
Pharmacogenetics of Antidepressant Response in Late Life Depression

Chi-Un Pae, MD, PhD

**Department of Psychiatry
The Catholic University of Korea College of Medicine**

**Department of Psychiatry and Behavioral Sciences
Duke University Medical Center**

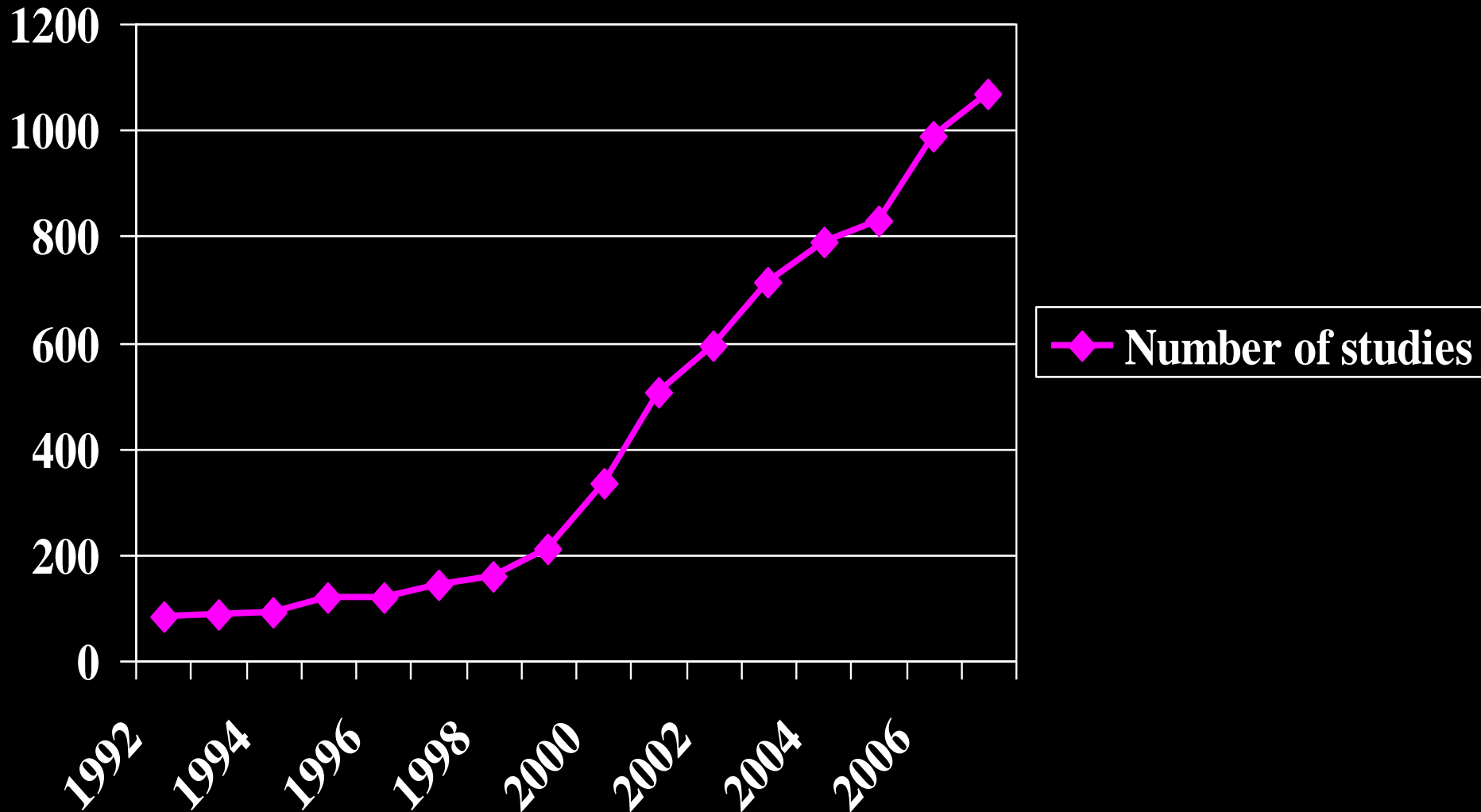
History

- 510 BC Pythagoras some people develop haemolytic anaemia after eating fava beans
- 1902 Garrod genetic factors direct chemical transformations
- 1932 Snyder phenylthiourea nontasting is inherited as an autosomal recessive trait
- 1957 Motulsky first demonstration of the relationship between adverse drug reaction and genetically determined variation
- 1959 Vogel "pharmacogenetics": the hereditary basis of variability in drug effects
- 1960 Evans speed of INH acetylation is under genetic control
- 1962 Kalow abnormal form of serum cholinesterase causes adverse reactions to succinylcholine
- 1977 Mahgoub polymorphism of CYP2D6 causes adverse effects to debrisoquine

Current trend of pharmacogenetics in Psychiatry

Pharmacogenetic studies

(Medline 1992-2008)



Finally FDA recommend submission of pharmacogenetic information on labeling

FDA News

Media Inquiries:

Sandy Walsh, 301-827-3418

Consumer Inquiries:

888-INFO-FDA

FOR IMMEDIATE RELEASE

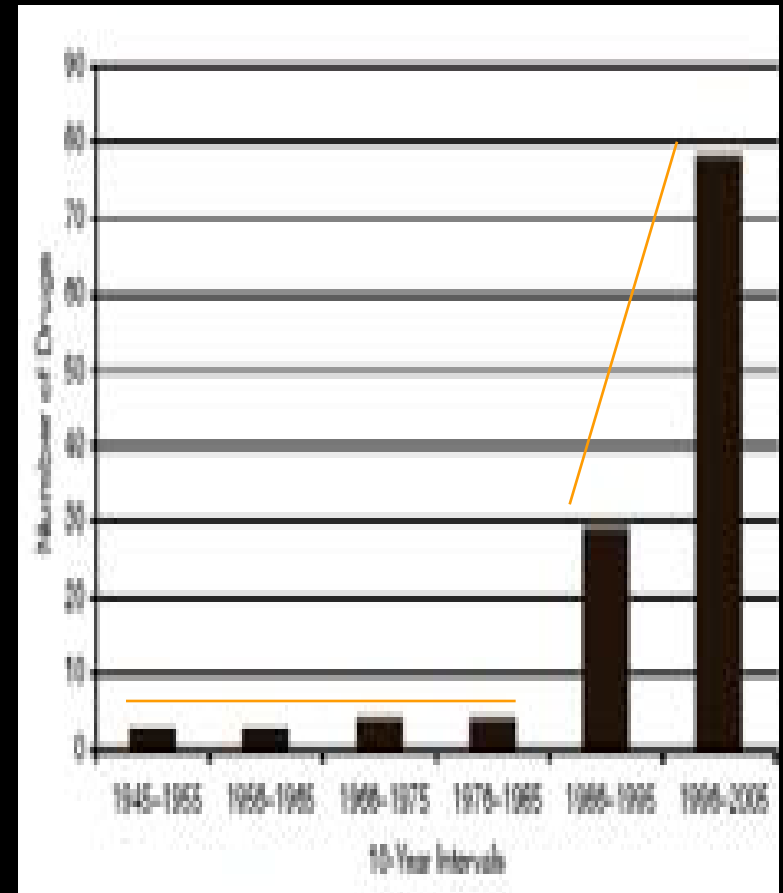
December 12, 2007

Carbamazepine Prescribing Information to Include Recommendation of Genetic Test for Patients with Asian Ancestry

Connection of genetic information with medication use can improve safe use of product

Number of drug approved with pharmacogenetic information

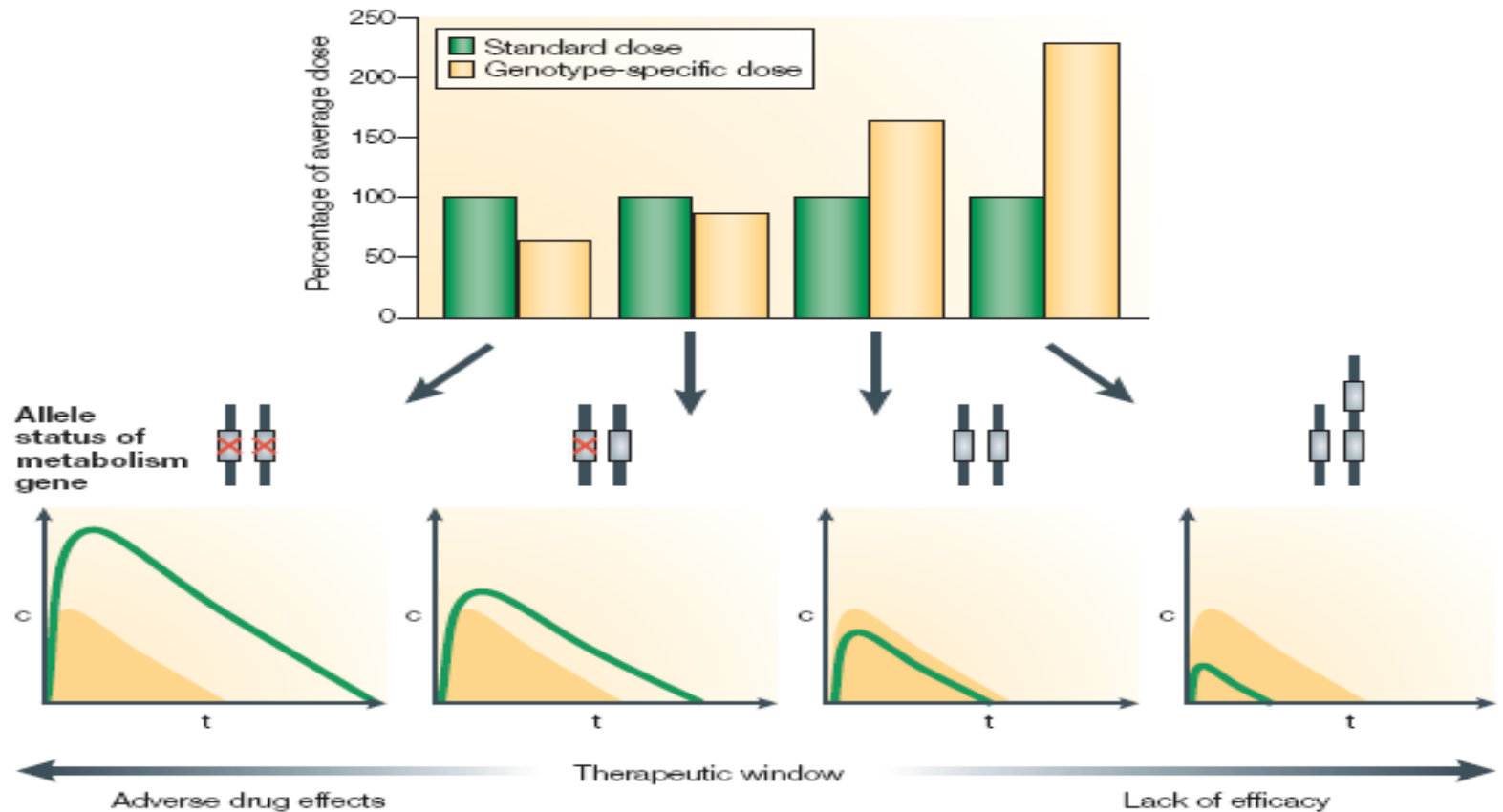
- Pharmacogenomic information is contained in about ten percent of labels for drugs approved by the FDA
- A significant increase of labels containing such information has been observed over the last decade



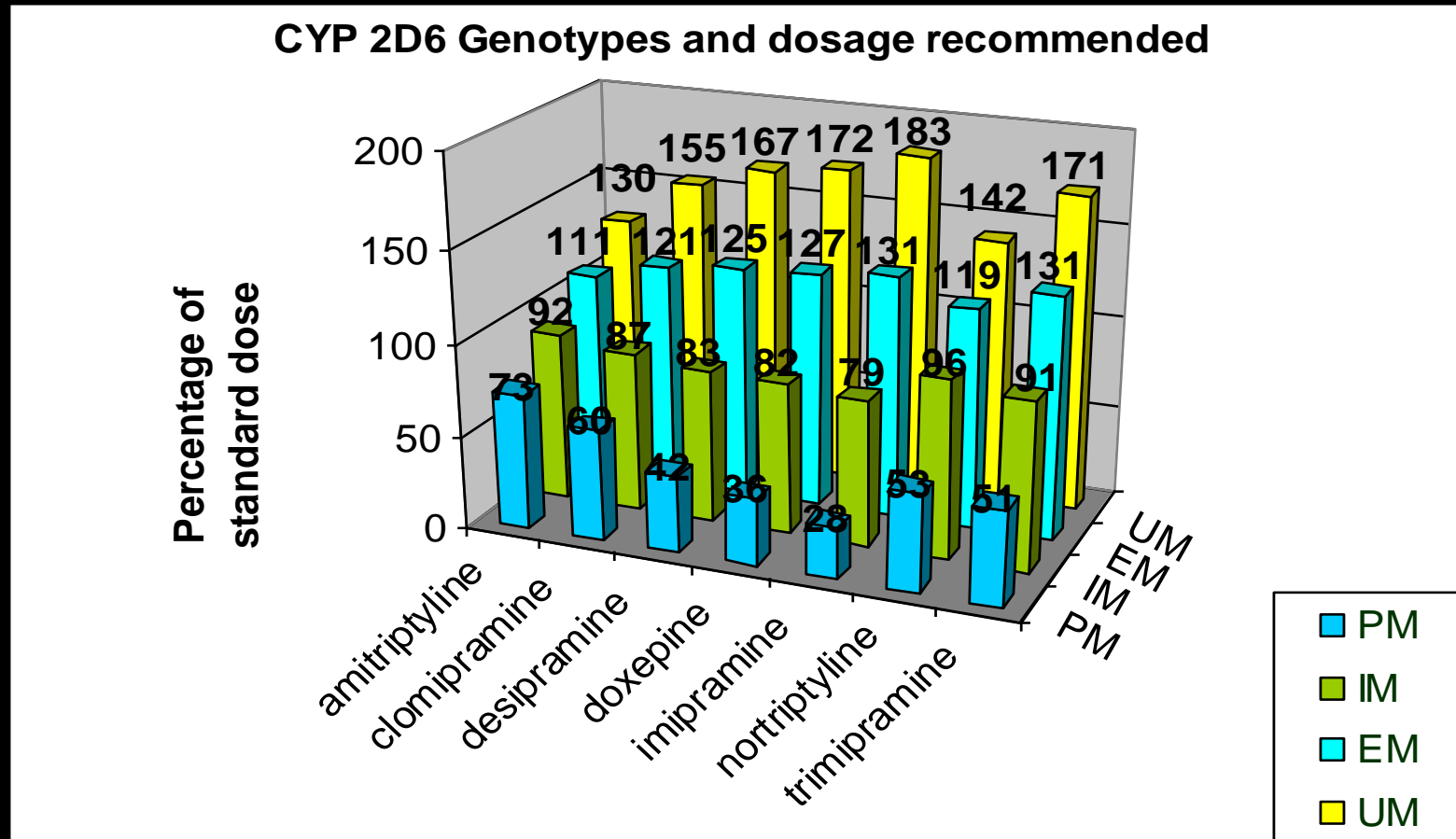
Pharmacogenetic Test Information on Drug Labels

- **Test required** (n=4); **Test recommended** (n=7) : **Information only** (more than 150)
- Currently, 4 drugs are required to have pharmacogenetic testing performed before they are prescribed: cetuximab, trastuzumab, maraviroc, and dasatinib
- HLA-B*1502: Carbamazepine, test recommended
- Urea cycle disorders : Valproic acid, test recommended

PGx in therapeutic decision-making: Dose adaptation

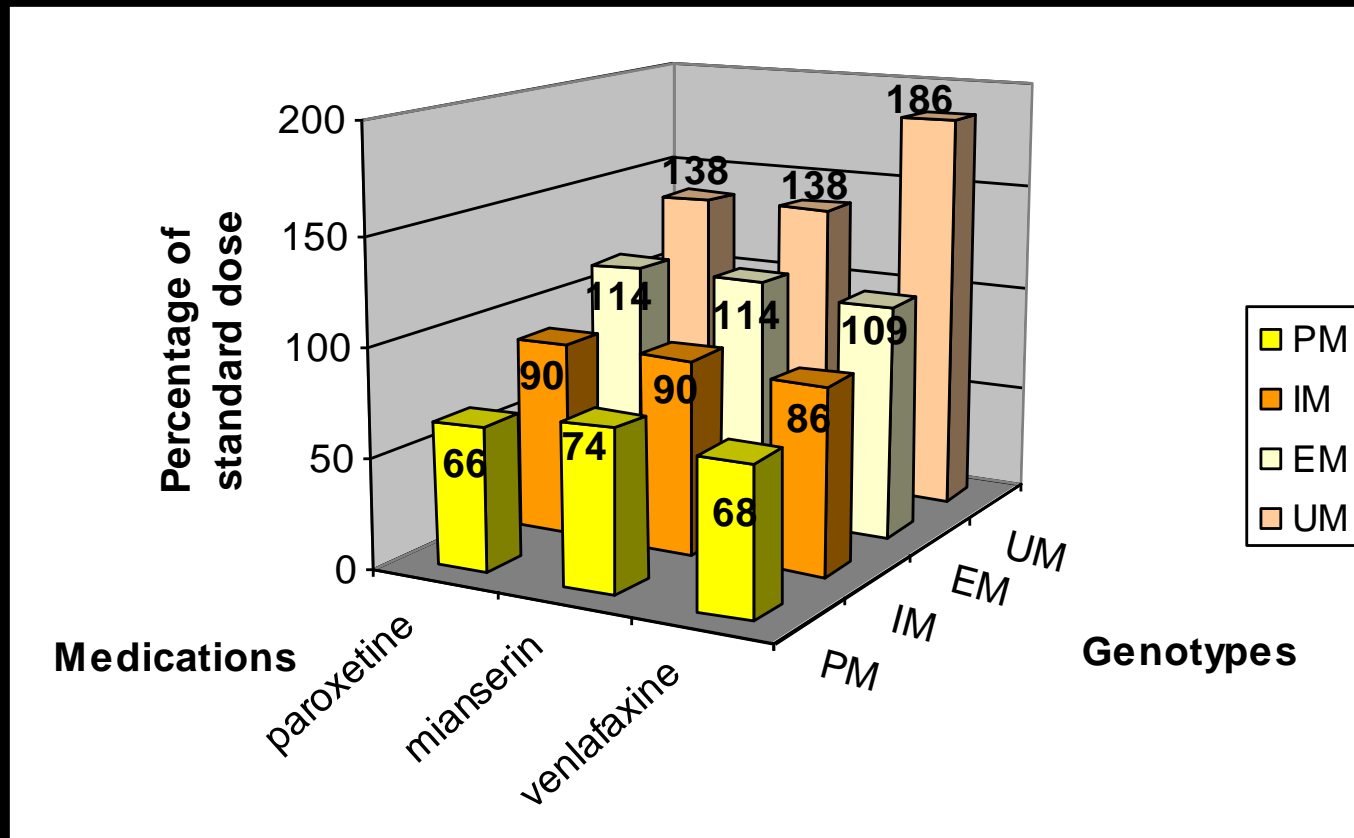


CYP 2D6 and tricyclic antidepressants

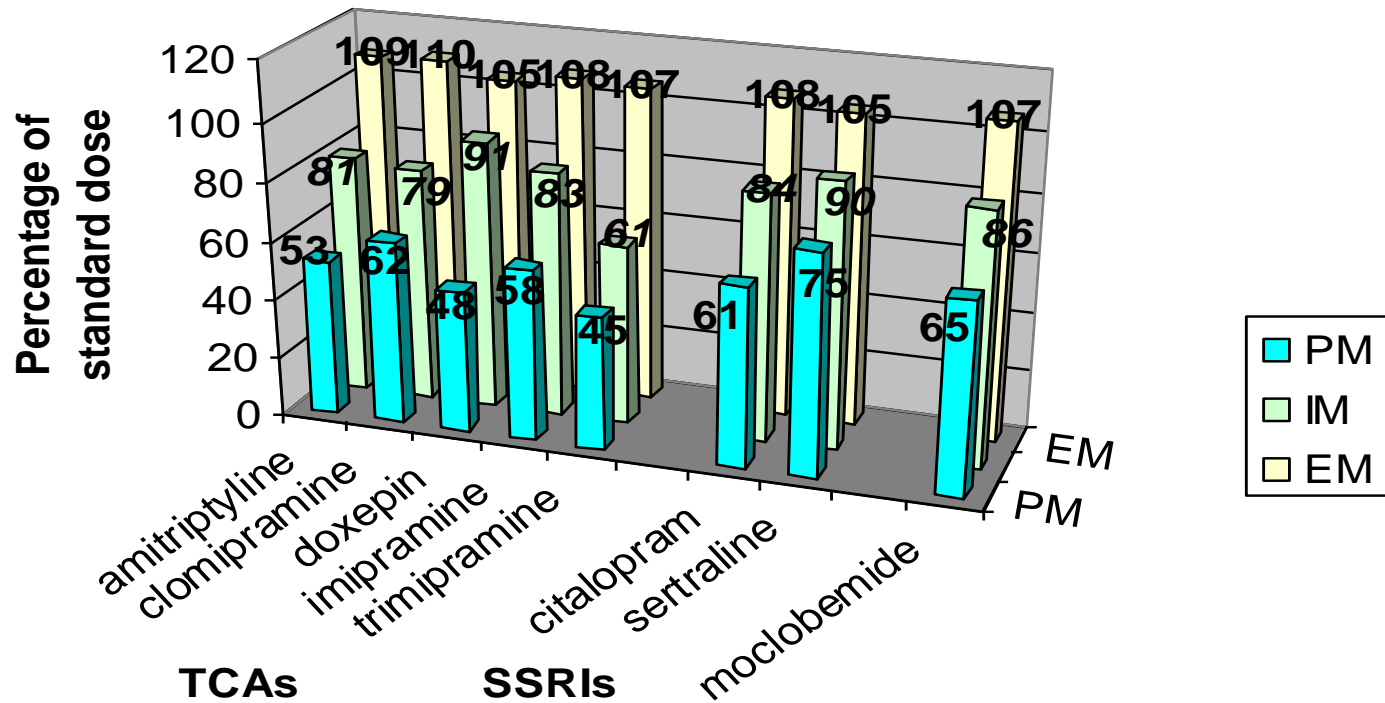


TCA dose adjustments are recommended for 2D6 PM and UM.

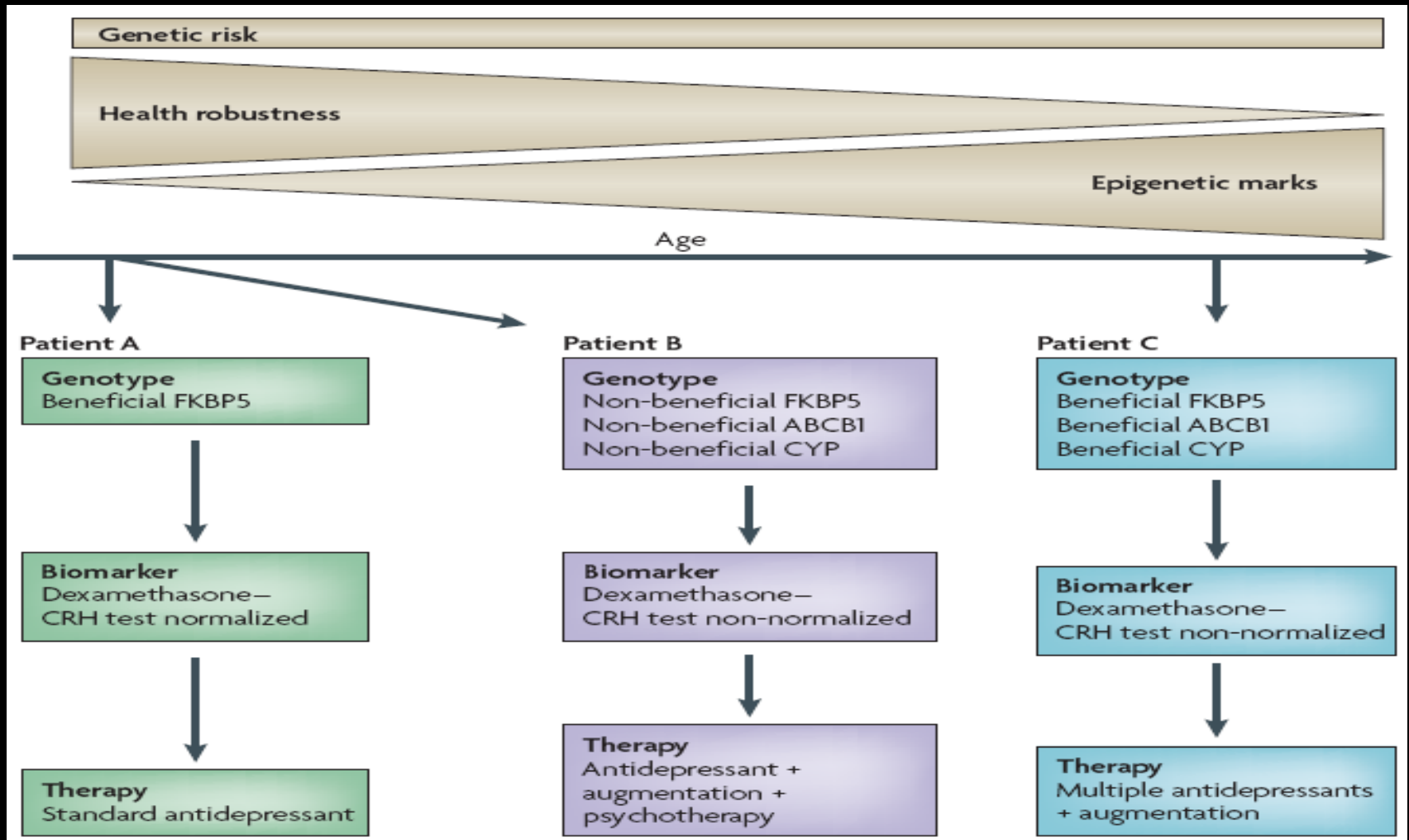
CYP 2D6 and other antidepressants



CYP 2C19 and antidepressants



Treatment strategies with biomarker and pharmacogenetics

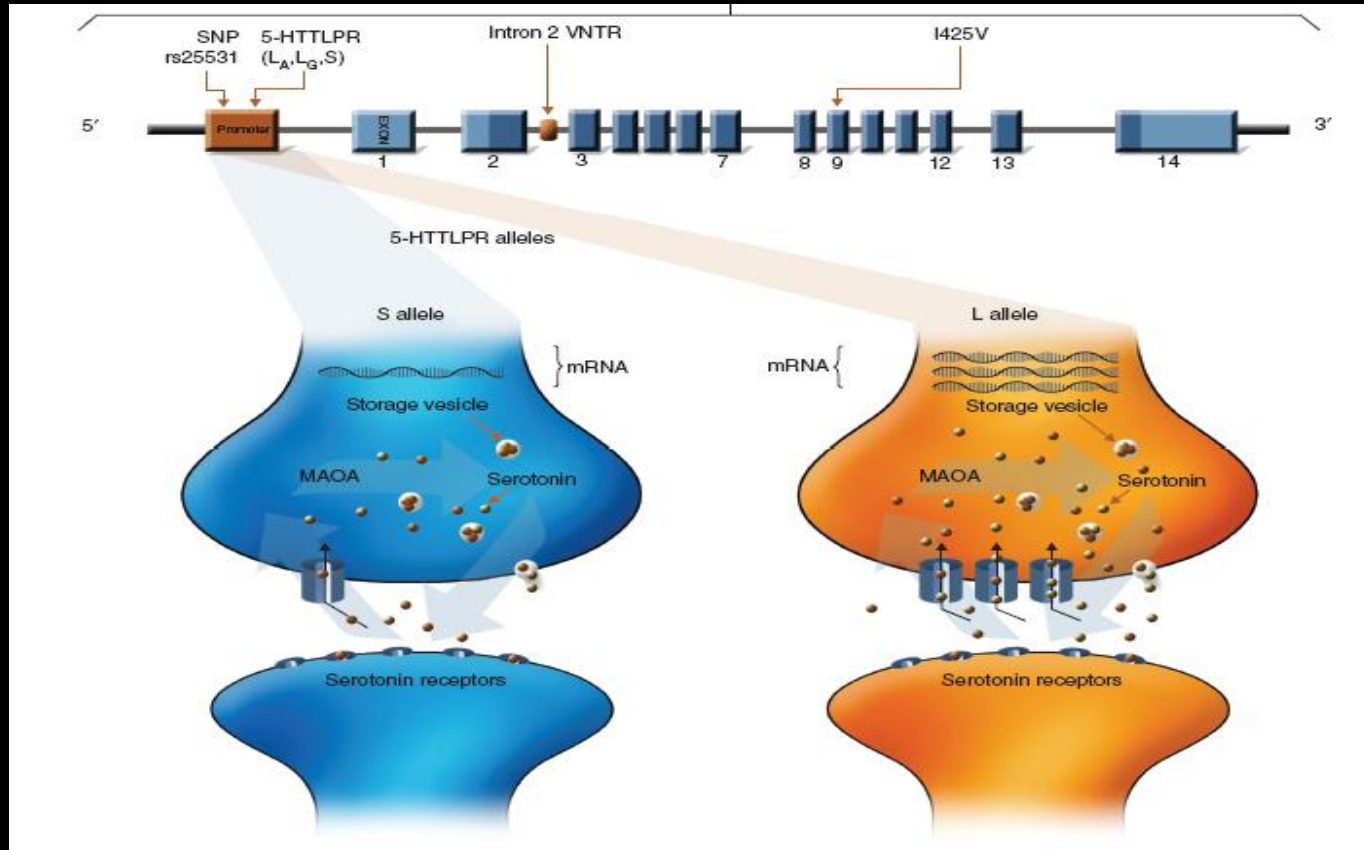


**Potential candidate
genes relating to
antidepressant
response**

Genes investigated

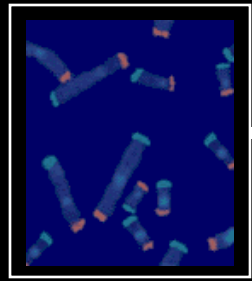
- HTTLPR
- SERT-STin2
- 5HT1A C-1019G
- 5-HT1B
- 5HT2A T102C
- 5HT2A G1438A
- 5HT2C
- 5HT6 C267T
- TPH1 A218C
- FKBP5
- NET T-182C
- NET G1287A
- COMT
- MAOA
- DRD2 S311C
- DRD4 VNTR
- ACE I/D polymorphism
- G-protein beta3 C825T
- ADRB1 G1165C
- CRHR1
- NOS C276T
- IL-1beta C511T
- CLOCK
- ◆ BDNF
- ◆ DTNBP1
- ◆ nNOS
- ◆ IL-1beta
- ◆ APOE
- ◆ MDR1P-gp
- ◆ GRIK4

Serotonin Transporter gene (SLC6A4)

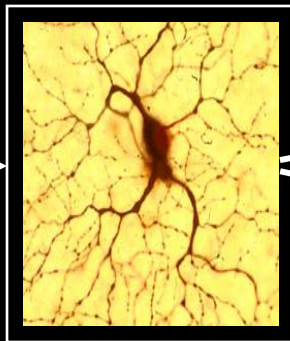


- Carriers of short allele have a poor outcome after treatment with SSRIs and a higher rate of adverse effects
- Antidepressant augmentation strategies with pindolol and lithium was beneficial to carriers of short allele

SLC6A4: How do we get there from here ?



SLC6A4:
5'HTTLPR
polymor
phism



Cells:
serotonin
mediated
excitability



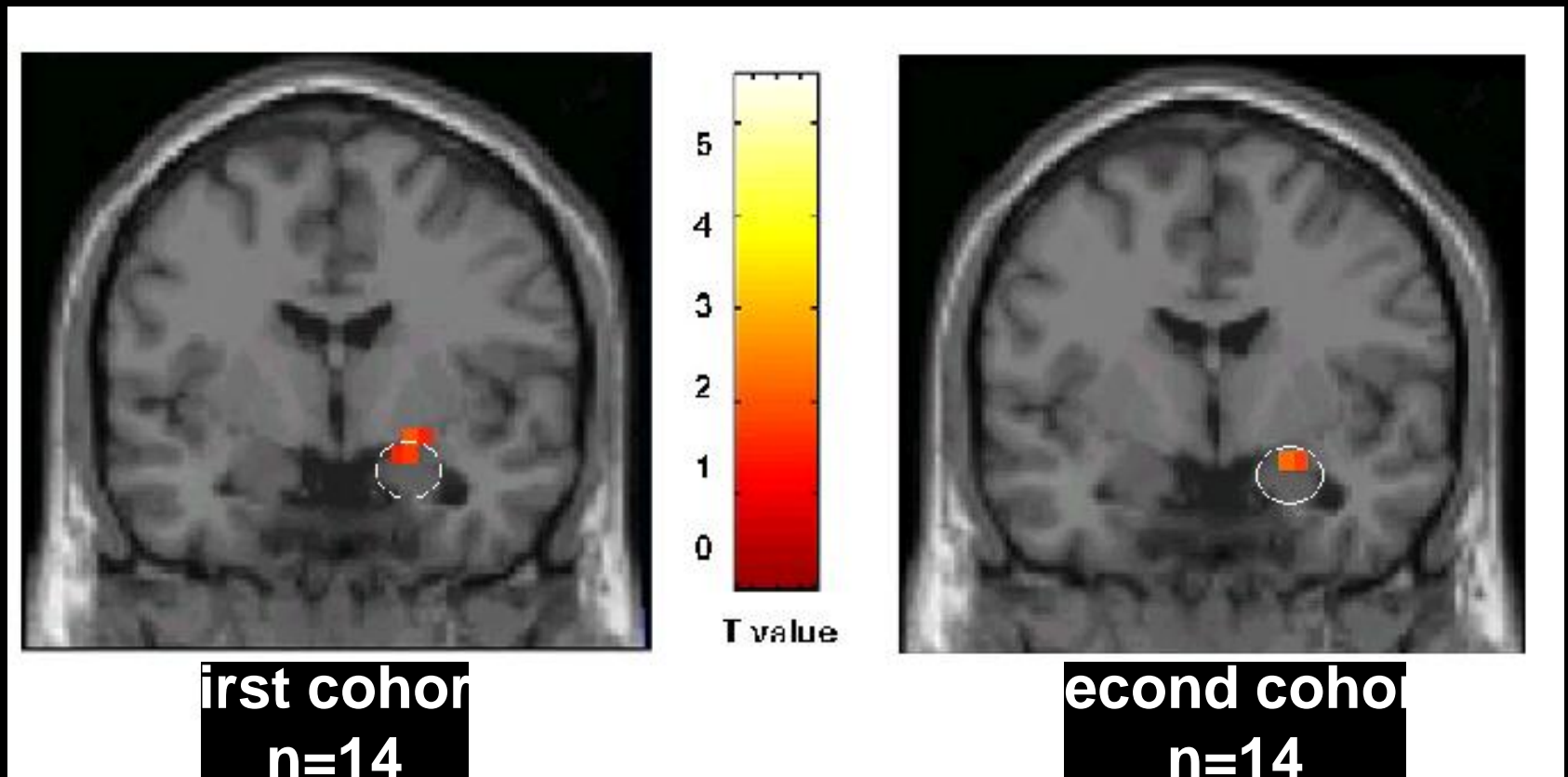
Systems:
amygdala
processing of
fearful stimuli

depression,
anxiety disorders,
neuroticism,
response to SSRIs,
substance abuse,
hallucinations

Behavior:
complex functional
interactions and
emergent phenomena



5'-HTTLPR genotype and fMRI during perceptual processing of fearful faces

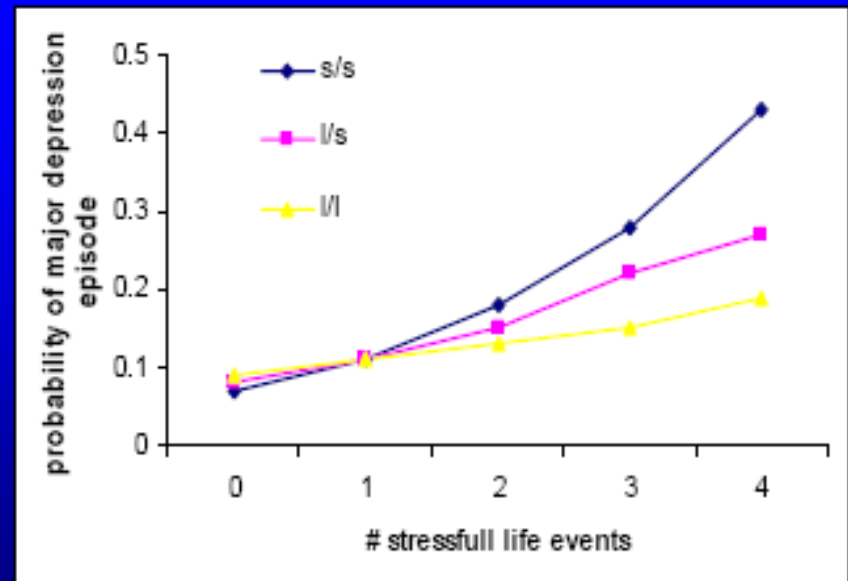


s allele carriers show a greater amygdala response than
// homozygous individuals

Genetic x environmental factors

SERT

Stressful life events and the number of short 5-HTTLPR alleles (l/l, l/s, or s/s) predicts occurrence of depression (Caspi et al, 2003)



5-HTTLPR variations.....

**Broad influence of a single gene
on a range of aspects**

Alteration of serotonin pathway plasticity

Anatomical change

Stress reactivity

Temperament

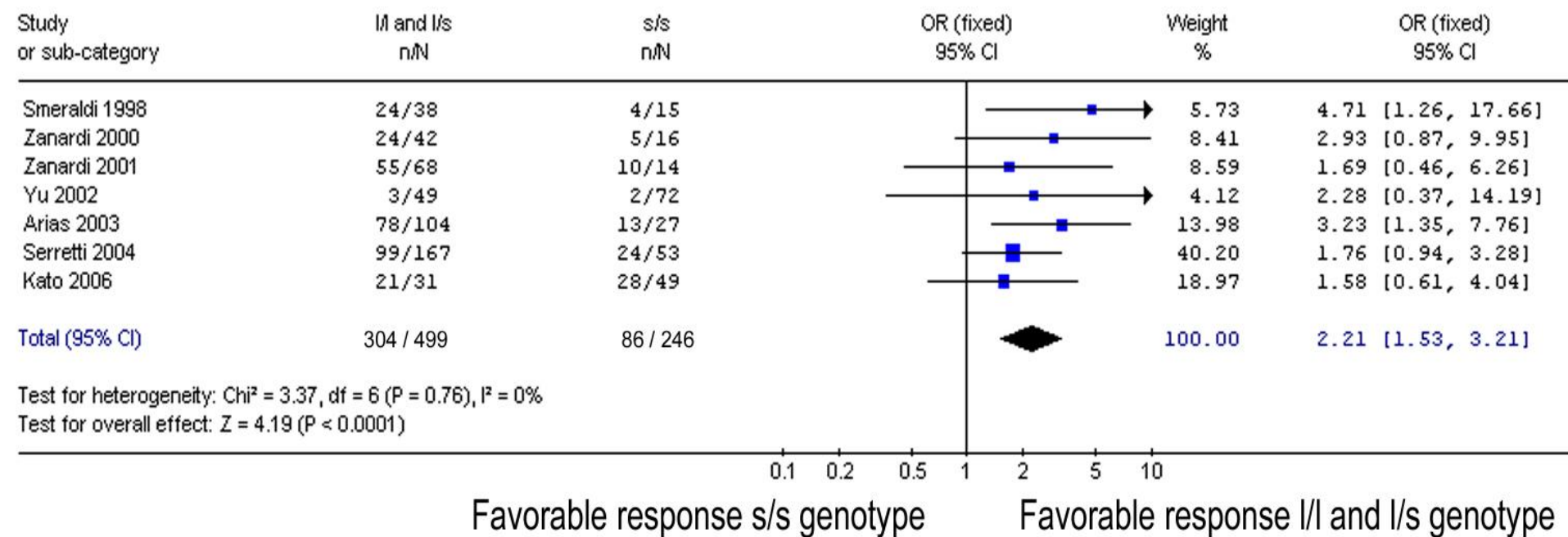
Response to antidepressants

Mood disorders

Meta-analysis for 5-HTTLPR

(n=15, 1435 subjects)

Remission



Late-Life Depression Demographics

- Community sample >age 65
 - 1% Major depression
 - 2% Dysthymia
 - 4% Adjustment Disorder with depressed mood
 - 15% Sub-syndromal Depression

Clinical Features of Late-life Depression

- “Depression” without sadness
- *Irritability*
- Prominent *Anxiety*
- Cognitive complaints
- Prominent vague *somatic complaints*
- Unexplained health worries
- Heightened pain complaints
- Loss of interest and pleasure
- Social withdrawal or avoidance of social interactions
- Multiple primary care visits without resolution of the problem
- Unexplained functional decline

Early-onset v. Late-onset

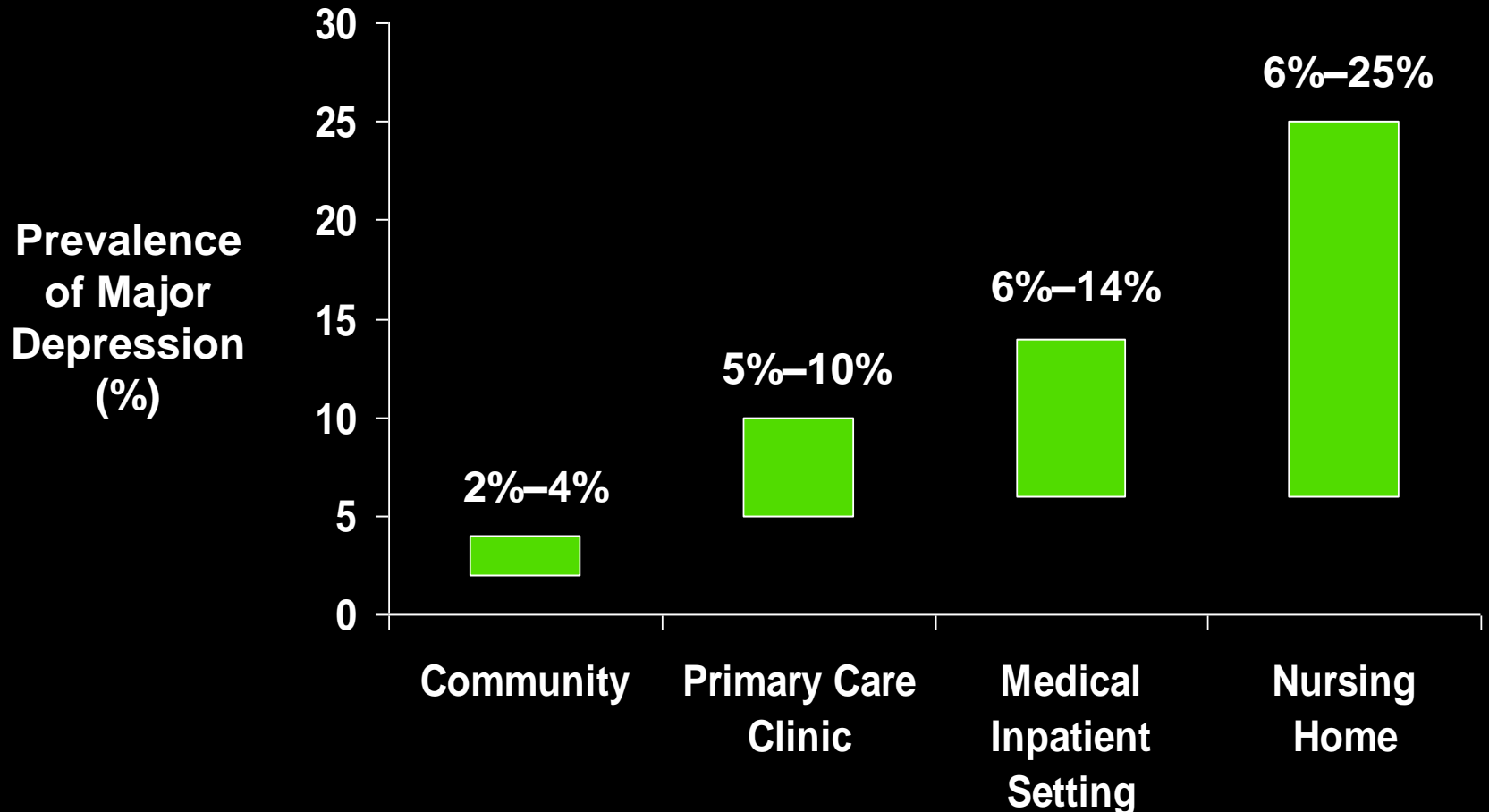
Early-onset

- Index episode in childhood or early adult life
- First degree relatives with depression
- Less physical illness
- More psychiatric comorbidity (SUD; personality disorders)
- Sad mood

Late-onset

- Index episode after age 50
- Less genetic predisposition
- **Chronic physical illness**
- Poorer treatment response with more chronic course
- Increased mortality
- Abnormal brain imaging
- Less psych comorbidity
- Apathy and anhedonia

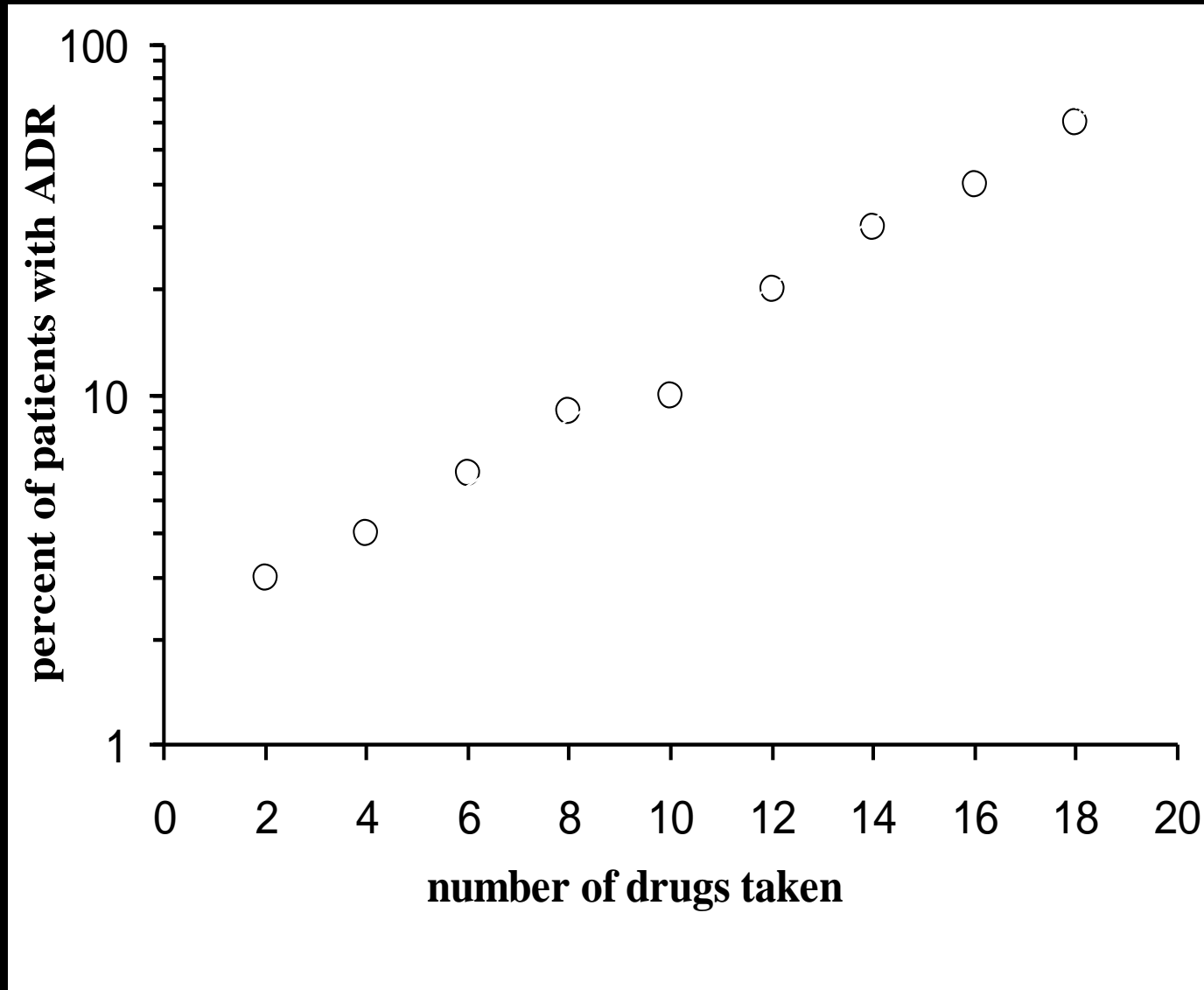
Major Depression Is Associated with Chronic Medical Illness



Katon W, Schulberg H. *Gen Hosp Psychiatry*. 1992;14:237-247.

Rosen J, Mulsant BH, Pollock BG. *Nursing Home Med*. 1997;5:156-165.

Medication and adverse reaction



Pharmacokinetic Changes in Aging

Parameter	Change	Effect
Absorption	Possible ↓	↓ Effectiveness
Protein binding	↓ if albumin is low	↑ free drug for protein bound
Volume of distribution	↑ for lipophilic	↑ accumulation
	↓ for hydrophilic?	↑ toxicity
Hepatic metabolism	↓ blood flow, 1 st pass, demethylation and hydroxylation	↑ accumulation ↑ toxicity ↓ prodrug activation
Renal excretion	Often ↓	↓ elimination

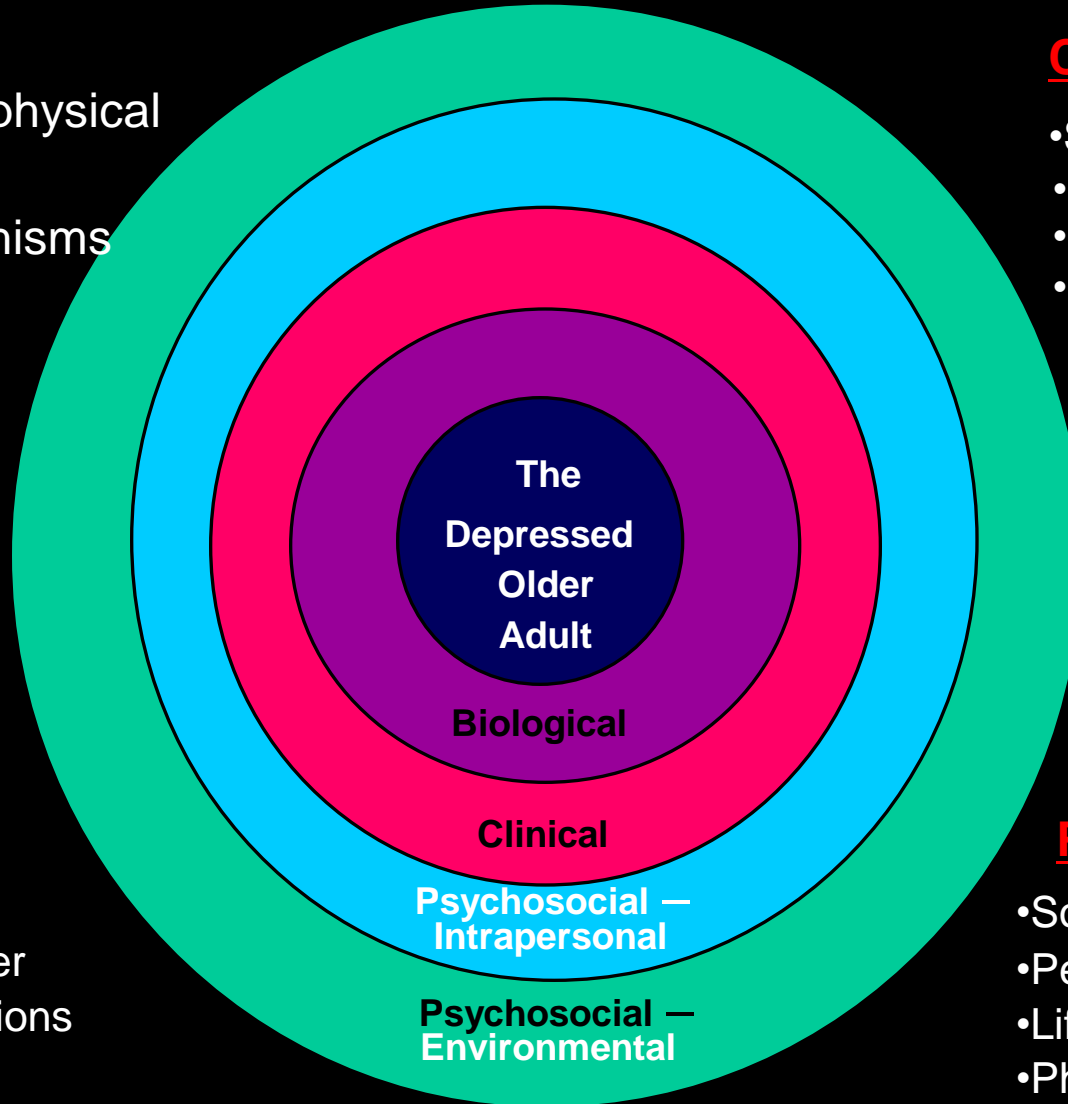
Nested Potential Predictors of Treatment Response in Late-Life Depression

Biological

- Nonpsychiatric physical illness
- Gene polymorphisms

Clinical

- Symptom severity
- Lifetime age of onset
- Comorbid anxiety
- Cognitive impairment



Psychosocial – Intrapersonal

- Demographics
- Personality disorder
- Traits and dispositions

Psychosocial

- Social supports
- Perceived chronic stress
- Life events/acute stress
- Physical environment

Individual pharmacogenetic trials

Monoamine transporter gene polymorphisms and antidepressant response in Koreans with late-life depression (mean age=60)

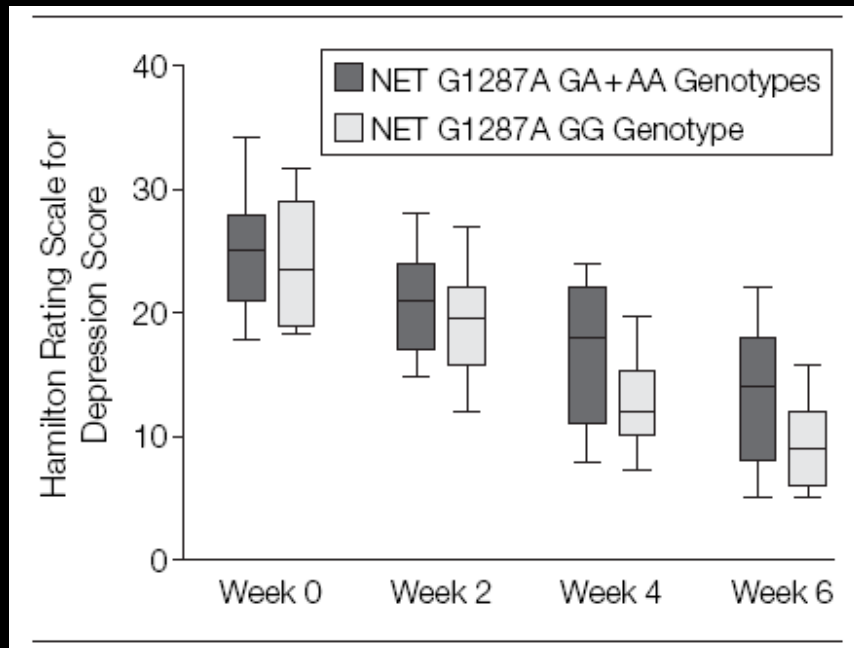
A 6-week naturalistic treatment study with blinded outcome evaluation of 241 Korean inpatients and outpatients with major depression

Treatment with an SSRI (fluoxetine or sertraline; n = 136) or an NRI (nortriptyline; n = 105) antidepressant. Adherence was assessed by measuring plasma concentration at 4 weeks

An SSRI and NRI response (defined as $\geq 50\%$ decrease in Hamilton Rating Scale for Depression score at 6 weeks)

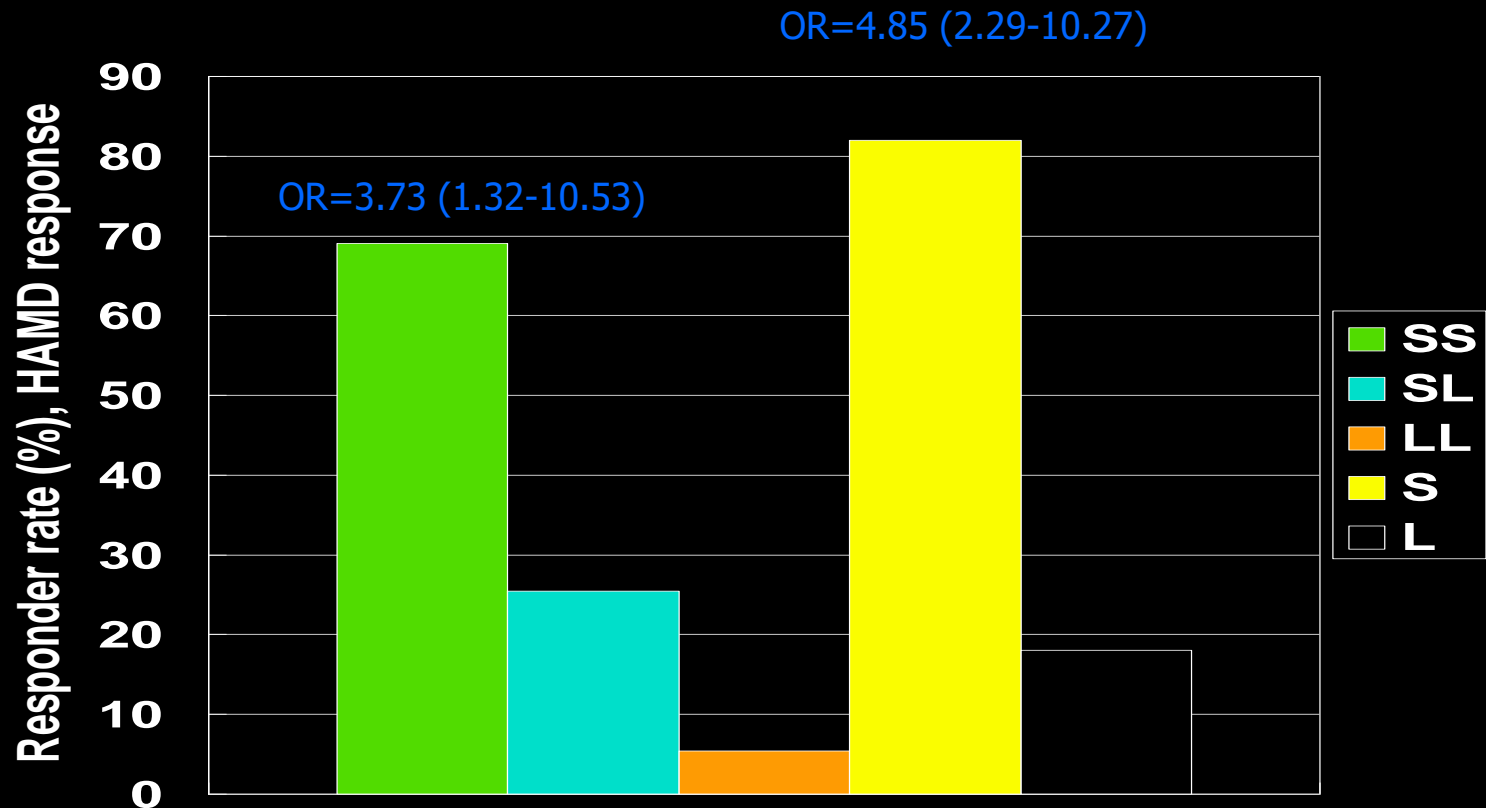
5-HTTLPR, 5-HTT VNTR in intron 2 and NET G1287A in exon 9

Changes in HAMD and responder rate: NET and NRI

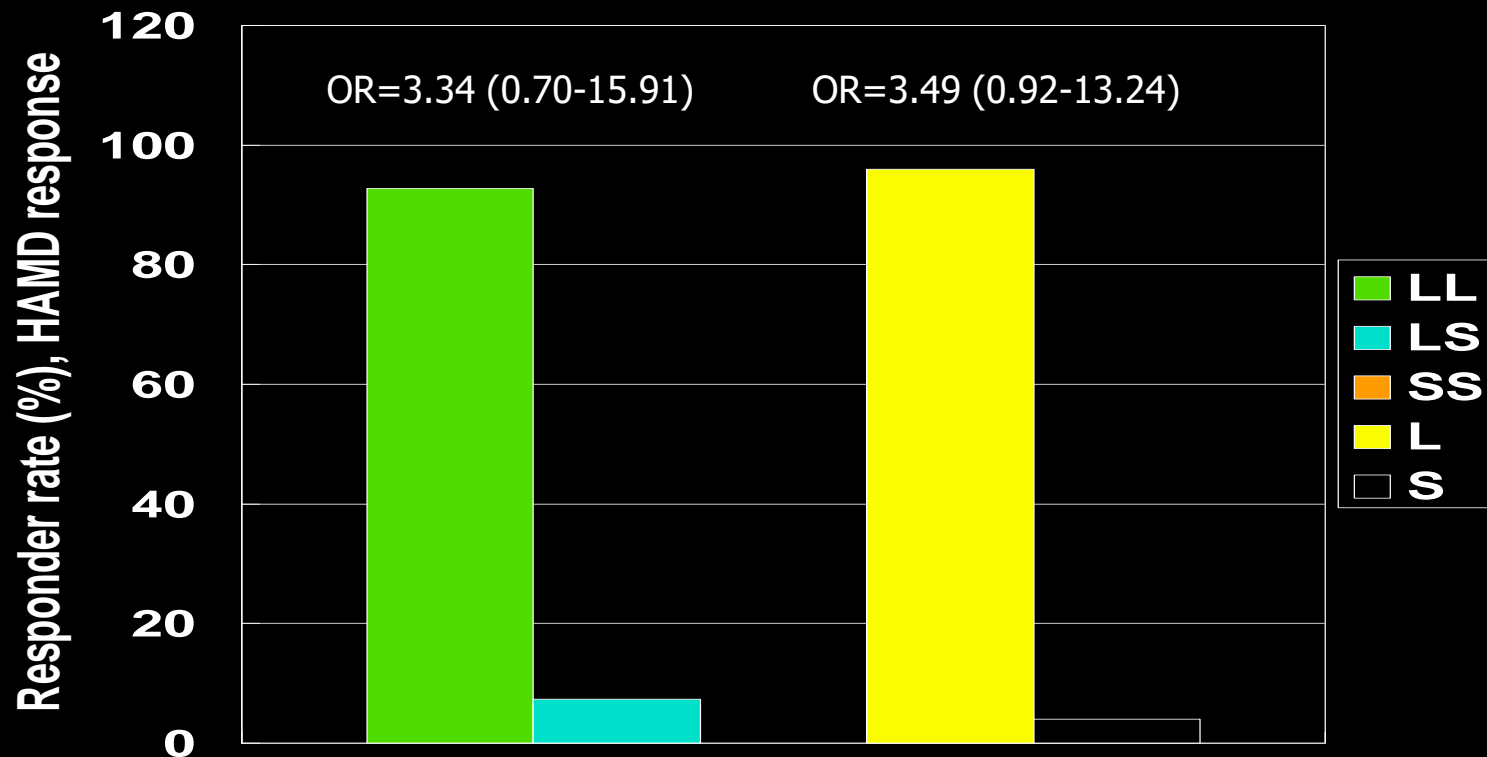


	Responder (%)	OR and p values
GG	63.6	7.54 (2.53-22.49)
GA	29.1	
AA	7.3	
G	78	3.48 (1.67-7.30)
A	22	

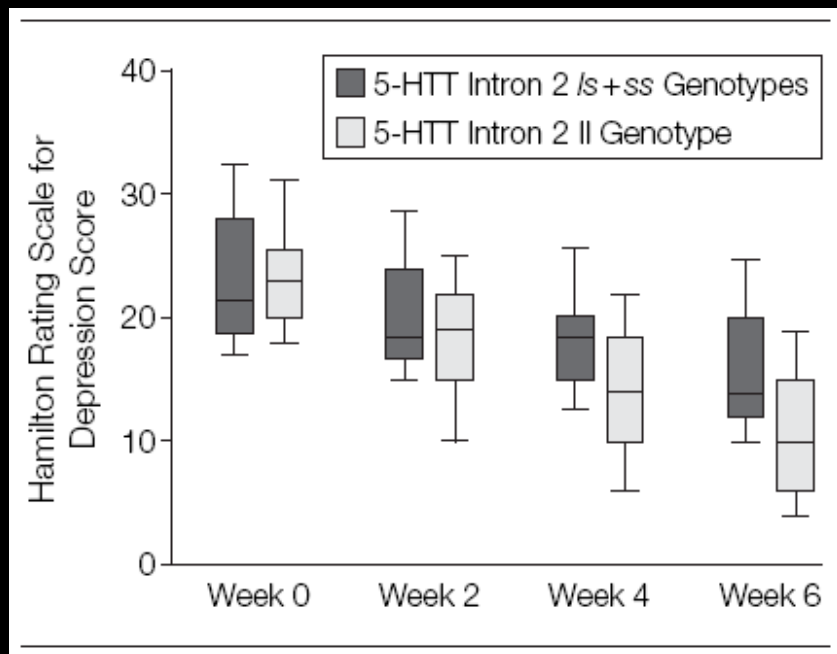
Responder rate: 5-HTTLPR and NRI



Responder rate: 5-HTT intron and NRI

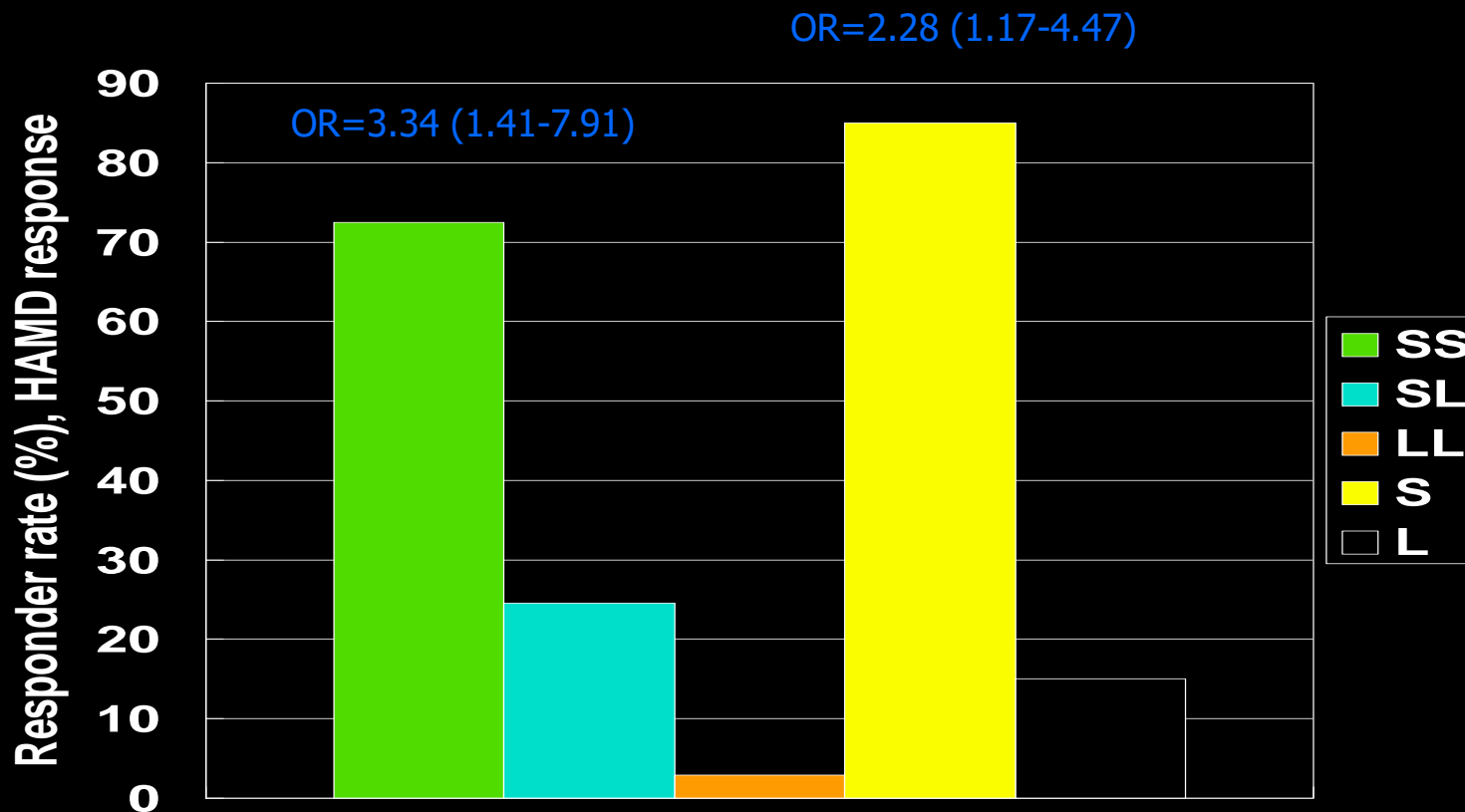


Changes in HAMD and responder rate: 5-HTT intron and SSRI

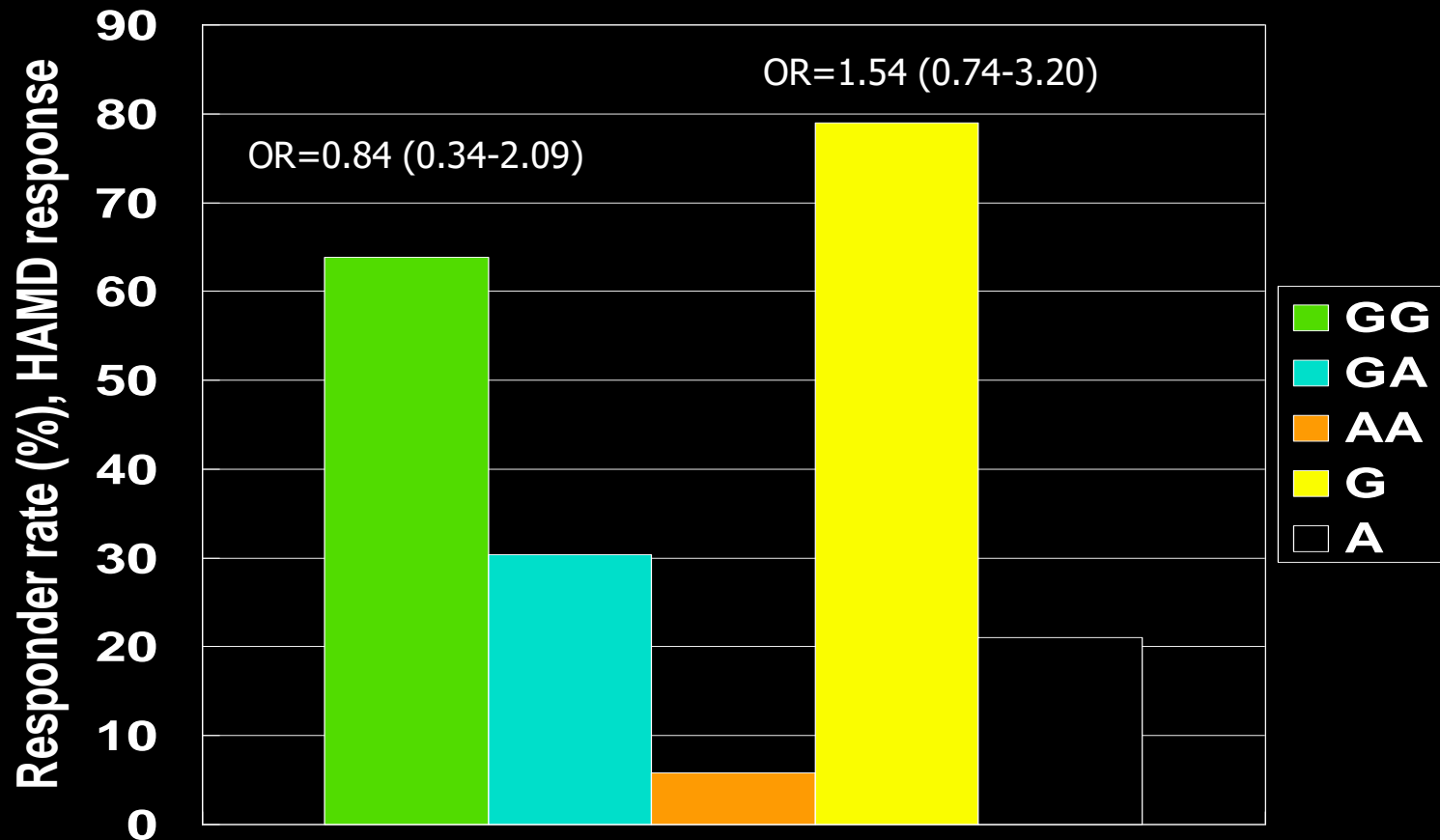


	Responder (%)	OR and p values
LL	97.1	20.11 (4.27-94.74)
LS	2.9	
SS	0	
L	99	15.87 (3.47-71.43)
S	1	

Responder rate: 5-HTTLPR and SSRI



Responder rate: NET and SSRI



Response Rates With Combinations of Monoamine Transporter Polymorphisms

NET G1287A	5-TTLPR	5-HTT Intron 2	Response Rate, No./Total (%)	P Value*
Norepinephrine Reuptake Inhibitor				
GG	ss	Any genotype	23/26 (88.5)	<.001
GG	I Carrier	Any genotype	12/16 (75.0)	<.008
A carrier	ss	Any genotype	15/24 (62.5)	<0.02
A carrier	I Carrier	Any genotype	5/23 (21.7)	Comparator
Selective Serotonin Reuptake Inhibitor				
Any genotype	ss	II	48/62 (77.4)	Comparator
Any genotype	I Carrier	II	19/35 (54.3)	<0.06
Any genotype	ss	s Allele carriers	2/8 (25.0)	<0.01
Any genotype	I Carrier	s Allele carriers	0/14	<.001

STAR*D, mean age=43

- 1,953 patients with major depressive disorder who were treated with the antidepressant citalopram in the Sequenced Treatment Alternatives for Depression (STAR*D) study and were prospectively assessed
- 68 candidate genes were genotyped, with 768 single-nucleotide-polymorphism markers chosen to detect common genetic variation
- significant and reproducible association between treatment outcome and a marker in *HTR2A*. *no evidence of association among any of the four genotyped SLC6A4 markers and treatment outcome in these data*

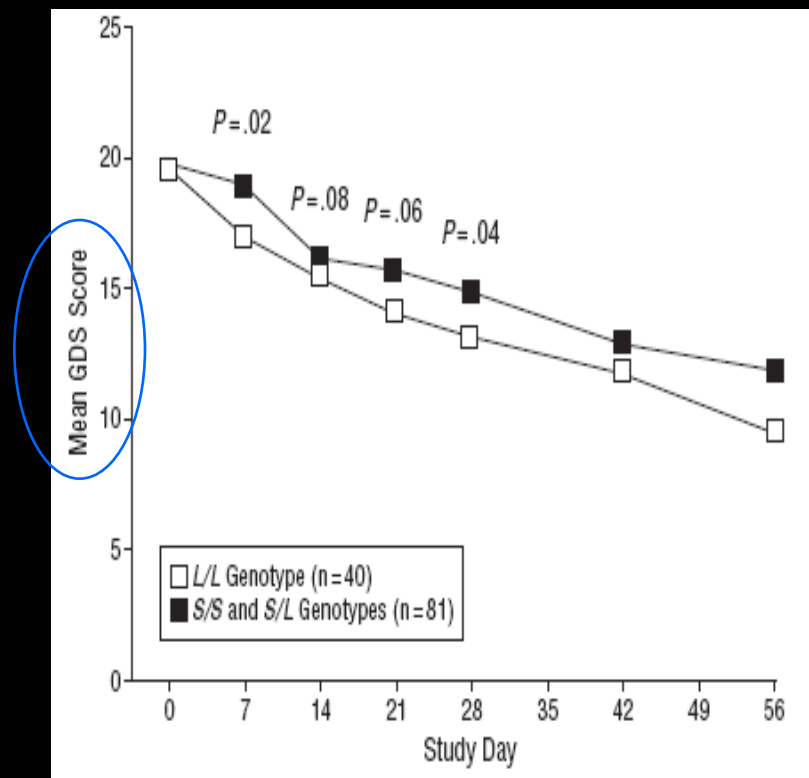
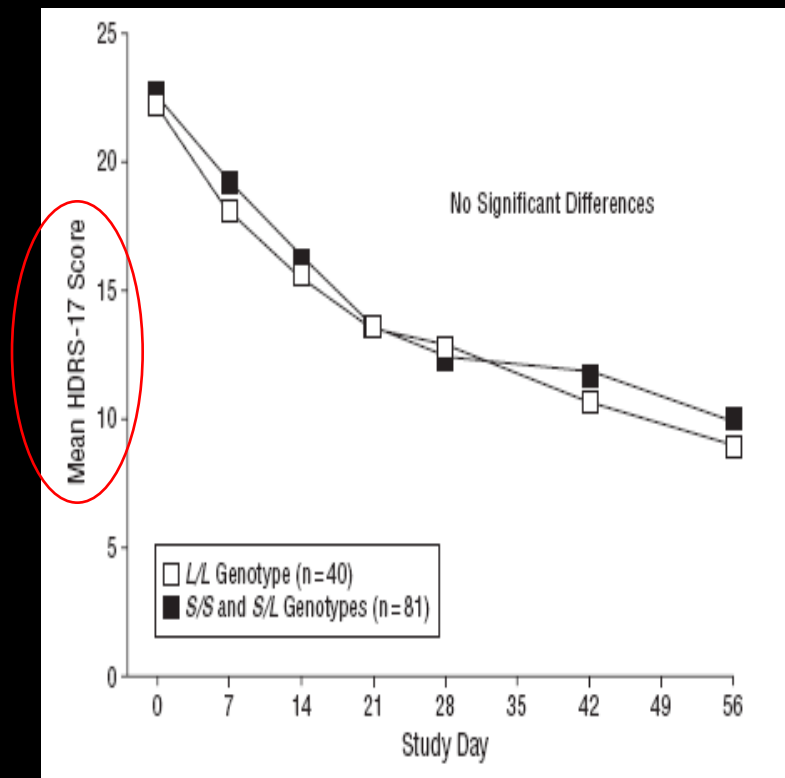
Association Analysis of Genotyped *HTR2A* SNPs, Stratified by Race

PHENOTYPE AND SNP	ALL			WHITE			BLACK		
	N	P		n	P		n	P	
		Allelewise	Genotypewise		Allelewise	Genotypewise		Allelewise	Genotypewise
Remission:									
<i>rs7997012</i>	1,149	.000024	.000035	911	.00107	.00183	170	NS	NS
<i>rs1928040</i>	1,148	.0446	.0701	910	.0626	NS	170	NS	NS
<i>rs6313</i>	1,183	NS	NS	942	NS	NS	172	NS	NS
<i>rs6311</i>	1,180	NS	NS	939	NS	NS	172	.0431	.0874
Response:									
<i>rs7997012</i>	1,329	.000037	.000002	1,049	.00183	.000157	199	NS	NS
<i>rs1928040</i>	1,327	.0709	NS	1,048	NS	NS	199	NS	NS
<i>rs6313</i>	1,372	NS	NS	1,086	NS	NS	202	NS	NS
<i>rs6311</i>	1,371	NS	NS	1,084	NS	NS	203	.0918	.0149
Change in QIDS-C ₁₆ :									
<i>rs7997012</i>	1,749	.000007	.00000146	1,380	.00123	.000516	261	NS	NS
<i>rs1928040</i>	1,747	.0214	.0072	1,378	.0738	.0887	261	NS	NS
<i>rs6313</i>	1,802	NS	.0878	1,425	NS	NS	264	NS	.0353
<i>rs6311</i>	1,804	.0599	.0494	1,426	NS	NS	265	.0094	.0261

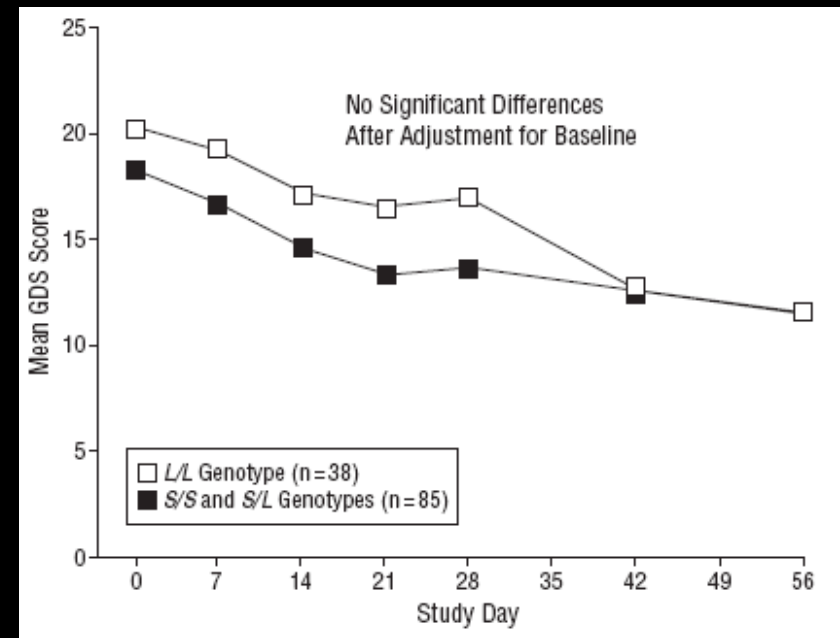
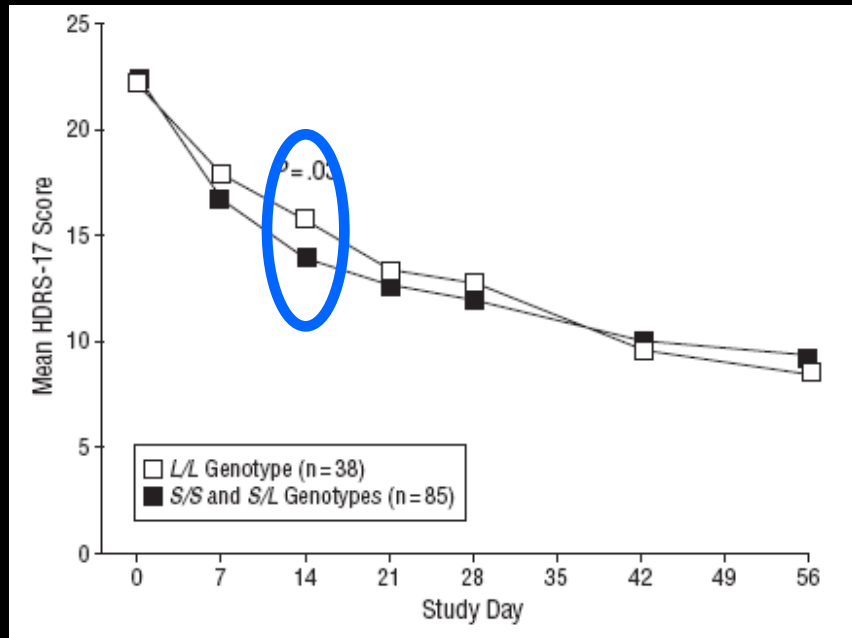
Association Results for SLC6A4 and Citalopram Response.

Marker	White (<i>n</i> = 799 vs. 509)				African American (<i>n</i> = 130 vs. 121)			
	MAF		Additive model <i>p</i> value	Dominant OR (95% CI)	MAF		Additive model <i>p</i> value	Dominant OR (95% CI)
	NonResp	Resp			NonResp	Resp		
rs25531	.07	.08	.22	1.25 (.90,1.73)	.25	.27	.54	1.19 (.72,1.96)
5-HTTLPR	.44	.42	.27	.89 (.70,1.13)	.20	.22	.60	1.19 (.71,2.01)
rs25533	.06	.06	.48	1.15 (.80,1.64)	.07	.13	.05	1.81 (.92,3.56)
rs2020933	.05	.05	.83	1.06 (.71,1.58)	.32	.36	.38	1.27 (.74,2.19)
rs2020934	.46	.49	.10	1.31 (.99,1.72)	.23	.18	.28	.68 (.37,1.25)
rs16965628	.06	.07	.66	1.10 (.77,1.57)	.33	.35	.63	.96 (.56,1.65)
rs2066713	.38	.42	.09	1.29 (1.00,1.67)	.26	.29	.42	1.41 (.82,2.41)
rs6354	.20	.20	.95	1.03 (.81,1.32)	.34	.34	.90	1.07 (.62,1.82)
rs140700	.07	.10	.03	1.48 (1.04,2.10)	.04	.09	.02	2.57 (1.07,6.19)
rs140701	.44	.42	.42	1.02 (.80,1.31)	.27	.28	.95	.97 (.56,1.68)
rs1042173	.45	.43	.28	.90 (.70,1.16)	.23	.22	.82	.94 (.55,1.61)

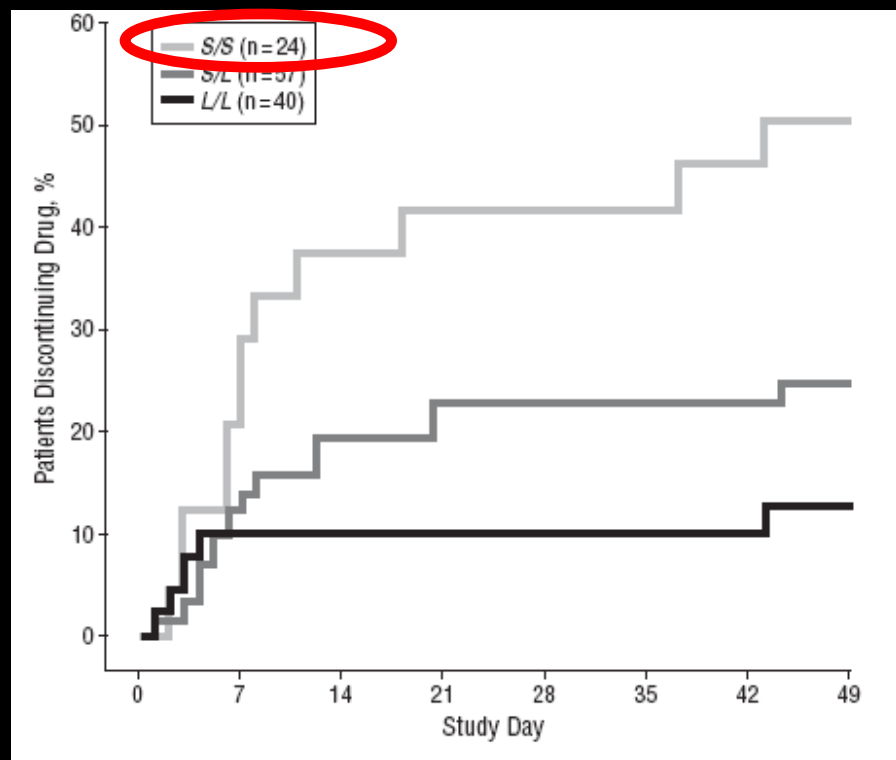
Effects of the 5HTTLPR polymorphism on the efficacy of paroxetine hydrochloride (mean age=72)



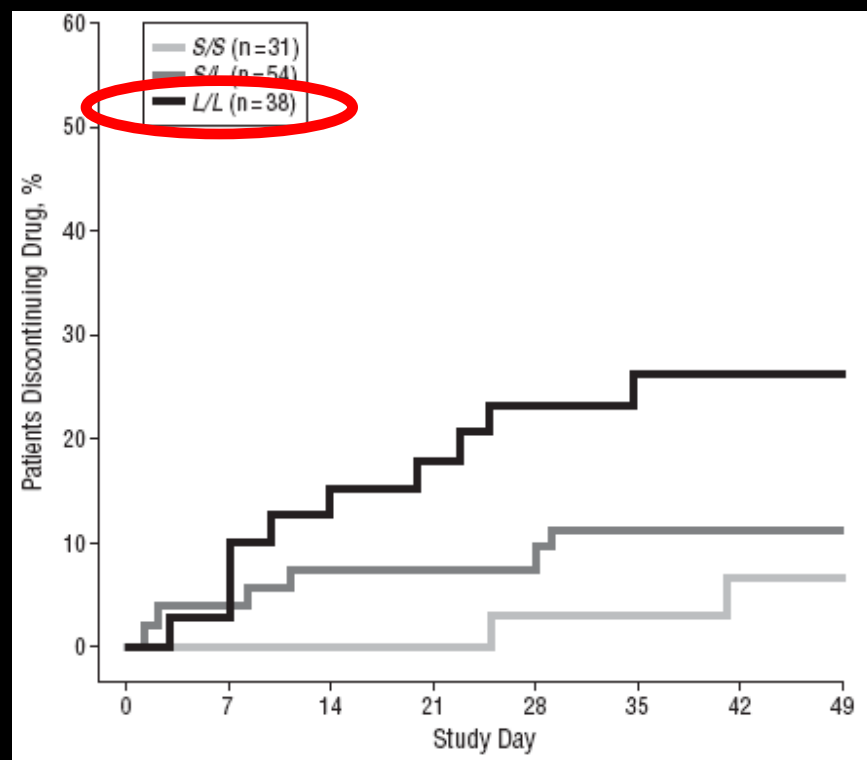
Effects of the *5HTTLPR* polymorphism on the efficacy of mirtazapine



Survival curves showing discontinuations due to adverse events

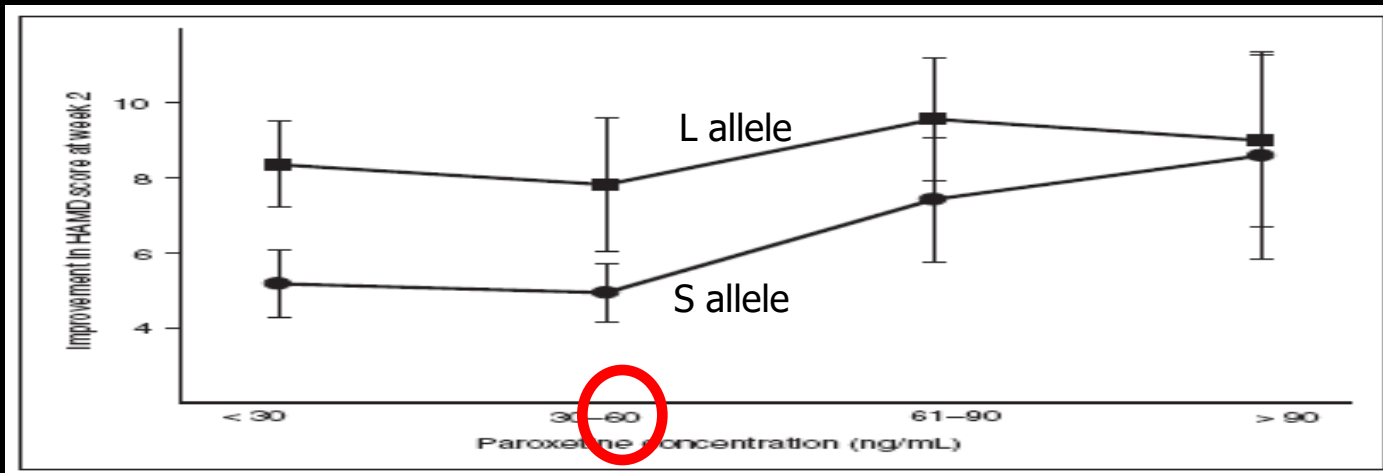


Paroxetine



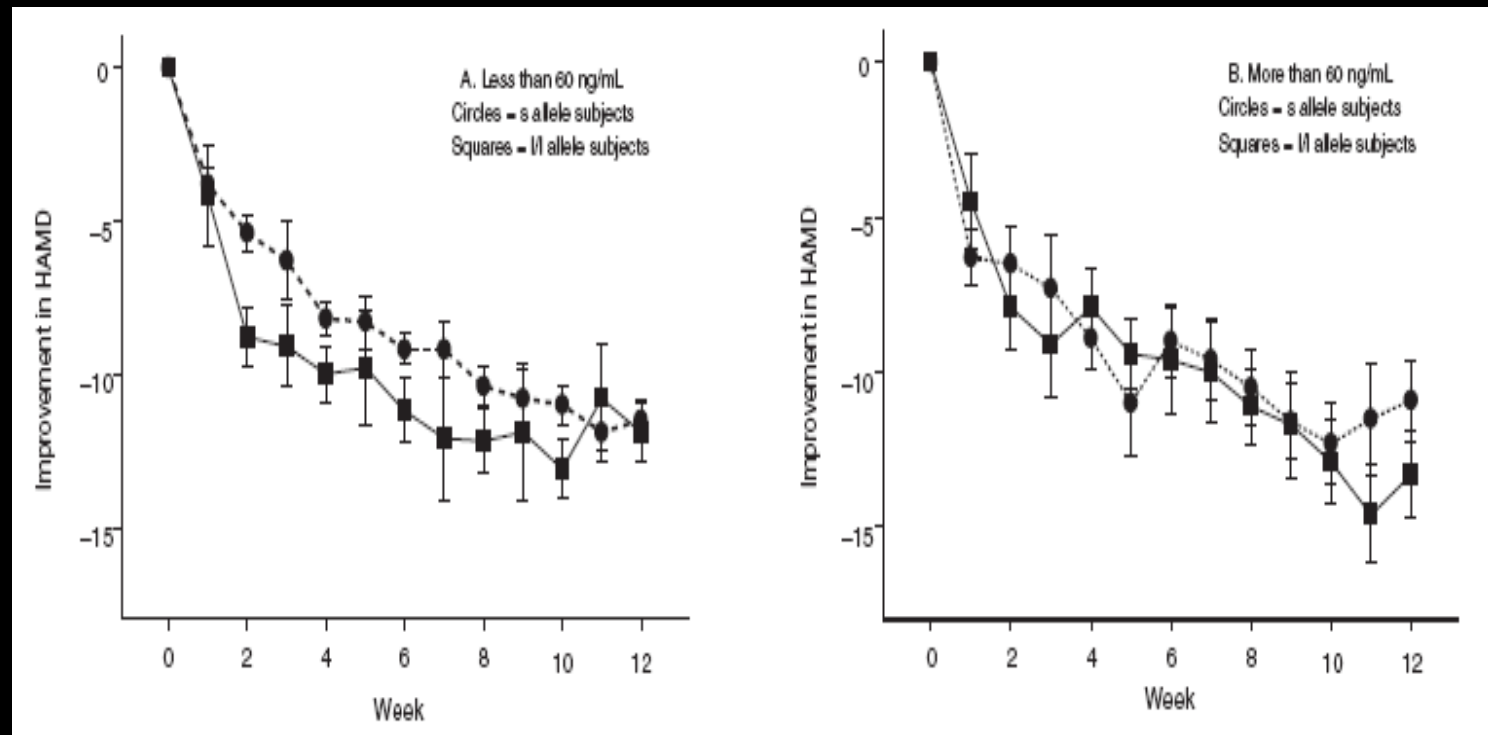
Mirtazapine

Serotonin transporter genotype interacts with paroxetine plasma levels to influence depression treatment response in geriatric patients



Subject group	Period; mean (and SEM)							
	Week 2 or 3		Week 4		Week 6		Week 10	
	Dose, mg	Level, ng/mL	Dose, mg	Level, ng/mL	Dose, mg	Level, ng/mL	Dose, mg	Level, ng/mL
l/l allele (n = 21)	16.1 (2.0)	43.9 (11.9)	21.8 (2.5)	77.9 (7.21)	24.5 (2.7)	89.7 (24.2)	30.0 (6.3)	154.6 (59.5)
s allele (n = 42)	19.2 (1.0)	55.6 (7.1)	23.5 (1.2)	100.8 (6.9)	26.4 (1.7)	110.7 (18.4)	36.7 (3.3)	185.7 (83.2)
Low exposure (n = 40)	17.7 (1.2)	28.8 (3.6)	20.9 (1.5)	46.6 (8.4)	23.5 (2.1)	62.8 (9.9)	25.0 (3.8)	92.4 (18.3)
High exposure (n = 23)	21.7 (2.1)	101.3 (10.1)	27.0 (2.1)	165.5 (20.5)	31.1 (2.6)	203.9 (21.2)	30.1 (3.7)	200.0 (29.6)

The interaction between genotype and nondichotomized paroxetine levels was significant

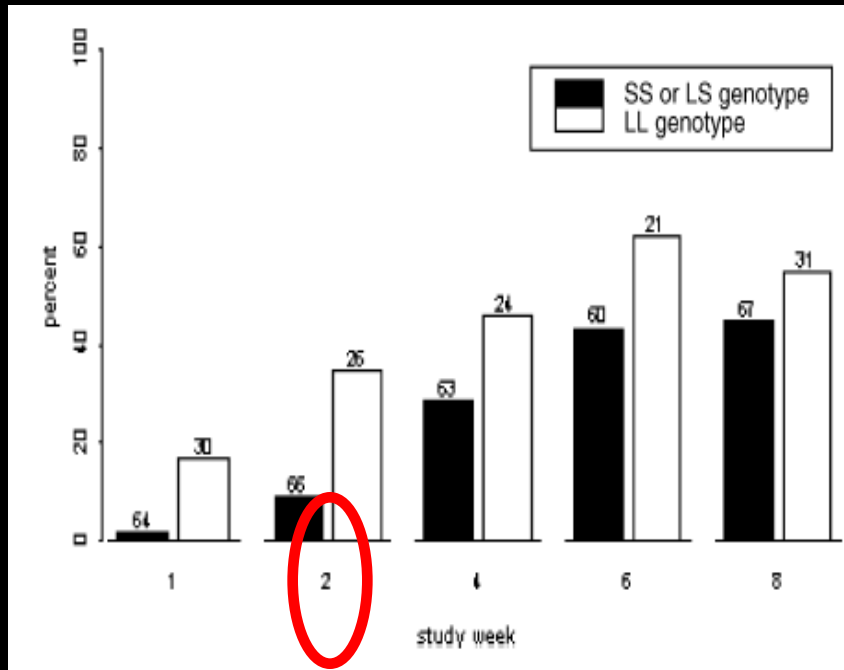


SERT and Remission: STAR*D (mean age=43)

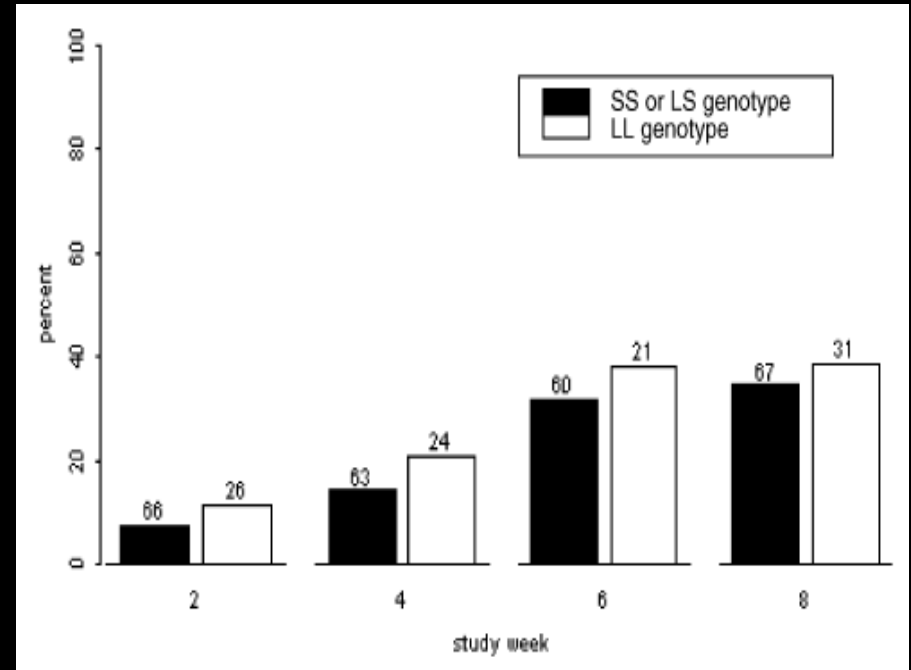
	White non-Hispanic				White Hispanic				Black			
	Remission	N	P-value ^a	P-value ^b	Remission	N	P-value ^a	P-value ^b	Remission	N	P-value ^a	P-value ^b
Intron 2 VNTR												
9/10	52.9%	17	0.810	1.000	0%	1	1.000	1.000				
9/12	70.6%	17	0.088	0.353	100%	1	0.374	1.000				
10/10	55.7%	158	0.070	0.282	29.2%	24	0.500	1.000	35.3%	17	1.000	1.000
10/12	49.9%	469	0.810	1.000	38.5%	78	0.880	1.000	37.9%	95	1.000	1.000
12/12	44.1%	81	0.017	0.069	38.5%	91	0.882	1.000	37.9%	116	1.000	1.000
Global P-value ^c		0.041				0.670				1.000		
Indel promoter												
L/L	53.7%	169	0.012	0.024	31.7%	60	0.267	0.533	37.8%	143	0.780	1.000
L/S	45.2%	504	0.038	0.115	39.8%	83	0.656	1.000	41.6%	77	0.567	1.000
S/S	46.2%	195	0.527	1.000	41.5%	53	0.512	1.000	30%	10	0.744	1.000
Global P-value ^c		0.039				0.490				0.787		
rs25531												
A/A	47.6%	925	0.134	0.267	38.5%	174	0.644	1.000	38.9%	126	0.893	1.000
A/G	54.5%	145	0.129	0.259	35%	20	1.000	1.000	38.3%	94	1.000	1.000
G/G	50%	4	1.000	1.000	0%	2	0.528	1.000	33.3%	12	1.000	1.000
Global P-value ^c		0.287				0.752				0.975		

Haplotype	Haplotype frequency			Haplotype simulation P-value	Maximum statistic simulation P
	WNH	WNH non-remitters	WNH remitters		
S-a-12	0.330	0.363	0.297	0.0007	0.0031
L-a-12	0.215	0.215	0.213	0.98	
L-g-12	0.052	0.048	0.057	0.32	
S-a-10	0.085	0.074	0.097	0.23	
L-a-10	0.291	0.279	0.304	0.14	

The serotonin transporter polymorphism, 5HTTLPR, is associated with a faster response time to sertraline in an elderly population (mean age=69)

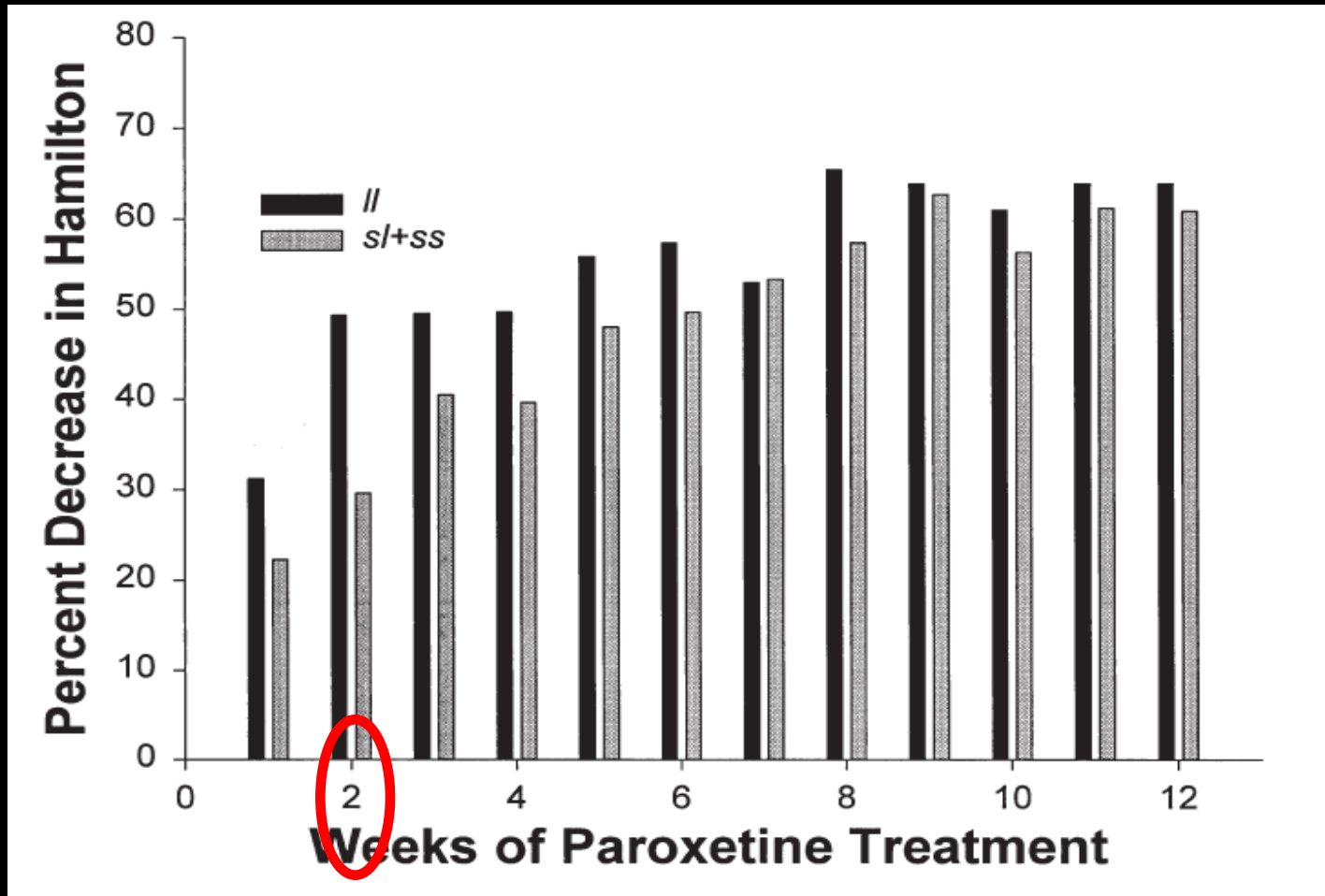


The percentage of responders (CGI-I=2 or <2)



The percentage of responders (50% or greater reduction in HAM-D)

Serotonin Transporter Promoter Affects Onset of Paroxetine Treatment Response in Late-Life Depression



Limitations to current pharmacogenetics studies

- Generally not multi-gene studies (or studies considering combinations of several genes)
- Little explanation of treatment variance by multiple small effect genes
- Statistically significant results are not necessarily clinically meaningful
- Many studies - few results replicated
- Gene-environment, gene-disease, gender, age and other hidden factors not controlled
- Candidate polymorphisms often associated with baseline disease severity
- Small samples
- Sensitivity of rating scales and response definition
- Ethnic difference in SNPs

Pharmacogenetics: problematic issues...and possible solutions

- Low variance explained by polymorphisms (HTTLPR=2.8%, TPH=2.7%, Gβ3=1.2%) → Other variables influence drug response: Life events, social support, temperament, hormones...*and should be included in the model! Neural Network?*
- Epigenetic factors, CNV, Splicing, Regional expression, gene interactions...*should be controlled with multivariate or neural network models.*
- Drug response may differ across episodes...*longer follow up*

Now and Outlook

- Genotyping and SSRI plasma concentrations can be recommended clinically
- How predictive is antidepressants' efficacy and toxicity for clinical endpoints?
- Drug target: transporter, Rs, signaling pathway
- Will genetic testing to predict response and toxicity be feasible and cost-effective?
 - Maybe, but expectations are probably too high
- Large studies with many genes are needed