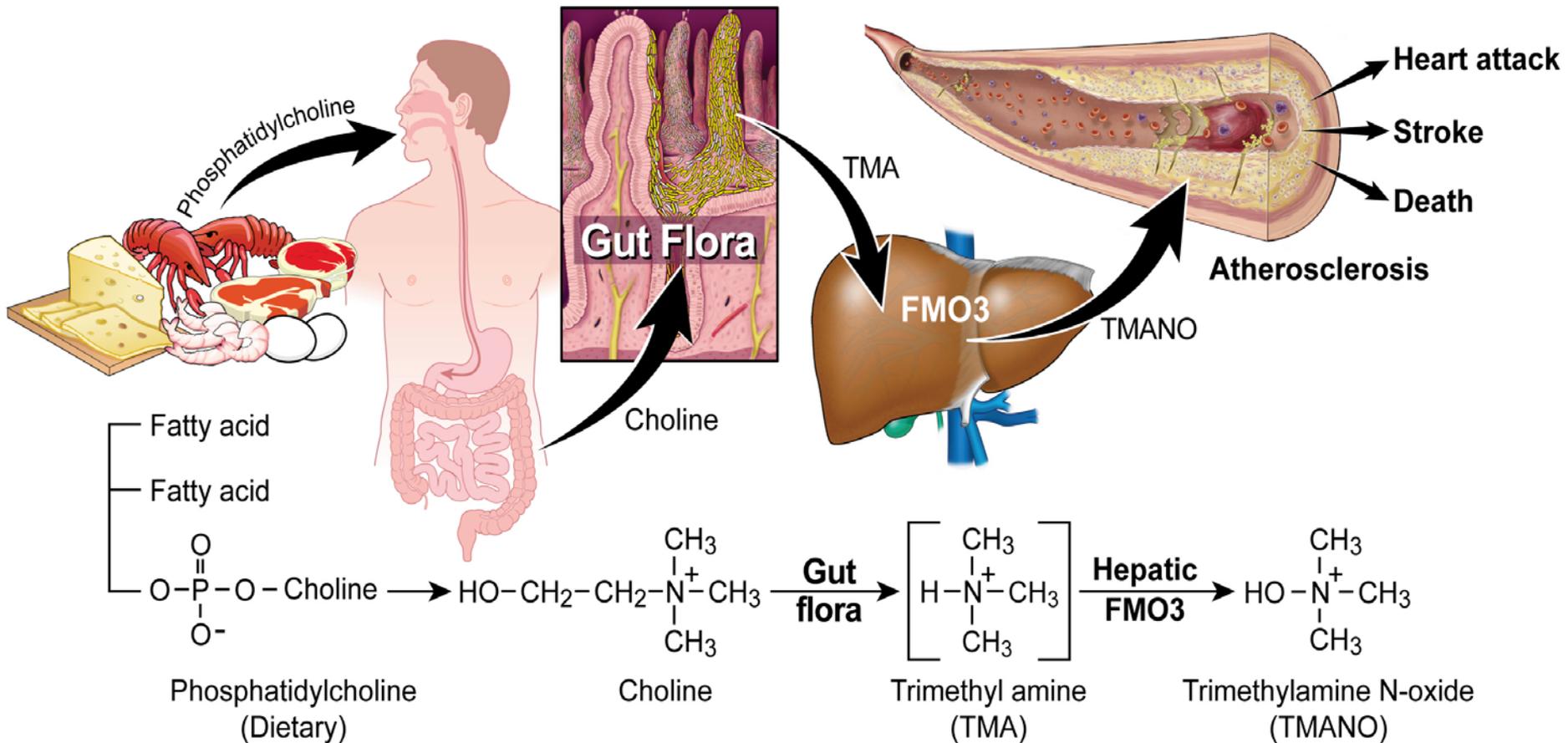


Gut flora  
dependent  
metabolism of  
phosphatidylcholine  
and the  
pathogenesis of  
cardiovascular  
disease

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

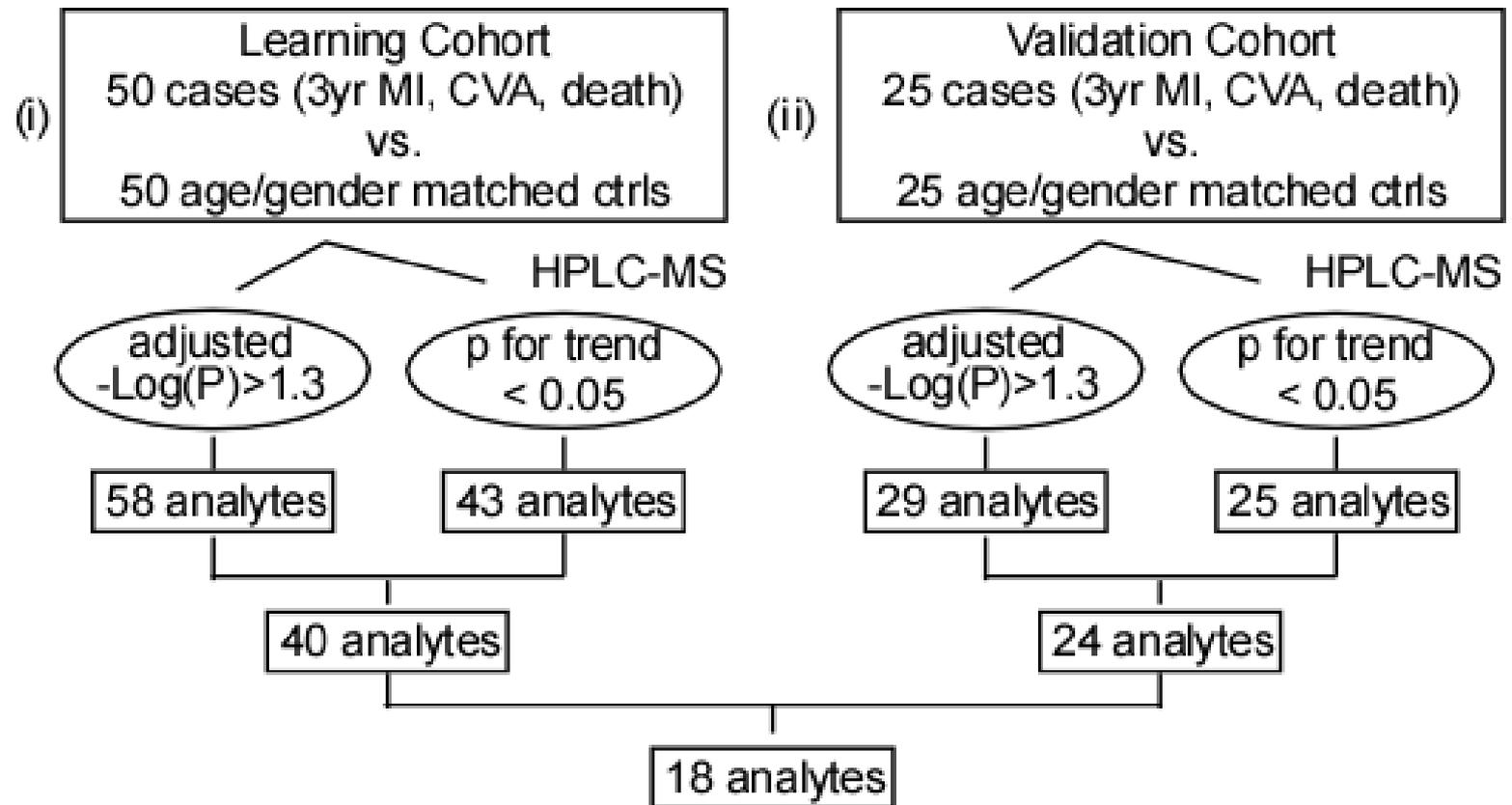
## Phase 2: Clinical validation

Demonstration of clinical utility

## Phase 3: Mechanistic studies

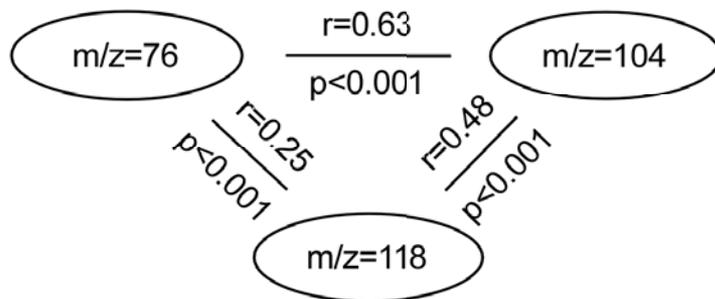
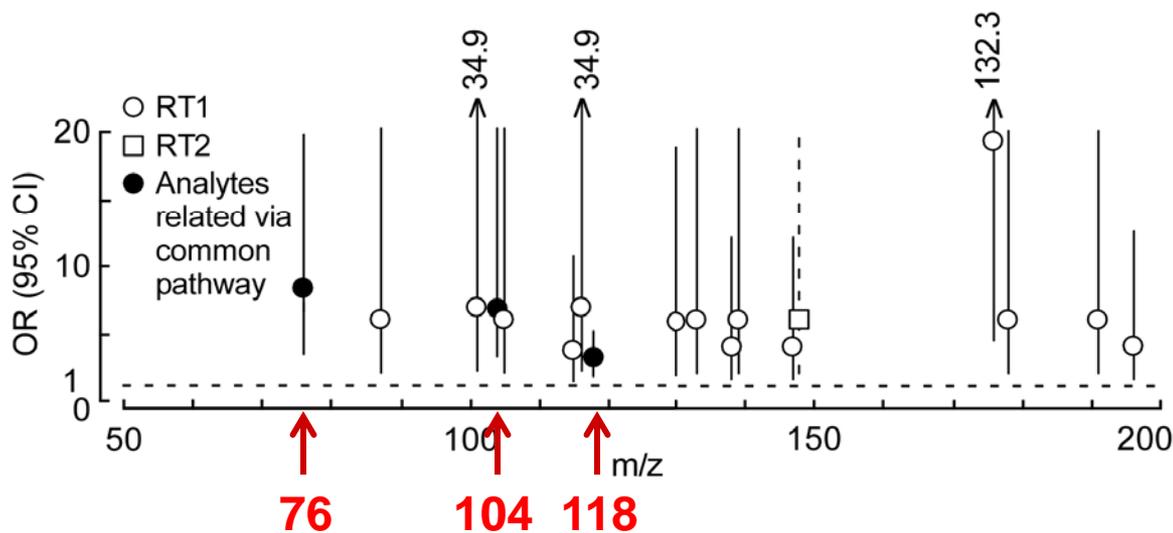
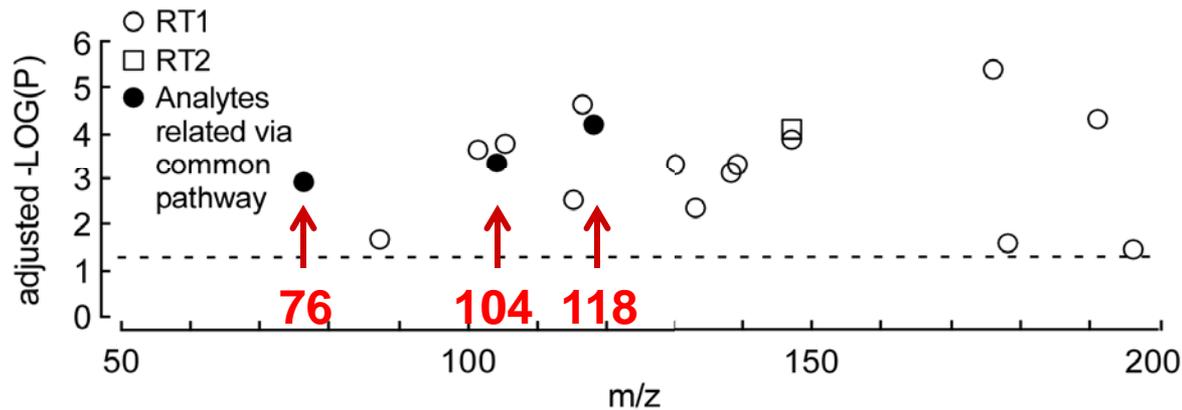
Demonstration of causality for novel pathway

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



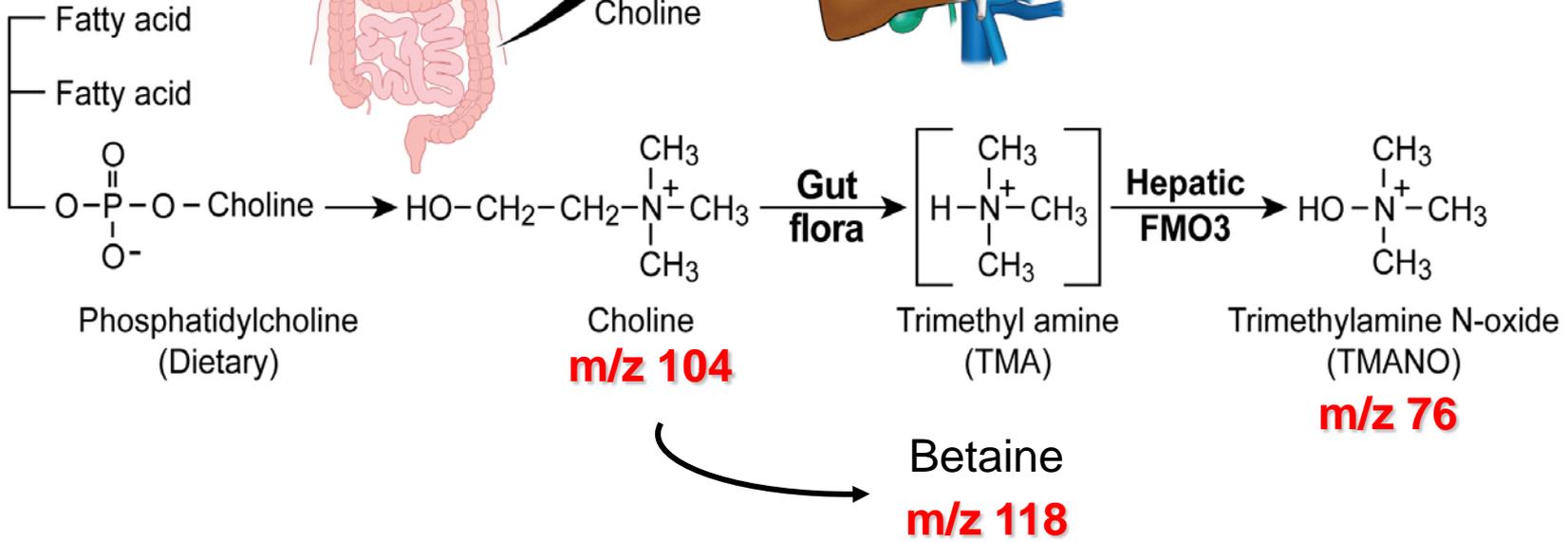
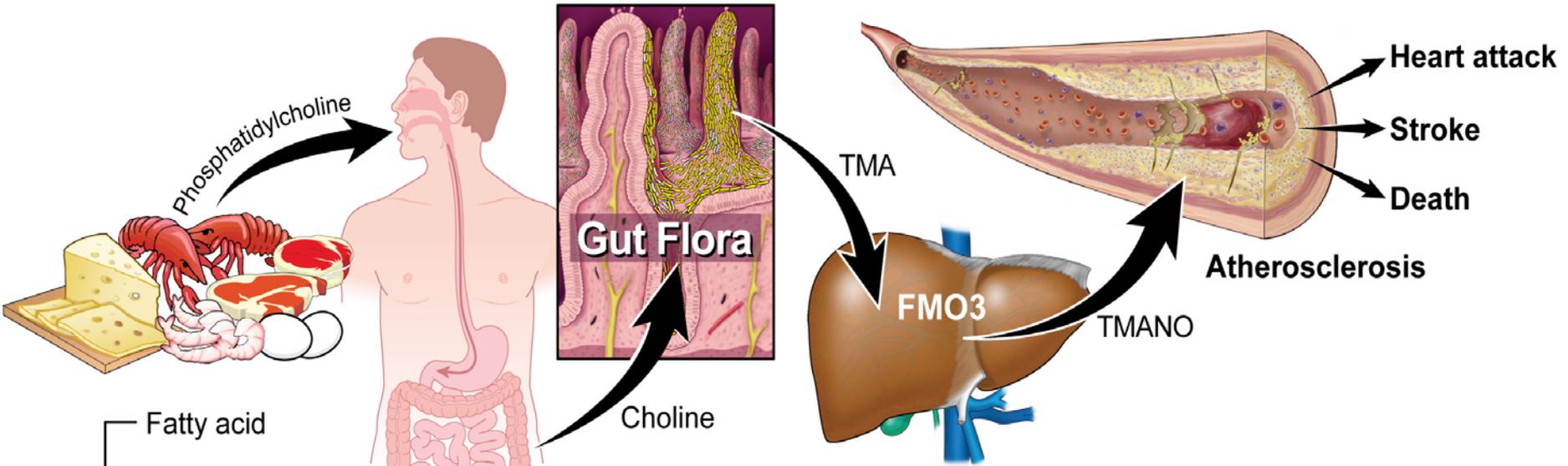
(ii) Structural identification of analytes

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



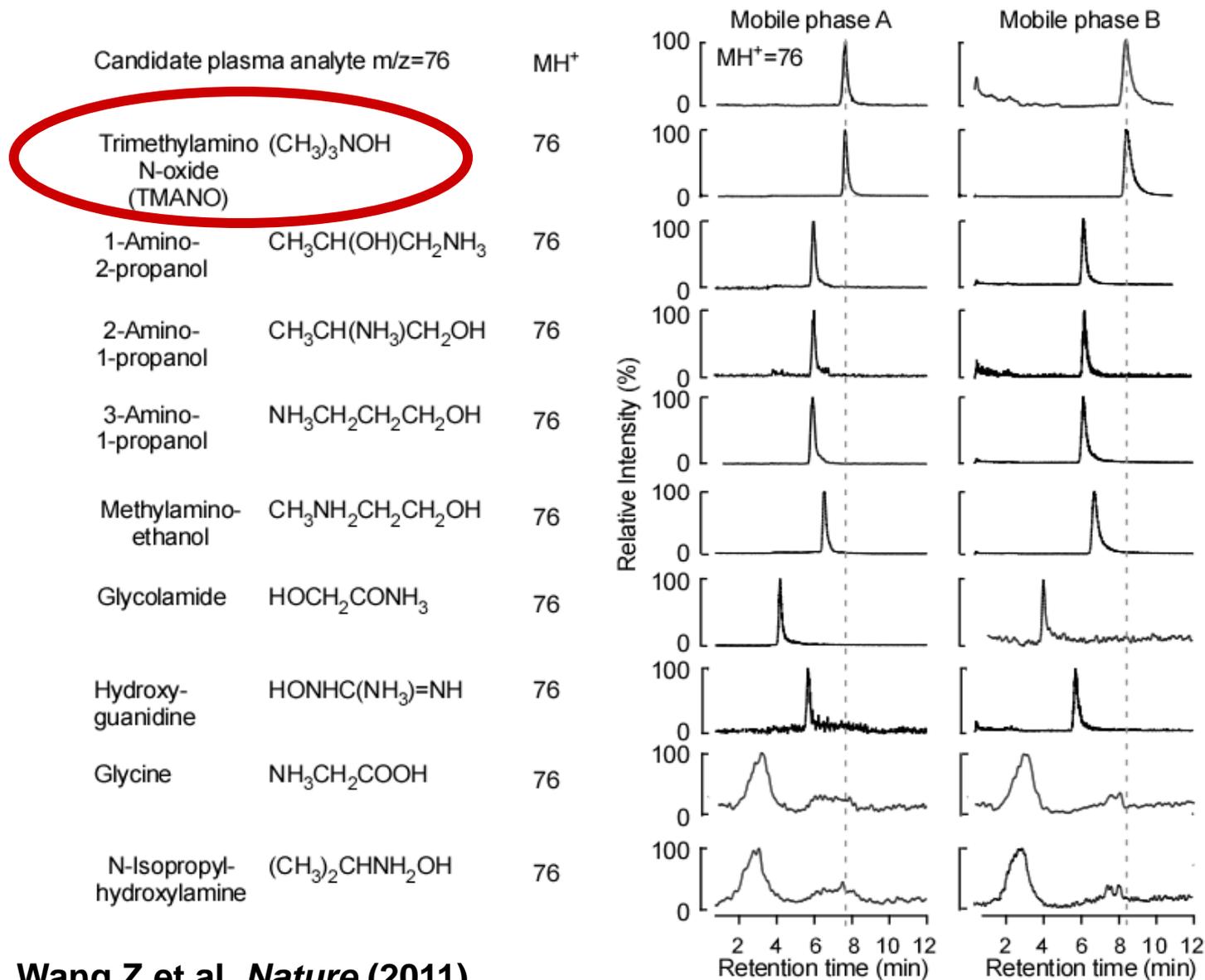
Plasma analytes with m/z 76, 104, and 118 are associated with CVD, show a dose-response relationship with MACE and are correlated, suggesting participation in a common pathway

# 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 $m/z=76$

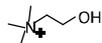


MH<sup>+</sup>

MS1, m/z=104

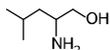
Plasma analyte m/z=104

choline



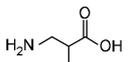
104

2-amino-3-methyl-1-butanol



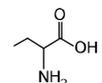
104

3-aminoisobutyric acid



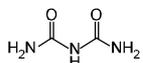
104

2-aminobutyric acid



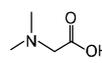
104

biuret



104

N,N-dimethyl glycine



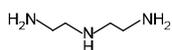
104

benzonitrile



104

diethylenetriamine



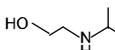
104

ethyl-N-hydroxyl acetimidate



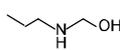
104

2-isopropyl-aminoethanol



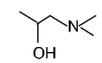
104

2-propyl-aminoethanol

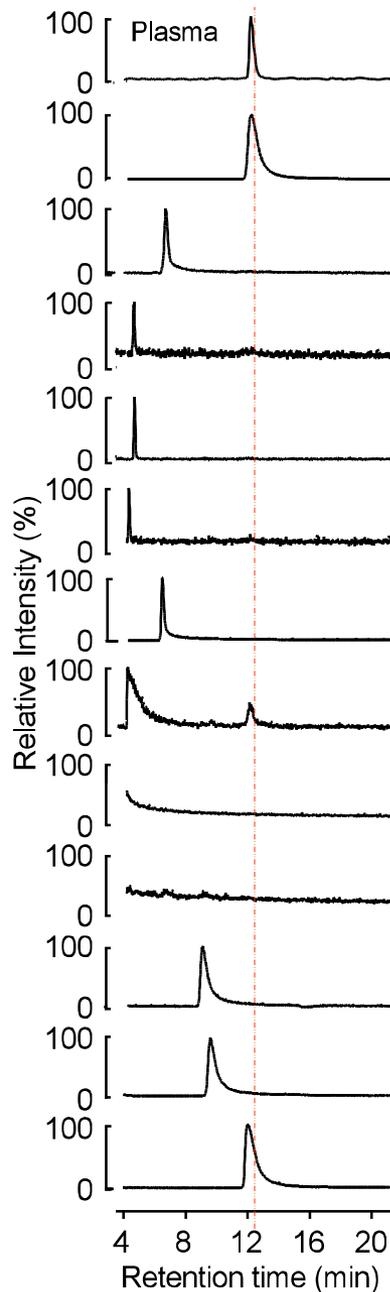


104

1-dimethyl-amino-2-propanol



104



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

**Identity as Choline was confirmed by:**

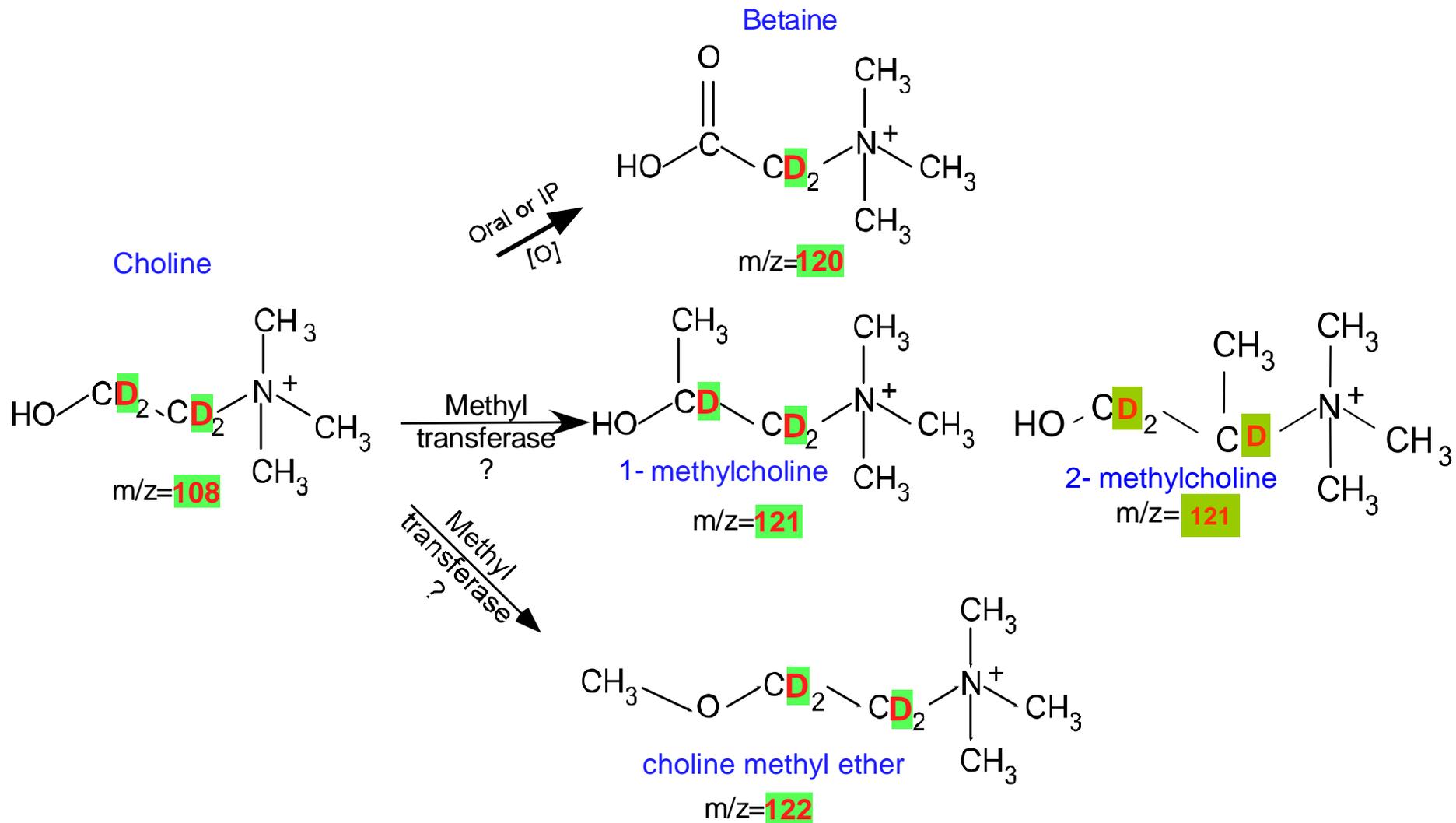
**LC-MS<sup>n</sup>**

**GC/MS/MS**

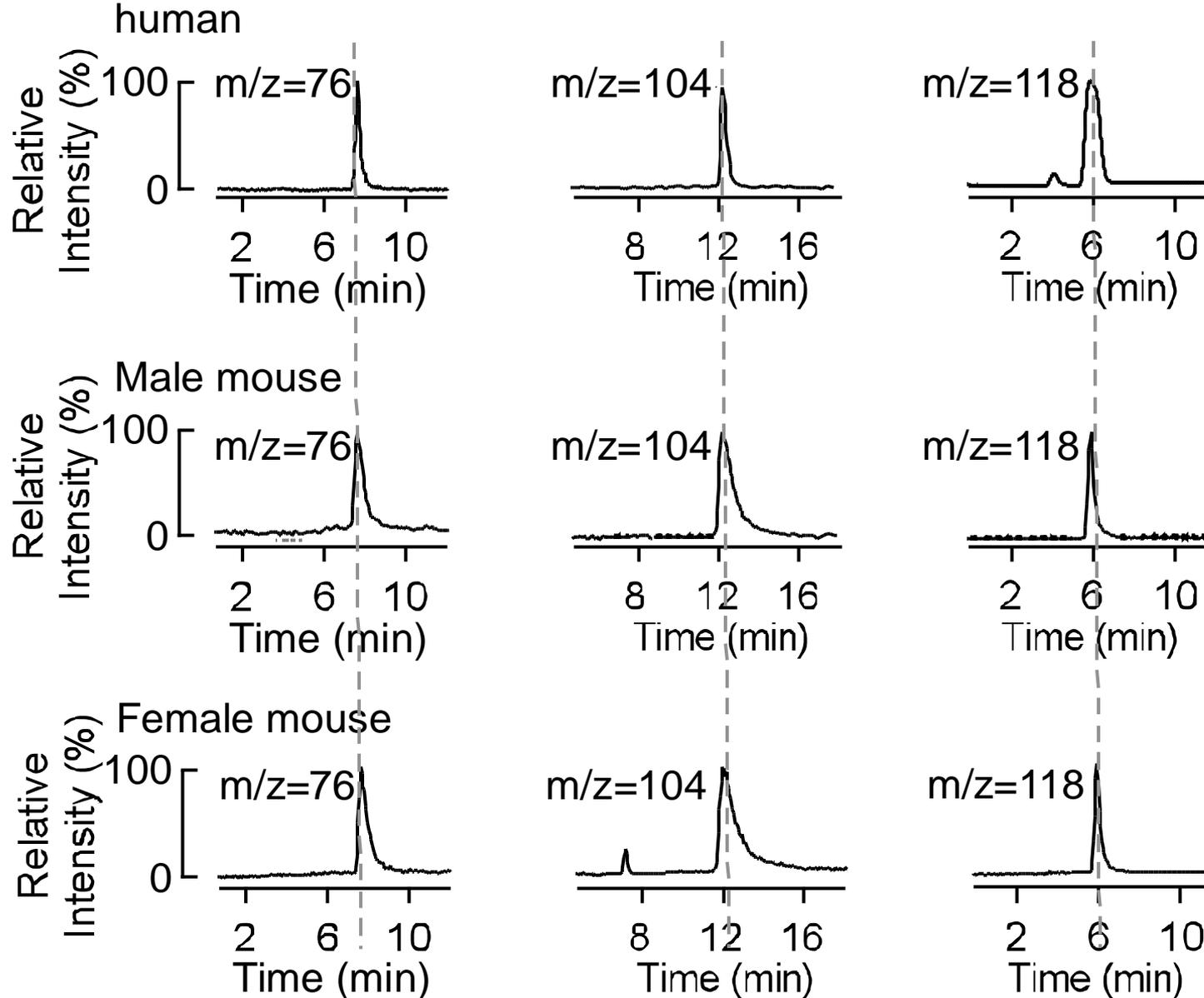
**<sup>1</sup>H, <sup>13</sup>C, and <sup>15</sup>N NMR**

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

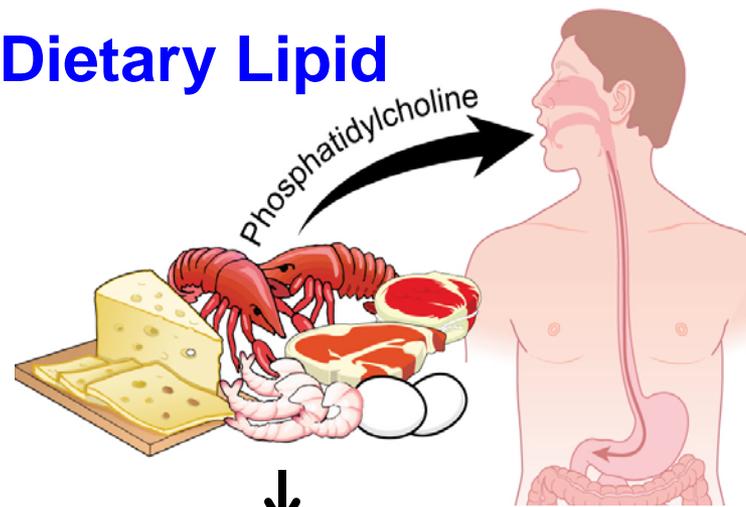
# Strategy to determine the analyte at m/z=118 by choline deuterated isotopologue feeding study



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

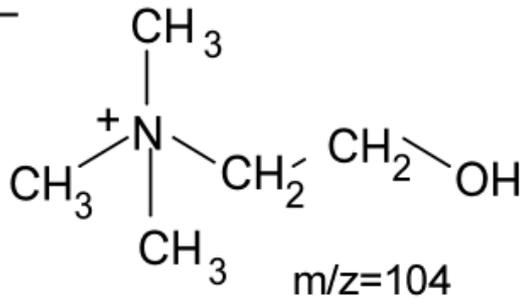


# Dietary Lipid



What is the role of gut flora?

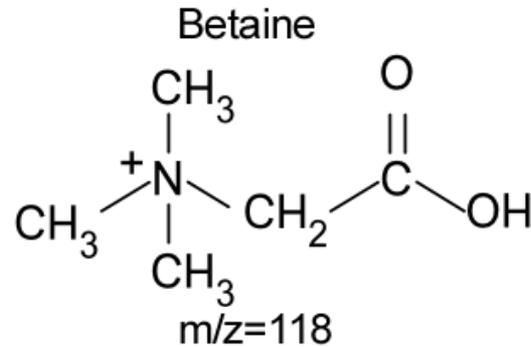
Choline



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

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

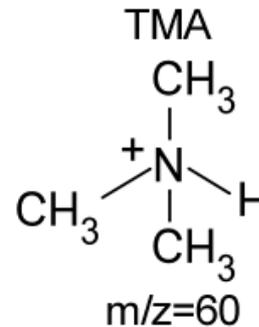
Oral or i.p.  
[O]



d4 metabolite m/z=120

d9 metabolite m/z=127

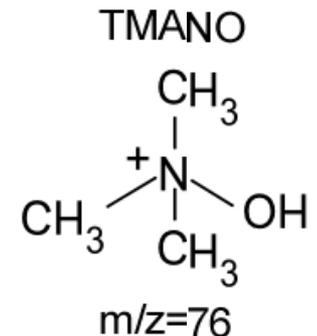
Oral only  
Gut Flora



d4 metabolite m/z=60

d9 metabolite m/z=69

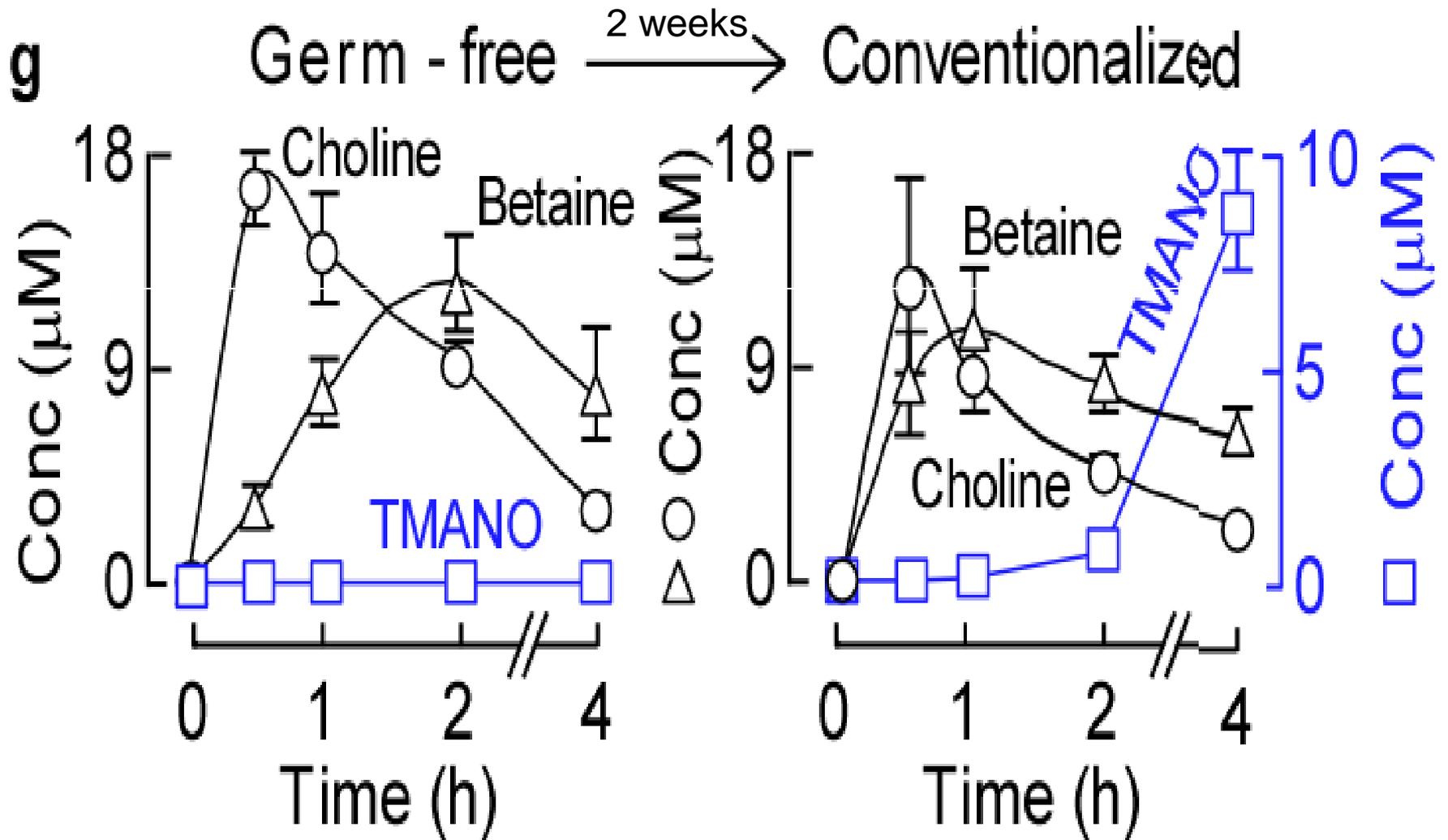
presumed  
FMO3  
[O]



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

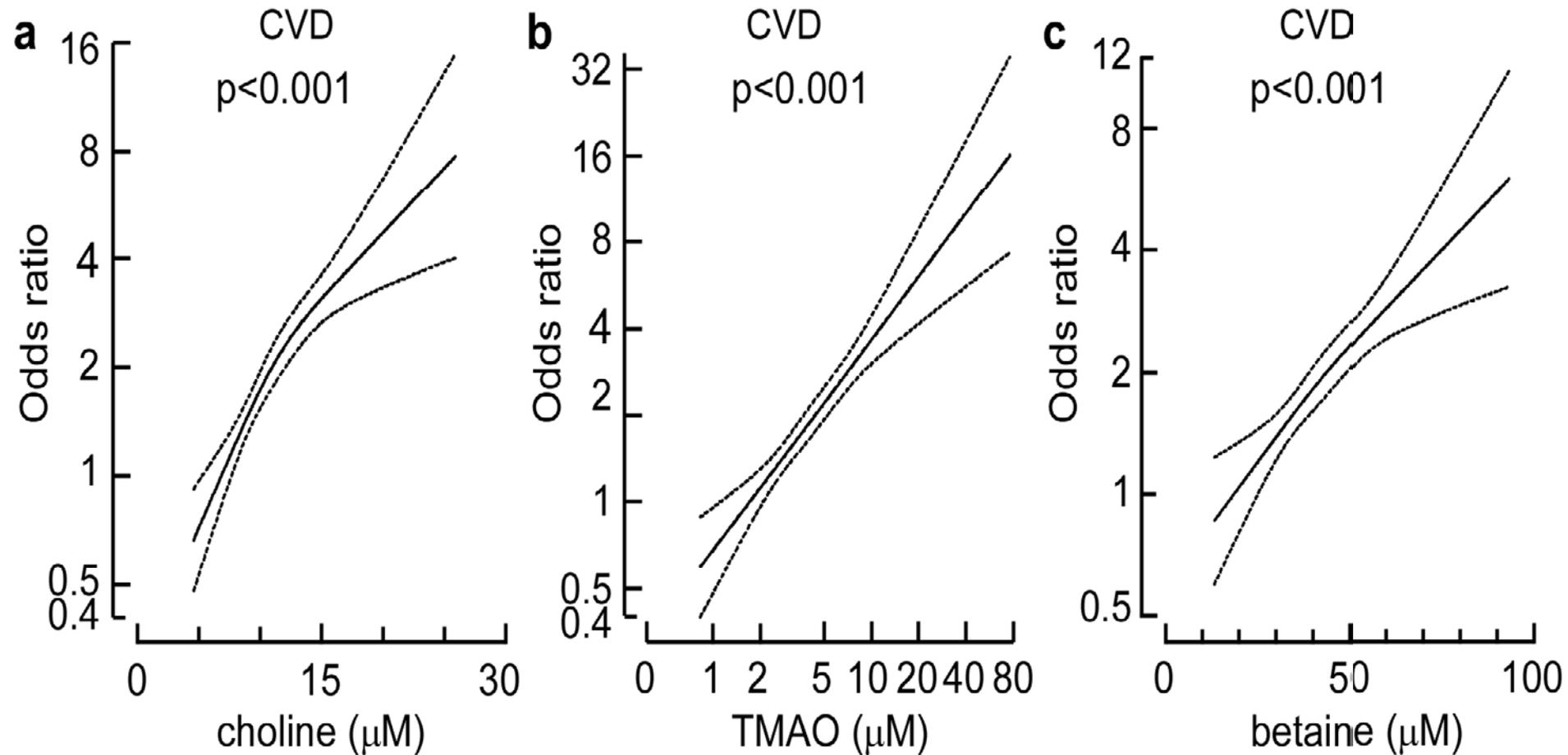
# Prospective Cohort: N=1865 Sequential Cardiology Patients

## 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), $\mu$ M	3.3 (2.2-5.0)	4.4 (2.8-7.4)	<0.001
Choline, median (IQR), $\mu$ M	10.9 (8.6-13.5)	12.3 (9.8-15.4)	<0.001
Betaine, median (IQR), $\mu$ 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

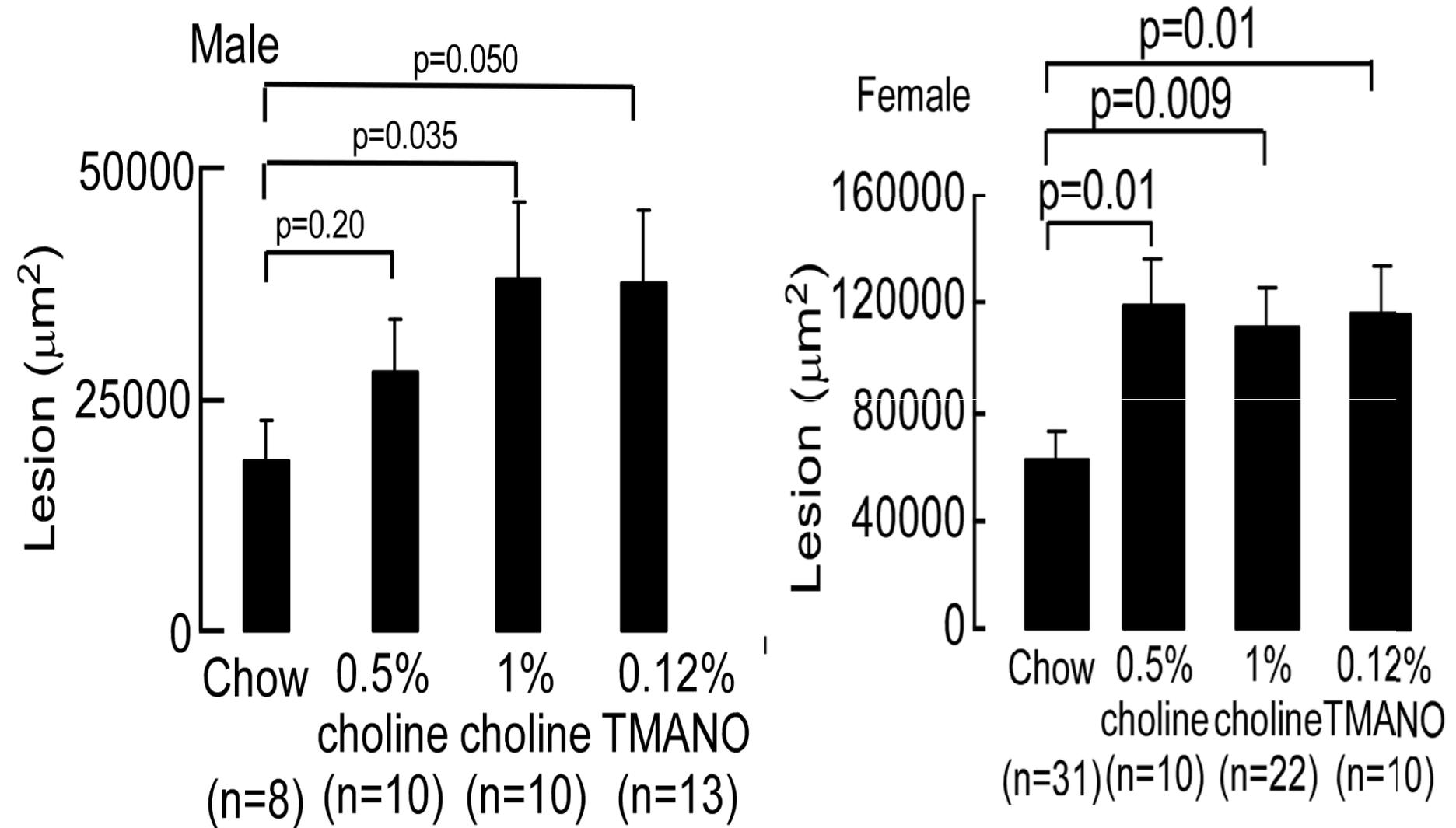
Demonstration of clinical utility

## Phase 3: Mechanistic studies

Demonstration of causality for a  
novel pathway

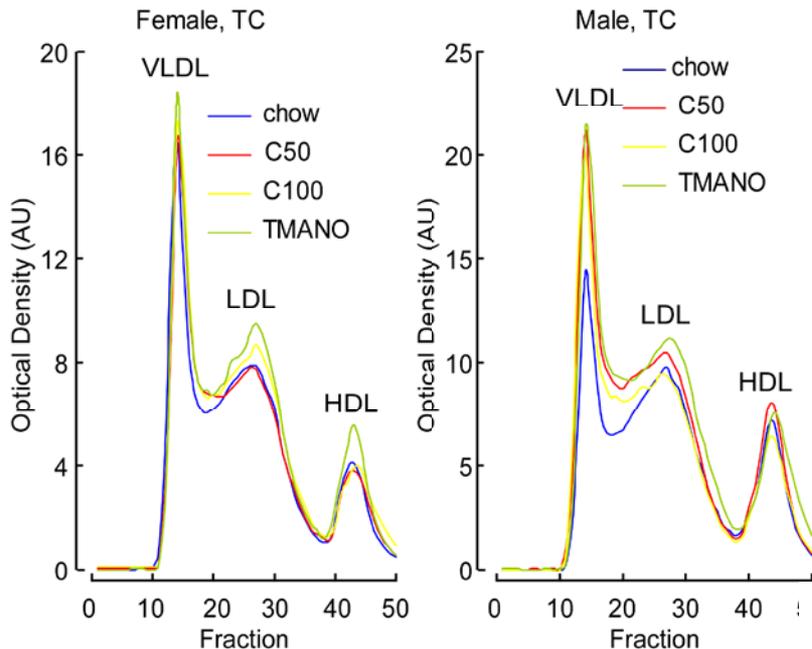
# Dietary supplementation with choline or TMANO promotes atherosclerosis ApoE<sup>-/-</sup> mice

Normal chow diet + indicated supplement weeks 4-20

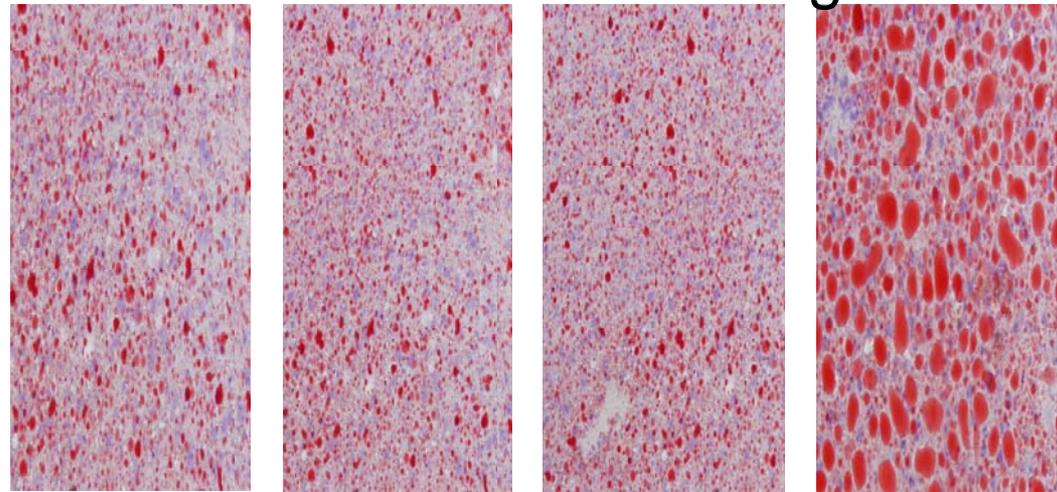


# Dietary supplementation with choline or TMAO promotes atherosclerosis without dyslipidemia or hepatic steatosis

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)

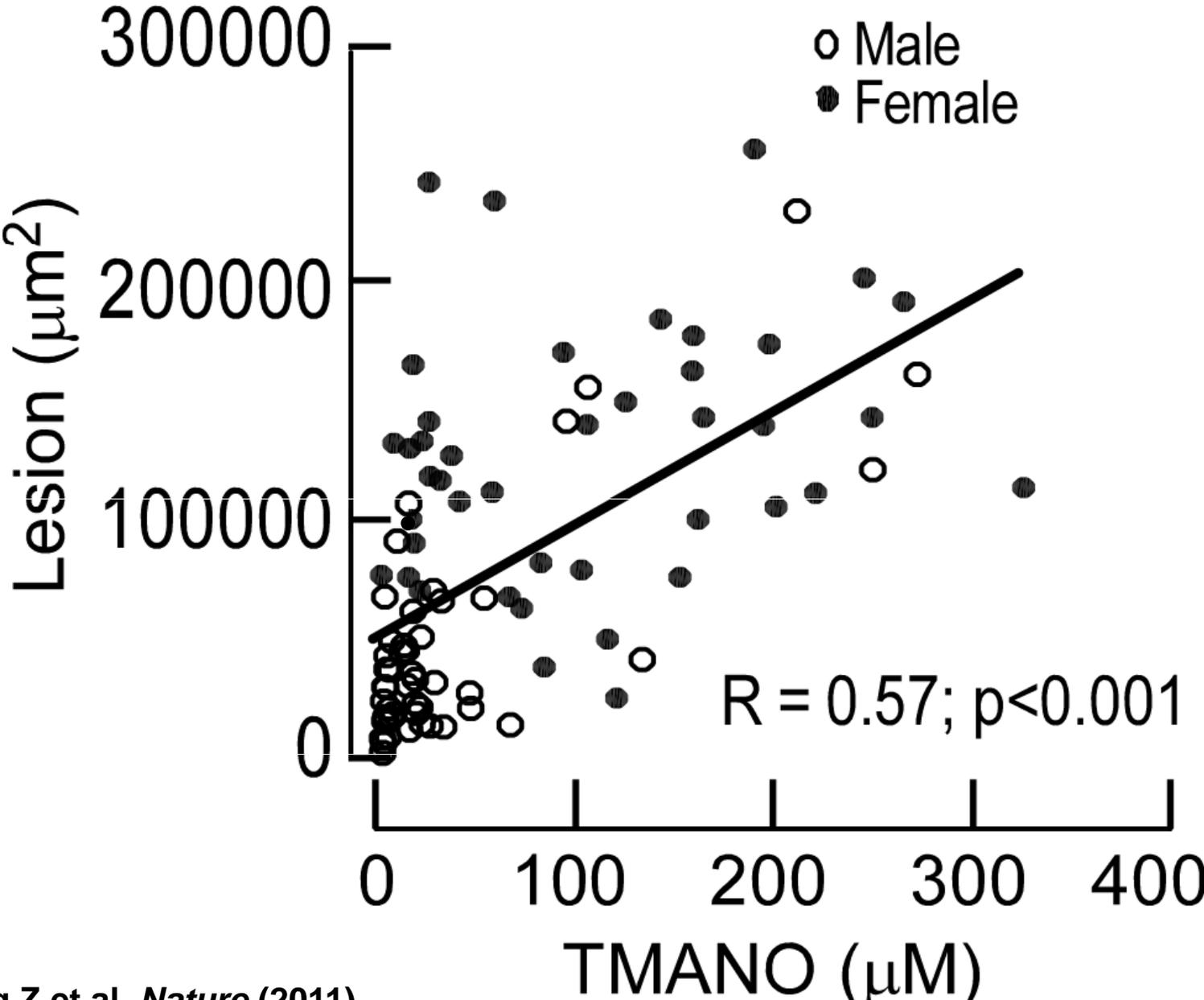


## Liver oil-red-o staining



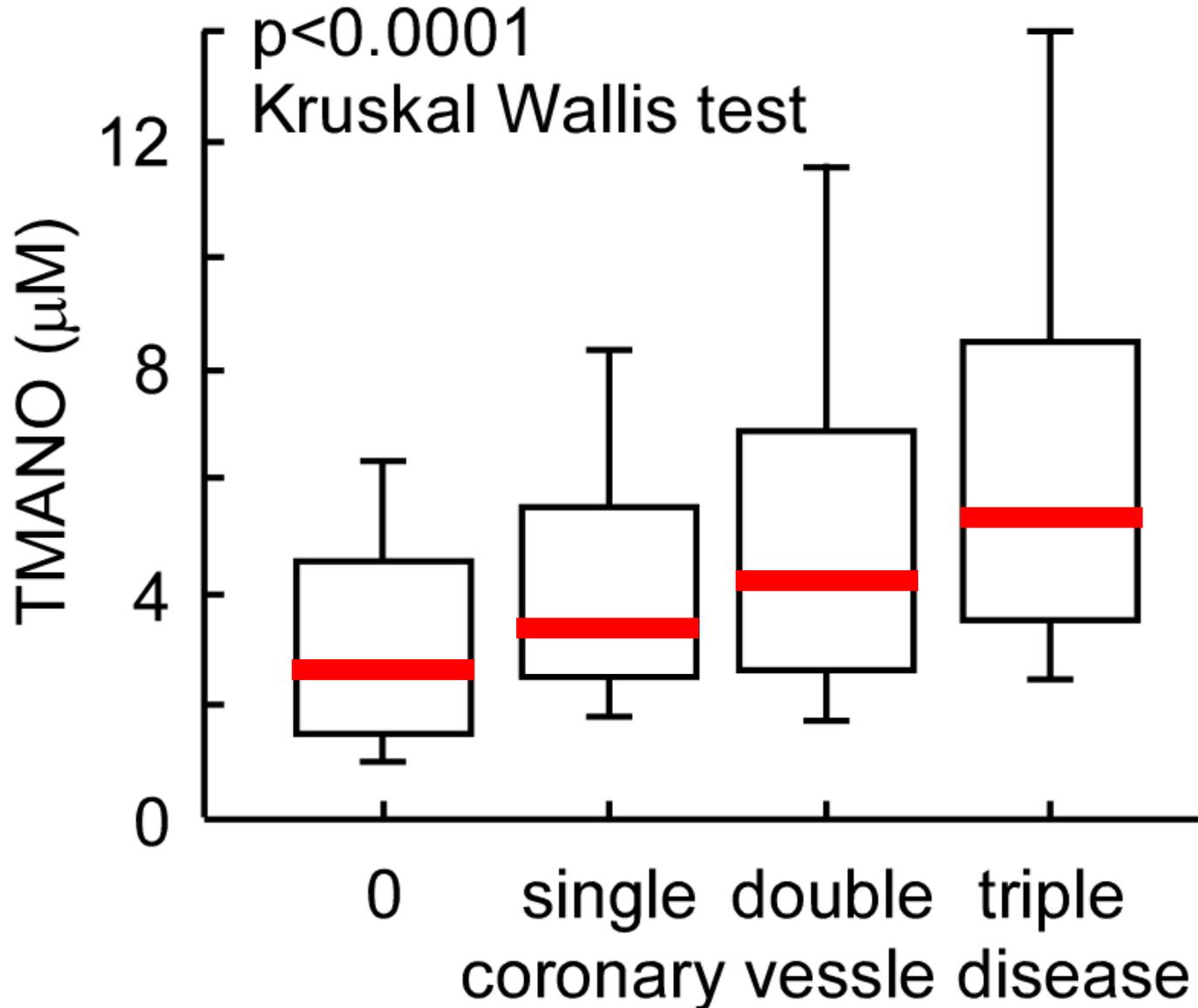
**Chow    1% Choline    0.12% TMAO    MCD**

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

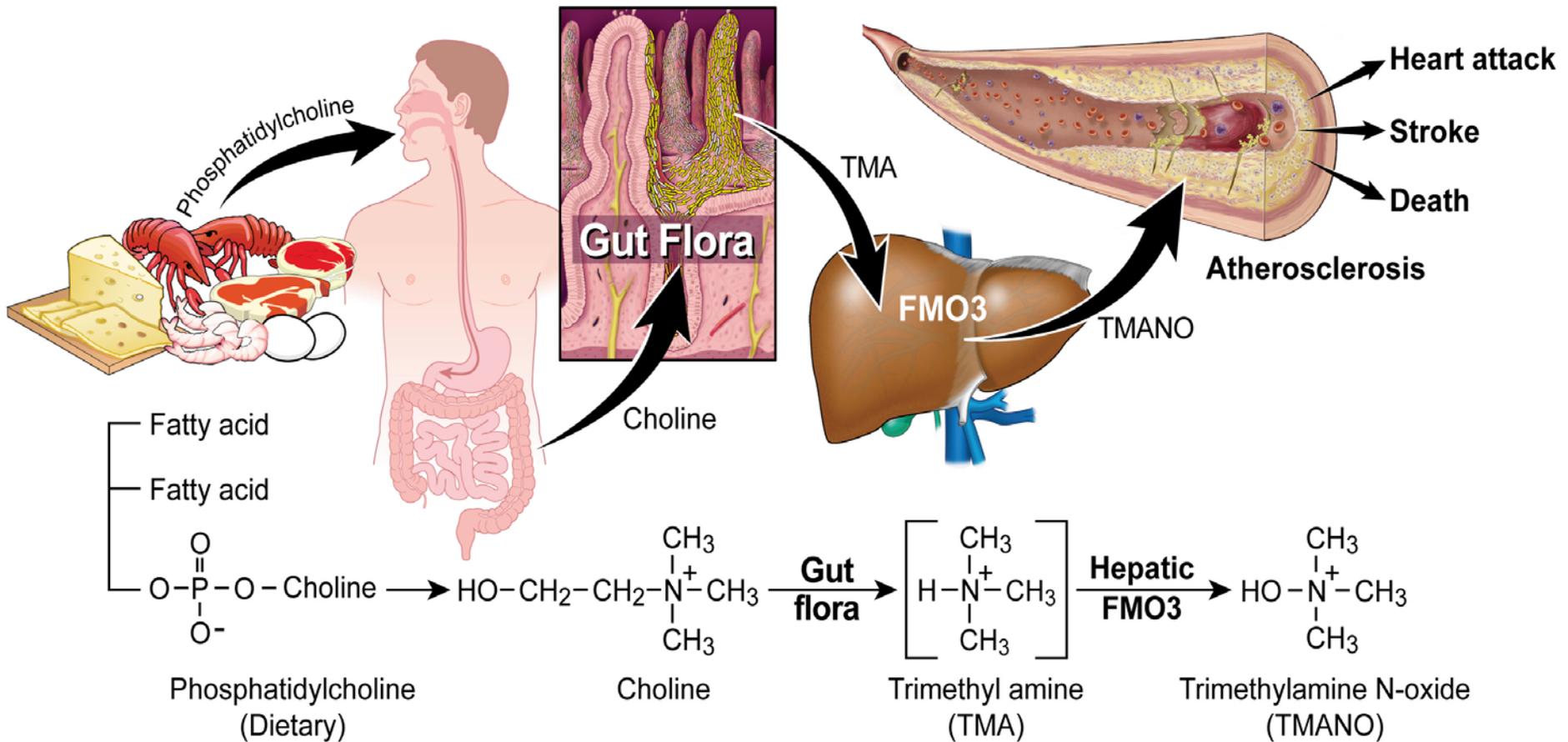


Wang Z et al, *Nature* (2011)

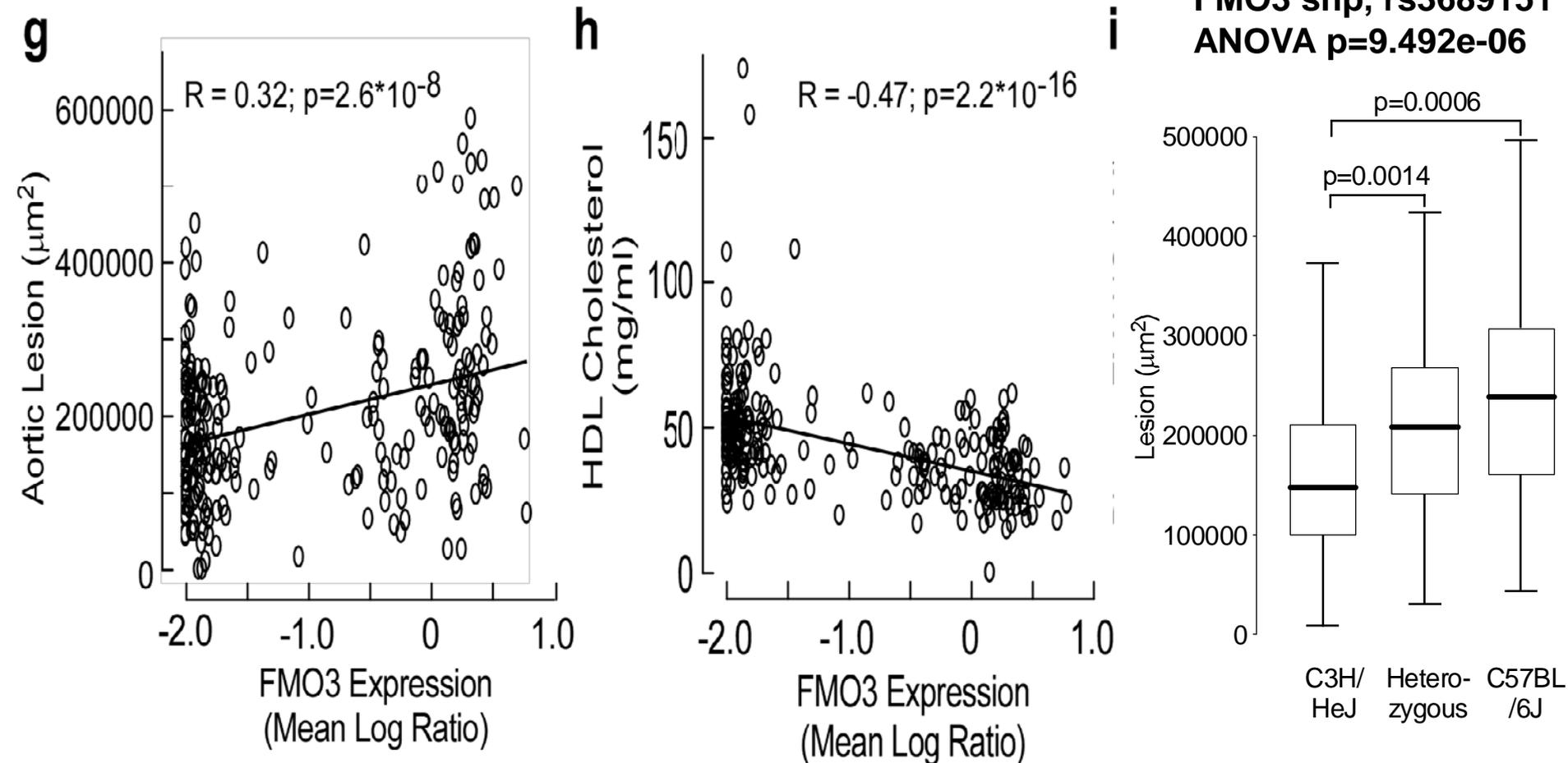
Plasma TMANO levels are correlated with angiographic coronary artery disease severity in subjects (N=1020)

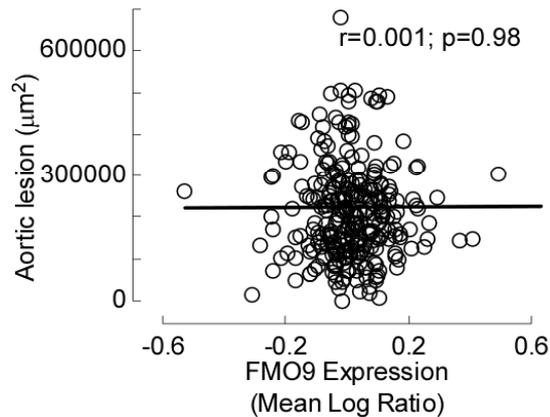
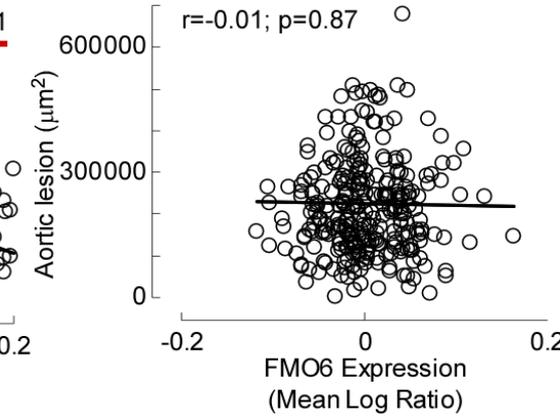
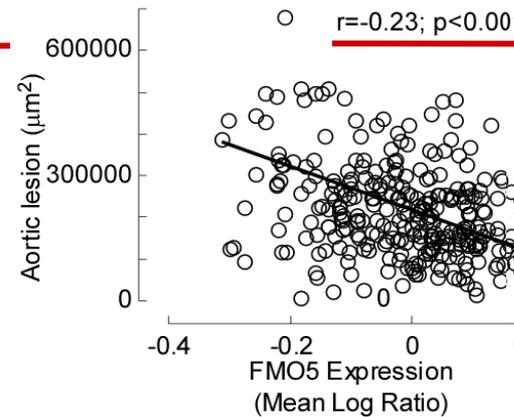
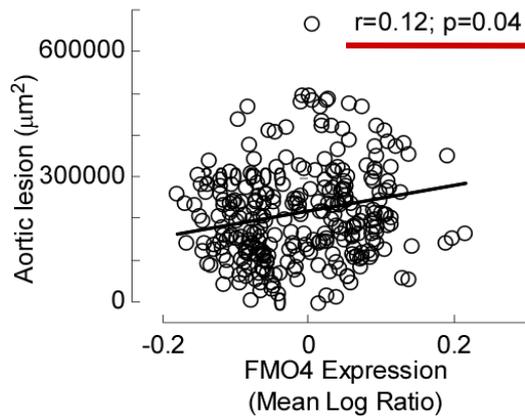
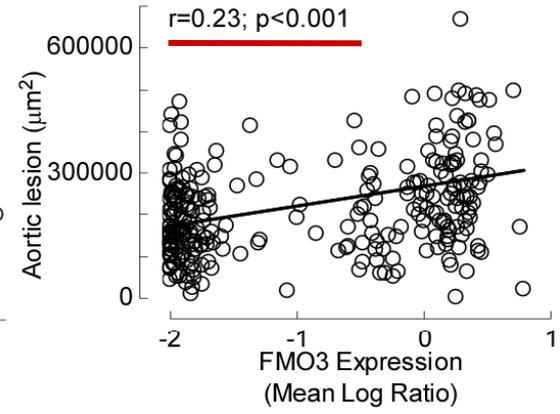
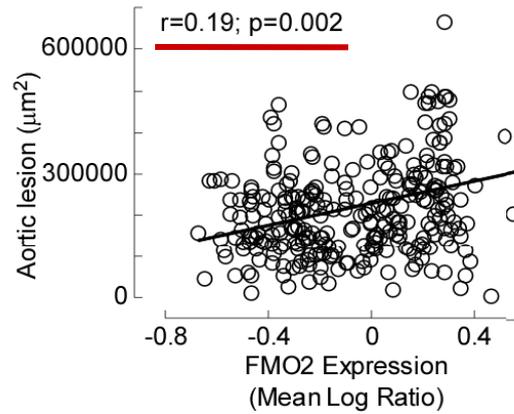
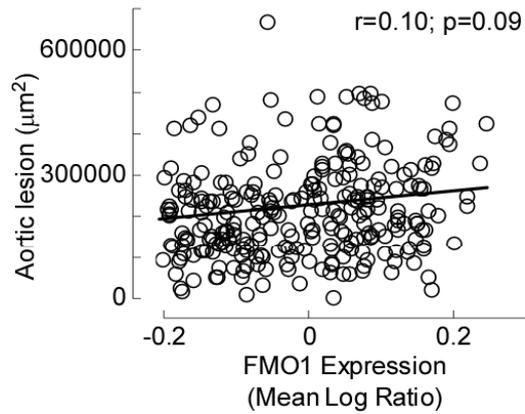


# What of FMO3 ?



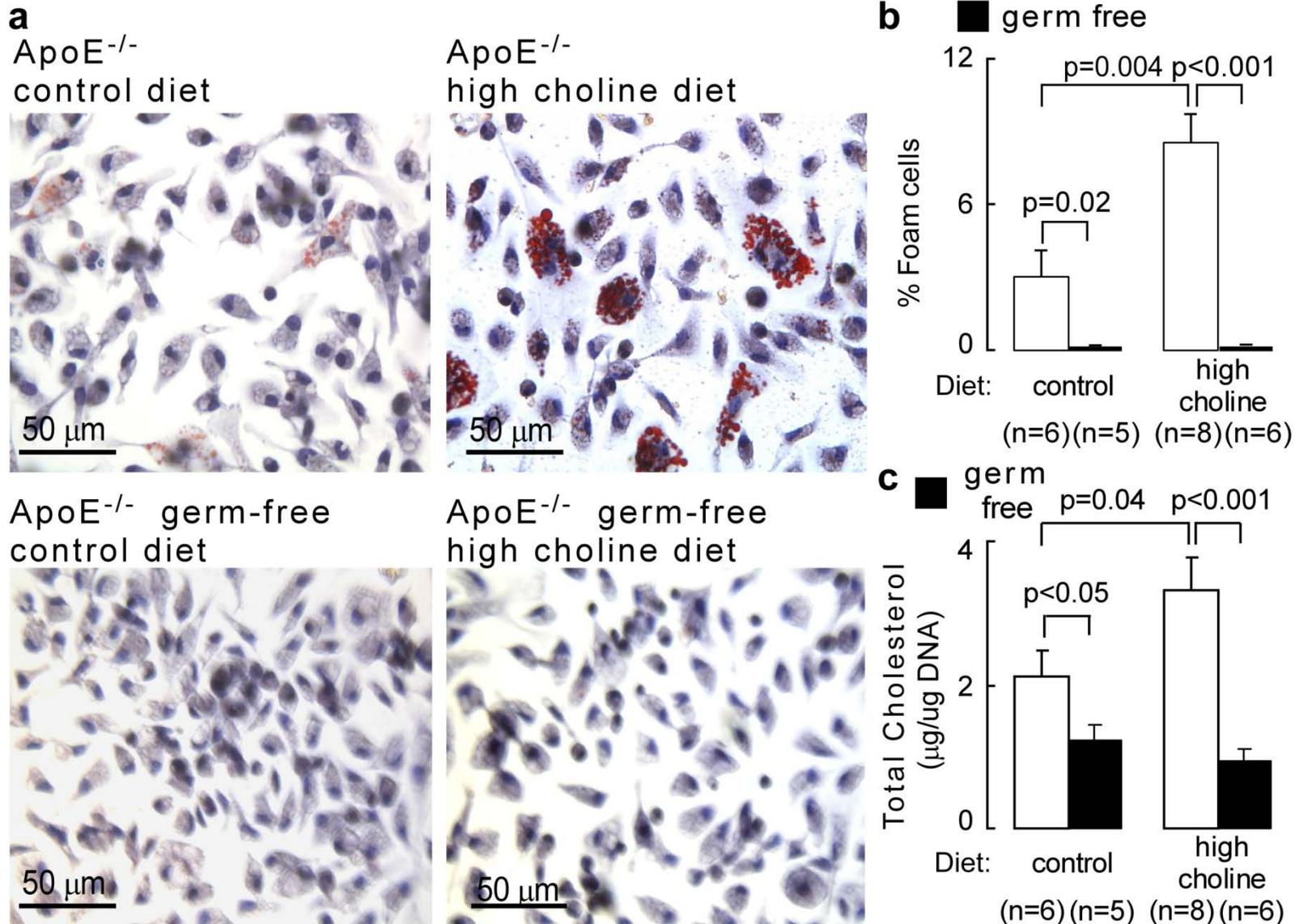
# Integrative genetic studies in mice show FMO3 is linked to atherosclerosis susceptibility



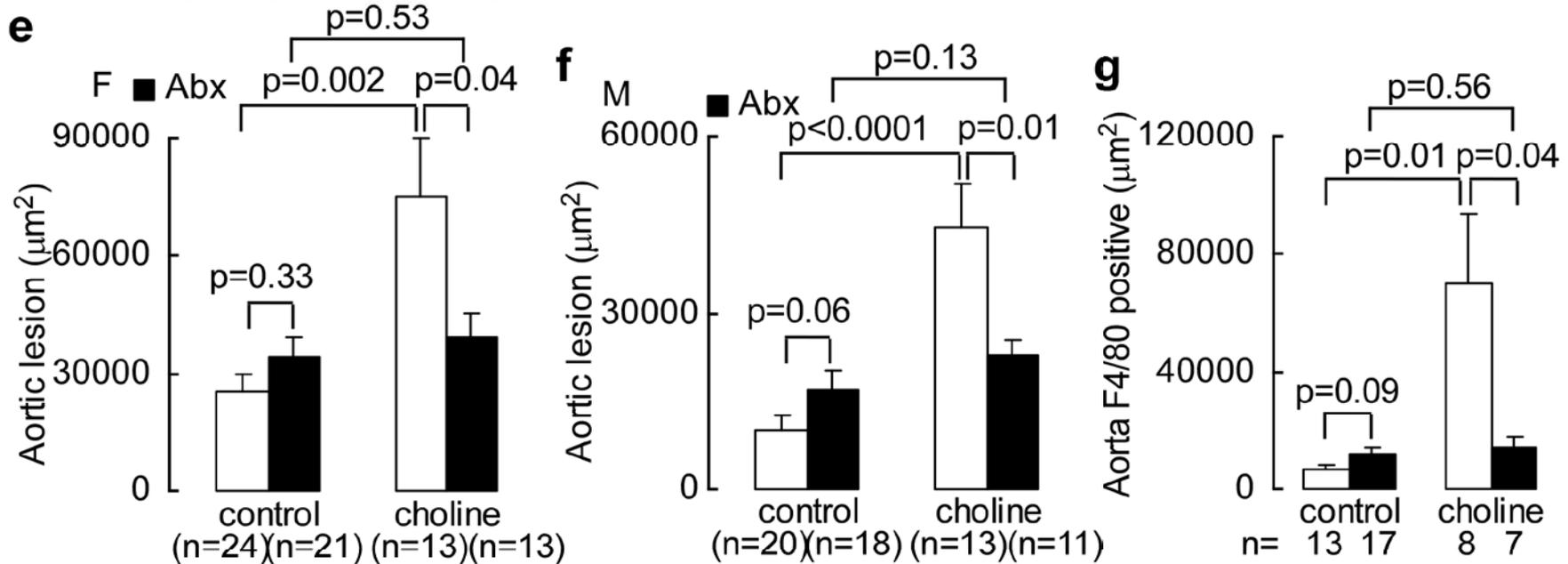
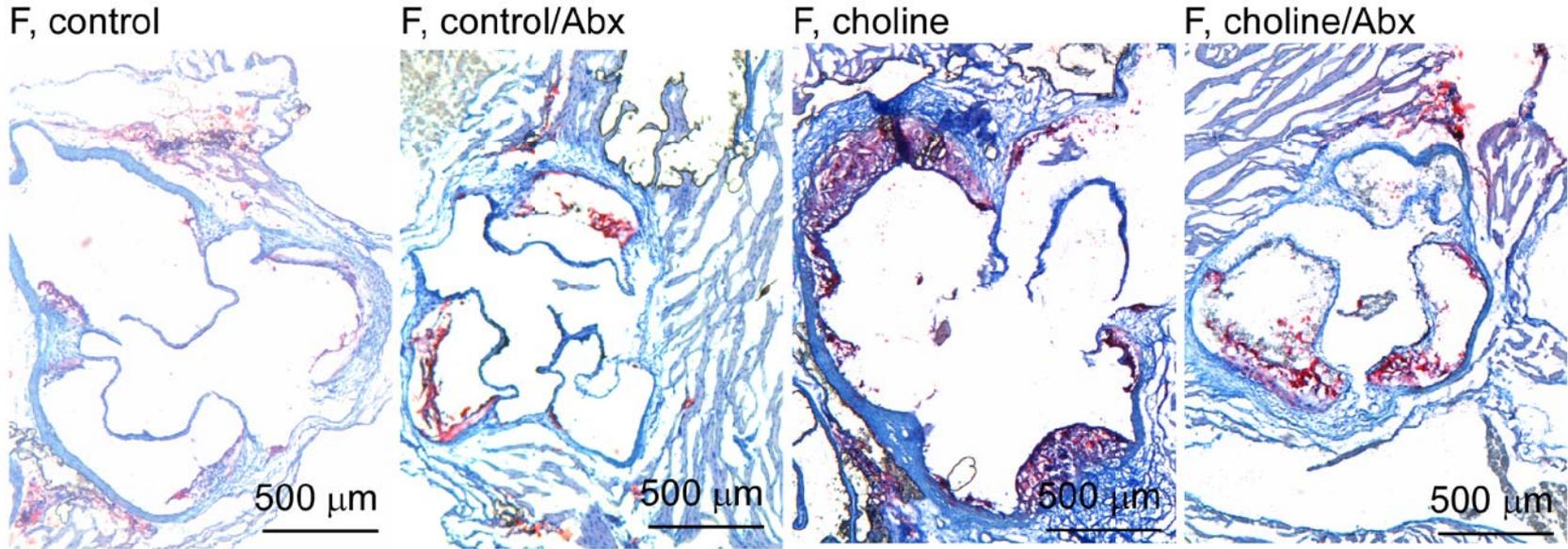


**Hepatic expression of multiple FMO in the FMO gene cluster are correlated with aortic atherosclerotic lesion area (and TMANO and HDL)**

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



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



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

