

Present & Future Role of Molecular Genetic Diagnostics in ADPKD

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**KDIGO Controversies Conference on Autosomal
Dominant Polycystic Kidney Disease
(ADPKD)**

Edinburgh, Scotland

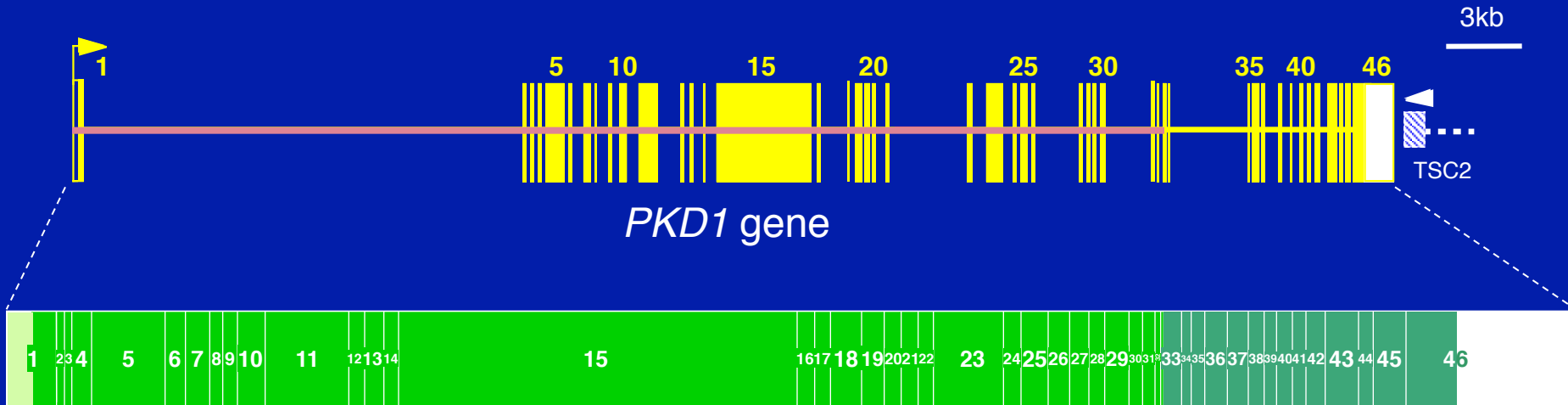
Friday, January 17, 2014

Disclosure of Interests

- Otsuka Pharmaceuticals: Research grant

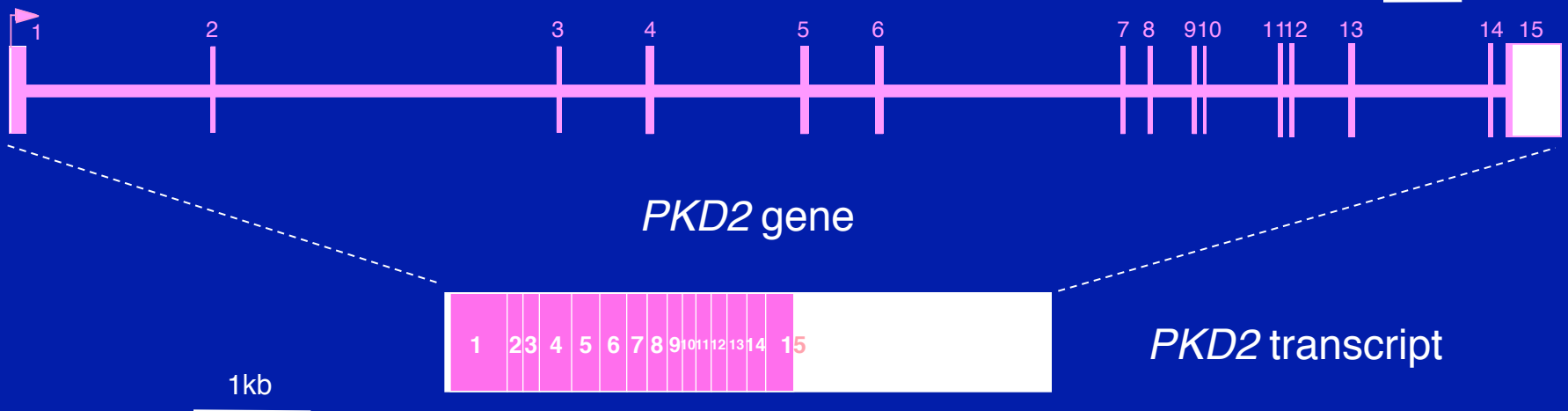
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Mutations to *PKD1* or *PKD2* cause ADPKD

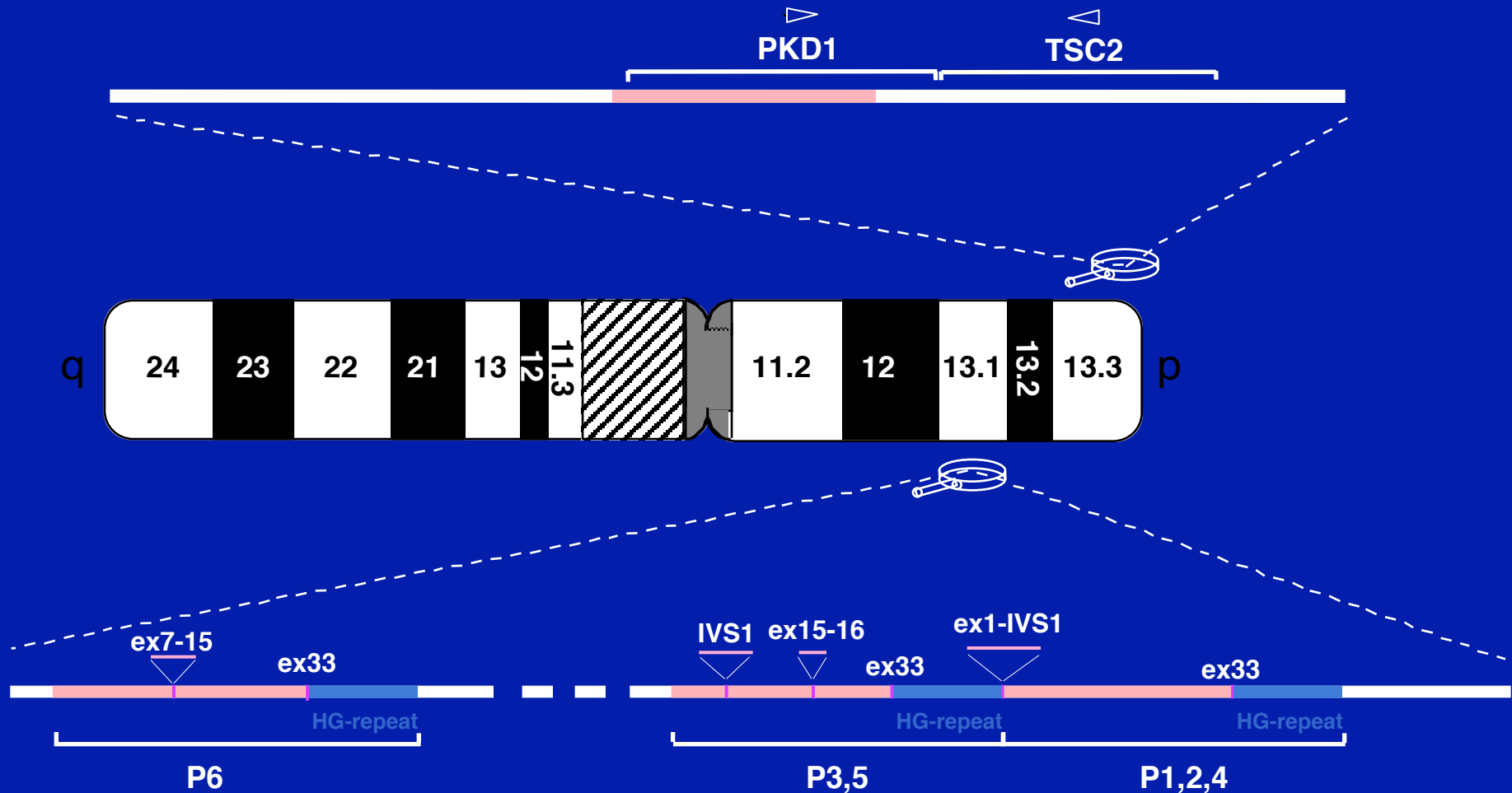


PKD1 ~85% families

PKD2 ~15% families



The *PKD1* region is duplicated in 16p13.1



6 *PKD1*-like pseudogenes; up to 99% identity to *PKD1*

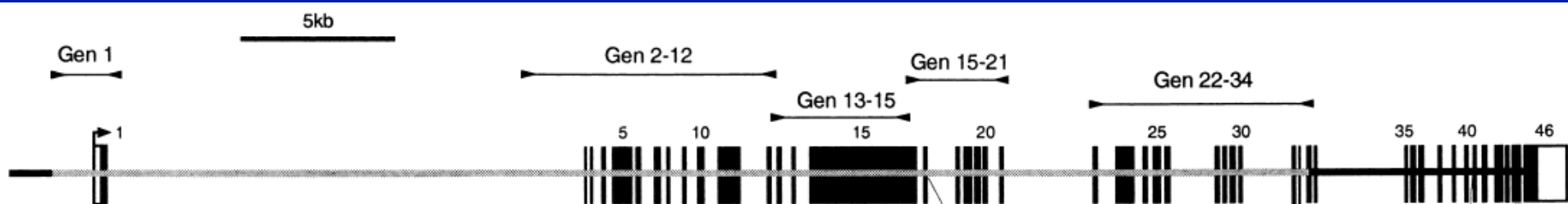
Sequence similarity between *PKD1* and the six pseudogenes

PKD1 exon23	201	ACCTGACCTCTGCCCTCATGCGCATCCTCA T GCGCTCCCGCGTGCTCAAC	250
PKD1P1	200	ACCTGACCTCTGCCCTCA C G C CATC G TCACGCGCTCCCGCGTGCTCAAC	249
PKD1P2	201	ACCTGACCTCTGCCCTCATGCGCATCCTCACGCGCTCCCGCGTGCTCAAC	250
PKD1P3	200	ACCTGACCTCTGCCCTCA C G C CATCCTCACGCGCTCCCGCGTGCTCAAC	249
PKD1P4	200	ACCTGACCTCTGCCCTCATGCGCATCCTCACGCGCTCCCGCGTGCTCAAC	249
PKD1P5	200	ACCTGACCTCTGCCCTCA C G C CATC G TCACGCGCTCCCGCGTGCTCAAC	249
PKD1P6	200	ACCTGACCTCTGCCCTCATGCGCATCCTCACGCGCTCCCGCGTGCTCAAC	249

PKD1 exon23	251	GAGGAGCCCCTGACGCTGGCGGGCGAGGAGATCGTGGCCCAAGGGCAAGCG	300
PKD1P1	250	GAGGAGCCCCTGACGCTGGCGGG T GAGGAGATCGTGGCCCAAGGGCAAGCG	299
PKD1P2	251	GAGGAGCCC G TGACGCTGGCGGGCGAGGAGATC A TGGCCCAAGGGCAAGCG	300
PKD1P3	250	GAGGAGCCCCTGACGCTGGCGGG T GAGGAGATCGTGGCCCAAGGGCAAGCG	299
PKD1P4	250	GAGGAGCCC G TGACGCTGGCGGGCGAGGAGATC A TGGCCCAAGGGCAAGCG	299
PKD1P5	250	GAGGAGCCCCTGACGCTGGCGGG T GAGGAGATCGTGGCCCAAGGGCAAGCG	299
PKD1P6	250	GAGGAGCCCCTGAC A CTGGCGGGCGAGGAGATCGTGGCCCAAGGGCAAGCG	299

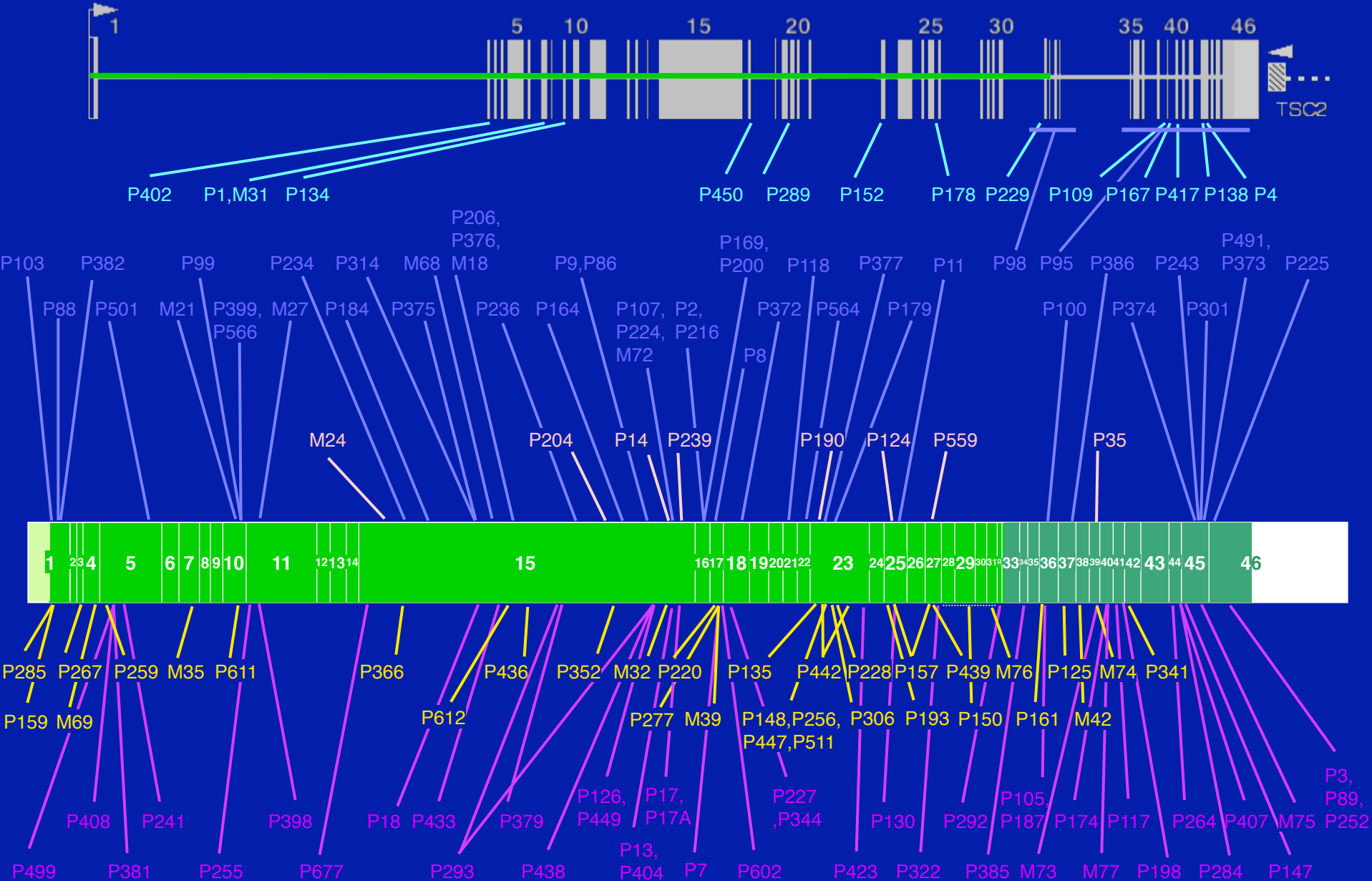
PKD1 exon23	301	CTCGGACCCGCGGAGCCTGCTGTGCTATGGCGGGCGCCCCAGGGCCTGGCT	350
PKD1P1	300	CTCGGACCCGCGGAGCCTGCTGTGCTATGGCGGGCGCCCCAGGGCCTGGCT	349
PKD1P2	301	CTCGGACCCGCGGAGCCTGCTGTGCTATGGCGGGCGCCCCAGGGCCTGGCT	350
PKD1P3	300	CTCGGACCCGCGGAGCCTGCTGTGCTATGGCGGGCGCCCCAGGGCCTGGCT	349
PKD1P4	300	CTCGGACCCGCGGAGCCTGCTGTGCTATGGCGGGCGCCCCAGGGCCTGGCT	349
PKD1P5	300	CTCGGACCCGCGGAGCCTGCTGTGCTATGGCGGGCGCCCCAGGGCCTGGCT	349
PKD1P6	300	CTCGGACCCGCGGAGCCTGCTGTGCTATGGCGGGCGCCCCAGGGCCTGGCT	349

Sanger screening protocol for ADPKD



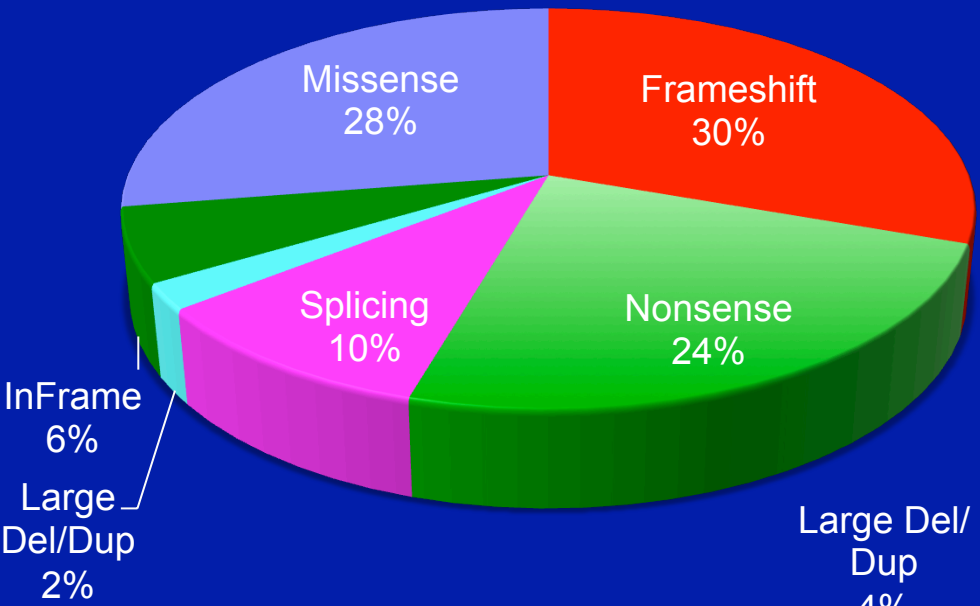
- 5 locus specific amplicons cover the duplicated part of *PKD1*
- *PKD1* exons and flanking intronic regions: 46 amplicons
- PKD2 exons and flanking intronic regions: 17 amplicons
 - Total 63 amplicons

High level of allelic heterogeneity in PKD1



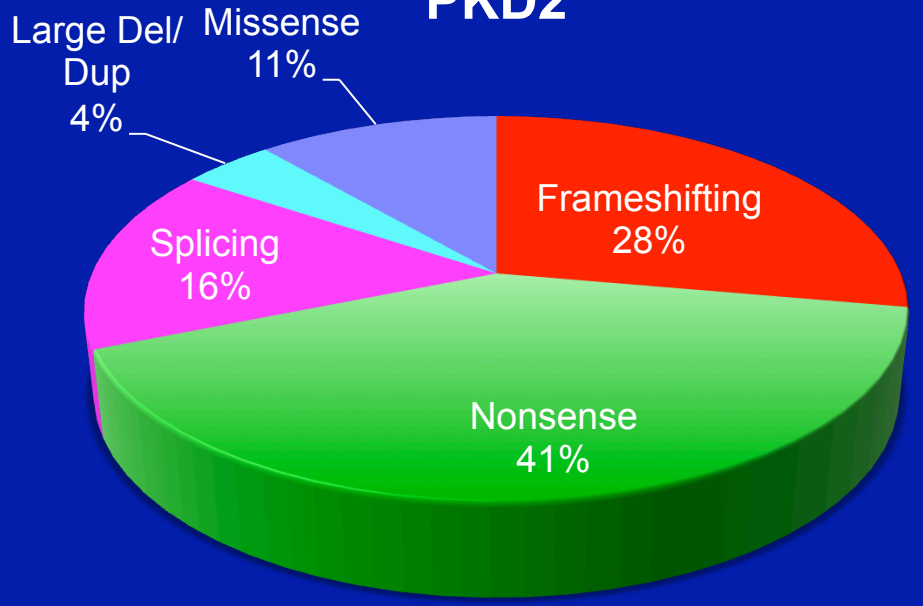
Mutation Types in ADPKD: HALT PKD population

PKD1



No mutation identified in ~8% families

PKD2



Non-definite mutations in 34% PKD1 and 11% PKD2

Update to ADPKD mutation database (PKDB): Version 2.95 <http://pkdb.mayo.edu>

- **Total 2544 variants**
 - **PKD1 = 2293** **PKD2 = 251**
- **1425 Likely Pathogenic mutations (2020 families)**
 - **PKD1 = 1250 (1677 families)**
 - **PKD2 = 175 (343 families)**
- **189 Indeterminate change**
 - **PKD1 = 181; PKD2 = 16**
- **867 Neutral Polymorphisms**
 - **PKD1 = 911 (94%); PKD2 = 58 (6%)**

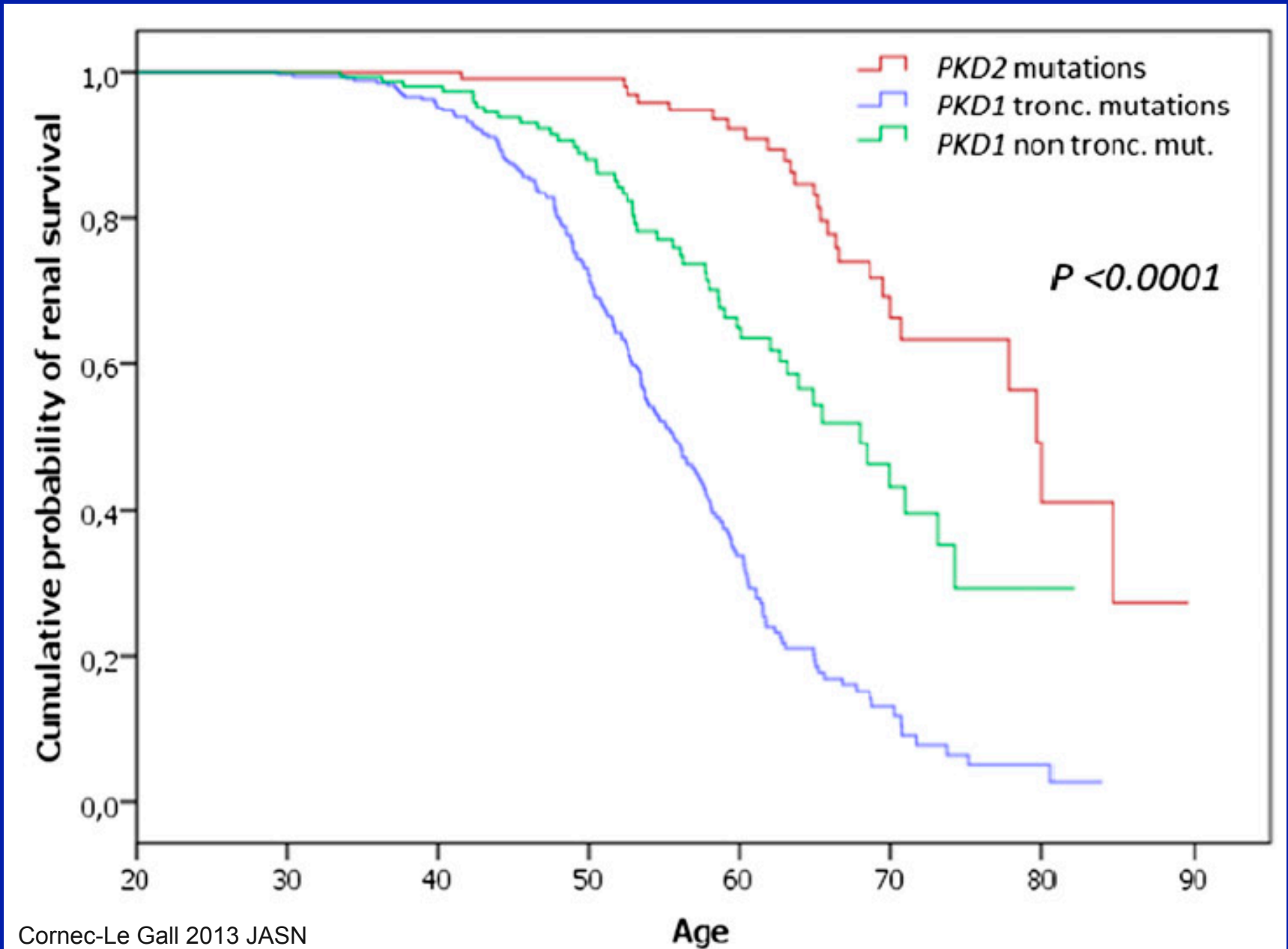
Diagnostics of ADPKD usually performed by renal imaging

- Imaging methods can usually accurately diagnose ADPKD in adults
- Imaging diagnostics less reliable in younger adults
 - Especially in families with less severe disease
 - *PKD2* and hypomorphic *PKD1* mutations

Gene-based diagnostics for ADPKD

- Genetic testing may be helpful when imaging results are equivocal and firm diagnosis required
 - Living related donors
 - Confirming negative diagnosis in young potential donor when imaging results may be unreliable
 - Clarifying the diagnosis in a potential donor when 1 or 2 cysts detected by imaging
- Individuals with a negative family history and/or an unusual disease presentation: to clarify the diagnosis
 - Early onset ADPKD
 - Mild PKD
 - Atypical radiological presentation
- Once therapies available: testing of young patients to obtain a firm diagnosis before starting treatment

Knowing the gene and the mutation are of prognostic value in ADPKD



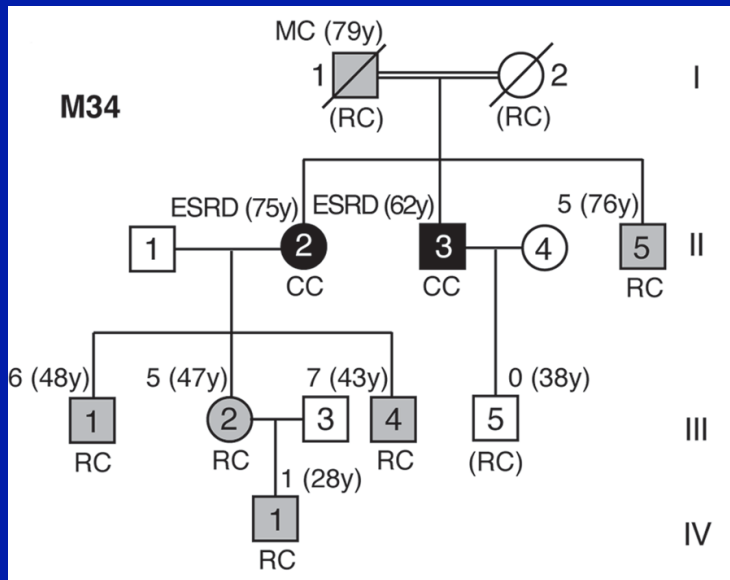
Gene-based prognostics in ADPKD

- Gene and mutation data can provide information about the severity of disease
 - Truncating *PKD1* mutations associated with more severe disease
 - *PKD2* mutations associated with milder disease
 - Hypomorphic *PKD1* mutations associated with milder disease
- Prognostic information can help patient management
 - Planning for ESRD
 - Provide reassurance for those with predicted less severe course
 - Select patients for clinical trials
 - Select patients treatments

Resolve complex ADPKD cases

- Negative family history
 - Determine if disease is ADPKD
 - Especially with mild disease
 - *De novo* mutation
 - Reduced risk in sibs
 - Mosaicism – can complicate risk prediction
- Early onset cases
 - Some due to combinations of ADPKD alleles
 - Of value for avoiding further early onset cases in a family
- Marked intrafamilial variation
 - Allelic/genic combinations
 - Mosaicism

Hypomorphic *PKD1* allele in homozygosity: extreme intrafamilial phenotypic variability



PC1 **C**

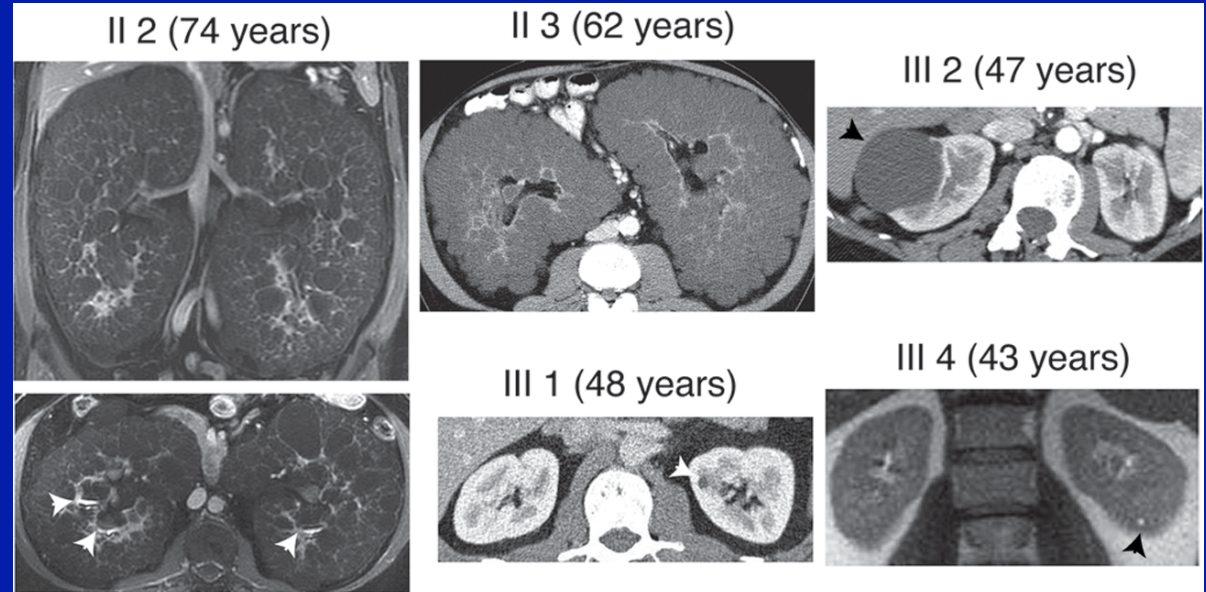
human	3267	W	D	R	P	P	R	S	R	F	T	R	I	Q	R	A	T	C	C	V	L	L	3287
mouse	3259	W	D	R	P	P	R	S	R	F	T	R	V	Q	R	V	T	C	C	V	L	L	3279
rat	3258	W	D	R	P	P	R	S	R	F	T	R	V	Q	R	V	T	C	C	V	L	L	3278
chicken	3276	W	D	R	P	P	R	S	R	F	T	R	V	Q	R	A	T	C	C	S	L	L	3296
xenopus	3247	W	D	R	P	P	R	S	R	F	T	R	V	Q	R	A	T	C	C	A	L	L	3267
fugu	3407	F	Q	R	P	P	R	S	P	F	T	R	L	Q	R	A	T	C	C	A	L	L	3427

PC1-like **C**

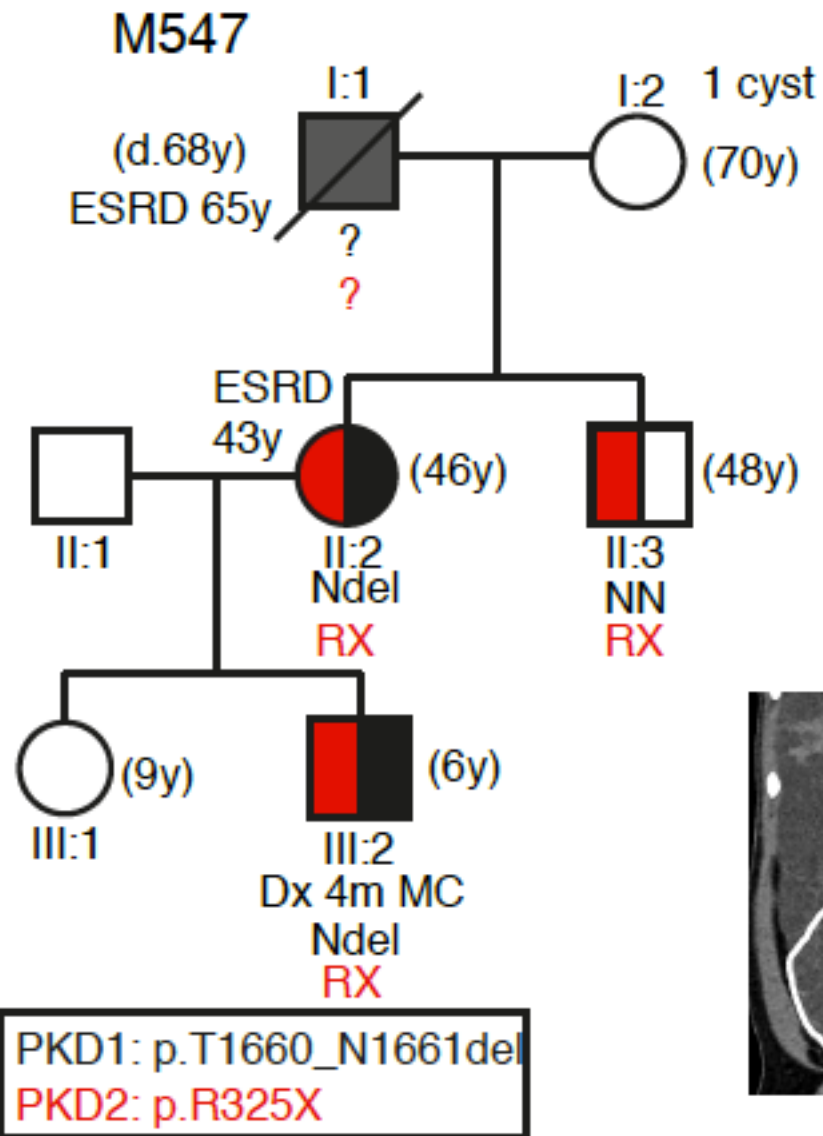
PC1	3267	W	D	R	P	P	R	S	R	F	T	R	I	Q	R	A	T	C	C	V	L	L	3287
PKDREJ	1376	F	A	S	V	V	A	K	T	F	N	R	L	Q	R	L	S	C	C	L	A	M	1396
PC1L1	1943	Y	S	R	P	S	S	S	R	Y	L	H	T	P	R	L	T	V	S	F	S	L	1963
PC1L2	1539	F	S	R	C	A	R	S	S	F	T	R	V	Q	R	V	S	C	C	F	S	L	1559
PC1L3	892	A	T	R	H	P	W	N	Q	F	T	R	V	Q	R	L	S	C	C	M	T	L	912
SpREJ4	1825	F	T	R	P	P	N	S	N	F	T	R	L	Q	R	I	T	C	C	F	T	L	1845
SpREJ5	2785	F	A	R	P	A	R	S	T	F	T	R	C	Q	R	A	L	C	C	L	S	L	2805
SpREJ7	3140	F	T	R	P	P	Y	S	I	F	T	R	V	Q	R	L	T	C	C	L	C	I	3160

Consanguineous US family of French origin homozygous or heterozygous for R3277C

Unusual, reniform kidneys with multiple small cysts in homozygotes

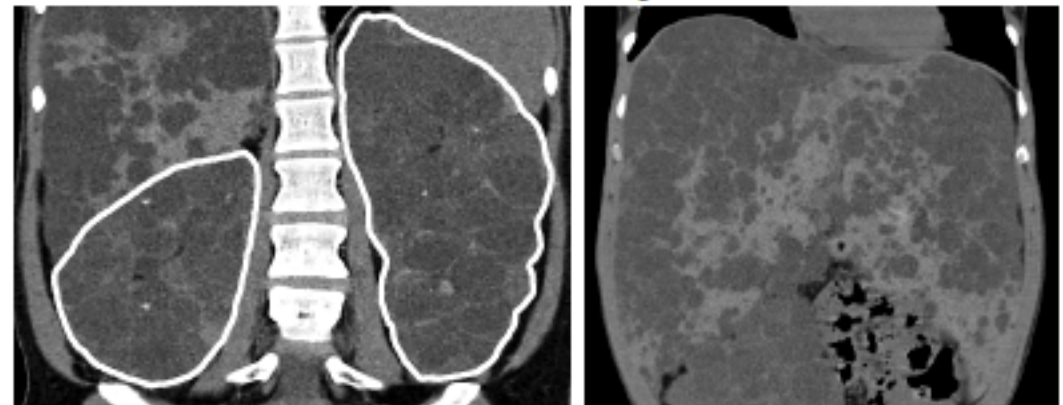


Family with *PKD1* and *PKD2* mutation: intrafamilial phenotypic variability



Patients with both mutations have more severe disease

II:2 42y

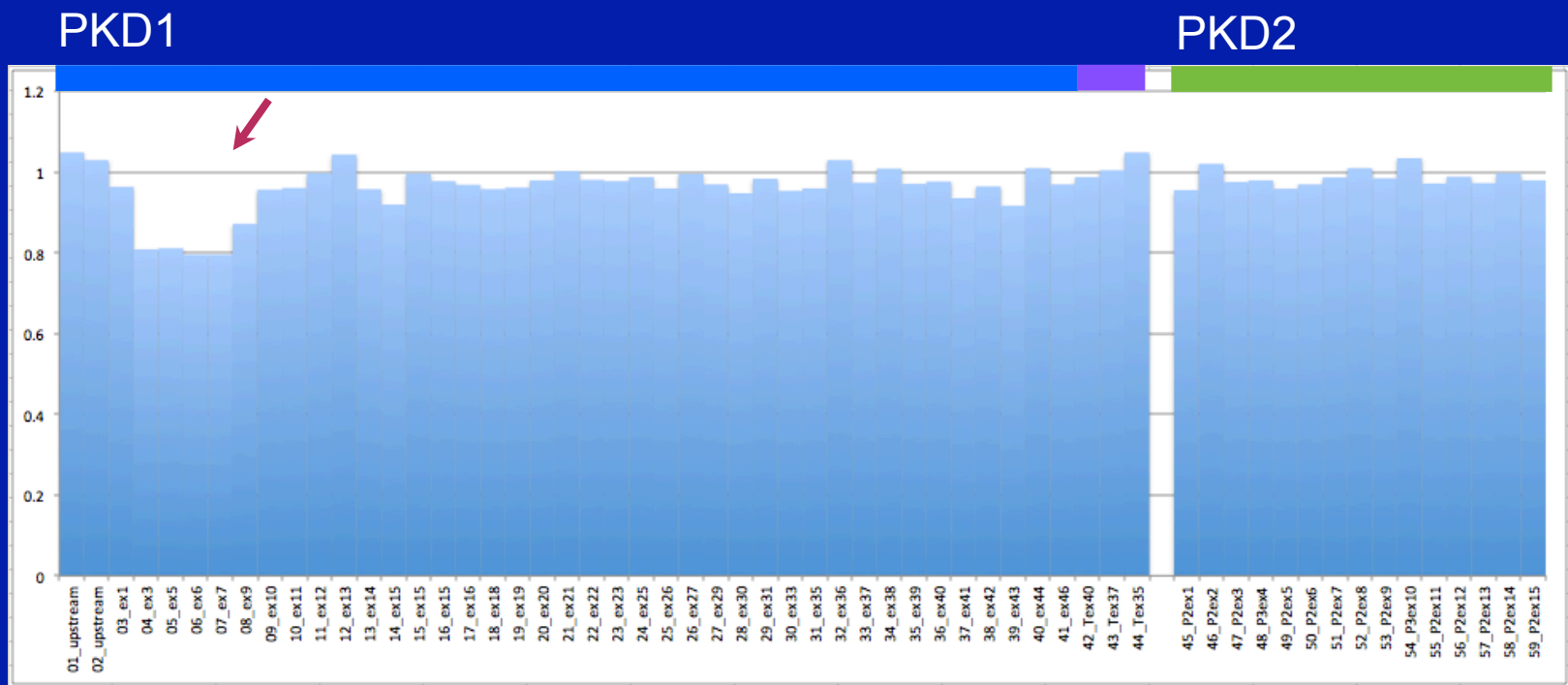


Kidney

Liver

MLPA assay for *PKD1* and *PKD2*

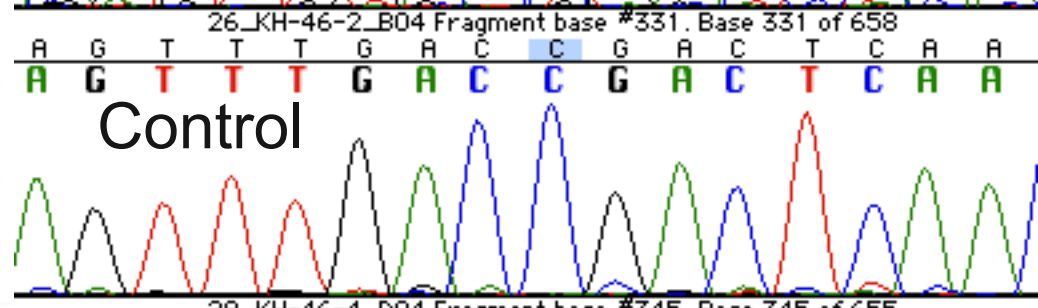
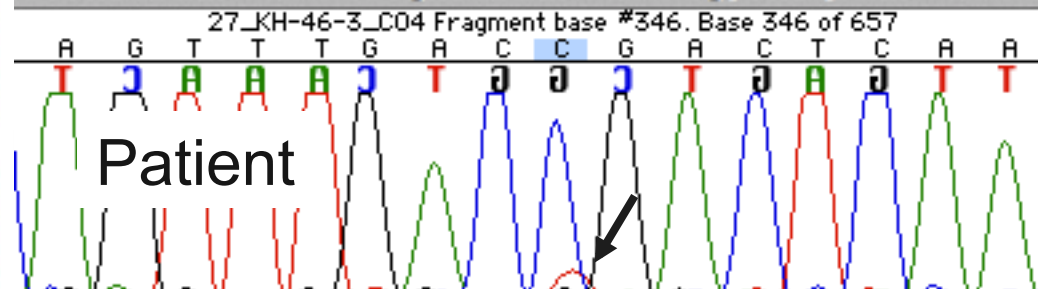
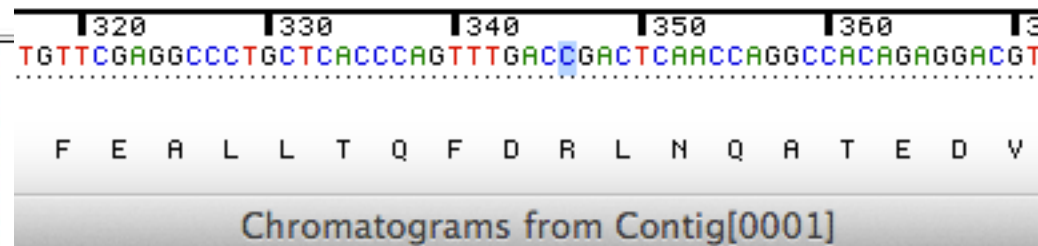
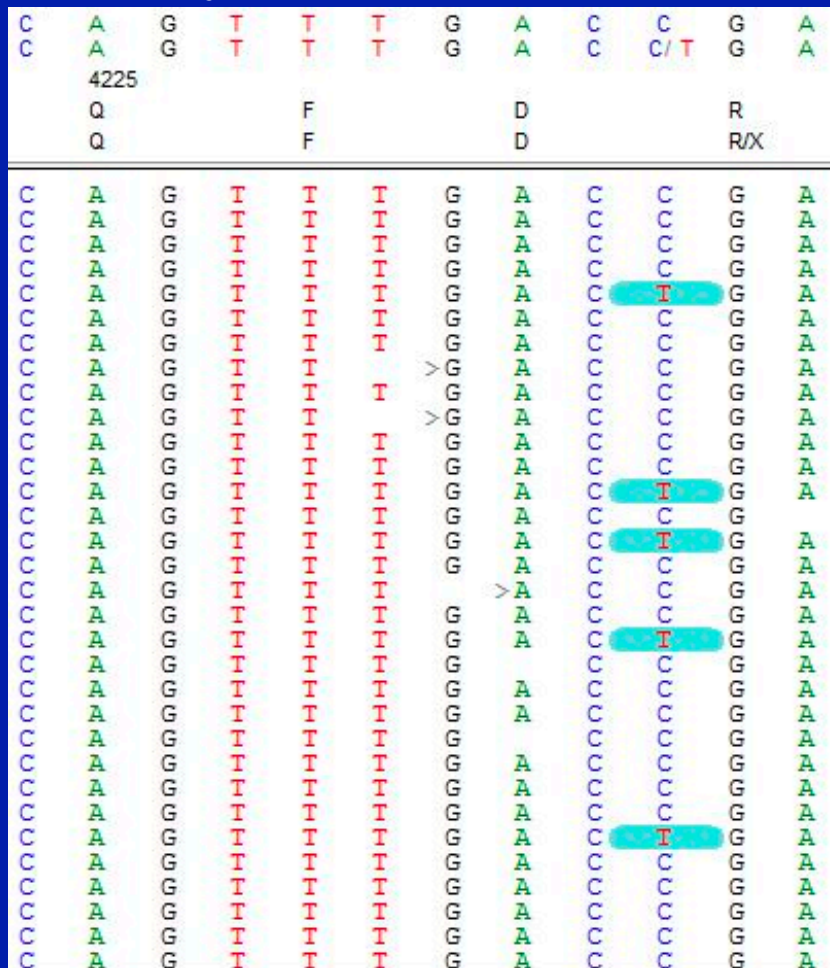
- 3-4% of mutations are large rearrangements
- PKD1 deletion of exons 3-9, 40% mosaic



Screening for mosaics employing next generation sequencing

ADPKD patient with mild disease (S.Cr. 1.9 at 77y) and a negative family history was mutation negative by Sanger sequencing

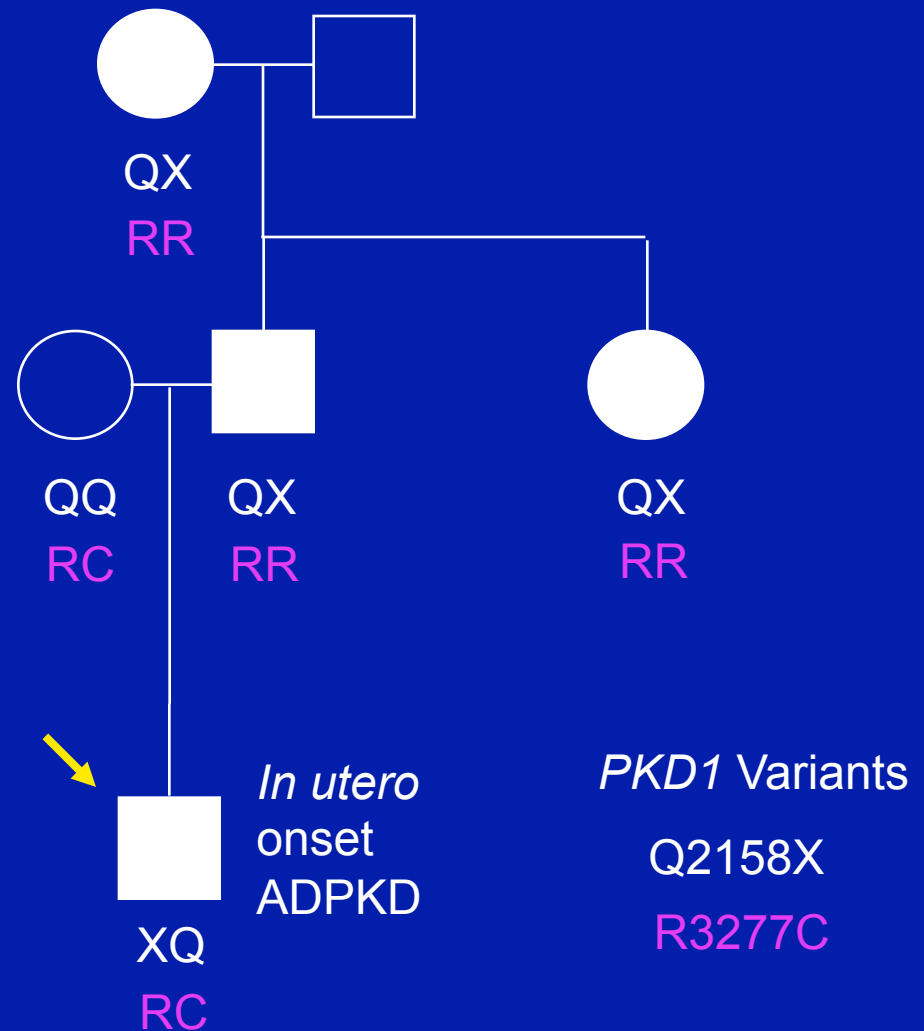
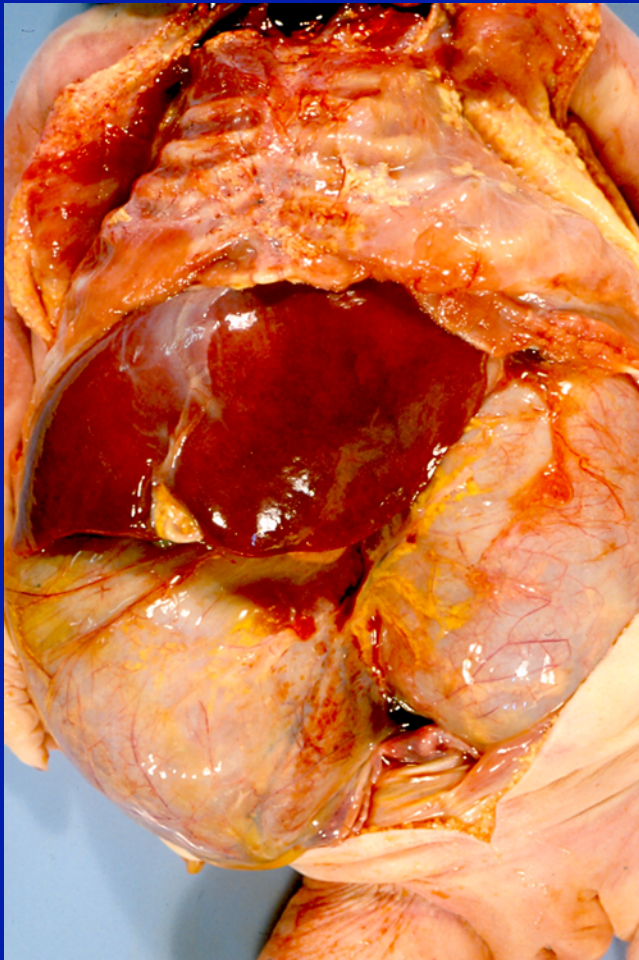
NGS analysis identified the nonsense mutation *PKD1*: p.R4228X at a low level



In utero onset ADPKD

- Rarely (<1%) ADPKD presents *in utero* with enlarged and echogenic kidneys in a family with otherwise typical ADPKD
- These cases can be confused with ARPDK
- Increased risk of recurrence in sibs
 - Suggests simple genetic mechanism

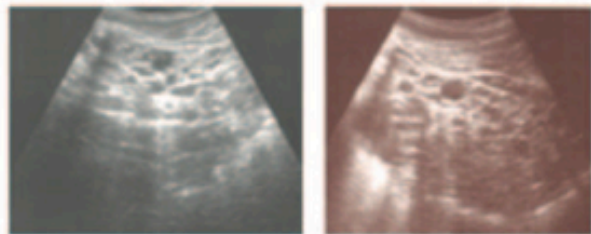
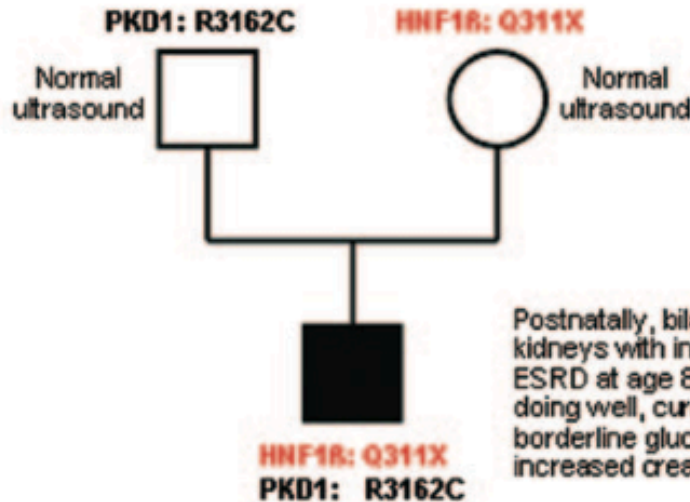
Early-onset disease associated with co-inheritance of a truncating and hypomorphic *PKD1* allele



Other explanations for early onset ADPKD

- Not all EO cases due to co-inheritance of ADPKD alleles
- Co-inheritance of mutations at other loci may cause EO PKD
 - *HNF1B*
 - *PKHD1*
- Analysis of candidate panel or whole exome screen may be appropriate in unresolved EO cases

Family D



Gene	Variant	GD ^a	SIFT ^d		Poly-Phen2 ^a		Align-GVGD ^b		Mutation-Taster ^e	Con-servation	Con-sensus	Com-ments ^f
			VS	MG	VS	MG	VS	MG	MG			
PKD1	R3162C	180	0.00	HLP	1.00	HLP	C65	HLP	HLP	0/6	HLP	novel

R3162C

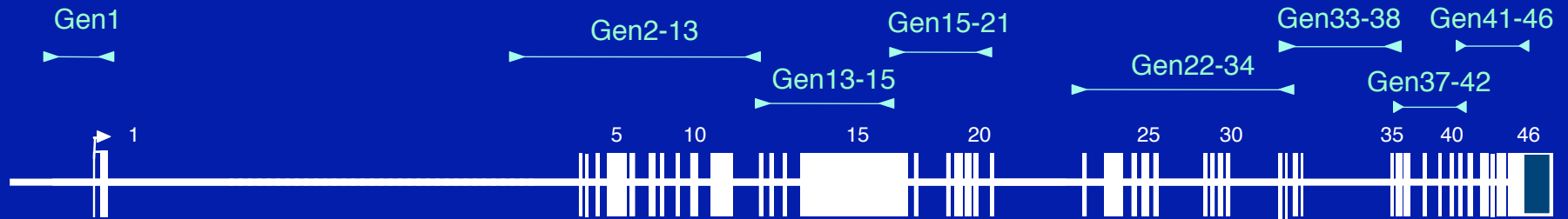
Human	3111	PCGKRGSPFYELLVKTGWGRSGTTAHVGIIMLYGVDNRSGHRHLDGDRAPHRNSLDIPQI
Dog	3117	PCGKRGSPFYELLVKTGWGRSGTTAHVGIIMLYGVDNRSGHRHLDGDRAPHRNSLDIPQI
Mouse	3103	PCGKRGSPFYELLVKTGWGRSGTTAHVGIIMLYGVDNRSGHRHLDGDRAPHRNSLDIPQI
Rat	3093	PCGKRGSPFYELLVKTGWGRSGTTAHVGIIMLYGVDNRSGHRHLDGDRAPHRNSLDIPQI
Chicken	3068	PCGKRGSPFYELLVKTGWGRSGTTAHVGIIMLYGVDNRSGHRHLDGDRAPHRNSLDIPQI
Xenopus	2696	PCGKRGSPFYELLVKTGWGRSGTTAHVGIIMLYGVDNRSGHRHLDGDRAPHRNSLDIPQI
Fugu	3238	LCGCDLRFYELLVKTGWGRGAGTTAHVGIIMLYGVDNRSGHRHLDGDRAPHRNSLDIPQI

Example of combination of *PKD1* and *HNF1B* allele causing EO PKD

Gene-based diagnostics in ADPKD is complex

- Genetic and extreme allelic heterogeneity
 - Completely screen *PKD1* and *PKD2* required
- Segmental duplication of *PKD1*
 - Locus specific enrichment required
 - Exon capture methods unreliable
- Many variants of uncertain significance
- Many *PKD1* non-truncating changes hypomorphic
 - Identification of hypomorphic alleles difficult
- Genetic test is expensive and not always informative
- Reports often uninformative and difficult to understand
 - Clinical testing only available through one vendor in US:
 - Athena Diagnostics - ~\$5000 with MLPA testing
 - Recent Supreme Court ruling may open US market

Next-generation sequencing allows rapid analysis of multiple patient samples



- *PKD1* and *PKD2* amplified as 14 long-range products
 - exon capture unreliable for *PKD1* because of genomic duplication
- Potential for higher throughput and reduced cost
- Mutation detection rate likely to be comparable
- Introns, UTRs and promoters could also be screened

Molecular Diagnosis of Autosomal Dominant Polycystic Kidney Disease Using Next-Generation Sequencing

Adrian Y. Tan,^{*} Alber Michael,^{*} Genyan Liu,^{*} Olivier Elemento,[†] Jon Blumenfeld,^{†§} Stephanie Donahue,[§] Tom Parker,[§] Daniel Levine,[§] and Hanna Rennert^{*}

Table 4 NGS Analytic Sensitivity and Specificity (Variants Detection)

NGS	Sanger sequencing		
	Variant alleles (positive)	Reference alleles (negative)	Total
Variant alleles (positive)	248	0	248
Reference alleles (negative)	2	1825	1827
Total	250	1825	2075

Table 6 Comparison of Reagents, Sequencing Costs, and Time of Labor for Sanger Sequencing and NGS

Method	Purpose	Quantity	Cost (\$)			Labor time (days)
			Per sample	Per run	Per subject	
Sanger sequencing (<i>N</i> = 25)	LR-PCR (<i>PKD1</i>)	250	2.40	600.00	24.00	5
	Standard PCR (<i>PKD2</i>)	400	1.50	600.00	24.00	4
	Purification	200	2.40	480.00	19.20	1
	Sequencing primers	3050	0.10	305.00	12.20	NA
	Sanger sequencing	1600	3.00	4800.00	192.00	5
	Data analysis	NA	NA	NA	NA	4
	Total				6785.00	271.40
NGS (<i>N</i> = 25)	LR-PCR (<i>PKD1</i> and <i>PKD2</i>)	250	1.45	362.50	14.50	2
	LR-PCR product quantification	250	0.12	30.00	1.20	0.5
	DNA fragmentation	25	6.50	162.50	6.50	0.5
	Library preparation	25	20.00	500.00	20.00	3
	Library quality assessment	25	0.20	5.00	0.20	0.25
	NGS sequencing (MiSeq)	1	990.00	990.00	39.60	1
	Data analysis	NA	NA	NA	NA	1
	Total				2050.00	82.00

Mutation-based diagnostics in ADPKD is likely to be more widely employed

- Mutation identified in 90% cases
 - Definite (truncating) mutations in ~65% families
- Bioinformatic scoring of non-definite mutations increasingly reliable
 - Recurrent mutations ~50% in recent studies
 - Mutation database of value
 - Identify pathogenic mutations
 - Highlight hypomorphic changes
- Of diagnostic and prognostic value
- Mutation type may in the future influence treatment options
 - Similar to cystic fibrosis
- Cost of test needs to decrease
- Reliability and interpretation of results needs to improve