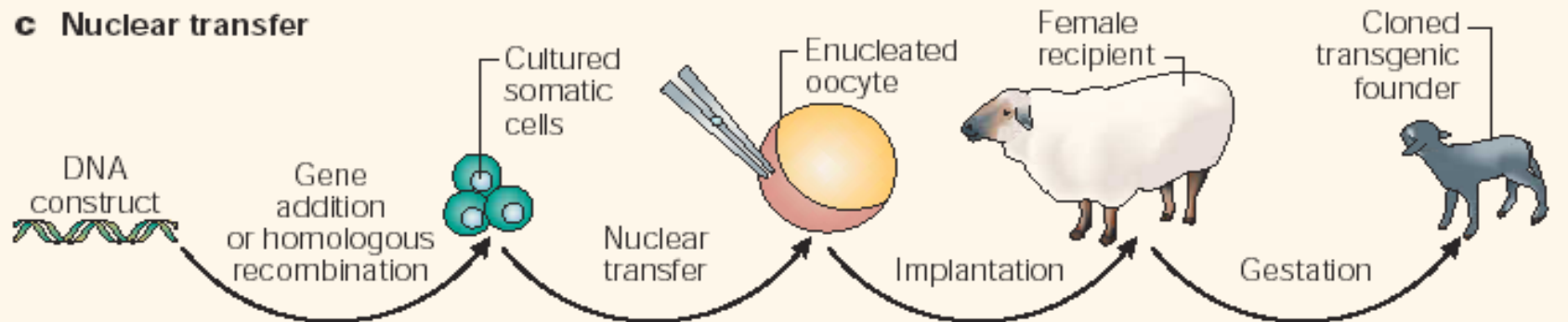
a Pro-nuclear injection



b Embryonic stem cells



c Nuclear transfer



Molecular Genetics of Human Blood Pressure Variation

Richard P. Lifton

Hypertension is a common multifactorial vascular disorder of largely unknown cause

- «As the era of focus on research on monogenic diseases comes to a close, the focus of genetics research is turning to studies of complex, multifactorial disorders such as cardiovascular diseases».
- *Markus Perola*



High Blood Pressure Rising, yet Often Ignored 60% of Americans Have or Are at Risk for High Blood Pressure

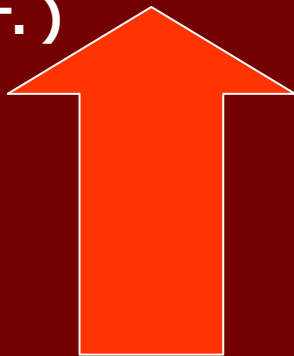
By Jennifer Warner

Reviewed By Brunjl

WebMD Medical News on Monday, October 25, 2004

<http://my.webmd.com/content/>

60%
(2004 г.)

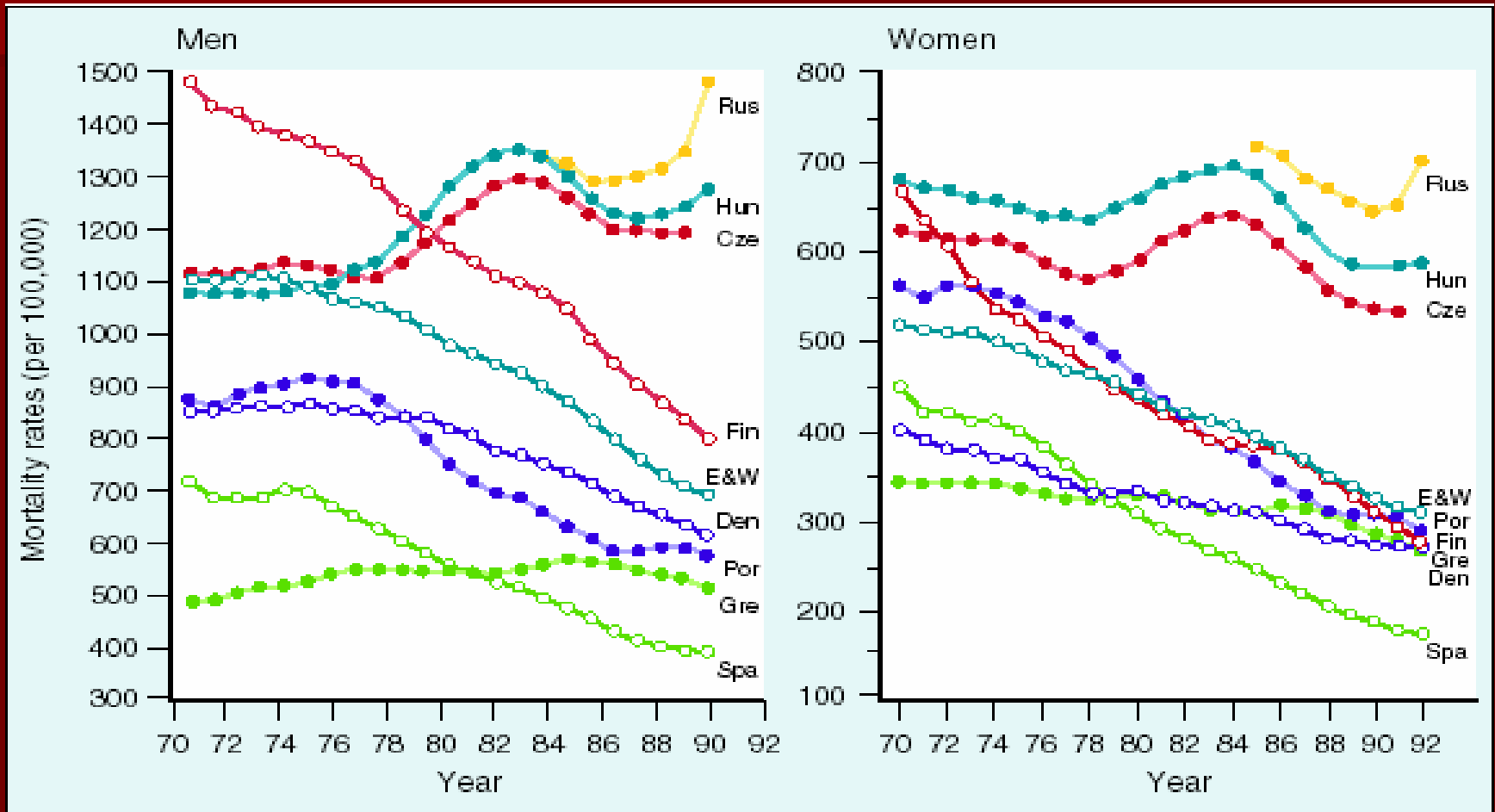


35%
(1995 г.)

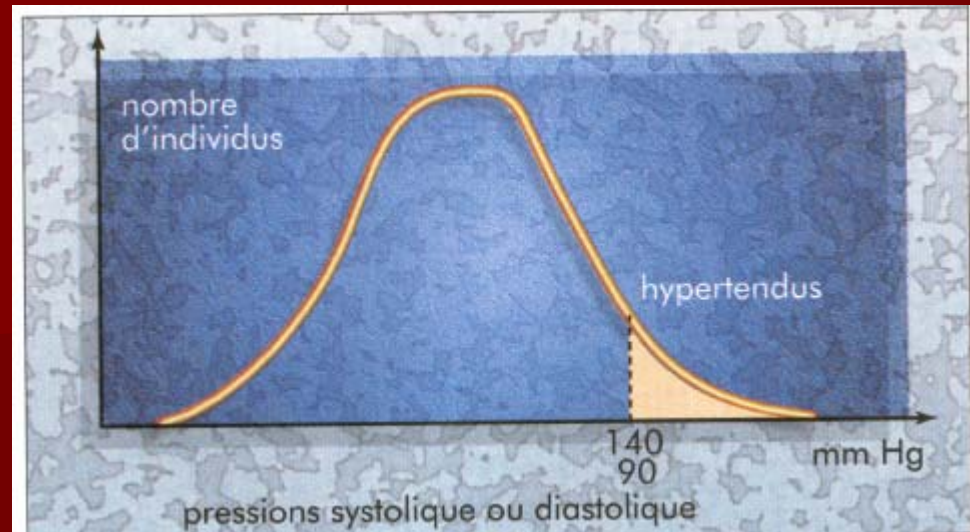


Global Burden of Cardiovascular Disease

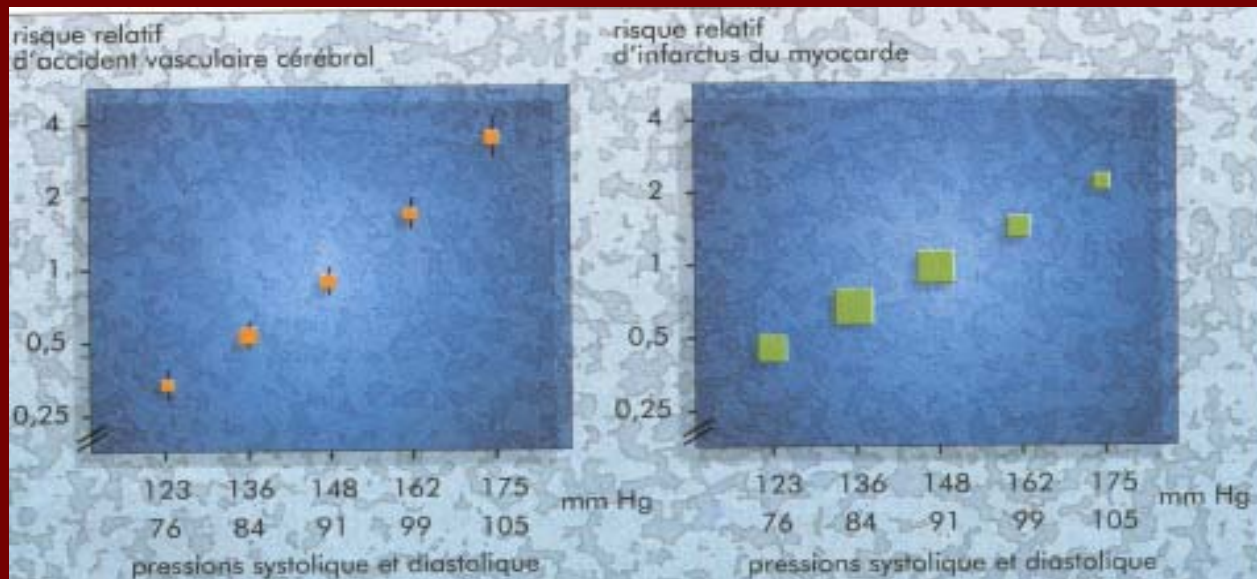
J. Michael Gaziano



140/90



**Какое давление можно считать нормальным
и какое - повышенным?**



125/75

From: Twins in Cardiovascular Genetic Research

Friedrich C. Luft, 2001

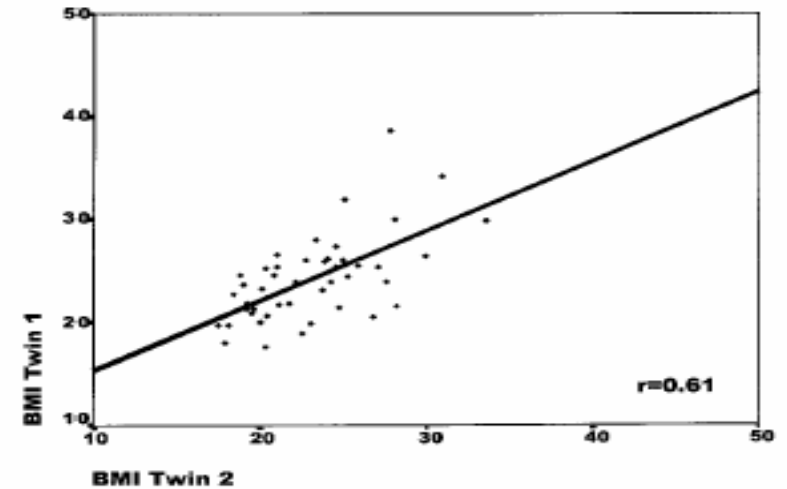
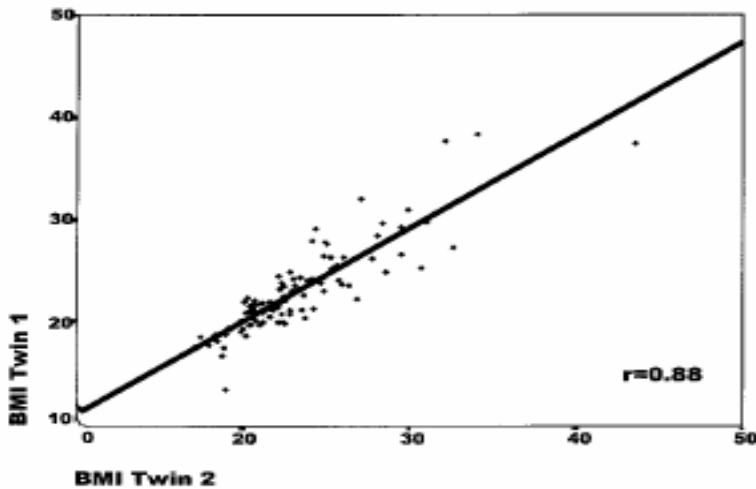
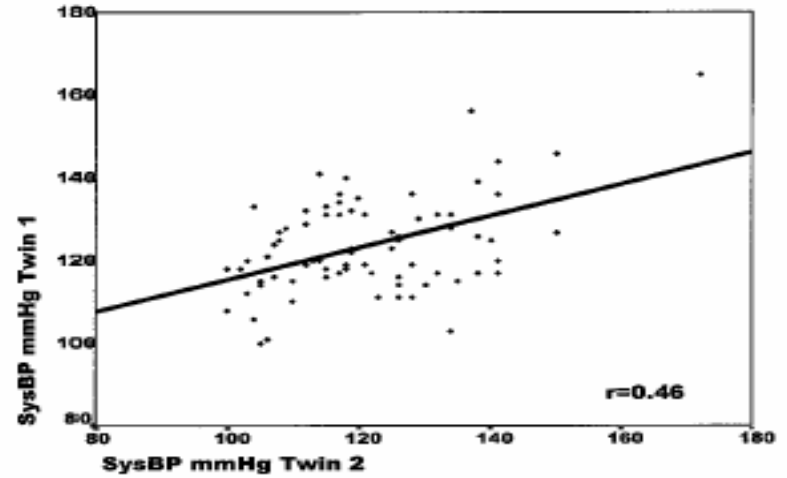
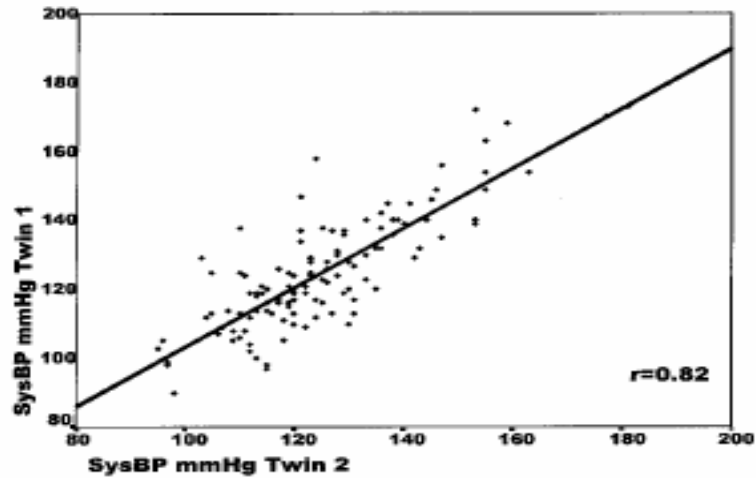
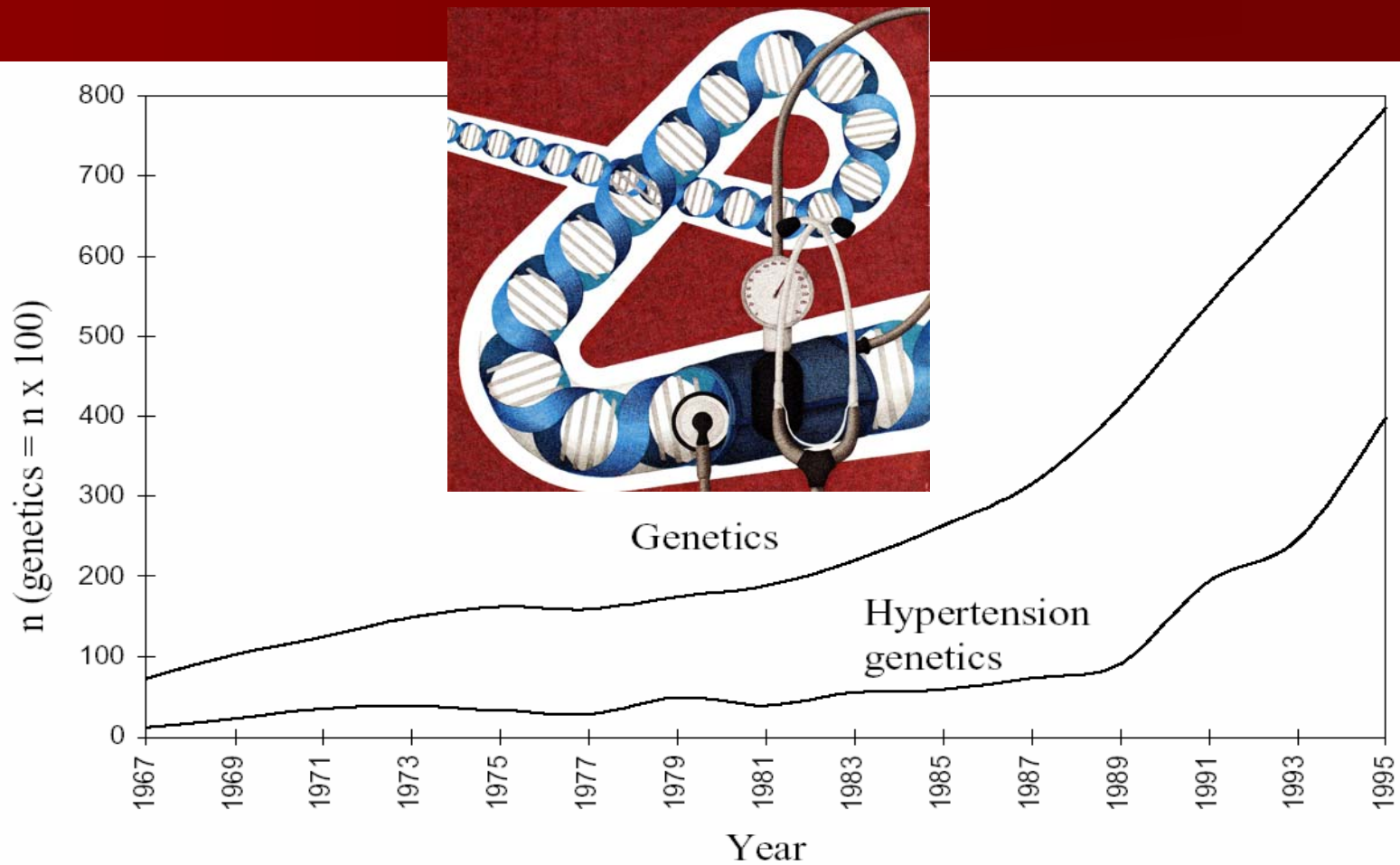


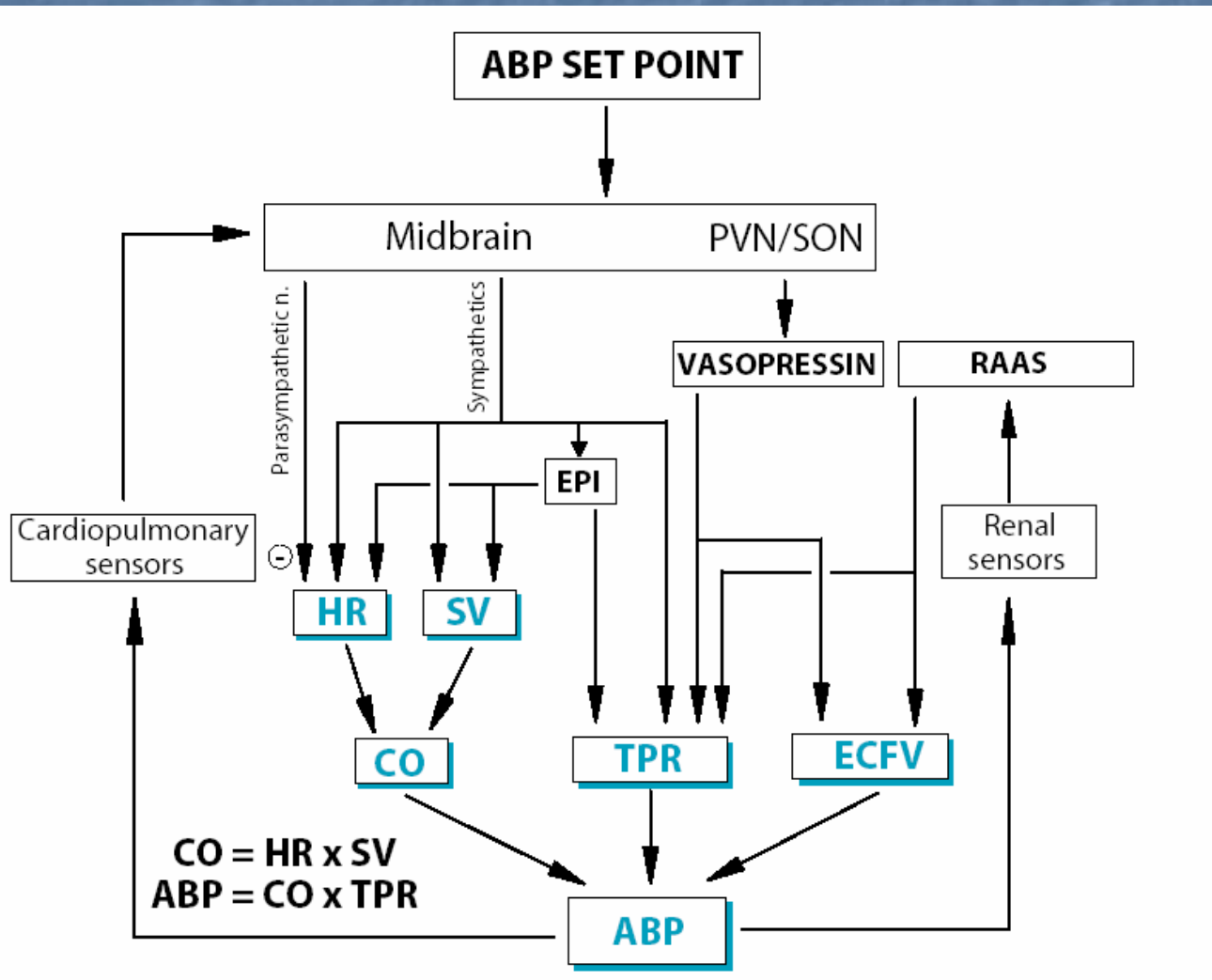
Figure 12. The annual number of publications on genetics or, more specifically, hypertension genetics, reported in MEDLINE© for the years 1967-1995.



From Marcus Perola, 1999

AD = OPG x MOC

$$V = R \times I$$



Tigersted and Bergman 1898 г. – экстракты из ткани почки («ренин») повышают АД

Ambard – 1904 – низкосолевая диета способствует снижению АД

Dole, Dahl, Cotzias – 1951 г. – снижение натрия в пище до уровня менее 230 мг/сутки приводит к понижению АД до 140/90 мм рт.ст. у 30-50% гипертоников

Skeggs – 1940 г. – идентификация ангиотензина II

Tait and Simpson – 1952-3 гг. – открытие альдостерона

Genest, Laragh, Davis, Ganong, Mulrow, Колпаков – 1959-70 гг – развитие представлений о **ренин-ангиотензин-альдостероновой системе** – регуляторе водно-солевого гомеостаза и АД

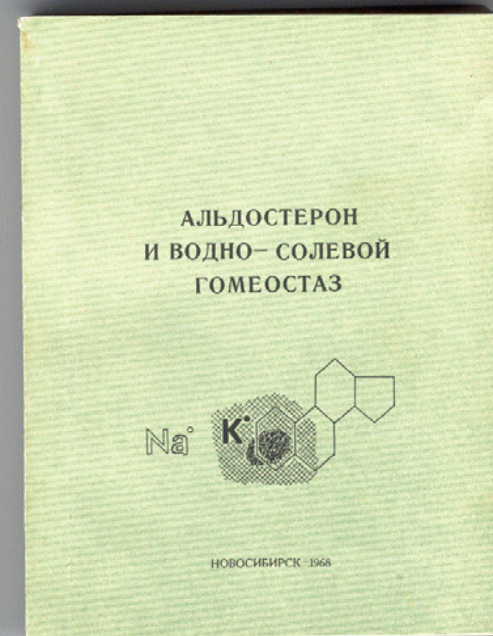


1968 г.

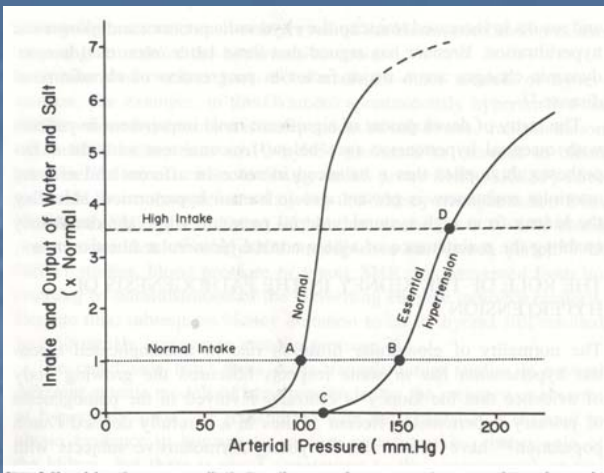
Новосибирск
Академгородок
Дом ученых

Симпозиум

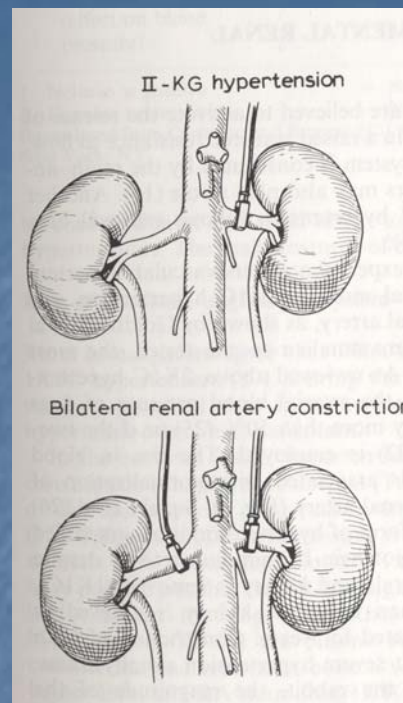
*Альдостерон и
водно-солевой
гомеостаз*



Почка и АД

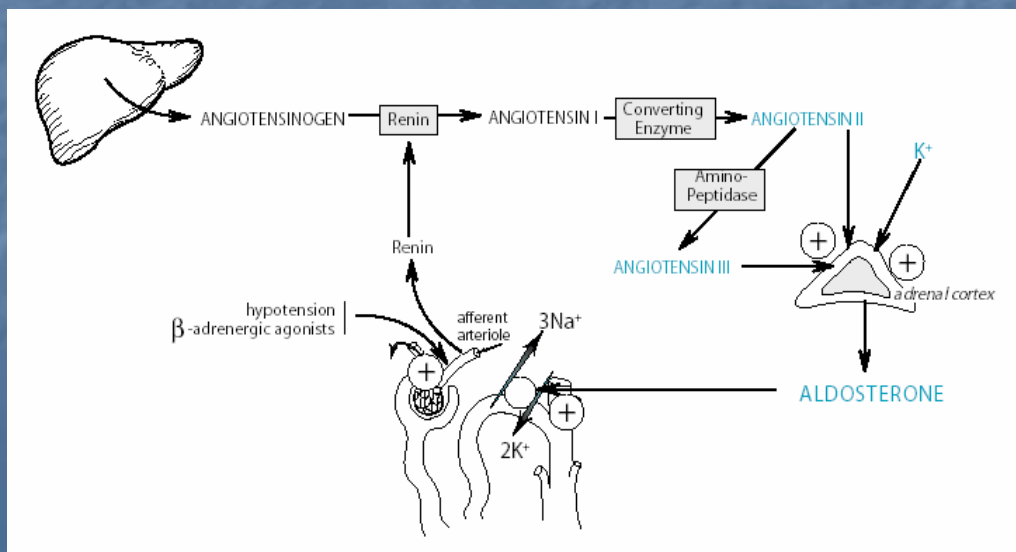
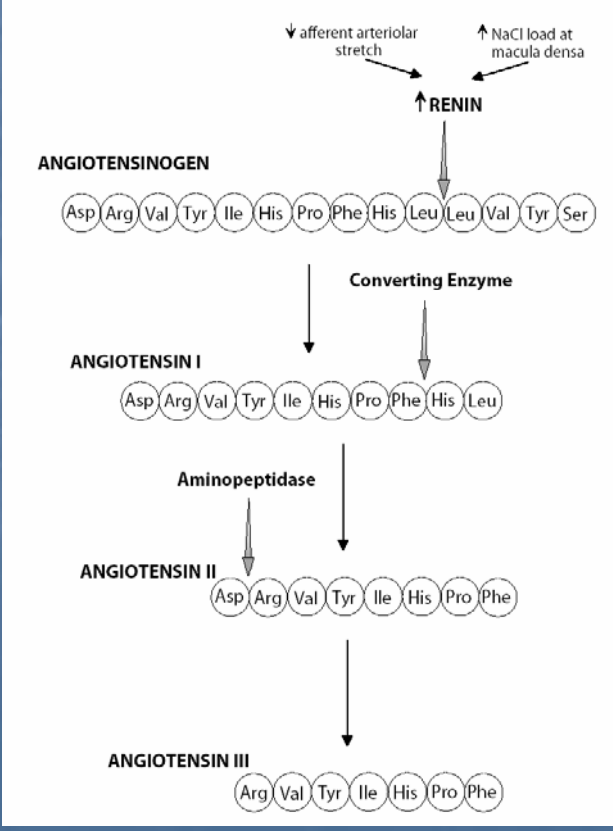


Переключение «установочной точки» (set point) при артериальной гипертонии

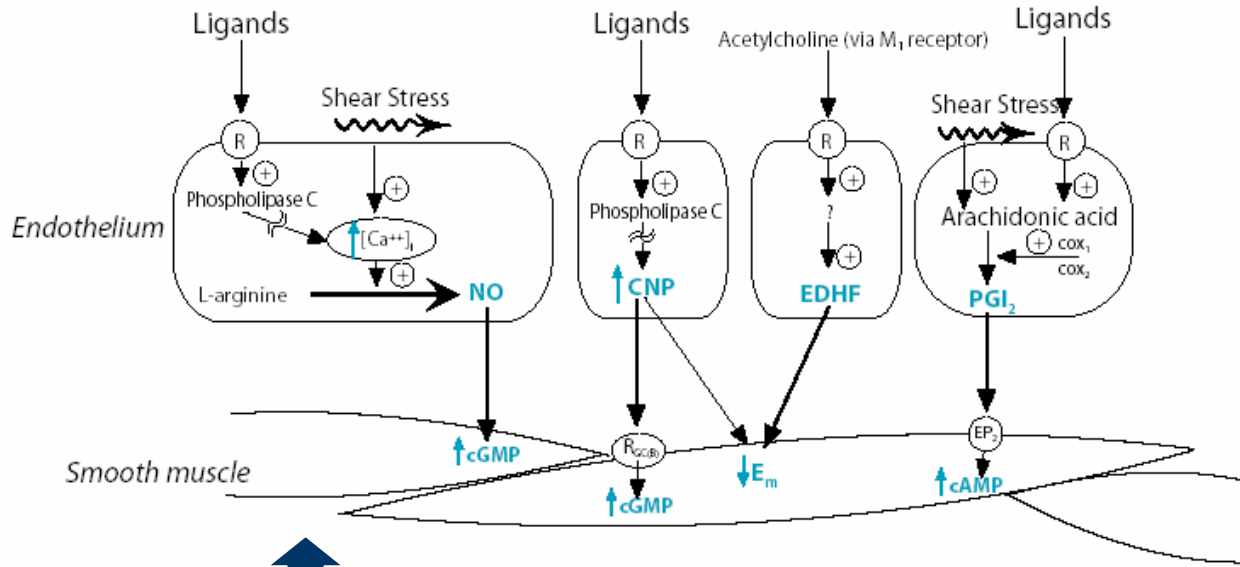


Сужение почечной артерии – модель гипертонии

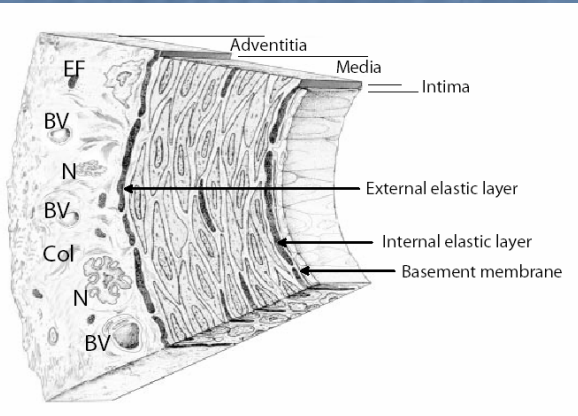
Ренин-ангиотензиновая СИСТЕМА



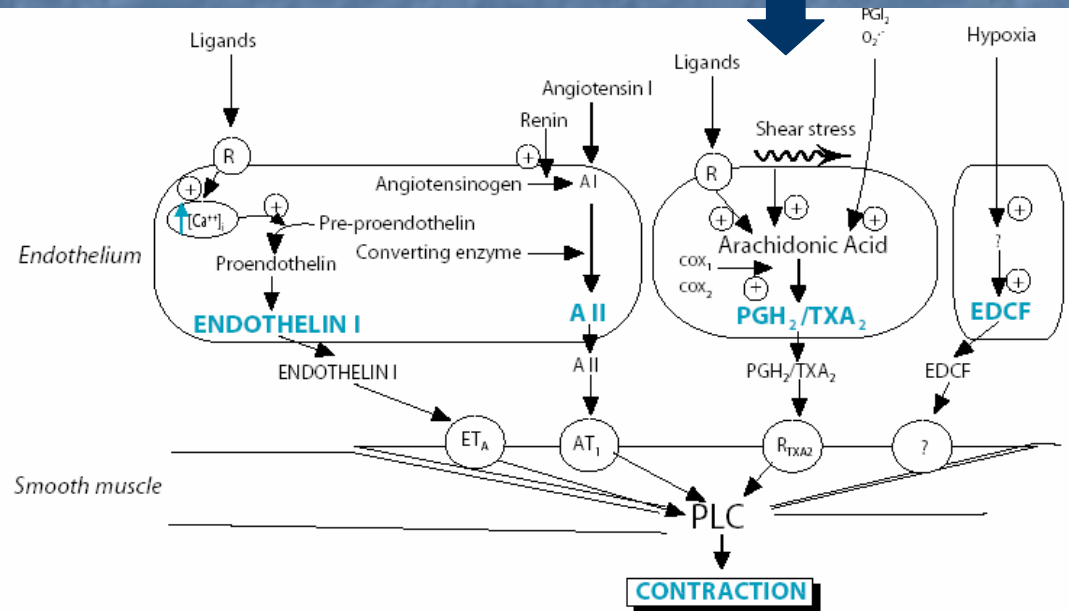
Региональные факторы регуляции АД



Вазодилатация



Вазоконстрикция





Hypertension Candidate Genes

Welcome to the on-line resource for Hypertension Candidate Genes. Below is a list of 150 candidate genes. Each gene is clickable and will open to a new web page showing all available data collected. Available data includes map location and genetic interval, OMIM number, sequence data, and expression data for each gene. If you would like a tab-delimited text file of this information, click [here](#).

List By Gene Names | List By Gene Symbols | GIST

Genes sorted by functional class

Apolipoproteins | Channels and Transporters | Cytoskeletal and Adhesion Molecules | Endothelins
 Fat and Lipid Regulation | Glucose Regulation | Growth Factors and Hormones
 Hypothalamus-Pituitary Axis | Intracellular Messengers | Kallikrein-Kinin pathway | Natriuretic Peptides
 Renin-Angiotensin-Aldosterone pathway | Steroids | Sympathetic Nervous System
 Thromboxanes and Prostaglandins | Miscellaneous

Apolipoproteins

Apolipoprotein A1 (APOA1)
 Apolipoprotein A2 (APOA2)
 Apolipoprotein C2 (APOC2)
 Apolipoprotein C3 (APOC3)
 Apolipoprotein C4 (APOC4)
 Apolipoprotein E (APOE)
 Apolipoprotein E Receptor 2 (APOER2/LRP8)

Channels / Transporters

Aquaporin 2 (AQP2)
 ATP-Sensitive K Channel (KCNJ6)
 Cardiac L type Calcium Channel, alpha-1 (CACNA1C/CACNL1A1)
 Chloride Bicarb. Exchanger 1 (AE1/SLC4A1)
 Chloride Bicarb. Exchanger 2 (AE2/SLC4A2)
 Chloride Bicarb. Exchanger 3 (AE3/SLC4A3)
 Chloride Channel, kidney, A (CLCNKA)
 Chloride Channel, kidney, B (CLCNKB)
 Glucose Transporter 1 (GLUT1/SLC2A1)
 Glucose Transporter 2 (GLUT2/SLC2A2)
 Glucose Transporter 3 (GLUT3/SLC2A3)
 Glucose Transporter 4 (GLUT4/SLC2A4)
 Potassium-chloride cotransporter (hKCC1)

file:///F:/Hypertension%20Candidate%20Genes.htm

21.11.2004

Sodium Bicarbonate Cotransporter (NBC/HNBC1)
 Sodium Calcium Exchanger 1 (NCX1/SLC8A1)
 Sodium Calcium Exchanger 2 (NCX2/SLC8A2)
 Sodium Channel (SCNN1A)
 Sodium Channel (SCNN1B)
 Sodium Channel (SCNN1D)
 Sodium Channel (SCNN1G)
 Sodium Channel - Heart (SCN5A)
 Sodium Chloride Cotransporter (SLC12A3/HTSC)
 Sodium Hydrogen Exchanger 1 (NHE1/SLC9A1)
 Sodium Hydrogen Exchanger 2 (NHE2/SLC9A2)
 Sodium Hydrogen Exchanger 5 (NHE5/SLC9A5)
 Sodium Potassium ATPase alpha1 (ATP1A1)
 Sodium-Glucose Cotransporter, renal (SGLT2/SLC5A2)
 B-cell K/ATP Channel (BIR/KCNJ11)

Cytoskeletal / Adhesion Molecules

Adducin - alpha (ADD1)
 Adducin - beta (ADD2)
 Adducin - gamma (ADD3)
 E-Selectin (ELAM/SELE)
 Intercellular Adhesion Molecule 1 (ICAM1)
 Intercellular Adhesion Molecule 2 (ICAM2)
 Intercellular Adhesion Molecule 3 (ICAM3)

Endothelins

Endothelin 1 (EDN1/ET1)
 Endothelin 2 (EDN2/ET2)
 Endothelin 3 (EDN3/ET3)
 Endothelin Convert. Enzyme1 (ECE1)
 Endothelin Receptor Type A (EDNRA)
 Endothelin Receptor Type B (EDNRB)

Fat and Lipid Regulation

Bombesin-like Receptor 3 (BRS3)
 LDL Receptor (LDLR)
 Leptin (LEP/OB)
 Leptin Receptor (LEPR)
 Lipoprotein Lipase (LPL)
 Uncoupling Protein 3 (UCP3)

Glucose Regulation

Aldose Reductase (ALDR1)
 Amylin (IAPP)
 Gastric Inhibitory Polypeptide Receptor (GIPR)
 Glucagon (GCG)
 Glucagon Receptor (GCGR)
 Glucagon-like Peptide 1 Receptor (GLP1R)

file:///F:/Hypertension%20Candidate%20Genes.htm

21.11.2004

Frustrating Search for Hypertension Genes Continues

By
Edward R.
Winstead

July 11, 2003

Despite years of searching, researchers have yet to discover a human gene that controls blood pressure or contributes to hypertension. The worldwide effort has generated a list of chromosome regions that may contain these genes but no specific genes or variants.

Now, a region of chromosome 6 has been added to the list. British researchers discovered the region by "scanning" the genomes of 2,000 pairs of siblings from families with severe hypertension for DNA they had in common. In addition to chromosome 6, several other areas were implicated, but the statistical evidence was modest.

The next step is to screen the regions for gene candidates, but the researchers acknowledge that whatever they find will not be the whole story on the genetics of hypertension.

Indeed, the story seems to be that many genes (or variants) contribute in small degrees to inherited differences in blood pressure, and that different genes may confer risk for hypertension in different populations.

"We really don't know how many blood pressure genes there are," says Mark J. Caulfield of Queen Mary's School of Medicine, London, who led the new study. "We hope they number 5 or 10, but there could be 20 or more."

As with other complex diseases, like schizophrenia and heart disease, pinpointing genetic risk factors for hypertension will not be easy.

Consider a recent analysis of data from genome scans for hypertension genes published in the scientific literature. This "meta-analysis" found that no single chromosome region consistently had large effects on blood pressure or hypertension in all the study populations. It was published in the *American Journal of Hypertension* in February.

The original genome scans were done at medical centers in the United States as part of the National Heart, Lung, and Blood Institute Family Blood Pressure Program (FBPP). The subjects came from different ethnic groups and represented the racial diversity of the U.S. population.

The new research is part of the Medical Research Council's British Genetics of HyperTension (BRIGHT) study. The findings appear in *The Lancet* along with a commentary entitled "Where are all the blood-pressure genes?"

Citing the challenges of finding genes underlying common diseases, the commentary's author, Stephen B. Harrap of the University of Melbourne, Australia, suggests that it simply may not be possible to discover gene variants for hypertension that would be valuable in the diagnosis of the disease.

Rather than search for every gene variant underlying hypertension, he argues, the field needs to "search for molecular clues to the common physiological mechanisms underlying disease."

The study in *The Lancet* may ultimately lead to new risk factors for severe hypertension. None of the chromosome regions in the study was associated with the disease in humans until now. But part of chromosome 2 corresponds to a region of the rat genome associated with elevated blood pressure.

"We are encouraged by this because it means we may have found a region that actually contains genes" for hypertension, says Caulfield. "We're making some progress, though perhaps not as much as we would like."

"If finding the genes were easy, we would not have undertaken such large studies."

...

Caulfield, M. *et al.* Genome-wide mapping of human loci for essential hypertension. *Lancet* **361**, 2118-2123 (June 21, 2003).

Harrap, S.B. Where are all the blood-pressure genes? *Lancet* **361**, 2149-2151 (June 21, 2003).

Province, M.A. *et al.* A meta-analysis of genome-wide linkage scans for hypertension: The National Heart, Lung and Blood Institute Family Blood Pressure Program. *American Journal of Hypertension* **16**, 144-147 (February 2003).

**'Where are all the
blood-pressure
genes?'**

Indeed, the story seems to be that many genes (or variants) contribute in small degrees to inherited differences in blood pressure, and that different genes may confer risk for hypertension in different populations



«ПОИСК ГЕНОВ ОТВЕТСТВЕННЫХ
ЗА ПАТОГЕНЕЗ
МУЛЬТИФАКТОРИАЛЬНЫХ
БОЛЕЗНЕЙ ИНОГДА ПОДОБЕН
ПОИСКУ ИГОЛКИ В СТОГЕ СЕНА»

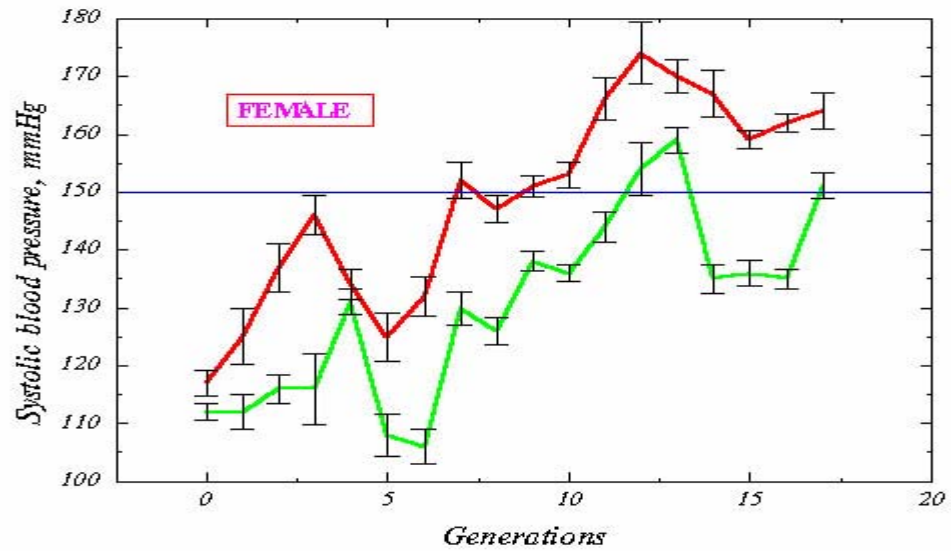
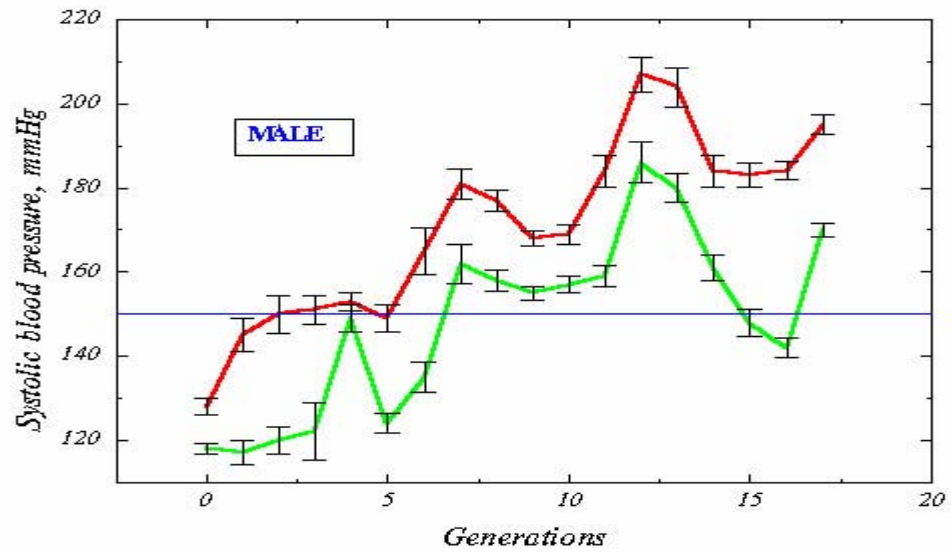
TABLE 1. *Rodent strains selectively bred for blood pressure*

| Strain | Lines | Original Stock | Year First Reported (Reference No.) |
|---|---------|--|--|
| <i>Rat</i> | | | |
| New Zealand (Dunedin): genetically hypertensive (GH) rats | H, C | Wistar derived | Smirk and Hall, 1958 (234) |
| USA (Brookhaven): Dahl salt-sensitive (S) and Dahl salt-resistant (R) rats | H, L | Sprague-Dawley | Dahl et al., 1962 (30, 31) |
| Japan (Kyoto): spontaneously hypertensive rats (SHR) | H | Wistar derived | Okamoto and Aoki, 1963 (175) |
| Japan (Kyoto): spontaneously hypertensive rats-stroke prone (SHRSP) | H | Wistar derived | Okamoto et al., 1974 (176) |
| Israel (Jerusalem): DOCA salt-sensitive (SBH) and resistant (SBN) rats | H, L | Unknown | Ben-Ishay et al., 1972 (7) |
| France: Lyon hypertensive (LH), Lyon normotensive (LN), and Lyon low blood pressure (LL) rats | H, C, L | Sprague-Dawley | Dupont et al., 1973 (53) |
| Italy: Milan hypertensive strain (MHS) and Milan normotensive (MNS) rats | H, C | Wistar | Bianchi et al., 1974 (8) |
| The Netherlands (Utrecht): fawn-hooded hypertensive (FHH) and fawn-hooded low blood pressure (FHL) rats | H, L | Greman brown × white Lashley | Kuijpers and Gruys, 1984 (124) |
| Russia (Novosibirsk): inherited stress-induced arterial hypertension (ISIAH) rats | H | Wistar derived | Markel, 1985 (158) |
| Czech Republic: Prague hypertensive rat (PHR), Prague normotensive rat (PNR) | H, L | Wistar derived | Heller et al., 1993 (80) |
| <i>Mouse</i> | | | |
| USA (University of Kansas) | H, C, L | Eight-way cross of inbred strains | Schlager, 1974 (217) |
| <i>Hamster</i> | | | |
| Canada (Halls Harbor) | H | Cardiomyopathic × gold Syrian hamsters | Thomas et al., 1997 (252) |

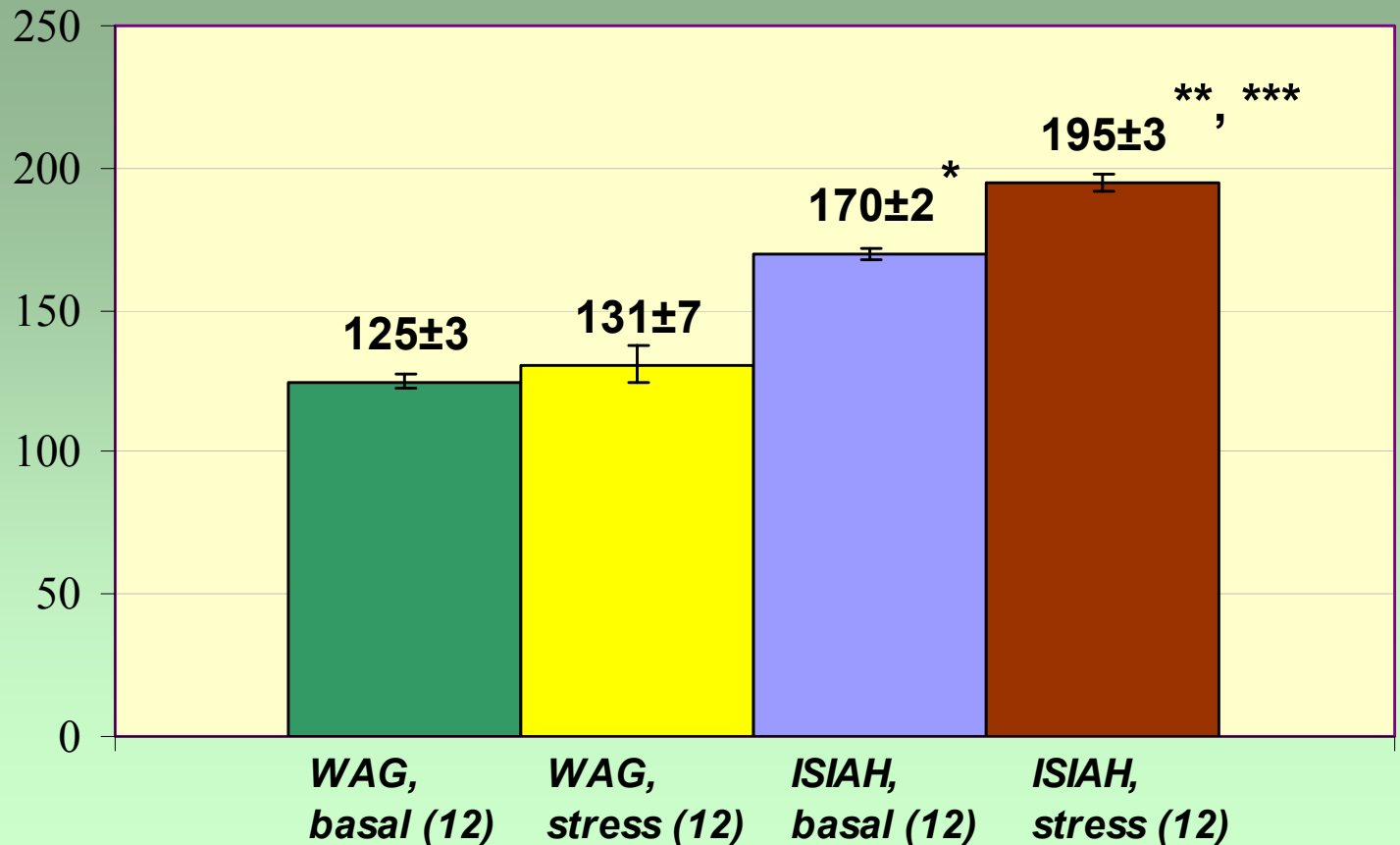
Kinds of lines developed in each model were as follows: H, line selected for high blood pressure; C, control line, unselected, random bred; L, line selected for low blood pressure.

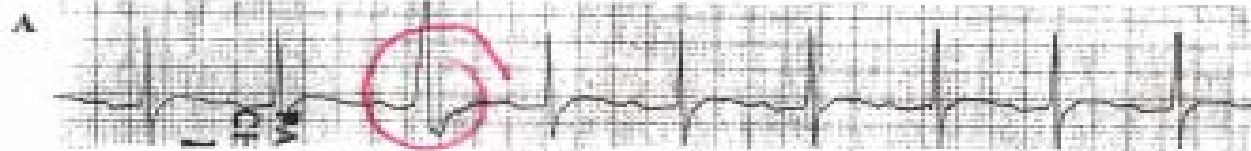
ISIAH rat strain: An experimental model of stress-sensitive arterial **hypertension**





Basal and stress-induced BP levels in WAG and ISIAH rat strains





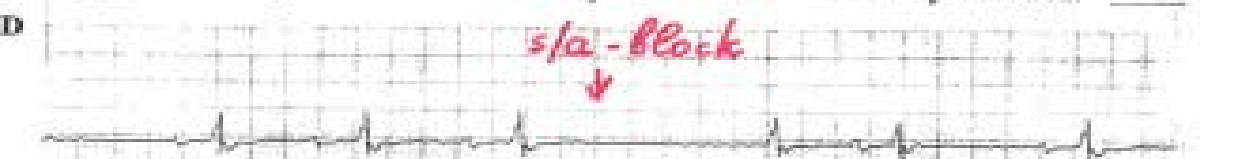
A. Единичная желудочковая экстрасистола.



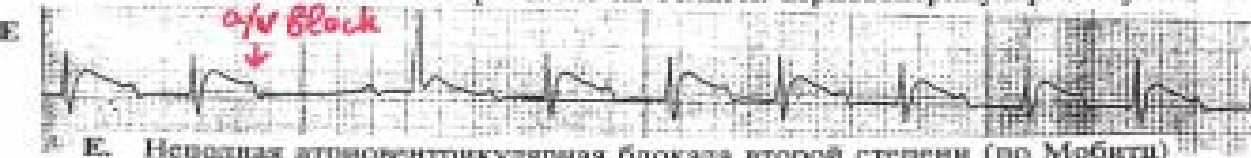
B. Нарушение внутрисердечной проводимости и спаренные желудочковые экстрасистолы.



C. Альтернация комплекса QRS по типу бигеминии, вызванная попеременной блокадой правой и левой ножек пучка Гиса.



D. Синусовая аритмия. Синоаурикулярный блок и единичное "выскакивающее" сокращение из области атриоventрикулярного узла.



E. Неполная атриоventрикулярная блокада второй степени (по Мобитц) и единичное "выскакивающее" сокращение из области желудочков.



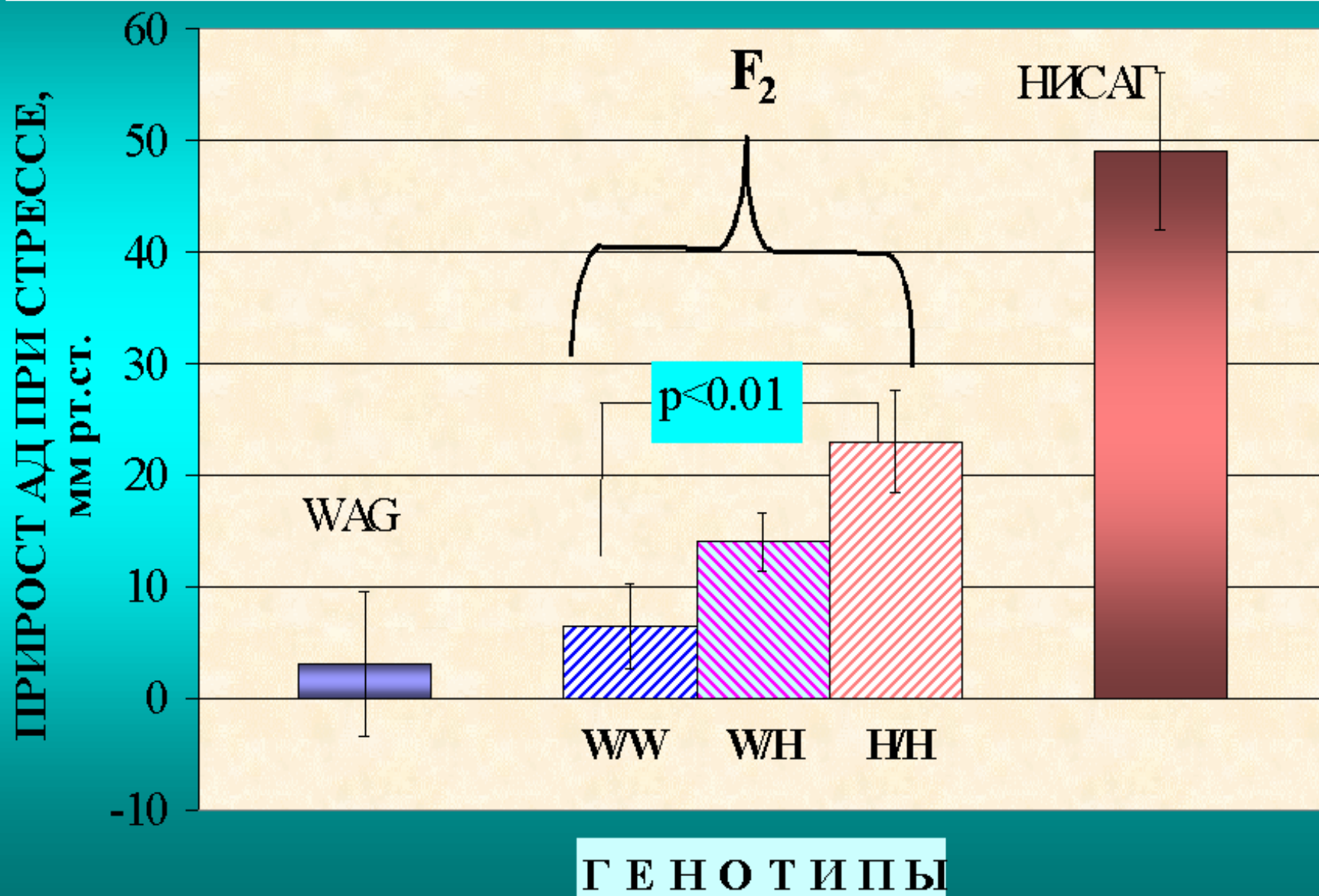
F. Неполная атриоventрикулярная блокада второй степени (периоды Венкебаха).

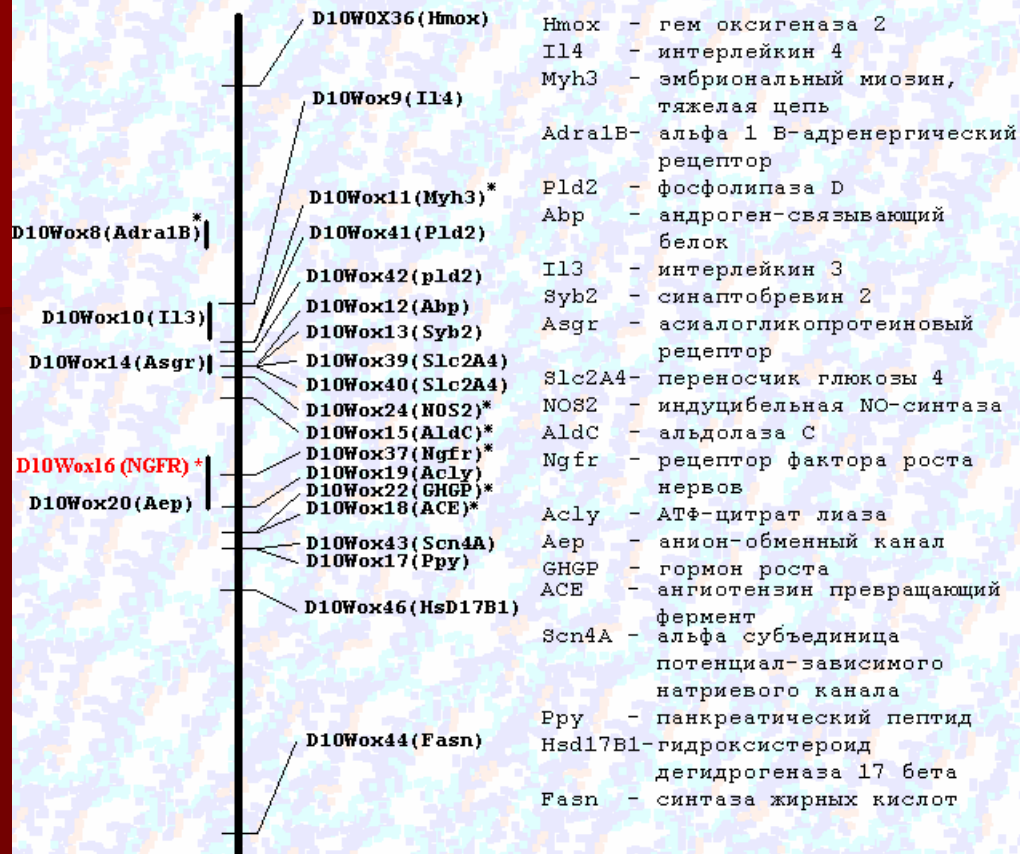


G. Атрио/ventрикулярная диссоциация на фоне низкого предсердного ритма.

Нарушения
сердечного
ритма у крыс
линии НИСАГ
по данным
ЭКГ исследо-
вания

Локус *NGFR* и величина прироста АД при стрессе





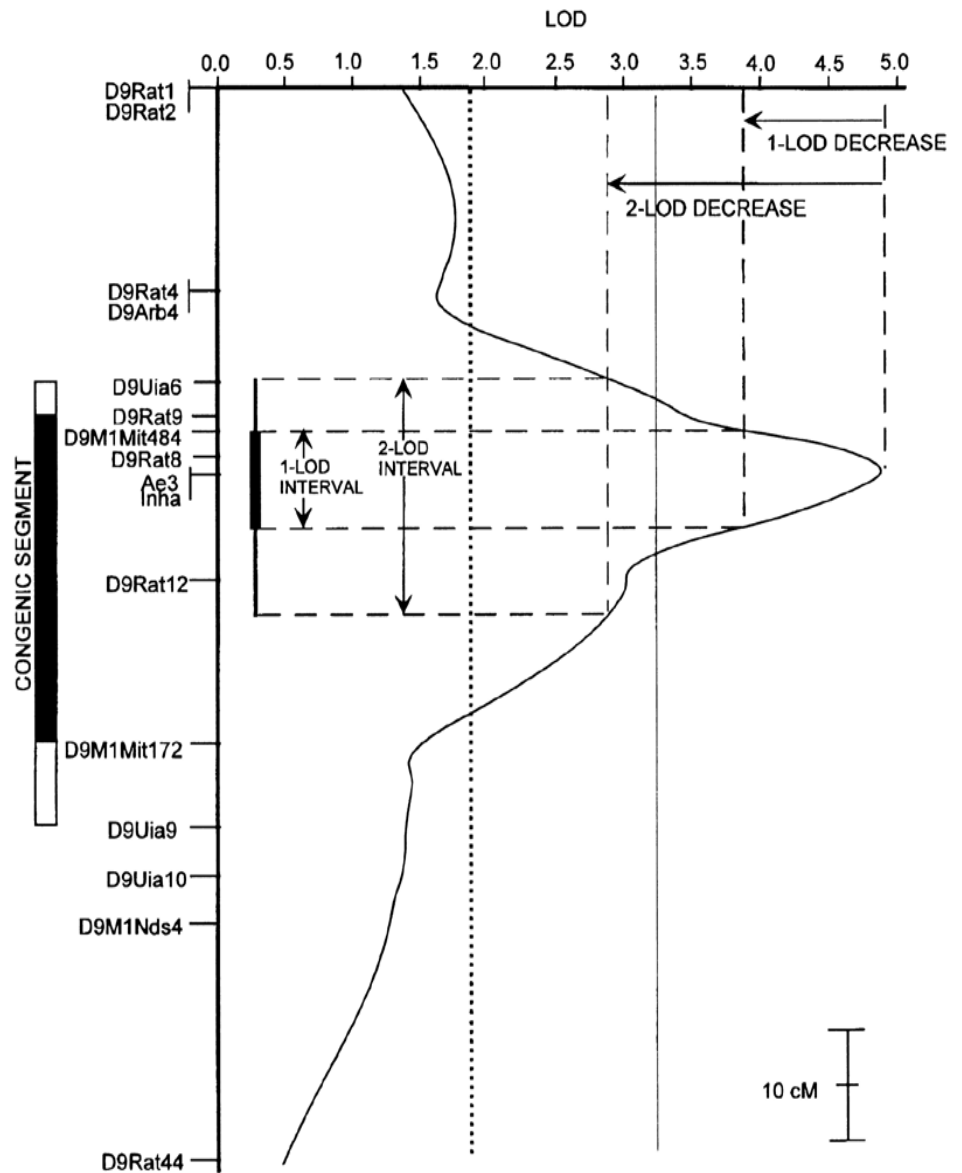
Звездочкой (*) обозначены локусы генов-кандидатов, для которых была найдена ассоциация с гипертонией у человека или на какой-либо из моделей экспериментальных животных

Карта сцепления 10-й хромосомы крысы

From: John P. Rapp, 2000

| CHROMOSOME 9 | Blood Pressure by Genotype | | | Oneway ANOVA P | Blood Pressure Effect = SS - RR (mm Hg) |
|---|----------------------------|----------------|---------------|----------------|---|
| | SS | SR | RR | | |
| D9Rat1 D9Rat2 | 183 (n=64) | 177 (n=95) | 170 (n=50) | 0.0690 | 13 |
| D9Rat4 D9Arb4 | 182 (n=61) | 180 (n=105) | 164 (n=40) | 0.0051 | 18 |
| D9Uia6 D9Rat9 D9M1Mit484 D9Rat8 Ae3 Inha | 186 (n=71) | 179 (n=114) | 160 (n=47) | <0.0001 | 26 |
| D9Rat12 | 181 (n=54) | 179 (n=92) | 160 (n=36) | 0.0006 | 21 |
| D9M1Mit172 | 176 (n=61) | 180 (n=97) | 164 (n=55) | 0.0505 | 12 |
| D9Uia9 D9Uia10 D9M1Nds4 | 181 (n=21) | 177 (n=65) | 176 (n=19) | 0.2552 | 5 |
| D9Rat44 | 179 (n=57) | 179 (n=105) | 171 (n=47) | 0.2659 | 8 |

10 cM



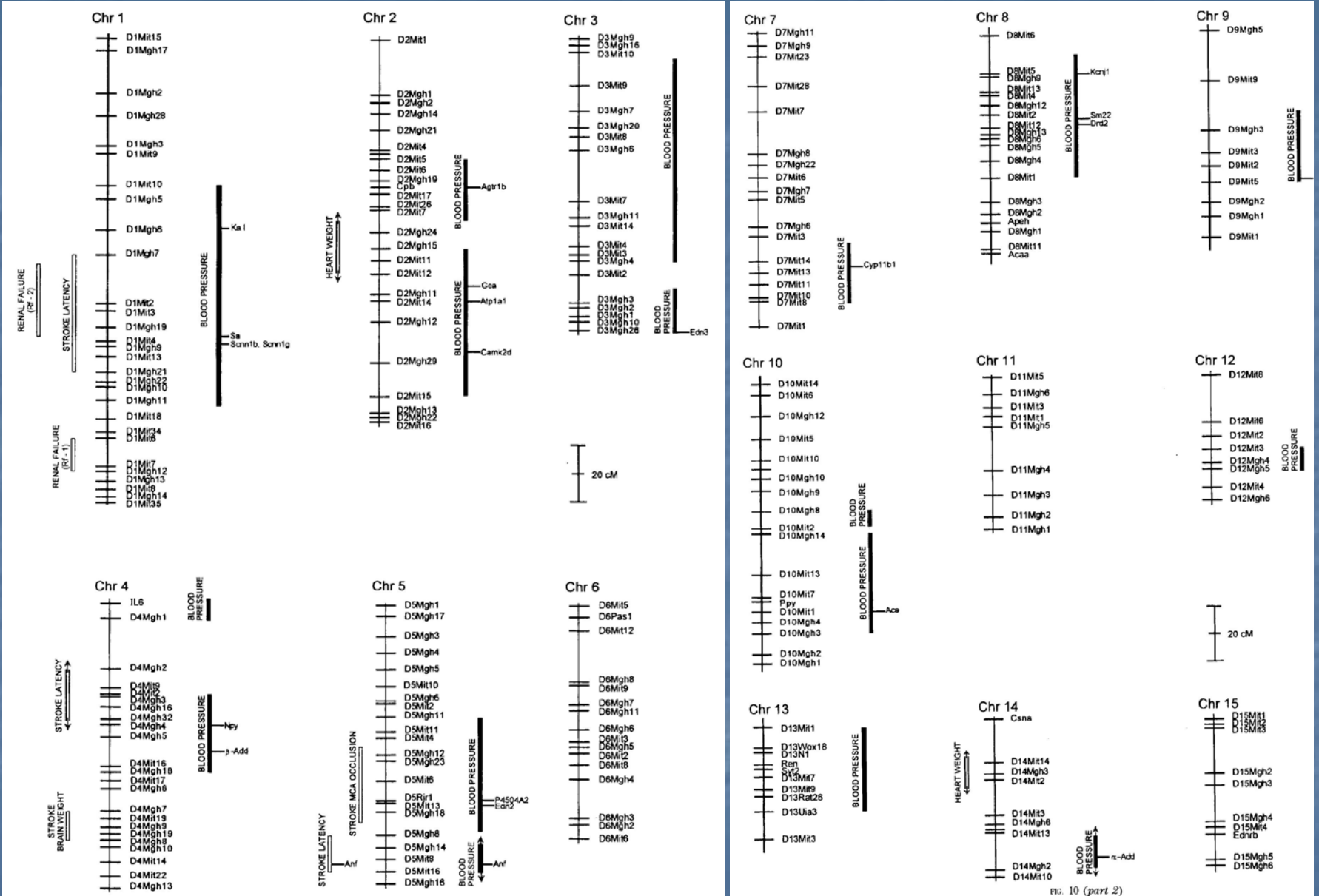
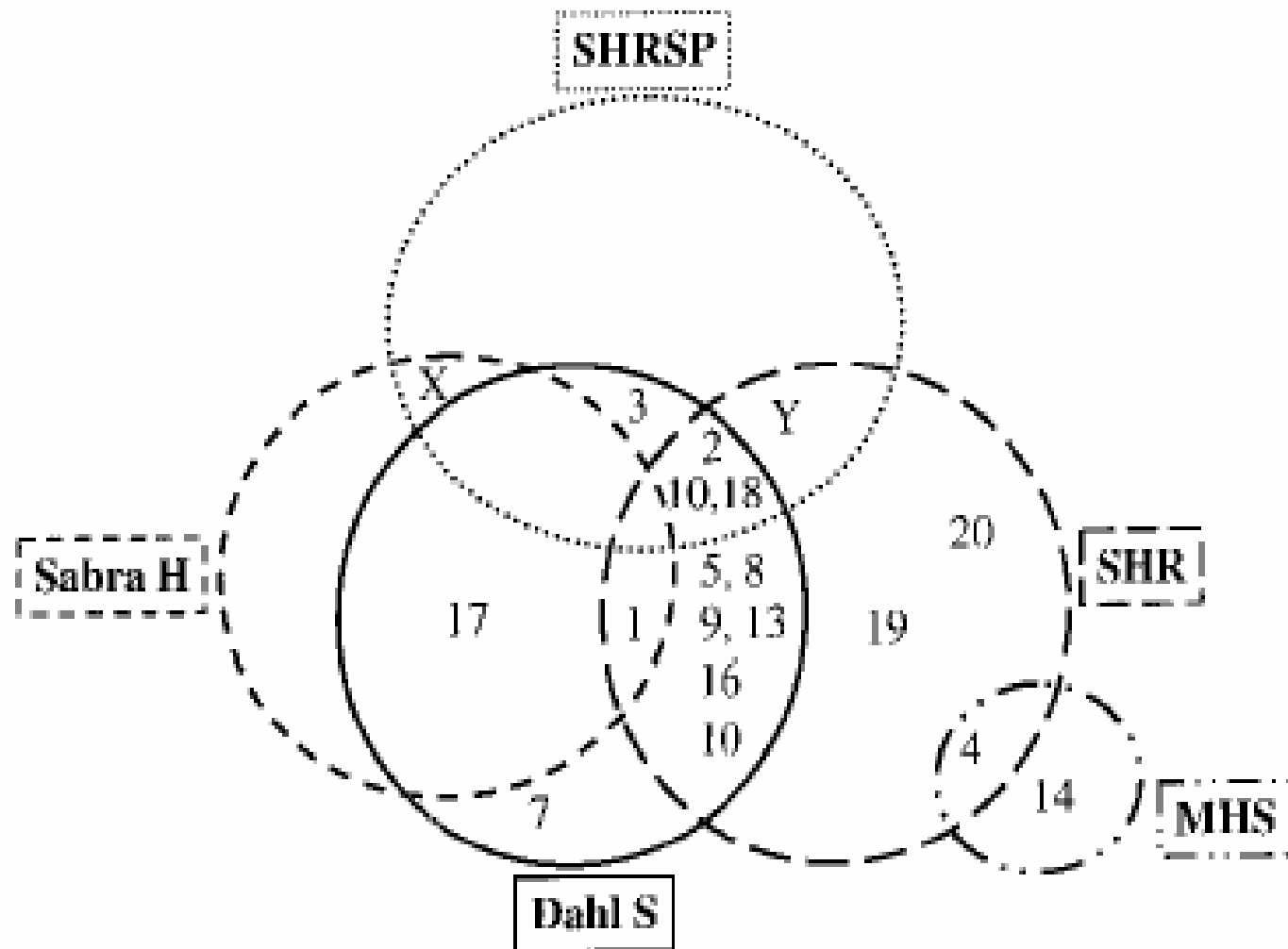


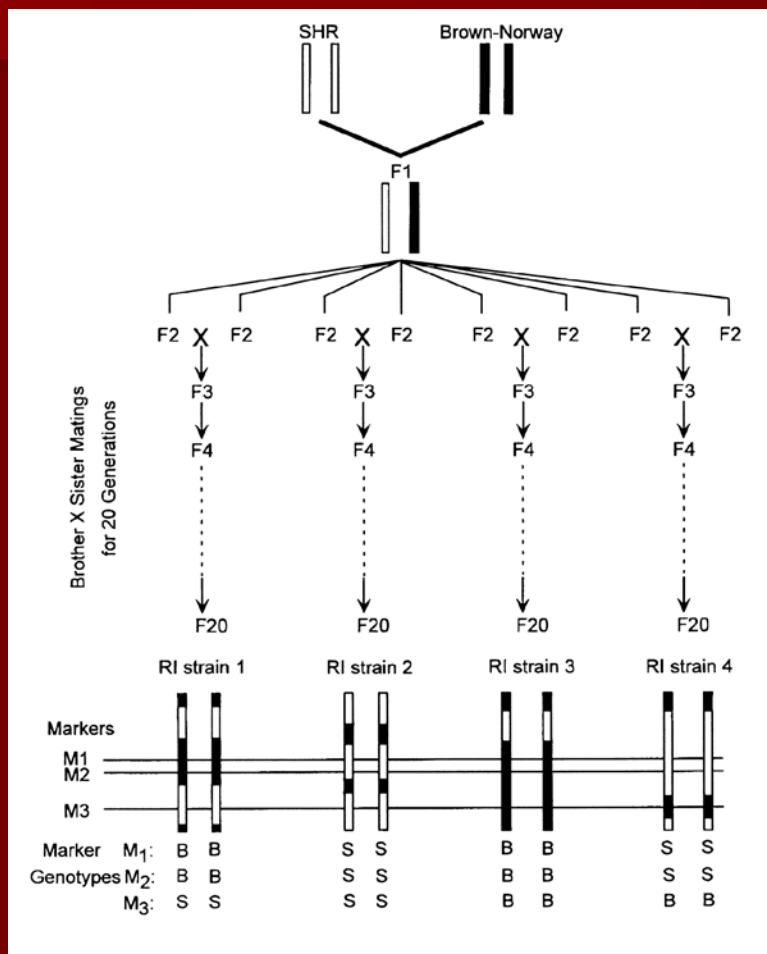
FIG. 10 (part 2)

Common or reproducible blood pressure QTL identified in several independent experiments

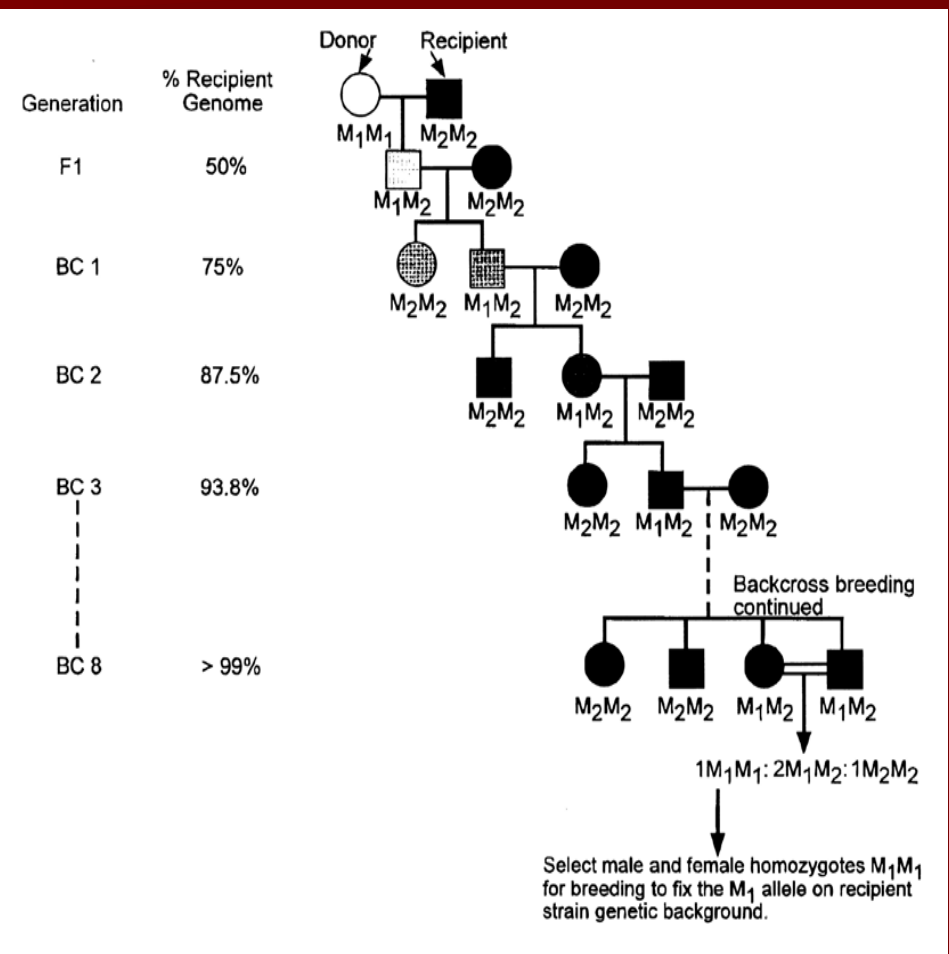


From: John P. Rapp, 2000

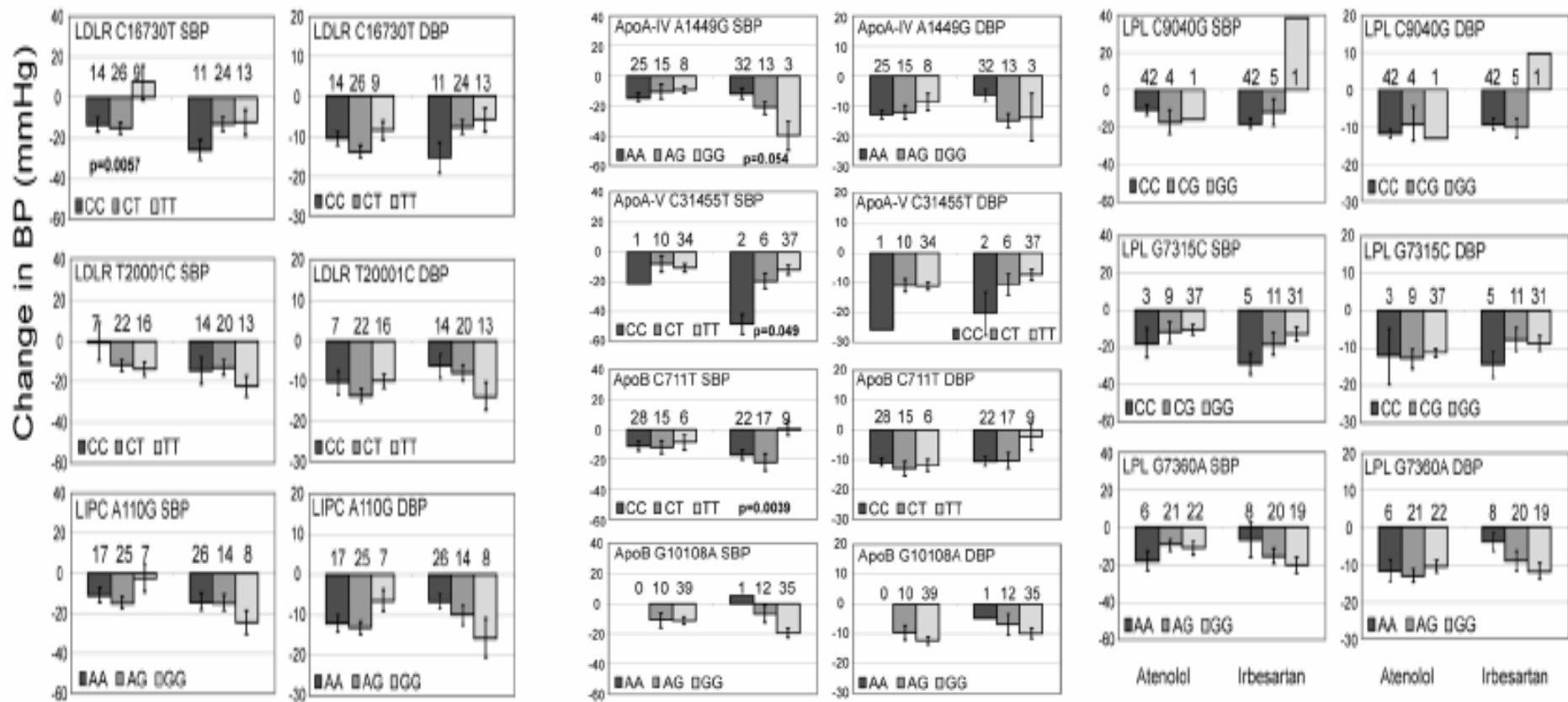
Рекомбинантные линии



Конгенные линии

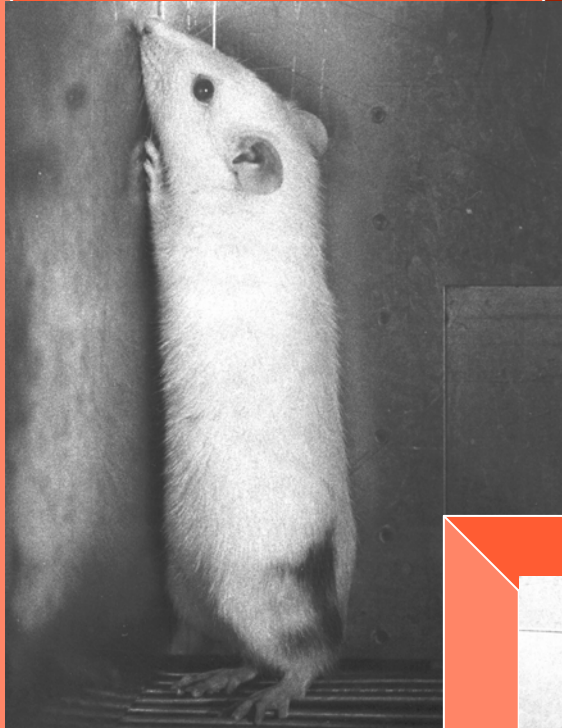


Effect of SNP genotype on the change in blood pressure after 12 weeks of treatment for the ten SNPs.



From: BMC Cardiovascular Disorders, 2004

Крысы линии ГК (генетическая каталепсия)



**Автор линии –
В.Г.КОЛПАКОВ
ИЦИГ СО РАН**



АГРЕССИЯ

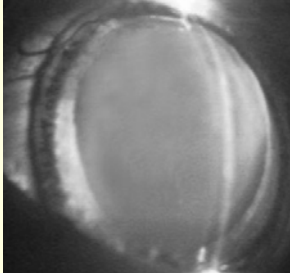
↓

ДЕПРЕССИЯ

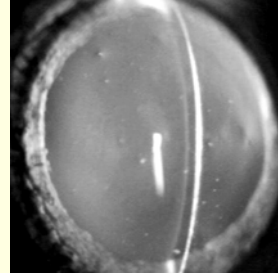


Создание и изучение
экспериментальной
модели депрессии
Д.б.н. Н.Н.Кудрявцева

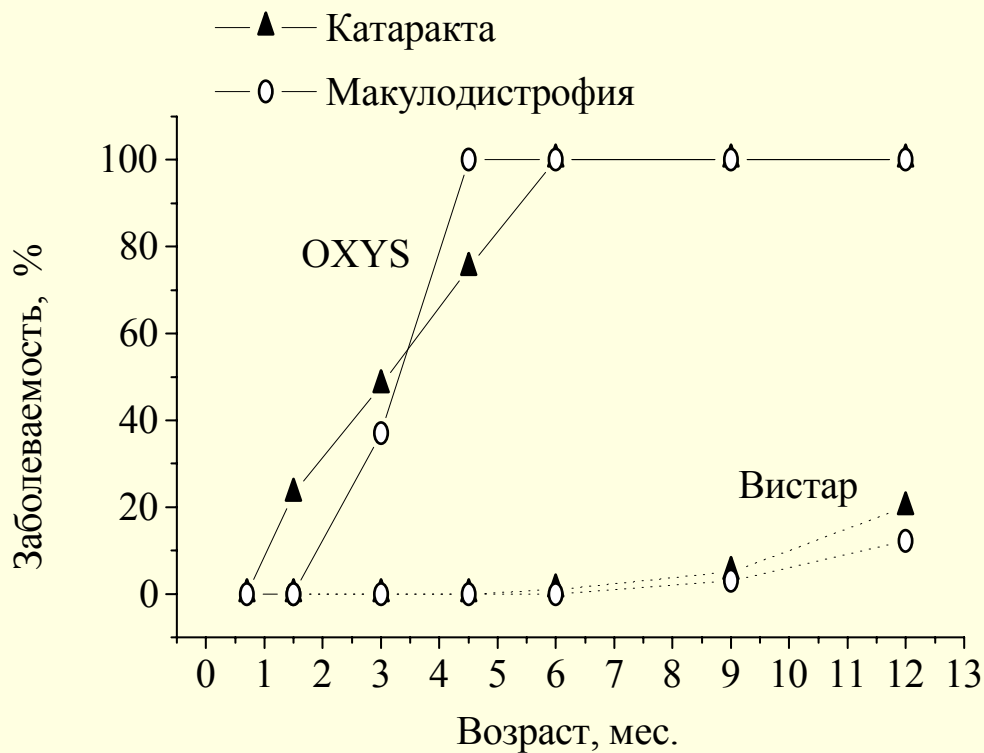




Здоровый хрусталик у крыс Wistar



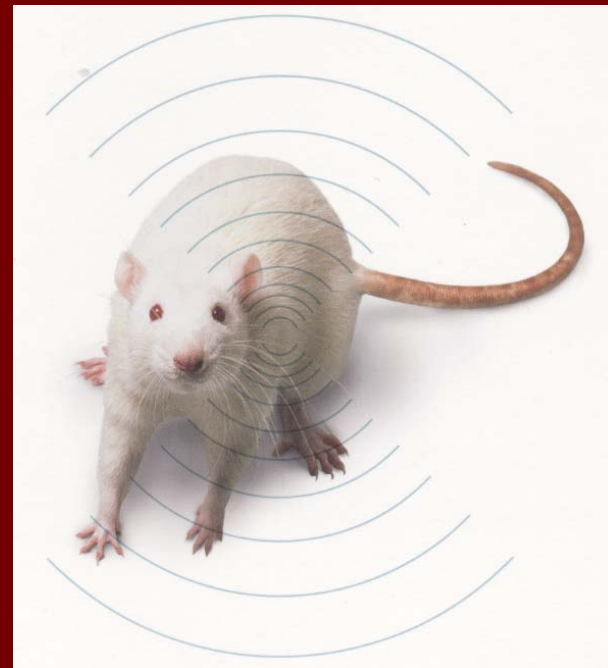
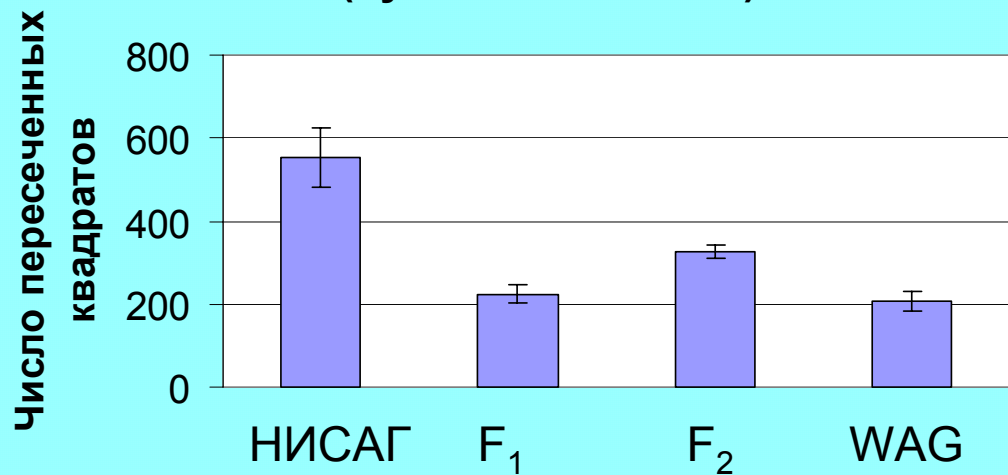
Точечная катаракта хрусталика у крыс OXYS



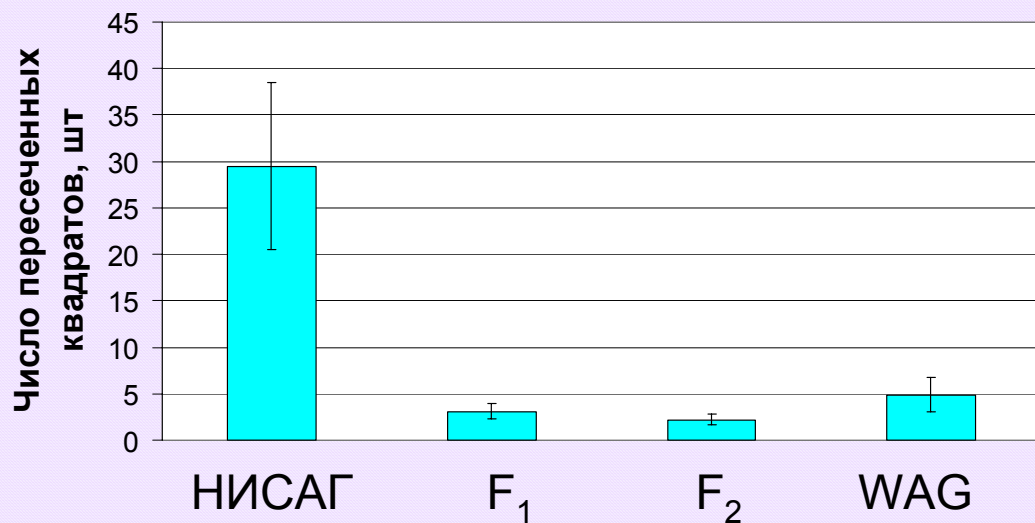
Содержание продуктов перекисного окисления липидов (ПОЛ) в плазме крови крыс OXYS повышено и связано с выраженностью патологических изменений. Проявления раннего старения органа зрения у крыс OXYS обусловлено окислительным стрессом, что подтверждается высокой эффективностью антиоксидантов в их профилактике.

2.1.1. Блок 3 (Н.Г. Колосова)

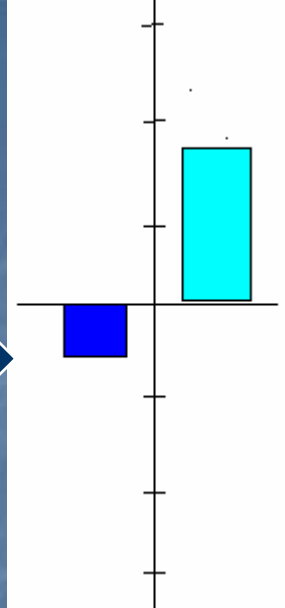
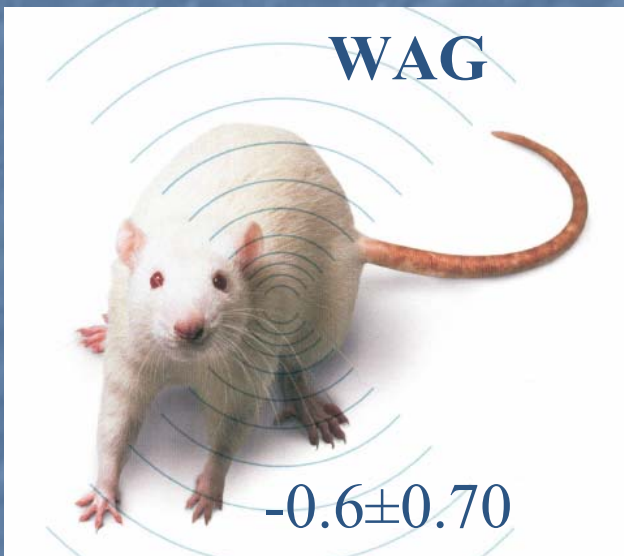
Двигательная активность на периферии площадки "открытого поля" (сумма за 4 теста)



Двигательная активность в центре площадки "открытого поля" (сумма за четыре теста)



Негативная реакция
(страх)



Позитивная реакция
(исследование)

$P < 0.05$



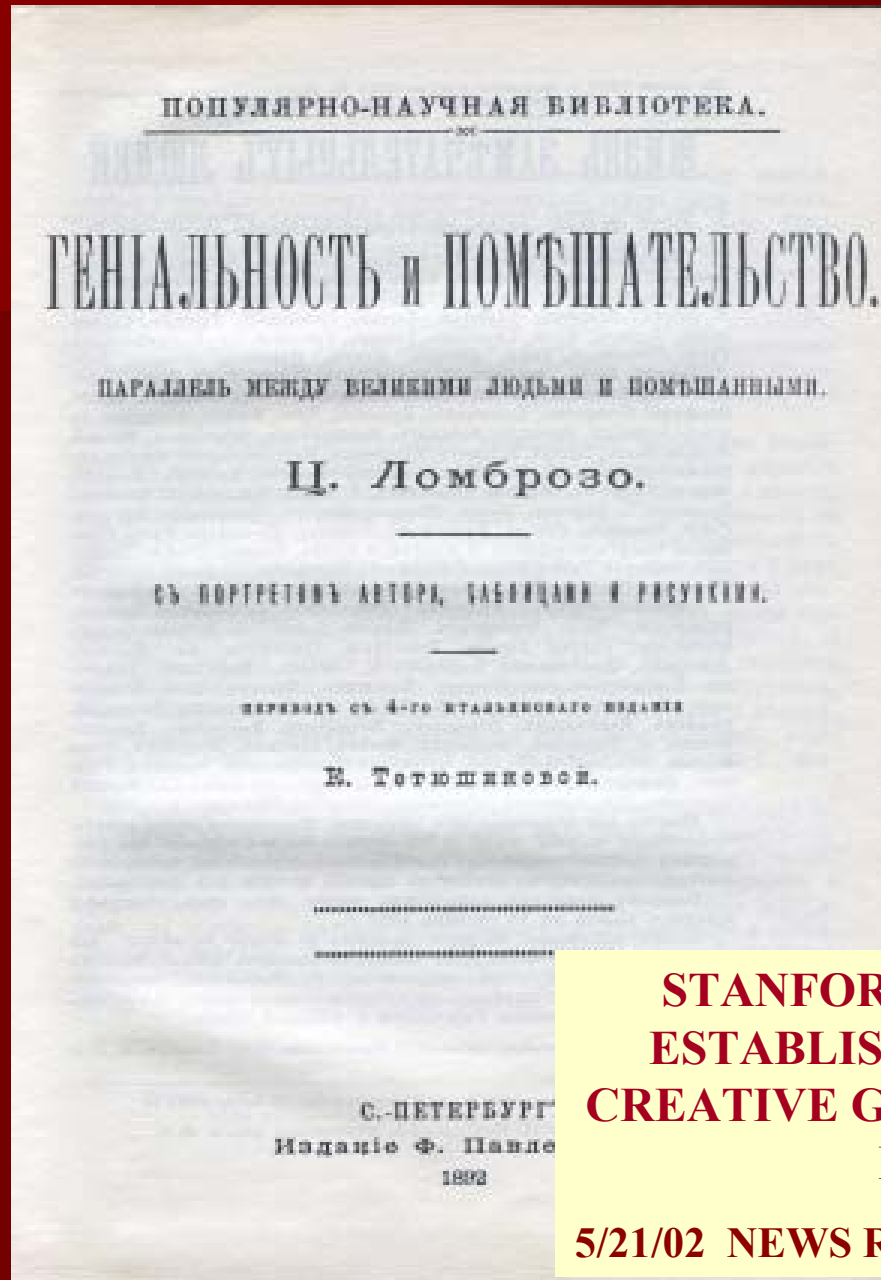
Внезапное появление звукового или
зрительного стимула

**«Не бывает
гения без
примеси
помешатель-
ства»**

Сенека

3 до н.э. — 65

н.э.



**“You don’t
have to be
mad to be a
genius... but
it helps”**

**Dr. Kenneth
Lyen**

2002 г.

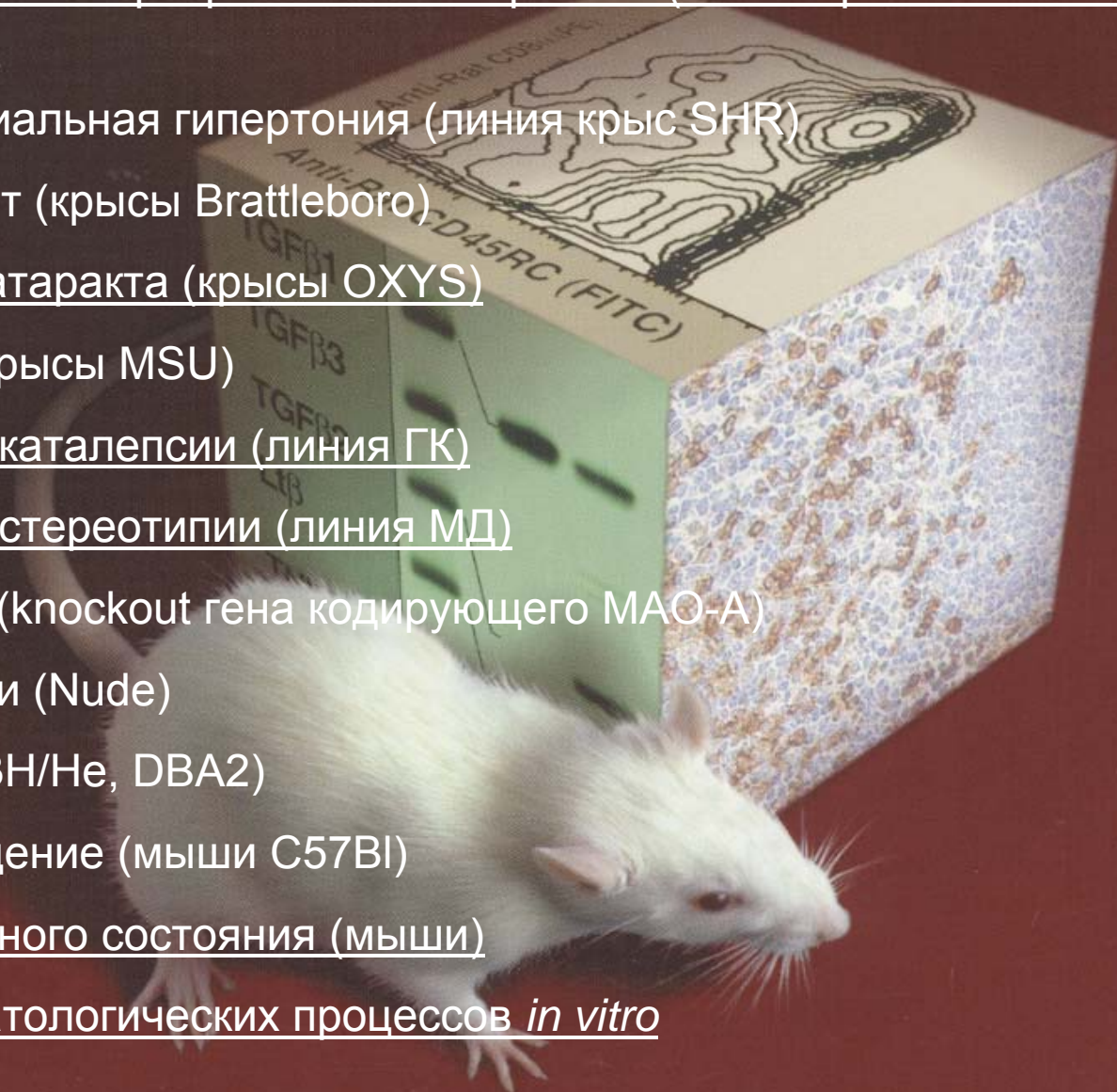
From an Essay

**Is there a link
between genius and
madness?**

**STANFORD RESEARCHERS
ESTABLISH LINK BETWEEN
CREATIVE GENIUS AND MENTAL
ILLNESS**

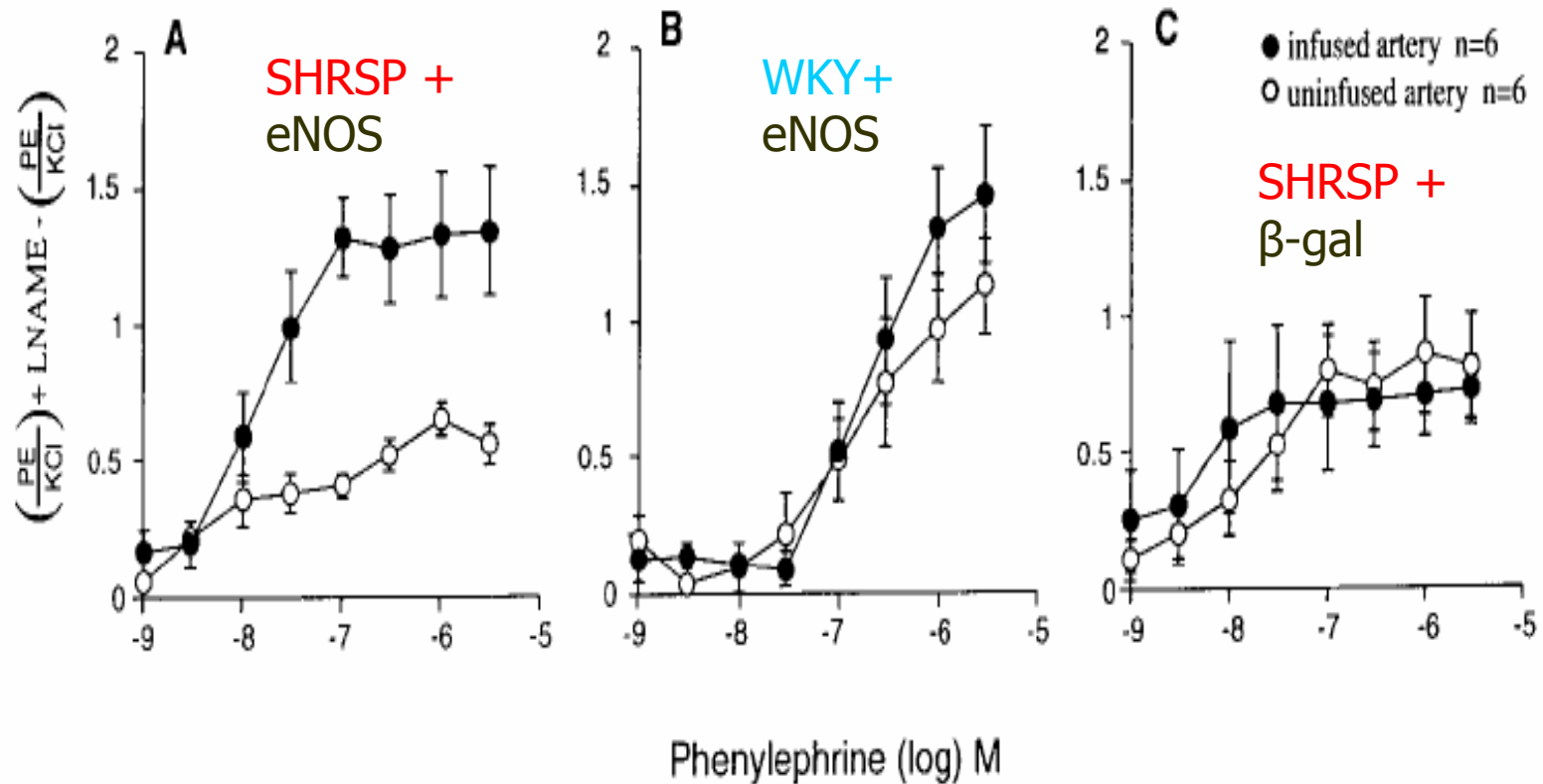
5/21/02 NEWS RELEASE

1. Стресс-чувствительная артериальная гипертония (линия крыс НИСАГ – ISIAH/lcgn)
2. Спонтанная артериальная гипертония (линия крыс SHR)
3. Несахарный диабет (крысы Brattleboro)
4. Наследственная катаракта (крысы OXYS)
5. Микрофтальмия (крысы MSU)
6. Крысы склонные к каталепсии (линия ГК)
7. Крысы склонные к стереотипии (линия МД)
8. Линия мышей Tg8 (knockout гена кодирующего MAO-A)
9. Бестимусные мыши (Nude)
10. Опухоли (мыши C3H/He, DBA2)
11. Агрессивное поведение (мыши C57Bl)
12. Модель депрессивного состояния (мыши)
13. Моделирование патологических процессов *in vitro*



Basal NO availability after infusion of the adenovirus encoding endogenous eNOS cDNA driven by cytomegalovirus promoter

(From Anna F. Dominiczak et al., 2000)



Gene Therapy for Common Acquired Diseases of the Heart The Sirens' Song

Eduardo Marbán, MD, PhD

*Then the queenly Circe spoke in words and address
come first of all to the Sirens, who are enchanters of a
whoever comes their way: and that man who un-
proaches them, and listens to the Sirens singing, has
coming home. You must drive straight on past...*

Homer

The prospect of using genes as therapy presents myriad opportunities. No physician can appreciate the power of this approach without abandoning the shackles of the limited pharmacologic device repertoire and focus instead on reengineering culprit tissue by somatic gene transfer? The genomic information leaves us with an embarrassment of riches in terms of potential therapeutic agents. We can mine nature's own genes expressed in their usual sites. Genes can readily be tailored to exhibit special properties found in nature, altering the function of their protein products for specific ends. Alternatively, wild-type genes can be expressed in tissues where they are normally silent.

See p 1578

In this issue, Weig et al¹ describe a clever application of the latter approach: V2 vasopressin receptor genes, usually expressed only in kidney, were delivered to myocardium packaged in recombinant adenoviruses. Expression of these adenyl cyclase-activating receptors in the myocardium converted the basal negative inotropic response to infused vasopressin into a positive one. Because vasopressin levels are elevated in heart failure, a situation in which β -receptors are uncoupled from cyclase,² ectopic expression of V2 receptors would logically be predicted to recruit cAMP-mediated contractility in failing myocardium while bypassing the desensitized β -adrenergic pathway. The work has conceptual beauty. It also presents 2 important methodological innovations: first, the use of *in situ* viral transduction followed by myocyte isolation to enable careful phenotypic characterization of genetically altered cells; second, the adaptation for *in vivo* use of previously described strategies for optimizing transcor-

The opinions expressed in this editorial are not necessarily those of the editors or of the American Heart Association.

From the Institute of Molecular Cardiobiology, Johns Hopkins University School of Medicine, Baltimore, MD.

Correspondence to: Eduardo Marbán, MD, PhD, Institute of Molecular

...faced with the Sirens. The song we hear is all the more seductive because it originates deep within ourselves. Nevertheless, our roles as physicians and as guardians of the public trust dictate that we do all we can to resist the temptation to jump into human studies until we have gathered overwhelming evidence for both safety and efficacy. For the

long-term studies with the same genes (and the same fundamental properties of excitation-contraction coupling in mice as humans) that conclusions so far reached in mice. Nevertheless, mice can help us evaluate candidate genes, and larger animals can provide a platform for pushing the envelope.

No matter how smart we think we are, our animal studies may be, when it comes to real therapy with well-reasoned but ultimately unproven interventions, notably with our patients⁹ and with phosphodiesterase inhibitors.⁵ In such cases, at the very least, therapy has the potential danger of causing dosing difficulties. Such concerns are particularly acute in the application of gene therapy, such as hemophilia; the culprit gene, even if expressed, may suffice to correct the defect.

Gene therapy differs fundamentally in its philosophy. Here the goal is not to correct a well-defined genetic disorder; instead, genes are used to manipulate the very workings of the diseased organ, in ways that may have no relationship to underlying changes

Sirens





WHL · NEWSLETTER

News from the World Hypertension League (WHL).
A division of the International Society of Hypertension, and in official relations with the World Health Organization.

No. 99, February 2005

Editorial

Korotkov sounds – what do we know about this 100 year old discovery?



Nikolai Sergeievich Korotkov (1874–1920)

This year, 2005, is the 100th anniversary of one of the most famous discoveries in the field of hypertension: the auscultative method of blood pressure measurement. The method, named after the Russian physician and scientist Nikolai Sergeievich Korotkov (1874–1920), has now been widely used for one century. Unfortunately, while the method itself and the name of its inventor are world famous, much less is known about the history of this outstanding discovery and the real role of Korotkov in the interpretation of the discovered sounds and the implementation of the new method into clinical practice.

Korotkov was only 31 years of age when he made a short presentation at the Scientific Meeting of the Military Hospital of the Imperial Military Academy about an easy, non-invasive method of blood pressure measurement. This was on November 5, 1905. The brief title of his presentation was "Concerning the problem of the methods for investigating blood pressure". He described the following sequence of sounds: first sound, ten murmurs, loud sounds, decreasing sounds and complete disappearance. These observations are now classified as different phases of the Korotkov sounds. Later, Russian scientists and investigators all over the world continued to find an explanation for the auscultative phenomenon. Although our modern understanding of the nature of the sounds has changed, the method remains the standard for hypertension diagnosis and management. Even today, 100 years later, devices are being devel-

continued on page 2

WHL News

Report from the 6th International Symposium on Hypertension and Related Diseases from October 15–18, 2004 in Beijing, China

On the occasion of this Symposium, the World Hypertension League (WHL) held a Regional Meeting on the Prevention of Hypertension. It was chaired by Prof. Liu Lisheng, Vice President of the WHL, Prof. Zhou Beifan, and Dr. Claude Lenfant, President of the WHL. There were 8 presentations describing research and public health activities in the respective countries.

Dr. Lenfant summarized 4 studies from the United States which exemplify the significance and effectiveness of programs to reduce excess weight, to encourage physical exercise, and to reduce salt and alcohol consumption to maintain, or reduce blood pressure (BP) to a normal level (i.e. <140/85 mmHg). He noted that to control hypertension we have the choice between pharmacological and preventive interventions: he concluded that preventive interventions are effective and far less costly than pharmacological regimens.

continued on page 2

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| – Calendar | 4 |

WORLD HYPERTENSION DAY May 14, 2005
Initiated by the World Hypertension League

Saturday, May 14, 2005



The objectives of the **World Hypertension League** are to promote the detection, control and prevention of arterial hypertension in populations. Bringing together and stimulating organizations committed to the control of hypertension is the goal and the raison d'être of the WHL.

HYPERTENSION - THE SILENT KILLER

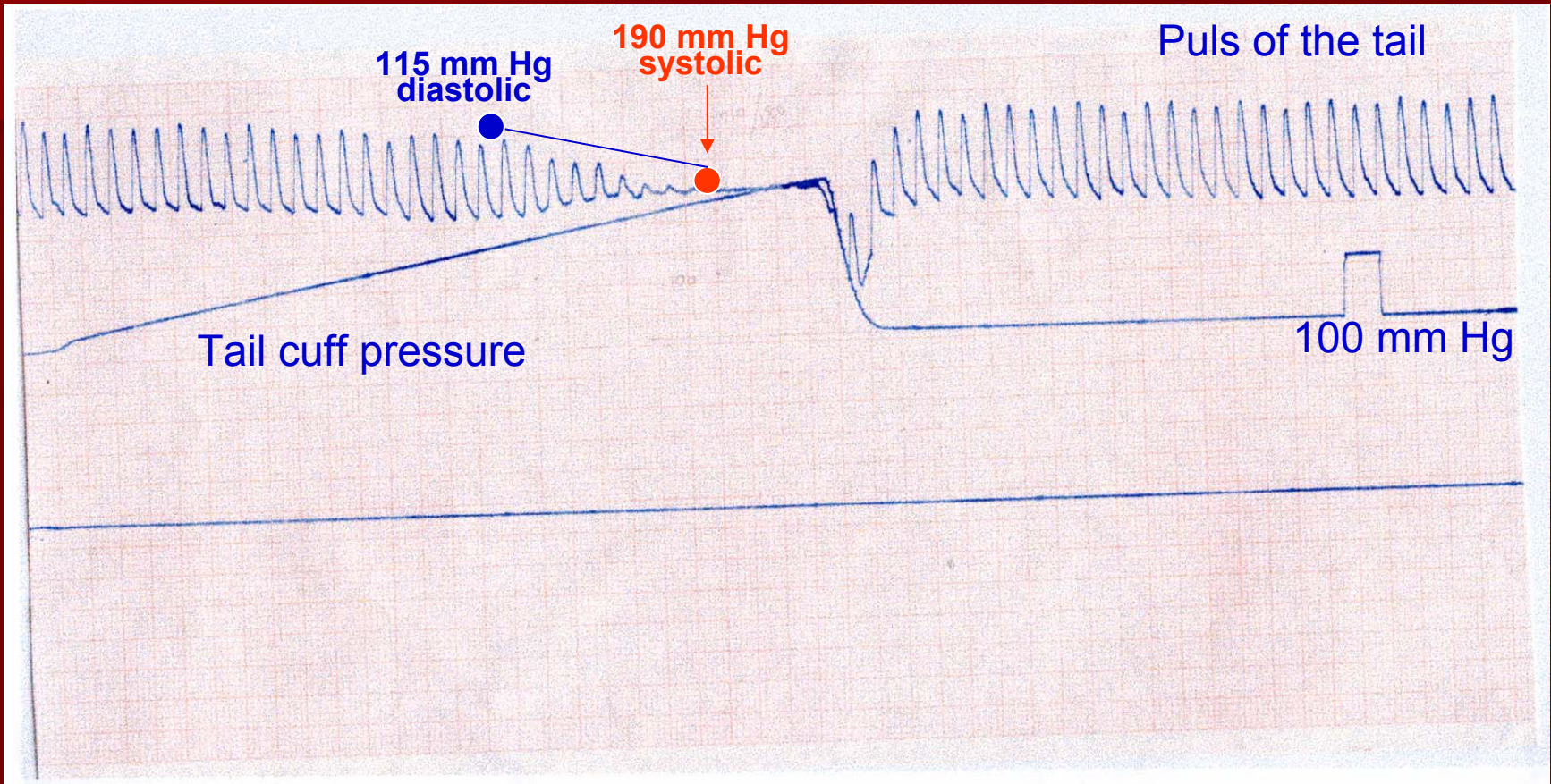
KNOW YOUR BLOOD PRESSURE

GENERAL RULES FOR BLOOD PRESSURE MEASUREMENT

- ▼ Rest for 5 minutes before measurement
- ▼ Refrain from smoking or ingesting caffeine for 30 minutes prior to measurement
- ▼ Be seated with feet flat on the floor, back and arm supported, and arm at heart level
- ▼ Use appropriate sized cuff
- ▼ Use recently validated device
- ▼ An average of two or more readings should be taken, at least two minutes apart

Under special circumstances measuring blood pressure in standing and supine positions may be indicated.

For further information see the WHL website at www.worldhypertensionleague.org



Tail-cuff blood pressure measurement

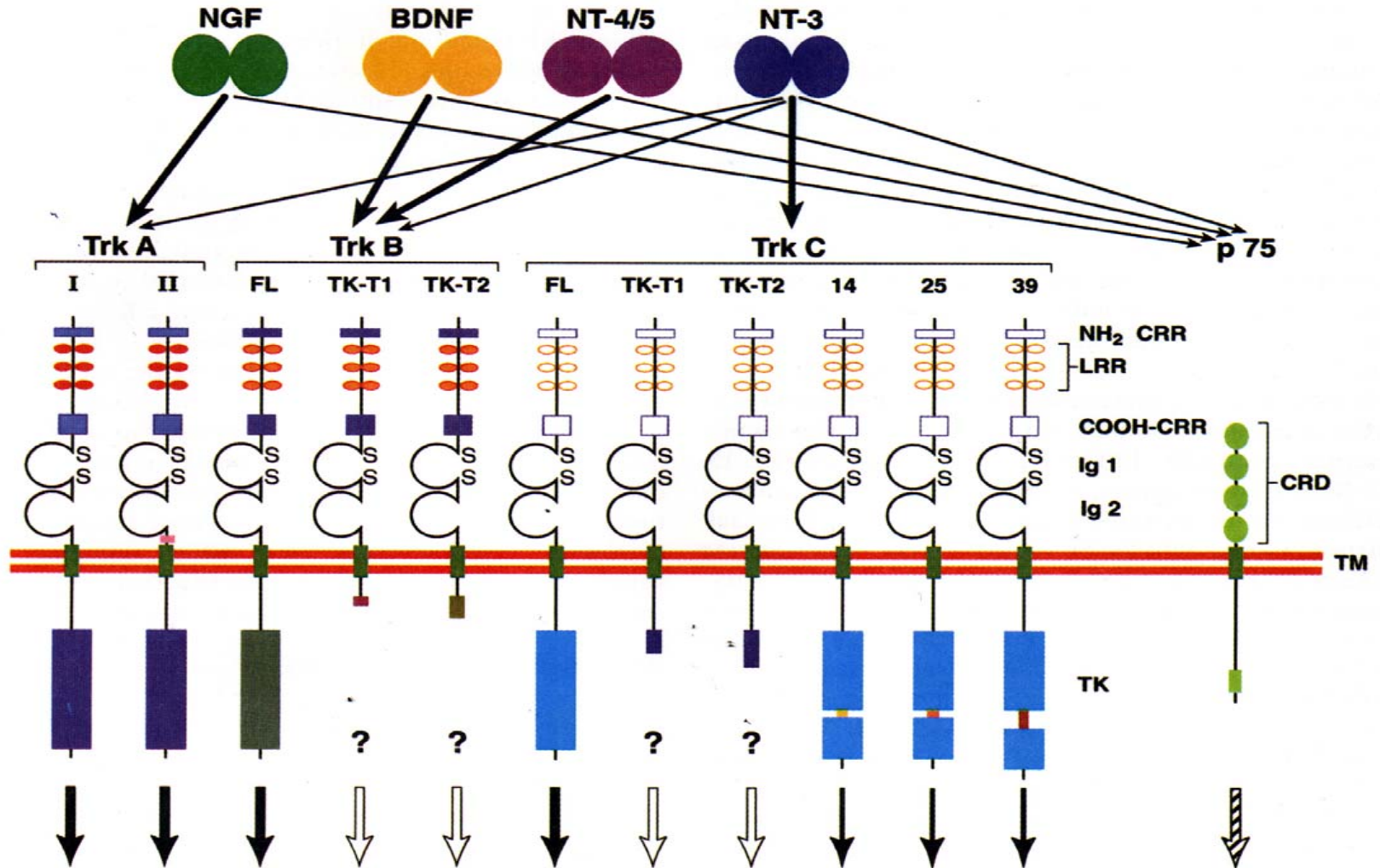
*Благодарю за
внимание!*



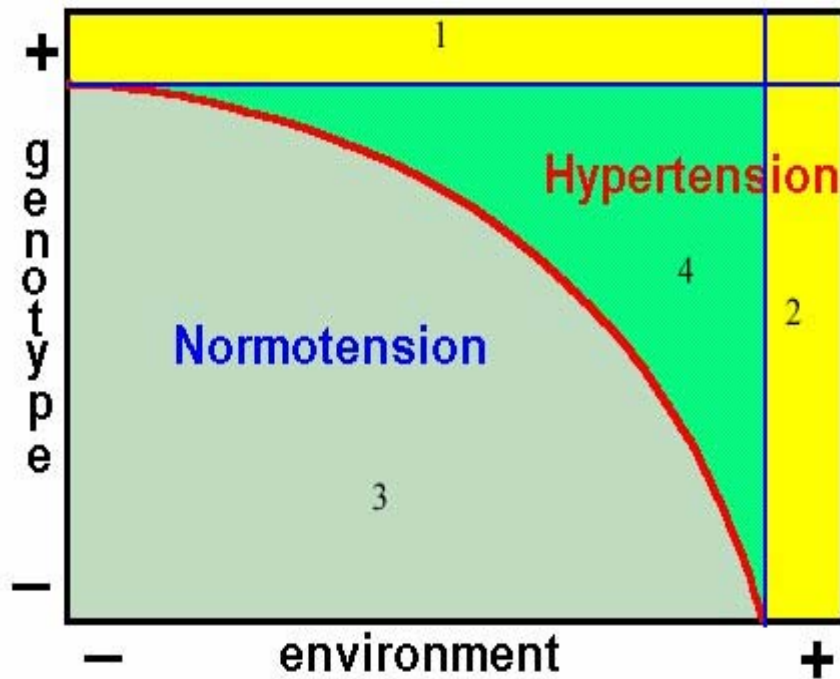
Nervous growth factor receptors family

128

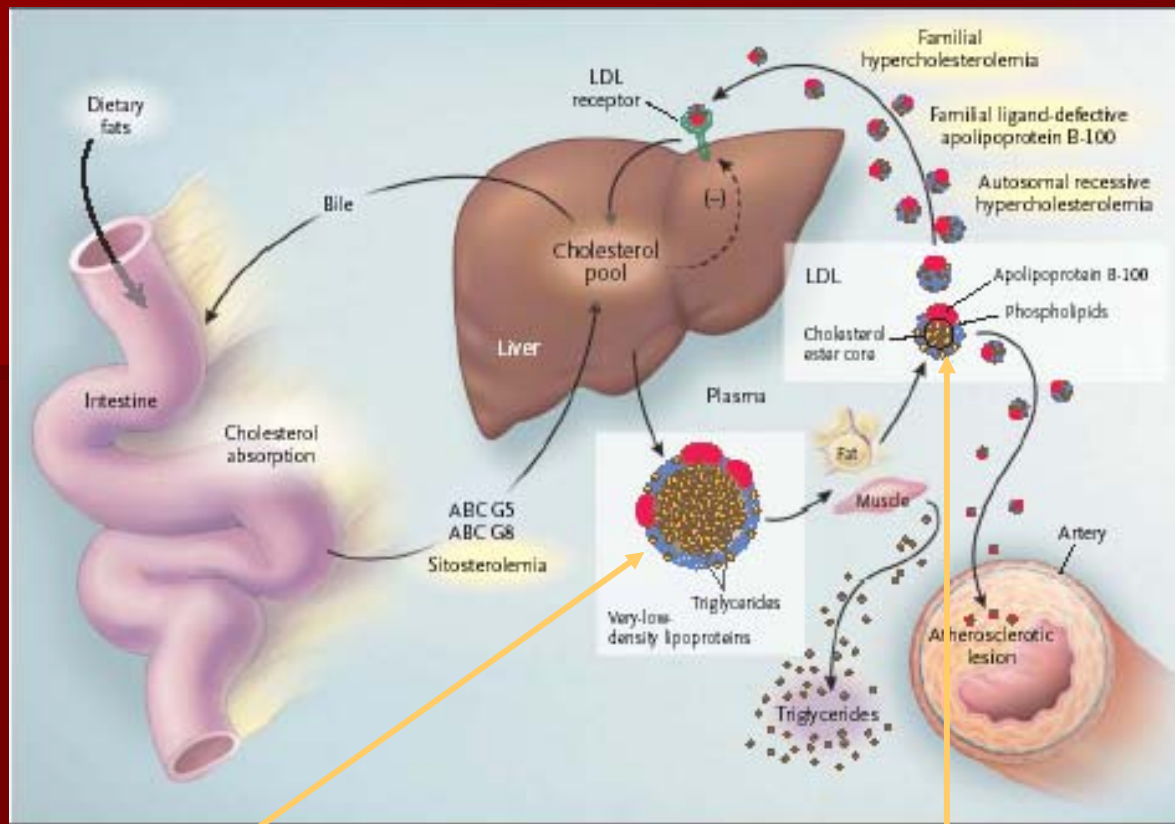
L. Tessarollo



GENOTYPE-ENVIRONMENT INTERACTION IN HYPERTENSION DEVELOPMENT



- 1 - Genotype-dependent hypertension
 - 2 - Environment-dependent hypertension
 - 3 - Normotension
 - 4 - Hypertension dependent on genotype-environment interaction
- + - "Hypertensive" genes or environments
- - "Normotensive" genes or environments



Мутации снижающие клиренс холестерина плазмы крови



Семейная гиперхолестеринемия: мутации (>600 вариантов) гена, кодирующего LDLR (гетерозиготы – 1:500, гомозиготы – 1:1 000 000)

Нарушения связывания LDLR с лигандом – APOB-100: нарушен клиренс холестерина плазмы крови вследствие мутаций гена *APOB-100* (гетерозиготы – 1:1000)

Ситостеролемиа – нарушение транспорта и абсорбции холестерина в просвет кишечника: мутации генов *ABCG5* и *ABCG8* (встречаются редко) – накопление холестерина в печени и подавление синтеза LDLR

Гиперхолестеринемия: редкая аутомальная мутация (1:10000000) гена *ARH* приводит к нарушению функции рецептора LDLR

Печень секретирует липопротеины очень низкой плотности, включающие эфиры холестерина и триглицериды в оболочке из фосфолипидов и аполипопротеина B-100

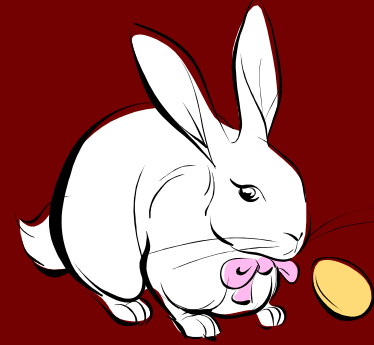
Липопротеин низкой плотности (LDL) – главный носитель холестерина в плазме крови и один из главных факторов сердечно-сосудистой патологии

НОСИТЕЛИ ХОЛЕСТЕРИНА

С последующим изучением
эффектов на уровне фенотипа

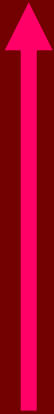


?



Два пути создания генетических моделей

Изменение ДНК

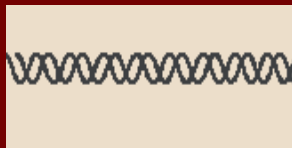


Селекция ДНК

Селекция
по фенотипу



С последующим анализом генома



?

