

Introduction

- Smoking and caffeine consumption are highly correlated
- The mechanism underlying this association is unknown
- Possibly explanation: shared genetic and/or environmental factors
- The *bivariate twin model* estimates genetic & environmental correlations
- Genome wide association (GWA) studies provide an additional method of estimating genetic overlap: *LD-Score regression*¹

Methods

Bivariate twin models

Data: survey data in 10,620 participants from the Netherlands Twin Register (mean age 32.5 [SD 14.5], 68.1% female)

Measures: current smoking (0=never/former smoking, 1=current smoking) | coffee use (0=low, 1=high) | total caffeine use (0=low, 1=high)

Analyses: monozygotic (MZ) twins share 100% and dizygotic (DZ) twins ~50%, of their segregating genes; both may share their environment

- if $r_{MZ} > r_{DZ}$ → additive genetic influences (A)
- if $r_{DZ} > 0.5 r_{MZ}$ → common environmental influences (C)
- If $r_{MZ} < 1$ → unique environmental influences (E)

LD-Score regression¹

Data: summary statistics from GWA meta-analyses of smoking² & coffee³

Measures: number of cigarettes per day | smoking initiation (0=never smoking, 1=former/current smoking) | smoking persistence (0=former smoking, 1=current smoking) | cups of coffee per day

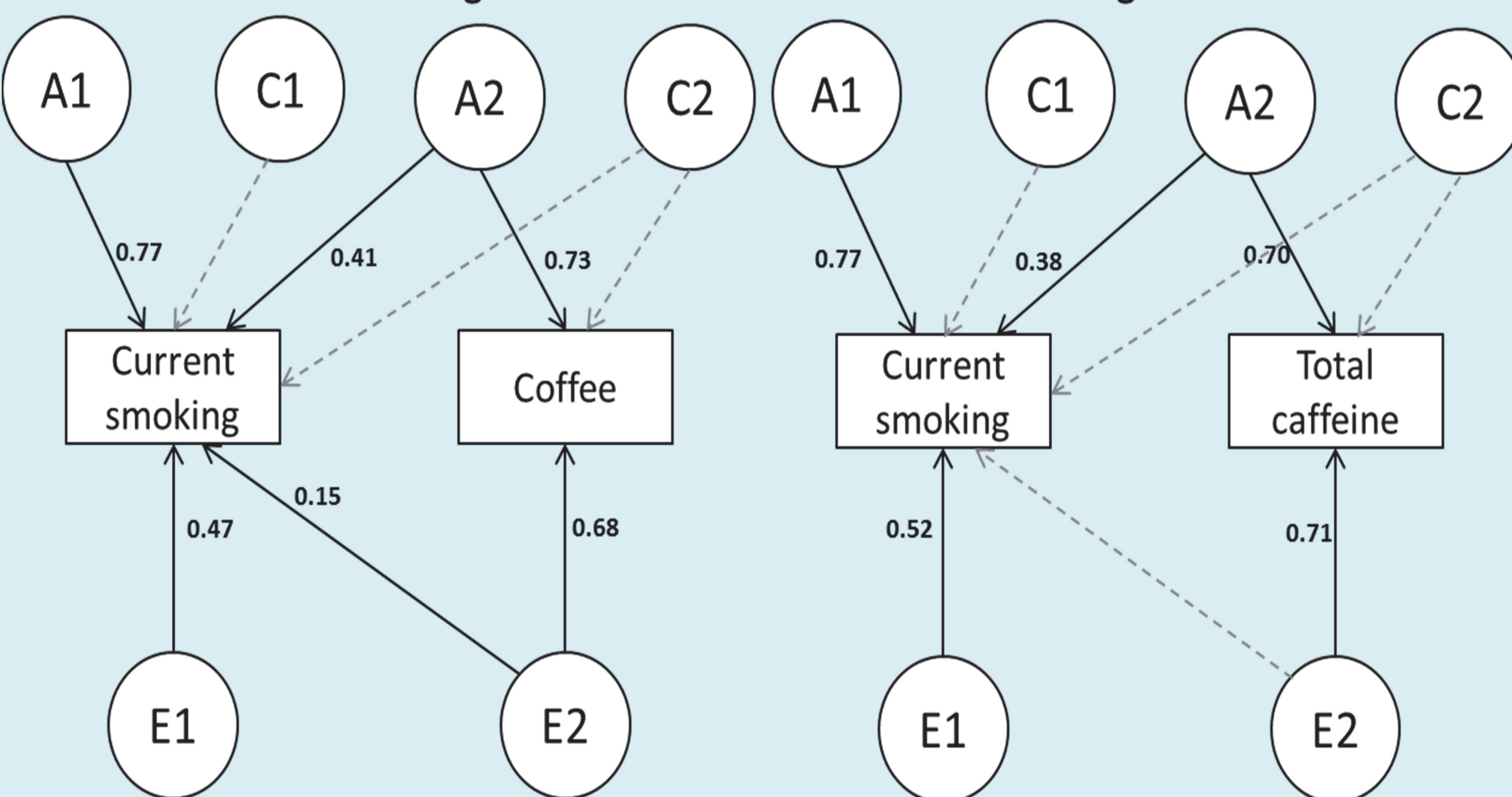
Analyses: the expected product for the Z scores of the association between a SNP and two phenotypes (smoking and coffee) is modelled as a function of LD-score (= degree of Linkage Disequilibrium a SNP has with all neighboring SNPs).

Results: bivariate twin models

ACE models (dotted lines were not significant)

Current smoking & coffee

Current smoking & total caffeine



Genetic correlation (A) = **r0.47**
 Unique environmental correlation (E) = **r0.30**

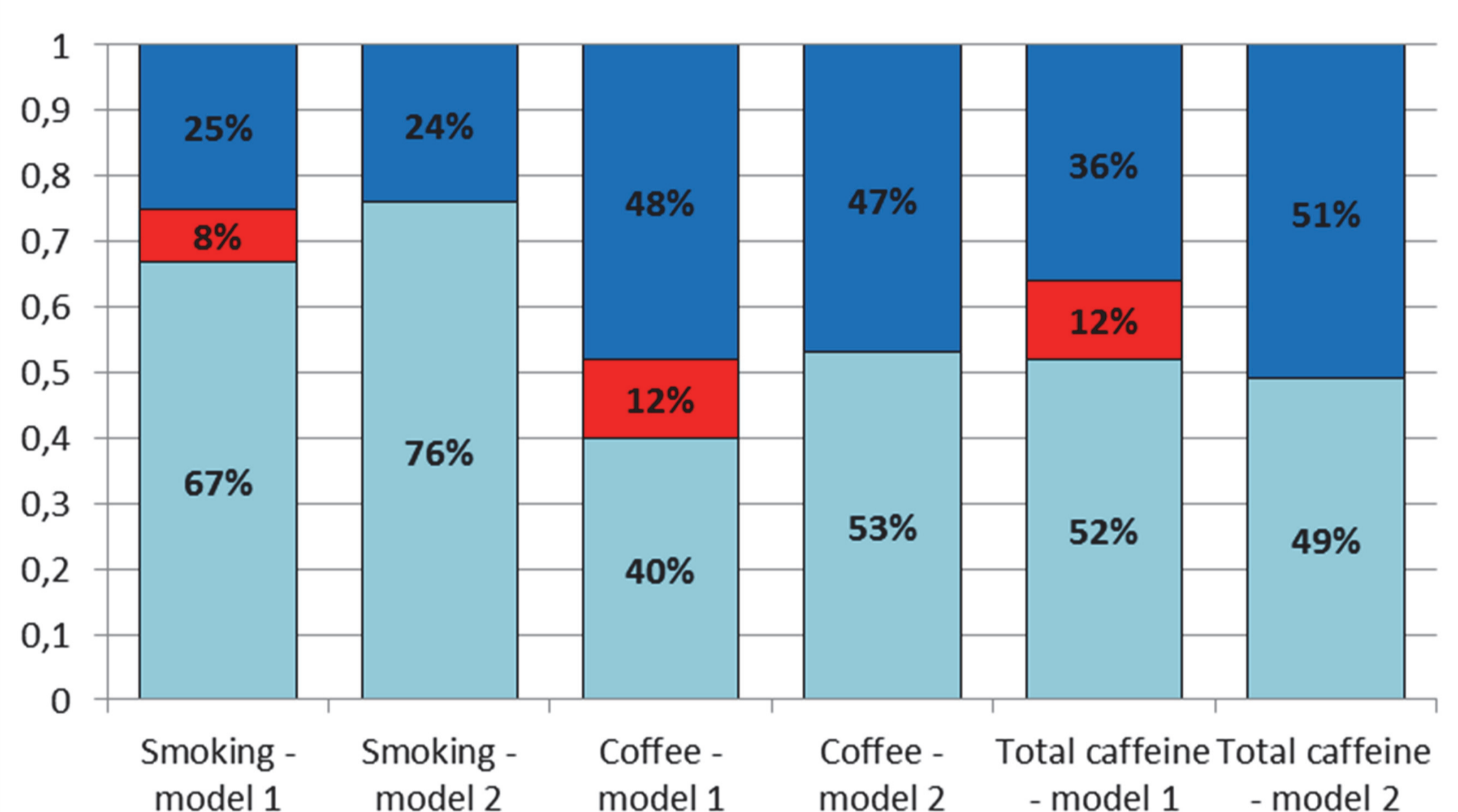
Genetic correlation (A) = **r0.44**

Results: LD Score regression (genetic correlations)

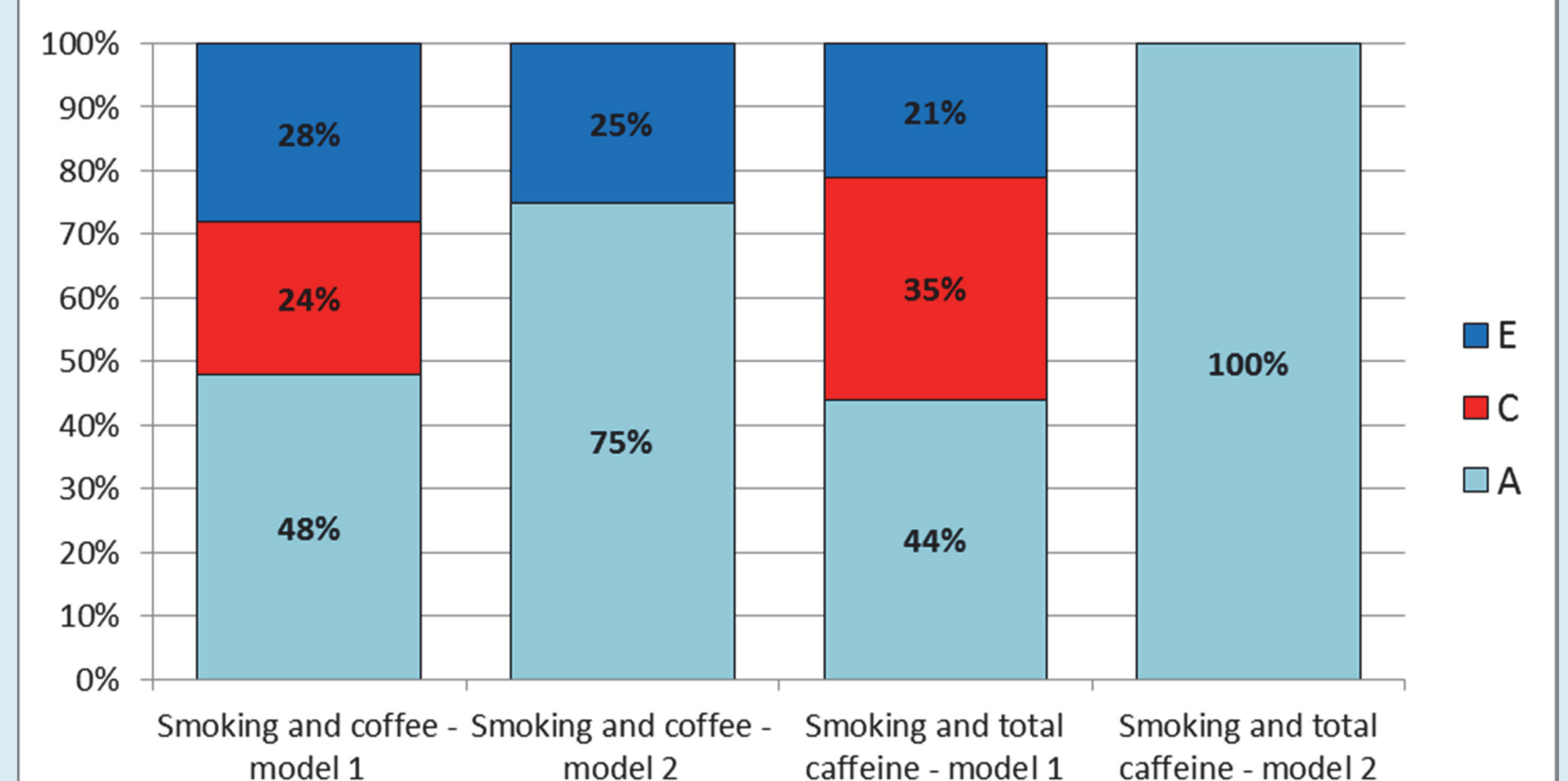
- Cigarettes per day & cups of coffee per day = **r0.44**
- Smoking initiation & cups of coffee per day = **r0.28**
- Smoking persistence & cups of coffee per day = **r0.25**

Results: bivariate twin models

Genetic/environmental influences on smoking & caffeine



Genetic/environmental influences on overlap



Model 1 = initial model, **Model 2** = best-fitting model
A = additive genetic influences, **C** = common environmental influences, **E** = unique environmental influences

Conclusions

- Consistent, moderate genetic correlation between smoking and caffeine consumption
- Genetic factors explain most of the association between smoking & caffeine (similar to previous US-based twin studies)
- Initiating smoking may be especially undesirable for heavy caffeine consumers, given their genetic susceptibility

References:

- Bulik-Sullivan *et al.* (2015)
- Tobacco and Genetics Consortium (2010)
- The Coffee and Caffeine Genetics Consortium (2014)

Contact: j.l.treur@vu.nl | nl.linkedin.com/in/jorientreur