Molecular Methods in Urology and Nephrology

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GENE, NUCLEOTIDE AND NUCLEOSIDE







DNA double helix





DNA replication





CHROMOSOME STRUCTURE







| Chromosome • | Genes ^[26] • | Total base pairs | % of bases • | Sequenced base pairs ^[27] |
|--------------------|-------------------------|------------------|-------------------|--------------------------------------|
| 1 | 2000 | 247,199,719 | 8.0 | 224,999,719 |
| 2 | 1300 | 242,751,149 | 7.9 | 237,712,649 |
| 3 | 1000 | 199,446,827 | 6.5 | 194,704,827 |
| 4 | 1000 | 191,263,063 | 6.2 | 187,297,063 |
| 5 | 900 | 180,837,866 | 5.9 | 177,702,766 |
| 6 | 1000 | 170,896,993 | <mark>5.5</mark> | 167,273,993 |
| 7 | 900 | 158,821,424 | 5.2 | 154,952,424 |
| 8 | 700 | 146,274,826 | 4.7 | 142,612,826 |
| 9 | 800 | 140,442,298 | <mark>4.6</mark> | 120,312,298 |
| 10 | 700 | 135,374,737 | 4.4 | 131,624,737 |
| 11 | 1300 | 134,452,384 | 4.4 | 131,130,853 |
| 12 | 1 100 | 132,289,534 | 4.3 | 130,303,534 |
| 13 | 300 | 114,127,980 | 3.7 | 95,559,980 |
| 14 | 800 | 106,360,585 | 3.5 | 88,290,585 |
| 15 | 600 | 100,338,915 | 3.3 | 81,341,915 |
| 16 | 800 | 88,822,254 | 2.9 | 78,884,754 |
| 17 | 1200 | 78,654,742 | 2.6 | 77,800,220 |
| 18 | 200 | 76,117,153 | 2.5 | 74,656,155 |
| 19 | 1500 | 63,806,651 | 2. <mark>1</mark> | 55,785,651 |
| 20 | 500 | 62,435,965 | 2.0 | 59,505,254 |
| 21 | 200 | 46,944,323 | 1.5 | 34,171,998 |
| 22 | 500 | 49,528,953 | 1.6 | 34,893,953 |
| X (sex chromosome) | 800 | 154,913,754 | 5.0 | 151,058,754 |
| Y (sex chromosome) | 50 | 57,741,652 | 1.9 | 25,121,652 |
| Total | 21,000 | 3,079,843,747 | 100.0 | 2,857,698,560 |

Base pairs of Genes





GENE STRUCTURE







REGULATION OF GENE EXPRESSION





Transcription, post-transcriptional processing, translation, and post-translational processing





ALLELE AND LOCUS





GENETIC DISORDERS



SINGLE-GENE DISORDER

CHROMOSOME ABNORMALITY

MULTFI ACTORIAL DI SORDER

polycystic kidney disease

| MIM | Name of Disease | Inheritance | Gene |
|--------|--|-------------|---------|
| 263100 | POLYCYSTIC KIDNEY, CATARACT, AND CONGENITAL BLINDNESS | | - |
| 600273 | - POLYCYSTIC KIDNEY DISEASE, INFANTILE SEVERE, WITH TUBEROUS SCLEROSIS; PKDTS | AD | |
| 600666 | POLYCYSTIC KIDNEY DISEASE 3 WITH OR WITHOUT POLYCYSTIC LIVER DISEASE; PKD3 | AD | GANAB |
| 618061 | POLYCYSTIC KIDNEY DISEASE 6 WITH OR WITHOUT POLYCYSTIC LIVER DISEASE; PKD6 | AD | DNAJB11 |
| 617874 | POLYCYSTIC LIVER DISEASE 3 WITH OR WITHOUT KIDNEY CYSTS; PCLD3 | AD | ALG8 |
| 617610 | POLYCYSTIC KIDNEY DISEASE 5; PKD5 | AR | DZIP1L |
| 617875 | POLYCYSTIC LIVER DISEASE 4 WITH OR WITHOUT KIDNEY CYSTS; PCLD4 | AD | LRP5 |
| 617004 | POLYCYSTIC LIVER DISEASE 2 WITH OR WITHOUT KIDNEY CYSTS; PCLD2 | AD | SEC63 |
| 263210 | GILLESSEN-KAESBACH-NISHIMURA SYNDROME; GIKANIS | AR | ALG9 |
| 173900 | POLYCYSTIC KIDNEY DISEASE 1 WITH OR WITHOUT POLYCYSTIC LIVER DISEASE; PKD1 | AD | PKD1 |
| 613095 | POLYCYSTIC KIDNEY DISEASE 2 WITH OR WITHOUT POLYCYSTIC LIVER DISEASE; PKD2 | AD | PKD2 |
| 606702 | PKHD1 GENE; PKHD1 FIBROCYSTIN; FCYT POLYDUCTIN POLYCYSTIC KIDNEY AND HEPATIC DISEASE 1 GENE | AR | |
| 263200 | POLYCYSTIC KIDNEY DISEASE 4 WITH OR WITHOUT POLYCYSTIC LIVER DISEASE; PKD4 | AR | FCYT |
| 174000 | MEDULLARY CYSTIC KIDNEY DISEASE 1; MCKD1 | AD | MUC1 |
| 102500 | HAJDU-CHENEY SYNDROME; HJCYS SERPENTINE FIBULA-POLYCYSTIC KIDNEY SYNDROME; SFPKS | AD | NOTCH2 |
| 609886 | GLOMERULOCYSTIC KIDNEY DISEASE WITH HYPERURICEMIA AND ISOSTHENURIA | | UMOD |

polycystic kidney disease

- PKD is a <u>genetic disorder</u> in which the <u>renal tubules</u> become structurally abnormal, resulting in the development and growth of multiple <u>cysts</u> within the kidney.
- These cysts may begin to develop in utero, in infancy, in childhood, or in adulthood.
- PKD is caused by abnormal genes which produce a specific abnormal protein which has an adverse effect on tubule development.
- PKD is a general term for two types, each having their own pathology and genetic cause:
- autosomal dominant polycystic kidney disease (ADPKD)
- ▶ and <u>autosomal recessive polycystic kidney disease</u> (ARPKD).

- Autosomal dominant polycystic kidney disease (ADPKD) is the most common of all the inherited cystic kidney diseases with an incidence of 1:500 live births.
- Studies show that 10% of <u>end-stage kidney disease</u> (ESKD) patients being treated with <u>dialysis</u> in Europe and the U.S. were initially diagnosed and treated for ADPKD.
- There are three <u>genetic</u> mutations in the <u>PKD-1</u>, <u>PKD-2</u>, and PKD3 <u>gene</u> with similar phenotypical presentations.

- Gene PKD1 is located on <u>chromosome 16</u> and codes for a protein involved in regulation of cell cycle and intracellular calcium transport in epithelial cells, and is responsible for 85% of the cases of ADPKD.
- A group of voltage-linked <u>calcium channels</u> are coded for by PKD2 on <u>chromosome 4</u>.
- PKD3 recently appeared in research papers as a postulated third gene. Fewer than 10% of cases of ADPKD appear in non-ADPKD families.

- Cyst formation begins in utero from any point along the <u>nephron</u>,
- although fewer than 5% of nephrons are thought to be involved.
- As the cysts accumulate fluid, they enlarge, separate entirely from the nephron, compress the neighboring kidney <u>parenchyma</u>, and progressively compromise <u>kidney function</u>.

- Autosomal recessive polycystic kidney disease (ARPKD) (OMIM #263200) is the lesser common of the two types of PKD, with an incidence of 1:20,000 live births.
- And is typically identified in the first few weeks after birth.
- Unfortunately, the kidneys are often <u>underdeveloped</u> resulting in a 30% death rate in newborns with ARPKD.

HYPEROXALURIA

- ► **Hyperoxaluria** is an excessive urinary excretion of <u>oxalate</u>.
- Individuals with hyperoxaluria often have <u>calcium oxalate kidney stones</u>.
- It is sometimes called Bird's disease, after <u>Golding Bird</u>, who first described the condition.
- ► Types
- Primary hyperoxaluria
- ► Enteric hyperoxaluria
- ► Idiopathic hyperoxaluria
- ► <u>Oxalate</u> poisoning

HYPEROXALURIA

| 259900 | HYPEROXALURIA, PRIMARY, TYPE I; HP1 | AR | AGXT |
|--------|---|----|-------|
| | OXALOSIS I | | |
| | GLYCOLIC ACIDURIA | | |
| | ALANINE-GLYOXYLATE AMINOTRANSFERASE DEFICIENCY | | |
| | PEROXISOMAL ALANINE:GLYOXYLATE AMINOTRANSFERASE | | |
| | DEFICIENCY | | |
| | HEPATIC AGT DEFICIENCY | | |
| | SERINE: PYRUVATE AMINOTRANSFERASE DEFICIENCY | | |
| 260000 | HYPEROXALURIA, PRIMARY, TYPE II; HP2 | AR | GRHPR |
| | OXALOSIS II | | |
| 1/4 | GLYCERIC ACIDURIA | | |
| | GLYOXYLATE REDUCTASE/HYDROXYPYRUVATE REDUCTASE | | |
| | DEFICIENCY | | |
| | D-GLYCERATE DEHYDROGENASE DEFICIENCY | | |
| 613616 | HYPEROXALURIA, PRIMARY, TYPE III; HP3 | - | HOGA1 |
| | | | |
| | | | |

- Type I (PH1) is associated with <u>AGXT</u> protein, a key enzyme involved in breakdown of <u>oxalate</u>.
- PH1 is also an example of a protein mistargeting disease, wherein AGXT shows a trafficking defect: instead of being trafficked to <u>peroxisomes</u>, it is targeted to <u>mitochondria</u>, where it is metabolically deficient despite being catalytically active.
- ► Type II is associated with <u>GRHPR</u>.
- It is also a complication of jejunoileal bypass, or in any patient who has lost much of the <u>ileum</u> with an intact <u>colon</u>.
- ▶ This is due to excessive absorption of oxalate from the colon.

Cystinuria

- Cystinuria is an inherited <u>autosomal recessive disease</u> that is characterized by high <u>concentrations</u> of the amino acid <u>cysteine</u> in the <u>urine</u>, leading to the formation of cystine <u>stones</u> in the <u>kidneys</u>, <u>ureter</u>, and <u>bladder</u>. It is a type of <u>aminoaciduria</u>.
- Cystinuria is caused by mutations in the <u>SLC3A1</u> and <u>SLC7A9</u> genes.
- These defects prevent proper reabsorption of basic, or positively charged, amino acids: <u>Cysteine</u>, <u>lysine</u>, <u>ornithine</u>, <u>arginine</u>.
- Under normal circumstances, this protein allows certain amino acids, including cysteine, to be reabsorbed into the blood from the filtered fluid that will become urine.

► CYSTINURIA

► Alternative titles; symbols

- CSNU CYSTINURIA, TYPE I, FORMERLY; CSNU1, FORMERLY CYSTINURIA, TYPE II, FORMERLY CYSTINURIA, TYPE III, FORMERLY; CSNU3, FORMERLY CYSTINURIA, TYPE NON-I, FORMERLY
- Other entities represented in this entry:
- CYSTINURIA, TYPE A, INCLUDED
- CYSTINURIA, TYPE B, INCLUDED CYSTINURIA, TYPE A/B, INCLUDED

Phenotype-Gene Relationships

| Location | Phenotype | Phenotype MIM number | Inheritance | Phenotype mapping key | Gene/Locus | Gene/Locus MIM number |
|----------|------------|-------------------------|-------------|--------------------------|------------|--------------------------|
| 2p21 | Cystinuria | 220100 | AR, AD | 3 | SLC3A1 | 104614 |
| 19q13.11 | Cystinuria | 220100 | AR, AD | 3 | SLC7A9 | 604144 |

- Mutations in either of these genes disrupt the ability of this transporter protein to reabsorb these amino acids, allowing them to become concentrated in the urine.
- As the levels of cysteine in the urine increase, it forms <u>cystine</u> crystals, resulting in kidney stones. Cystine crystals form hexagonal-shaped crystals that can be viewed upon microscopic analysis of the urine.
- The other amino acids that are not reabsorbed do not create crystals in urine.
- ► The overall <u>prevalence</u> of cystinuria is approximately 1 in 7,000 neonates (from 1 in 2,500 neonates in <u>Libyan</u> <u>Jews</u> to 1 in 100,000 among <u>Swedes</u>).

Male infertility

Genetic factors cause account for 10–15% of male infertility, including chromosomal aberrations and single gene mutations.

Klinefelter Syndrome

Y chromosome deletions

Klinefelter Syndrome

One of the most commonly known causes of infertility is Klinefelter Syndrome, affecting 1 out of 500-1000 newborn males.

Klinefelter Syndrome is a chromosomal defect that occurs during gamete formation due to a non-disjunction error during cell division.

Resulting in males having smaller testes, reducing the amount of testosterone and sperm production.

Y chromosome deletions

Y chromosomal infertility is a direct cause of male infertility due to its affects on sperm production, occurring in 1 out of every 2000 males.

Usually affected men show no sign of symptoms other than at times can exhibit smaller teste size.

Men with this condition can exhibit <u>azoospermia</u> (no sperm production), <u>oligospermia</u> (small number of sperm production), or they will produce abnormally shaped sperm.

These individuals are thereby "Y-linked", although daughters are not affected due to the lack of the Y chromosome.

Microdeletions in the Y chromosome

- Microdeletions in the Y chromosome have been found at a much higher rate in infertile men than in fertile controls and the correlation found may still go up as improved genetic testing techniques for the Y chromosome are developed.
- Much study has been focused upon the "azoospermia factor locus" (AZF), at Yq11.
- A specific partial deletion of AZFc called gr/gr deletion is significantly associated with <u>male infertility</u> among Caucasians in Europe and the Western Pacific region.
- Additional genes associated with <u>spermatogenesis</u> in men and reduced fertility upon Y chromosome deletions include <u>RBM</u>, <u>DAZ</u>, <u>SPGY</u>, and <u>TSPY</u>.



Lab Methods

Karyotyping

PCR/Sanger Sequencing

Deletion/Duplication Analysis

Next-Gen Sequencing

Karyotyping



| Human male G-bands | 2 | and succession of the successi | | | | s second |
|------------------------|------------|--|----|---|----|------------|
| 2 | | a change | S. | | | |
| P ₁₃ | 1 4 | | 16 | 3 | 17 | 218 |
| 19 | 20 | 21 | 22 | 8 | × | B _ |



PCR and Sanger sequencing



AGXT gene

| GCCTGCCAAGCCTCAATTATCTGTGCTCCGACCCTTTAAGAAACACTTCTCTCACCCCTG |
|---|
| AGCTRACCAGARTAAGAGGGCCTGGACCTCCAGGACTCAGAGTGGGACCGAGGGGCTG |
| GOOLDAGGACAGCITIGICACACICIGITITICIGICACICAGCCCCIGGGGGCICCCICI GOOTGAGGGCTTGTTGTCGCTGTGTCGCGTGACGGCAGCAGCAGCAGGGGGCICCCICI |
| TCAGEGAACAAAAGECAGGGCTGCCACEGAACCCCATCCACCACTCTCACCTCAC |
| CTGTGTCCCCCCTGCTGGGAAATATTCCAGGCTTTGGCCAAGGCCAGTGCAGCCCCAGGT |
| TCCCGAGCGCAGCTGGGTGCGGACCA <mark>TG</mark> GCCTCTC <mark>A</mark> CA <mark>AGC</mark> TGCT <mark>GG</mark> TG <mark>ACCCC</mark> C <mark>QCC</mark> |
| RAGCCCTCCTCAACCCCTCTCCATCCCAACCAGCTCCTCCTCCCCCCCC |
| AACCTCCCTCCCCCACCACCCCCCCCCCCCCCCCCCCC |
| GATATGTACCAG |
| gtaggagtgggggtcactcggggggctcaccctataccacccgcatgcag |
| ATCATEGACGAGATCAAGGAAGGCATCCAGTAGTGTTCAAGACCAGGAAGCCACTCACA |
| CTGSTCATCTCTGGCTCGGGACACTGTGCCCTGGAGGCCGCCCTGGTCAAHGTGCTGGAG |
| CCTGGGGACTCCTTCCTGGTTGGGGCCAATGGCATTTGGGGCCAGCGAGCCGTGGACATC |
| GGCEAGCATAG |
| qtaaqqqaqaqqcccaqqtqqqqatacqqcccactctqtcctqcacccaq |
| |
| AGGA |
| |
| gtaggggacccggggtggggggtcagcccacagccgtccctgcttcctcag |
| GGCCT <mark>G</mark> GCCA <mark>GCA<mark>CA</mark>AG<mark>CAET<mark>GETGCTGTT</mark>CTTA<mark>AC</mark>CGA<mark>GG</mark>GG<mark>GAGT<mark>GT</mark>CCAC<mark>G</mark>GG</mark></mark></mark> |
| GTGCTGCAGCCCTTGATGGCTTCGGGGGAACTCTGCCACAG |
| qtqaqcctqqccccaqqqcqqtqqatccattctqtcccccacctctccaq |
| |
| GRAND COLOURS COLOURS COLOURS COLOURS |
| |
| gtaagggtgggetetgagageeetateeegageaaaceaeeeatetaeag |
| CCARCCACATCCTGTACTCCGCGCTCCCAGAAGGCCCTGAACGCCCCACCAGGACCTCGC |
| TCA <mark>TCTCC</mark> TTCAGTG <mark>A</mark> C <mark>AAG</mark> CC <mark>A</mark> 2 |
| gtgagtgacccacagacceteacetgeteageetgettettteteeceag |
| 22 Z C 2 Z C 2 T C 2 Z C C C C C 2 C 2 C 2 C C C C C C C |
| GCC AACTTCTCCCCCTCTCACCACCACCACCACCACCACCACC |
| |
| grgæggerrggeægggærgggæægggæeeæægeeeeregrgrerreeæg |
| GTACCATCACACAATCCCCCGTCATCCCCCCGTACAGCCTCCAGAGAGAG |
| TOCOGRA |
| gtgcatgggctgcactccacaggagcccaccagegccatctcccacacag |
| GECTGEAGAACAGCTGGCGACCAGCGCGAGGCGCGCGCGAGGCGCGCGC |
| CAGGCACTGGGGCTCCAGCTCTTCGTGAAGG <mark>ACCCG</mark> |
| |
| g værggeggerer vygen vogggen i finne i gene vyngerer vervyeg |
| GC (CTCCGG CTTCCCA 04) GTCA CCACTGT CCTCCTCCG 0 G (C) Au GACTG 04) GAGAG |
| ATCOLOGENAG |
| |
| gtgagagggagegeetegagggeetegteageeegeeetgtgeeeeeag |
| GTGCTGCGGATCGGCCTGCTGGGCTGCAATGCCACCCGCGAGAATGTGGACCGCGTGACG |
| GAG <mark>GCCCTGAGGGCG</mark> GCCC <mark>TGCAGC</mark> ACTGCCCCAAGAAGCTG <mark>A</mark> GACCTGCCCACTGG |
| CARACASCI GECART GERALAR CARACATERIC CATERIC CAREGO TO A GERAL CAREGO A CARACATERIA CAREGO A CARACATERIA CAREGO A CARACATERIA |
| AGAACCAGGCAGCCTCCCTGGCCCCAGGCAGCCCTTTTCCCTCCAGTGGCACCCCTGGA |
| AACAGTCCACTTEGGCGCAAAACCCAGTGCCTTCCAAATCAGCTGCACTCCCCAGGCCAT |
| GAGCCTCCCGGGAATGTTTAATAAAGGGCCTGGCCAACTCTCCTCAC |
| |

..tgetttgeaageaegeageagggttgagttgttgeaeeatagaeeaeagg







MLPA

Whole exome sequencing (NGS)









آزمایش های ngs و پانل

موارد زیر مثال هایی از اختلالاتی هستند که به روش NGSو با بررسی تعدادی ژن یا کل ژنوم مورد بررسی قرار می گیرند.

- سرطانها
- ناتوانی ذهنی و تاخیر تکاملی
 - ناشنوایی های ارثی
 - نابینایی های ارثی
 - بیماری های متابولیک ارثی
- بیماری های سیستم اسکلتی و استخوانی
 معلولیت های حرکتی
 - بیماری های سیستم اعصاب مرکزی
- بیماری های ژنتیک پوست (ژنودر ماتوز ها)
 - بیماری های ژنتیک قلب (آریتمی ها و...)



Diagnosis



Prenatal testing and prenatal expectations

Prenatal testing

► <u>in vitro fertilisation</u>, via <u>preimplantation genetic diagnosis</u>

Chorionic villus sampling

► <u>amniocentesis</u>



آمنیوسنتز، CVS و NIPT







🗸 مشاورہ ژنتیک

🗸 ژنتیک سرطان

🗸 مشاوره پیش از ازدواج، پیش از بارداری و حین بارداری 🗸 مشاوره موارد مربوط به سقط، ناباروی و مرده زایی 🗸 انجام تمامی آزمایشات تخصصی ژنتیک پزشکی 🗸 سینوژنتیک و کاریوتیپ





اصفهان - خیابان شریعتی - بین چهارراه پلیس و حکیم نظامی - پلاک ۲۰۸ 🌈 موسس و مسئول فنی 1 TETERANY - TETERANE ·9187719118

دكتر مجيد خيراللهي







GENE AZMA Laboratory



⊘ مش_اورہ ژنتےک

- 📀 مشاوره پیش از ازدواج و پیش از بارداری و حین بارداری
- 📀 مشاوره مـوارد مـربوط به سقـط ، نا باروری و مـرده زایی
- 📀 انجــام همــه آزمایشـات تخصصـی ژنتیـک پزشـکـی
- 🔗 آزمایشات تشخیص ژنتیک بر روی بند ناف نوزادان

فوت شده با علت نا مشخص ۵ سیت وژنتیک وکاری وتیپ ۵ ژنتیک مولک ولی تشخیصی ۵ تشخیص پیش از تولد و PND ۱۹ آمنی وسنت ز، CVS و NIPT ۵ آزمایش های NGS و پانل

⊗ ژنتیک س_رط_ان

موسس و مسئول فنی: ه که دکتـــر مجیــــد خیراللهـــی

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خیابان شریعتی، بین چهارراه پلیس و حکیم نظامی، پلاک ۲۰۸





