

RaDaR: an Integrated Strategy for Rare Disease

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Understanding Rare Diseases is challenging

- Many different rare diseases, each affecting 1 in 1000
 - Rare diseases in total affect an estimated 1 in 10 population and a large proportion of renal failure
 - Each clinician manages few patients with some diseases so may lack expertise
 - Most patients do not know much about their disease although networks easier to form in social media
- Potential market for therapies is seen as small
 - Historically unattractive to commercial and academic research funders
- Hard to assemble large enough cohorts to perform studies that are powerful enough to answer important questions robustly
 - Especially true for non-monogenic rare diseases

Rare \neq Monogenic

2 UK initiatives to address these challenges

- **NIHR BioResource for Rare Diseases**



- **RaDaR – the Rare Renal Disease Registry**



- Different approaches but some commonality:
 - Recruitment from multiple centres across a country
 - Aggregation of large datasets
 - Use of routinely collected hospital record data to enrich dataset



Rare Diseases WGS Study

13,037 DNA samples from patients at 57 UK hospitals

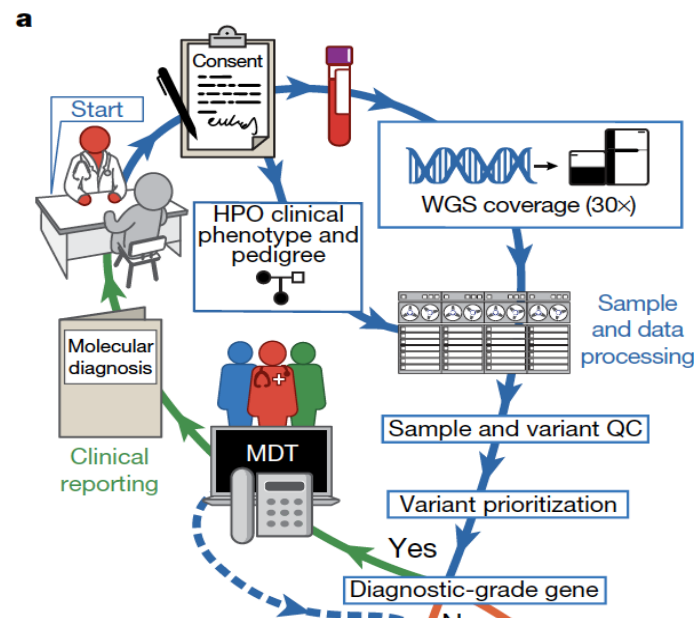
Recruitment 2012-2015

NIHR | BioResource

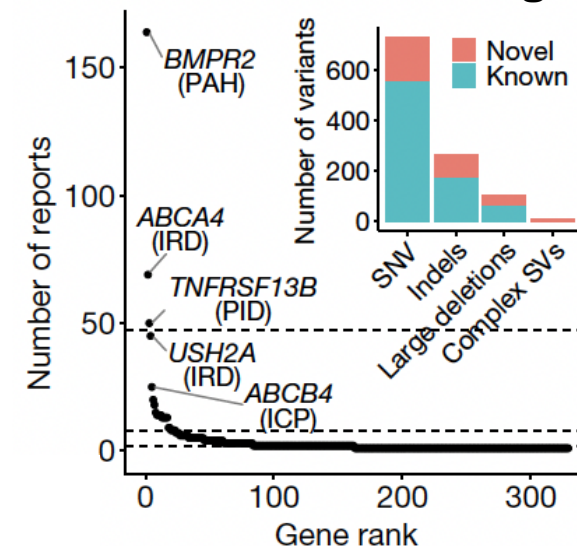
Genomics
england



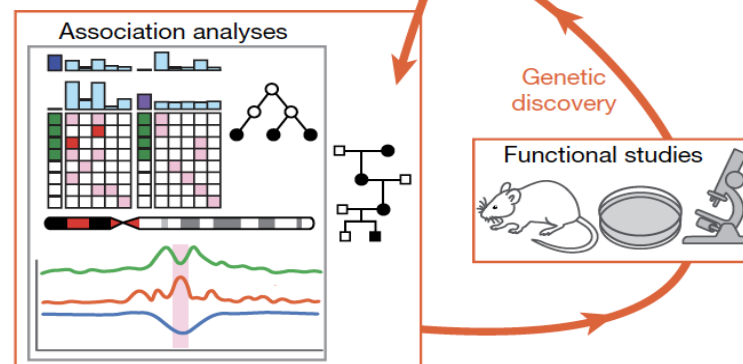
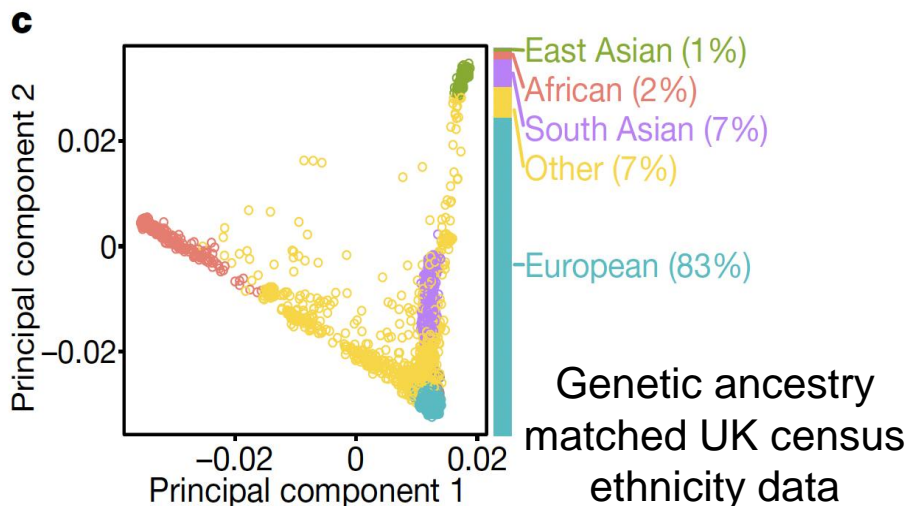
Bleeding, thrombotic and Platelet Disorders	BPD	Multiple Primary Malignant Tumours	MPMT
Cerebral Small Vessel Disease	CSVD	Neurological and Developmental Disorders	NDD
Ehler-Danlos Syndromes	EDS	Neuropathic Pain Disorders	NPD
Rare Diseases Pilot-II	GEL	Pulmonary Arterial Hypertension	PAH
Hypertrophic Cardiomyopathy	HCM	Primary Immune Disorders	PID
Intrahepatic Cholestasis of Pregnancy	ICP	Primary Membranoproliferative Glomerulonephritis	PMG
Inherited Retinal Disorders	IRD	Stem cell and Myeloid Disorders	SMD
Leber Hereditary Optic Neuropathy	LHON	Steroid Resistant Nephrotic Syndrome	SRNS
biobank ^{uk} recruiting the nation's future generations		UK Biobank – Extreme Red Cell Traits	
		UKBio	



Diagnostic Grade Reports issued for 1100 variants in 329 genes



Turro et al Nature 2020



Variation in coverage at disease genes was much greater for WES platforms that would have missed 2.67-10.5% SNVs reported in this study

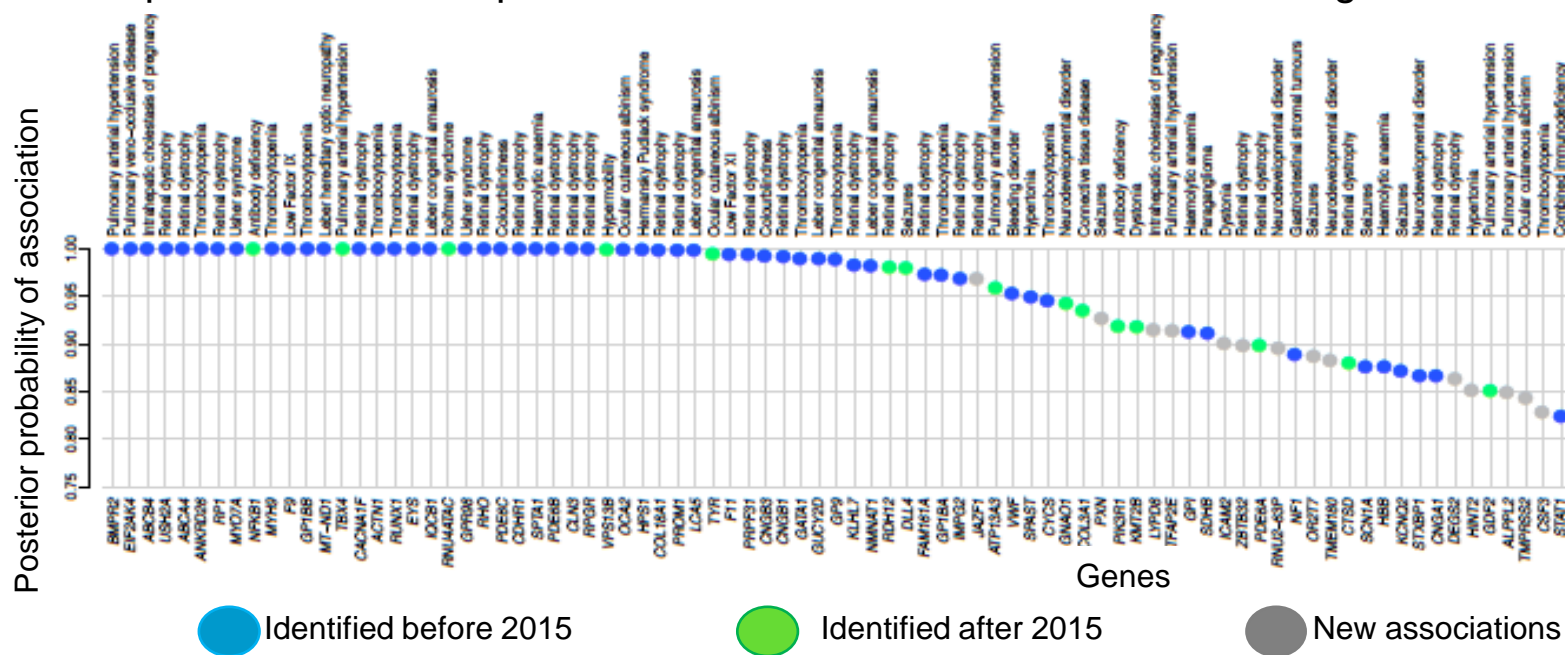


NIHR | BioResource WGS Study

13,037 DNA samples from patients at 57 hospitals

Strong evidence for 99 genetic associations

Groups of rare disease patients based on HPO term driven clustering



Of these 99 associations, 61 are consistent with firmly established evidence and a further 18 have been reported in the literature 2015-20, either by us or by others

Article

Whole-genome sequencing of patients with rare diseases in a national health system

<https://doi.org/10.1038/s41586-020-2265-1>

Received: 23 December 2018

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Check for updates

Article

Whole-genome sequencing of a sporadic primary immunodeficiency cohort

<https://doi.org/10.1038/s41586-020-2265-1>

Received: 1 December 2018

A list of authors and affiliations appears at the end of the paper

Science

Germline selection shapes human mitochondrial DNA diversity

WEI WEI, SALIH TUNA, MICHAEL J. KEOGH, KATHERINE MARIA A. K. BITNER-GUINDICZ, PATRICK F. CHINNERY

THE LANCET Respiratory Medicine

Volume 7, Issue 3, March 2019, Pages 227-238

Genetic determinants of risk in pulmonary arterial hypertension: international genome-wide association study

Christopher J. R. Philpott, Marie Barbara Girard, Marika Kaakinen, A. B. Kathleen E. Styrud, Salih Tuna, Julie M. MGN/DD/C3 Glomerulopathy Rare Dis

REPORT AJHG

De Novo Truncating Cause Intellectual D

Yoko Ito, Keren J. Carr, Isabelle Marey, Perinne Char, Alba Sanchis-Juan, Hans van Bokhoven, NIHR BioResource, CareRare Canada Cor Sarah Dyack, and E.

REPORT AJHG

A Fast Association Test for Pathogenic Variants Invol

Daniel Gree

REPORT AJHG

Bi-allelic Loss-of-Function in Progressive Epilepsy

Kathleen M. Gorman, Alba Sanchis-Juan, Ronit M. Pressler, Jenny Morton, Mary D. King, J. Helen Cross, Ingrid E. Schiffer, Tobias H. Study, Consortium, N. J. Clayton-Smith, Eamonn J.

REPORT AJHG

Large-Scale Whole-Genom Genetic Architecture of F Membranoproliferative G

Adam P. Levine, Melanie M.Y. Chan, H. Terence Cook, Sofie Ashford, Keren Claire Louise Harris, Paul McAlinden, Heather Maxwell, Karyn Megy, Kathleen E. Styrud, Salih Tuna, Julie MGN/DD/C3 Glomerulopathy Rare Dis

REPORT AJHG

Bi-allelic Mutation

Determination of

REPORT AJHG

Whole genome sequ that genetic conditio in intensively ill child

Courtney E. French, Isabelle Delon, Helen Dollin, Stephen Abbo, Topun Austin, Sarah Bowdin, R Disease, Next Generation Children Project, David H. Rowitch, and F. Lucy Raymond

REPORT AJHG

Telomerecat: A ploidy-agr method for estimating tel length from whole genom sequencing data

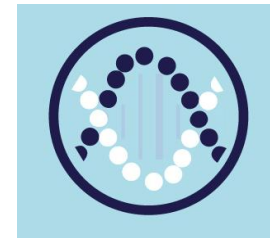
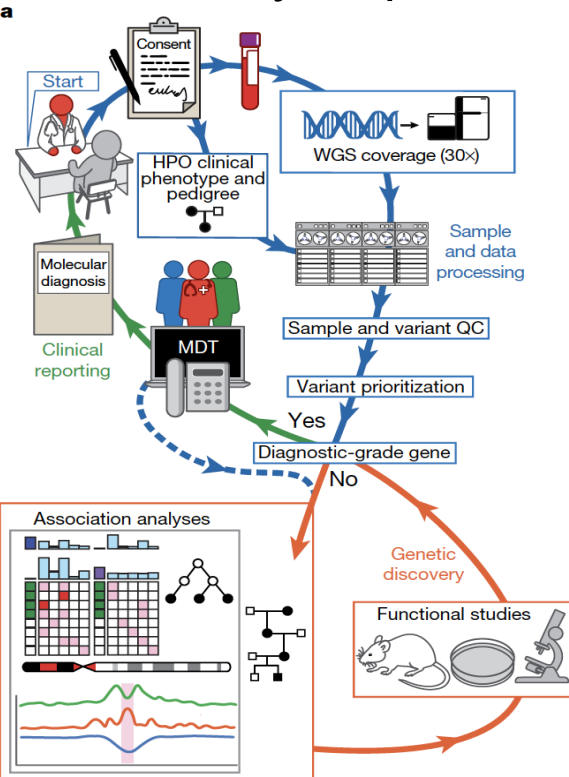
James H. R. Farmer, Mike L. Smith, & NIHR BioResource - Rare Dis

The 100,000 Genomes Project and beyond

NIHR | BioResource

'infinity loop'

88,597 participants with rare disease or cancer, and their families



120,239 whole
genomes
> 30x coverage

+



NHS
patient
records

+



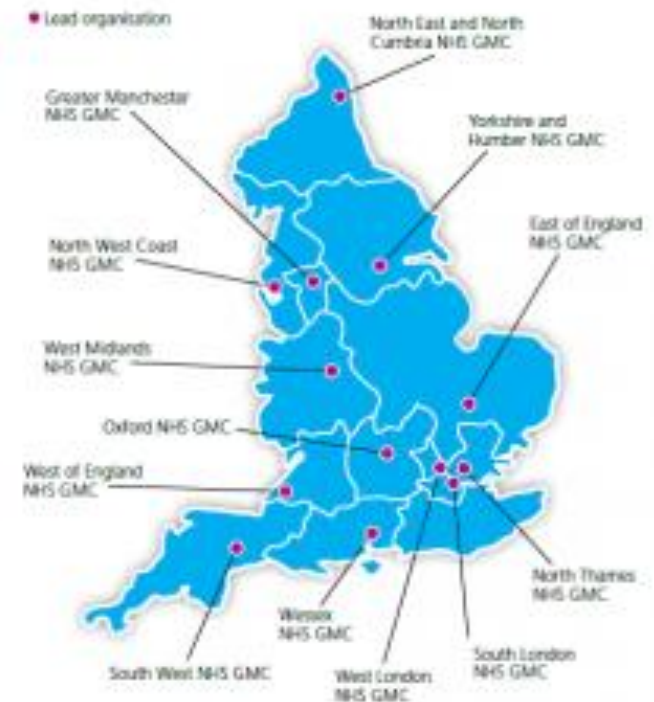
Deep
phenotyping
with HPO codes

Data from 100KG beginning to emerge:

- Numerous novel ultrarare disorders
- Genetic basis of Posterior Urethral Valves
- *IFT140* as a cause of mild ADPKD
- etc

**NHS Genomic
Medicine Centres**

Paving the way to personalised medicine



Basis for UK-wide clinical
Genomic Medicine Service

RaDaR: Rare Renal Disease Registry



- Established by the RA in 2010 and hosted by UKRR with funding from MRC/KRUK/BKPA



- ~100 disorders in 30 Rare Disease Groups (RDGs)
 - Clinicians – Research and clinical collaborations
 - Patients – Clinical provision, education, drive research
 - Scientists



- Aims to connect clinicians/scientists and patients and to facilitate clinical and academic research

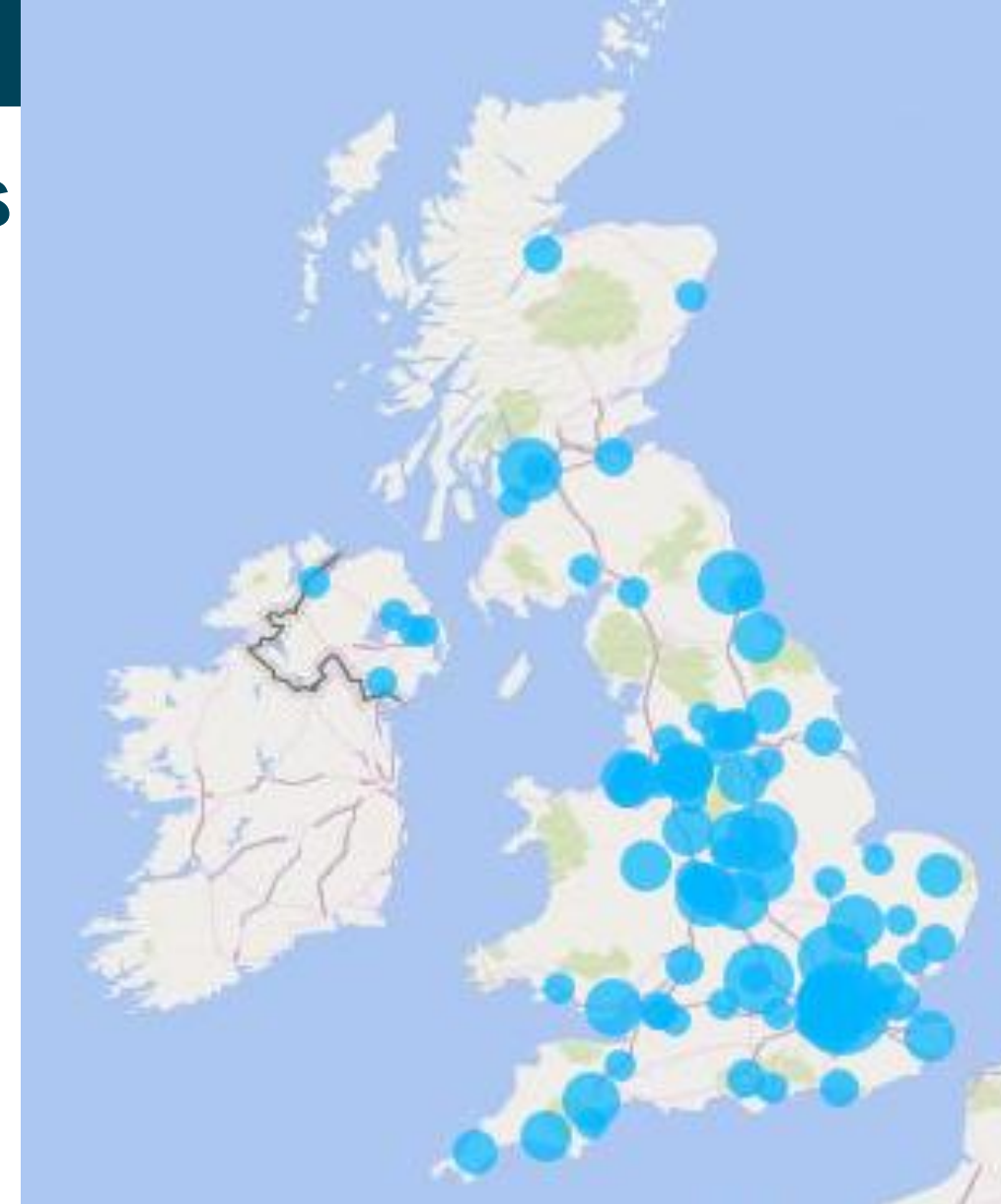
Components of RaDaR

- Comprises a support/clinical management network
 - <https://rarerenal.org>
 - Clinical guidelines and information for clinicians
 - Patient education, communication and details of patient meetings
 - Registry summary
 - Social media presence (@RenalRadar)
- Patient research register with clinical data
 - All participants give **written informed consent** for analysis of their clinical records and linkage to their data held in any other research or clinical database
 - Also agree to be contacted about studies they might be eligible for in the future



30,000 participants from 105 hospitals

Rare Disease Cohort	Cohort Size	Lab result Link
Alport Syndrome	901	711
APRT Deficiency	<10	<10
Atypical Haemolytic Uraemic Syndrome	282	221
Autosomal Dominant Polycystic Kidney Disease	7637	5840
Autosomal Dominant Tubulointerstitial Kidney Disease (FUAN)	209	157
Autosomal Recessive Polycystic Kidney Disease/Nephronophthisis	228	170
BK Nephropathy	51	42
Calciophylaxis	57	39
CKD due to Genetic Factors in people of African ancestry	64	33
Cystinosis	150	127
Cystinuria	466	307
Dent Disease and Lowe Syndrome	61	31
Fabry Disease	46	34
Fibromuscular Dysplasia	40	25
HNF1b Mutations	84	52
Hyperoxaluria	121	94
Idiopathic Nephrotic Syndrome	4106	3012
IgA Nephropathy	4134	3502
Inherited Renal Cancer Syndromes	99	<10
Membranoproliferative Glomerulonephritis / Dense Deposit Disease	1122	794
Membranous Nephropathy	2381	1614
Mitochondrial Renal Disease	<10	<10
Monoclonal Gammopathy of Renal Significance	173	121
Pregnancy	672	523
Pure Red Cell Aplasia	<10	<10
Retroperitoneal Fibrosis	141	91
STEC-associated HUS	169	88
Tuberous Sclerosis	240	129
Tubulopathy	352	193
Vasculitis	4621	2890



<https://ukkidney.org/rare-renal/metadata>

RaDaR data and the Renal Registry

- All data held and processed at the UK Renal Registry
 - Established 1995 to collect data from all UK RRT patients with Section 251 (NHS Act 2006) approval for patient data without consent
 - Aim is to assure quality and consistency of dialysis/transplantation
- RaDaR data is all from research-consenting patients so less stringent controls on use/access/linkage within this robust information governance framework
- UKRR database and informatics infrastructure
 - Automated datafeeds from all renal units in the country
 - Very good capture of initiation of RRT and death

Assembly of a large registry

- Broad, publicly available eligibility criteria for each RDG
- Small payment to each site for each subject recruited to study
 - National Institute of Health Research (NIHR) fund to support recruitment to ethically approved research at NHS sites
 - Sites incentivised to recruit but not resourced to input data
- Manual entry of (**very few**) mandatory fields at recruitment
 - Development of site-based mechanisms to enrich dataset by automated data transfer
 - Used existing UK Renal Registry Infrastructure to do this (esp lab results/medications)
 - **Avoid lengthy wish-list of clinical data entered manually by recruiting sites**

Automatic data feed

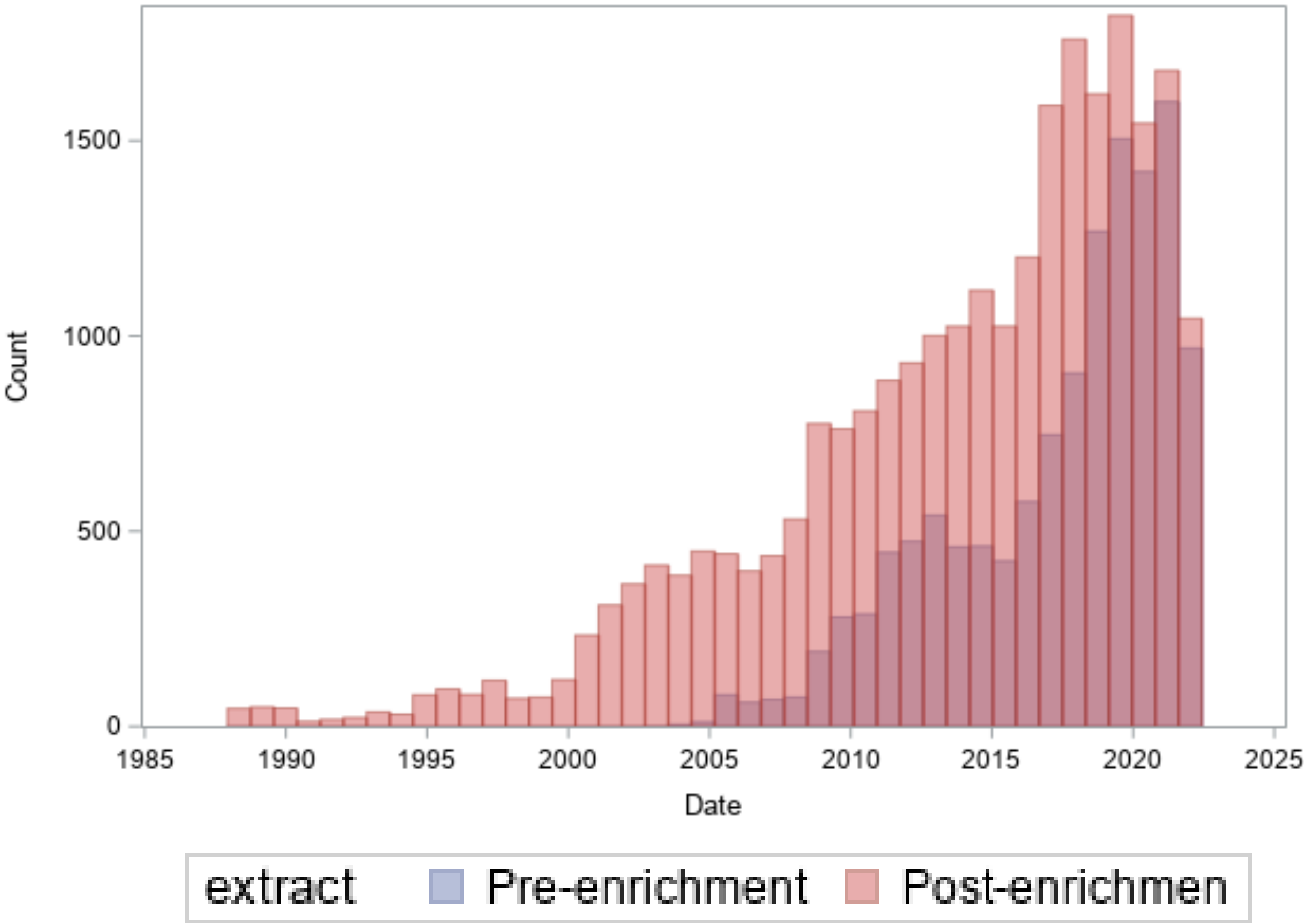


- RaDaR receives daily data feed from renal units
 - All or last 10 historic and all future values for each field
 - >60% of participants linked (increasing)
- Biochemistry and other laboratory results
- Clinic letters, imaging and other reports (variable)
- **This feed also goes to Patient View/Patients Know Best to allow participants to access their own data**

Data enrichment

With help of informatics staff at sites (especially those recruiting large numbers of patients) laboratory dataset was substantially enriched

Creatinine values at one site

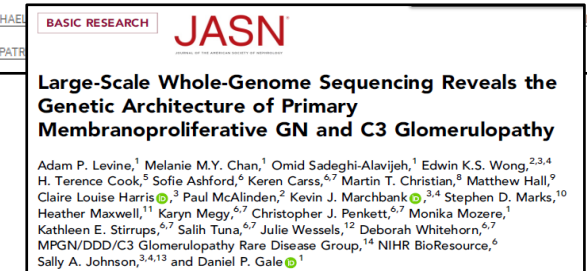


RaDaR data allows capture of large-scale clinical and renal outcome data

- Age at onset/presentation
- Presenting clinical features
- Medication use
- Rate of progression
- Outcomes including Death and initiation of Renal Replacement Therapy
- Secondary care use (with Hospital Episode Statistics data link)

Types of studies

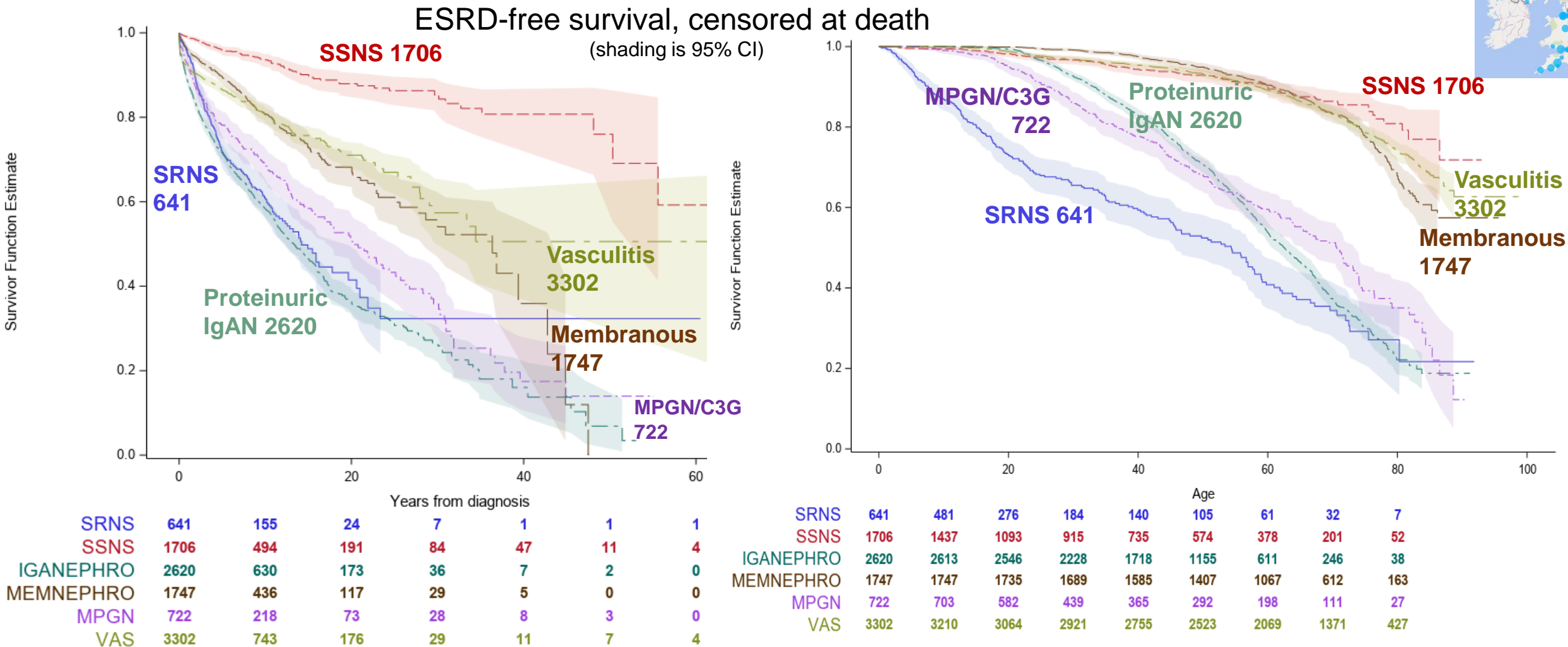
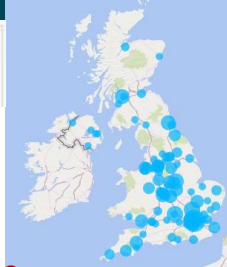
- Registry study
 - Epidemiology of disease in a defined area
- Genetic association studies
 - Collaboration with **NIHR | BioResource**
- Retrospective and prospective longitudinal Cohort studies: Natural History, outcomes and clinical practice
- Randomized Controlled Therapeutic Trials



RaDaR Natural History Analysis

- Ascertainment and data collection similar across all disorders so comparisons between them are possible
 - Prevalence/distribution
 - Treatments – effect and uptake
 - Outcomes and predictive factors
- Engagement with academic and commercial collaborators
 - In general individual patient-level data is not shared unless compelling scientific case for doing so
 - For commercial partners our statistician answers their questions using our data

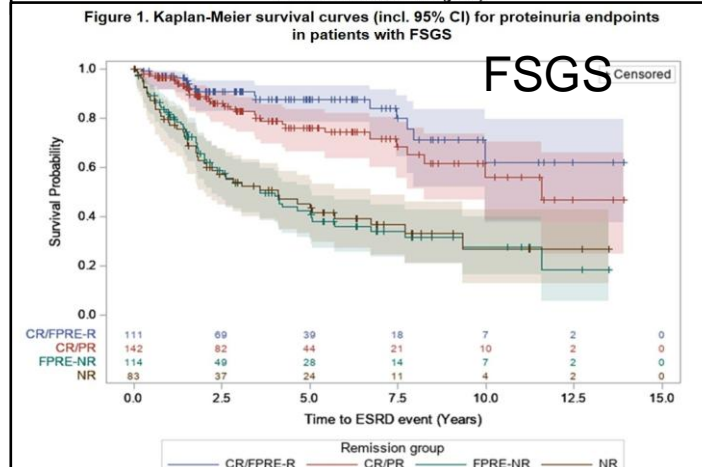
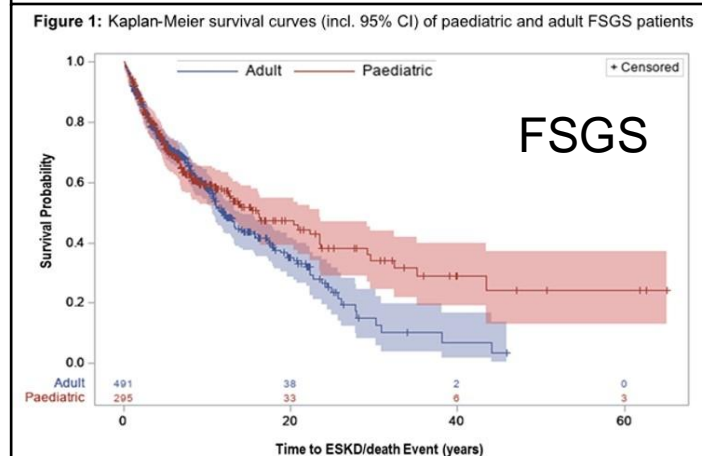
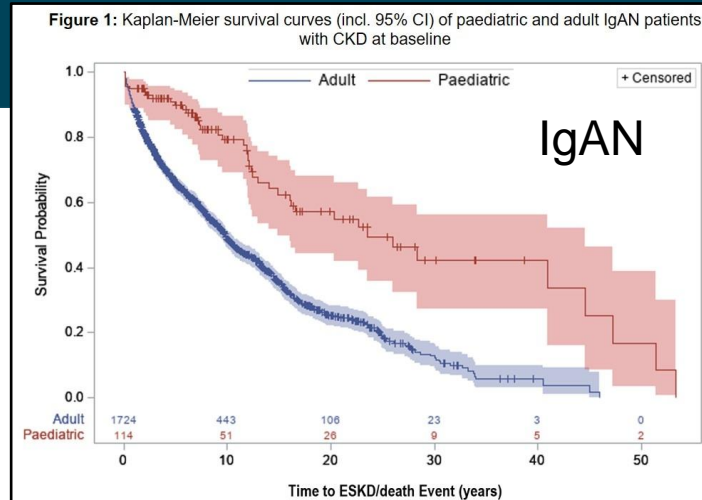
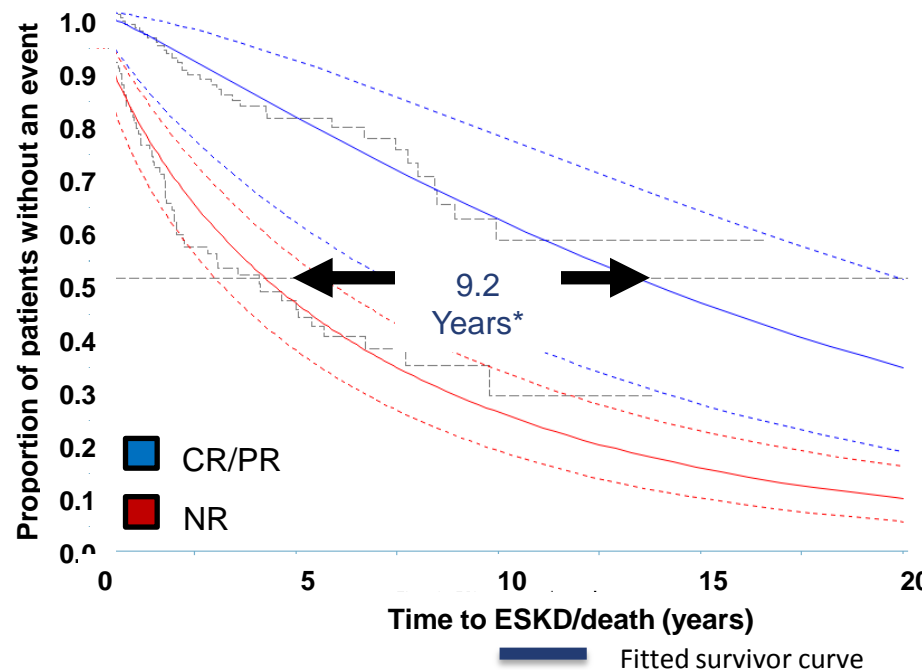
Renal outcomes – glomerular diseases



NB Ascertainment and survivor biases operate - recruitment 2010 onwards
RAS blockade/IS but not SGLT2i in common use over this timeframe

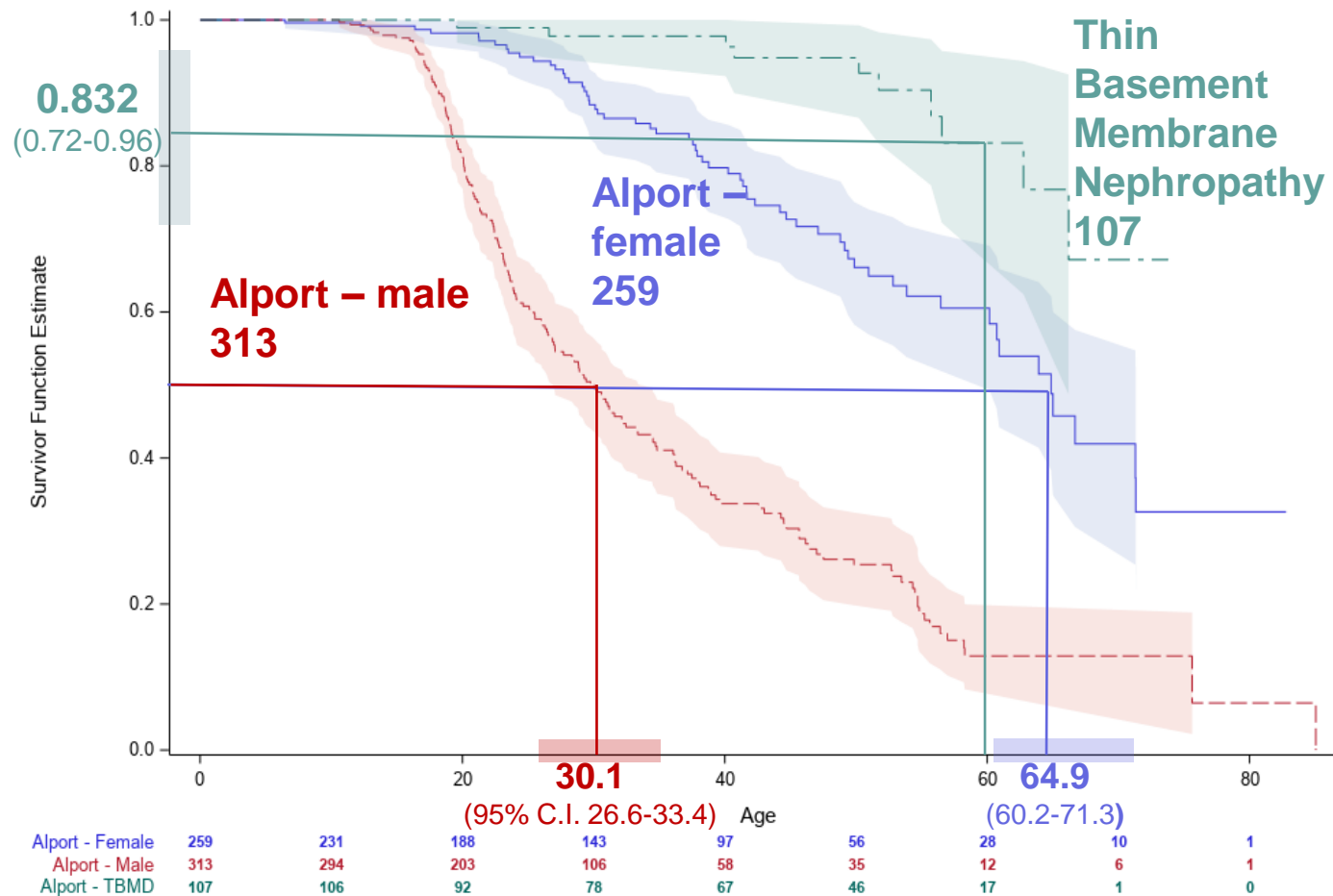
Commercial collaboration

- Looked at renal survival in children and adults with IgAN and NS with biopsy-proven FSGS
- Also looked at association of proteinuria response and renal survival in FSGS
 - Presented at ASN in 2021: PO1529, PO1530, PO1577



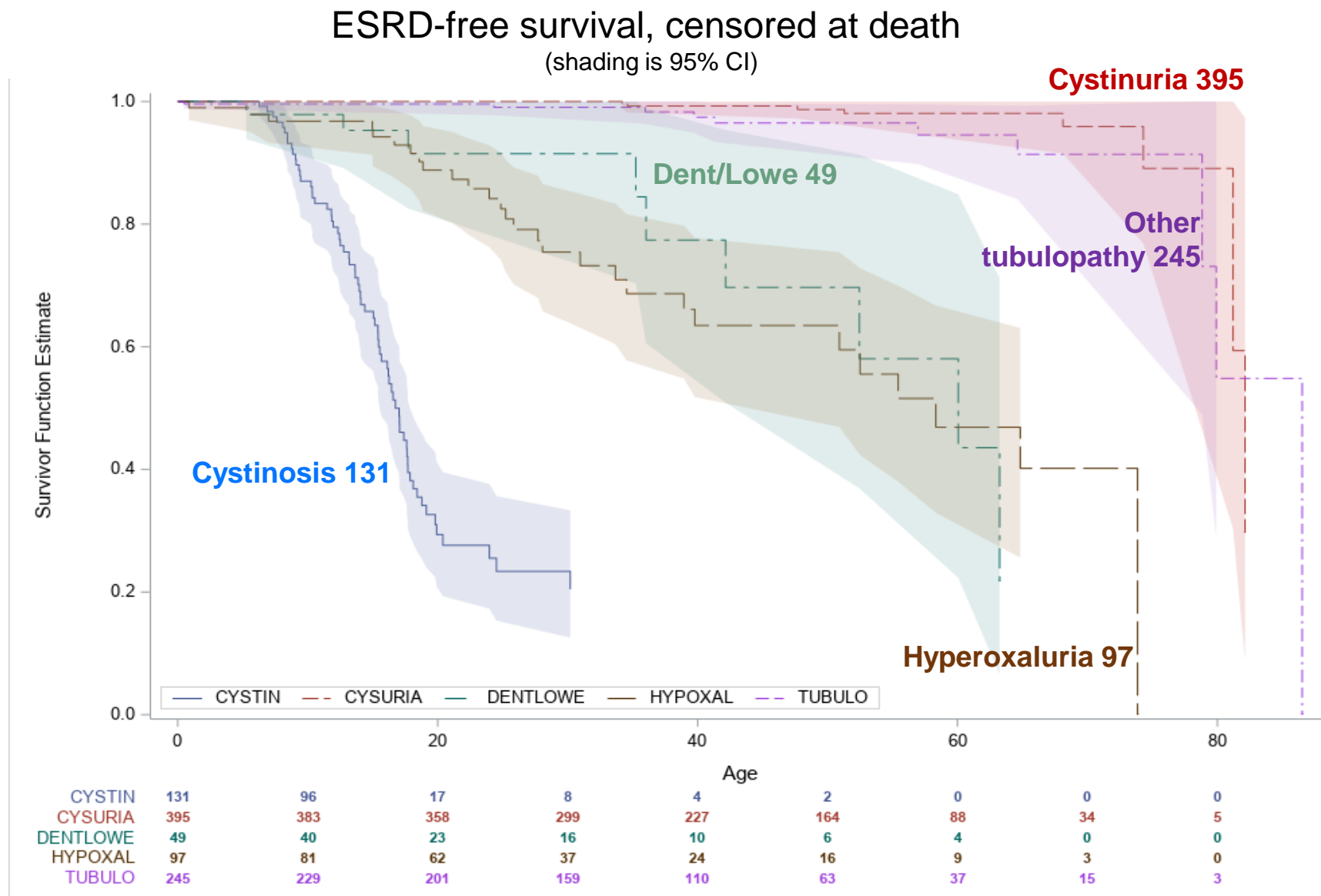
Alport Syndrome

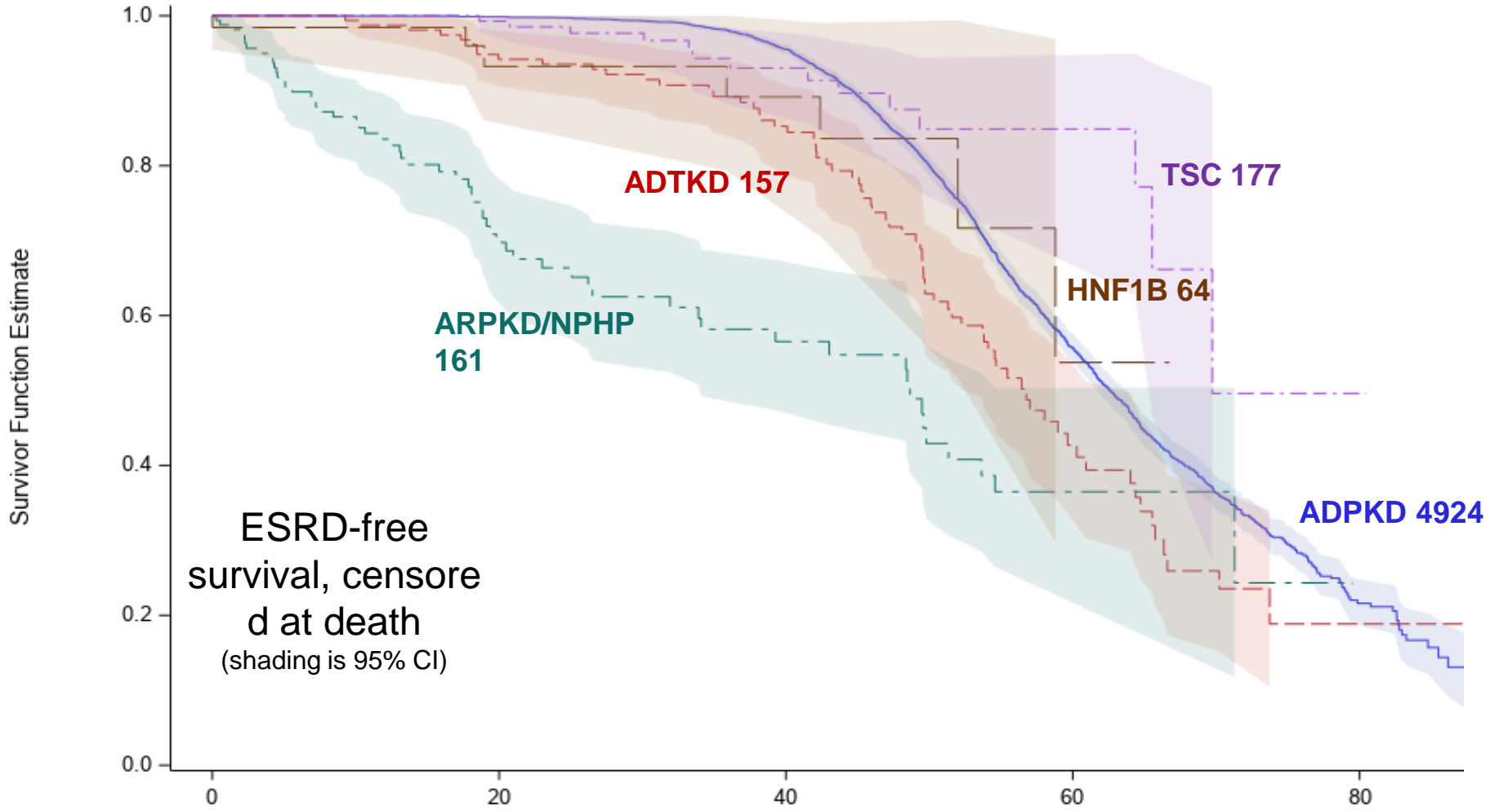
ESRD-free survival, censored at death
(shading is 95% CI)



- Median age at RRT in Alport Syndrome: 30.1(male) and 64.9 (female)
- For TBMN 83.2% have not required RRT before age 60

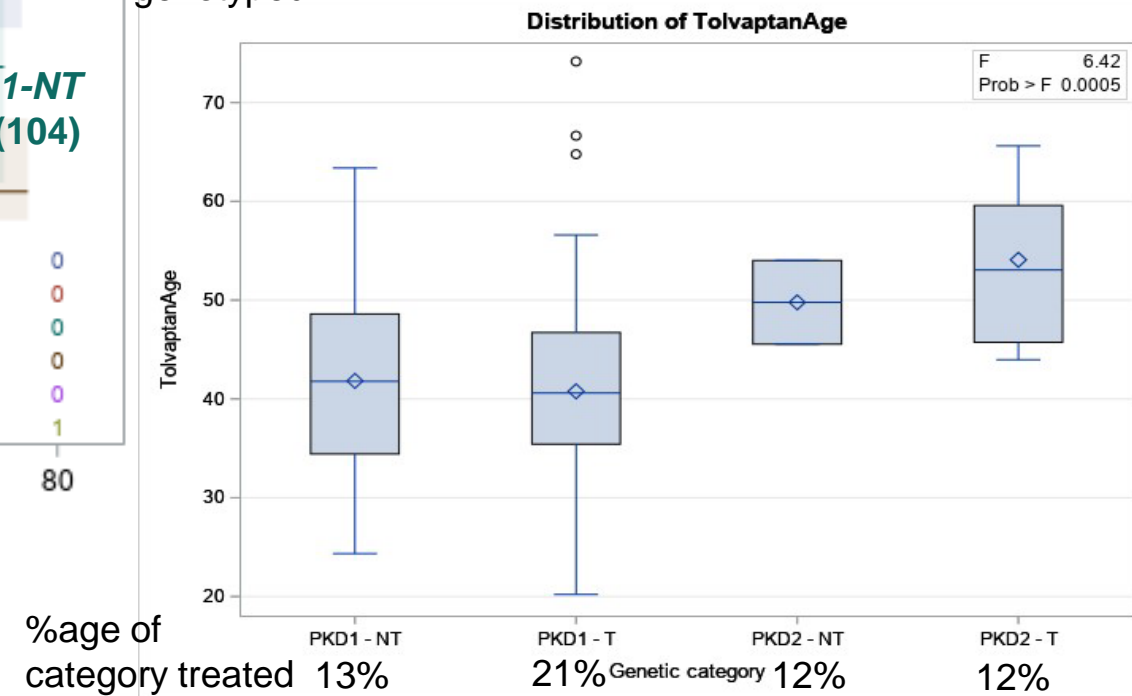
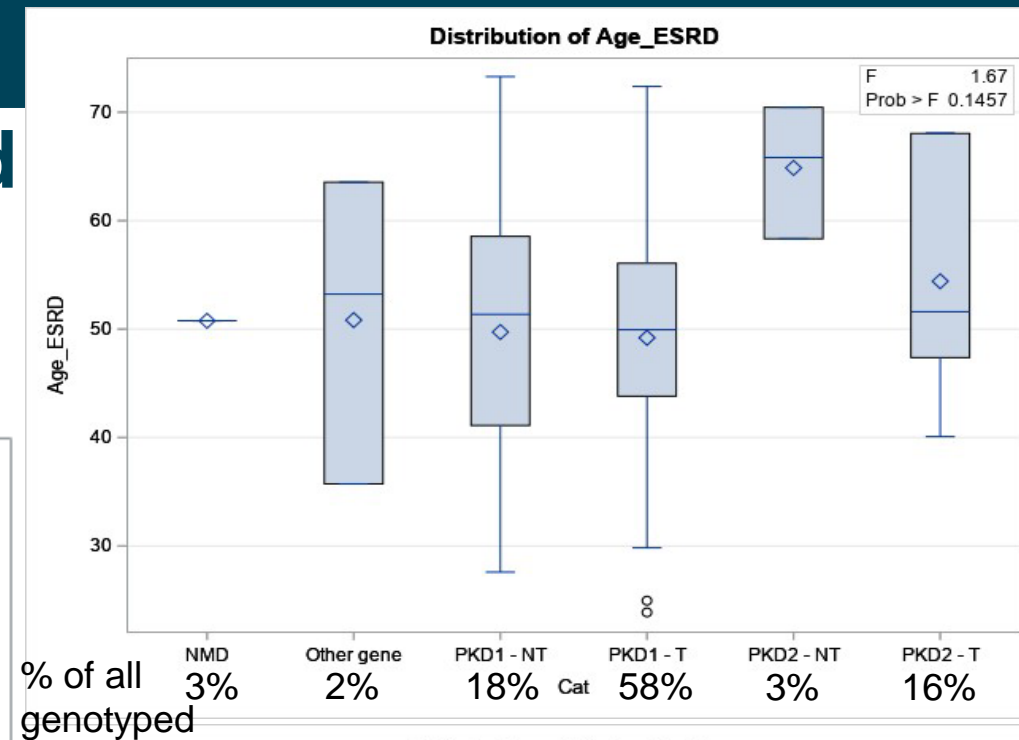
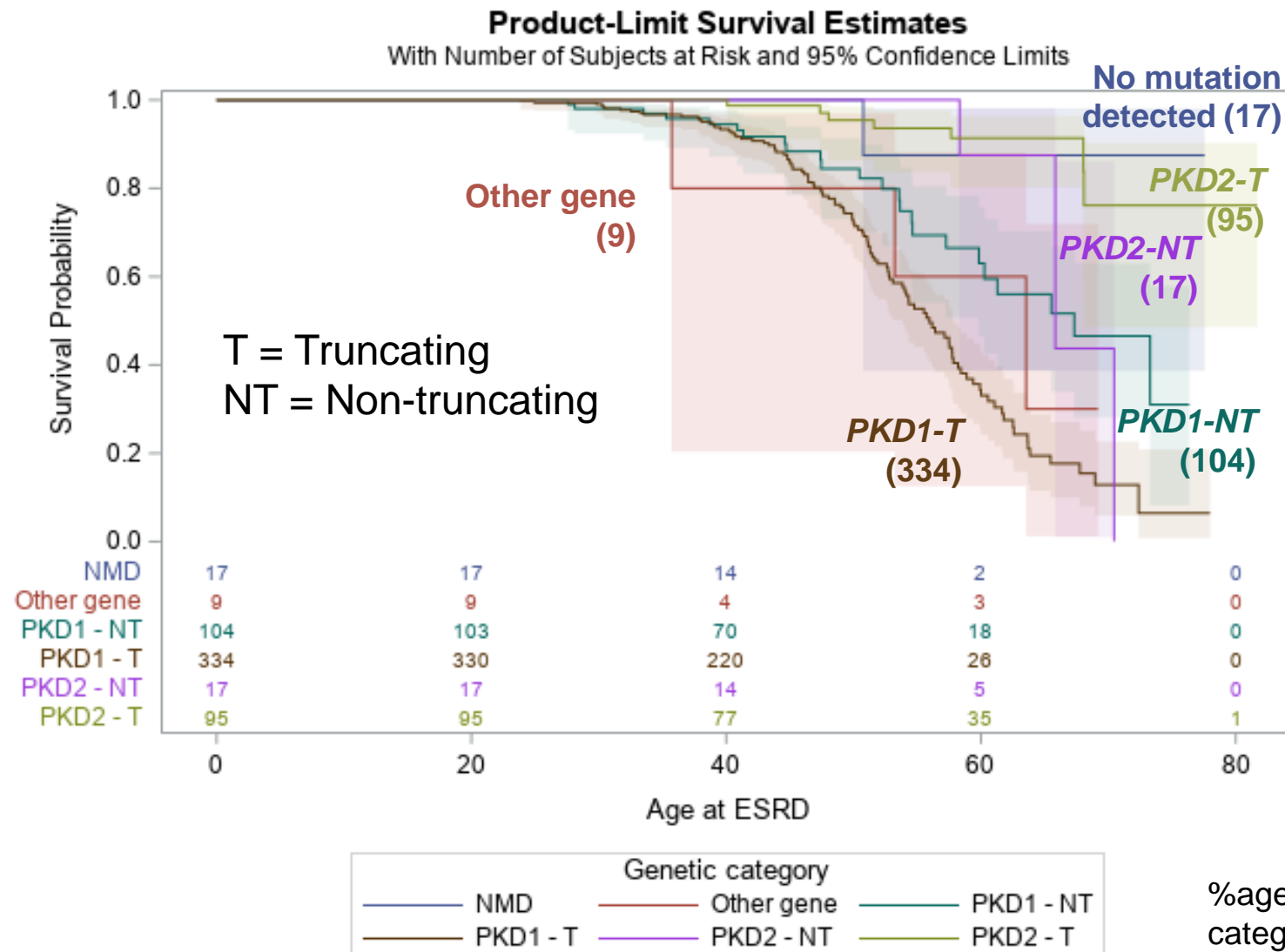
Ascertainment and survivor biases both possible





ADPKD	4924	4896	4774	4478	3681	2406	1081	382	54
ADTKD	157	156	145	131	107	62	27	11	1
ARPKD NPHP	161	120	64	47	32	21	9	3	0
HNF1B	64	49	32	26	20	9	3	0	0
TSC	177	164	131	99	59	31	14	3	1

ADPKD analyses: genotype, ESRD and treatment – work in progress



Rate of eGFR loss (ml/min/year) from first reading below 75 ml/min

Adult at time of diagnosis

Child at time of diagnosis

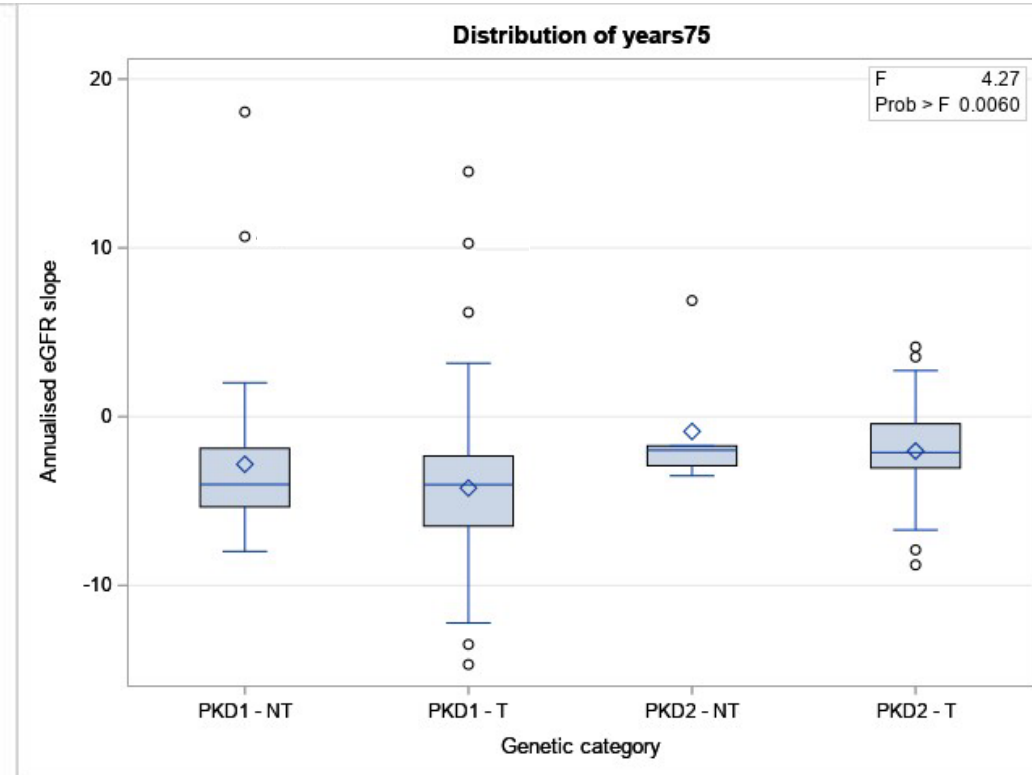
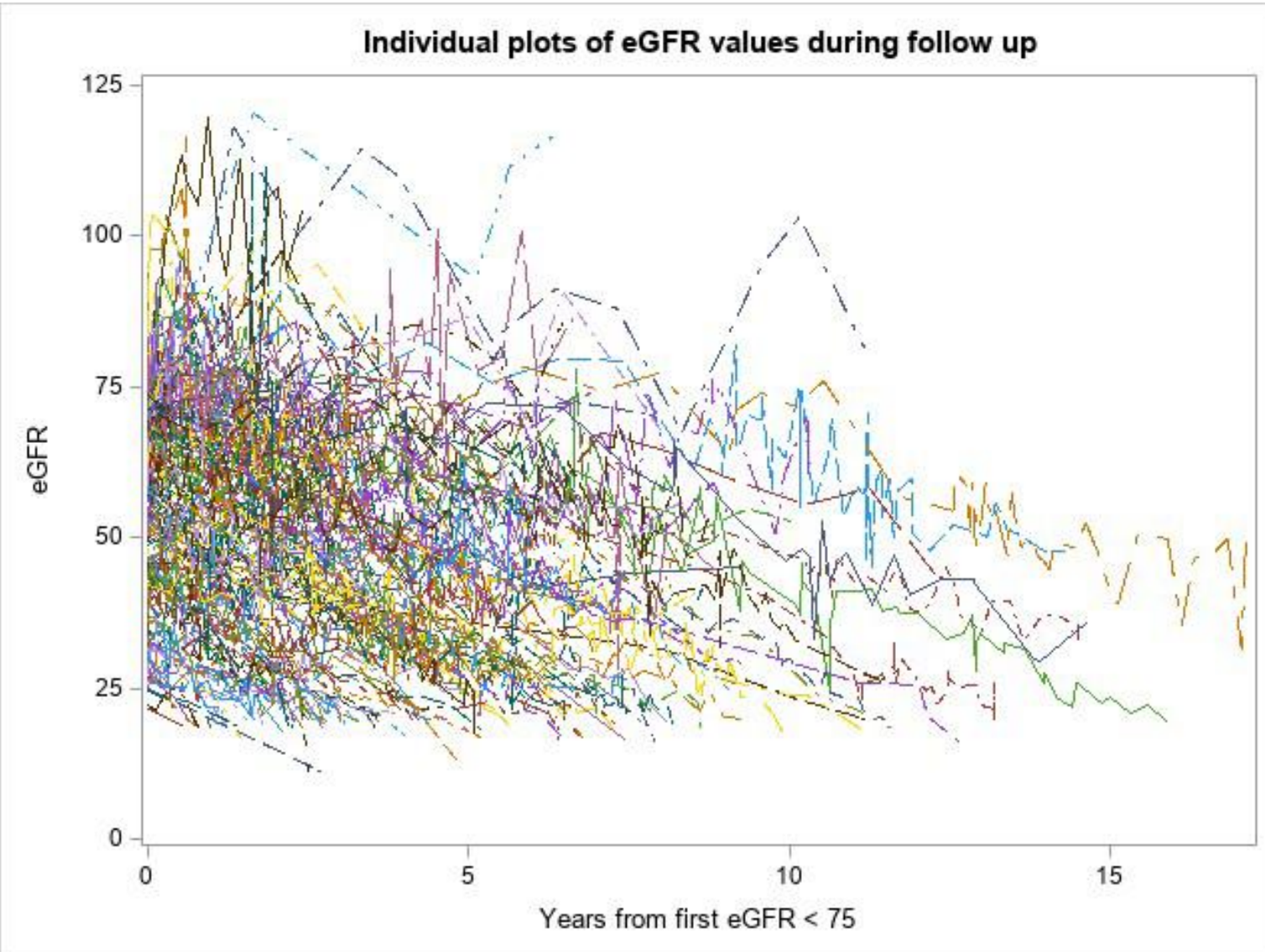
Disease	N	Median
Autosomal Dominant Polycystic Kidney Disease	1622	-3.62
IgA Nephropathy	931	-2.86
Vasculitis	929	-0.15
Membranous Nephropathy	725	-1.83
Steroid Sensitive Nephrotic Syndrome	384	-0.40
Steroid Resistant Nephrotic Syndrome	91	-2.99
MPGN/C3G	167	-2.50
Cystinuria	119	0.36
Alport Syndrome	107	-2.38
Pregnancy	98	-1.72
ADTKD	42	-2.81
Retroperitoneal Fibrosis	42	0.33
Monoclonal Gammopathy of Renal Significance	39	-5.81
ARPKD/Nephronophthisis	20	-2.08
Atypical Haemolytic Uraemic Syndrome	19	1.59
Tubulopathy	19	-0.71
HNF1b Mutations	14	-0.90
Tuberous Sclerosis	13	-1.49

Disease	N	Median
Steroid Resistant Nephrotic Syndrome	144	-11.6
Steroid Sensitive Nephrotic Syndrome	86	2.52
MPGN/C3G	59	-0.67
ADPKD	57	-3.17
Cystinosis	38	-4.82
ARPKD/Nephronophthisis	18	-1.51
Vasculitis	16	-2.01
IgA Nephropathy	14	-7.44
Alport Syndrome	12	-8.95
Tuberous Sclerosis	12	-1.98

NB In SHARP CKD study rate of eGFR loss was <2ml/min/year
(Haynes et al JASN 2014)

Unpublished RaDaR data

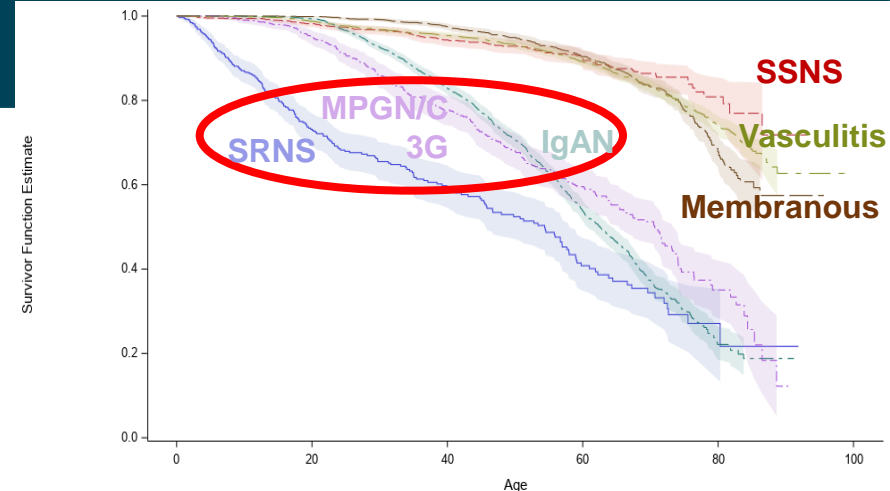
ADPKD: eGFR loss between 75 and 20 ml/min



Unpublished RaDaR data

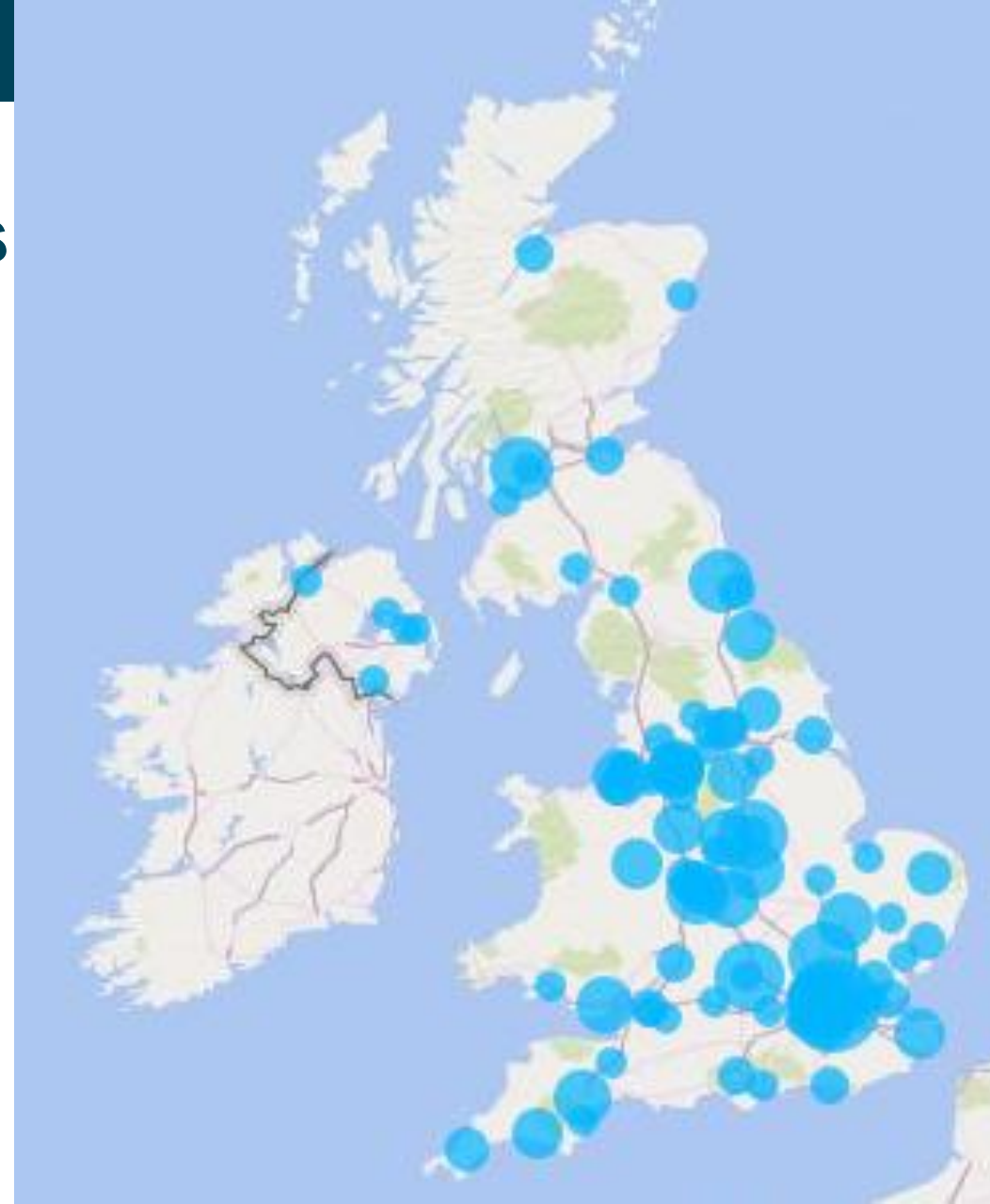
RaDaR can help design clinical trials

- Reveals diseases with particularly great unmet medical need
- Allows large-scale studies of natural history/outcomes
 - Where eGFR decline is rapid this might be a feasible trial endpoint
 - This information particularly important to regulators considering new therapies
- Identify number and proportion of RaDaR participants with particular genetic/phenotypic features, allowing feasibility of clinical trial to be assessed by biotech/pharma companies
 - Also allows estimation of eventual market size
 - Can reveal gaps in knowledge of natural history



RaDaR can help deliver clinical trials

- RaDaR holds geographical distributions
 - Integration with clinical data allows identification of sites with patients likely to be eligible for clinical trials
- RaDaR consent includes participants' permission to be contacted to inform them of trials that are relevant to them
 - Patients informed how to contact sites recruiting to studies they are potentially eligible for
 - This is an important motivator for many people to sign up to RaDaR
- Active collaborations to improve recruitment to interventional studies are in place and proving successful in enabling recruitment



RaDaR Summary

- Powerful source of real-world data about patients with rare renal diseases
- Linking RaDaR data with clinical/research genomic data can reveal robust associations in rare diseases
- Catalyst for clinical trials and therapeutic development
- Much more to do:
 - Genomic, multi-omic/biomarker and interventional studies
 - Linking with Hospital Episode Statistics and other data sources to enrich data on non-renal co-morbidities
 - Socio-economic deprivation associations
 - Ancestry, ethnicity and pharmacogenomic associations
 - International collaborations will be needed for insights into rarer disorders

Take home messages

- Studying rare diseases in an era when powerful molecular techniques (eg WGS) are widely available requires large, adequately controlled cohorts to gain important new insights
- Early and full engagement with patient groups has provided powerful momentum
- Assembling cohorts and collecting rich enough phenotypic dataset has required distributed recruitment and participation across disease areas and on a National scale
 - One day the scale required will be Continental, then Global
- Big data collection/processing and analysis presents its own challenges

<https://rarerenal.org>

@RenalRadar

Acknowledgements



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