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Reporting Summary

Nature Research wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Research policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.

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n/a	Confirmed					
	The exact	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement				
	🔀 A stateme	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly				
	The statistical test(s) used AND whether they are one- or two-sided Only common tests should be described solely by name; describe more complex techniques in the Methods section.					
	A description of all covariates tested					
	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons					
	A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)					
	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i>) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted Give <i>P</i> values as exact values whenever suitable.					
\boxtimes	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings					
\boxtimes	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes					
	Estimates of effect sizes (e.g. Cohen's d, Pearson's r), indicating how they were calculated					
Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.						
Software and code						
Poli	Policy information about <u>availability of computer code</u>					
Da	ata collection	No software was used.				
Data analysis		The development version of R package QDRS and documentation of quantitative disease risk scores from eMERGE dataset are available online (https://github.com/danqingxu/QDRS).				
For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.						

Data

Policy information about availability of data

All manuscripts must include a <u>data availability statement</u>. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A list of figures that have associated raw data
- A description of any restrictions on data availability

The eMERGE- III genetic datasets with linked phenotypes are accessible through dbGAP (accession number: phs001584.v1.p1).

Field-specific reporting						
Please select the o	Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.					
∑ Life sciences	Behavioural & social sciences Ecological, evolutionary & environmental sciences					
For a reference copy of	f the document with all sections, see <u>nature.com/documen</u>	ts/nr-reporting-summary-flat.pdf				
Life sciences study design						
All studies must dis	isclose on these points even when the disclosi	ure is negative.				
Sample size	No sample size calculation is performed. The sample size is the size of cohort constructed based on kidney function data and chronic kidney disease stage status.					
Data exclusions	Patients with end-stage renal disease (ESRD) are excluded since the end stage is not of our primary interest.					
Replication	Not applicable.					
Randomization	Not applicable.					
Blinding	Not applicable.					
Reporting for specific materials, systems and methods						
We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.						
Materials & experimental systems Methods						
n/a Involved in th	he study n/a Invo	lved in the study				
Antibodies	es 🖂 🗀 (ChIP-seq				
Eukaryotic	c cell lines	low cytometry				
Palaeontol	Palaeontology and archaeology MRI-based neuroimaging					

Palaeontology and archaeology
Animals and other organisms
Human research participants

 \boxtimes

Clinical data

Dual use research of concern