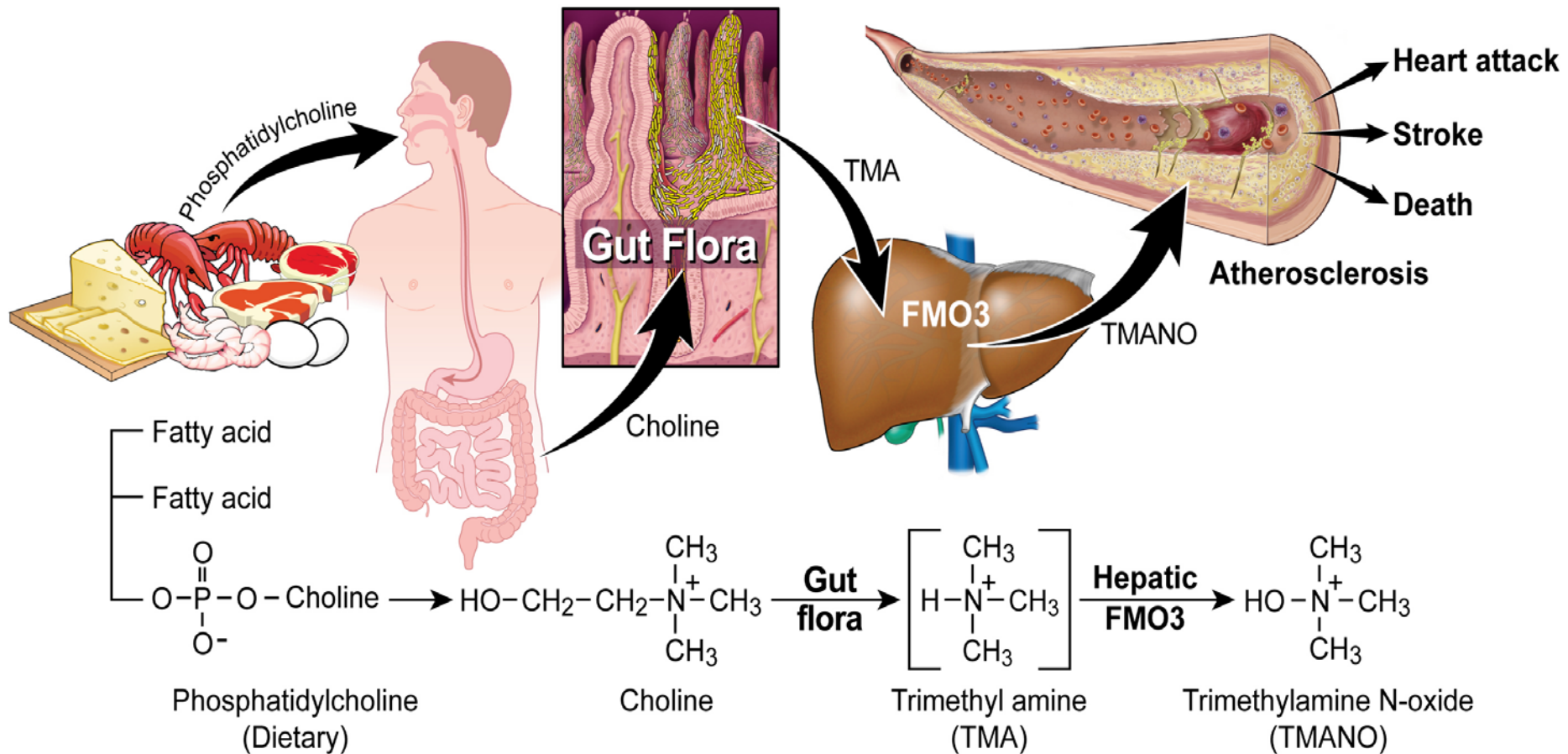


Gut flora
dependent
metabolism of
phosphatidylcholine
and the
pathogenesis of
cardiovascular
disease

Stanley Hazen MD PhD

Take home summary:

Gut flora participates in atherosclerosis in the presence of specific dietary exposures



Phase 1: Discovery-based investigations

Metabolomics screening and structural
identification

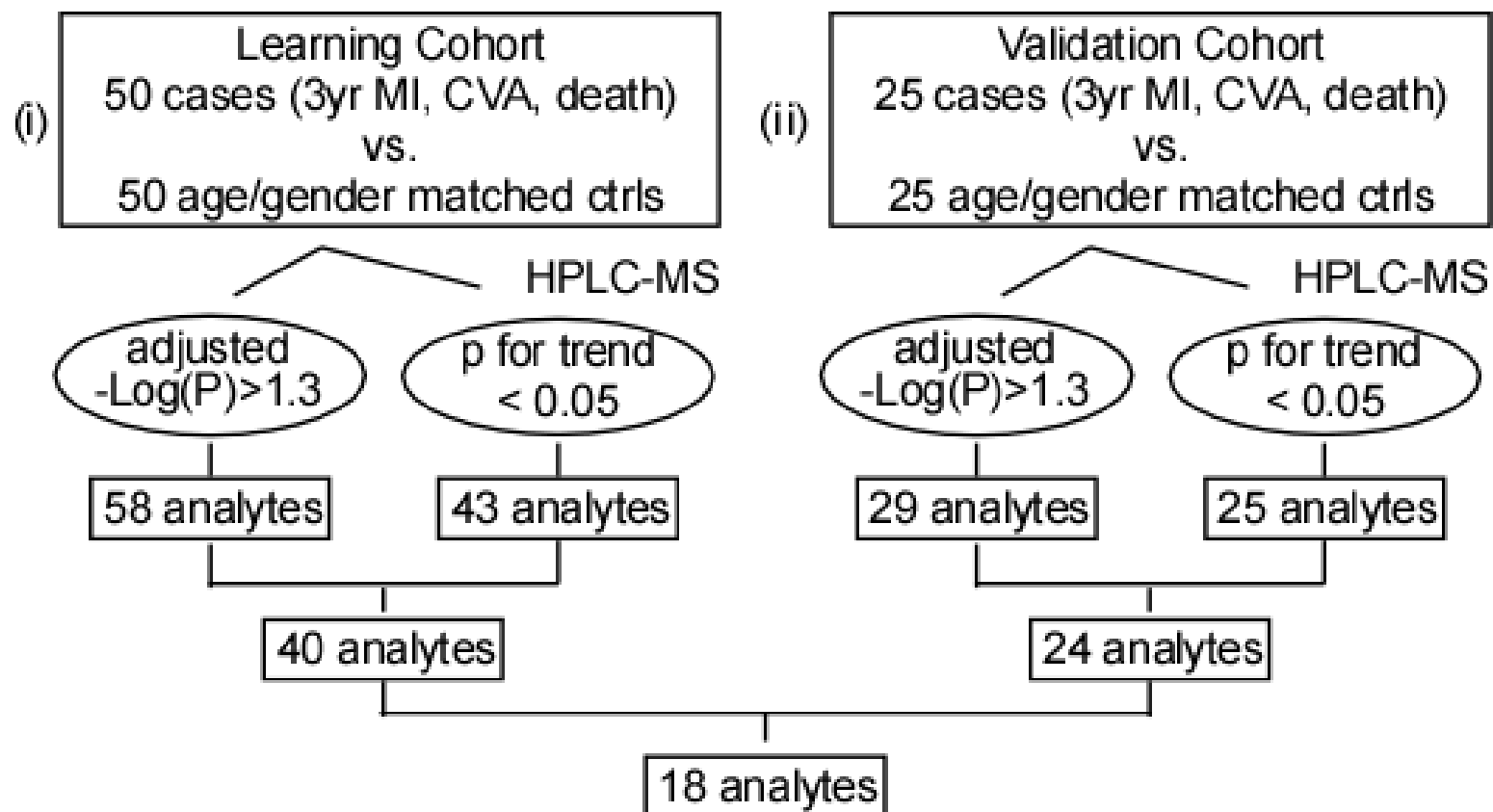
Phase 2: Clinical validation

Demonstration of clinical utility

Phase 3: Mechanistic studies

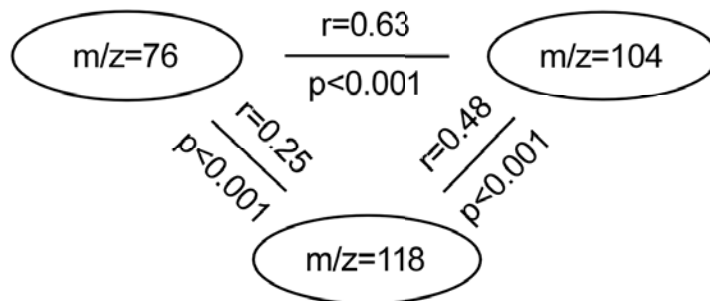
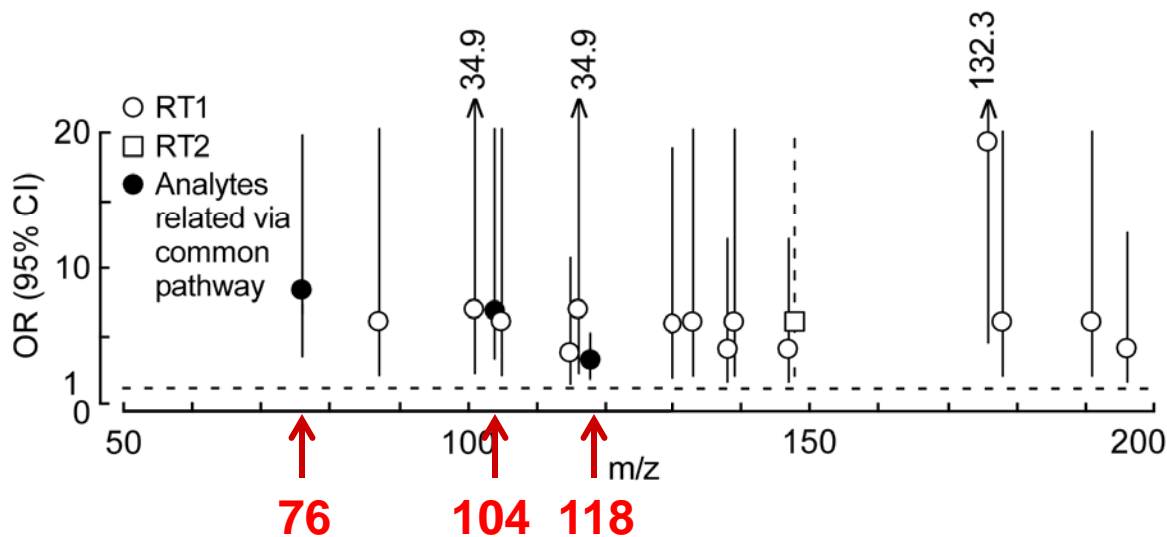
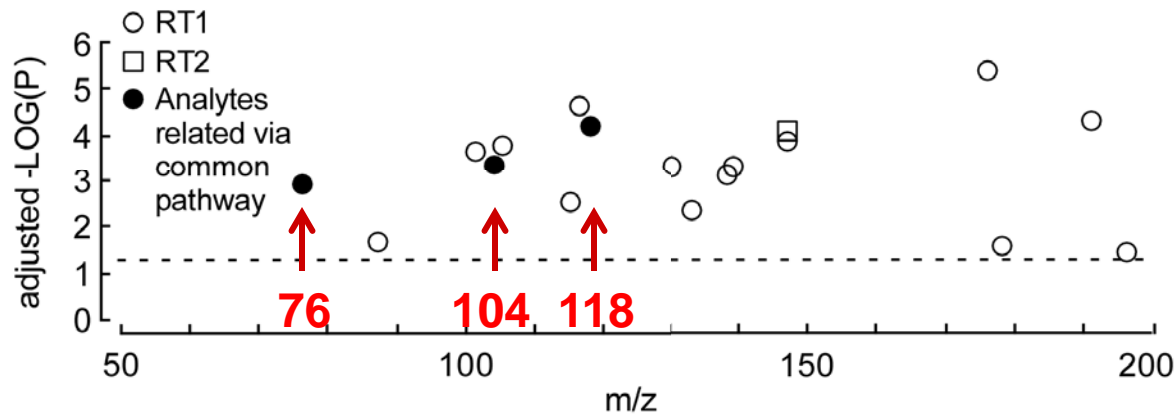
Demonstration of causality for novel pathway

Strategy of metabolomics study design for identifying unbiased small molecule profiles predictive of incident risks for major adverse cardiovascular events



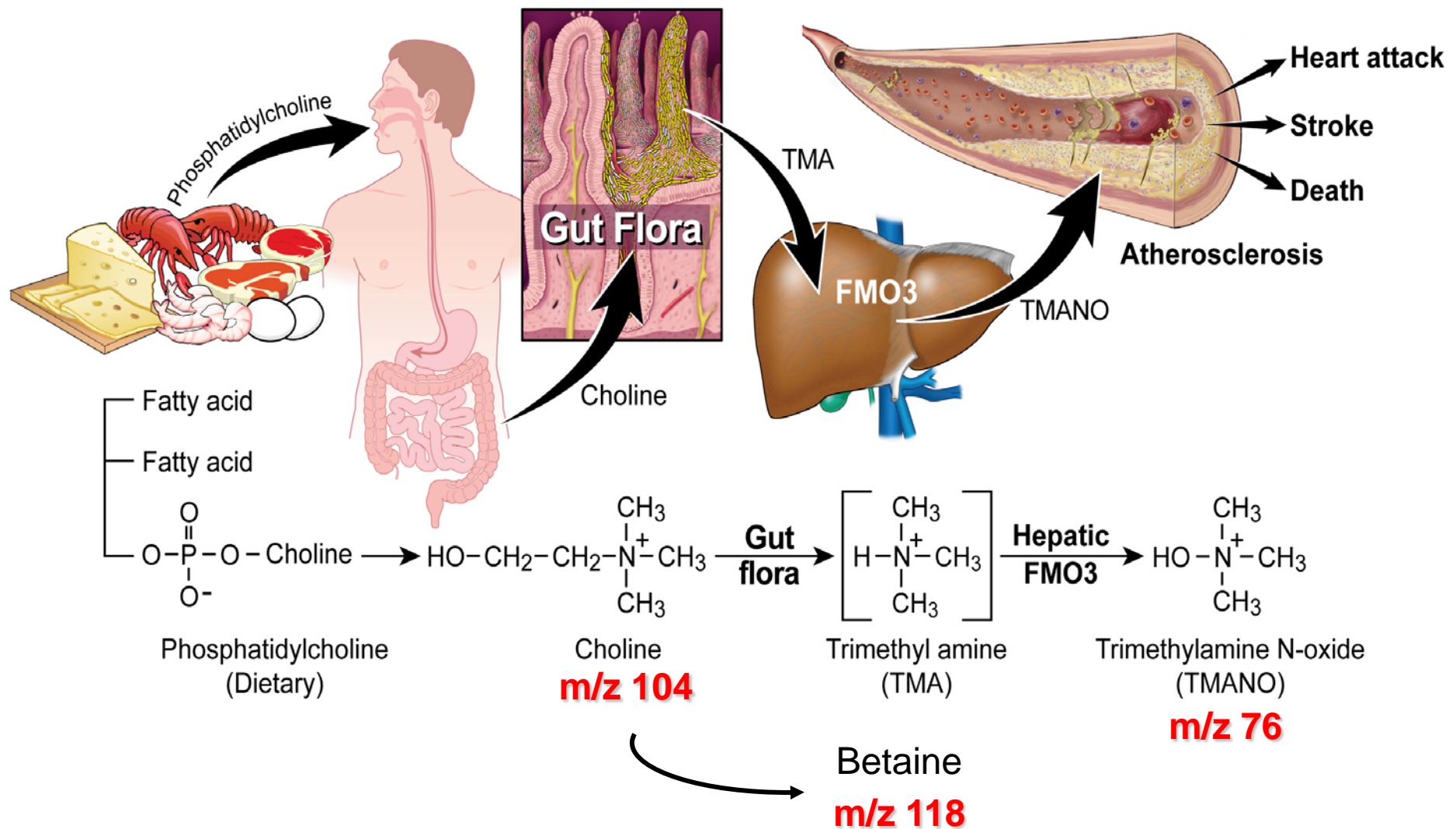
(ii) Structural identification of analytes

(iv) Confirm clinical prognostic utility in Independent Prospective Cohort (N>1000)



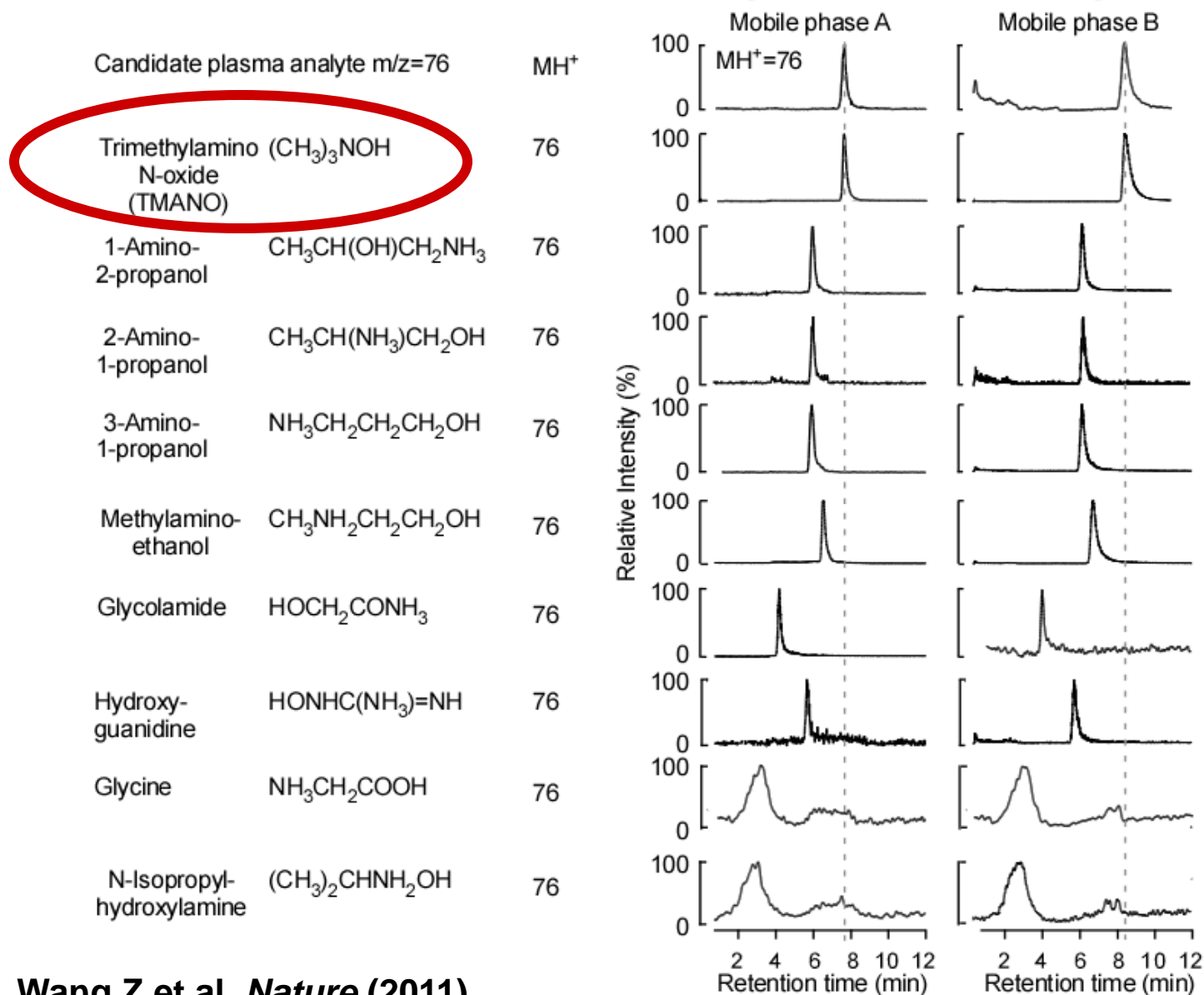
Plasma analytes with m/z 76, 104, and 118 are associated with CVD, show a dose-response relationship with MACE and are correlated, suggesting participation in a common pathway

Choline, betaine and trimethylamine N-oxide are the plasma analytes associated with CVD



But the metabolomics library says its "glycine"...

Candidate structures of plasma analyte $m/z=76$

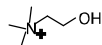


MH⁺

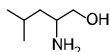
MS1, m/z=104

Plasma analyte m/z=104

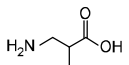
choline



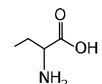
104

2-amino-3-methyl-
1-butanol

104

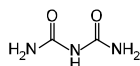
3-aminoisobutyric
acid

104

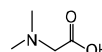
2-aminobutyric
acid

104

biuret



104

N,N-dimethyl
glycine

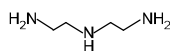
104

benzonitrile



104

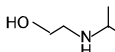
diethylenetriamine



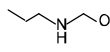
104

ethyl-N-hydroxyl
acetimidate

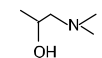
104

2-isopropyl-
aminoethanol

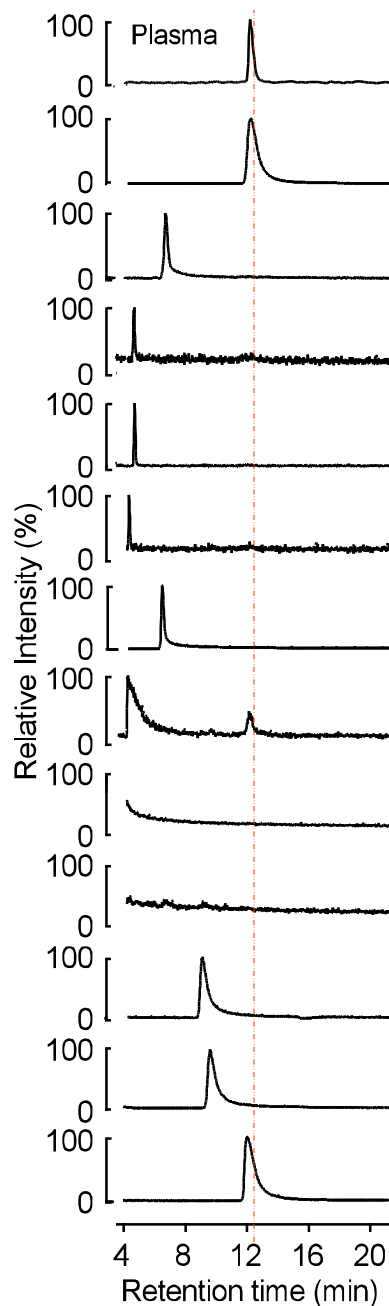
104

2-propyl-
aminoethanol

104

1-dimethyl-amino-
2-propanol

104



**Candidate plasma
analytes linked to CVD
risks with m/z 104**

**Identity as Choline was
confirmed by:**

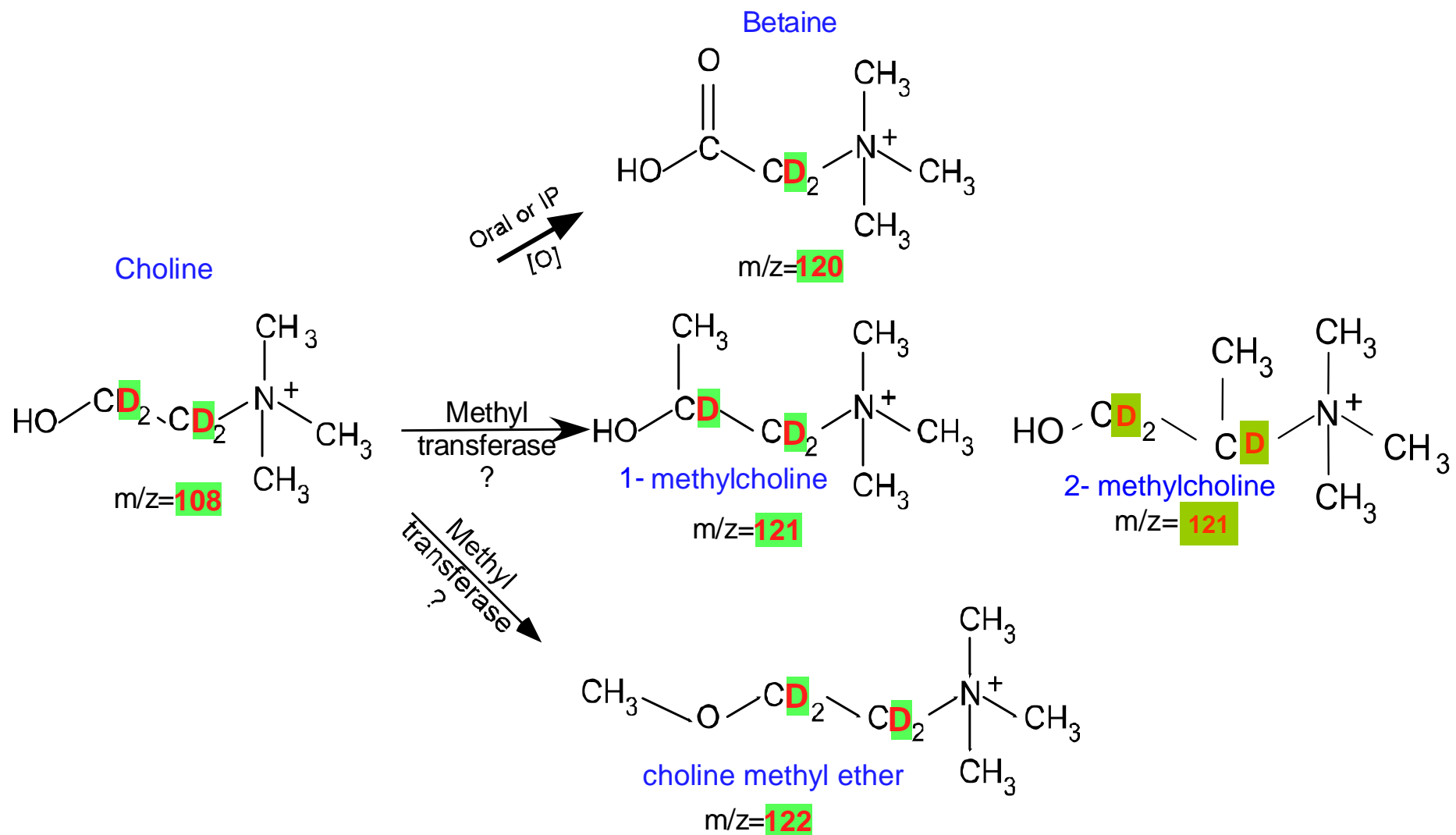
LC-MSⁿ

GC/MS/MS

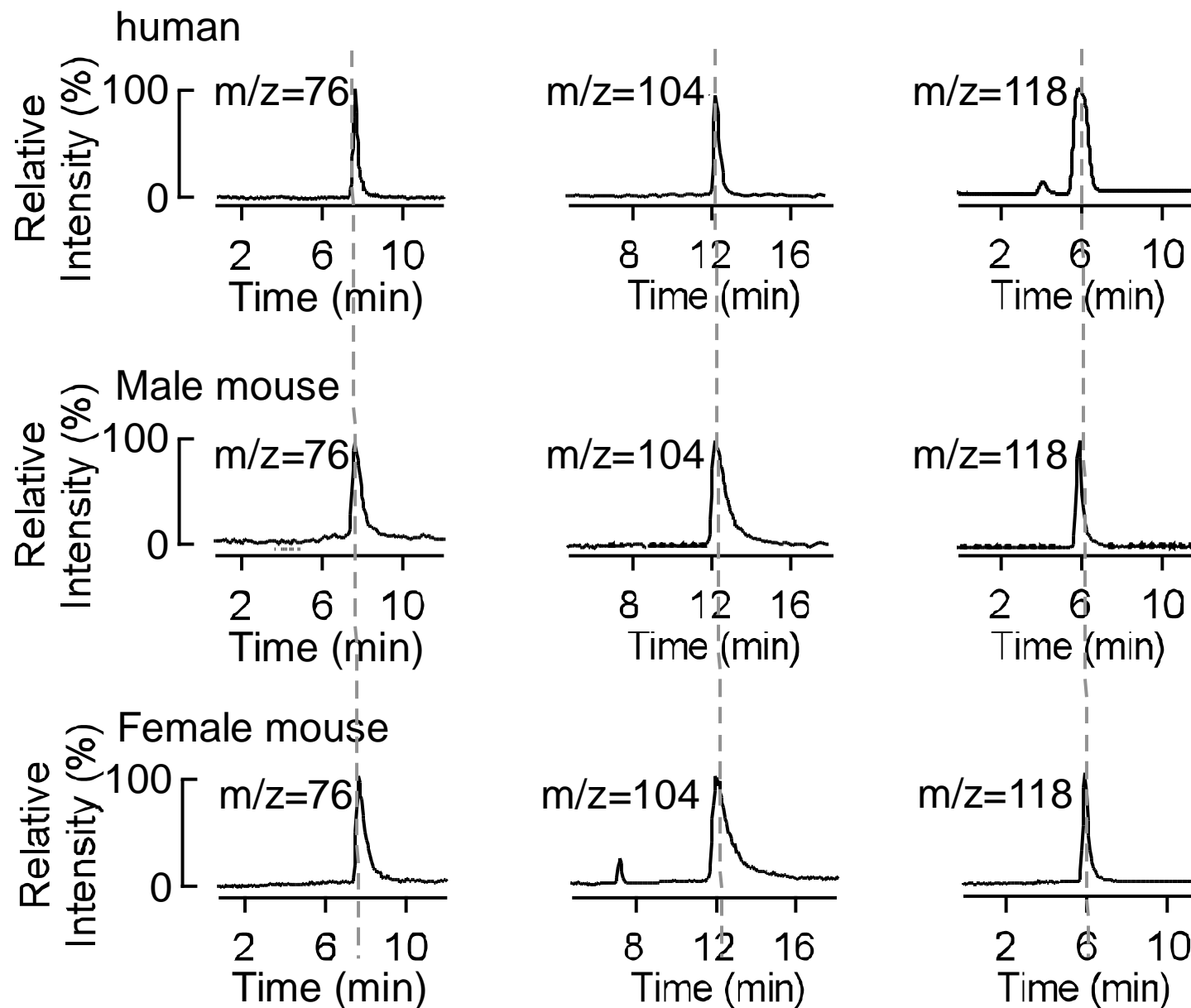
¹H, ¹³C, and ¹⁵N NMR

**Isotope tracer studies:
d₉-choline and
d₄-choline**

Strategy to determine the analyte at $m/z=118$ by choline deuterated isotopologue feeding study

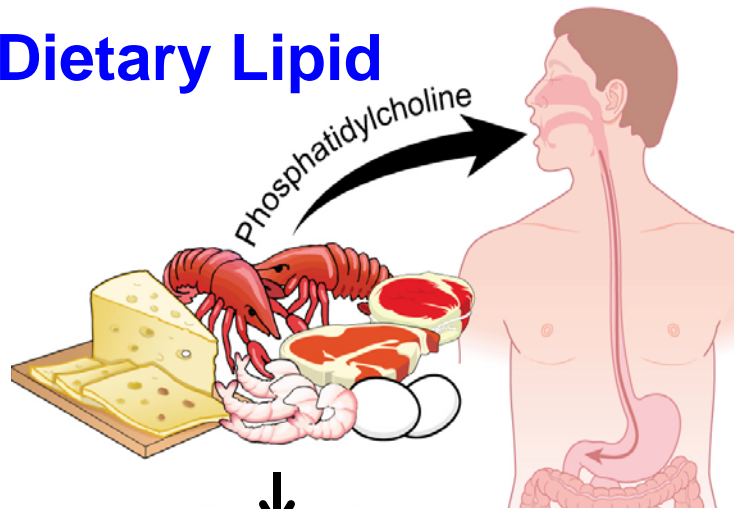


Dietary egg yolk PC produces increases in analytes with m/z 76, 104, and 118 in both human and mouse plasma

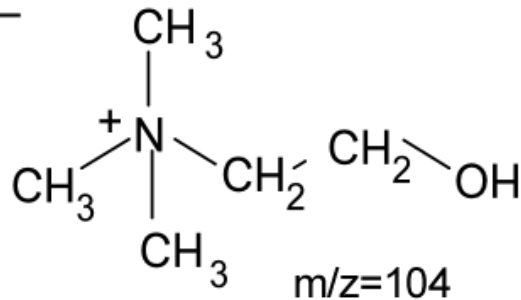


Dietary Lipid

What is the role of gut flora?



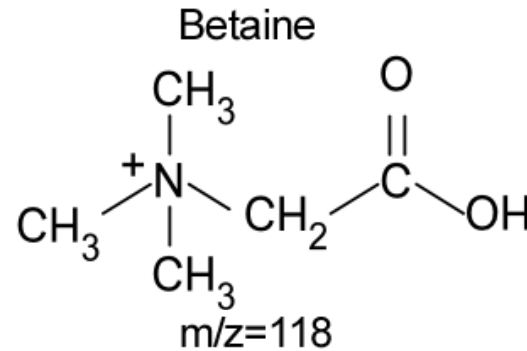
Choline



d4-choline: $(\text{CH}_3)_3^+\text{NCD}_2\text{CD}_2\text{OH}$

d9-choline: $(\text{CD}_3)_3^+\text{NCH}_2\text{CH}_2\text{OH}$

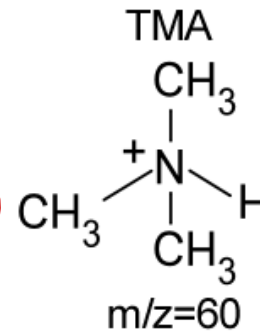
Oral or i.p.
[O]



d4 metabolite $m/z=120$

d9 metabolite $m/z=127$

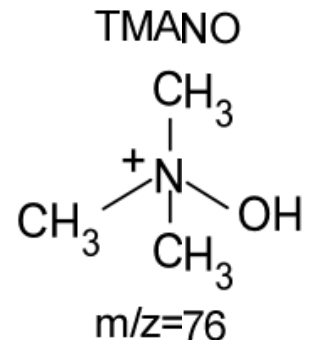
Oral only
Gut Flora



d4 metabolite $m/z=60$

d9 metabolite $m/z=69$

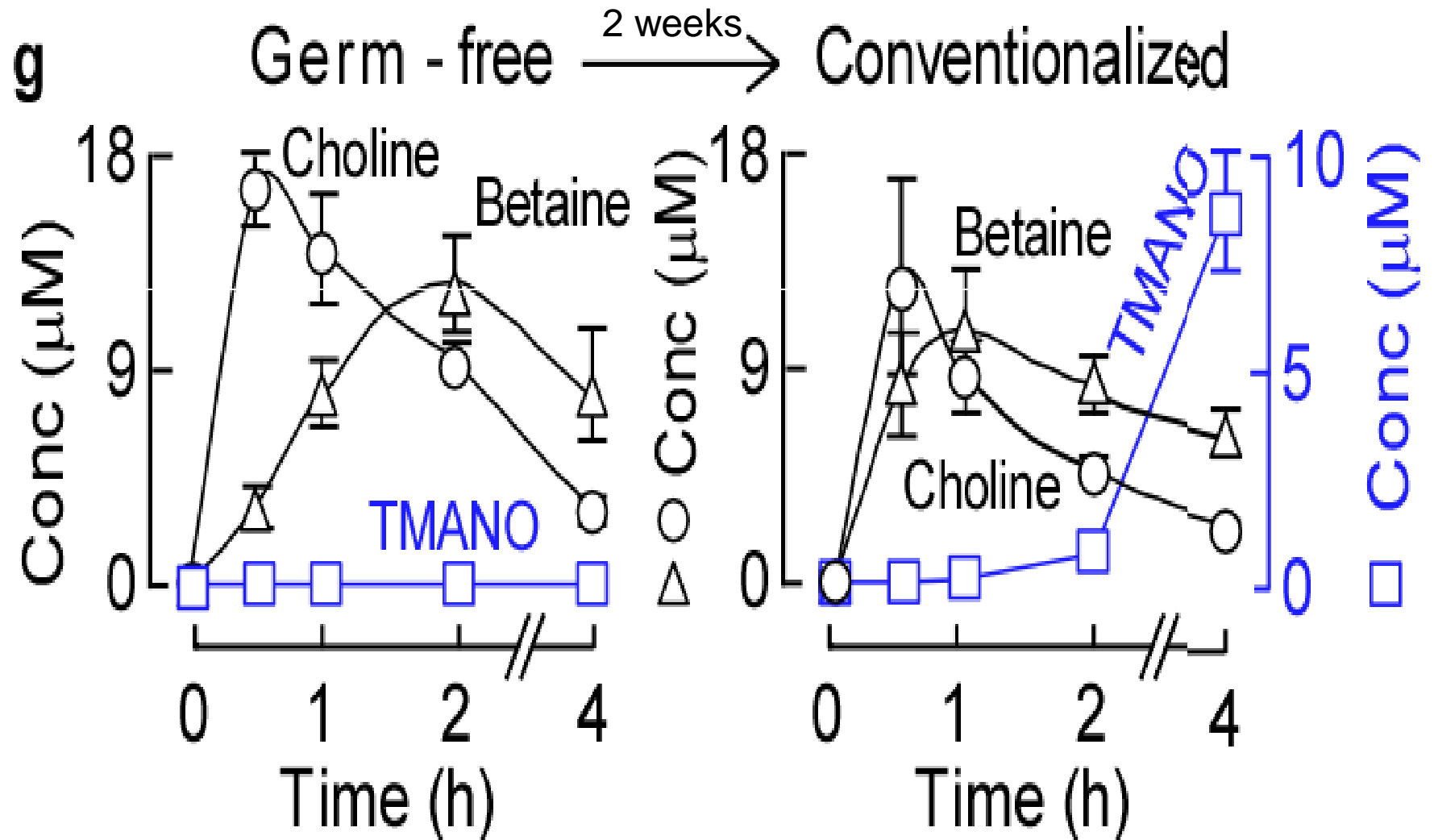
presumed
FMO3
[O]



d4 metabolite $m/z=76$

d9 metabolite $m/z=85$

Intestinal Microbial Organisms Play an Obligatory Role in TMANO Generation from Dietary Egg Yolk PC



Phase 1: Discovery-based investigations

Metabolomics screening and structural identification

Phase 2: Clinical validation

Demonstration of clinical
prognostic utility

Phase 3: Mechanistic studies

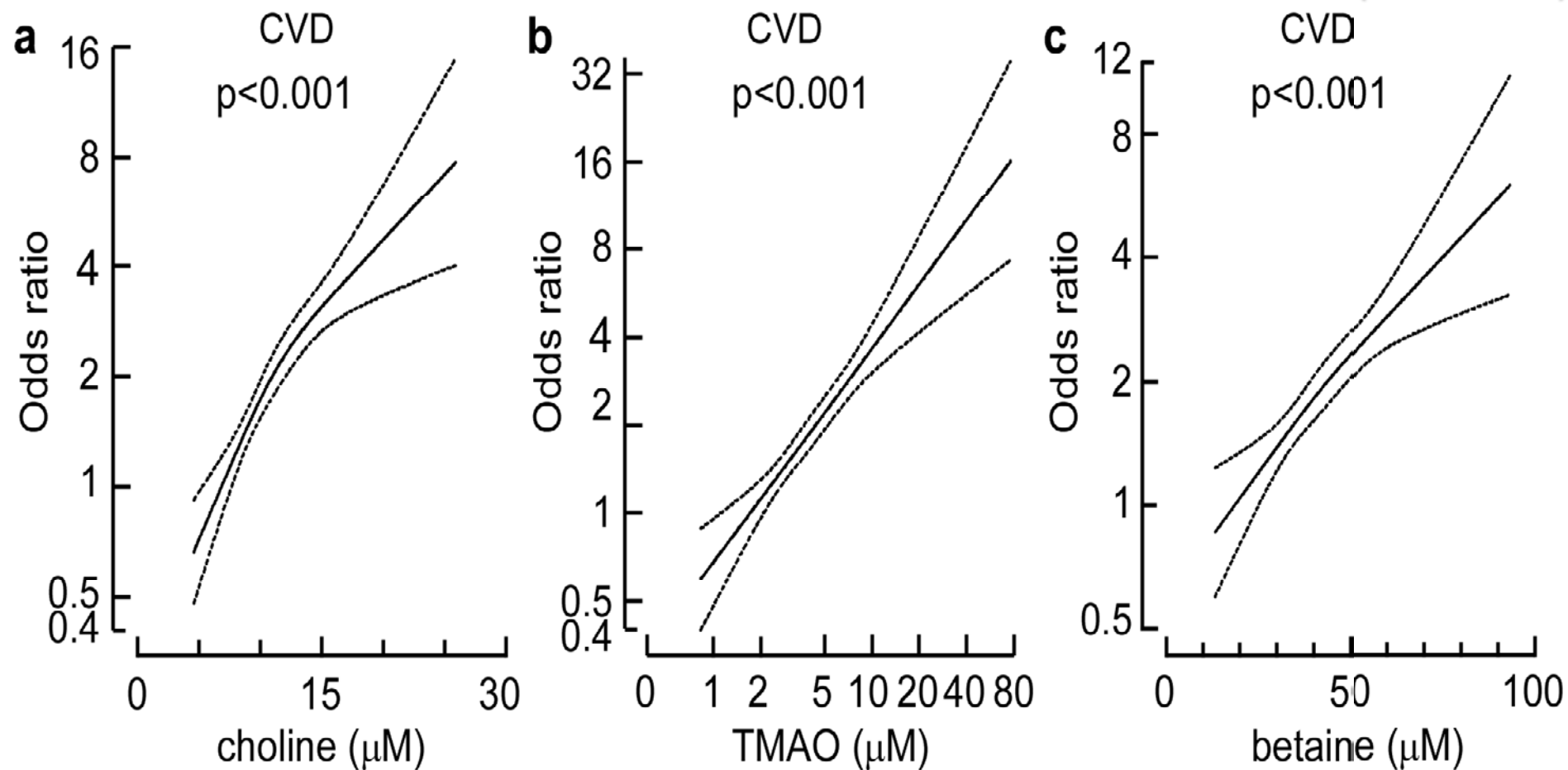
Demonstration of causality for novel pathway

Prospective Cohort: N=1865 Sequential Cardiology Patients

a. Demographics of CVD prevalence

Characteristic	Non-CVD	CVD	P value
Age, mean (SD), y	61.2 (7.8)	65.3 (9.8)	<0.001
Women, %	52.5	49.8	0.59
Diabetes, %	11.1	32.8	<0.001
Hypertension, %	65.3	81.7	<0.001
History of smoking, %	52.2	56.7	0.36
Current smoking, %	5.3	5.9	0.80
LDL cholesterol, median (IQR), mg/dL	108 (85-132)	95 (78-122)	<0.001
HDL cholesterol, median (IQR), mg/dL	50 (40- 63)	42 (34-53)	<0.001
Triglycerides, median (IQR), mg/dL	116 (81-163)	139 (103-200)	<0.001
CRP, median (IQR), mg/dL	2.2 (1.2-4.6)	3.3 (1.5-7.3)	<0.001
Framingham Risk Score, median (IQR)	7.0 (5.0-9.0)	9.0 (6.0-12.0)	<0.001
MDRD (GFR), median (IQR)	75.5 (64.7-87.9)	71.6 (56.8-84.8)	0.004
ACE, % / Statin, % / Aspirin, %	34.7/29.8/55.1	56.4/64.9/75.5	each <0.001
TMANO, median (IQR), μ M	3.3 (2.2-5.0)	4.4 (2.8-7.4)	<0.001
Choline, median (IQR), μ M	10.9 (8.6-13.5)	12.3 (9.8-15.4)	<0.001
Betaine, median (IQR), μ M	5.6 (4.4-7.0)	5.9 (4.6-7.2)	0.02

Plasma choline, TMAO and betaine levels predict CVD risks (N=1865)



Odds ratio (95%CI) adjusted for age, sex, DM, HTN, smoking, LDL, HDL, TG, CRP, eGFR

Phase 1: Discovery-based investigations

Metabolomics screening and structural identification

Phase 2: Clinical validation

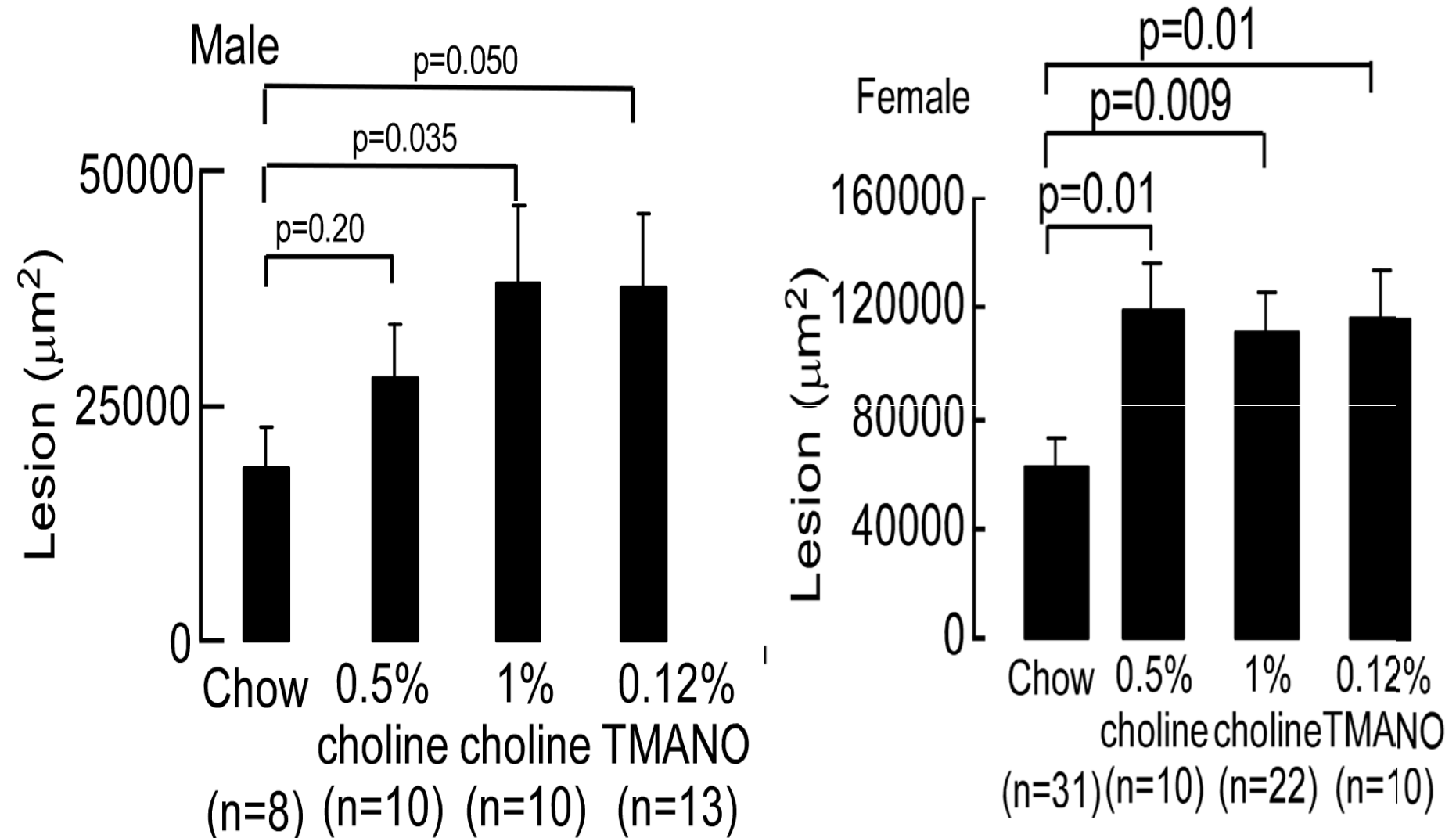
Demonstration of clinical utility

Phase 3: Mechanistic studies

Demonstration of causality for a
novel pathway

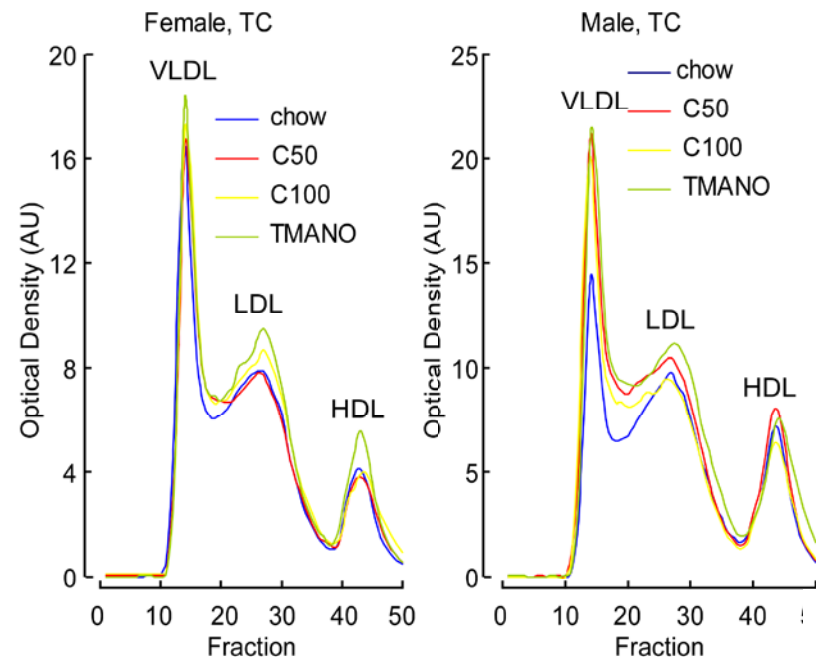
Dietary supplementation with choline or TMANO promotes atherosclerosis ApoE^{-/-} mice

Normal chow diet + indicated supplement weeks 4-20

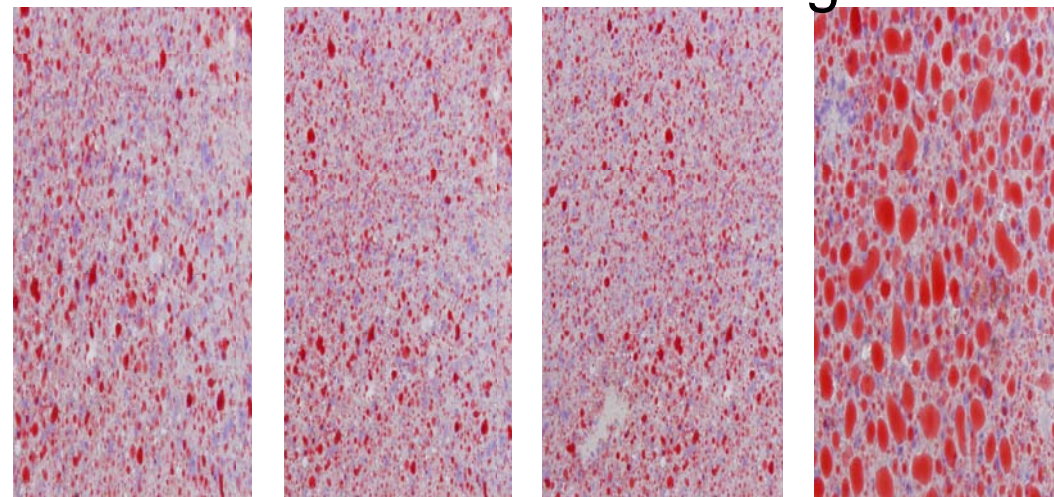


Dietary supplementation with choline or TMAO promotes atherosclerosis without dyslipidemia or hepatic steatosis

Lipids		Chow (n=9)	0.5% choline (n=10)	1.0% choline (n=10)	0.12% TMAO (n=13)
Plasma	Triglyceride (mg/dL)	114±16	150±22 (p=0.20)	99±13 (p=0.46)	104±8 (p=0.58)
	Cholesterol (mg/dL)	426±28	387±25 (p=0.32)	346±10 (p=0.02)	352±12 (p=0.04)
	Glucose (mg/dL)	238±28	290±18 (p=0.15)	191±33 (p=0.29)	248±20 (p=0.78)
Liver	Triglyceride (mg/g protein)	34.9±4.4	42.9±8.2 (p=0.47)	48.0±5.3 (p=0.07)	47.2±8.5 (p=0.22)
	Cholesterol (mg/g protein)	10.1±0.8	9.6±0.6 (p=0.64)	10.4±0.4 (p=0.72)	12.0±1.2 (p=0.18)

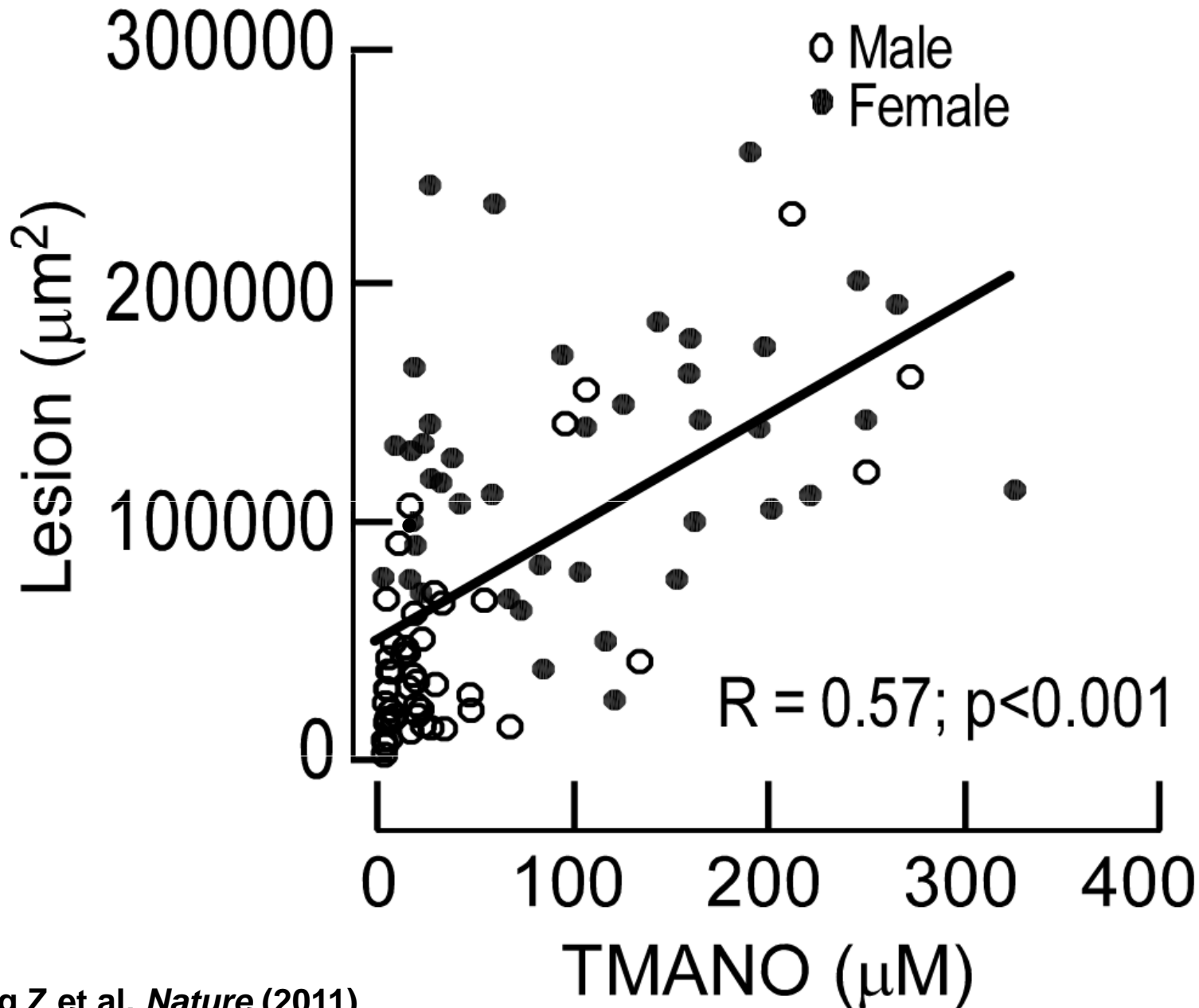


Liver oil-red-o staining

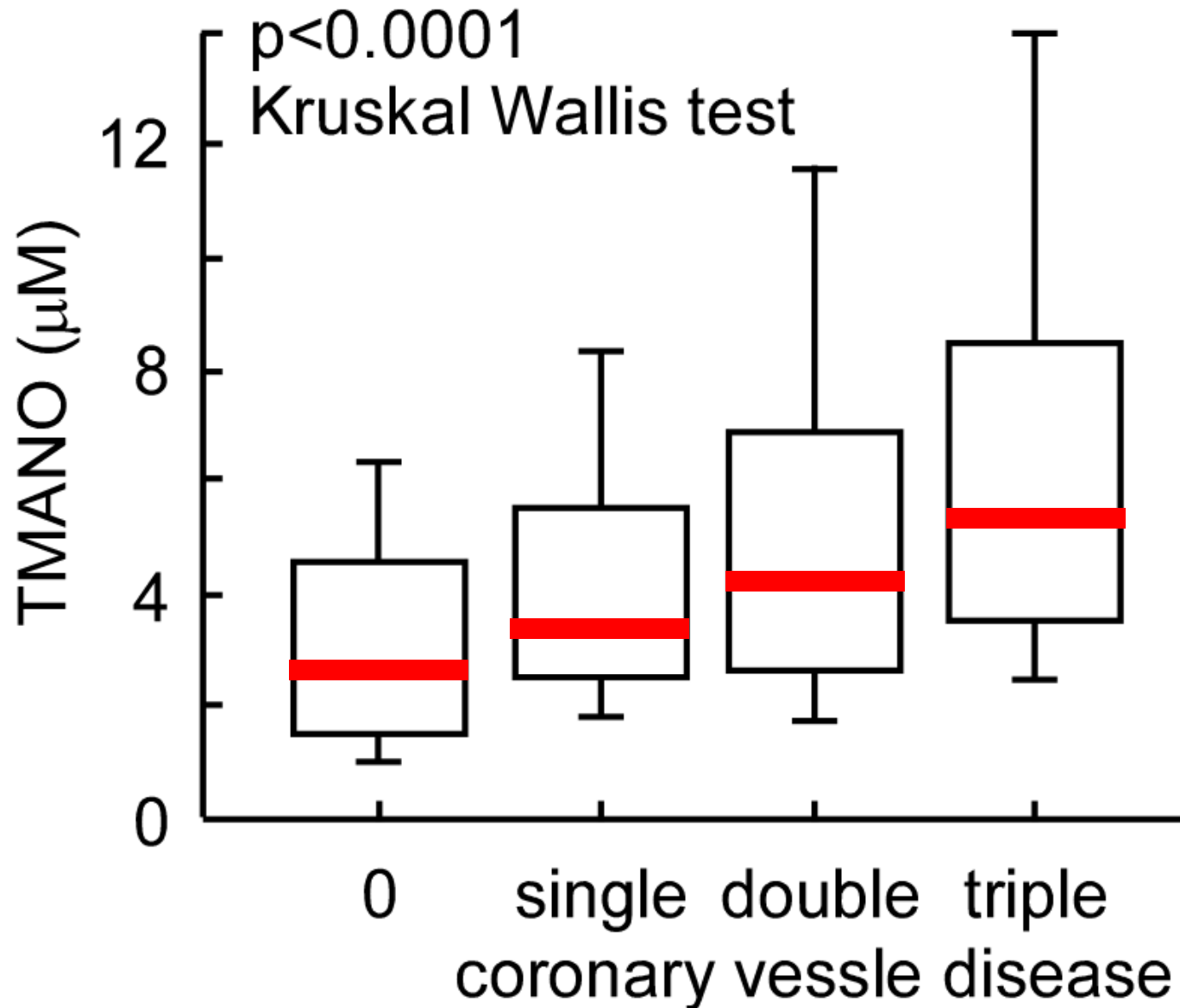


Chow 1% Choline 0.12% TMAO MCD

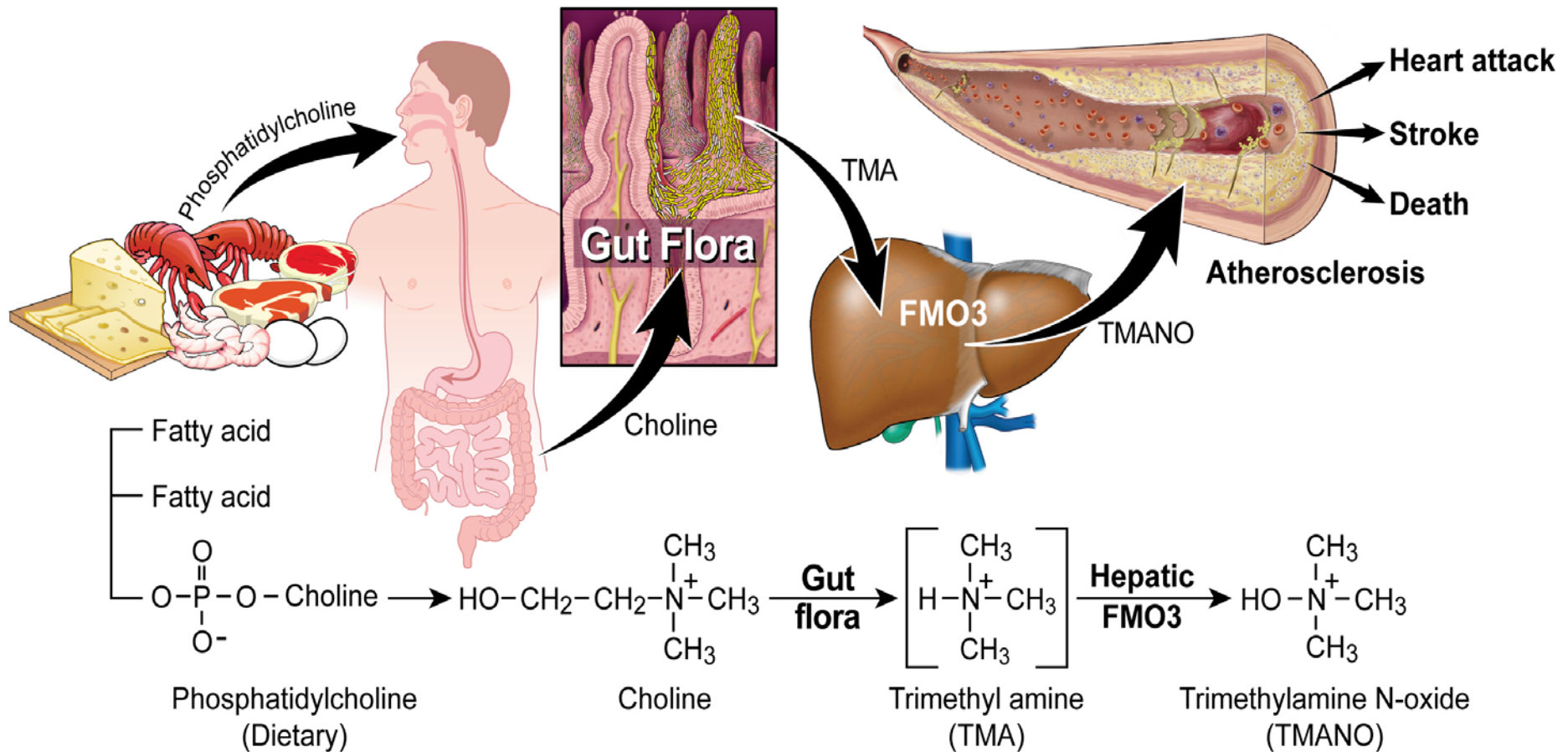
Plasma TMANO levels are correlated with aortic plaque in apoE-/-



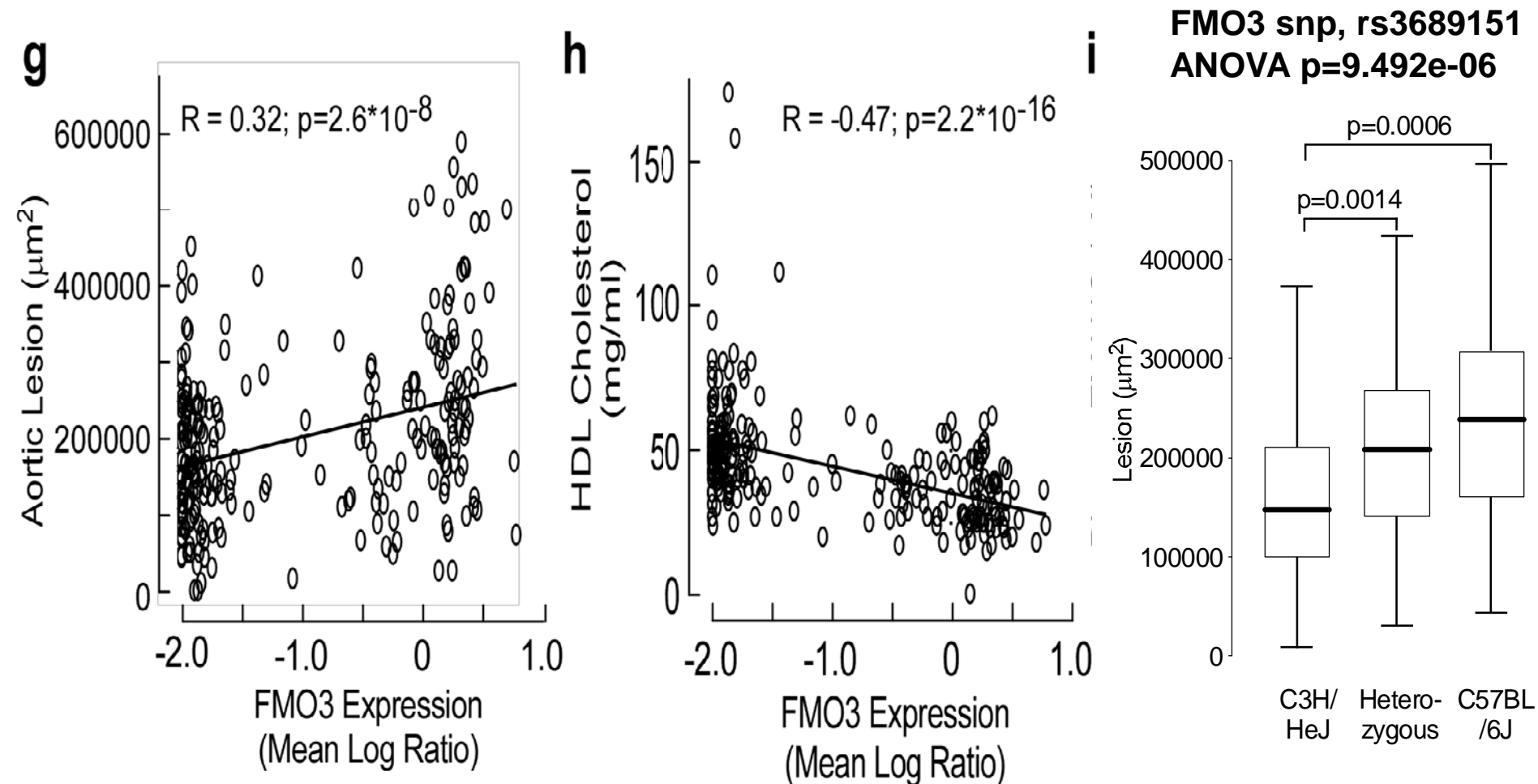
Plasma TMANO levels are correlated with angiographic coronary artery disease severity in subjects (N=1020)

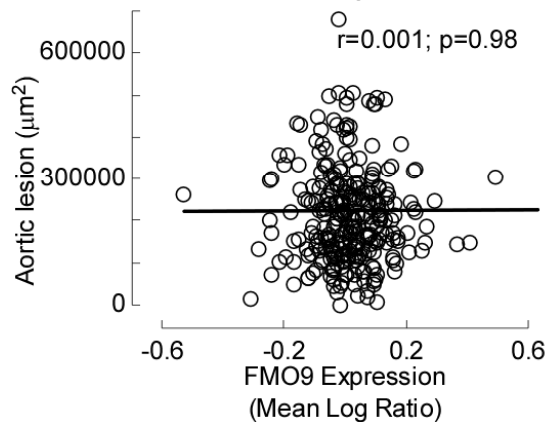
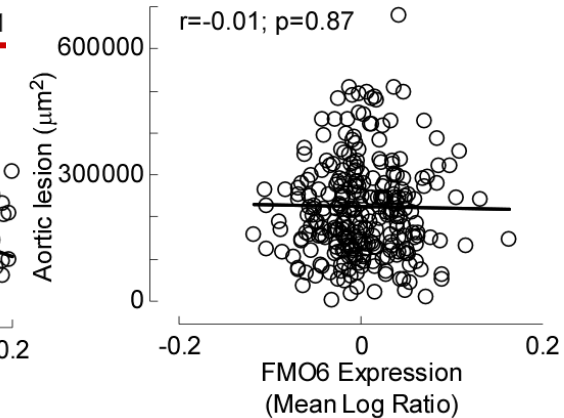
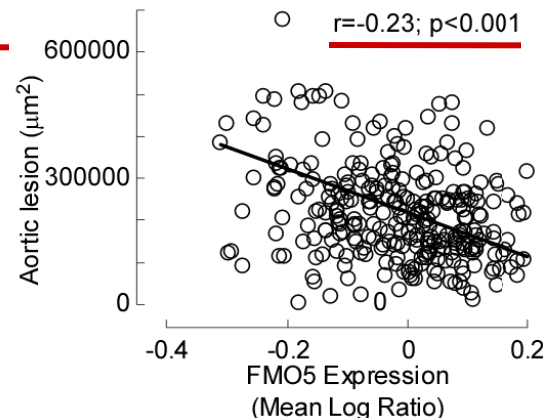
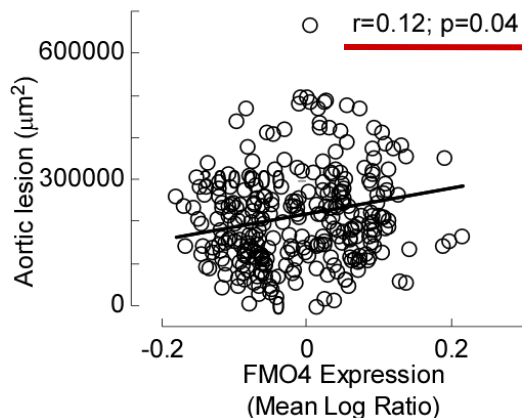
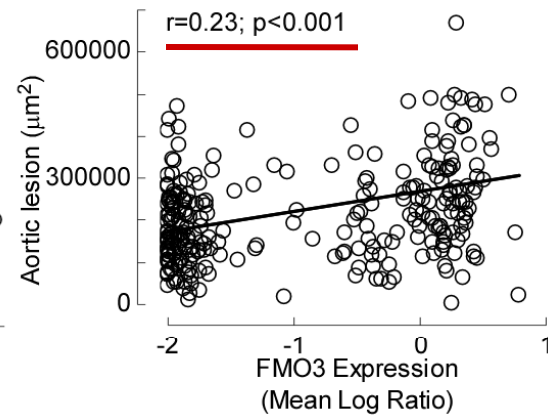
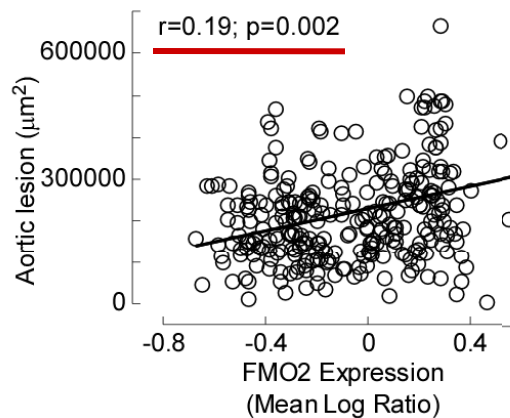
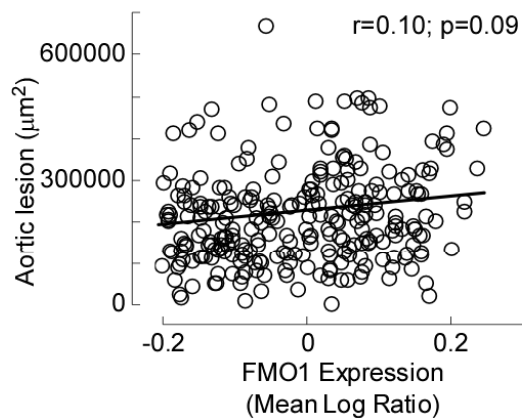


What of FMO3 ?



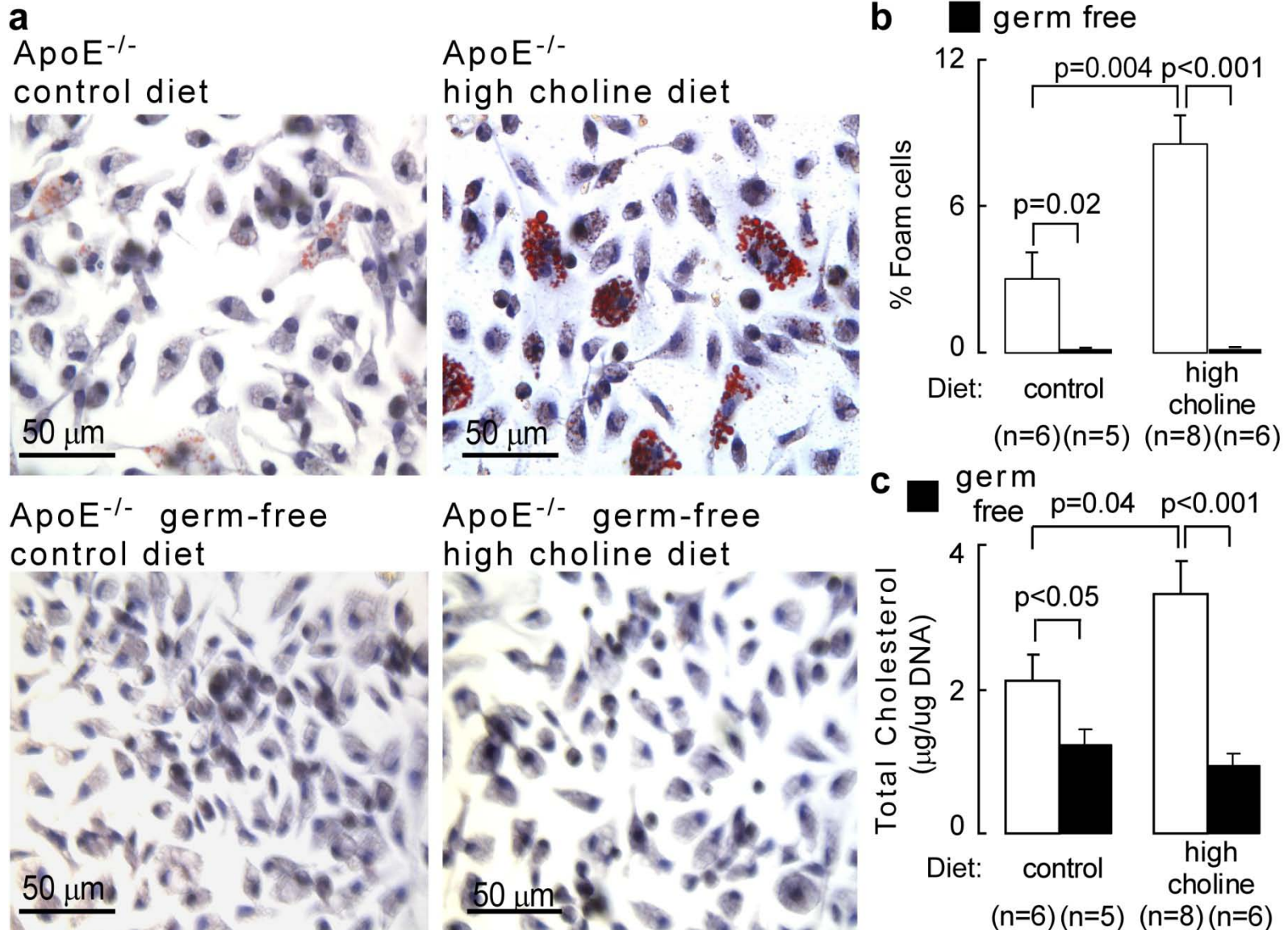
Integrative genetic studies in mice show FMO3 is linked to atherosclerosis susceptibility





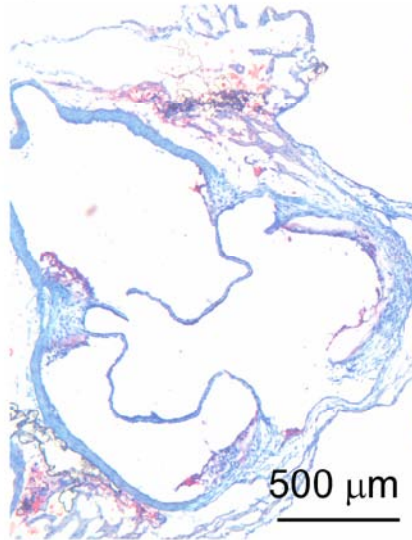
Hepatic expression of multiple FMO in the FMO gene cluster are correlated with aortic atherosclerotic lesion area (and TMANO and HDL)

Dietary choline and gut flora promote a pro-atherogenic macrophage phenotype *in vivo*

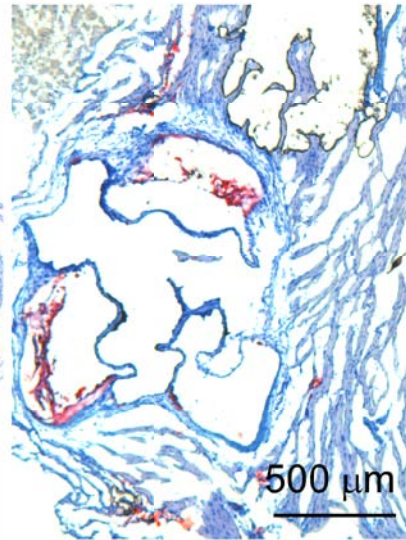


Suppression of gut flora inhibits dietary choline induced pro-atherogenic phenotype

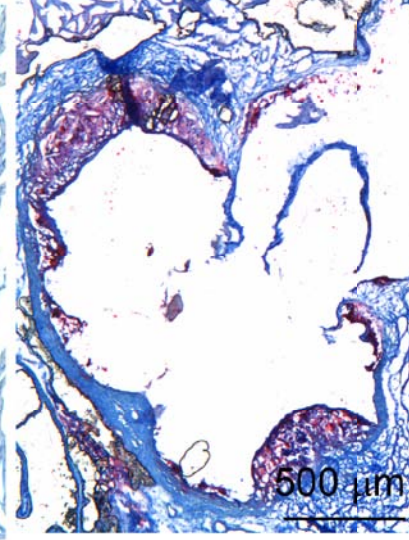
F, control



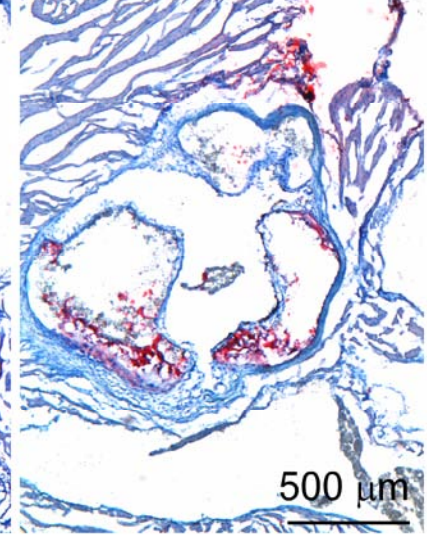
F, control/Abx



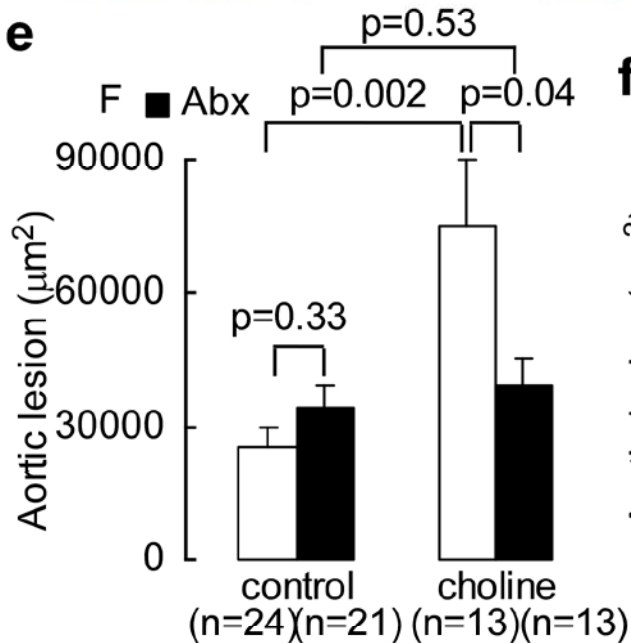
F, choline



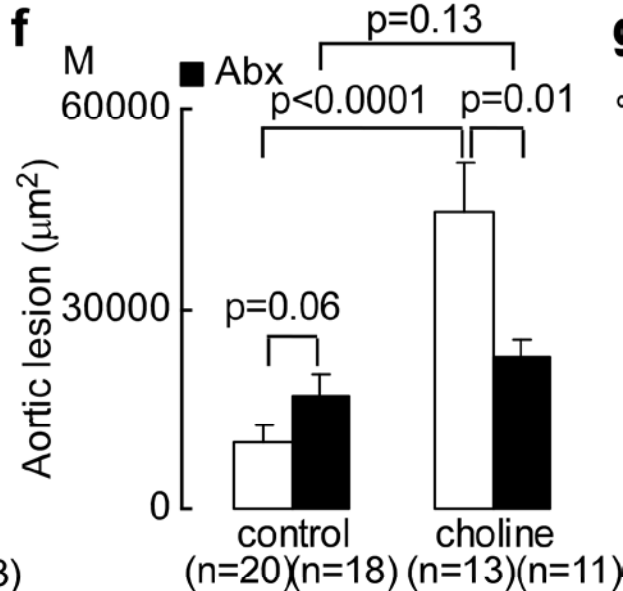
F, choline/Abx



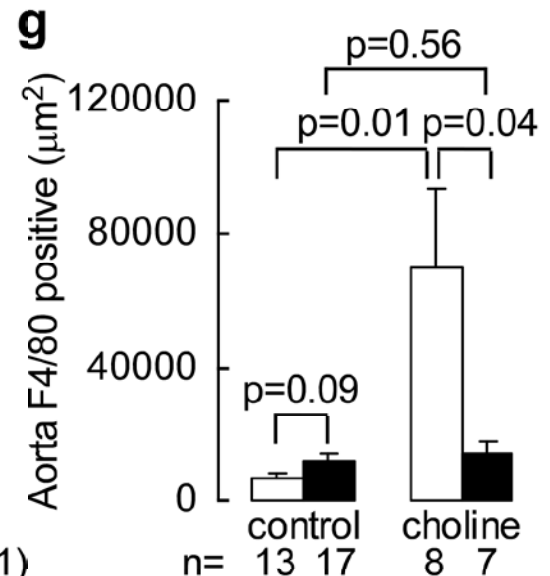
e



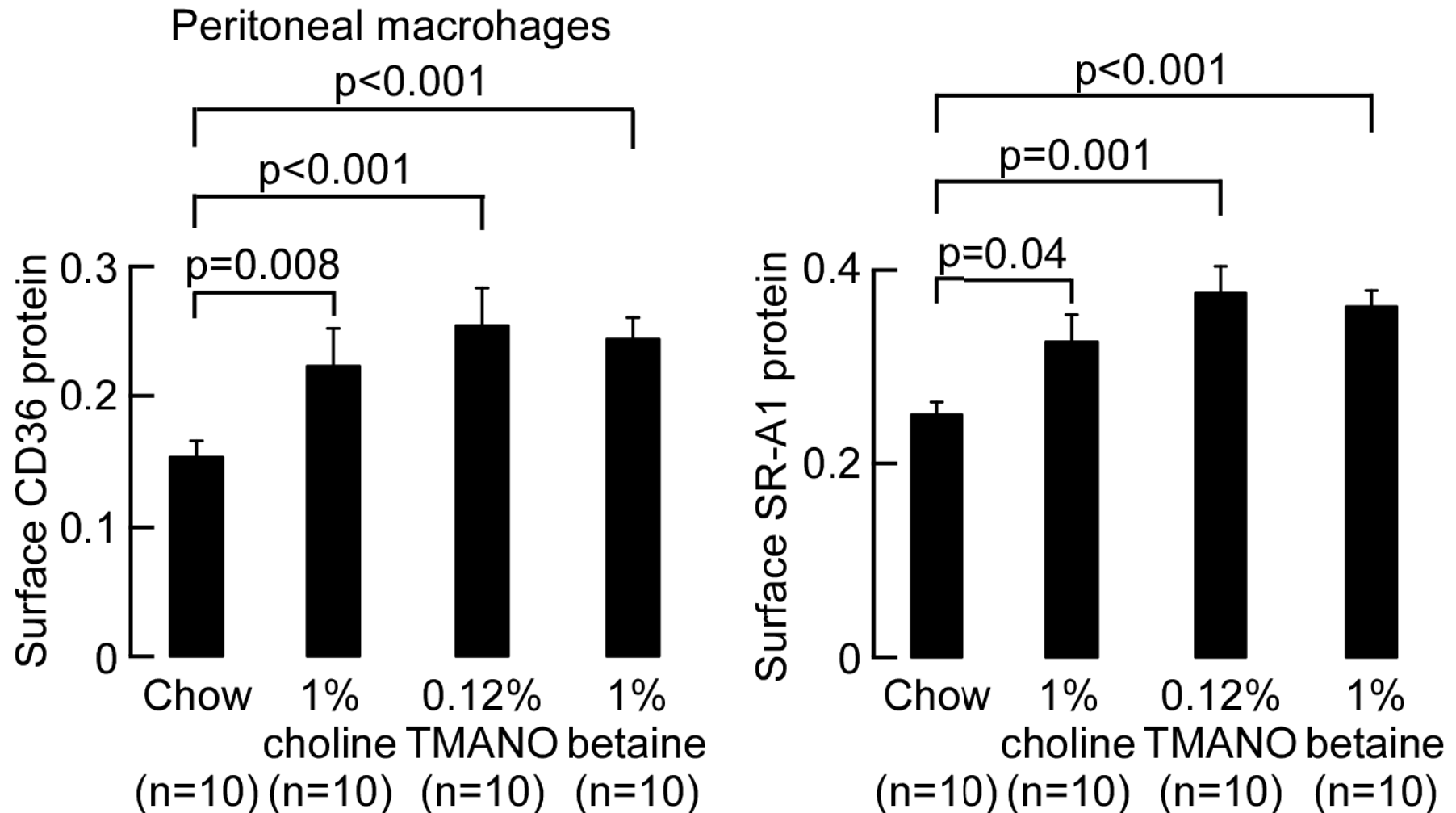
f



g



Effect of trimethylamine compound supplementation on macrophage scavenger receptor surface levels *in vivo*



A new (additional) nutritional basis of atherosclerotic heart disease.....suggests new diagnostic and therapeutic approaches for the detection/treatment of CVD

