Molecular pathology and genetics of congenital hepatorenal fibrocystic syndromes

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The hepatorenal fibrocystic (HRFC) syndromes are a heterogeneous group of severe monogenic conditions that may be detected before birth. Commonly, HRFC syndromes present in the neonatal and paediatric age, with consistent developmental abnormalities mostly involving the liver and kidney. The changes include the proliferation and dilatation of epithelial ducts in these tissues with abnormal deposition of extracellular matrix. In this review, we examine the clinical features and differential diagnoses of this group of syndromes, including autosomal recessive polycystic kidney disease (ARPKD), juvenile nephronophthisis (JNPH), Meckel-Gruber syndrome (MKS), Bardet-Biedl syndrome (BBS), and Jeune asphyxiating thoracic dystrophy (JATD). Extrahepatic manifestations include mostly bone and central nervous system abnormalities, dysmorphic features, and developmental delay. Previously, it has been suggested that ARPKD, JATD, and Ellis-van Creveld syndrome (EvC) may arise from defects in differentiation in a common developmental pathway. We review recent molecular advances in the recessive HRFC syndromes and discuss this hypothesis.

Hepatic fibrosis may manifest either as a consequence of ductal plate malformation within the congenital hepatorenal fibrocystic syndromes or as a separate entity. In this review we provide a brief update on the molecular pathology and genetics of these disorders. We examine how recent molecular genetic advances in the characterisation of the ARPKD and JNPH genes provide insights into the "common developmental pathway" hypothesis for the aetiology of these clinically heterogeneous group of disorders. The molecular basis of a number of rarer HRFC syndromes remains unknown, and it is hoped that these insights might provide a rationale for the selection of candidate genes in the future, on the basis of protein function.

PATHOLOGY OF AUTOSOMAL RECESSIVE FIBROCYSTIC DISEASES

HRFC disease is characterised by changes in the parenchymal tissues of the liver, kidney, and sometimes pancreas or other organs. They include the proliferation and dilatation of epithelial ducts, and proliferative changes in the extracellular matrix of stromal connective tissue. One of the manifestations that has received intensive study are related to the ductal plate malformation of the liver and comprise proliferation and dilatation of the intrahepatic bile ductules, with a variable degree of fibrosis and cyst formation. 

The intrahepatic bile ducts develop out of sheets of primitive hepatic epithelial cells in a pattern determined by the branching of the portal vein and surrounding mesenchyme. The primitive epithelial cells that are in direct contact with the portal vein mesenchyme transform into bile duct type cells by forming a single layer and later two layers (fig 1A). Development progresses by the formation of a cleft between the two layers of cells known as the ductal plate (at gestational weeks 9 to 12). In response to unknown developmental signals, the biliary cells migrate from the ductal plate into the portal mesenchyme, with biliary structures still in partial contact with the ductal plate (the remodelling ductal plate stage at gestational weeks 13 to 17, fig 1A).

Finally, remodelled bile ducts are formed that are centrally located in the portal tracts, without contact with ductal plate remnants (the remodelled bile duct stage at gestational weeks 18 to 40, fig 1A). The unincorporated epithelium of the ductal plate then disappears. Any interruption in the remodelling of the ductal plate can result in the persistence of embryonic duct structures, termed the ductal plate malformation (fig 1B). The initial lesion in the liver is thought to be excessive ductular proliferation, characterised by the formation of multiple dilated perportal biliary ductules, presumably resulting from a defect in complex epithelial-mesenchymal interactions. The ductal plate malformation is a histopathological characteristic of a number of fibrocystic syndromes, such as MKS (fig 2A, B), hepatic-pancreato-renal or Ivemark syndrome (HPRS), and JATD (table 1). The ductal plate malformation can also result in subsequent liver damage, and inflammation can induce the formation of fibrosis, and in some cases, cirrhosis.

Hepatic fibrosis may manifest either as a consequence of ductal plate malformation within
a monogenic fibrocystic disorder, in response to hepatocyte damage and subsequent necrosis, or as necrosis independent hyperstimulation of matrix producing cells. In the liver, the major cellular sources of extracellular matrix are activated Ito or hepatic stellate cells. These are fat storing cells that resemble fibroblasts, and in response to chronic hepatocyte damage they differentiate into myofibroblasts. These cells then secrete matrix proteins, such as interstitial collagen types I, III, IV, and laminin, and are responsible for the deposition and accumulation of the majority of the excess extracellular matrix in the fibrotic liver. In most cases, fibrosis consists of broad, collagenous fibrous bands surrounding normal hepatic lobules. The increase of the periportal connective tissue may compress portal vein radicles, leading to hypertension in the portal system. Cirrhosis is a common end stage for a number of monogenic disorders that result in hepatic insult owing, in part, to fibrotic and cystic changes. Renal cysts can occur as manifestations of both non-genetic and genetic disorders. In the former case, non-hereditary lesions, consisting of an enlarged kidney filled with large cystic structures, can occur in response to uterine atresia or obstruction. "Renal dysplasia" is the usual term to describe the abnormal differentiation of renal parenchyma, with the occurrence of microscopic abnormalities such as cystic dilata-
tions and primitive ducts surrounded by connective tissue. In some conditions, renal dysplasia has been attributed to abnormal embryonic differentiation or developmental arrest leading to the persistence of mesonephric tissue and primitive or fetal structures. Epithelial cell proliferation is thought to be a prerequisite for cyst expansion, with some cysts retaining some of the functional properties of the tubular epithelium, such as active ion transport, from which they were derived. Cysts that lose their connection with their tubule of origin are thought to expand by a mechanism of transepithelial fluid secretion, which cause compression and eventual atrophy of surrounding parenchyma. It is not clear if only a proportion of nephrons contribute to cyst formation, or if all nephrons are involved with varying degrees of dilatation.

**FIBROCYSTIC DISEASES OF VISCERAL ORGANS**

**Polycystic kidney disease in children**

"Polycystic disease" was first described as long ago as 1856, and a spectrum of disease is now known to exist. Polycystic kidney disease includes autosomal dominant polycystic kidney disease (ADPKD) and autosomal recessive polycystic kidney disease (ARPKD). Congenital hepatic fibrosis is a characteristic feature of ARPKD, but may also be associated with other conditions. Liver histology shows portal spaces enlarged by periportal fibrosis, proliferation of biliary ducts, which can be dilated and tortuous, and ductal plate malformation or insufficient remodelling of the primitive intrahepatic biliary system. In some cases, abnormal multiple bile ductules can lose connection with the biliary system and dilate to form large cysts. Patients with ARPKD tend to present with congenital hepatic fibrosis and Caroli disease (known as the "combined" form of the disease or Caroli syndrome). The "pure" form of Caroli disease is characterised by multifocal dilatations of the larger, segmental intrahepatic bile ducts. Subsequently, patients can present with recurrent bacterial cholangitis because of bile stagnation at the ectopias of the intrahepatic biliary tree.

The main clinical manifestation of ARPKD is both ectasia and cystic dilatation of the renal collecting tubules in the kidney that prevents the normal development of nephrons, so that they remain as primitive ducts lined by undifferentiated epithelium and sheathed with thick layers of connective tissue. The kidney appears spongy, and there is no clear separation of the cortex and medulla (fig 2D). The prognosis of ARPKD is dependent on the degree of renal involvement and speed of disease progression to renal insufficiency and end stage renal failure (ESRF). A number of long term survivors have been reported.

The PCK rat model of polycystic kidney disease also develops renal and hepatic cysts, but in addition also displays ductal plate malformation of the intrahepatic bile ducts and overgrowth of portal connective tissue that resembles human ARPKD. Genetic analysis of the PCK rat has recently led to the identification of the human orthologue, PKHD1, of the rat gene. The human PKHD1 gene is predicted to encode a large protein of approximately 450 kDa, named fibrocystin, of unknown function (fig 3). Fibrocystin contains at least 10 copies of an immunoglobin fold-like domain (the IPT/TIG domain) found in plexins, transcription factors, and extracellular regions of receptor proteins that appear to regulate cell proliferation and cellular adhesion. Another group reported recent similar findings, with the identification of a PKHD1 gene product essentially identical in both size and sequence, to fibrocystin that they name “polyductin”. It will be interesting to elucidate the normal role of fibrocystin in the development and differentiation of collecting ducts in the kidney and bile ducts in the liver. Fibrocystin may be a receptor of growth signals because of the predicted extracellular domain, but could also participate in mediating cell adhesion. The unusual multiplicity of splicing variants for PKHD1 may be an important mechanism in regulating these functions by different protein isoforms.

In contrast, patients with ADPKD tend to present with large hepatic cysts (fig 2C), and the number of hepatic cysts can...
<table>
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<tr>
<th>Inherited disorder</th>
<th>Gene(s) or loci, (protein product); mode of inheritance</th>
<th>Renal abnormalities: imaging and histopathology</th>
<th>Hepatic abnormalities</th>
<th>Other associated features</th>
<th>Prognosis</th>
<th>Differential diagnosis</th>
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<tr>
<td>JAR or juvenile polycystic kidney disease, congenital hepatic fibrosis</td>
<td>PKHD1 (fibrocystin); AR</td>
<td>Antenatally diagnosed by ultrasonography with grossly enlarged bilaterally echogenic kidneys, absent urine, oligohydramnios and pulmonary hypoplasia. Cysts less than 3 mm in diameter extend from papillary tips to the surface of the cortex; lined by flattened epithelium representing dilated collecting tubules. Adjacent parenchyma develops interstitial fibrosis and glomerulosclerosis</td>
<td>Congenital hepatic fibrosis, Caroli’s disease, and biliary dysgenesis. Hepatic disease more prominent if patients present as older children, and results in cirrhosis and portal hypertension</td>
<td>Neonatal presentation often with pulmonary hypoplasia</td>
<td>Majority of neonatally presenting patients die within 6 weeks of pulmonary disease and renal failure; 80% of survivors live until 15 years of age. All will develop ESRF by 20 years</td>
<td>Caroli’s disease</td>
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<td>Nephronophthisis</td>
<td>NPHP1 (nephrocystin), loci NPHP2-4; AR</td>
<td>Ultrasound will detect small kidneys, uniformly increased echogenicity and cysts at the corticomedullary junction. Histology shows cortical atrophy with variable numbers of cysts at the corticomedullary junction (up to 1 cm in size). Histology shows tubular atrophy, interstitial fibrosis, and chronic inflammation; progression to glomerulosclerosis with markedly thickened basement membrane</td>
<td>Congenital hepatic fibrosis</td>
<td>Cone shaped epiphyses, retinal degeneration [Senior-Løken syndrome]</td>
<td>Patients develop ESRF within 5–10 MCKD years after presentation</td>
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<tr>
<td>Renal-hepatic-pancreatic dysplasia (Ivemark syndrome)</td>
<td>Autosomal recessive</td>
<td>May be detected on antenatal ultrasound showing bilaterally enlarged polycystic kidneys. Histology usually reveals cystic dysplasia</td>
<td>Congenital hepatic fibrosis and Caroli’s disease</td>
<td>Dilated pancreatic ducts, short ribs, polydactyly, polydactyly, cardiac anomalies, situs inversus, CNS abnormalities</td>
<td>Prognosis depends on the severity of renal disease</td>
<td>ARPKD, isolated MCKD</td>
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<tr>
<td>Joubert syndrome</td>
<td>Loci at 9q34, 17p11; AR</td>
<td>Renal changes of nephronophthisis often present in the subset of patients with retinal degeneration. Also associated with multicystic dysplastic kidney</td>
<td>Congenital hepatic fibrosis</td>
<td></td>
<td>Some patients die in infancy from breathing abnormalities; survivors have developmental delay with 25% developing renal insufficiency</td>
<td>SLS (NPHP)</td>
</tr>
<tr>
<td>Meckel-Gruber syndrome</td>
<td>Loci at 13q13, 17q21-24; AR</td>
<td>Antenatal ultrasound diagnosis: bilateral gross renal enlargement, variable size cysts. In the peripheral cortex the cysts are very small, thin walled, with a thin zone of normal glomerul, more centrally the cysts are larger. The largest cysts are in the medulla, up to several millimetres with thick fibromuscular walls separating them from collageneous stroma</td>
<td>Ductal plate malformation with periperal fibrosis and biliary dysgenesis</td>
<td>Occipital encephalocele, other CNS malformations, polydactyly, cardiac malformations</td>
<td>Lethal in the neonatal period from respiratory and renal insufficiency</td>
<td>Goldstain syndrome, ARPKD, Joubert syndrome</td>
</tr>
<tr>
<td>Bardet-Biedl syndrome</td>
<td>Genes BBS2, BBS4, BBS5; loci at 11q13, 3p13, 2q31; AR or possible tripellic inheritance</td>
<td>Persistent fetal labalation, small kidneys with clubbed calyces, cystic dysplasia; may be diagnosed on antenatal ultrasound scanning. Histology changes of nephronophthisis</td>
<td>Congenital hepatic fibrosis</td>
<td></td>
<td>Renal disease leads to a need for dialysis and transplantation during adolescence and early adulthood</td>
<td>Prader-Willi syndrome</td>
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<td>Jeune asphyxiating thoracic dysplasia</td>
<td>Possible locus at 12p; AR</td>
<td>Histology may show cystic renal tubular dysplasia, with or without glomerular sclerosis; could also have similiar findings to nephronophthisis</td>
<td>May have neonatal cholestasis, Caroli’s disease, and congenital hepatic fibrosis</td>
<td>Short stature, narrow, long thorax with short ribs, micromelia, other skeletal malformations, lung hypoplasia, retinal degeneration, pancreatic cysts and insufficiency</td>
<td>Severity of thoracic involvement variable and improves with age; patients who survive beyond infancy may develop renal involvement and ESRF, which determines prognosis</td>
<td>Ellis-van Creveld syndrome, short rib-polydactyly syndrome (SRPS) types I, II, III</td>
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AR, autosomal recessive; MCKD, medullary cystic kidney disease; MCDK, multicystic dysplastic kidney; SLS, Senior-Løken syndrome; ESRF, end stage renal failure.
severe form of ADPKD in comparison to chromosome 16p13, mainly adults, has an autosomal dominant inheritance and different age of onset, but otherwise resemble one another. Corticomedullary cysts, fibrosis, and progressive tubular atrophy are common features of MCKD. MCKD1 affects mainly adults, has an autosomal dominant inheritance pattern, and maps to chromosome 1q21. The juvenile form, MCKD2, has recently been shown to arise from mutations in the UMOD gene, encoding the glycoprotein uromodulin. In contrast, juvenile nephronophthisis is recessive and two genes (NPHP1 and NPHP4) have been identified recently. Two other loci have been described for autosomal recessive nephronophthisis that affect children of different age groups. It is interesting to note that patients with Senior-Loken syndrome (SLS), which is an association of nephronophthisis and retinal dystrophy, were found to have identical mutations in either the NPHP1 or NPHP4 genes to the affected children with purely renal involvement. This variable phenotype may indicate the action of some modifier locus, although the presence of intrafamilial variation suggests that all patients may have some degree of retinal involvement.

Both NPHP and MCKD share characteristic features, such as tubular atrophy, cyst development, and fibrosis of interstitial cells, although the kidneys are usually of normal size. Cyst formation tends to occur close to the papillary tips in MCKD and at the corticomedullary border in NPHP. Nephrocystin, the NPHP1 gene product (fig 3), appears to have a role in mediating cell adhesion. Functional studies of the murine homologue of nephrocystin indicate that it can interact with Crk associated substrate, p130(Cas), a protein known to participate in integrin mediated signal transduction and organisation of the actin cytoskeleton at sites of cell adhesion. Nephrocystin also contains a c-src homology 2 (SH2) domain, that is implicated in signal transduction (fig 3). In addition, nephrocystin has been shown to interact with nephrocystin-4, the protein product of the NPHP4 gene.
FIBROCYSTIC CHANGES AND MALFORMATION SYNDROMES

Skeletal malformation syndromes

Renal, hepatic, and pancreatic abnormalities are common features of Jeune asphyxiating thoracic dystrophy (JATD), although the consistent features are skeletal abnormalities, including a long and narrow thorax, metaphyseal irregularities, and shortness of the ribs and long bones. Polydactyly of both hands and feet is an occasional feature. Most severely affected cases have a fatal outcome in the perinatal period owing to asphyxia arising from a small thorax and hypoplastic lungs. Chronic renal failure occurs in the 20% of patients with JATD who survive beyond the neonatal period. Hepatic involvement can be severe, and can manifest as neonatal cholestasis or hepatic fibrosis, which leads to biliary cirrhosis and portal hypertension. JATD appears to overlap with other chondrodysplasias in a phenotypic spectrum (table 1).

There are particularly striking similarities between JATD and Ellis-van Creveld syndrome (EvC), which is characterised by short limbs, short ribs, postaxial polydactyly, and dysplastic nails and teeth. A differential diagnosis of JATD or EvC, on the basis of radiology alone, may not be possible. However, polydactyly, fingernail dysplasia, and congenital cardiac defects, usually a defect of primary atrial septation that forms a common atrium, are the primary features of EvC. In contrast, the main abnormality of JATD is renal. As for JATD, there are occasional developmental defects and dysplasias in the kidneys and liver for EvC. It has been suggested previously that JATD, EvC, and hepatic-pancreato-renal (HPRS) syndrome have overlapping features from one disease spectrum. The EvC gene has been mapped recently to chromosome 4p16 and the EvC protein has distant homology to the serine/threonine protein kinase family of proteins and contains a potential transmembrane region, but has at the present time no known function.

Bardet-Biedl syndrome (BBS) is an autosomal recessive disorder that displays pronounced phenotypic variability, with both developmental and progressive defects. The clinical manifestations are difficult to classify, but are characterised primarily by retinal dystrophy, obesity, and postaxial polydactyly. Renal structural abnormalities, such as multicystic dysplasia, are a minor clinical feature of BBS and may not give rise to clinical evidence of renal disease. At least seven BBS loci have been identified, and the BBS6, BBS2, and BBS4 genes have been identified. The BBS4 gene product has significant homology to O-linked N-acetylglucosamine transferases, and the BBS4 protein may glycosylate proteins to modulate signal transduction during normal development.

Central nervous system malformation syndromes

Johann F Meckel first described an unusual association of renal cysts, polydactyly, and posterior encephalocele in 1822. Subsequently, this association was studied further by Georg B Gruber who used the label “dysencephalia splanchnocystica”. Meckel-Gruber syndrome (MKS) is a lethal malformation syndrome characterised by a frequent

Figure 3  Schematic diagrams of the protein domain structures of polycystins, fibrocystin, and nephrocystin. Coloured boxes or circles on the main backbone of the proteins (thick black line) indicate conserved domains or regions that have a high degree of homology with known proteins. Annotation of domains and regions is based on primary published reports and conceptual translations of entries in the SwissProt and Genbank databases for the following accession numbers: NP_000287 (polycystin-1), Q13563 (polycystin-2), O15259 (nephrocystin), and AAL74290 (fibrocystin). The boundaries of selected regions are given as the number of amino acid residues from the N-terminus of the protein (N). The approximate location of other protein domains and putative N-glycosylation sites are based on output from the CDART tool at <www.ncbi.nlm.nih.gov/Structure/lexington/>. Putative transmembrane domains are indicated by grey rectangles, spanning the cell membrane (wavy horizontal line). Regions that have no apparent domains or conserved regions have been left blank.
FIBROCYSTIC CHANGES AND METABOLIC CONDITIONS

The association of renal cystic changes and liver disease is also a feature of a number of inherited metabolic disorders (table 2), and it is therefore important to differentiate these syndromes, which can be readily tested by biochemical means, from recessive HRFC syndromes. For example, Smith-Lemli-Opitz syndrome (SLOS) is caused by the deficiency of 7-dehydrocholesterol reductase (DHCR7), and the clinical features of SLOS include structural brain anomalies such as holoprosencephaly, microcephaly, cleft palate, syndactyly and postaxial polydactyly, cholestatic liver disease, and renal multicystic disease. Renal disease is characterised by microcytosis or tubular dilatations. It is particularly interesting that the pattern of anomalies seen for Smith-Lemli-Opitz syndrome (SLOS), such as cerebellar vermis aplasia and renal cystic changes, are similar to those for MKS. SLOS forms frustes subjects appear to have an even closer overlap with MKS. The discovery of the biochemical defect in this condition has led to the improved management of affected patients and emphasised the importance of sterols in embryogenesis, namely that cholesterol has a direct involvement in the hedgehog embryonic signalling pathway.75

Glutaric acidemia type II (GA2 or multiple acyl CoA dehydrogenase deficiency) is a mitochondrial disorder that results from a deficiency of either electron transfer flavoprotein (ETF)
or ETF-ubiquinone oxidoreductase (ETF-QO). Both are intermediate electron carriers to ubiquinone in the mitochondrial respiratory chain. The complete deficiency of ETF-QO is associated with multiple congenital anomalies, including cystic dysplasia of the kidneys, which has led to the misdiagnosis of MKS in the past. CNS anomalies include dysplasia of the cerebral cortex and abnormal neuronal migration, and liver histology typically shows microvesicular steatosis and fibrosis. Bile duct hypoplasia and cholestasis may also occur.

Zellweger syndrome (ZS), or cerebrohepatorenal syndrome, is the most severe variant of the Zellweger spectrum of disorders. Patients have severe neurological deficit, progressive hepatic and renal dysfunction, and skeletal abnormalities. Renal cysts can vary from glomerular microcysts to large cortical cysts of glomerular or tubular origin. The liver includes architectural abnormalities that disrupt the arrangement and spacing of portal and central areas, with multiple central veins in each hepatic lobule. There may be a paucity of the intrahepatic bile ducts, and either severe fibrosis or micronodular cirrhosis in more advanced cases.

**ANIMAL MODELS OF FIBROCYSTIC SYNDROMES**

A number of animal models of fibrocystic diseases have been developed, including models of polycystic diseases that involve the liver and biliary tree. Mice with an insertional mutation or gene targeting of the Pkd1 gene have a phenotype similar to human autosomal recessive polycystic kidney disease (ARPKD). The c pk (congenital polycystic kidneys) mutation on chromosome 12 in mice produces a lethal recessive form of PKD that is associated with hepatic cysts, and has been used as an animal model of human polycystic disease. The mouse c pk mutation is also associated with biliary ductal plate malformations, although the expression of the defect was modulated by the genetic background. A second, spontaneous recessive polycystic kidney mutation is jck (juvenile cystic kidneys), and a third, the bpk mutation in BALB/c mice, associates with renal cysts and epithelial hyperplasia of the intra- and extrahepatic biliary tracts.

Transgenic mouse embryos that overexpress keratinocyte growth factor (KGF) develop hepatomegaly, biliary hyperplasia, and cystic dilatation of renal collecting tubules resulting in polycystic kidneys. The correct expression of KGF is thought to be important for the mesenchymal-epithelial signalling required during normal embryogenesis. However, transgenic mice that overexpress the c-myc proto-oncogene, a regulator of cell proliferation, develop polycystic kidneys, as do knock out mice homozygous for deletions of the bcl-2 gene, which normally regulates apoptosis. In the latter two model systems, disruption of apoptosis or proliferation would be a reasonable primary cause of cystogenesis. However, the renal cystic abnormalities could equally arise as a pleiotropic effect of the genetic lesion in the mouse model. Until the molecular mechanisms that regulate epithelial differentiation and the balance of cell proliferation or cell death are known in more detail, it is difficult to assess the importance of the phenotypes in these models.

**DISCUSSION**

In this review, we have compared the variable pathological processes and phenotypes that are associated with several recessive HRFC syndromes. Several reviews have postulated a spectrum of “fibrocystic syndromes” that have a more or less similar aetiology, but that then subsequently diverge in terms of disease progression and clinical outcome, with variable presentations of fibrosis, hepatocyte damage, cirrhosis, nephritis, and end stage organ failure. The suggestion has been made that the initial phenotypic similarity of HRFC syndromes, at least in terms of histopathology, may arise from similar causative defects in the embryological pathways of liver and kidney development that define tubular structures in epithelial cells. Developmental defects could therefore involve the disruption of cell-cell or cell-matrix interactions of epithelium and mesenchyme. In particular, biliary dysgenesis appears to be a key manifestation that distinguishes the HRFC syndromes (table 1) from the fibrocystic changes of metabolic conditions. Although renal cystic dysplasia is a common (but not obligate) feature of many conditions (table 1), the co-occurrence of biliary dysgenesis, in a form suggestive of ARPKD, appears to be limited to the group of HRFC syndromes that are inherited in an autosomal recessive pattern. In contrast, biliary dysgenesis is not a feature of metabolic conditions, but congenital hepatic fibrosis, cirrhosis, and renal cystic dysplasia are common end points in the progression of metabolic conditions (table 2). This probably indicates that the latter features are the result of any form of insult to the liver and kidney, whether it is congenital or acquired. Renal cyst formation, for example, could be induced by an energy deficit in the tubular epithelial cells, perhaps by disturbing the balance of cell proliferation and apoptosis, rather than by a specific defect in a developmental pathway. Clinical manifestations in Smith-Lemli-Opitz syndrome (table 2) could be a result of cholesterol deficiency during fetal development.

Three plausible molecular mechanisms could explain the common features of HRFC syndromes: (1) the diverse proteins participate in a default developmental pathway for tubulo-epithelial differentiation during embryogenesis of different renal systems, and that this process is perturbed by a number of genetic lesions; (2) different proteins interact directly with each other to mediate either adhesion or signal transduction events, or to form a multi-subunit complex; (3) different proteins participate in distinct but complementary pathways. These hypotheses are now testable following the recent identification of the Pkh6/1, Pkd1, Pkd2, Nphp1, and Nphp4 genes. The encoded proteins (fibrocystin, polycystin-1 and polycystin-2, nephrocystin and nephrocystin-4, fig 3) appear to have pivotal functional roles in renal epithelial differentiation and organisation by mediating interactions between epithelial cells and interstitial components.

Defects in direct protein-protein interactions are now thought to be the primary cause of the cystic changes of the various types of polycystic kidney diseases and juvenile nephronophthisis (NPHP), but the precise mechanisms remain unclear at present. The loss of contact points and signalling events between cells and either the extracellular matrix (at focal adhesions) or other cells (at adherens junctions) may be the defect that underlies NPHP. Nephrocystin, the product of the NPHP1 gene (NPHP1), appears to have a fundamental role in mediating cell adhesion. In addition, the C-terminal region of polycystin-1 has been shown to form multi-subunit complexes with focal adhesion proteins that include the structural and actin binding proteins vinculin, talin, tensin, and α-actinin, the adaptor proteins paxillin and p130(Cas), and the signalling kinase c-src. Polycystin-1 is also thought to form a heterodimeric ion channel with polycystin-2 (see above). The complex of focal adhesion proteins links the extracellular matrix to the actin cytoskeleton through the cell membrane. It is possible that mutations in other proteins of the polycystin-1 complex may lead to either cystic kidney disease or additional extrarenal fibrocystic changes. Familial focal segmental glomerulosclerosis, for example, is thought to be caused by mutations in the focal adhesion protein, α-actinin-4. The very recent demonstration, by a series of immunoprecipitation experiments, that nephrocystin and nephrocystin-4 interact directly suggests that these proteins are either in the same multi-subunit complex, or interact in a developmental pathway to mediate either adhesion or signal transduction events. Defects in either of the proteins will therefore result in NPHP. A similar molecular explanation has been suggested for the existence of at least six
loci or genes for Bardet-Biedl syndrome.\textsuperscript{1} BB\textsubscript{56} encodes a protein that has significant sequence homologies to a class of chaperonins that facilitate protein folding, and other BB\textsubscript{5} proteins could either form part of a chaperonin complex or interact with BB\textsubscript{56} while still unfolded polypeptides. It will be interesting to determine the causes of genetic heterogeneity in Meckel-Gruber syndrome and Joubert syndrome: do the different encoded proteins interact or participate in the same developmental pathways?

In the future, further study of the structure and function of known proteins, such as fibrocytin and nephrocytin, will contribute to our knowledge of renal cystic dysplasia and biliary dysgenesis. It will then be clear whether or not a common pathogenetic pathway is involved in the range of congenital fibrocystic syndromes, and perhaps provide molecular mechanisms to explain the variability in the rates of disease progression and the severity of clinical phenotype. Understanding these mechanisms may allow novel pharmacological interventions or gene therapy procedures to be designed that would prevent renal cyst development, dysgenesis of the intrahepatic biliary tree, and hepatic fibrosis.

CONCLUSION

This review has identified a number of hepatoportal fibrocystic (HRFC) syndromes that are a group of severe malformation syndromes that cause fibrocystic changes in the liver and kidney, and that are inherited in an autosomal recessive manner. Biliary dysgenesis is a common feature of these conditions, which appears to differentiate them from metabolic conditions with fibrocystic changes. This diversity emphasises the common nature of the embryogenesis in different organ systems, and suggests that tubulo-epithelial differentiation in the kidney and intrahepatic biliary system has a common molecular mechanism. The possible functions of the gene products in these pathways may involve signal transduction by both the extracellular matrix and soluble growth factors, and regulation of epithelial cell proliferation and complex epithelial-mesenchymal interactions that are disrupted in HRFC syndromes. The identification of these genes, and the assessment of the role of mutations, will be an important step in describing critical developmental pathways in molecular detail. Recent work has identified a number of integral membrane proteins that appear to participate in these pathways, but their exact molecular functions remain to be elucidated.

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