

Erasmus MC
University Medical Center Rotterdam



Genetics of PCOS: The insulin receptor pathway

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Mark McCarty, Steven Franks, Joop Laven

PCOS consensus meeting. Women's health aspects of PCOS. November 18th

PCOS is a complex genetic trait

Oligo- or Amenorrhea (cycle length > 35 days)
Hyperandrogenism (hirsutism and / or FAI > 4.5)
PCO (> 12 follicles, 2 – 9 mm, in one or both ovaries
and/or ovarian volume > 10 mL)

} 2 out of 3: **PCOS**

Rule out:

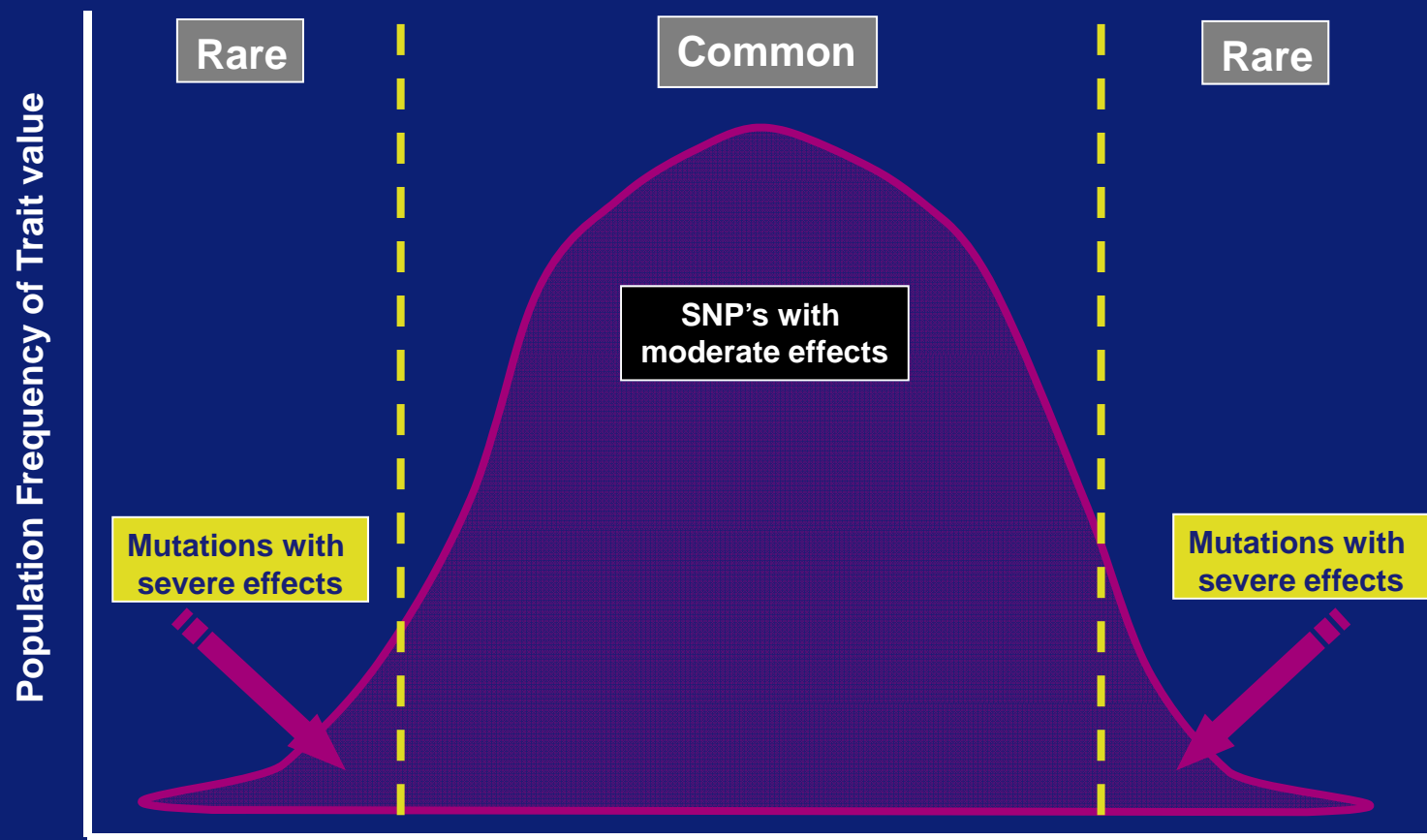
Hyperprolactinemia, NCAH, Cushing's syndrome,
Androgen Secreting Neoplasm, Acromegaly



ESHRE / ASRM sponsored PCOS consensus meeting 2003, Rotterdam, The Netherlands

PCOS is a heterogeneous phenotype indicating a complex genetic background,
which might be altered by the environment

Genetics of population variation of traits



PCOS; BMD; Hypertension; Glucose levels; Height; Menopause

Disease

Uitterlinden et al, 2004

Genetics of PCOS

Two candidate gene studies:

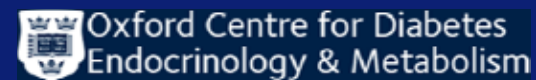
- The insulin receptor pathway: replication of association

In collaboration with: professor Goodarzi and professor Azziz, Cedars-Sinai Medical Center

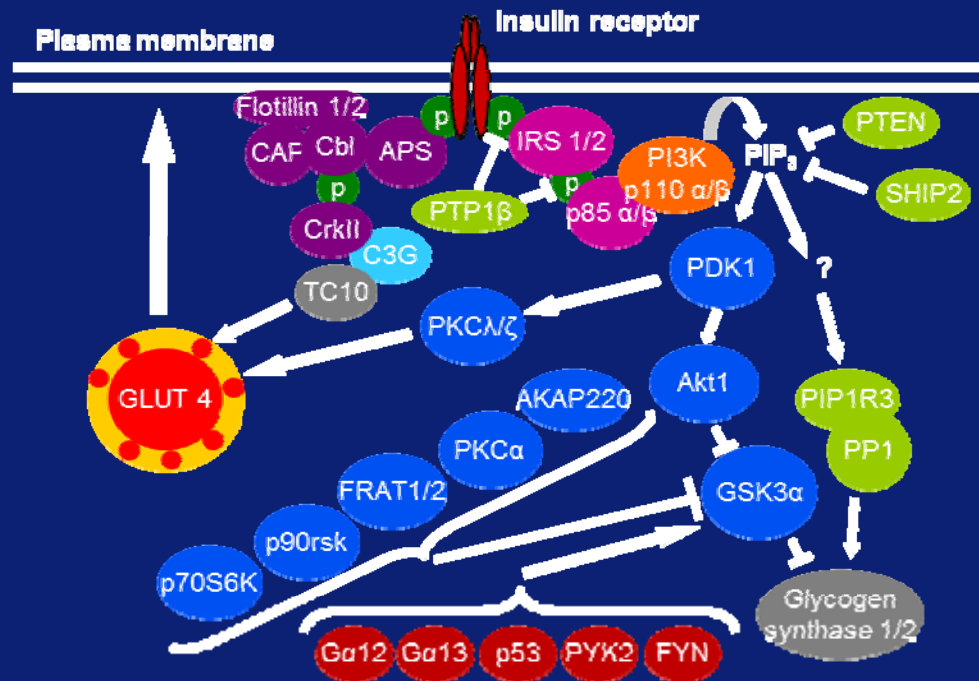


- BMI susceptibility genes: genetic risk score

In collaboration with: professor McCarthy and professor Franks, Oxford University



Insulin signaling pathway



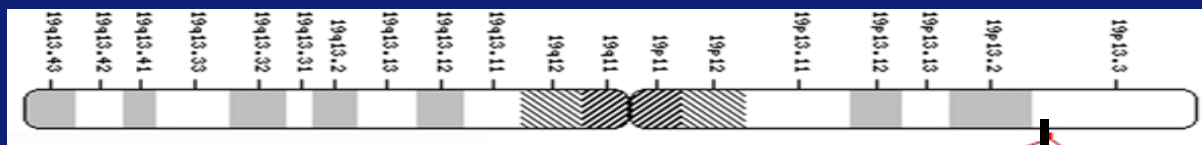
Discovery stage
295 variants in 39 genes

Insulin receptor (INSR)
4 SNPs associated with PCOS

Replication

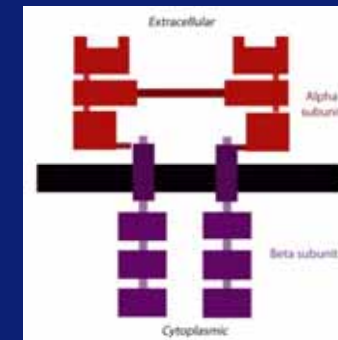
In collaboration with Goodarzi and Azziz et al, unpublished data

Insulin receptor polymorphisms and PCOS



chromosome 19

Insulin receptor (INSR)



SNP	Discovery cohort			Replication cohort			Meta-analysis
	OR	95% CI	P-value	OR	95% CI	P-value	P-value
rs12459488	0.62	0.40-1.0	0.05	1.11	0.92-1.34	0.30	0.71
rs12971499	0.68	0.42-1.11	0.12	1.05	0.86-1.28	0.067	0.21
rs2252673	1.99	1.17-3.58	0.011	1.32	1.08-1.60	0.006	0.00066
rs10401628	0.45	0.255-0.801	0.0065	1.00	0.80-1.25	0.99	0.40

In collaboration with Goodarzi and Azziz et al, unpublished data

Genetics of PCOS

Two candidate gene studies:

- The insulin receptor pathway: replication of association

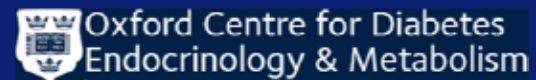
In collaboration with: professor Goodarzi and professor Azziz, Cedars-Sinai Medical Center



A tagging SNP approach identified novel INSR SNPs associated with PCOS, one of which confirmed association with PCOS in a replication cohort

- BMI susceptibility genes: genetic risk score

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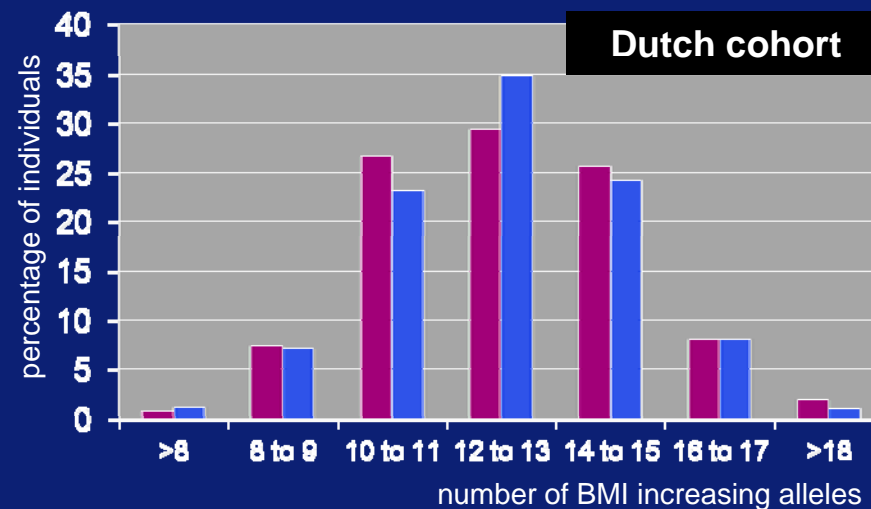
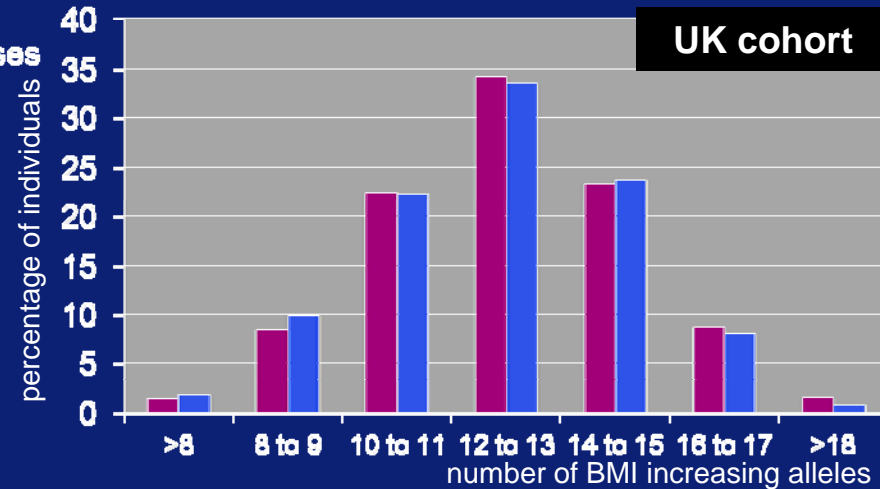


BMI susceptibility genes and PCOS (1)

Genetic Risk Score (GRS)

■ PCOS cases
■ controls

nearest gene	SNP	chromosome
BDNFOS	rs4074134	11
FAIM2	rs7138803	12
ETV5	rs7647305	3
FTO	rs9939609	16
GNPDA2	rs10938397	4
KCTD15	rs11084753	19
MC4R	rs17782313	18
MTCH2	rs10838738	11
NEGR1	rs2815752	1
SEC16B	rs10913469	1
SH2B1	rs7498665	16
TMEM18	rs6548238	2

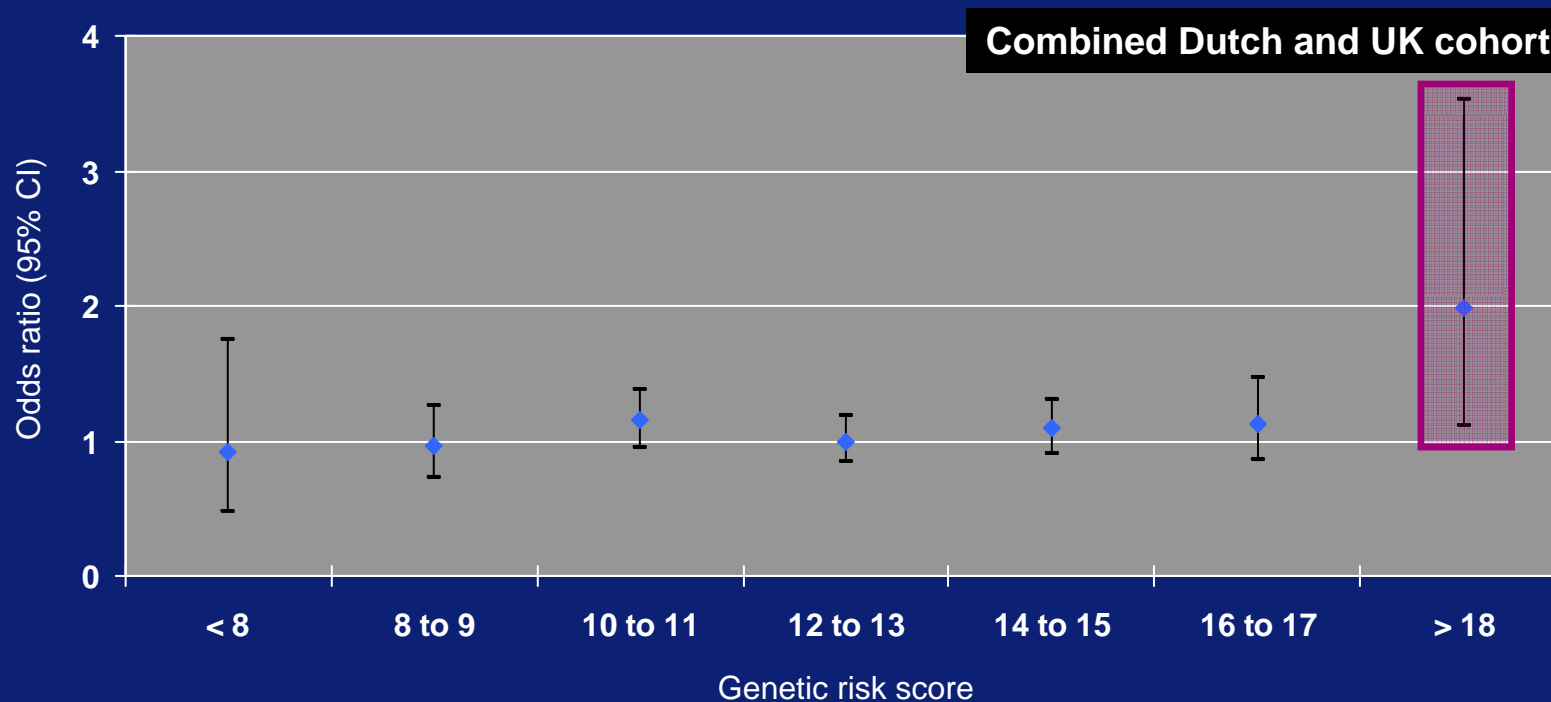


In collaboration with McCarthy et al, unpublished data



BMI susceptibility genes and PCOS (2)

GRS for BMI increasing alleles and risk for PCOS



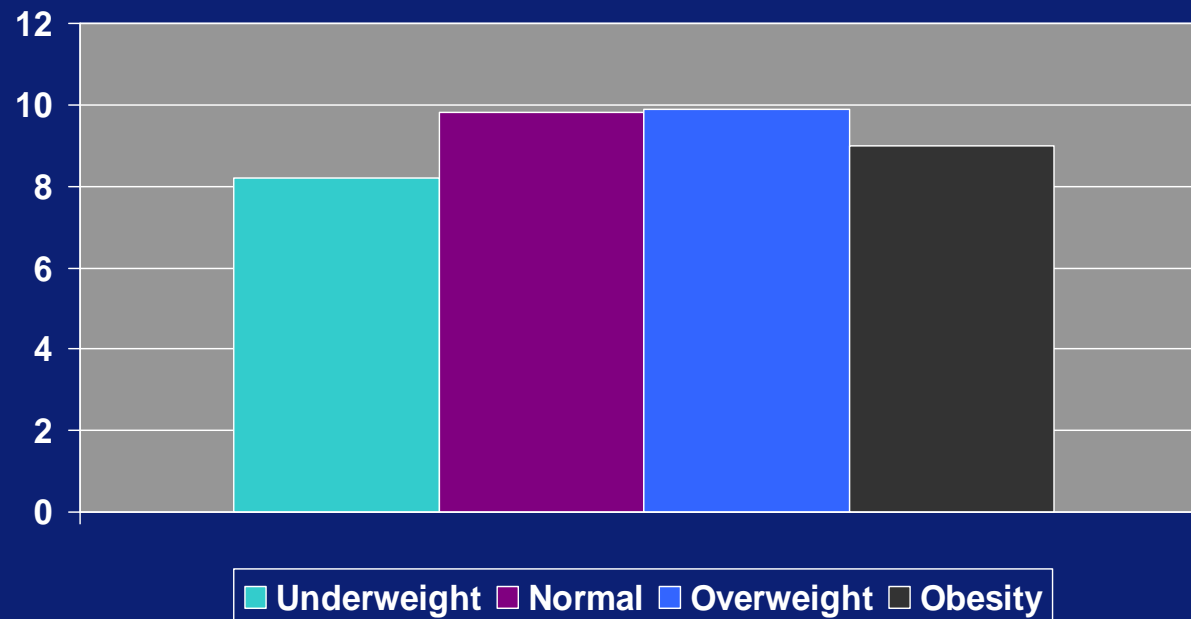
Number of BMI risk alleles carried is not associated with an increased risk of PCOS

In collaboration with McCarthy et al, unpublished data



BMI susceptibility genes and PCOS (3)

Incidence of PCOS



Genetics of PCOS

Two candidate gene studies:

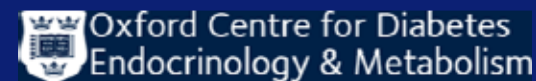
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