

# Max-Planck-Institut für Psychiatrie

## Stress und Depression – Einfluss genetischer Faktoren

12. Jahrestagung der DGBS in Hannover  
29. September 2012



MAX-PLANCK-GESELLSCHAFT

Marcus Ising

# Stress und Depression – Einfluss genetischer Faktoren

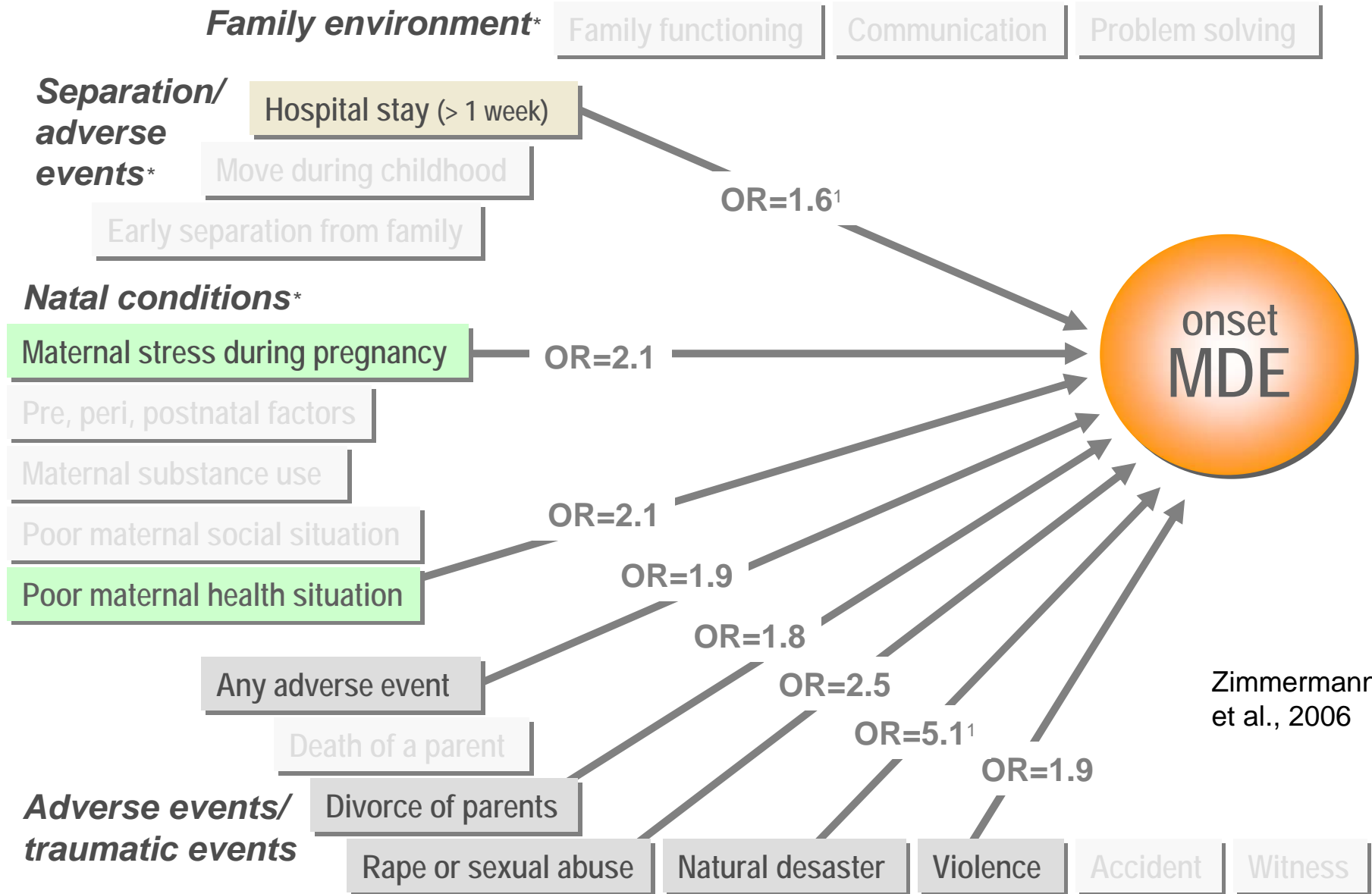
## ***1) Depression – Eine Stresserkrankung***

# Early Developmental Stages of Psychopathology

## Study: Risk Factors for MDE Onset



Max-Planck-Institut für Psychiatrie

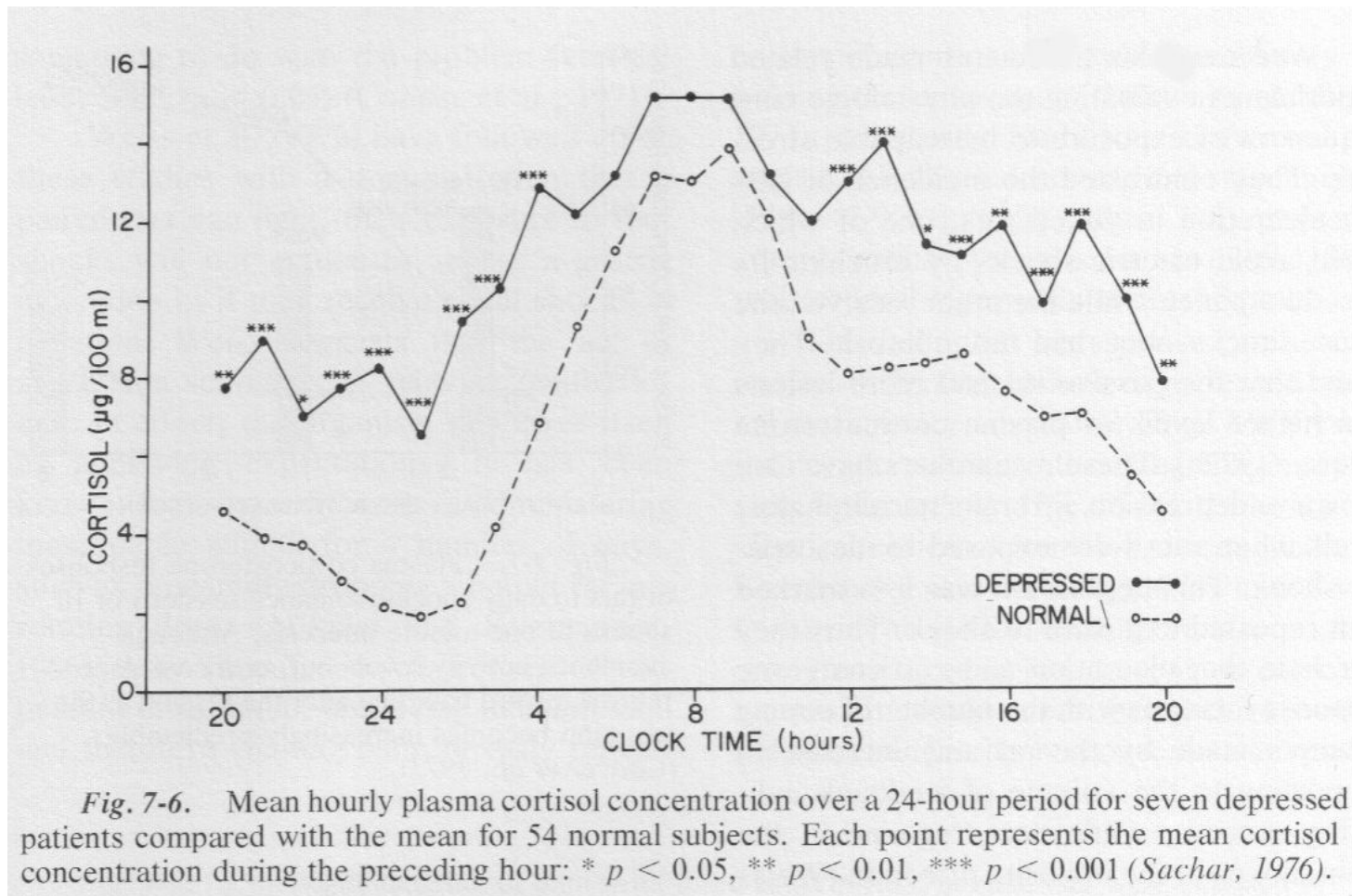


# Plasma Cortisol Levels and Circadian Rhythm



Max-Planck-Institut für Psychiatrie

- Elevated plasma cortisol and attenuated circadian cortisol rhythms in depressed patients compared with normal subjects (Sachar, 1976).



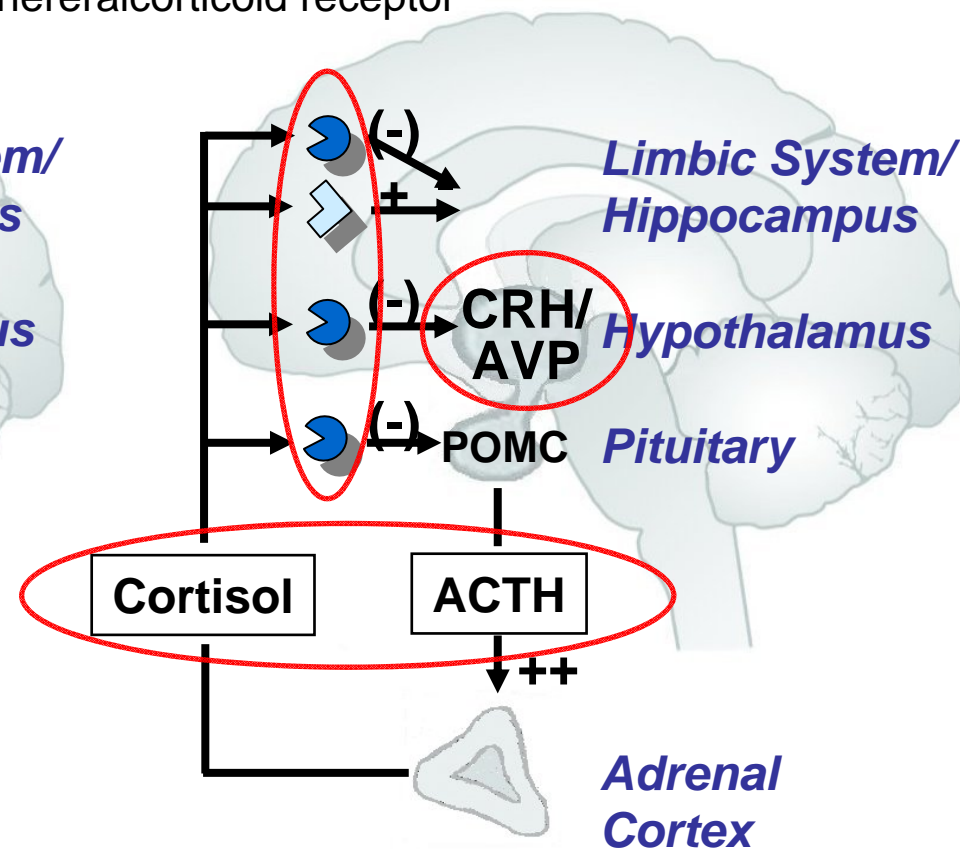
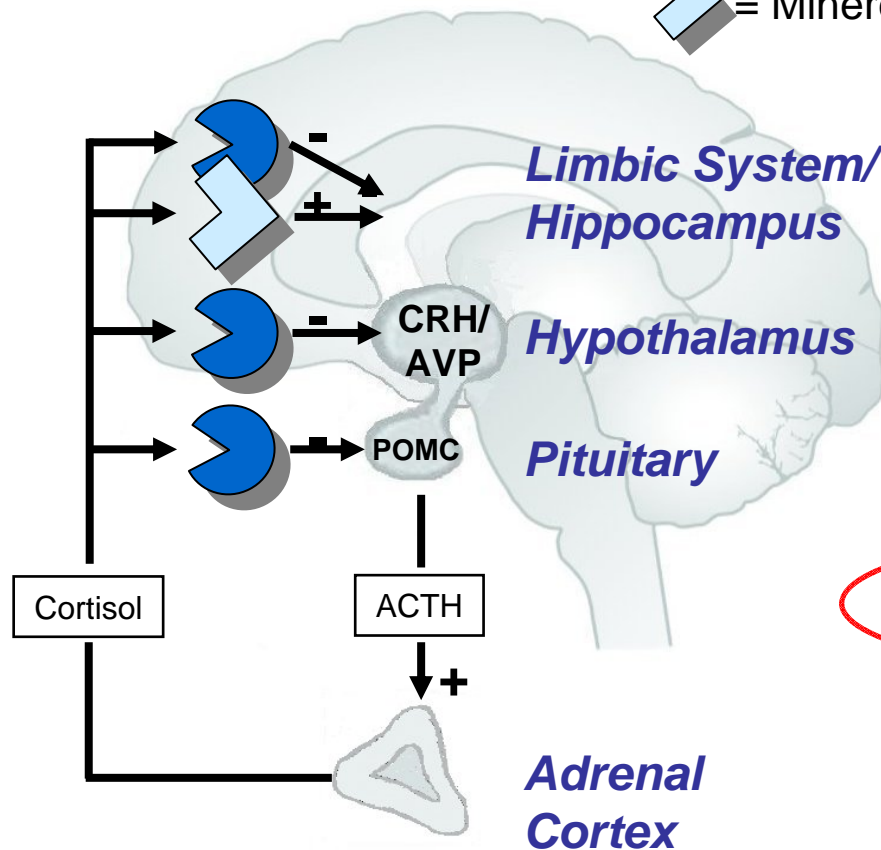
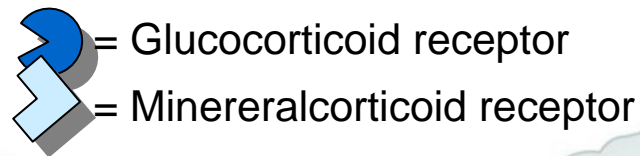
# Impaired Stress Hormone Regulation in Acute Depression



Max-Planck-Institut für Psychiatrie

## Normal Regulation

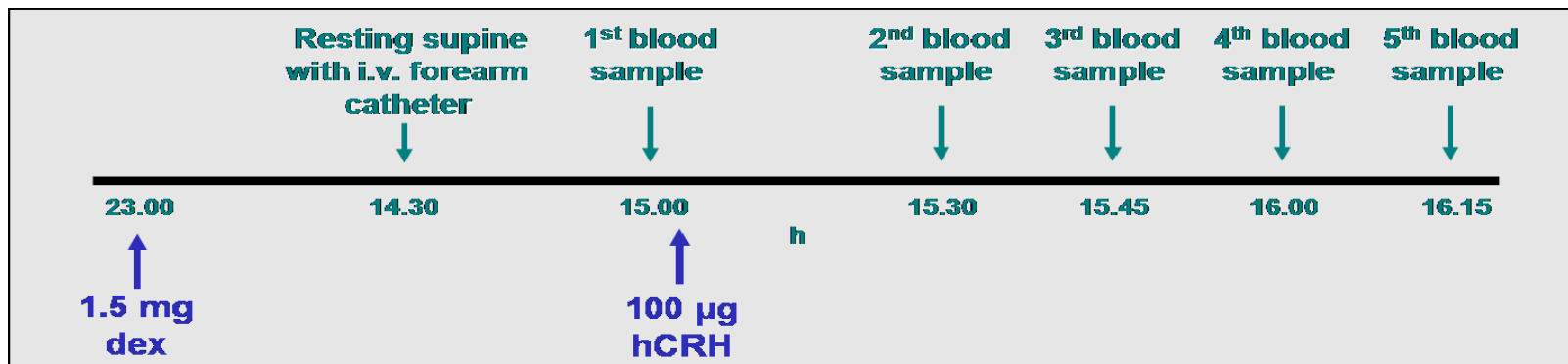
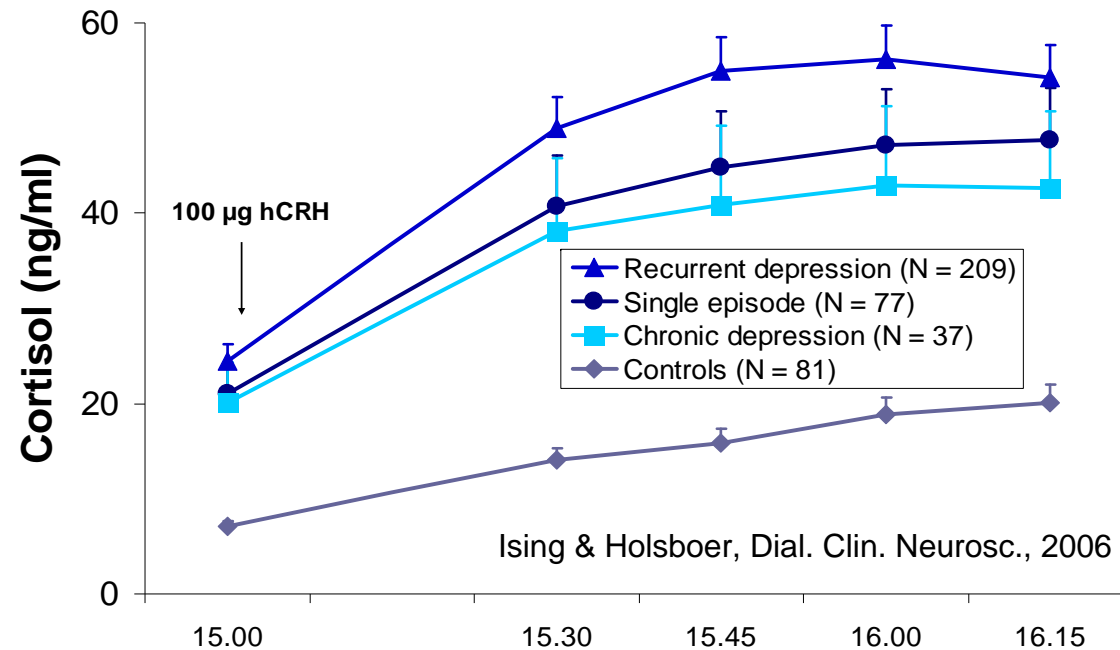
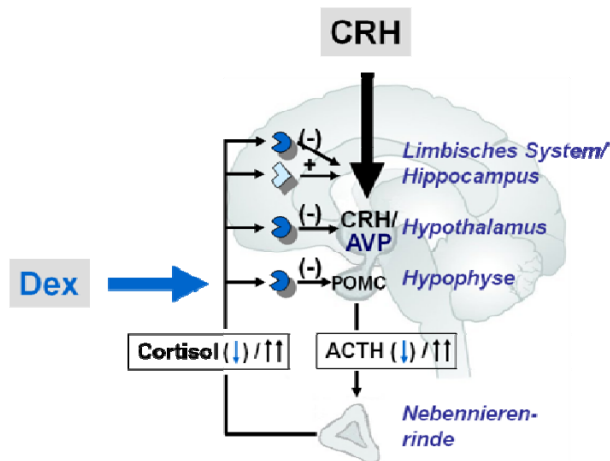
## Impaired Regulation



# Combined Dex/CRH Test to Evaluate Impaired Stress Hormone Regulation



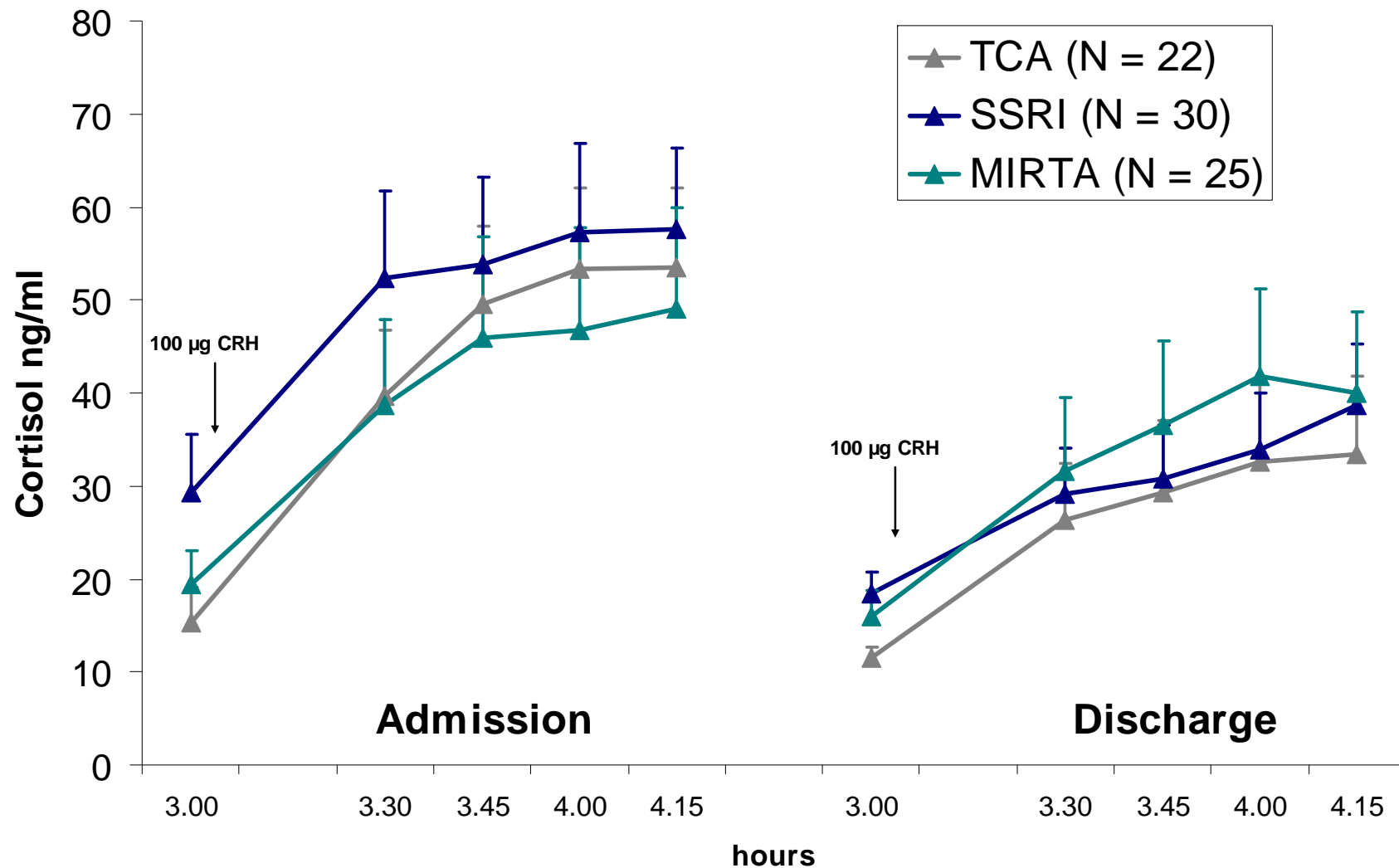
Max-Planck-Institut für Psychiatrie



# Impaired Stress Hormone Regulation Resolves under Monoaminergic Treatment ...



Max-Planck-Institut für Psychiatrie



Ising & Holsboer, Dial. Clin. Neurosc., 2006

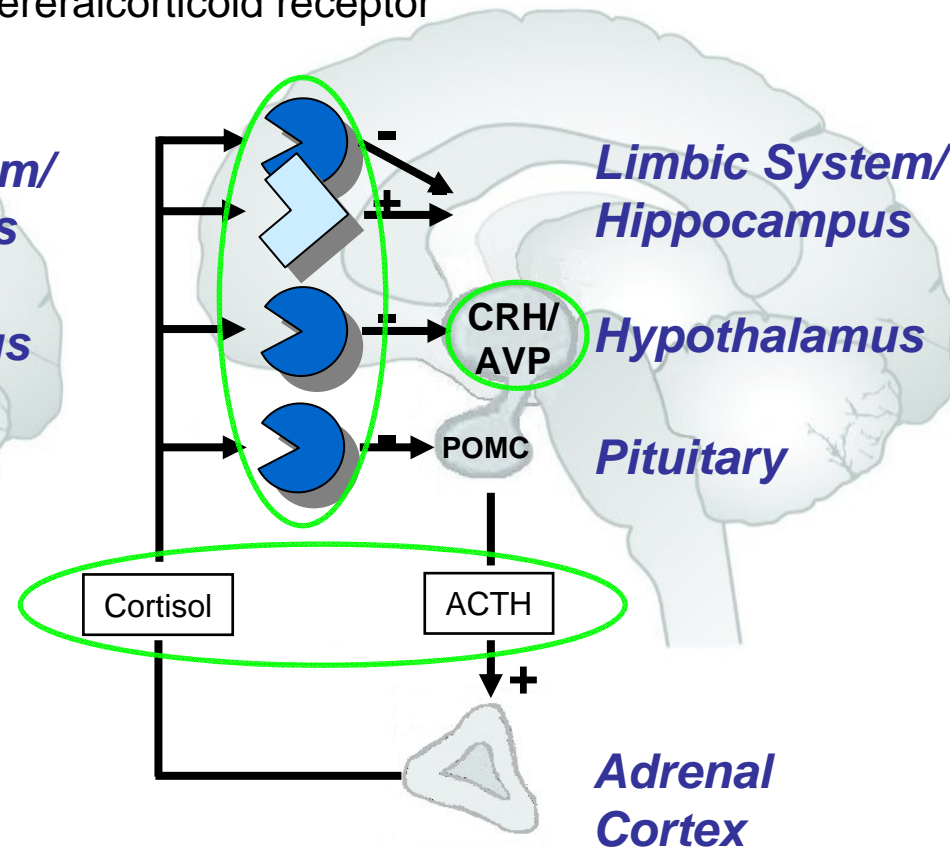
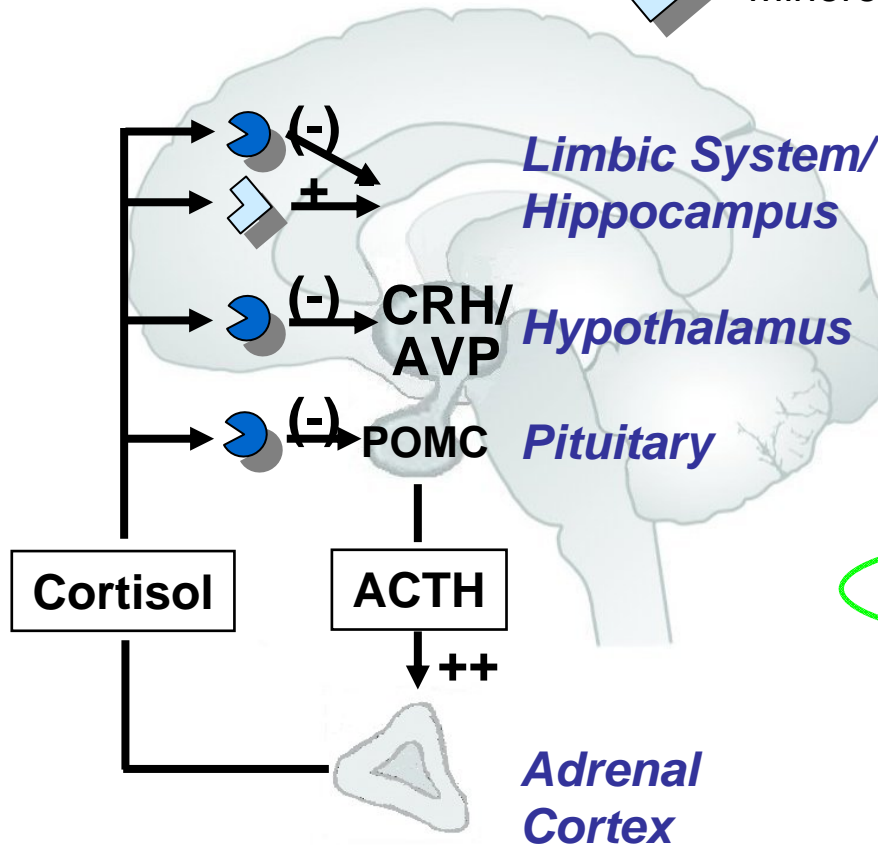
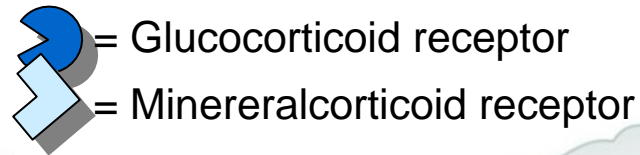
# ... Suggesting a Restoration of the Glucocorticoid Receptor Function



Max-Planck-Institut für Psychiatrie

## Impaired Regulation

## Restored Regulation





# Stress und Depression – Einfluss genetischer Faktoren

## ***2) Genetik der Stresshormonachse und der Circadianen Rhythmik***

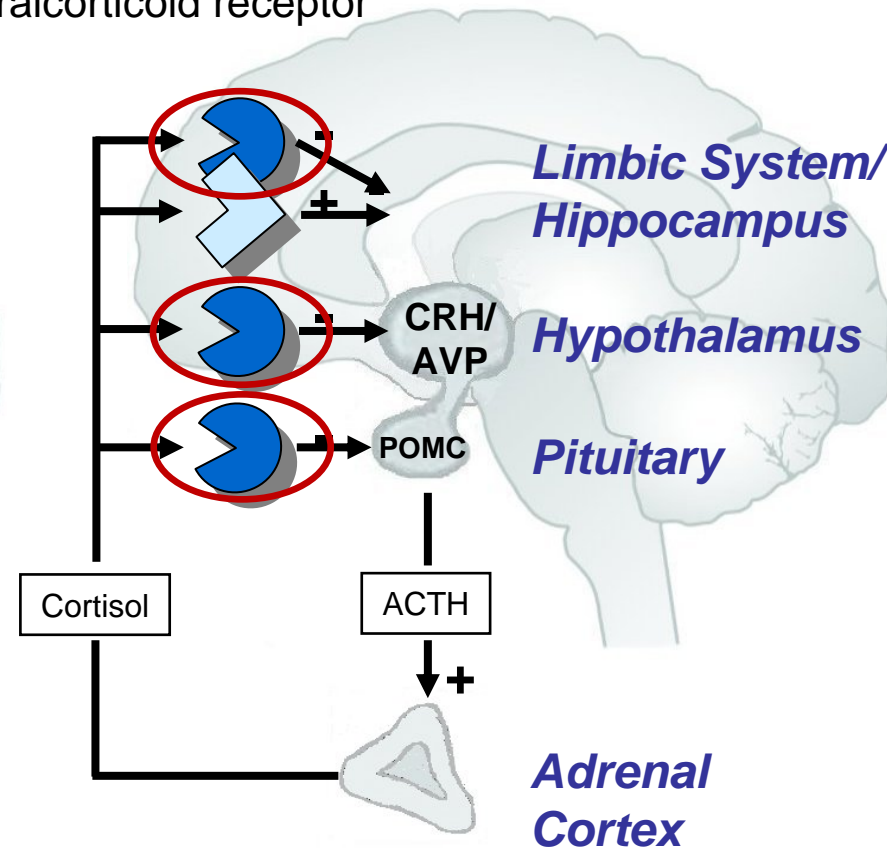
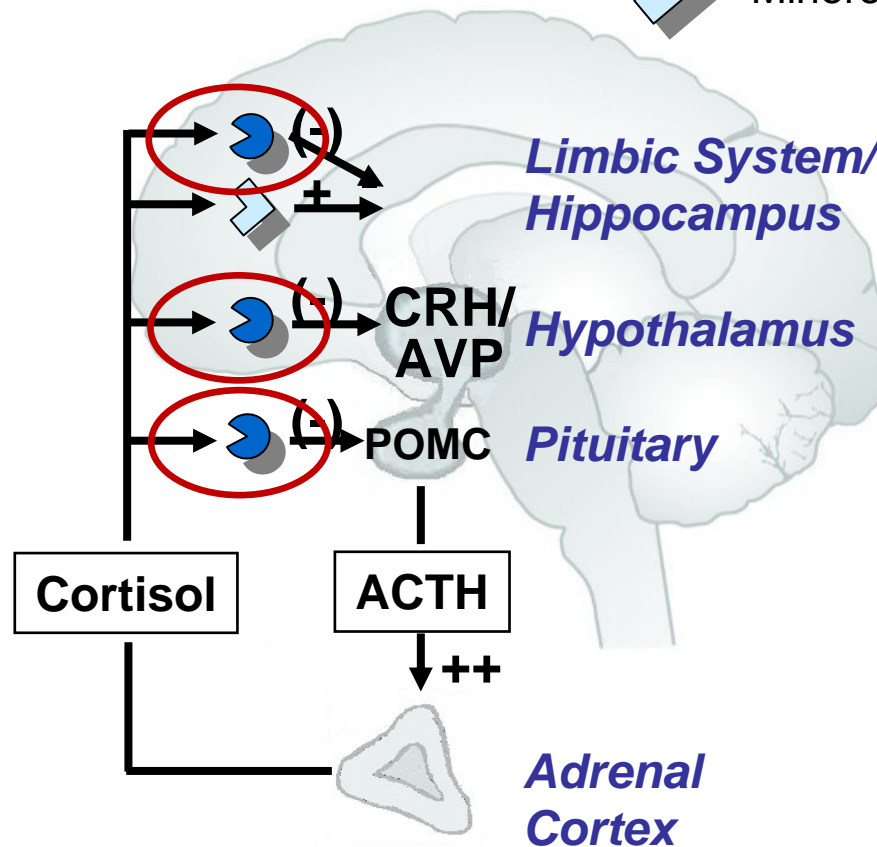
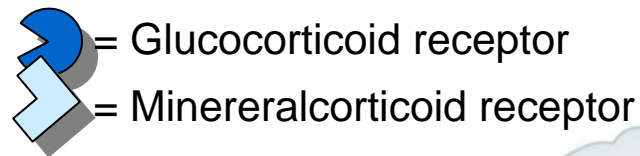
# Genes Involved in Stress Hormone Regulation



Max-Planck-Institut für Psychiatrie

## Impaired Regulation

## Restored Regulation

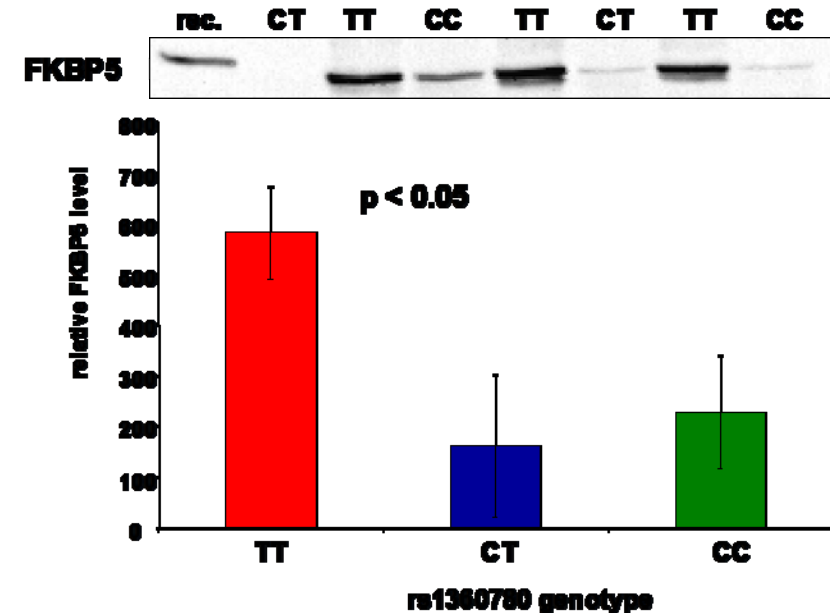
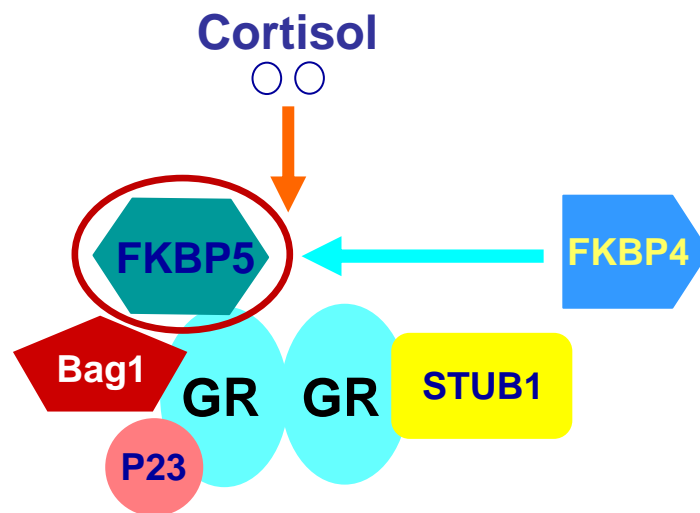


# Glucocorticoid Receptor Complex



Max-Planck-Institut für Psychiatrie

- The glucocorticoid receptor (GR) is a complex structure involving a number of “assistant” proteins, so-called chaperones, moderating glucocorticoid binding and GR translocation into the nucleus. Among all chaperones, FKBP5 seems to be of special importance preventing GR translocation.
- The FKBP5 gene comprises only a single major LD block including mainly intronic SNPs. Nevertheless, these SNPs influence FKBP5 protein levels, presumably, due to linkage with a yet unknown functional variant.



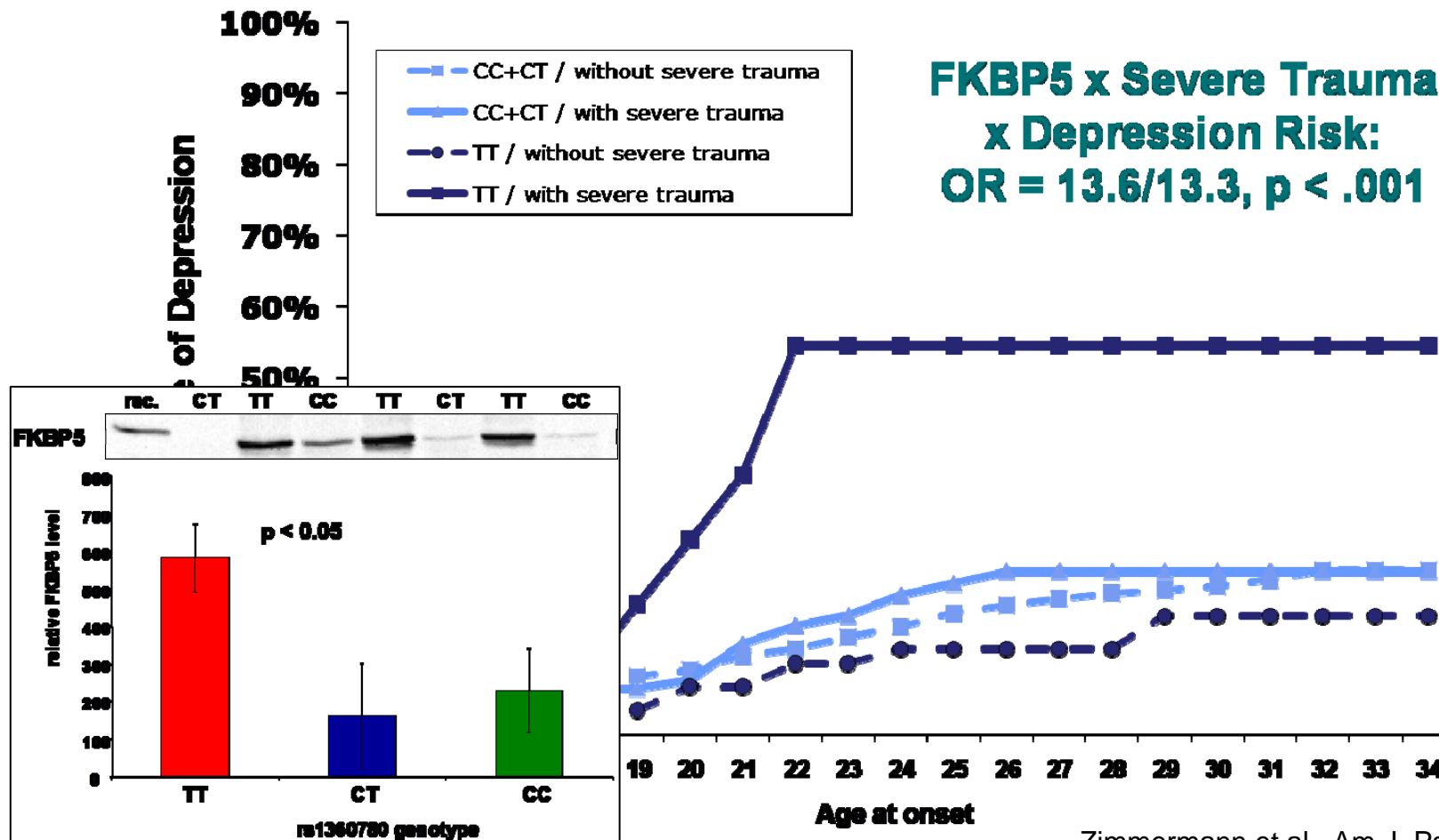
Binder et al., Nat. Gen., 2004

# Polymorphisms in the FKBP5 Gene Affect Risk for Depression in Traumatized Subjects



Max-Planck-Institut für Psychiatrie

- The FKBP5 genotype (rs1360780) associated with increased FKBP5 activity predicts an increased depression risk, but only for those subjects previously exposed to traumatic events.



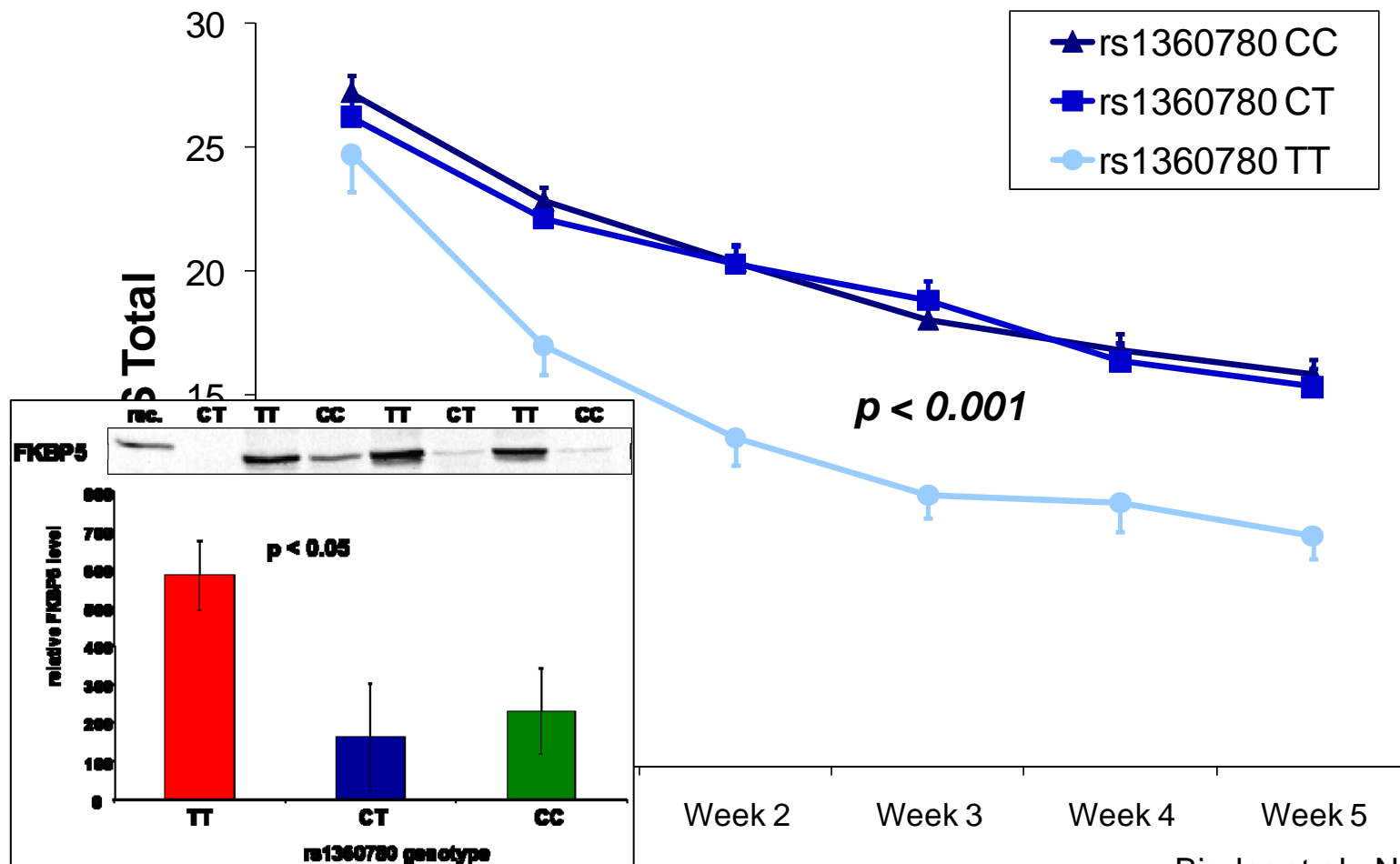
Zimmermann et al., Am J. Psychiat., 2011

# Polymorphisms in the FKBP5 Gene Affect Antidepressant Treatment Outcome



Max-Planck-Institut für Psychiatrie

- The same FKBP5 genotype (rs1360780) associated with increased depression risk also predicts antidepressant treatment outcome.

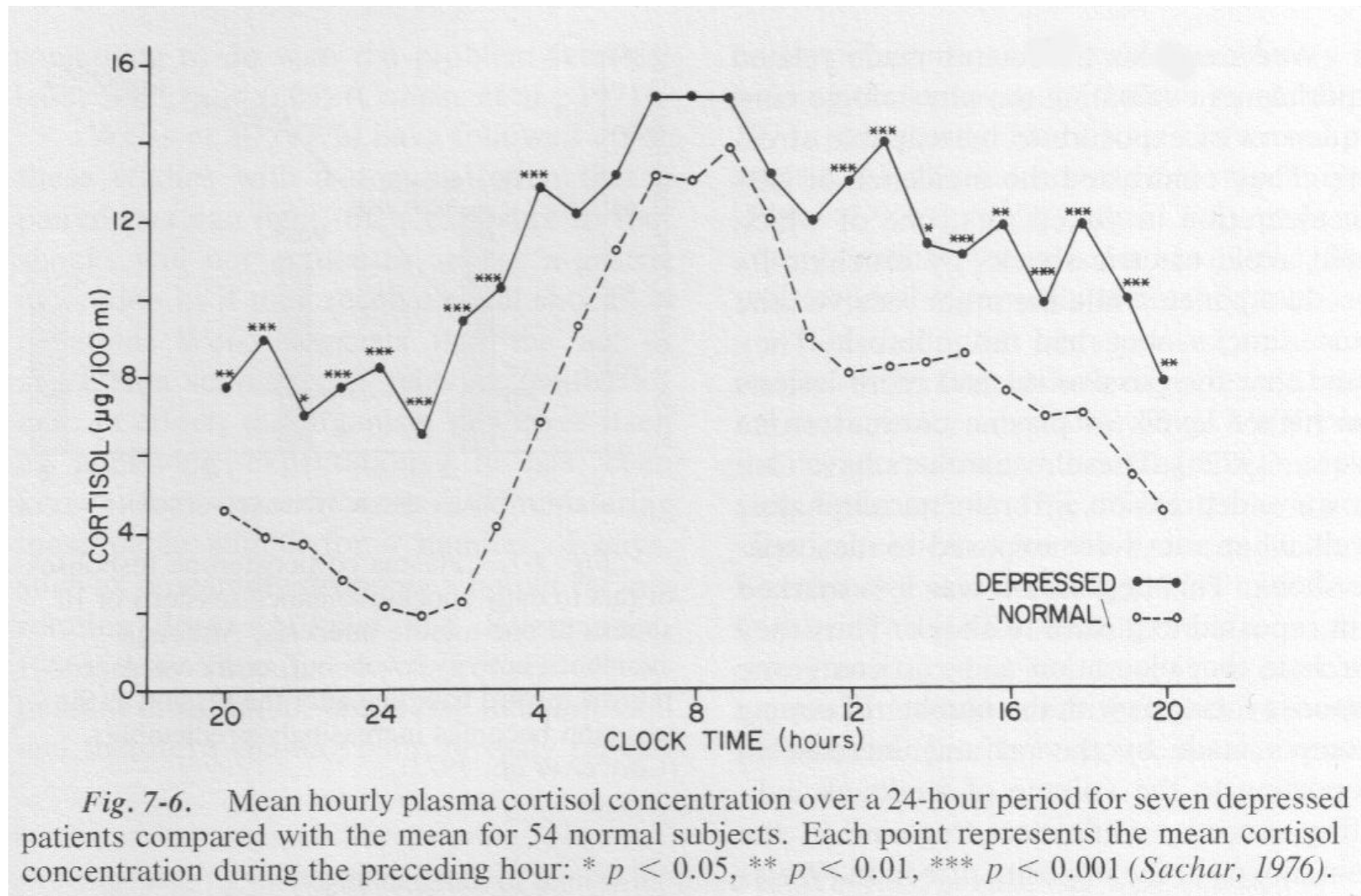


Binder et al., Nat. Gen., 2004

# Plasma Cortisol Levels and Circadian Rhythm



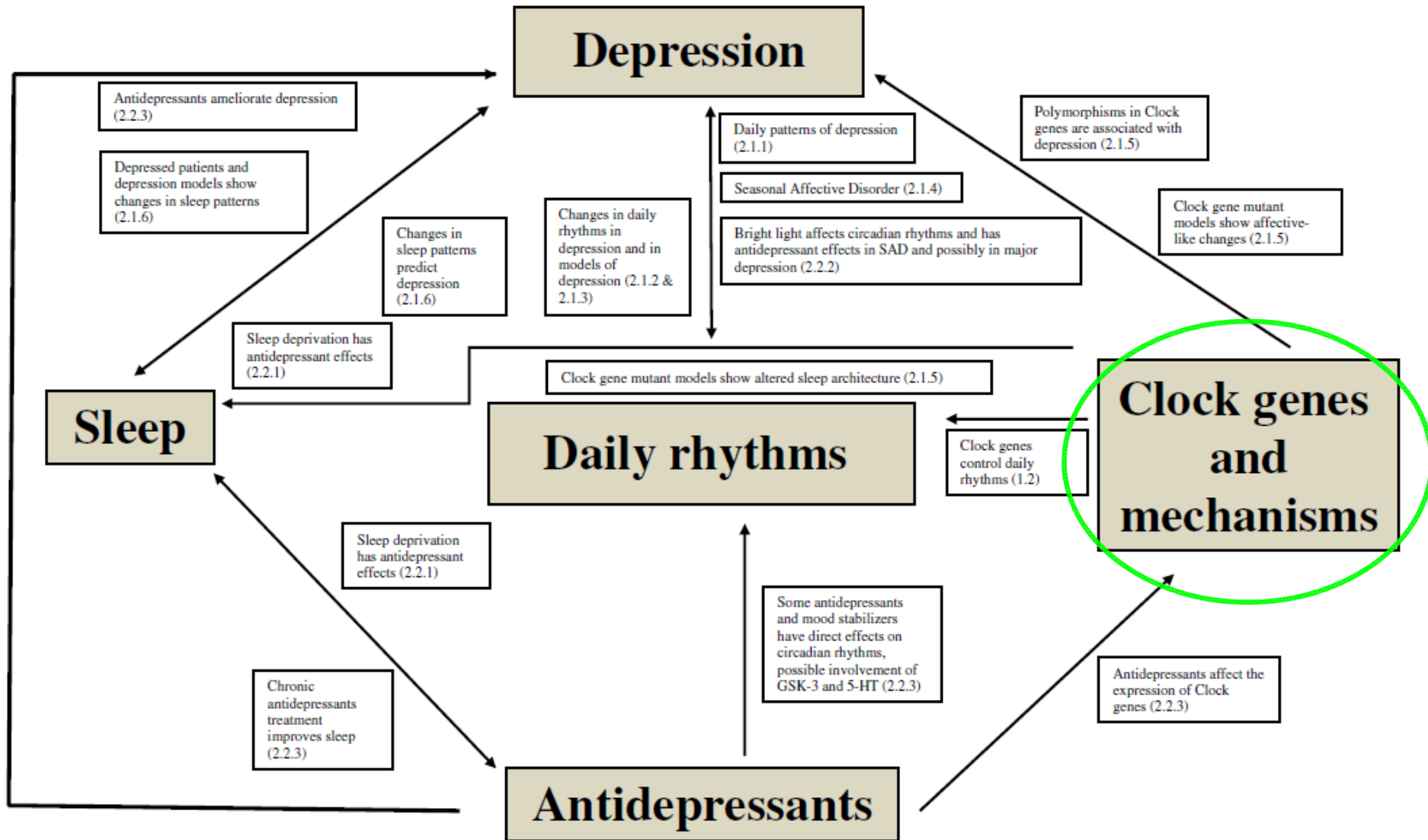
Max-Planck-Institut für Psychiatrie



# Circadian Rhythms in Depression



Max-Planck-Institut für Psychiatrie

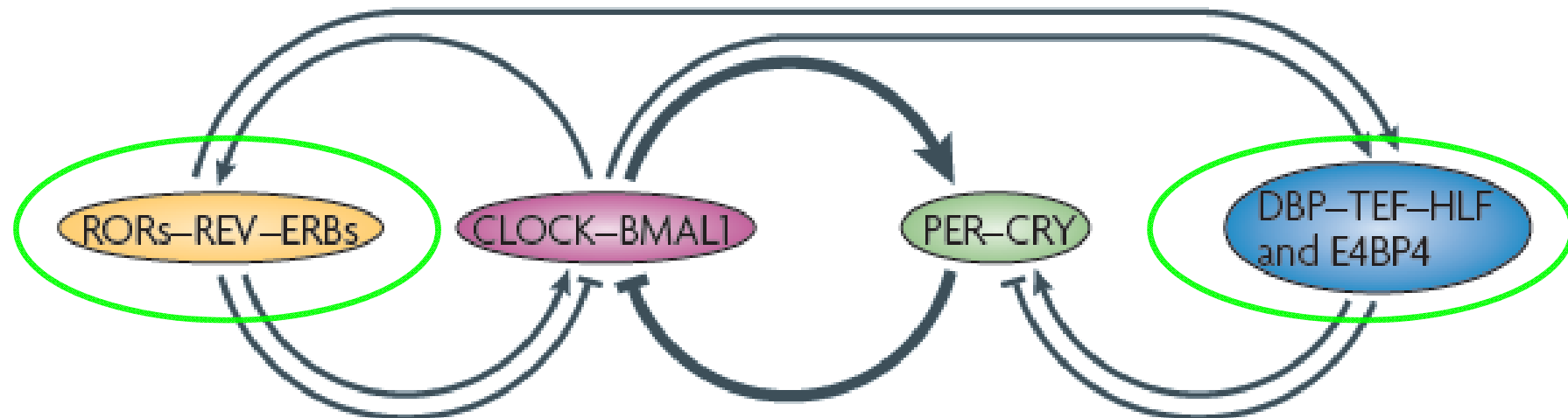


# Primary “Clock Gene” Loops



Max-Planck-Institut für Psychiatrie

- Clock proteins compose regulatory loops, with CLOCK, BMAL1, PER and CRY constituting the primary feedback loop.
- CLOCK and BMAL1 activate PER and CRY, which in turn suppress CLOCK and BMAL1. Transcriptional activators RORs and DBP–TEF–HLF and repressors REV–ERBs and E4BP4 synergistically modify the expression of primary clock proteins, conferring a robust transcriptional regulation to the clock



Zhang & Kay, Nat. Rev. MolCellBio., 2010



## Stress und Depression – Einfluss genetischer Faktoren

### **3) „Clock Genes“, Stresshormonregulation und Depressionsrisiko**

# “Clock Genes” and Affective Disorders



Max-Planck-Institut für Psychiatrie

**Table 2** Studies reporting suggestive associations between circadian genes and mood spectrum disorders BD: bipolar disorder, UD: unipolar disorder, SAD: seasonal affective disorder.

Gene	OMIM nomenclature	Studies reporting suggestive associations		
		BD	UD Major depression	SAD
<i>CLOCK</i>	CIRCADIAN LOCOMOTOR OUTPUT CYCLES KAPUT	(Kripke et al., 2009; Lee et al., 2010; Shi et al., 2008; Soria et al., 2010b)	(Soria et al., 2010b)	
<i>NPAS2</i>	NEURONAL PAS DOMAIN PROTEIN 2	(Kripke et al., 2009; Mansour et al., 2009; Soria et al., 2010b)	(Soria et al., 2010b)	(Johansson et al., 2003; Partonen et al., 2007)
<i>ARNTL1</i>	ARYL HYDROCARBON RECEPTOR NUCLEAR TRANSLOCATOR-LIKE 1 (BMAL1)	(Mansour et al., 2006, 2009; Nievergelt et al., 2006; Soria et al., 2010b)	(Soria et al., 2010b; Utge et al., 2010)	(Partonen et al., 2007)
<i>ARNTL2</i>	ARYL HYDROCARBON RECEPTOR NUCLEAR TRANSLOCATOR-LIKE 2 (BMAL2)	(Soria et al., 2010b)	(Soria et al., 2010b)	
<i>PER1</i>	PERIOD HOMOLOG 1 (DROSOPHILA)	(Kripke et al., 2009)		
<i>PER2</i>	PERIOD HOMOLOG 2 (DROSOPHILA)	(Kripke et al., 2009)	(Soria et al., 2010b)	(Partonen et al., 2007)
<i>PER3</i>	PERIOD HOMOLOG 3 (DROSOPHILA)	(Mansour et al., 2006; Nievergelt et al., 2006; Soria et al., 2010b)	(Soria et al., 2010b)	
<i>CRY1</i>	CRYPTOCHROME 1	(Soria et al., 2010b)	(Soria et al., 2010b)	
<i>CRY2</i>	CRYPTOCHROME 2	(Mansour et al., 2009)		(Lavebratt et al., 2010)
<i>TIMELESS</i>	TIMELESS HOMOLOG (DROSOPHILA)	(Mansour et al., 2006)	(Utge et al., 2010)	
<i>NR1D1</i>	NUCLEAR RECEPTOR SUBFAMILY 1, GROUP D, MEMBER 1	(Kishi et al., 2008; Kripke et al., 2009; Severino et al., 2009) *	(Soria et al., 2010b)	
<i>RORA</i>	RAR-RELATED ORPHAN RECEPTOR A	(Soria et al., 2010b)	(Utge et al., 2010)	
<i>RORB</i>	RAR-RELATED ORPHAN RECEPTOR B	(Mansour et al., 2009; McGrath et al., 2009)		
<i>CSNK1δ</i>	CASEINE KINASE I DELTA	(Kripke et al., 2009)		
<i>CSNK1ε</i>	CASEINE KINASE I EPSILON	(Mansour et al., 2009; Shi et al., 2008; Soria et al., 2010b)	(Utge et al., 2010)	
<i>GSK3β</i>	GLYCOGEN SYNTHASE KINASE 3-BETA	(Szczechankiewicz et al., 2006) *		
<i>DBP</i>	D SITE OF ALBUMIN PROMOTER-BINDING PROTEIN		(Soria et al., 2010b)	
<i>BHLHB2</i>	BASIC HELIX-LOOP-HELIX DOMAIN CONTAINING, CLASS B, 2	(Shi et al., 2008)		
<i>BHLHB3</i>	BASIC HELIX-LOOP-HELIX DOMAIN CONTAINING, CLASS B, 3	(Soria et al., 2010b)		
<i>ASMT</i>	ACETYLSEROTONIN METHYLTRANSFERASE, X-CHROMOSOMAL		(Galecki et al., 2010)	
<i>MTNR1B</i>	MELATONIN RECEPTOR 1B		(Galecka et al., in press)	
<i>AANAT</i>	ARYLALKYLAMINE N-ACETYLTRANSFERASE		(Soria et al., 2010a)	
<i>PPARGC1B</i>	PEROXISOME PROLIFERATOR-ACTIVATED RECEPTOR-GAMMA, COACTIVATOR 1, BETA	(Kripke et al., 2009)		

\*: association observed only for females with bipolar disorders (Kishi et al., 2009; Szczechankiewicz et al., 2006).

Etain et al.,  
Europ. Neuropsychopharm., 2011

# Clock Gene Associations with the Combined Dex/CRH Test



Max-Planck-Institut für Psychiatrie

- 350 acutely depressed inpatients of the Munich Antidepressant Response Signature (MARS) project participating in a combined dex/CRH test during the first 10 days after clinic admission have been genotyped for the following 18 clock genes.

➤ **ARNTL2**

➤ **BHLHB2**

➤ **CLOCK**

➤ **CRY1, CRY2**

➤ **CSNK1D, E**

➤ **GSK3B**

➤ **MTNR1A, 1B**

➤ **NPAS2**

➤ **NR1D1**

➤ **PER1, 2, 3**

➤ **RORA, RORB**

➤ **TIMELESS**

# Results of the Genetic Association Analysis



Max-Planck-Institut für Psychiatrie

- Controlled for population stratification, 37 SNPs within eight genes showed nominal significant association with the cortisol response in the combined dex/CRH test.

➤ **BHLHB2**: 1x 5'UTR

➤ **CLOCK**: 1x intronic

➤ **CSNK1E**: 2x 5'UTR

➤ **GSK3B**: 2x intronic, 1x 5'UTR

➤ **MTNR1A**: 1x intronic

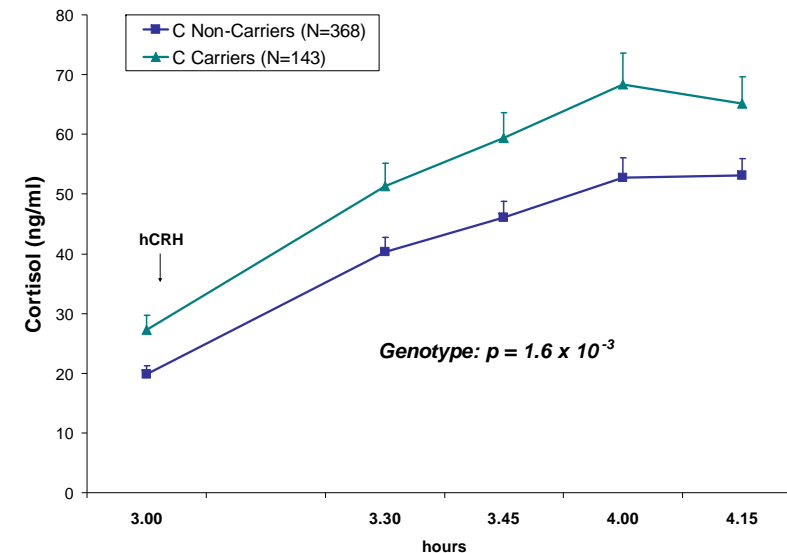
➤ **NPAS2**: 1x intronic

➤ **RORA**: 24x intronic, 2x 5'UTR

➤ **RORB**: 1x intronic, 1x 5'UTR

- Most consistent associations were found for **RORA**, expressing RAR-related orphan receptor alpha, which is a transcriptional activator of the primary clock gene loop.

- The strongest association was observed for the intronic SNP rs17204910 located in **RORA**:  $p = 1.6 \times 10^{-3}$ .

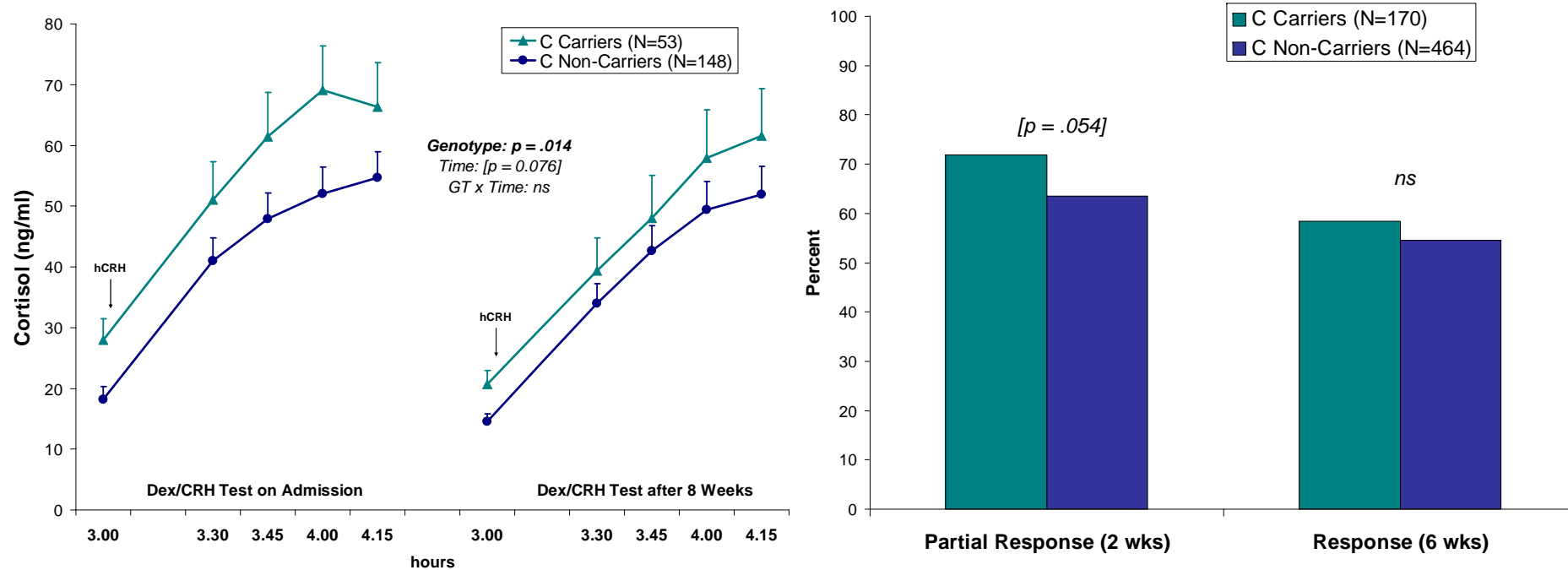


# Effects of RORA (rs17204910) on Depression Risk and Antidepressant Treatment Outcome



Max-Planck-Institut für Psychiatrie

- Besides its effect on stress hormone regulation, rs17204910 (RORA) shows substantial effects on depression risk: ***RR = .71 (protective C allele), p = .005 (N=711/540).***
- However: no effect on change in stress hormone regulation and [almost] no effect on antidepressant treatment outcome.

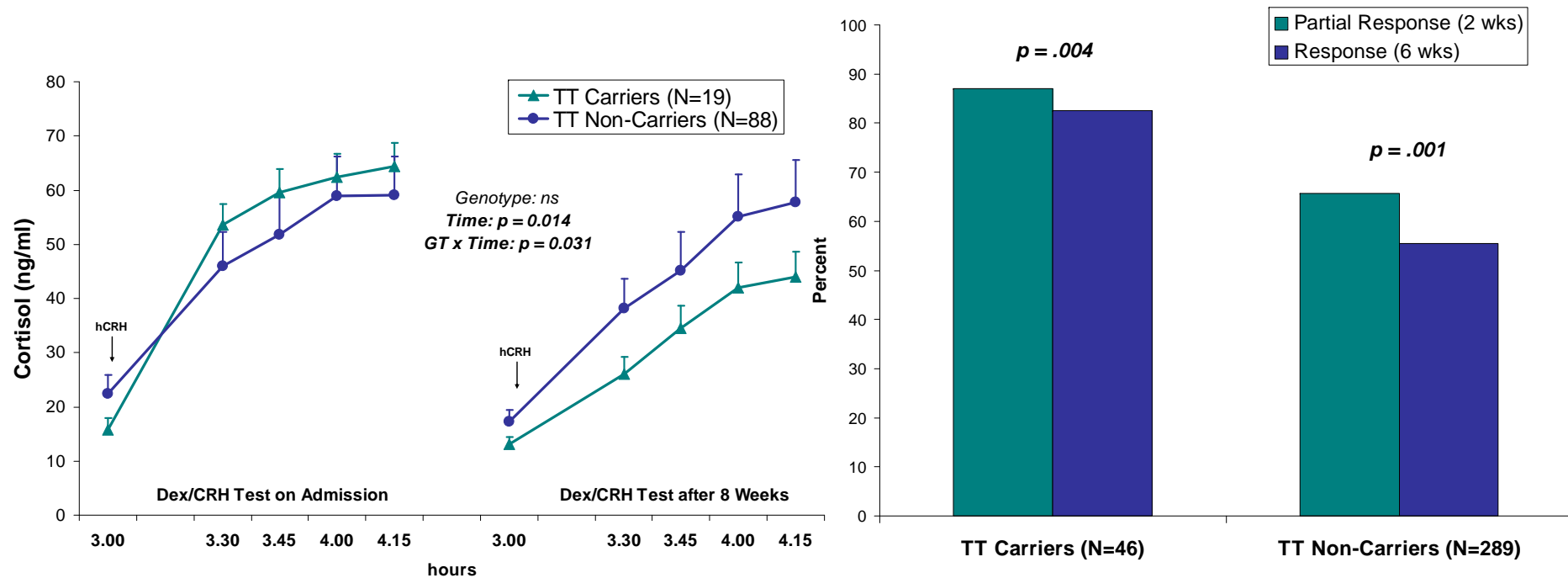


# Effects of FKBP5 on Stress Hormone Regulation and Antidepressant Treatment Outcome



Max-Planck-Institut für Psychiatrie

- In contrary to RORA, the FKBP5 variant rs1360780 shows profound effects on both, change in stress hormone regulation and antidepressant treatment outcome.



# Stress und Depression – Einfluss genetischer Faktoren

## ***4) Zusammenfassung und Einordnung der Ergebnisse***

# Key Messages



Max-Planck-Institut für Psychiatrie

## 1) *Depression – A Stress-related Disorder*

- Chronic stress and intensive trauma are risk factors for depression.
- Acute depression is characterized by an impaired stress hormone regulation and a disturbance of circadian rhythms.

## 2) *Genetics of Stress Hormone Regulation and Circadian Rhythms*

- Strongest and most consistent genetic associations with stress hormone regulation and antidepressant treatment outcome have been shown for the FKBP5 gene moderating stress sensitivity.
- Circadian rhythms are primarily regulated by CLOCK, BMAL1, PER and CRY constituting the circadian feedback loop. These genes are regulated by further “clock genes” including the ROR gene family.

## 3) *“Clock Genes”, Stress Hormone Regulation, and Depression Risk*

- Most consistent associations with disturbed stress hormone regulation were observed for RORA.
- RORA variants are further associated with depression risk, but do not majorly predict monoaminergic antidepressant treatment outcome.



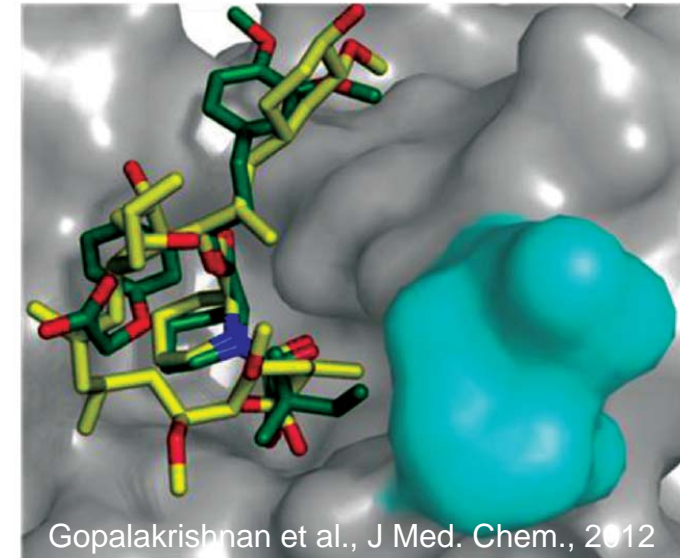
# Perspective of a Personalized Medicine Approach in Depression



Max-Planck-Institut für Psychiatrie

## 1) *FKBP5 driven Stress Hormone Dysregulation in Depression*

- Stress- associated impairment of stress hormone regulation
- Improves under monoaminergic antidepressant treatment
- Potentially best treated with treatments directed to improve stress hormone regulation



## 2) *Clock Genes driven Stress Hormone Dysregulation in Depression*

- Impaired stress hormone regulation due to disturbed chronobiology
- Does not improve under monoaminergic antidepressant treatment
- Potentially best treated with antidepressant medication directed against chronobiological disturbance

# Studien-Team



Max-Planck-Institut für Psychiatrie

Stress und  
genet



Einfluss



Study  
Coordinators

Lab &  
Genotyping

Statistical  
Genetics  
Group



Director

