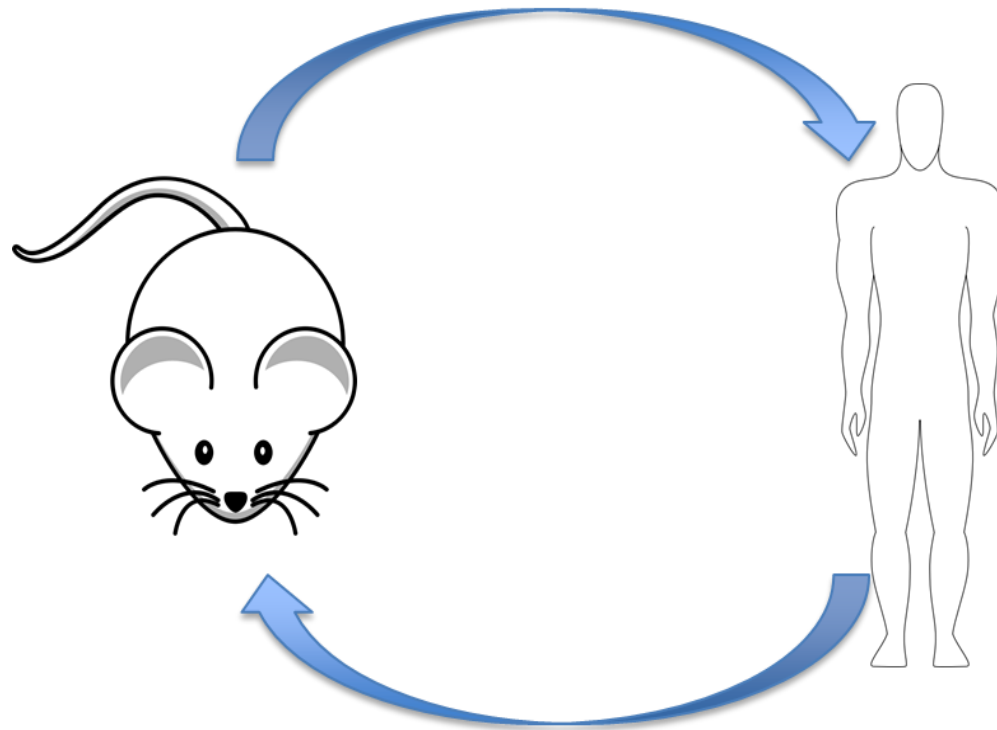


Epigenetic Programming by Early-Life Stress: Translational approaches from Rodent to Human and back

Chris Murgatroyd

Manchester Metropolitan University

Translational animal models



Animal models for stress: testing/developing hypotheses

Control for environment and genetics



Generation of different animal model paradigms

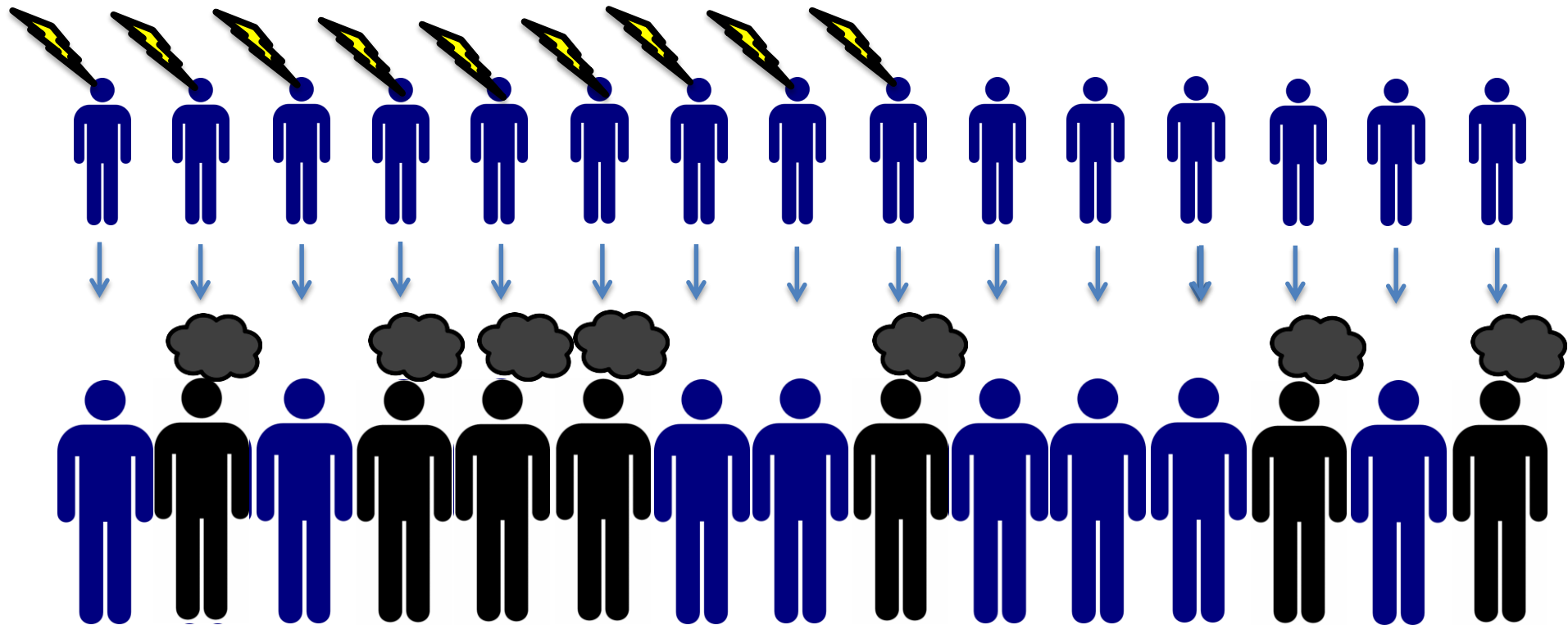
Different tests

Access to tissues, e.g. brain

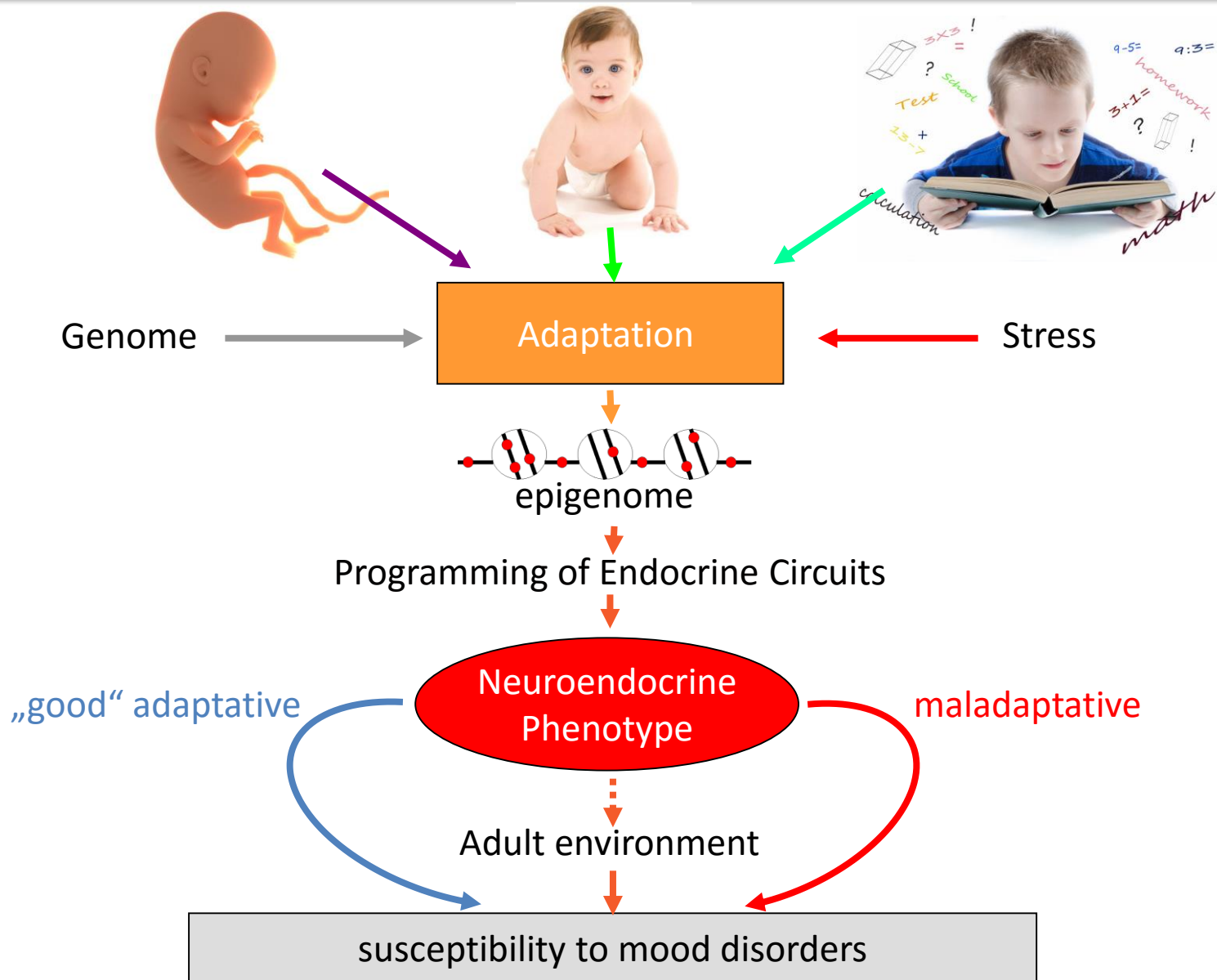
Transgenics

Selective breeding

Childhood trauma leads to increased risk of depression in adulthood



Adaption

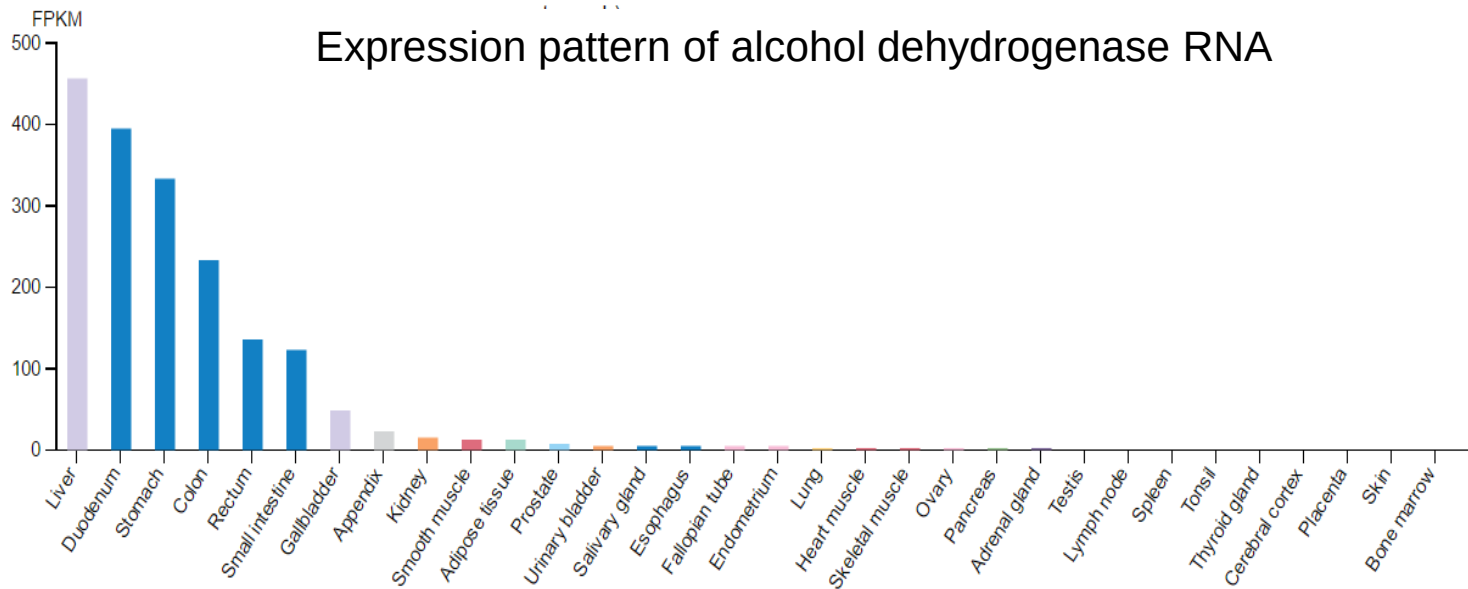
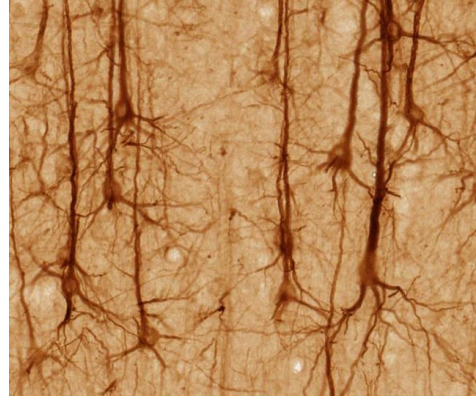


Epigenetic programming of early-life stress

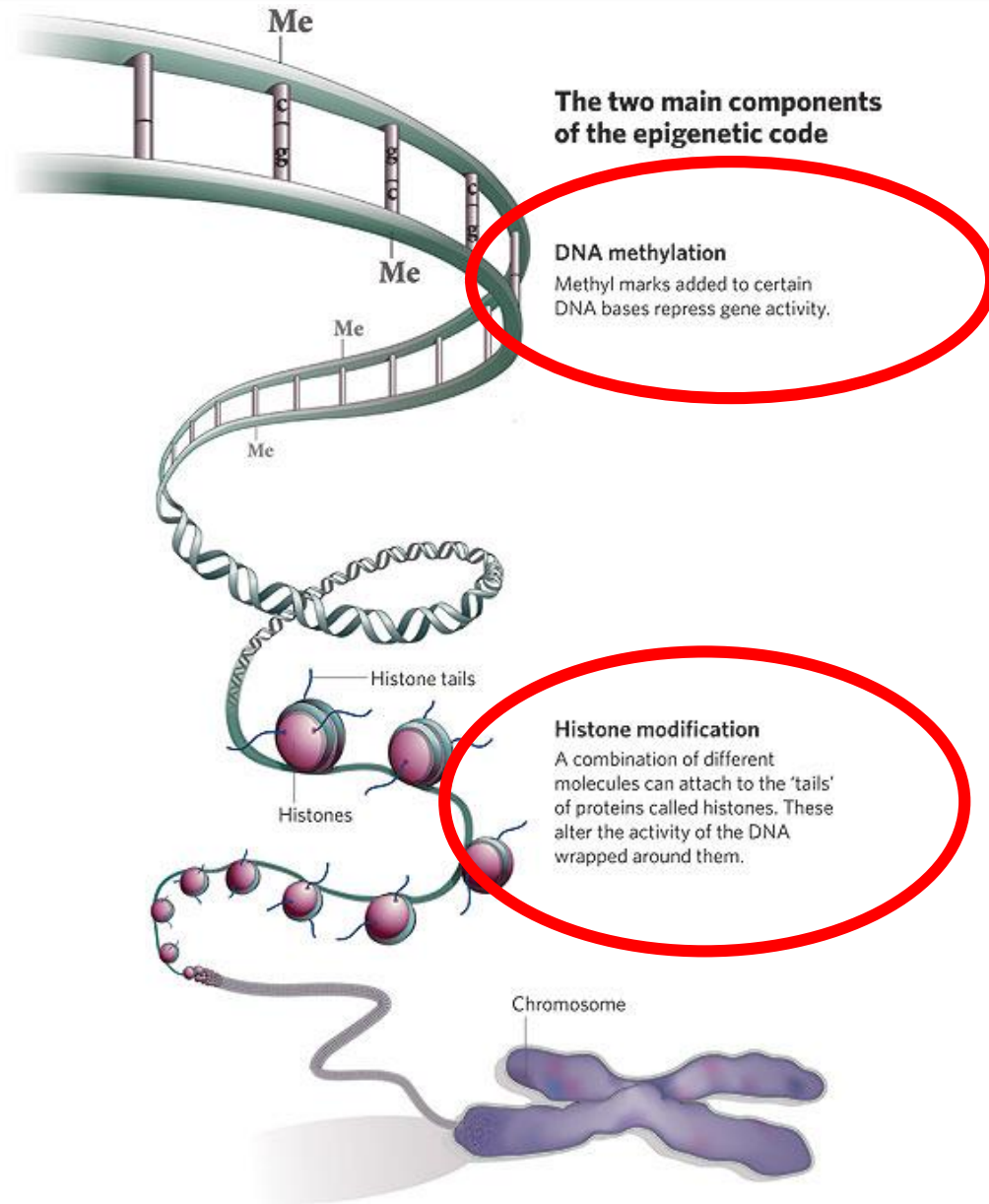
Early-life stress epigenetically programs the expression of genes regulating stress

epi = above/upon, genetics

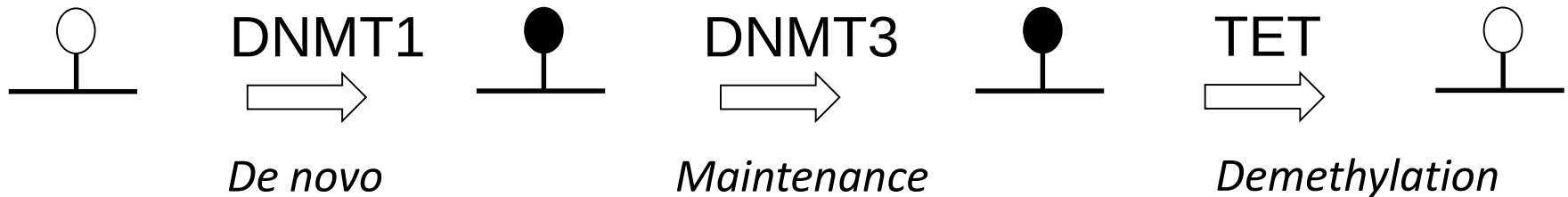
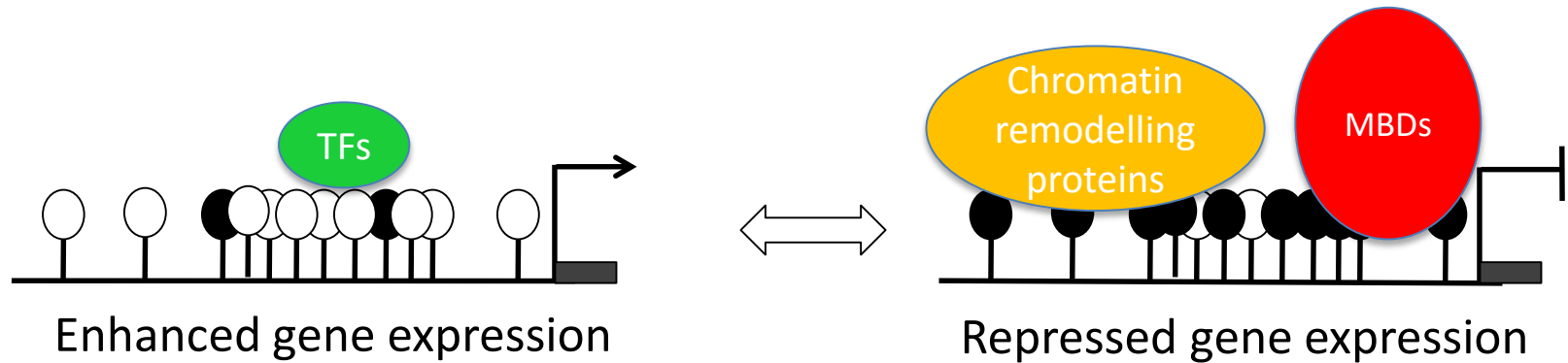
Only 10-20% of genes are active in any cell.



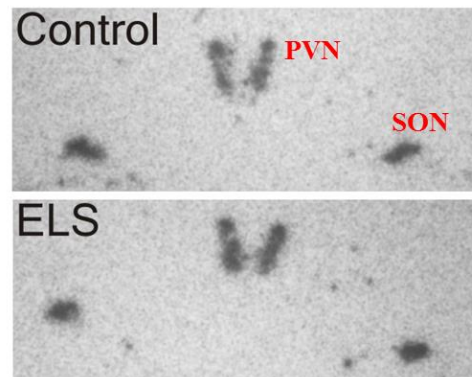
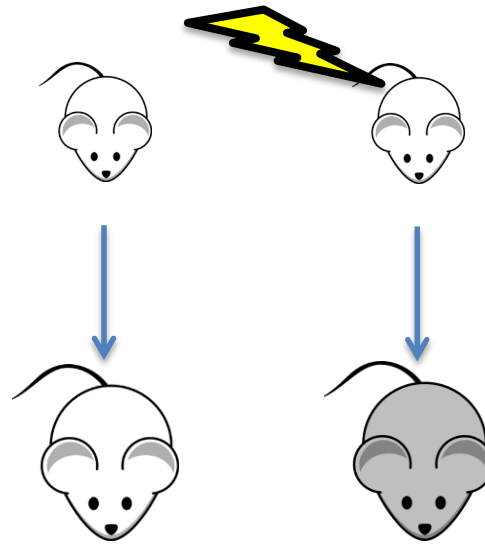
Gene silencing that occurs without changes in the DNA sequence



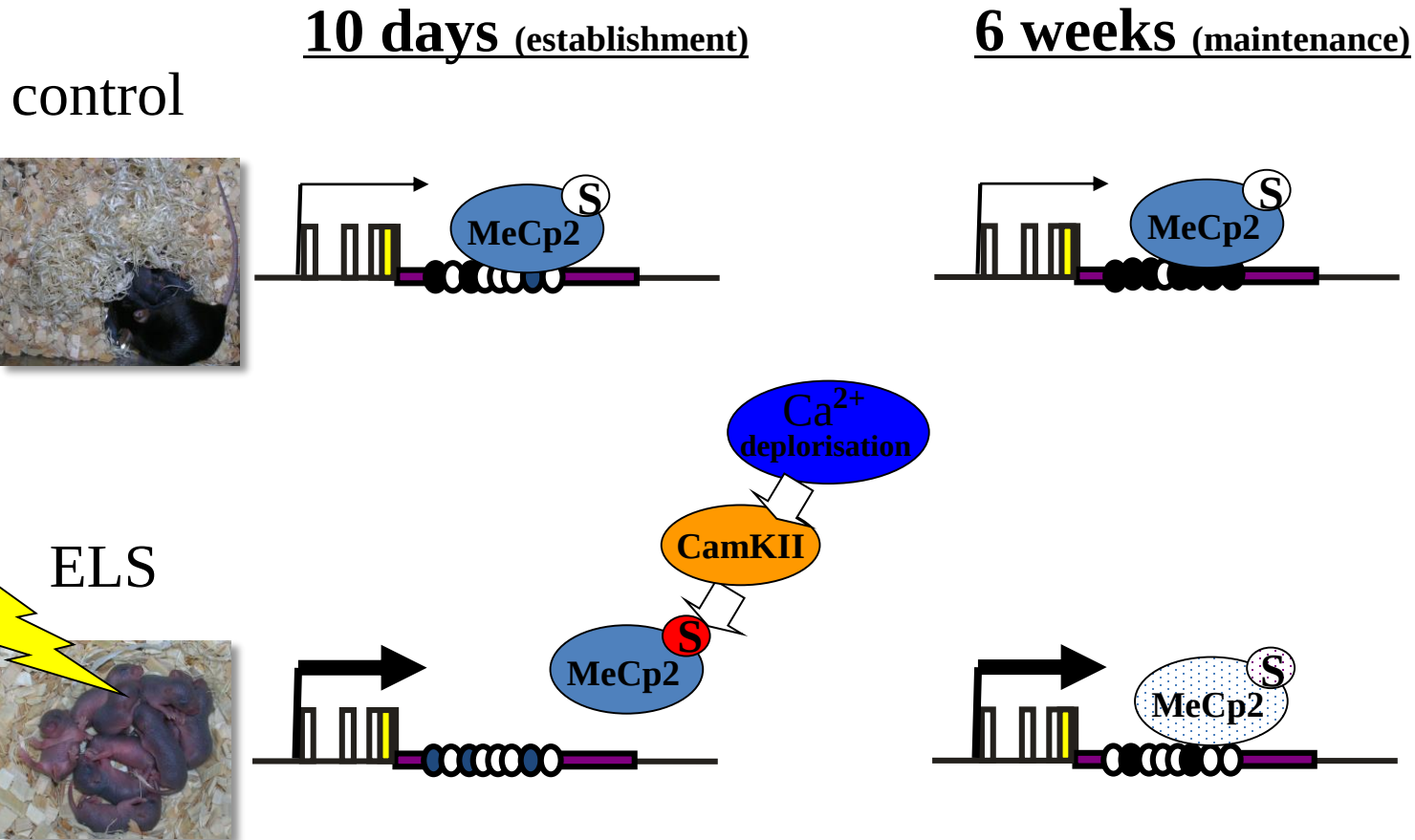
DNA methylation at gene regulatory regions can inhibit transcription



Early-life programming of anxiety & depression

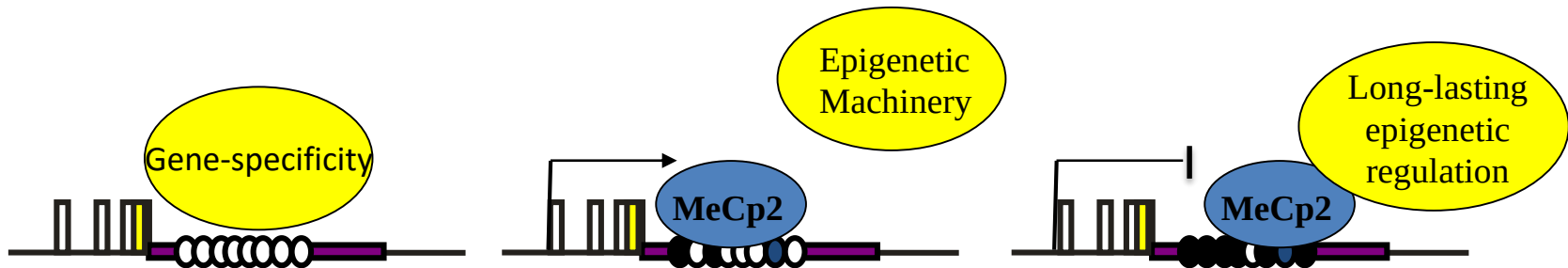


Molecular model for early-life stress programming

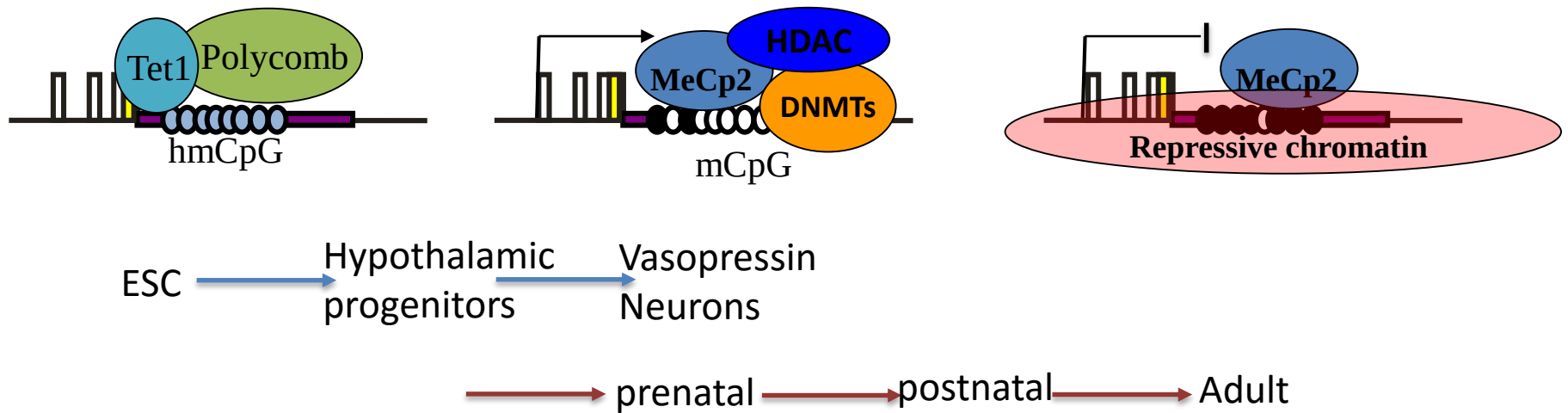


What are the mechanisms for the gene-specific marking

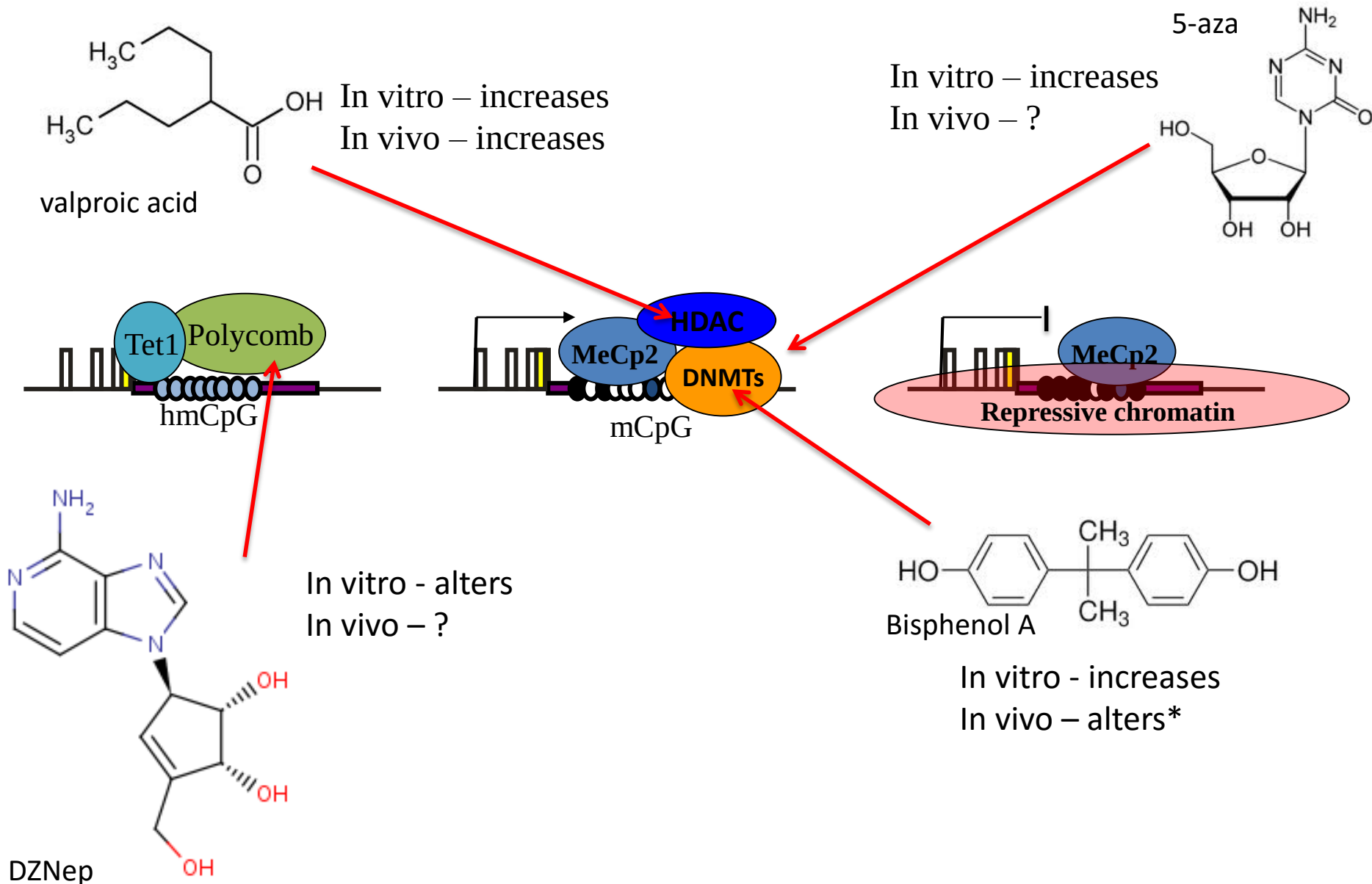
“Factor(s)”?



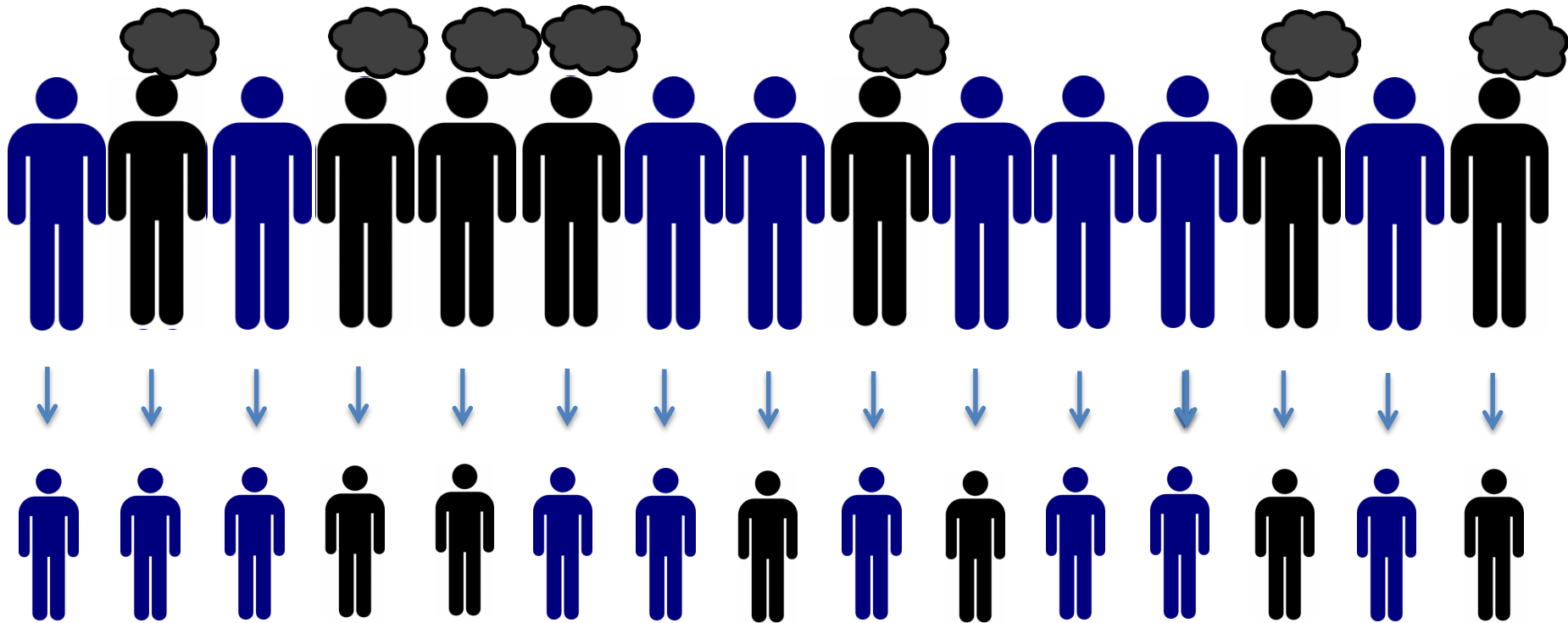
„some“ of the factors...



Pharmacologically targeting epigenetic regulation of AVP



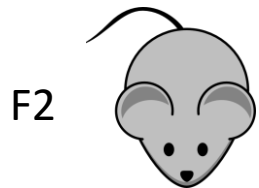
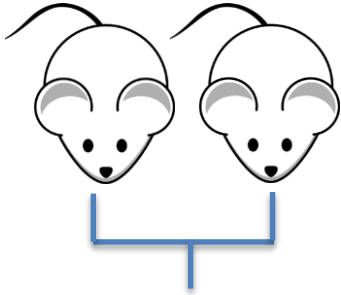
Children of depressed parents are at high risk for depression themselves



Inheritance of acquired stress



Inheritance of acquired stress



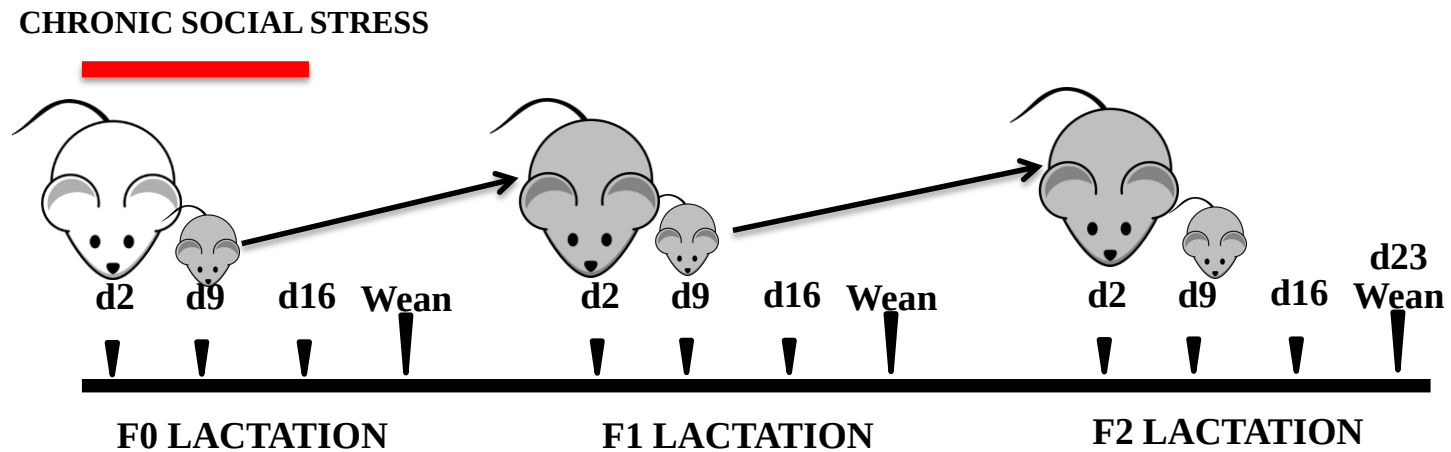
Epigenetic

Behaviour

Endocrine & Immune

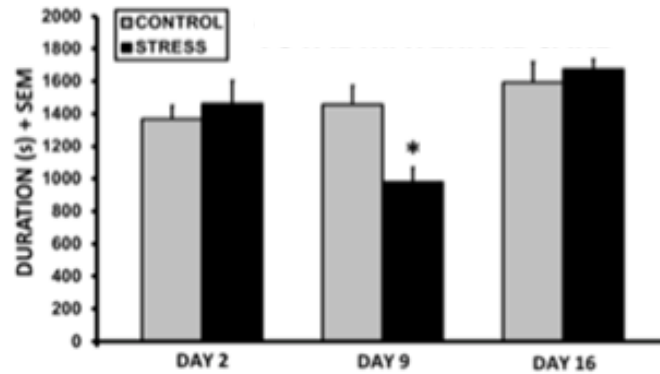
Milk (lactocrine programming)

Generational chronic social stress model for maternal behaviour

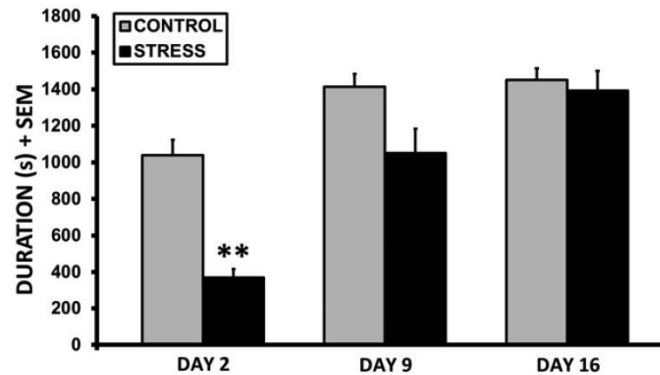


Reduced maternal care (including pup grooming)

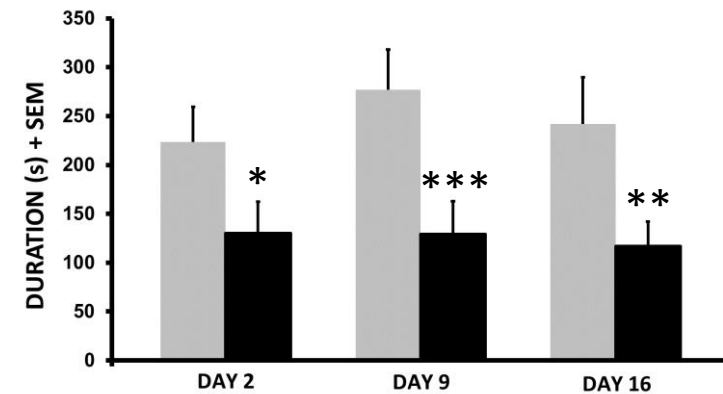
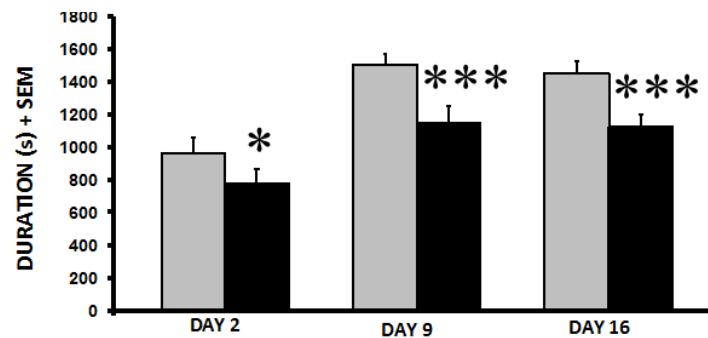
F0



F1



F2



Intergenerational transmission of maternal measures

MATERNAL FACTORS	F0			F1			F2		
	Day 2	Day 9	Day 16	Day 2	Day 9	Day 16	Day 2	Day 9	Day 16
MATERNAL CARE	-----	↓	-----	↓	-----	-----	↓	↓	↓
MATERNAL RESTLESSNESS	-----	↑	-----	↑	-----	-----	↑	-----	↑
MATERNAL AGGRESSION	-----	↑	↑	↑	-----	-----	-----	-----	-----
MILK INTAKE	-----	↓	↓	↓	-----	-----	-----	-----	-----
Corticosterone (Day 23)	-----			↑			↓		
Prolactin (Day 23)	-----			↓			-----		
Gene Regulation	PVN: ↑ GR MeA: ↓ OXT			PVN, SON, CeA, MeA: ↓ OXT PVN: ↓ OXTR, PRLR, AVP			-----		
Nr3c1 exon 1.7 Methylation	PVN: ↑ CpG2			PVN: ↑ CpG3 Hipp: ↓ CpG1-6			-----		
ICAM-1 (Day 23)	-----			-----			↓		

Milk production (intake), Milk Corticosterone

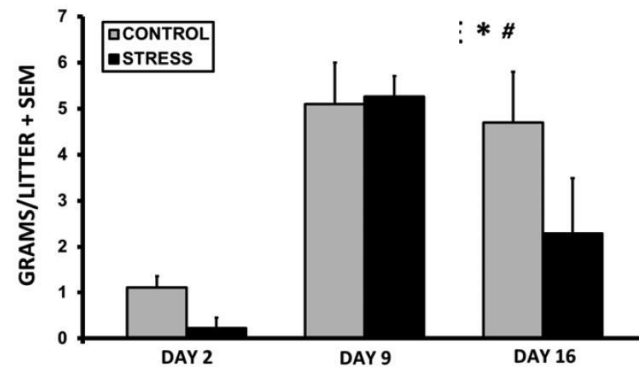
Milk production

Milk corticosterone

F0

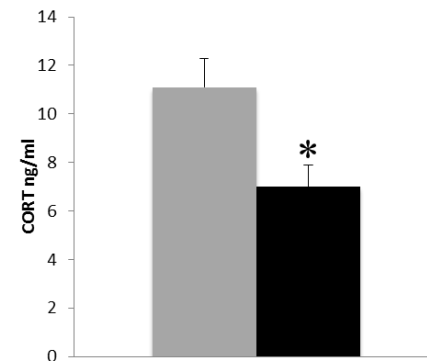
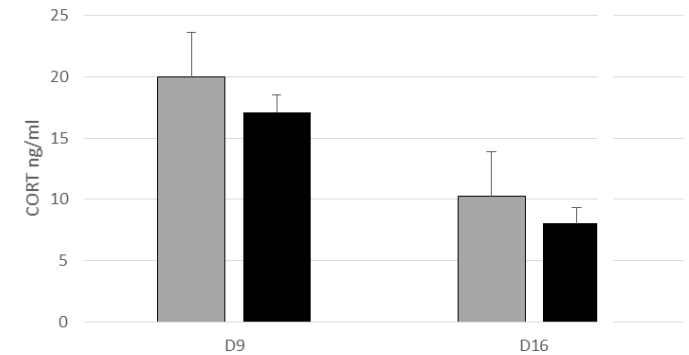


F1



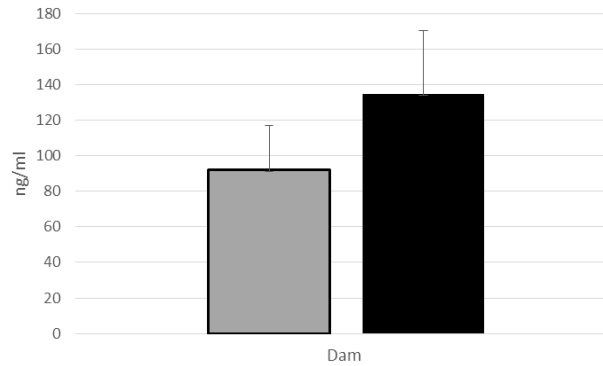
F2

No differences

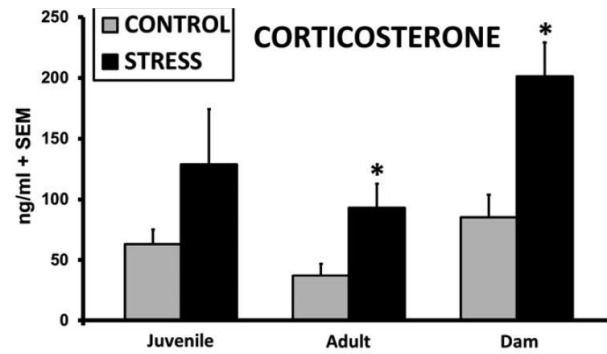


Serum Corticosterone

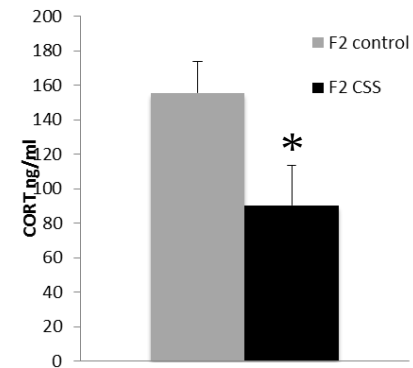
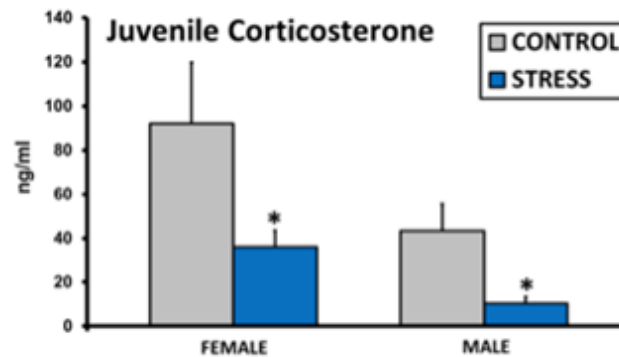
F0



F1



F2



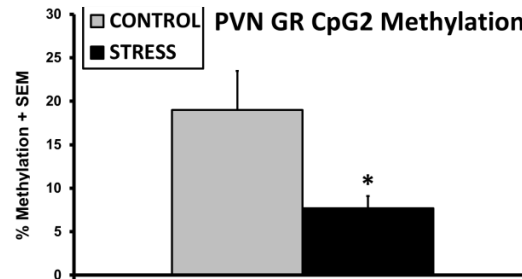
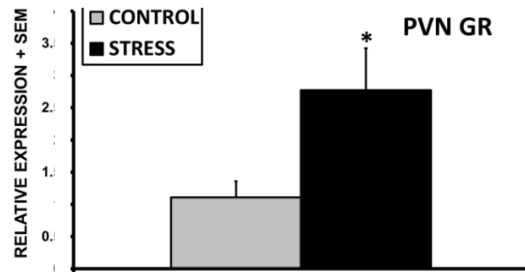
GR regulation

PVN expression

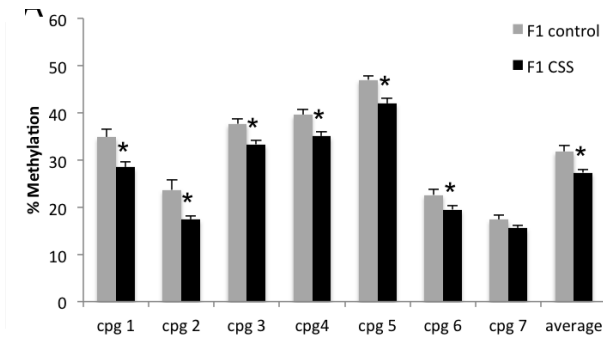
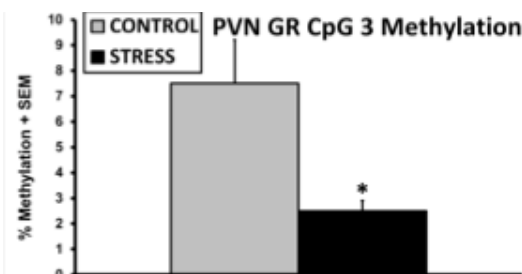
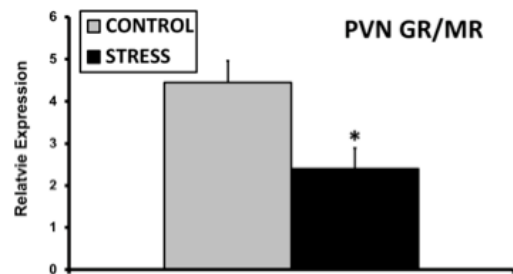
PVN Methylation

Hippocampus Methylation

F0

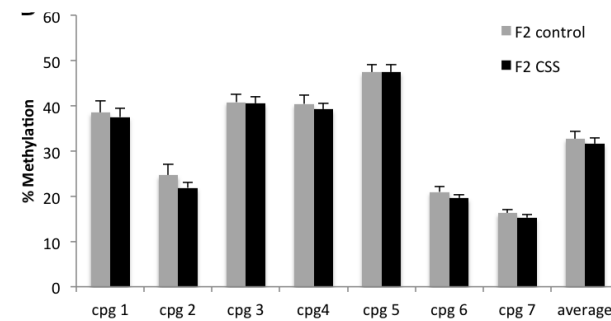


F1



F2

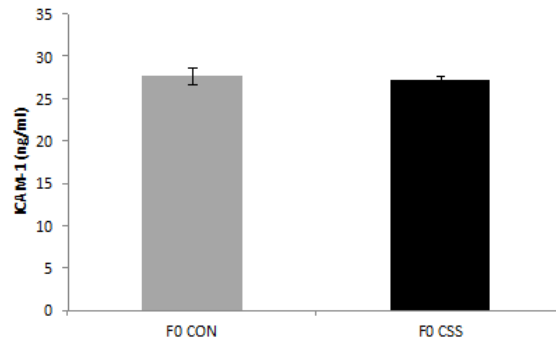
No differences



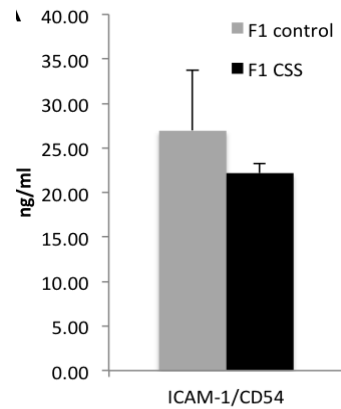
*Intergenerational genomewide analysis ongoing

Immune markers (ICAM-1)

F0

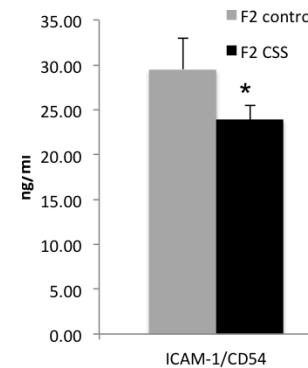
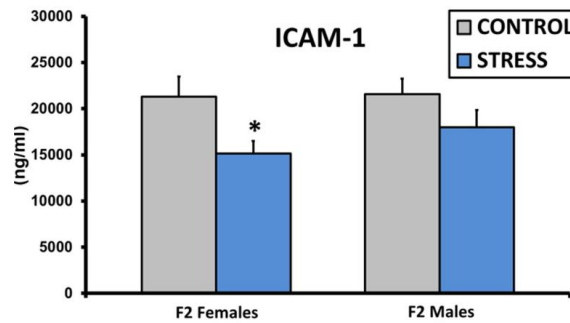


F1

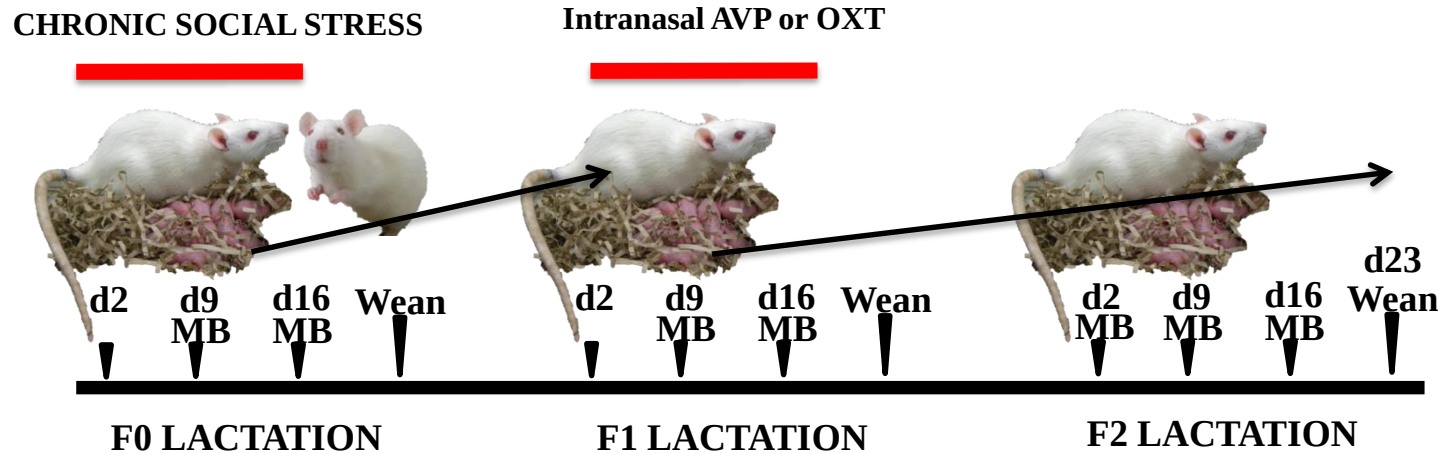


F2

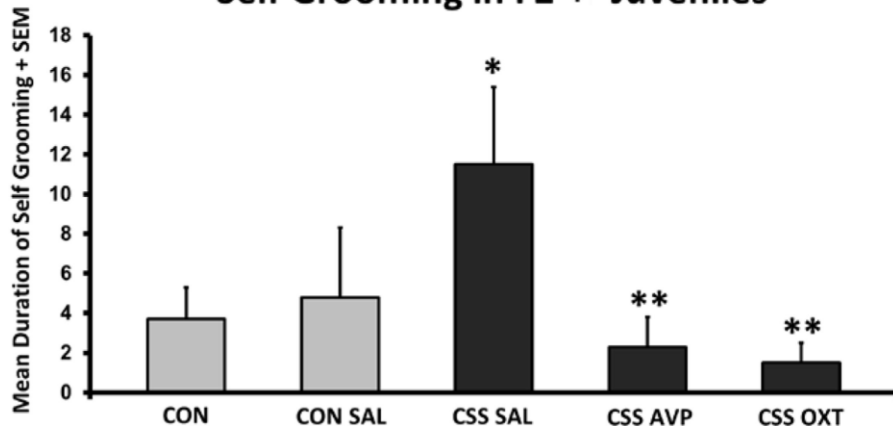
IFN γ
IL18,
a1AGP,
BDNF



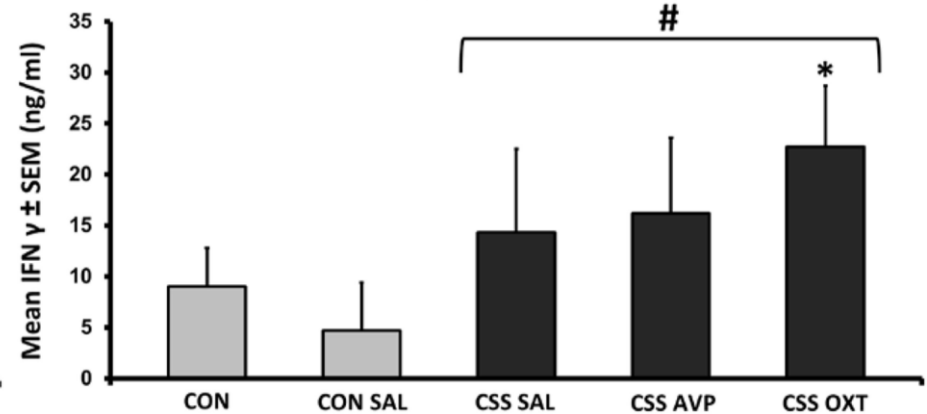
Effects of AVP and OXT treatment of CSS dams on female offspring



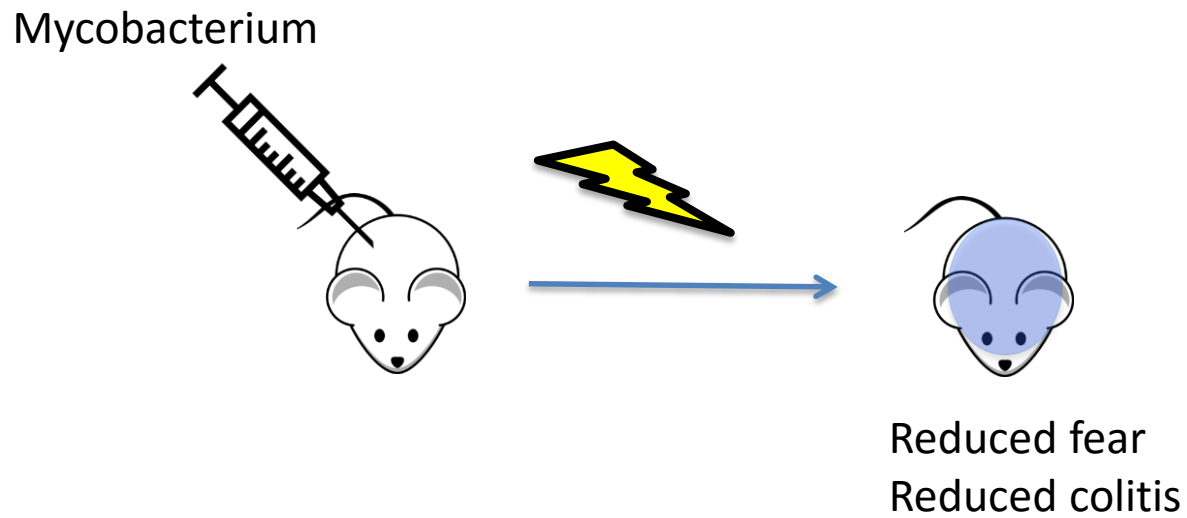
Self Grooming in F2 ♀ Juveniles



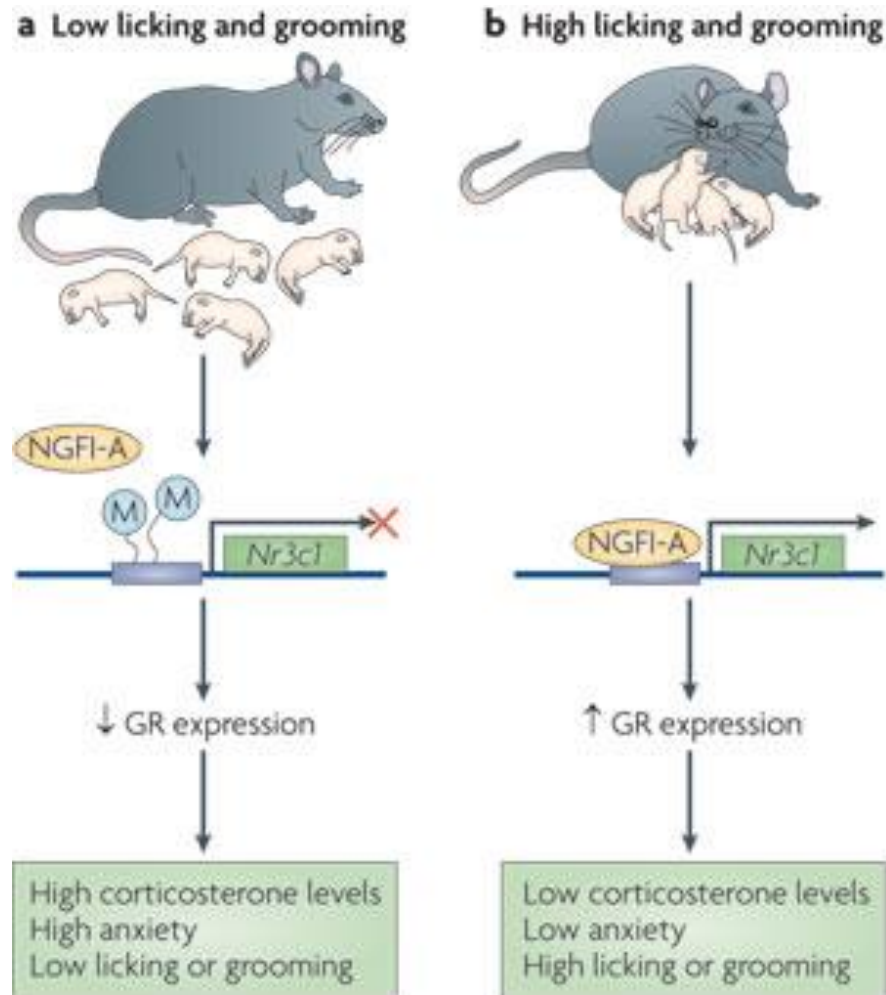
Basal IFN γ in ♀ F2 juveniles



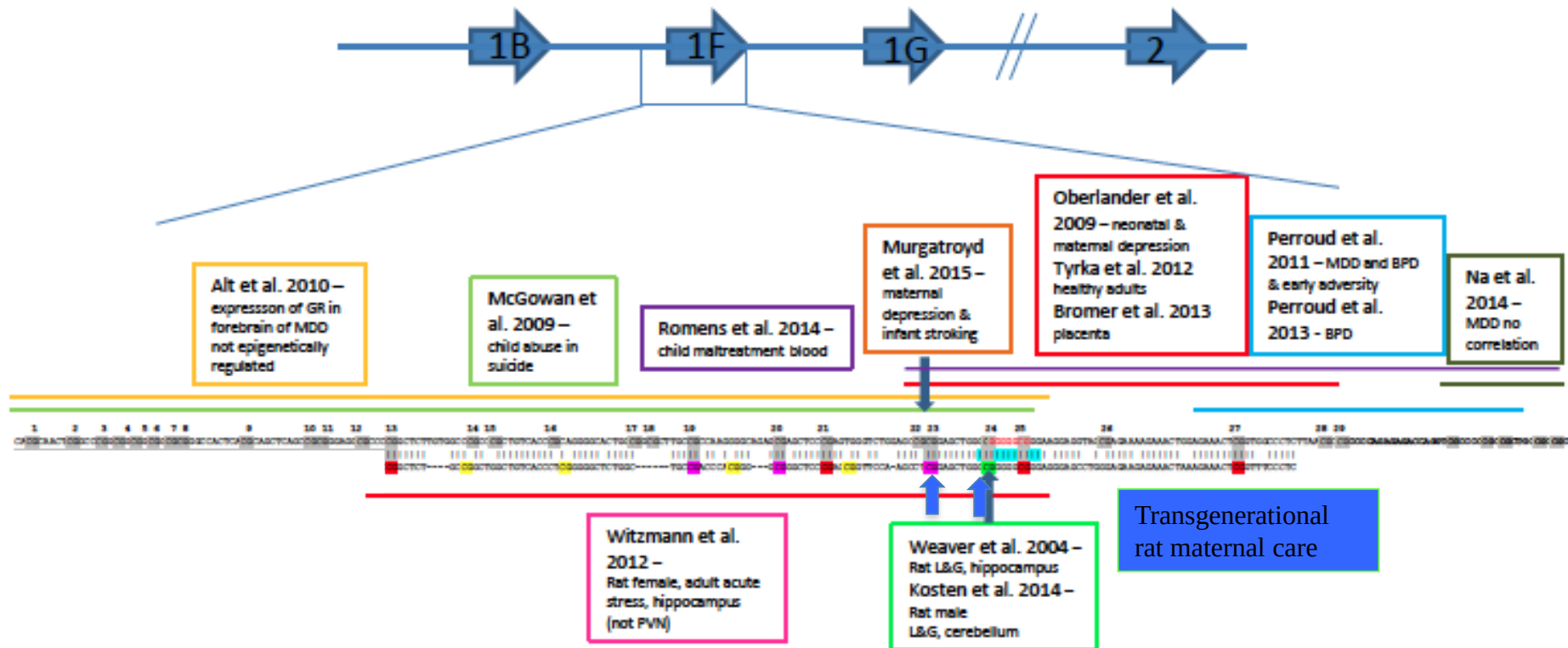
Immunisation promotes stress resilience



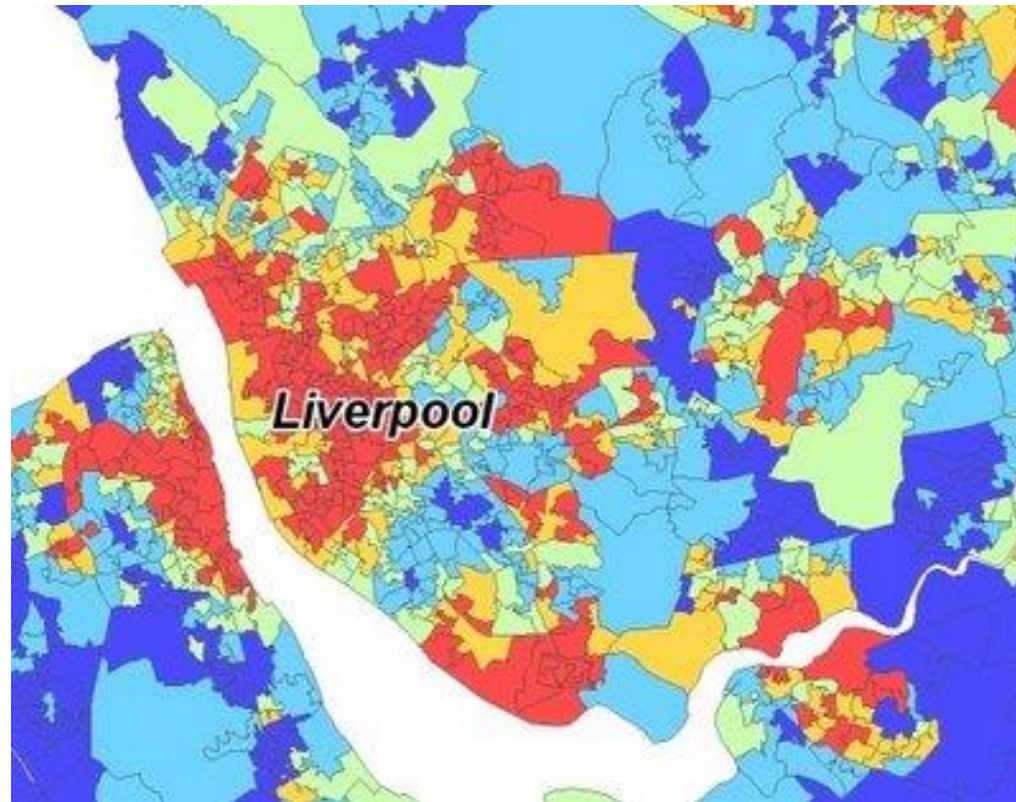
Nr3c1 methylation and early life licking/grooming



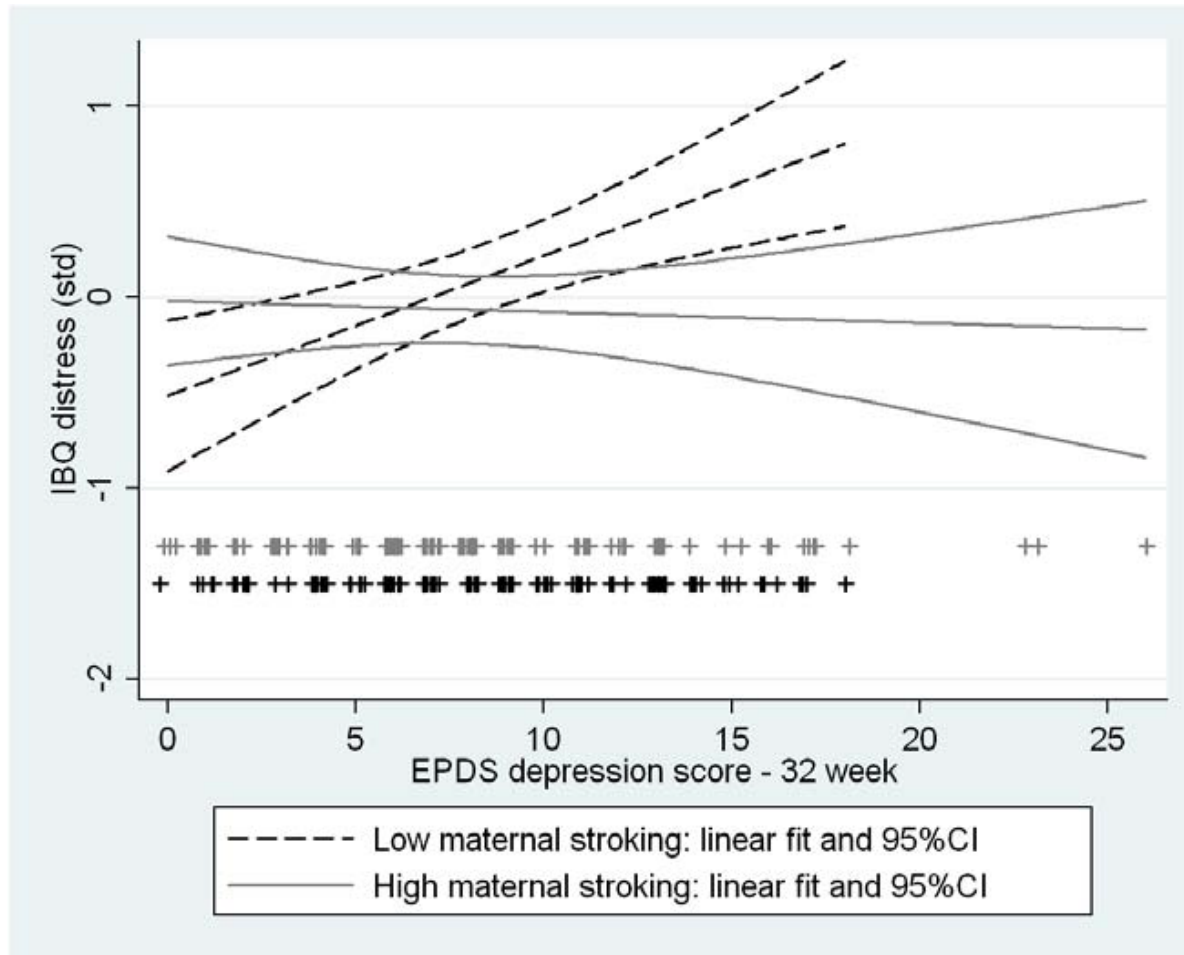
Methylation of human & rat GR exon 1F/7 in the literature



WCHADS - Wirral Child health and Development Study



Maternal stroking moderates effect of prenatal depression on infant behaviour





IASAT

International Association for
the Study of Affective Touch



IASAT2017

2nd Congress Programme

Friday 1st September - Sunday 3rd September

in Liverpool, UK at Liverpool John Moores University
Derwent Parkside Building, Byrom Street, Liverpool, L3 3NP, UK



Håkan Olausson · Johan Wessberg
India Morrison · Francis McGlone
Editors

Affective Touch and the Neurophysiology of CT Afferents

 Springer

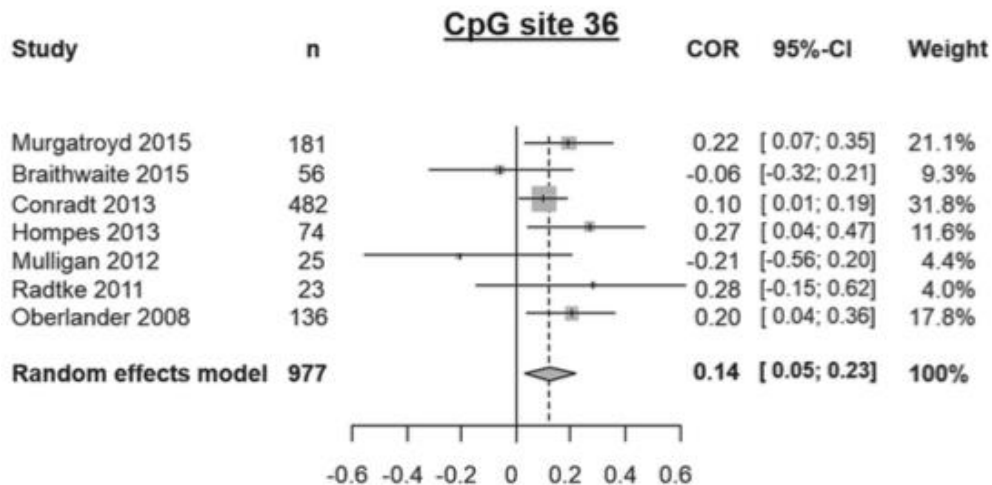
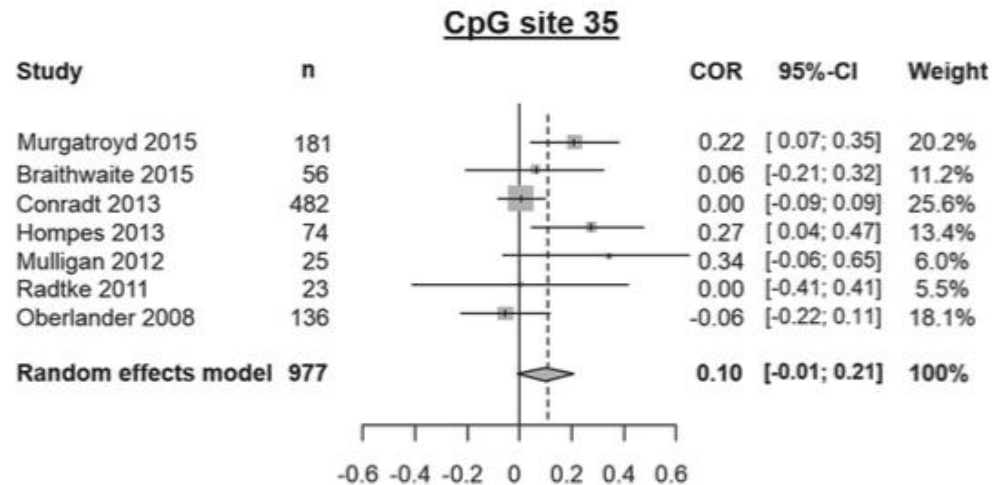
C-tactile afferents: Cutaneous mediators of oxytocin release during affiliative tactile interactions?



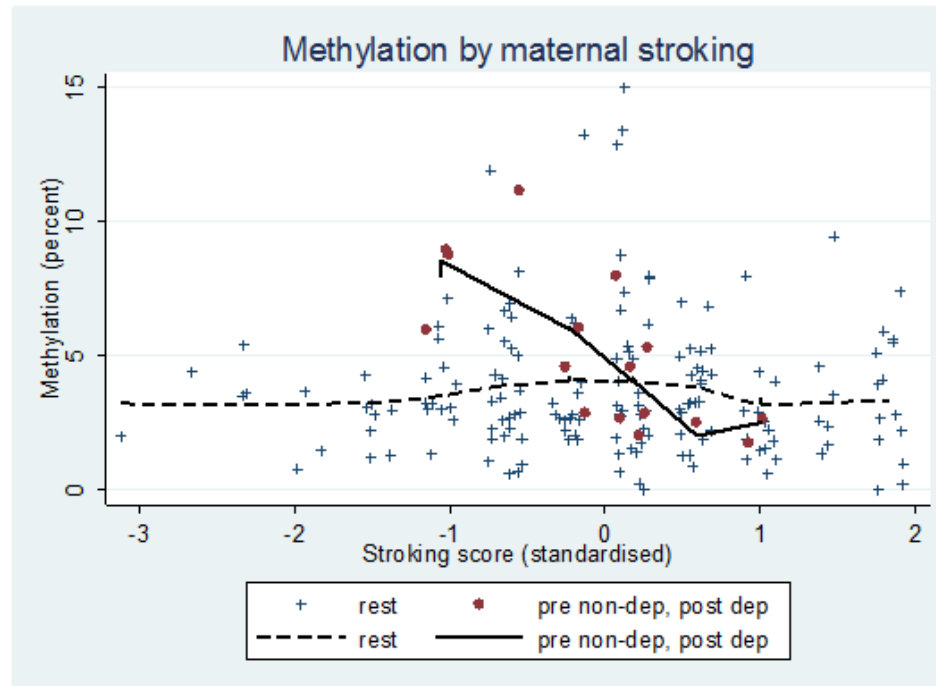
Stroking and DNA methylation in human

*Early life stroking reduces GR methylation
and stress behaviour in infants*

Specific CpGs associated with maternal depression and childhood stress



Maternal stroking (5 weeks age) reduces GR methylation



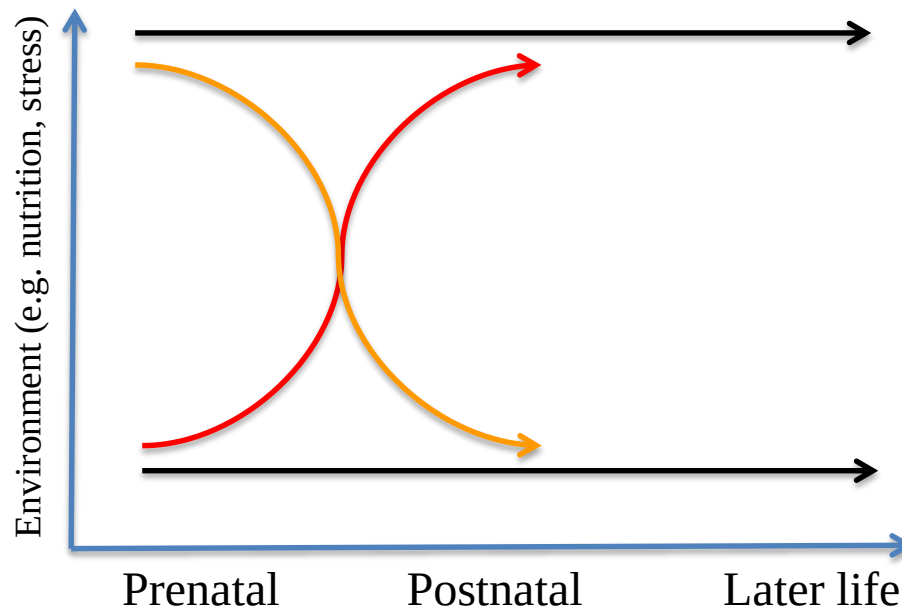
no effect of maternal stroking at 9 weeks of age

Predictive Adaptive Response & Mismatch

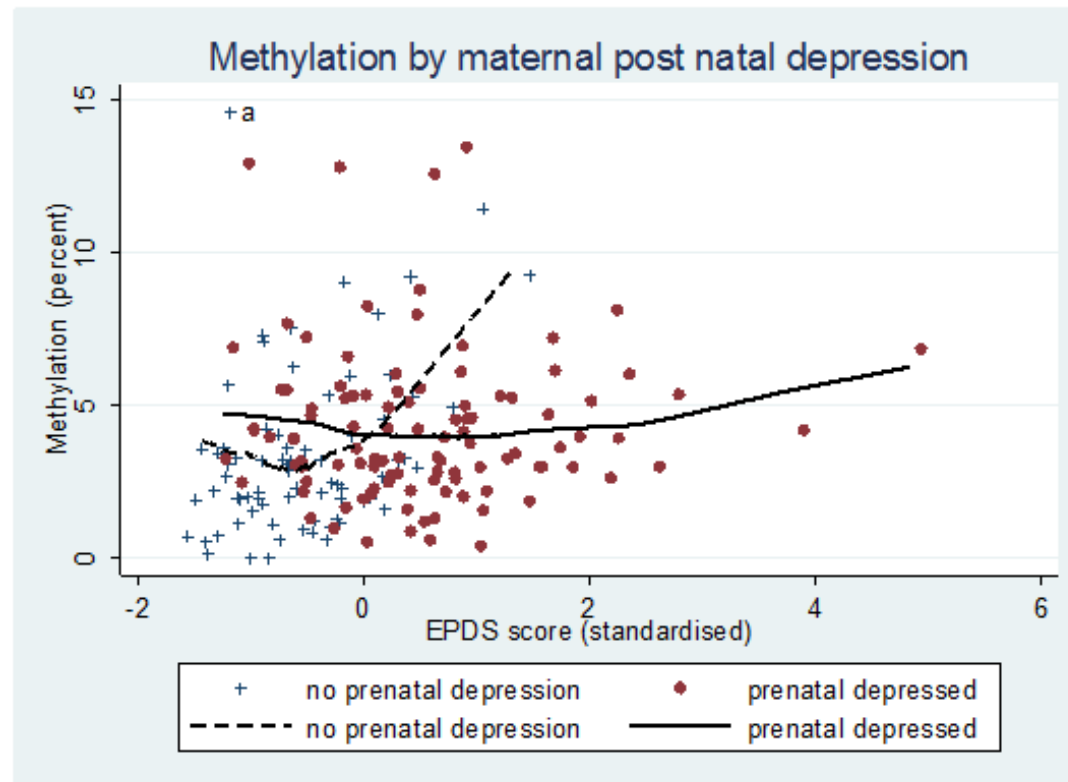
PAR hypothesis

Barker, Br Med Bull. 1997 53:96-108

An adaptive response to environmental cue acting early in life, where the advantage of the induced phenotype is primarily manifest in later life.



Infants of mothers with low prenatal depression were more vulnerable to the effects of postnatal depression

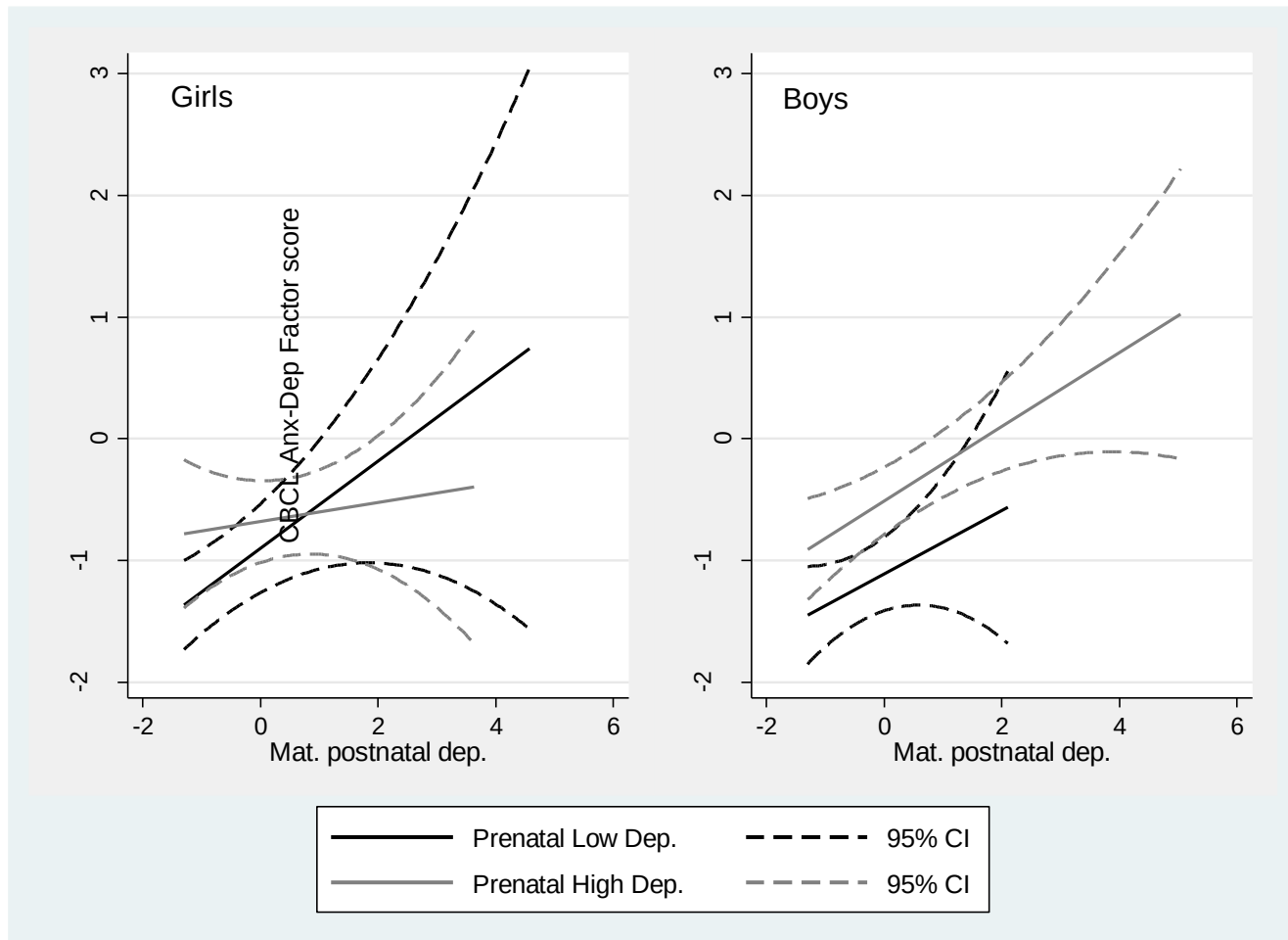


Sex differences in interactions of pre- and postnatal depression

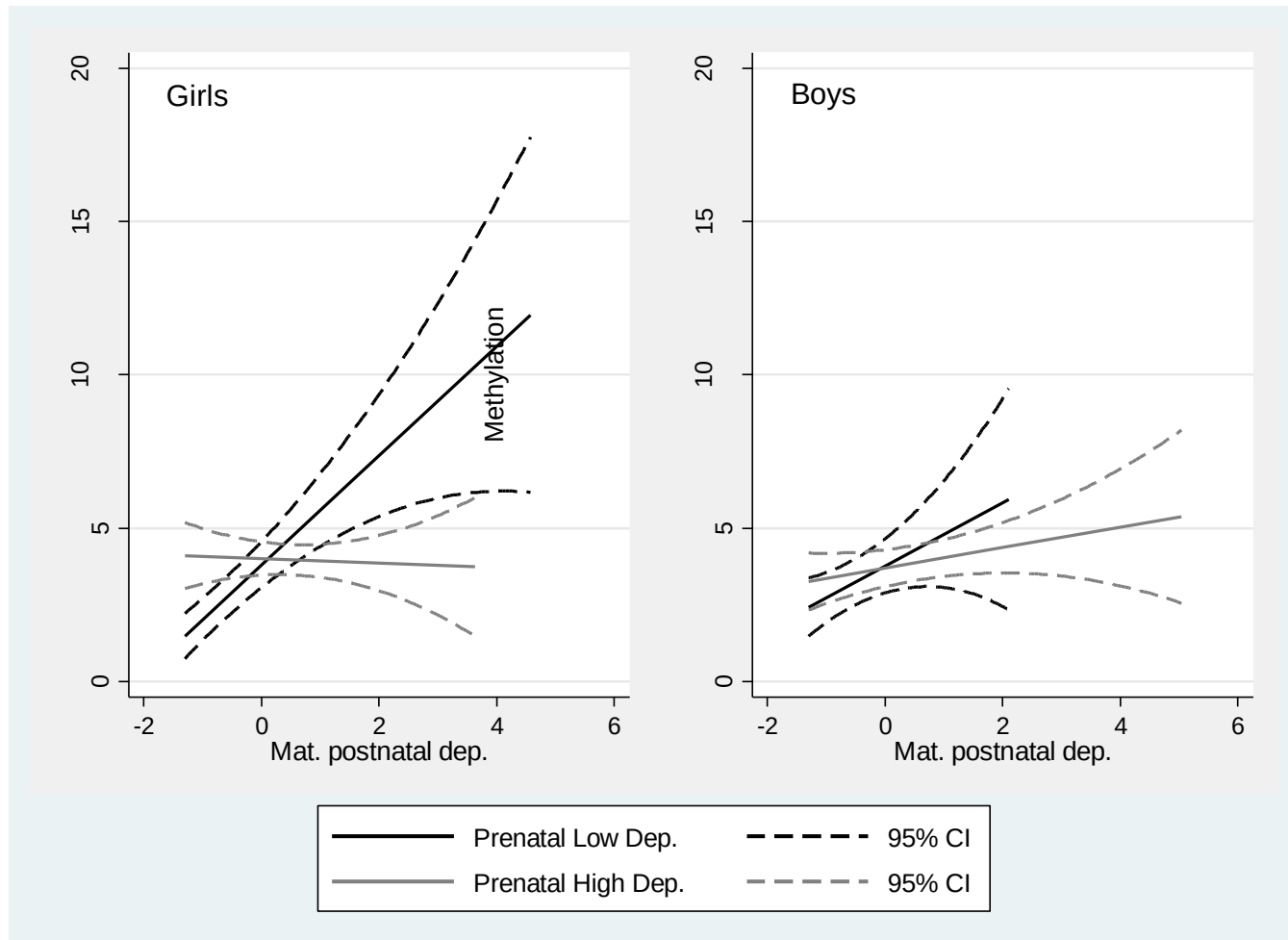
Trivers–Willard hypothesis
1973, Science 179 (4068): 90–9

A good conditioned mother transfers competitive ability to male offspring for greater reproductive payoff

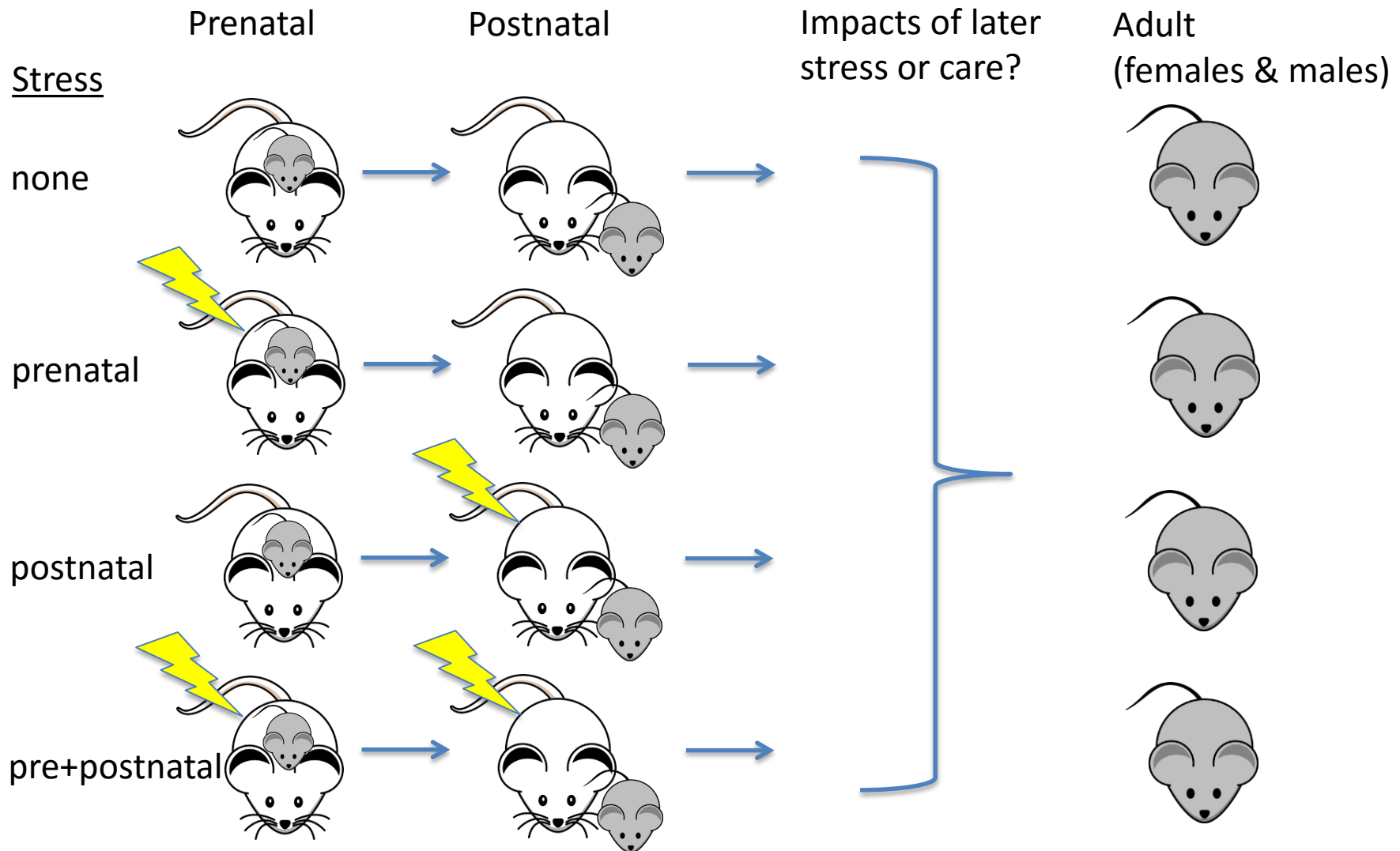
Prenatal-postnatal mismatch conditions associate with child symptoms in girls only



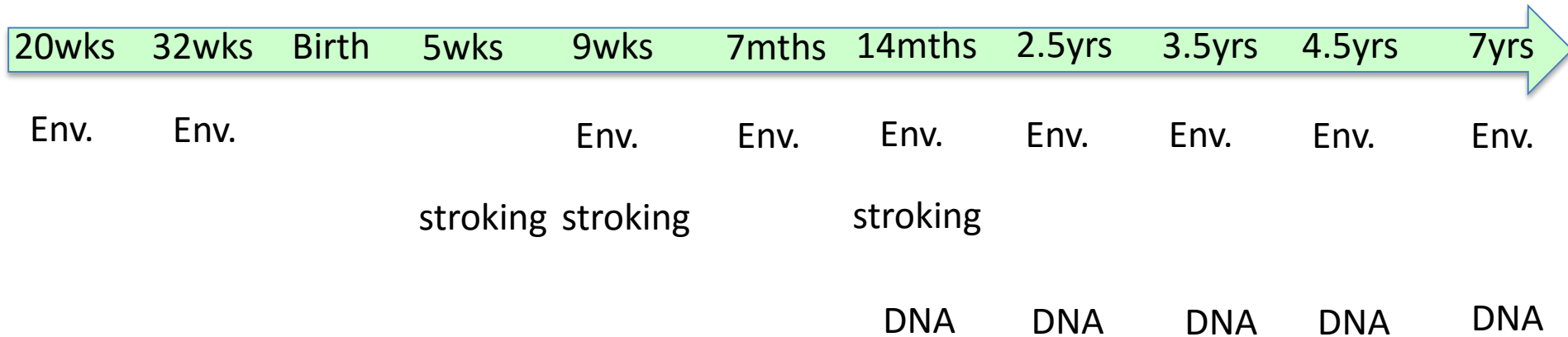
Prenatal-postnatal mismatch conditions significantly associate with methylation in girls only



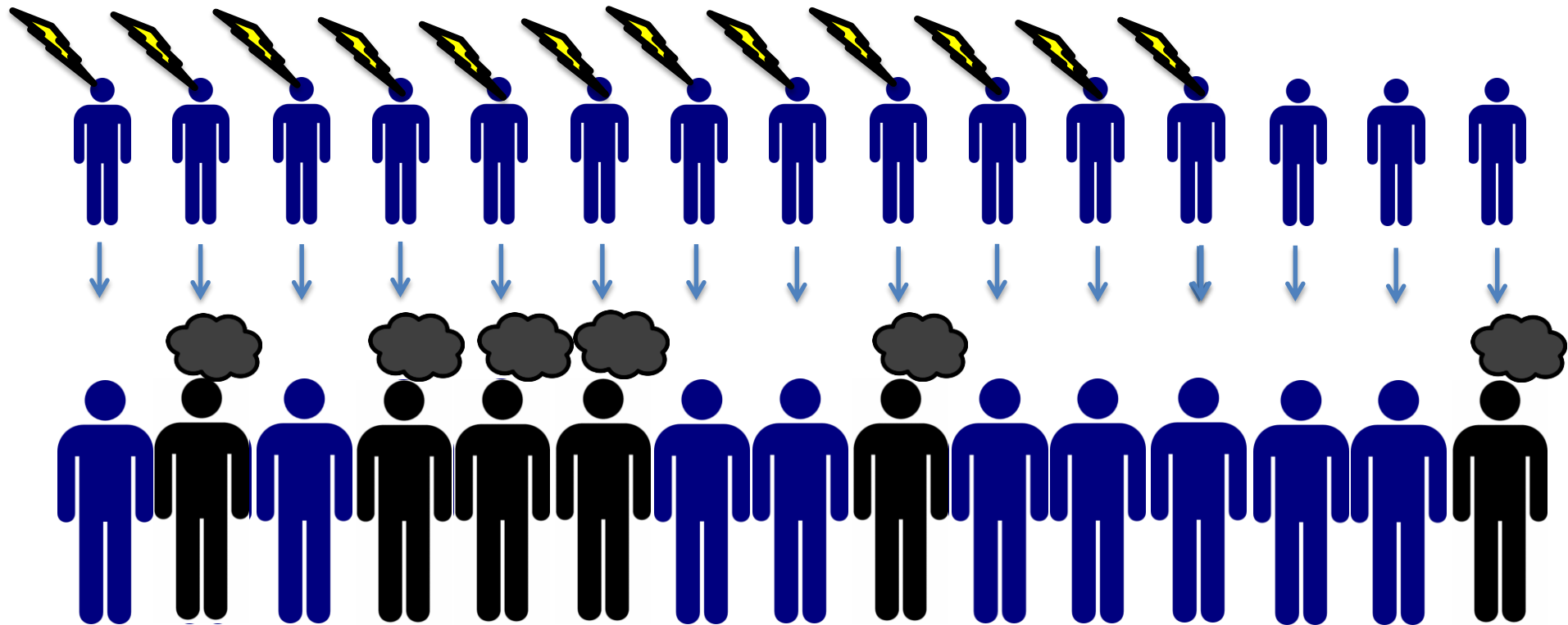
Animal model for Prenatal-postnatal mismatch conditions



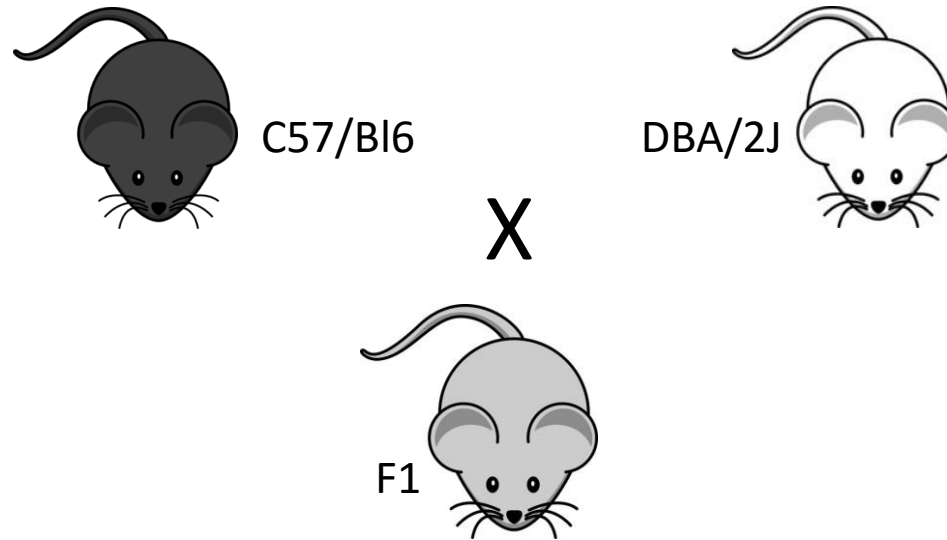
WCHADS - Longitudinal changes in DNA methylation



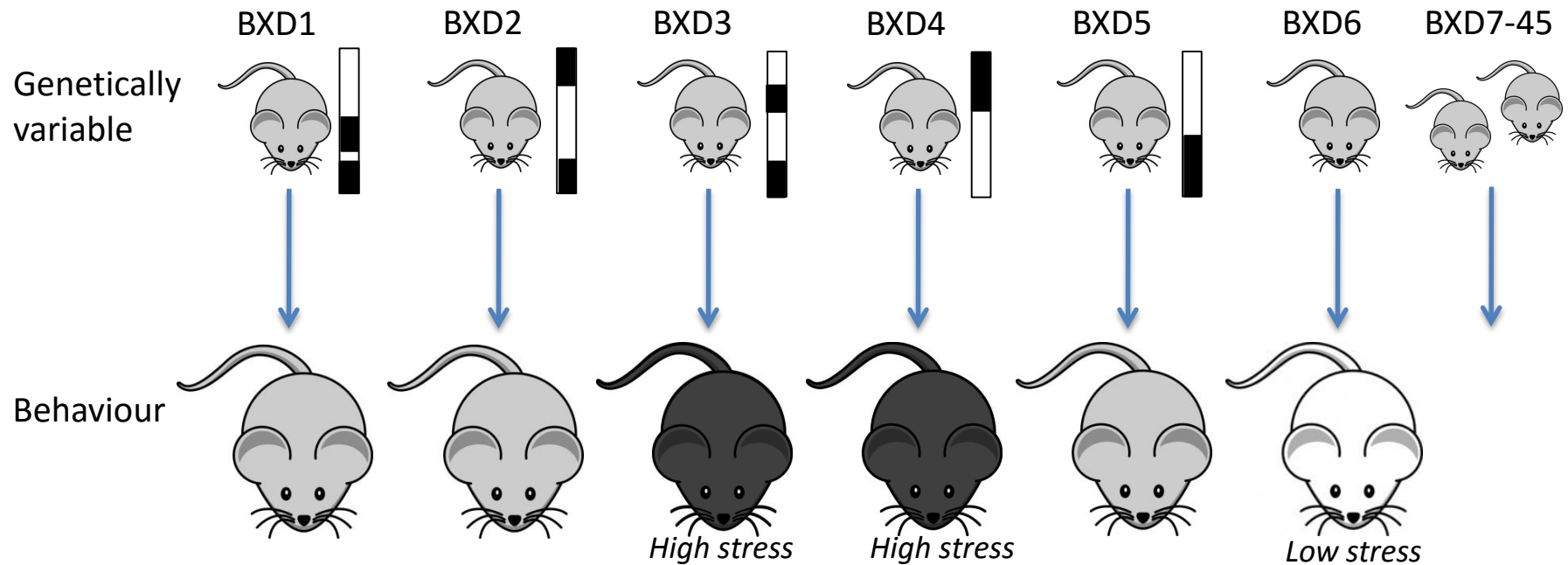
Not everyone with early-life stress develops depression
Not everyone with depression suffered early life stress



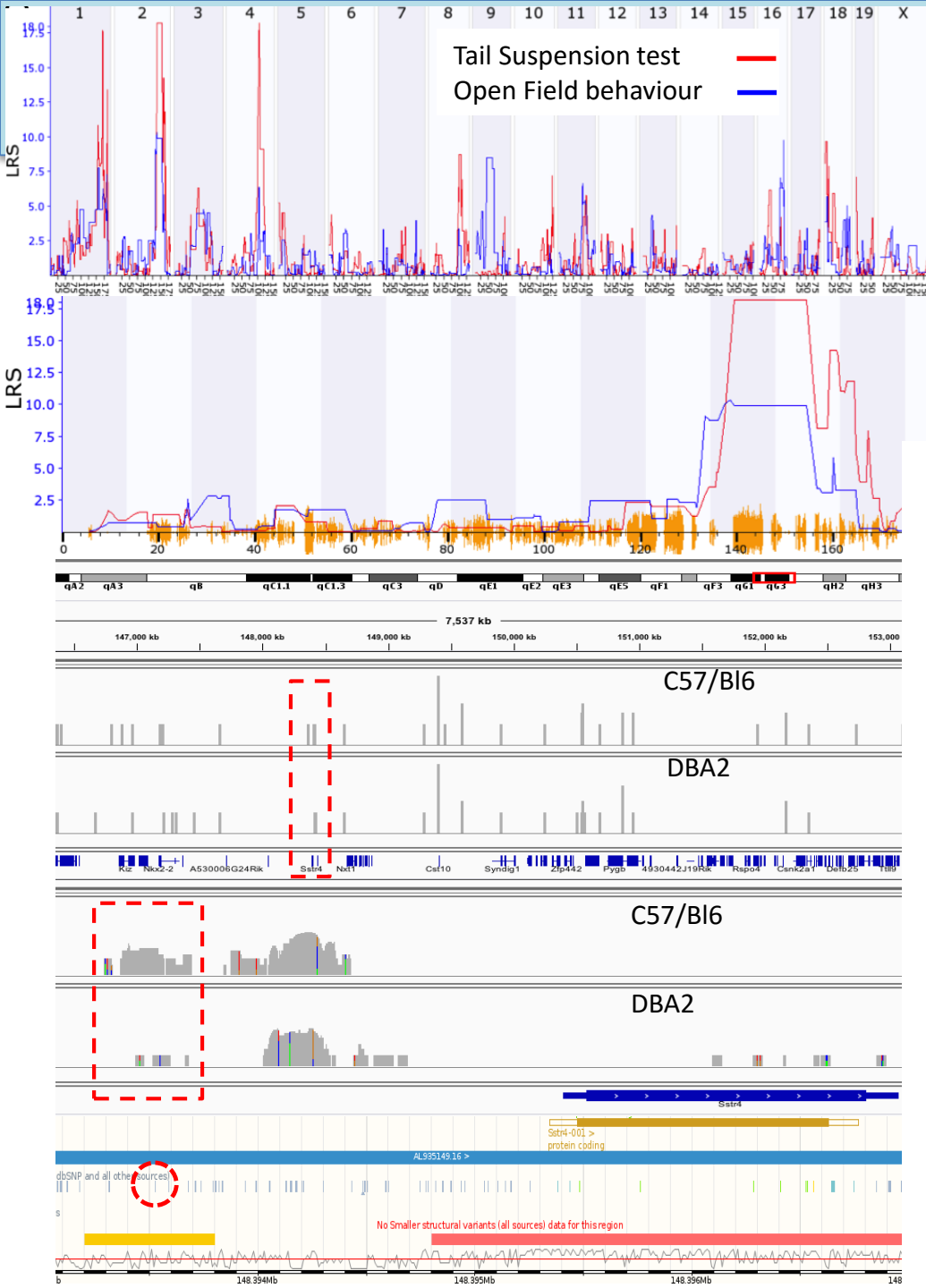
Recombinant inbred mouse strains



Animal model for genomewide analyses

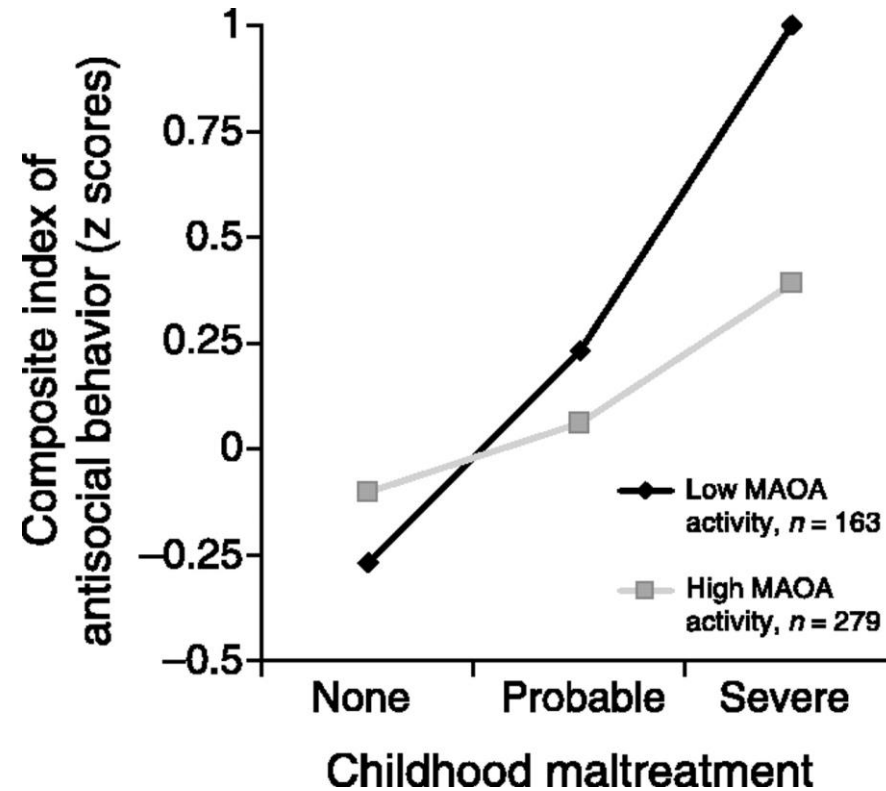


epiQTL for Stress-related behaviour

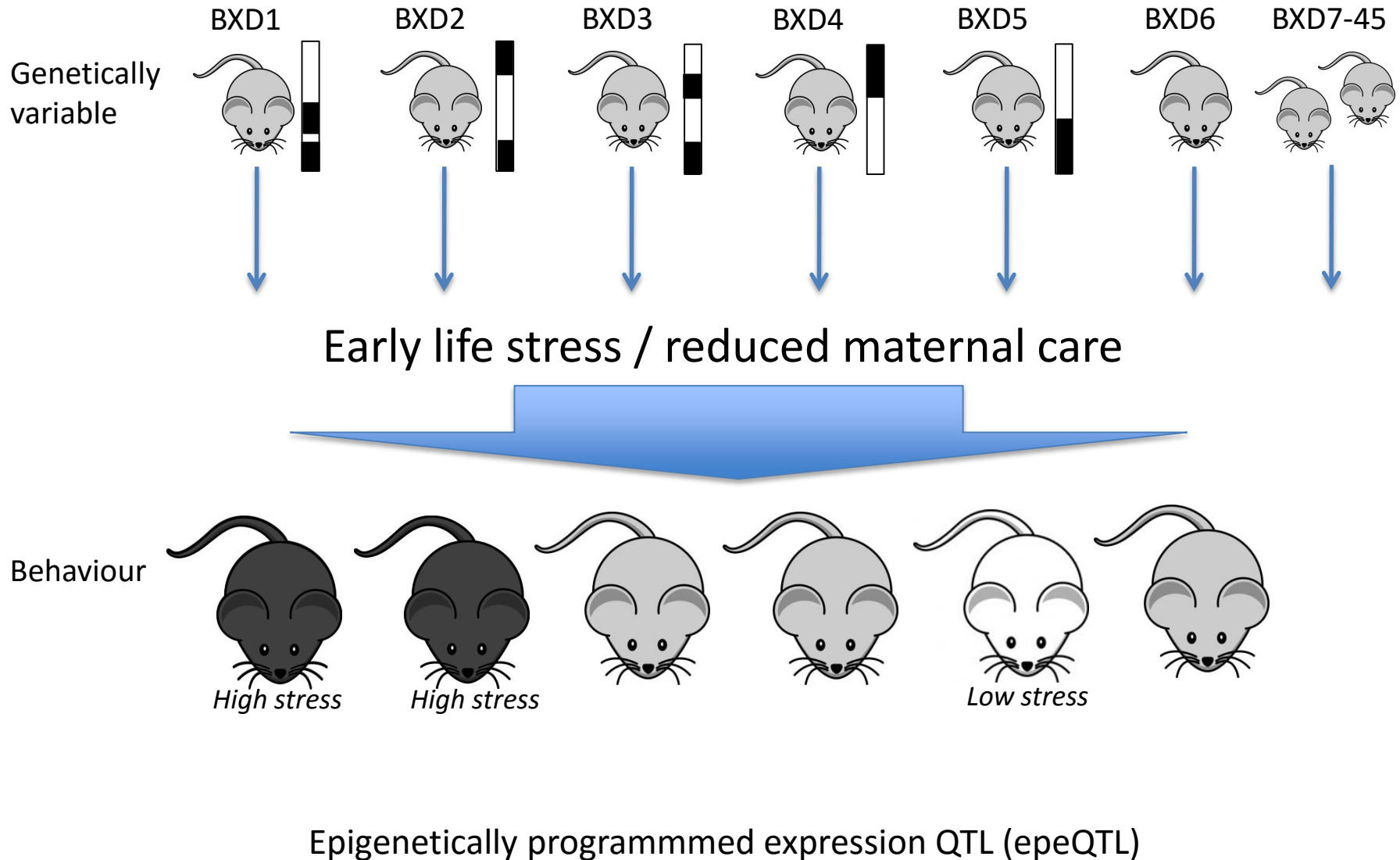


Scheich et al. 2016 Neuropharmacology. 101:204-15. Somatostatin receptor subtype 4 activation is involved in anxiety and depression-like behavior in mouse models.

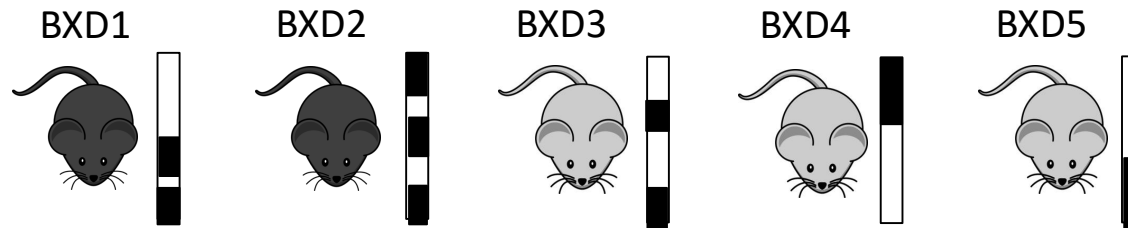
Gene x Environment (GxE)



Animal model for genomewide GxE analyses



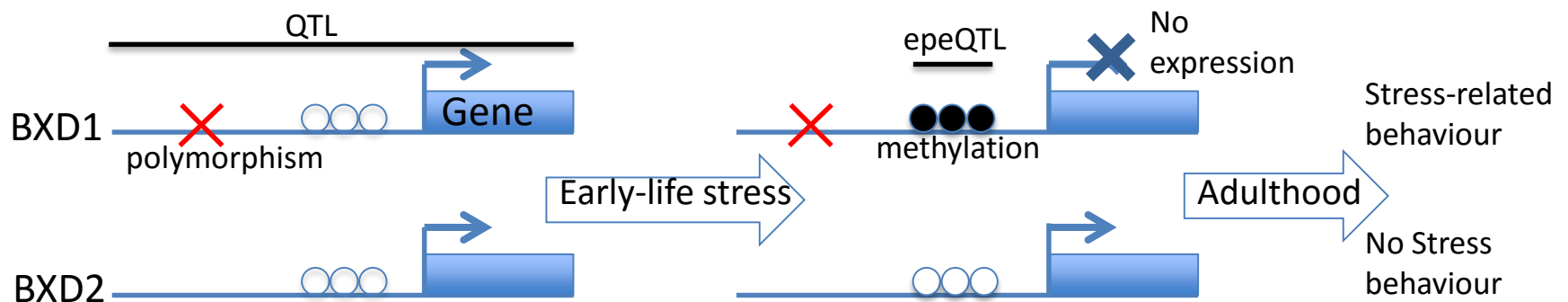
An epigenetically programmed expression QTL (epeQTL)



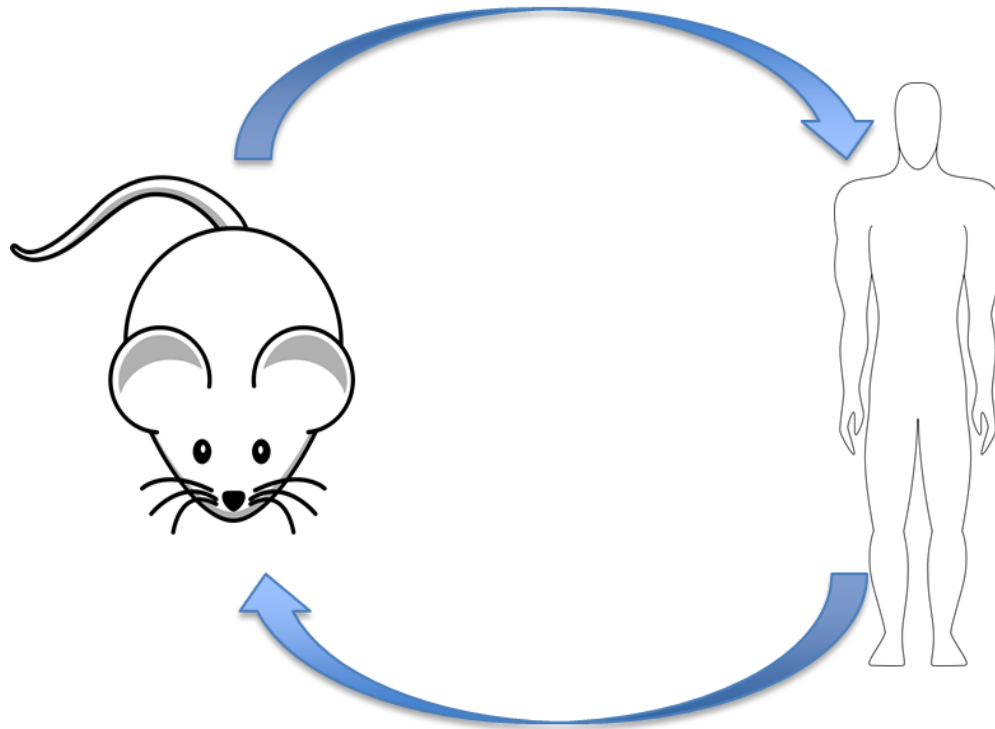
Which genetic regions are shared between sensitive (or insensitive) BXD lines (QTL)?

Which regions in QTL show differential mDNA following early-life stress (epeQTL)?

Which genes are differentially expressed in association with epeQTL



Translational animal models



Acknowledgements



WCHADS, Helen Sharp, John Quinn

**Institute of
Psychiatry**

at The Maudsley

KING'S
College
LONDON

Andrew Pickles

Jonathan Hill



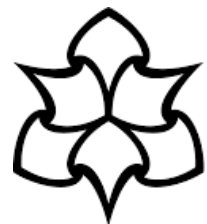
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Reading**



The University of Manchester

Jo Neill, Mike Hearte, Reinmar Hager

Ben Nephew



**Manchester
Metropolitan
University**

MRC

Medical
Research
Council



National Institutes
of Health