

# DISCLOSURES AND CONFLICT OF INTEREST

MC has received support from Boehringer Ingelheim

Honoraria for educational meetings performed on behalf of Boehringer Ingelheim, Lilly, Novartis, AstraZeneca and Sanofi

Advisory Board member for AstraZeneca & MSD



**IDENTIFYING**  
**CKD**

**BEFORE IT'S TOO LATE**

**Professor Mark Cooper**

**THERE IS NO REVERSE!**

36 WEEKS

**ONLY SLOWER**



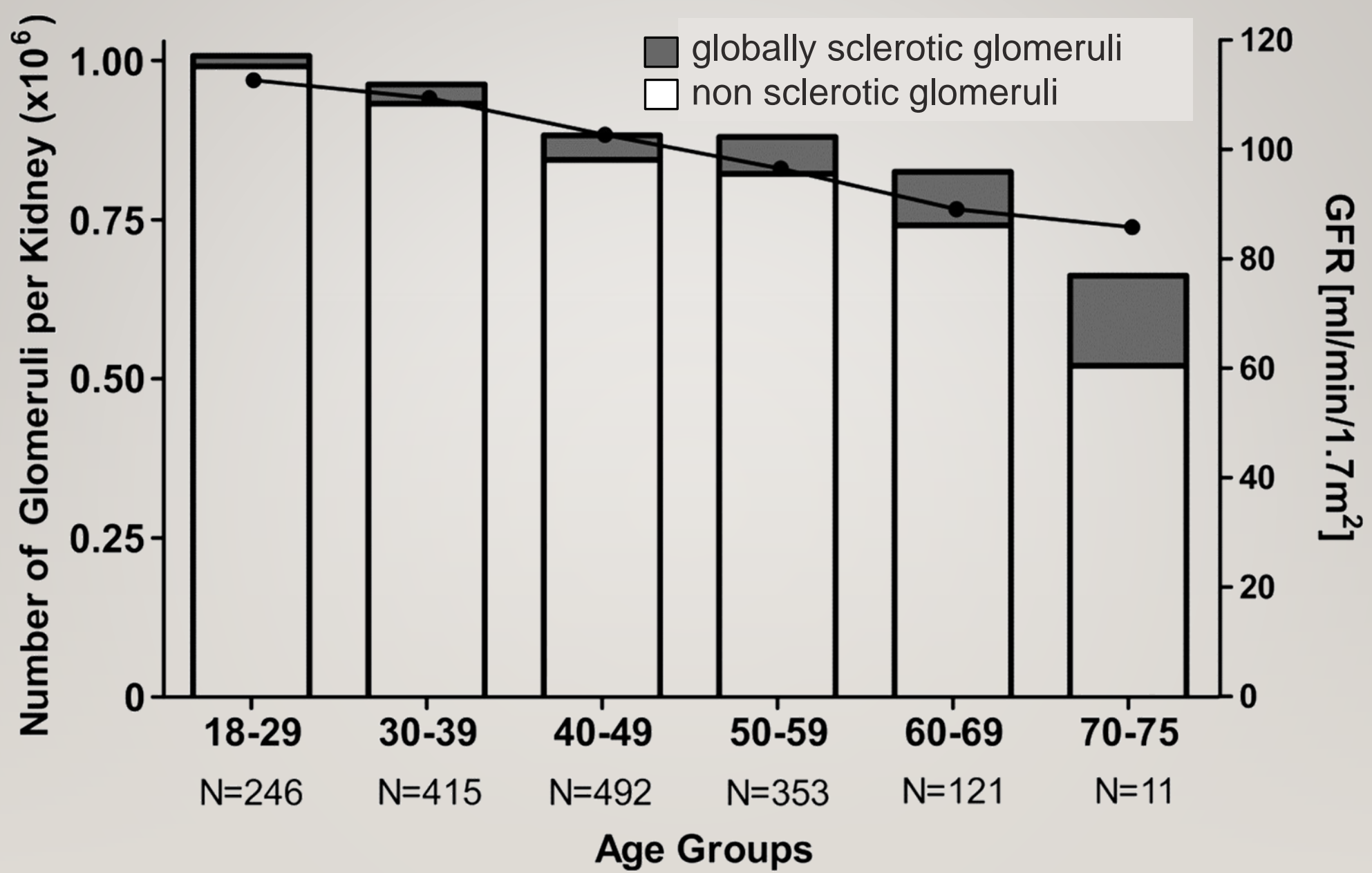
**OR FASTER**

**NEPHRON NUMBERS**



By the time you discover that  
the eGFR is abnormal ( $< 60$ )

Subjects have irreversibly lost  
half of their functioning nephrons

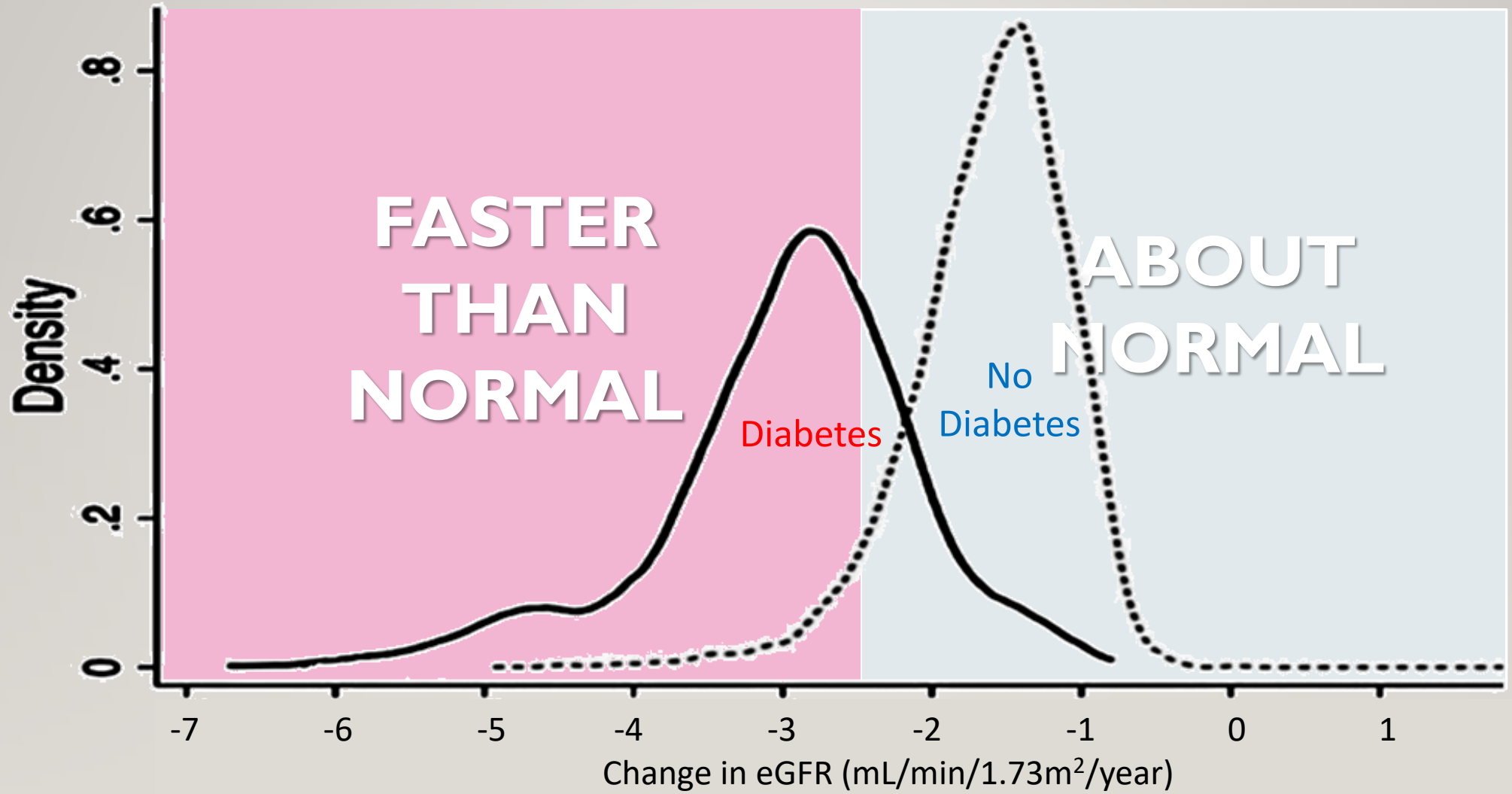




**HOW CAN WE KNOW  
THAT GFR DECLINE IS GOING TOO FAST  
(SO THAT WE MIGHT SLOW THIS GFR DECLINE  
BEFORE IT'S TOO LATE)**

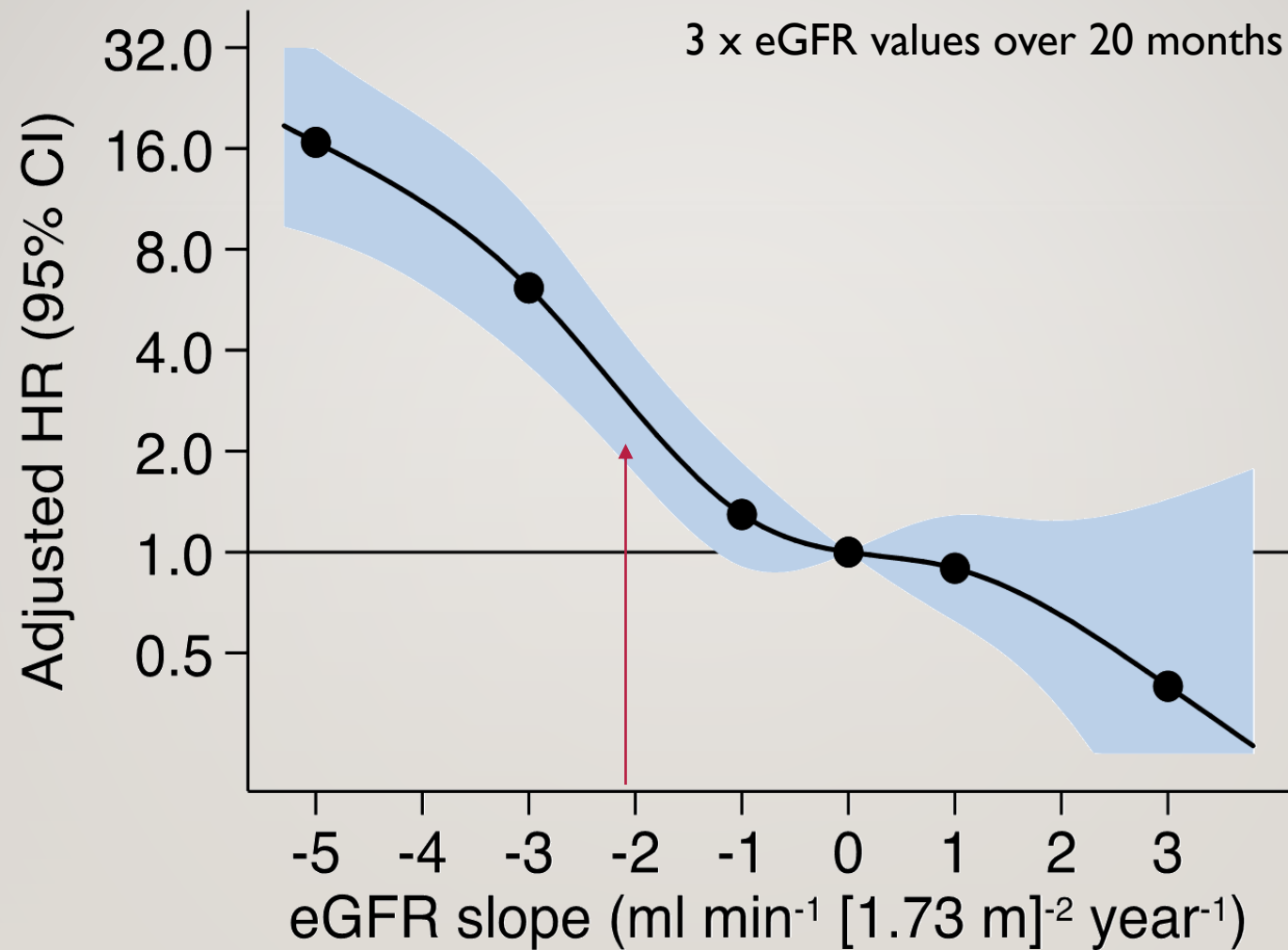
# Some patients are more likely to develop CKD

- they are already “speeding” (rapid decline in GFR)
- elevated albuminuria
- other co-morbidity (e.g. CVD, PVD, NAFLD, retinopathy)
- poorly controlled BP
- other risk factors (e.g. Indigenous, FHx, etc)





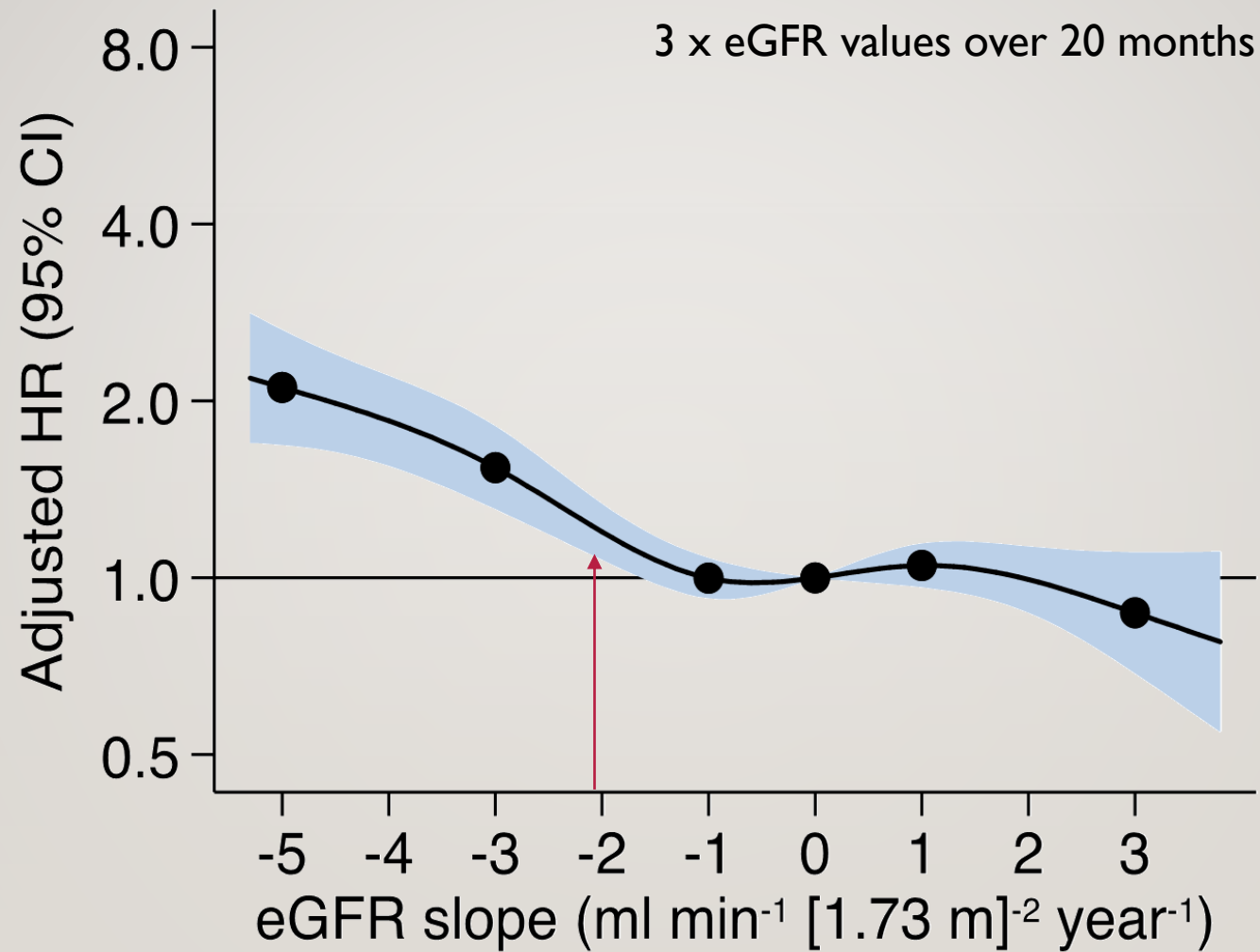
# Change in eGFR and combined major renal events



Oshima et al. Diabetologia (2019)

**ADVANCE Study.** eGFR slope estimated using three measurements of eGFR at 4, 12 and 24 months after randomisation over 20 months. Values were trimmed at a slope of  $<-5.4$  and  $>3.8 \text{ ml min}^{-1} (1.73 \text{ m})^{-2} \text{ year}^{-1}$  (each included 1.0% of participants). Knots were placed at -5, -3, -1, 1 and 3  $\text{ml min}^{-1} (1.73 \text{ m})^{-2} \text{ year}^{-1}$ , using 0  $\text{ml min}^{-1} (1.73 \text{ m})^{-2} \text{ year}^{-1}$  as the reference point. Covariates: Registration values of age, sex, region of residence, duration of diabetes, log-transformed UACR, systolic BP, diastolic BP, a history of macrovascular disease, smoking, drinking, treated hypertension,  $\text{HbA}_{1c}$ , HDL-cholesterol, LDL-cholesterol, log-transformed triacylglycerol and BMI, 4-month eGFR and randomised treatment allocation (BP and glucose treatment)

# Change in eGFR and all cause mortality



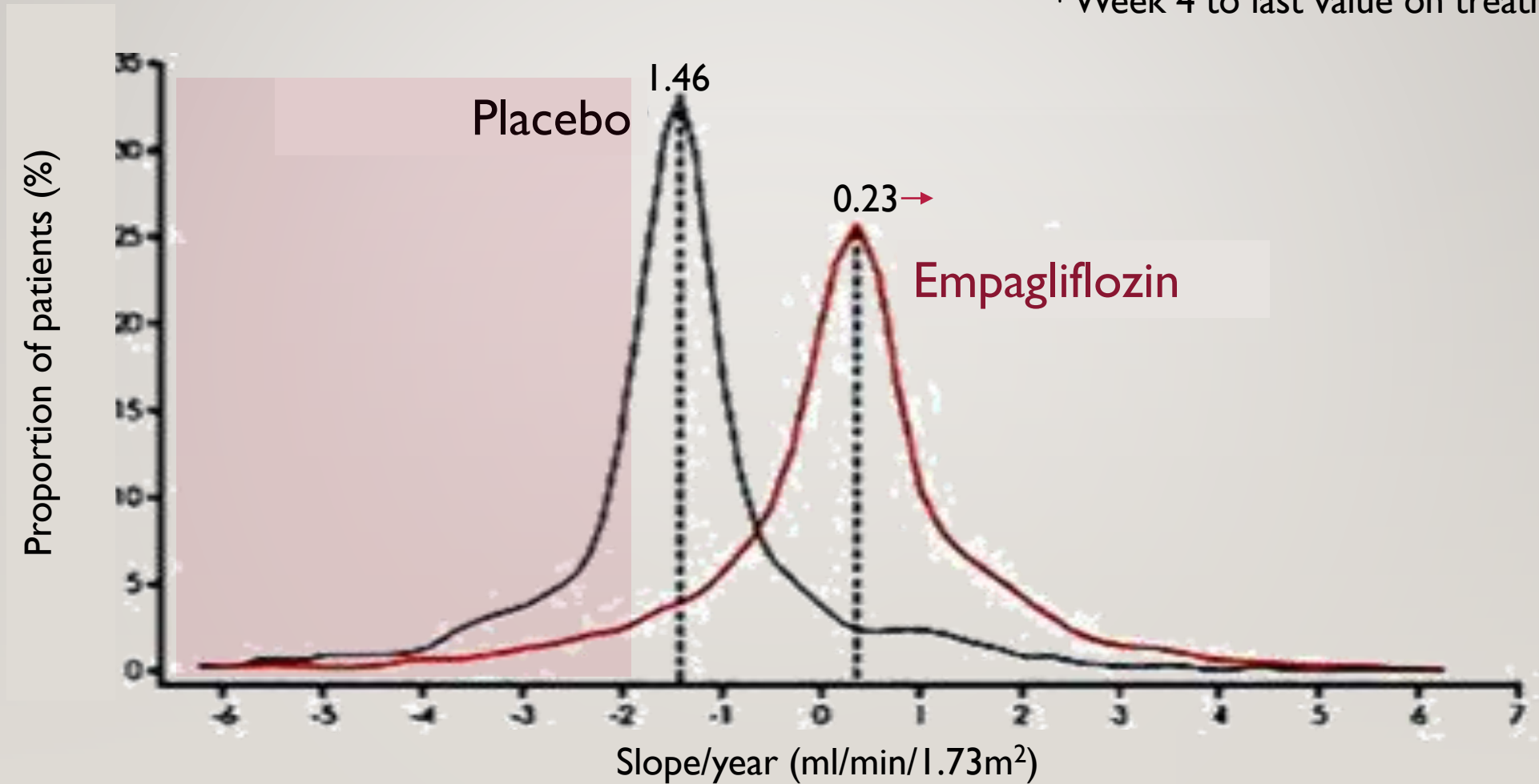
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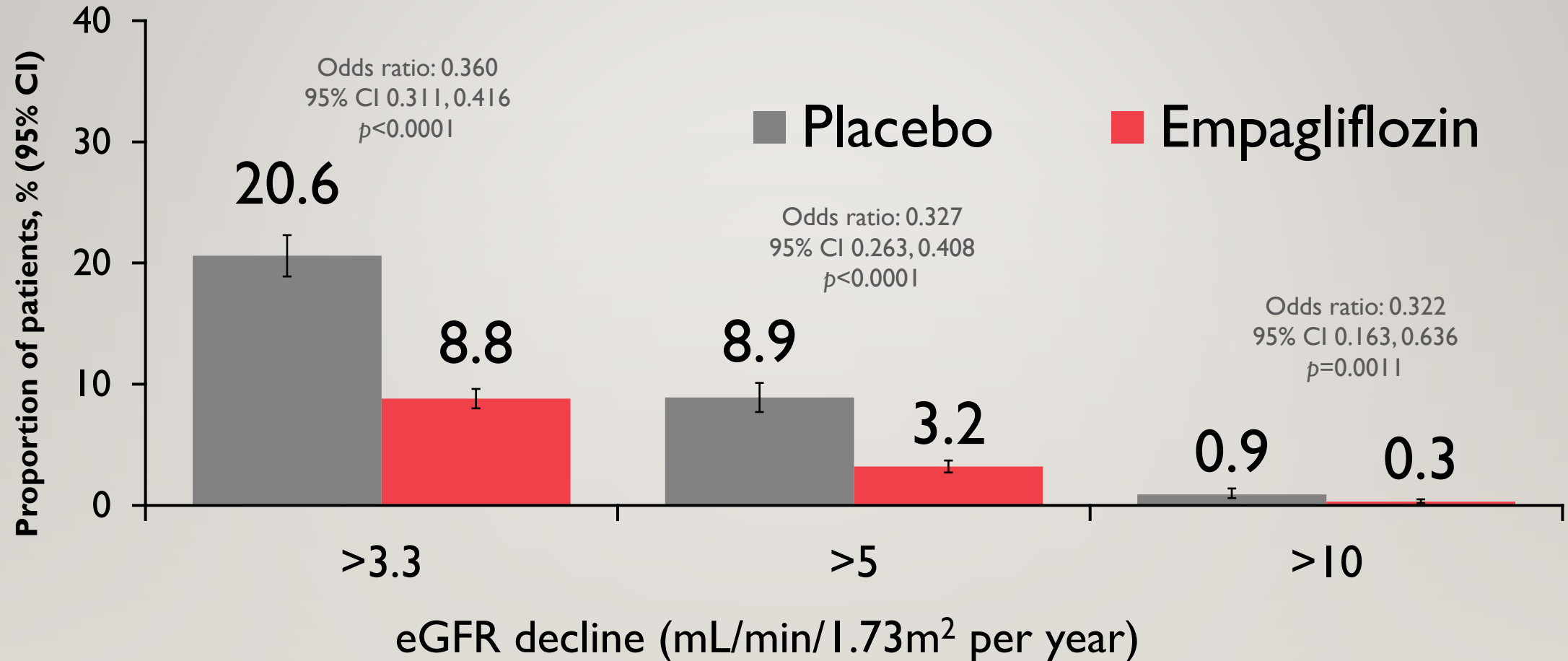
# Slower decline in eGFR when treated with empagliflozin



\* Week 4 to last value on treatment



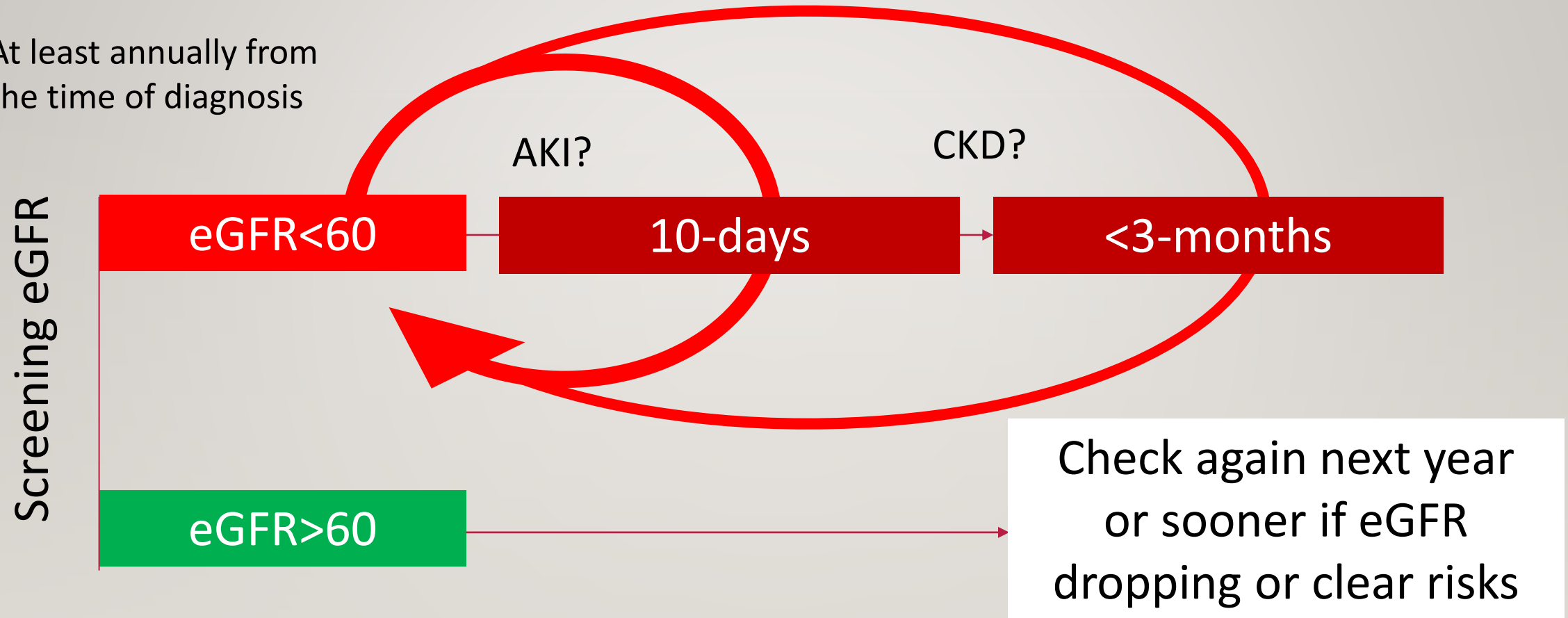
# Fewer fast progressors when treated with empagliflozin



Logistic regression analysis included treatment, sex, baseline BMI category, baseline HbA1c category, baseline eGFR category, geographical region and age, in patients treated with  $\geq 1$  dose of study drug. eGFR assessed by MDRD formula. Baseline eGFR values were available for 6967 participants. Median treatment duration was 2.6 years. Median observation time was 3.1 years. BMI, body mass index; eGFR, estimated glomerular filtration rate; HbA1c, glycated hemoglobin; MDRD, Modification of Diet in Renal Disease.

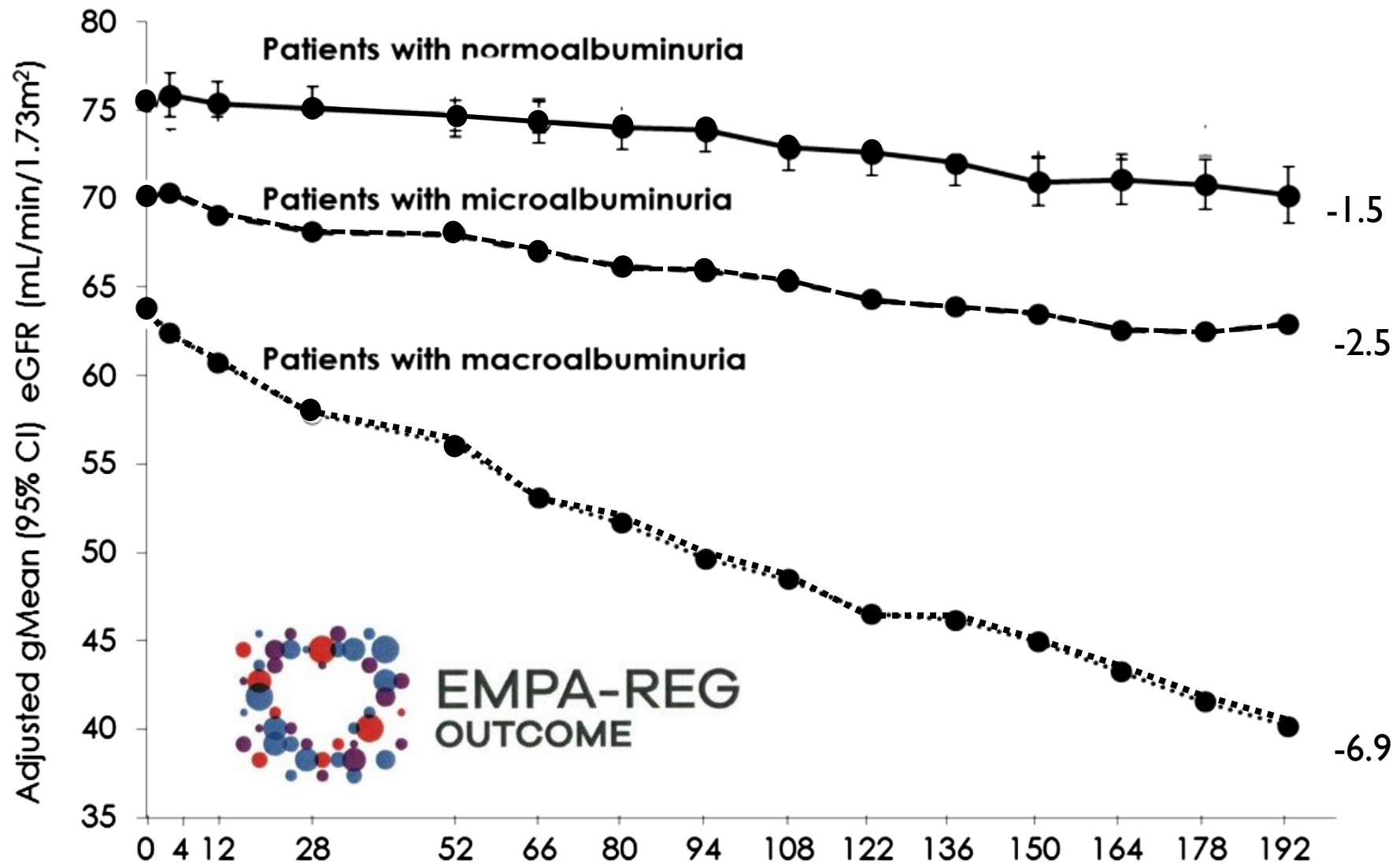
# Annual eGFR screening

At least annually from the time of diagnosis



# But some patients are more likely to get CKD sooner

- fast declining GFR
- **elevated albuminuria**
- other co-morbidity (e.g. CVD, PVD, NAFLD, retinopathy)
- poorly controlled BP
- other risk factors (e.g. Indigenous, FHx, etc)





# Identifying CKD in Type 2 Diabetes<sup>1</sup>

Test **AER** in all patients with T2D (without CKD)

**From the time of diagnosis & annually thereafter**

It can be most easily quantified by measuring the urine albumin-to-creatinine ratio (UACR)<sup>1,2</sup>

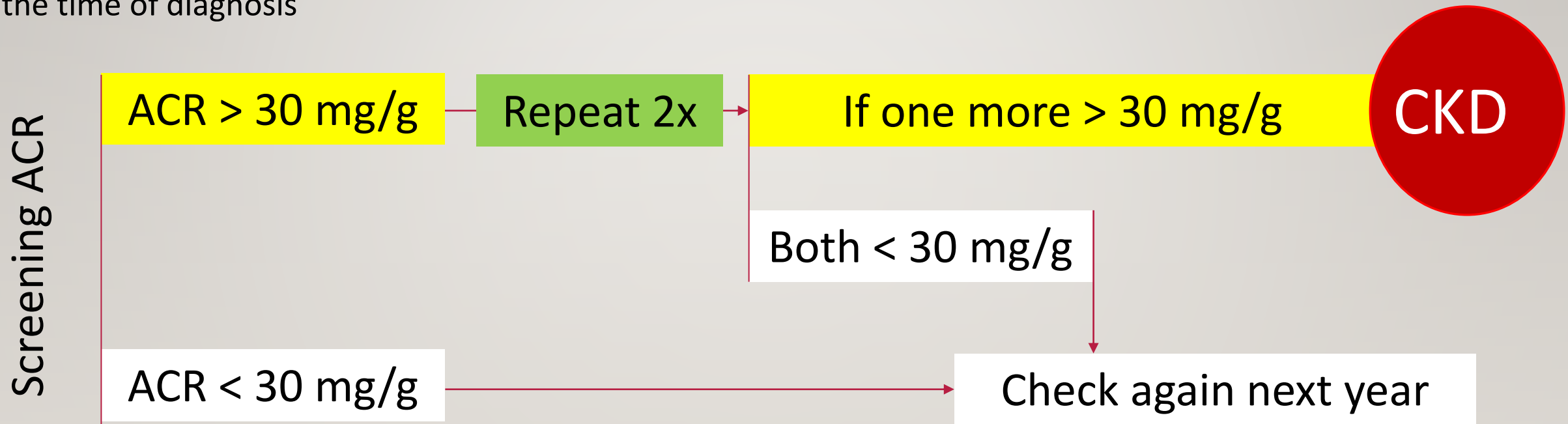
UACR = albumin concentration (mg)  
creatinine concentration (g)

- Measured using a single spot urine test (morning sample preferred).
- Albuminuria can also be quantified by collecting 24-h urine<sup>1,3,4</sup>



# SCREENING FOR AN ELEVATED AER

At least annually from  
the time of diagnosis

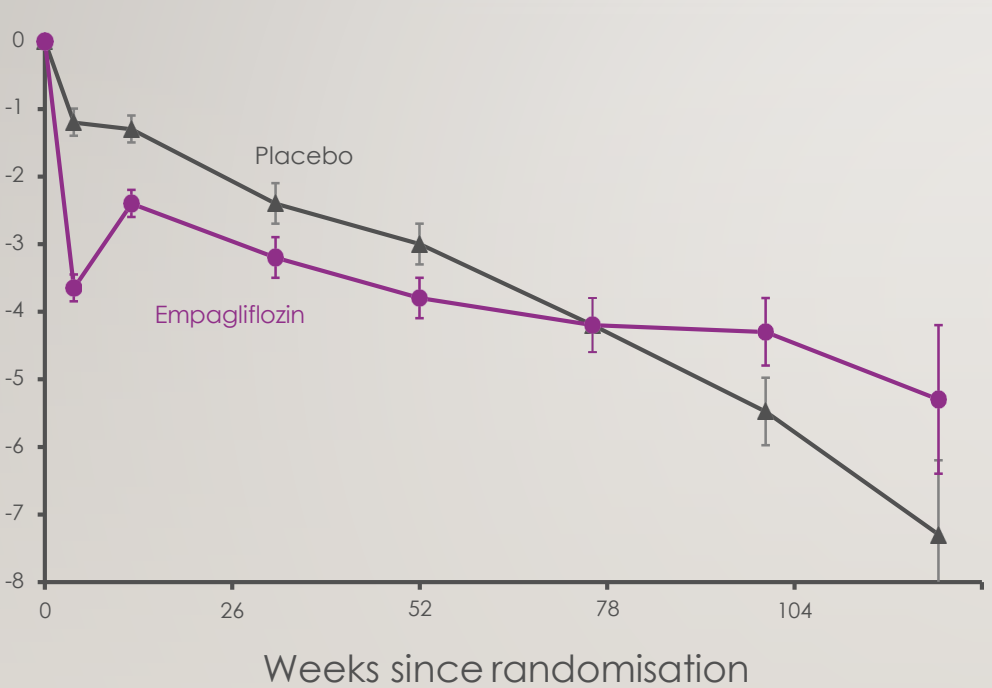


# **But some patients are more likely to get CKD sooner**

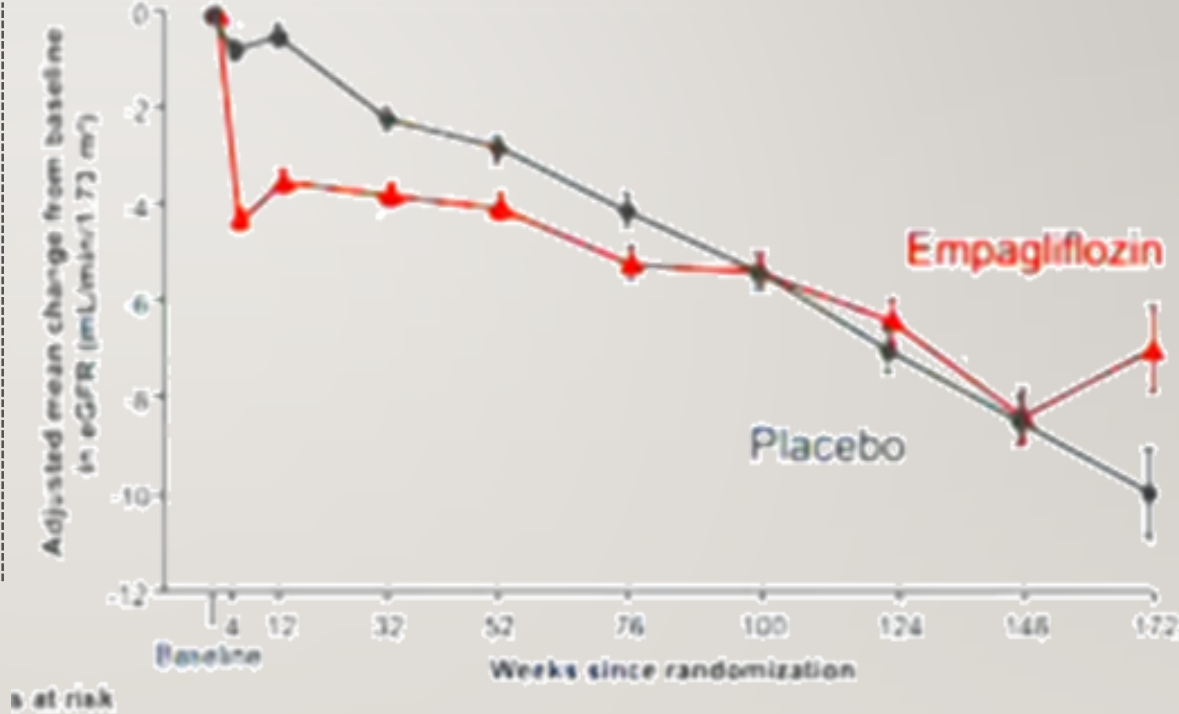
- fast declining GFR
- elevated albuminuria
- **other co-morbidity (e.g. CVD, CHF, NAFLD, OSA)**
- poorly controlled BP
- other risk factors (e.g. Indigenous, FHx, etc)

# In patients with Heart Failure SGLT2 inhibition slows their decline in eGFR

 **EMPEROR-Reduced**

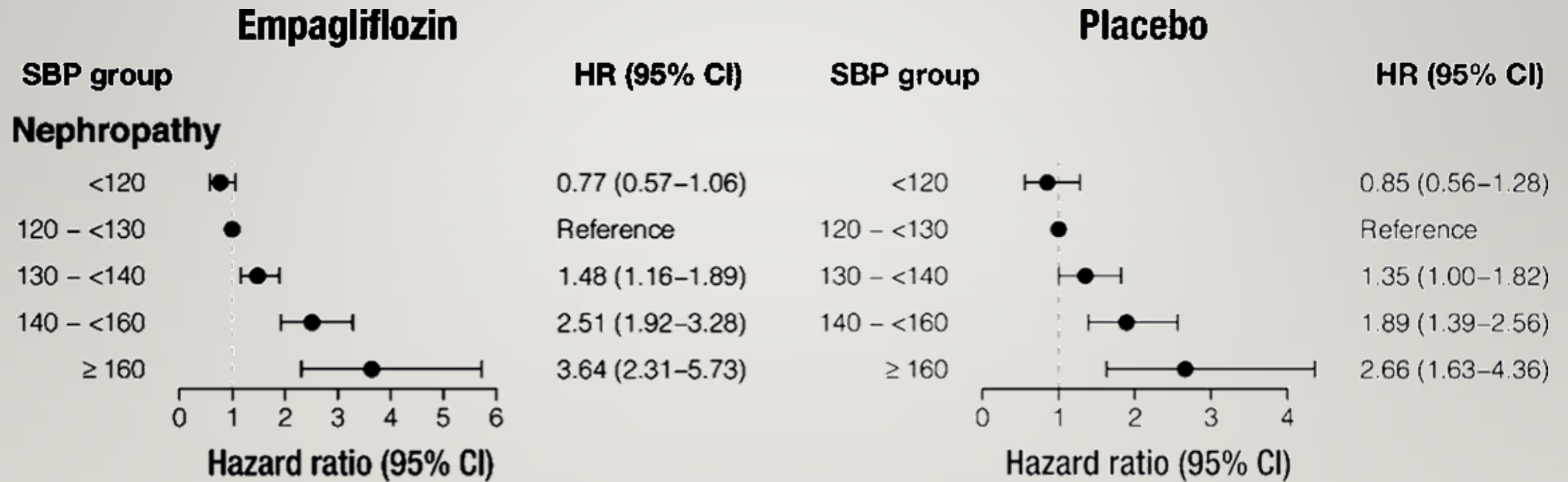


 **EMPEROR-Preserved**



# But some patients are more likely to get CKD sooner

- fast declining GFR
- elevated albuminuria
- other co-morbidity (e.g. CVD, CHF, MAFLD, retinopathy)
- **Poorly controlled BP**
- other risk factors (e.g. Indigenous, FHx, etc)

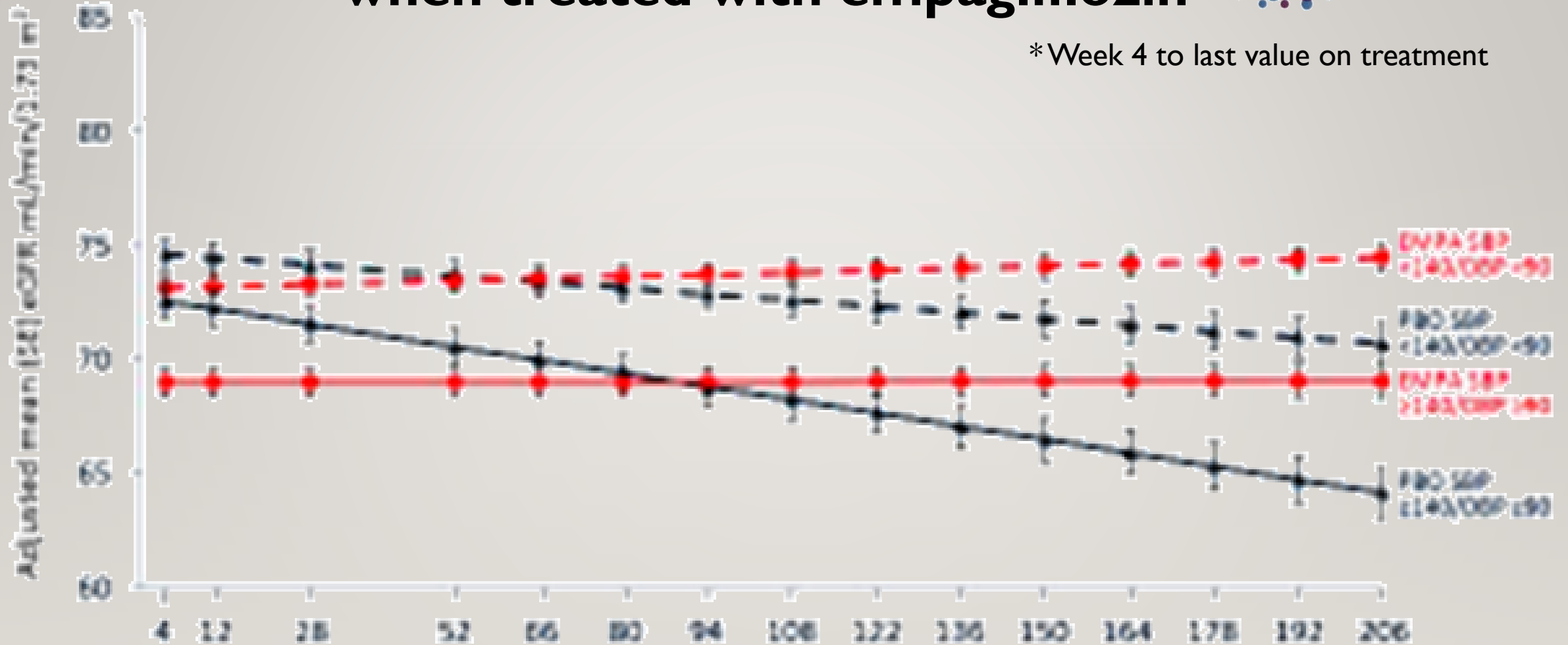


Blood pressure control is associated with incident CKD in patients with T2D + CVD

# Slower decline in eGFR when treated with empagliflozin



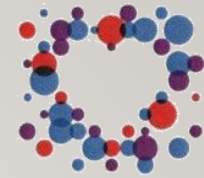
\* Week 4 to last value on treatment



# **But some patients are more likely to get CKD sooner**

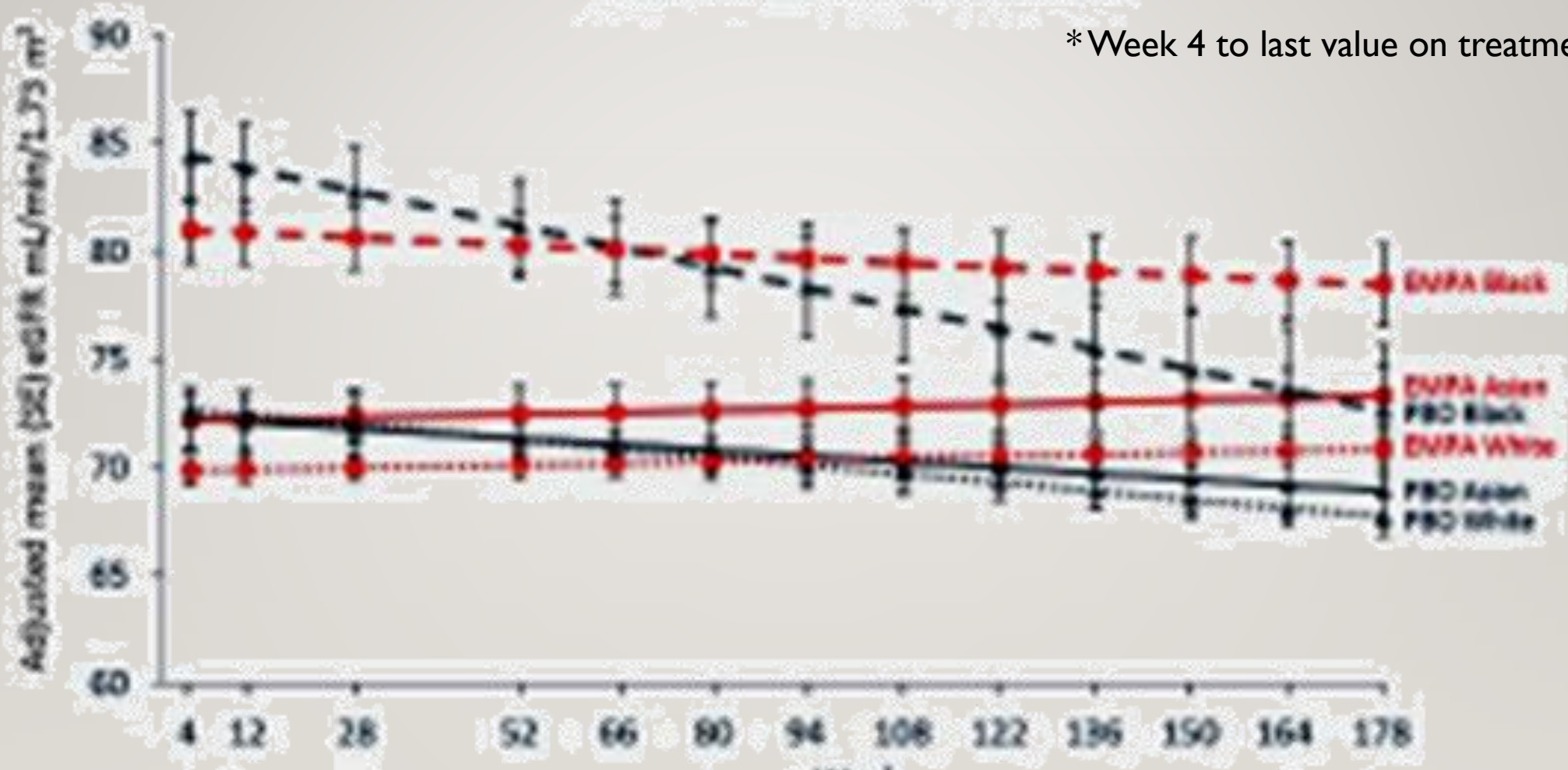
- fast declining GFR
- elevated albuminuria
- other co-morbidity (e.g. CVD, CHF, MAFLD, retinopathy)
- poorly controlled BP
- **other risk factors (e.g. Indigenous, Race, FHx, etc)**

# Slower decline in eGFR when treated with empagliflozin



EMPA-REG  
OUTCOME

\* Week 4 to last value on treatment



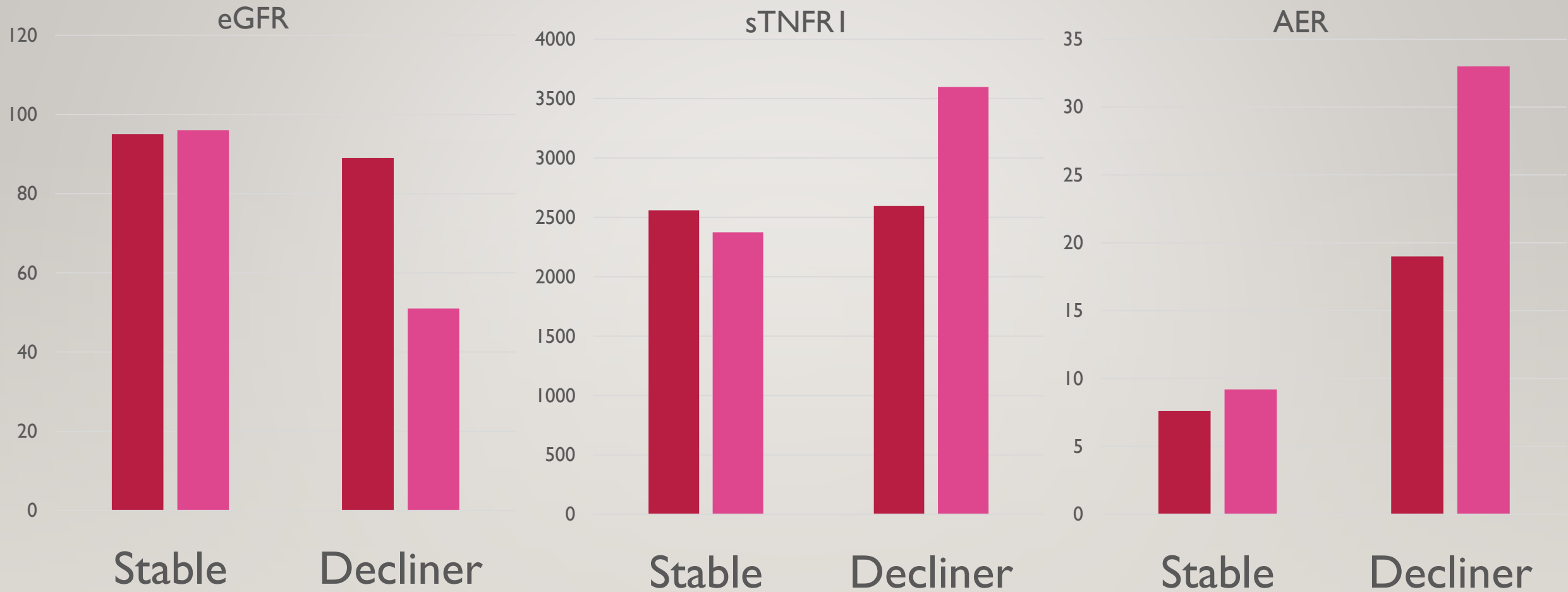


# NOVEL BIOMARKERS FOR PRE-CKD?

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- Soluble TNFR?
- Promarker D
- Urine biomarkers?
- Epigenetic markers?
- Artificial Intelligence?

# SOLUBLE TUMOR NECROSIS FACTOR RECEPTOR TYPE 1 (TNFR1) PREDICTS EARLY DECLINE IN PATIENTS WITH T2



\* from baseline in patients with an early decline in renal function ( $n = 30$ ) after 8 years of follow up

# Multi-Ethnic Study of Atherosclerosis (MESA) and Reasons for Geographic and Racial Differences in Stroke (REGARDS) cohorts.

## Case Cohort & Methods



**MESA** (n = 497 in the random subcohort and 163 incident CKD cases)

**REGARDS** (n = 497 in the random subcohort and 497 incident CKD cases)



**Patients without Diabetes**



**eGFR > 60ml/min/1.73m<sup>2</sup>**

## Outcome

**Incident chronic kidney disease**  
(eGFR < 60ml/min AND ≥ 40% decline from baseline)



**MESA**  
9.2-year median follow-up

*For each 2-fold increase*

**HR**

- **KIM-1** 1.38 (1.05- 1.81)
- **SuPAR** 1.96 (1.10- 3.49)
- **TNFR-1** 1.65 (1.04 -2.62)
- **TNFR-2** 2.02 (1.21- 3.38)
- **YKL-40** 1.38 (1.09- 1.75)



**REGARDS**  
9.4-year median follow-up

- **TNFR-1** 1.99 (1.43- 2.76)
- **TNFR-2** 1.76 (1.22- 2.54)

# PromarkerD

- Panel of three novel protein biomarkers (ApoA4, CD5L and IGFBP3)
- Combined with three clinical risk factors (HDL, age, eGFR)

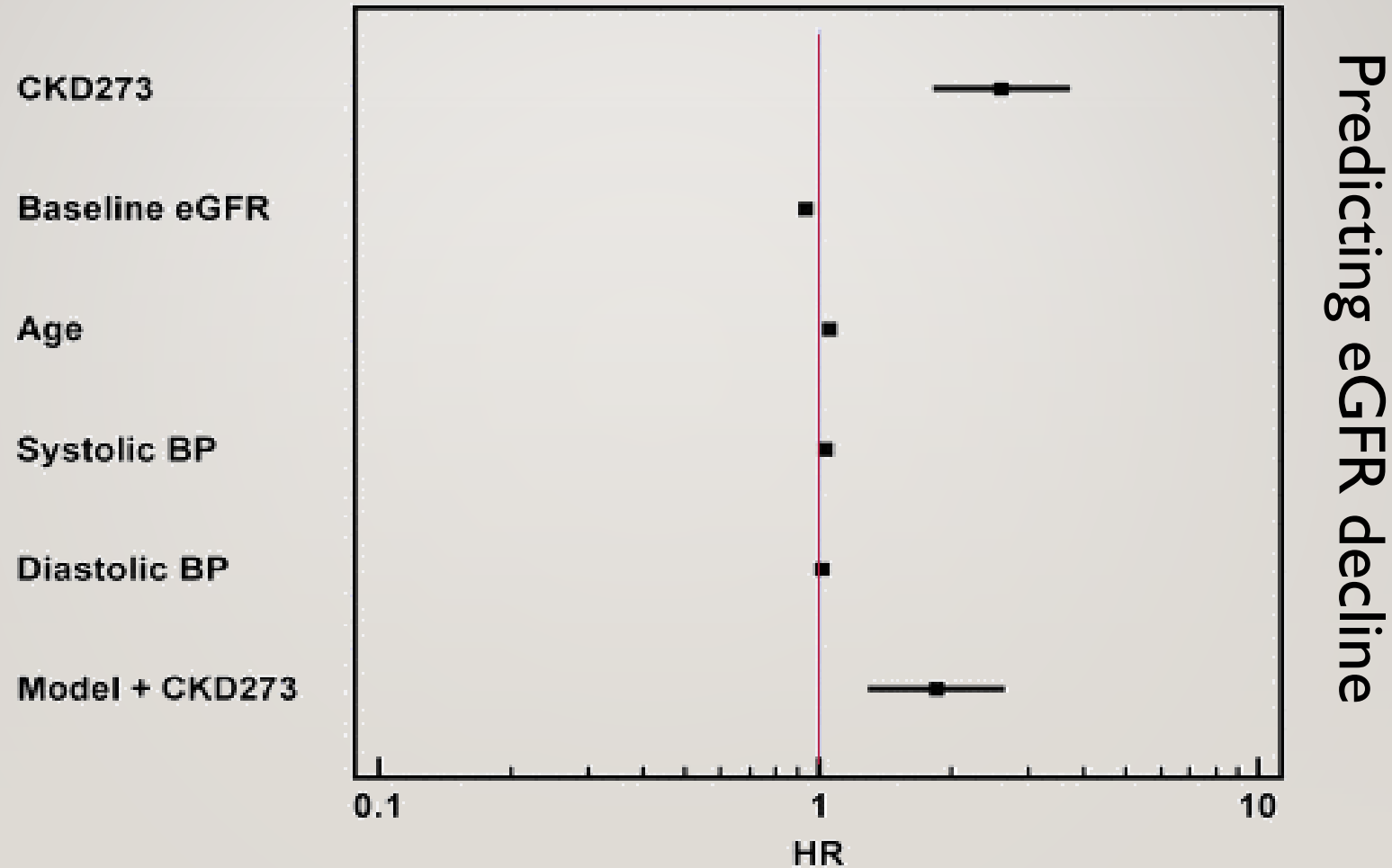
LOW RISK	MODERATE RISK	HIGH RISK
<b>0% to &lt;10%</b> <b>Low four-year risk of developing DKD.</b>	<b>10% to &lt;20%</b> <b>Moderate four-year risk of developing DKD.</b>	<b>20% to 100%</b> <b>High four-year risk of developing DKD.</b>
Standard diabetes monitoring. Retest annually.†	Consider more frequent monitoring. Retest every 3-6 months.†	Consider very close monitoring. Retest every 3 months.†

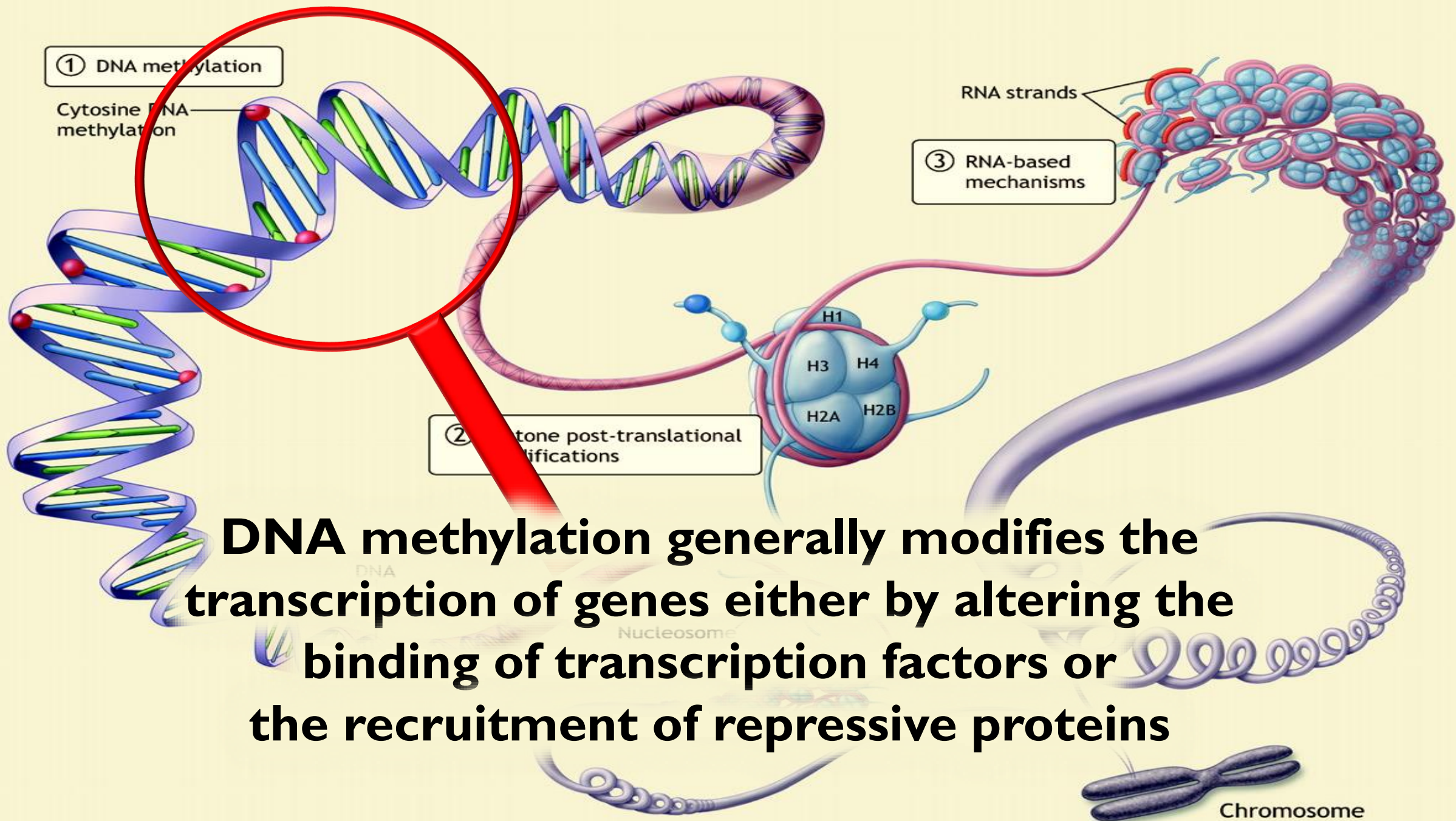
## Promarker D – predicting loss of kidney function in the Fremantle Diabetes Study

Rapid eGFR decline:	≥30% decline	Incident CKD	eGFR trajectory	≥5mL/yr decline
<b>Development Cohort</b>				
AUC (95%CI)	0.81 (0.75-0.87)	<b>0.89 (0.85-0.94)</b>	0.86 (0.80-0.93)	0.70 (0.61-0.80)
Sn/Sp (%)	97/62	<b>95/68</b>	84/82	61/73
Calibration <i>P</i>	0.06	<b>0.39</b>	0.41	0.07
<b>Validation Cohort</b>				
AUC (95%CI)	0.72 (0.63-0.82)	<b>0.88 (0.84-0.93)</b>	NI	0.62 (0.53-0.72)
Sn/Sp (%)	65/71	<b>86/78</b>	NI	69/57
Calibration <i>P</i>	0.68	<b>0.77</b>	NI	0.61

# CKD273 – urine proteomic profile

based on 273 protein fragments (mostly collagen peptides)



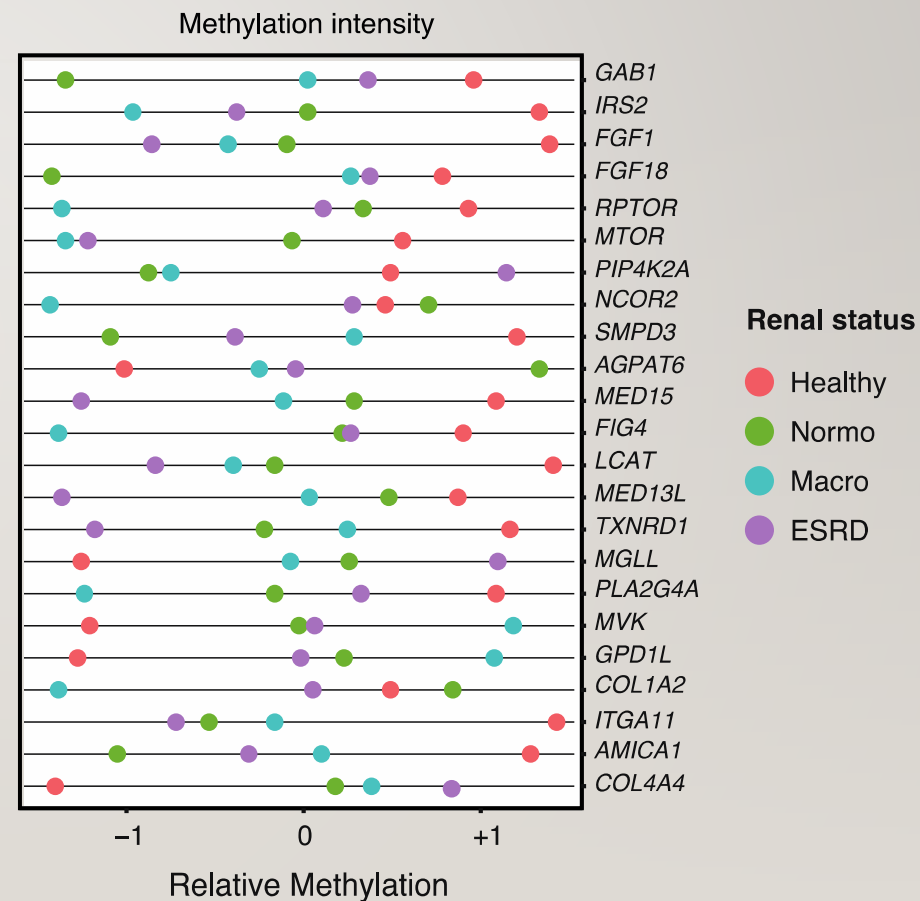


**DNA methylation generally modifies the transcription of genes either by altering the binding of transcription factors or the recruitment of repressive proteins**

# Convergence of DNA methylation and Diabetes-Associated pathways

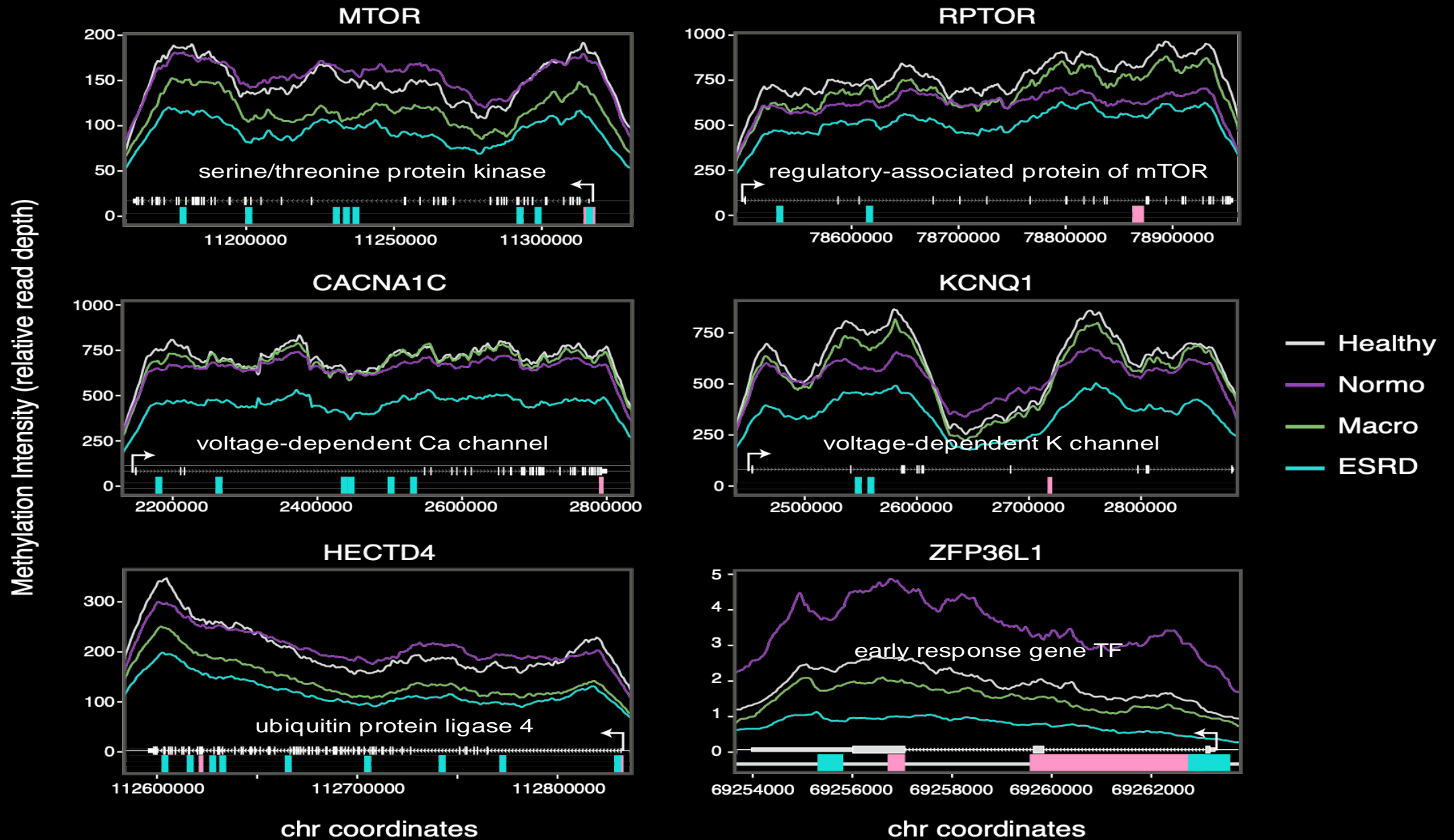
## Differentially methylated genes associated with diabetic kidney disease

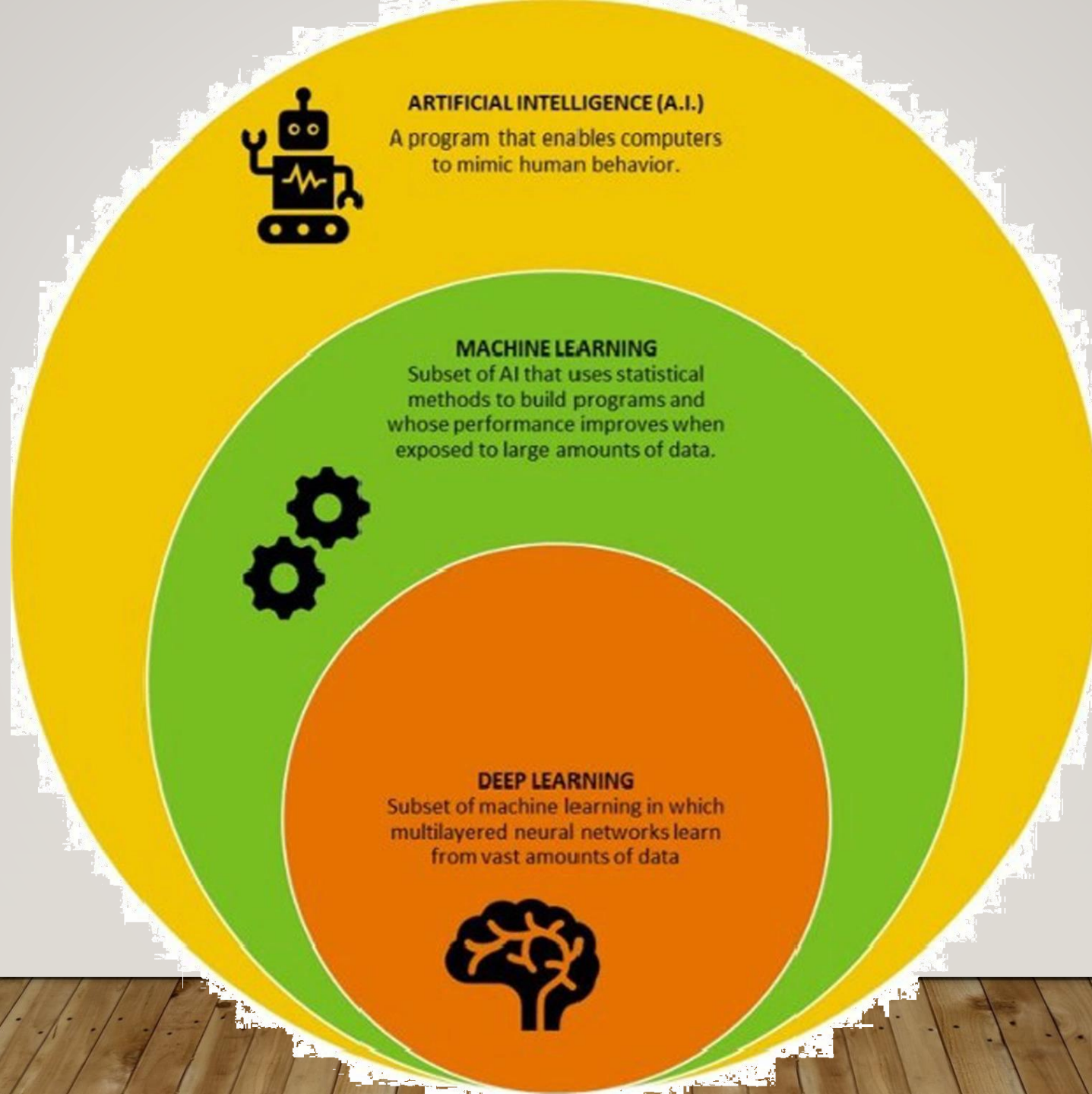
Gene	Ensembl ID	P value	DMR coordinates	Diabetes associated pathways
<i>GAB1</i>	ENST00000262995	1.69E-04	4:144404420-144404861	Insulin signaling
<i>IRS2</i>	ENST00000375856	1.58E-03	13:110443347-110443786	Insulin signaling
<i>FGF1</i>	ENST00000494579	6.39E-03	5:142078293-142078636	Insulin signaling
<i>FGF18</i>	ENST00000274625	6.63E-03	5:170909699-170911557	Insulin signaling
<i>RPTOR</i>	ENST00000537330	7.71E-03	17:78549200-78549666	Regulates insulin release and signaling
<i>MTOR</i>	ENST00000361445	8.79E-03	1:11307862-11309107	Regulates insulin release and signaling
<i>PIP4K2A</i>	ENST00000376573	9.05E-05	10:22874619-22875118	Lipid metabolism
<i>NCOR2</i>	ENST00000405201	2.82E-04	12:124864792-124865295	Lipid metabolism
<i>SMPD3</i>	ENST00000219334	5.86E-04	16:68399936-68400660	Lipid metabolism
<i>AGPAT6</i>	ENST00000396987	4.00E-03	8:41470262-41470706	Lipid metabolism
<i>MED15</i>	ENST00000425759	4.69E-03	22:20918886-20919334	Lipid metabolism
<i>FIG4</i>	ENST00000441478	5.14E-03	6:110104322-110104932	Lipid metabolism
<i>LCAT</i>	ENST00000264005	5.37E-03	16:67975360-67975797	Lipid metabolism
<i>MED13L</i>	ENST00000281928	5.99E-03	12:116716013-116716425	Lipid metabolism
<i>TXNRD1</i>	ENST00000525566	6.44E-03	12:104687048-104687537	Lipid metabolism
<i>MGLL</i>	ENST00000434178	7.76E-03	3:127416330-127417102	Lipid metabolism
<i>PLA2G4A</i>	ENST00000367466	7.77E-03	1:186955914-186956703	Lipid metabolism
<i>MVK</i>	ENST00000539696	8.82E-03	12:110033681-110035595	Lipid metabolism
<i>GPD1L</i>	ENST00000282541	9.67E-03	3:32185089-32185514	Lipid metabolism
<i>COL1A2</i>	ENST00000297268	1.34E-04	7:94025128-94026206	Integrin cell interaction
<i>ITGA11</i>	ENST00000315757	5.44E-03	15:68608377-68609157	Integrin cell interaction
<i>AMICA1</i>	ENST00000356289	6.90E-03	11:118069845-118070106	Integrin cell interaction
<i>COL4A4</i>	ENST00000396625	8.02E-03	2:227972698-227973120	Integrin cell interaction





# Genes implicated in DN are differentially methylated





- Indicated for DKD Stages 1-3b (excluding G1&A1 and G2&A1)
- NY state-approved for clinical use
- Granted Breakthrough Device designation by the US Food and Drug Administration (FDA)



### Standard blood draw

Biomarkers *sTNFR-1, sTNFR-2, and KIM-1*



### Standard Clinical Data Elements

*eGFR, UACR, serum calcium, HbA1c, systolic BP, platelets, AST*

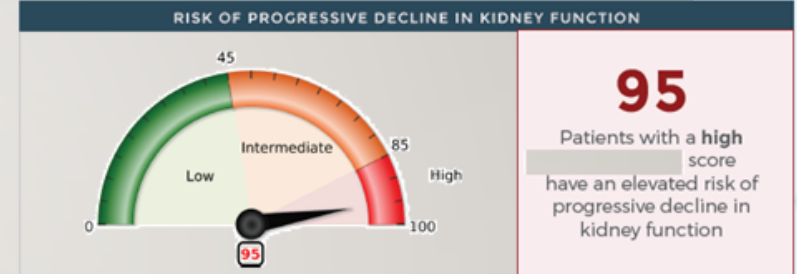
**Machine Learning Algorithm  
Harmonizes Disparate Data**

# Actionable Risk Assessment

Ordered by Dr. Fran Lake  
Collection Date 8/4/2020  
Report Date 8/9/2020  
Specimen ID 665544

## Test Report

PATIENT INFORMATION			
NAME	SEX	DATE OF BIRTH	MEDICAL RECORD #
Jane Lee	F	1/1/1960	00998877



score ranges from 0-100 and correlates with the probability of progressive decline in kidney function in the study population. Risk classification is provided to guide interpretation of the risk score using cut-offs related to clinical outcomes.

SIGNED	DATE	TIME
Laboratory Director Michael J. Donovan PhD, MD, CLIA, Renalytix AI, 1016th Ave 3rd Floor, Room 334, New York, NY 10018, CLIA Number 330256875		

This test was developed and its performance characteristics determined by Renalytix AI Inc. It has not been cleared or approved by the FDA nor is it currently required to be. The laboratory is regulated under CLIA as qualified to perform high-complexity testing. The test is used for clinical purposes. It should not be regarded as investigational or for research. See page 2 for further details.

EXAMPLE OF CLINICAL PATHWAY			
Frequency of Monitoring / Referral <sup>1</sup>		Comprehensive Strategy to Maximize Protection for Diabetic Kidney Disease Progression and Cardiovascular Disease <sup>2,3</sup>	
Monitoring 3x/year	Nephrology Referral	Titrate ACEi or ARB to maximally tolerated dose	Strongly consider SGLT2 inhibitor therapy unless contraindicated

<sup>1</sup> KDIGO 2012 Clinical Practice Guideline for the Evaluation and Management of Chronic Kidney Disease [https://kdigo.org/wp-content/uploads/2012/02/KDIGO\\_2012\\_CKD\\_CPG.pdf](https://kdigo.org/wp-content/uploads/2012/02/KDIGO_2012_CKD_CPG.pdf)  
<sup>2</sup> Executive Summary of the 2020 KDIGO Diabetes Management in CKD Guideline <https://kdigo.org/2020/08/04/>  
<sup>3</sup> ACEi guideline: <https://pubs.nidDKD.org/doi/10.1001/jama.2013.278133>

RENALYTIX AI