REVIEW ARTICLE

A. Mortell · S. Montedonico · P. Puri Animal models in pediatric surgery

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Introduction

The advent of animal models of pediatric surgical diseases has not only allowed us to study the etiology and pathogenesis of complex congenital anomalies but has also led to major advances in the surgical and therapeutic management of these conditions. Even though representative animal models do not exist for every human disease, the animal models available to us today are still of great importance for medical research. Animal models allow us to comprehend the molecular and biochemical basis of diseases, and possibly find new drugs and forms of therapy, both medical and surgical. Different animal models have been and may be used. Because of their easy availability, smaller animals such as rodents are often preferred as models. Mice (Mus musculus) offer many advantages. They are similar to humans in a biological and genetic manner, so that much, but not all, of the knowledge gained from studies in mice may be applied to humans. The genome of the mouse is similar to the human genome, with regard to the size and number of genes. Mice have a very high reproduction rate, a short life span and mature quickly. This makes it possible to easily follow the effects of changed genes over many generations. At the present time, mice have become the most important and common animal models for human diseases. For many years the mouse genome has been closely investigated, and many mutations, which have either spontaneously developed in a population or were induced through outer influences (for example chemical mutagenesis) have been demonstrated.

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A. Mortell · S. Montedonico · P. Puri (⊠) Children's Research Centre, Our Lady's Hospital for Sick Children, Crumlin, Dublin 12, Ireland E-mail: prem.puri@ucd.ie Tel.: + 353-1-4096420 Fax: + 353-1-4550201 Many international co-operations work on methods of describing the function of each single genome with different strategies. Rats (*Rattus norvegicus*) are also widely used in research, as they are particularly suitable for physiological, pharmacological and behavioral research. They have also contributed greatly to research involving surgical techniques and the study of many teratogens.

Animal models are of great importance for the analysis of genes. With their help it is possible to set up models for the examination of gene expression and gene regulation. Genes may be specifically inactivated within animals (knockout model), and the effect on the organism can be studied. By inserting an additional recombinant gene (transgenic animal model), its expression and influence on the metabolism of the organism can be examined. The setting up of expression analysis is only part of solving the functions of genes in an organism or in tissues. Since genes generally work in coordination, thus defining the characteristics of an organism, it is necessary to examine the function of the genes in connection with the organism under physiological conditions.

Broadly speaking, there are four main types of animal models in use today: naturally occurring, teratogen induced, surgically created and transgenic animals. Obviously, the animals in which diseases occur naturally are the ideal models with which to study disease pathogenesis, as there is little or no interference to the animal prior to the study. The ground-breaking development of transgenic animals has not only allowed researchers to almost mimic the natural occurrence of certain conditions but also gives insight into the specific genes involved in the regulation of the disease process and how their modification might alter the course of the disease. Teratogen-induced models, although useful, have the drawback of exposing the animal to a generalized noxious stimulus, which can result in widespread detrimental effects rather than simply targeting a specific organ system. Finally, surgical models can closely simulate surgical procedures and conditions found in hu-

Following from these models, research has sought to develop potential therapeutic approaches, including surgical interventions, which can be employed in the management of many conditions, and as a result many different models for several important human diseases have been developed over many years of pediatric surgical research. Lesser known animal models not discussed in this review include those for spina bifida (curly tail and delayed splotch mouse models, surgical rat and lamb models), hydrocephalus (HTX rat, HSV and L1 knockout mouse models), cloacal exstrophy (suramin chick and surgical lamb model), cryptorchidism (Insulin-3 and HOXA 11 knockout mice, flutamide rat, surgical rat and rabbit models), biliary atresia (rotavirus mouse model), necrotizing enterocolitis (hypoxia/lipopolysaccharide and casein piglet model, surgical rabbit model, hypoxic rat model), short bowel syndrome (surgical mouse model), vesico-ureteral reflux (surgical piglet model) and pyloric stenosis (mutant hph-1 mouse model). Research into important and challenging neonatal conditions such as gastroschisis (GS), Hirschsprung's disease (HD), VACTERL association and congenital diaphragmatic hernia (CDH) has been greatly advanced by the ready availability of animal models and will form the basis of the discussion in this review.

Animal models of the VATER association

The vertebral, anorectal, tracheo-esophageal, renal (VATER) association is a spectrum of clinical conditions in the human neonate, which involves multiple anomalies first reported by Quan [1]. The exact incidence of VATER is difficult to quantify in view of the fact that there is such variation in clinical presentation, but it affects approximately 1 in 5,000 live births and often requires urgent surgical intervention after birth due to the foregut and hindgut anomalies. VATER has not been recognized as a specific syndrome in humans but rather represents a non-random association of congenital anomalies of poorly understood etiology and pathogenesis and its components have been variable. The most important features of the VATER association, which will be discussed in this paper from a pediatric surgical point of view, include anorectal malformations, esophageal atresia (EA), tracheoesophageal fistula (TEF), and renal anomalies. Etiologically, the VATER association may be a feature of some chromosomal anomalies [2] but the majority of cases have no recognized cause. The higher incidence of malformations occurring in association with each other than might be thought to occur randomly prompted some to suggest that the VATER association arose as a result of an insult to the developing embryo early in its development which led to abnormalities in diffuse organ systems at birth. Khoury [3] suggested that the VATER association arose as a result of defective mesodermal development during early embryogenesis, leading to abnormalities in diffuse organ systems at birth. Others have suggested that teratogens, such as the maternal use of hormones during pregnancy may have a role in bringing about the association [4].

Four types of animal model of the main surgical anomalies found in the VATER association have been developed: teratogen induced, surgically created, knockout models and naturally occurring.

Teratogen-induced models of VATER

The Adriamycin rat model (ARM) of the VATER association is a well-established model, which has provided us with a unique opportunity to investigate the mechanisms behind the origins of this congenital anomaly and its associated conditions. Similarities with the human pattern made the Adriamycin rat a useful model for pathogenetic analysis of these defects. Adriamycin (doxorubicin hydrochloride) is an anthracycline, anti-neoplastic drug that has been widely used in the treatment of many malignancies, including sarcomas, lymphomas, neuroblastoma and breast cancer [5]. One proposed mechanism of action lies in its ability to inhibit nucleic acid synthesis through intercalation with DNA [6].

The teratogenicity of Adriamycin was first established by Thompson et al. [7] in 1978, when the intraperitoneal administration of 1-2 mg/kg/dayof Adriamycin to pregnant rats from gestational days (GD) 6-9 resulted in characteristic malformations, including esophageal and intestinal atresia, TEF, hypoplasia of the urinary bladder and cardiovascular malformations. The first report of the Adriamycin rat as a pediatric surgical model for anorectal malformations, EA and TEF and renal anomalies came in 1996 from Diez-Pardo et al. [8]. With the administration of 1.5-2 mg/kg/day on GD 6-9 they achieved significant rates of EA with TEF (45%), anorectal anomalies (50%) and renal anomalies (100%) and this has prompted a huge volume of research to be carried out since then.

The role of neuropeptides and innervation in the esophagus of the ARM was looked at early on as a possible cause for esophageal dysmotility. The vagus and recurrent laryngeal nerves in ARM embryos were found to have inherent abnormalities in their course and culminated in a deficient extrinsic nerve fiber plexus in the lower esophagus [9]. Elevated levels of neuropeptides of the enteric nervous system, such as S100 and galanin were found in the distal esophagus of experimental embryos compared to controls [10]. Also the density of the nerve plexus, ganglia and number of cell bodies per ganglion immunostained by neuron-specific enolase, vasoactive intestinal peptide and substance P was significantly reduced in EA-TEF fetuses [11]. These findings were all suggested as contributing factors to dysmotility in patients with EA.

The role of apoptosis has been studied as a potential mechanism for the development of tracheoesophageal abnormalities in the ARM. Initial studies demonstrated a specific and consistent pattern of apoptosis or cell death in the region of the so-called "tracheoesophageal septum" that might have contributed to the normal process of tracheoesophageal separation [12, 13]. Subsequent studies did not demonstrate significant apoptosis in the immediate period following Adriamycin administration, suggesting that the teratogenic effect of Adriamycin is not caused by cell death [14, 15].

The contribution of the notochord to normal embryonic development has been well established and it has been shown to play a role in the patterning of mesoderm as well as regulating endodermal development [16, 17]. The notochord has been found to be grossly abnormal in ARM embryos with foregut anomalies, such as EA and TEF [18]. The various abnormalities of the notochord include accessory ventral branching, bending, splitting and tethering to foregut structures with ventral displacement of the notochord and dorsal displacement of the foregut [19-21]. Abnormalities of Sonic hedgehog (Shh), an important genetic signaling molecule, have also been implicated in the pathogenesis of the anomalies seen in the rat model of VATER. Studies have shown high levels of Shh protein expression in the normal rat foregut during embryogenesis, which slowly declines as the embryo approaches term [22]. In contrast, Adriamycin-treated embryos demonstrated a low level of Shh expression throughout the same time period. In a quantitative morphological study of the notochord, Mortell et al. [23] demonstrated a significant increase in the relative notochord volume of Adriamycin-treated embryos compared to controls. Immunohistochemically, the accessory branches of notochord, which contributed to this increased volume, expressed Shh similar to control notochord. Interestingly, the increase in notochord volume, relative to embryo volume, was found to be maximal soon after Adriamycin administration in experimental embryos [24]. This early alteration in notochord volume, with associated Shh expression, during a critical phase of embryonic development was thought to contribute to the malformations found in the VATER association.

The development of anorectal malformations in the ARM has also been of great interest to pediatric surgeons engaged in research. Although less frequently seen than EA with TEF, anorectal malformations occur in 24–28% of Adriamycin-treated embryos [25, 26]. Anorectal malformations have also been produced with the application of other teratogens to either rodent or murine models. Rat embryos prenatally exposed to a single dose of oral retinoic acid on GD 10 have been shown to develop anorectal anomalies in 94% of the cases [27]. The most common defect noted was anal atresia with or without cloaca and in association with these defects, abnormalities of the tail were commonly found. Interestingly, spina bifida was also noted in 55% of the embryos studied. In pregnant mice treated with oral retinoic acid as a single dose on one of the GD 8 or 9, a large number of their offspring developed multiple anomalies [28]. These consisted of urogenital malformations, anal agenesis and rectourethral fistula. More than 95% of murine embryos treated with retinoic acid developed anorectal malformations, with rectoprostatic urethral fistula, rectocloacal fistula and a short tail being commonly found [29].

As a teratogen, ethylenethiourea (ETU) has been used to induce anorectal malformations since the early 1990s [30]. The offspring of pregnant rats, which are gavage fed 1% ETU (125 mg/kg) on GD 10, develop anal atresia with a rectourethral fistula in up to 85% of embryos [31, 32]. These abnormalities seem to be caused by aberrations in the development of several components of the hindgut and cloaca. Abnormalities of the notochord, similar to those noted in the ARM, were found in this model and suggest that abnormal notochord development may be pivotal in producing anorectal malformations [32].

Renal defects are seen in approximately 50% of patients with the VACTERL association. The most common renal defects seen in humans with VATER/ VACTERL association include renal agenesis, cystic dysplasia, renal ectopia, horseshoe kidney, persistent urachus, and urethral agenesis [33]. These problems can lead to significant morbidity and progressive renal deterioration early in life and may sometimes require renal transplantation. Renal anomalies form a significant part of the spectrum of abnormalities seen in the ARM of the VATER association. These findings have allowed researchers to intensively study the pathogenesis of urinary tract anomalies, which can affect human neonates. Although bladder agenesis is rare in humans, it can be developed experimentally in up to 100% of embryos receiving 2 mg/kg/day of Adriamycin on GD 6-9. This study suggests that GD 7 is the critical embryological window for bladder development in rats [34]. In association with bladder agenesis, all embryos had significant unilateral or bilateral hydronephrosis/ hydroureter, presumably as a result of distal obstruction [35]. Further work determined that primary agenesis rather than secondary resorption of the bladder occurred in this model [36]. Microscopically, affected kidneys have abnormal architecture, with an abnormal nephron configuration, decreased tubular differentiation and a thinned medulla [37]. Dilatation was noted mainly in the collecting system, ducts and tubules, with no evidence of collagen deposition or fibrosis. A further development of this animal model, using a lower dose (1.75 mg/kg/day) of Adriamycin on GD 7–9, led to the findings of bilateral hydronephroureter with preservation of the bladder and this has allowed for a more physiological study of the renal tract, as might be found in the human setting [38].

The availability of murine molecular probes and genetic strains of knockout mouse models of EA and TEF

has prompted researchers to try and develop a new murine model incorporating the known teratogenic effects of Adriamycin [39]. Although this model has not been fully developed to yield the same spectrum of anomalies seen in the ARM, early results have shown tracheoesophageal anomalies in 47% of embryos from pregnant CBA/Ca mice treated with 4 mg/kg/dav of Adriamycin on GD 7.5 and 8.5. Further work demonstrated immunohistochemical evidence of maintained Nkx2.1 expression (similar to that seen in controls) in the respiratory element of Adriamycin-treated embryos despite their abnormal morphogenesis [40]. The authors concluded that the dorsal fistula retains its non-respiratory commitment, in the absence of tracheoesophageal separation, suggesting that dorso-ventral patterning of foregut development is undisturbed by Adriamycin exposure. The authors then studied the expression of Sonic hedgehog (Shh) and Patched (Ptc1) in the foregut of developing mouse embryos, which had been treated with Adriamycin [40]. Shh is a secreted glycoprotein, which is known to play an important patterning role in the development of many organ systems including the foregut. *Ptc1* is a transmembrane receptor, which acts as a target gene for *Shh* [41]. Their results showed a reversal in the dorso-ventral pattern of Shh expression during the brief period of tracheoesophageal separation, with the foregut of Adriamycin-treated embryos diffusely expressing Shh [40]. They suggested that a transient disturbance in *Shh* expression at this critical time might contribute to the abnormal organogenesis in this model.

Surgical models of EA and TEF

Although the chick embryo has not provided us with a reliable model of EA and TEF, its contribution to our knowledge of the normal development of the esophagus and trachea cannot be ignored. The theory of foregut hyperflexion had been proposed as a cause of EA and distal TEF as far back as 1976 [42]. However, this theory has never been proven. Kleckner et al. [43] performed hyperflexion studies on chick embryos in order to test this theory. Although one case of an "H-type" TEF resulted from their study, no cases of EA developed. This isolated case was felt to possibly represent a chance finding of little significance and therefore the theory was disregarded.

Embryological studies in the chick have endeavored to look at the primitive foregut during the period of differentiation, as previous work has suggested that the tracheoesophageal septum plays a significant role in the division of the foregut into trachea and esophagus. Kluth et al. [44] studied the foregut region of chick embryos from Hamburger and Hamilton [45] stages 14–26 by scanning electron microscopy and light microscopy. They found no evidence of lateral folds or fusing foregut wall components, which might have contributed to a tracheoesophageal septum. Their findings suggested that esophageal and tracheal development results from a simple reduction of the size of the primitive foregut.

Although the vast majority of animals used for research purposes today are rodents, canine models of EA have been studied as far back as the 1970s when Oh [46] described a microsurgical technique for esophageal reconstruction in puppies. Subsequently, beagles were widely used to study the post-operative results following esophageal resection or transection with subsequent reanastomosis, utilizing an autologous jejunal mucosa transplant [47], a tubular musculopleural pedicle graft [48] or a variety of myotomies with or without delayed esophageal reconstruction [49–51]. The results from these studies were variable with no clear advantage for one particular technique. Although early contrast and manometric studies suggested a good outcome from spiral myotomy, long-term follow-up in this group demonstrated esophageal dysmotility, as observed in children following esophageal surgery. However, further studies in canine models of EA compared manometric findings following esophageal transection and re-anastomosis versus esophageal vagotomy alone [52]. Their results showed coordinated peristaltic contractions between the proximal and distal esophagus in the first group with abnormal simultaneous contractions in the vagotomized group. This suggested that post-operative dysmotility might arise from disruption of the vagus nerve as part of the congenital abnormality or secondary to surgical trauma.

An interesting paper by Carachi et al. [53] looked at the effect of an indwelling silicone transanastomotic tube (TAT) on the healing esophageal anastomosis in puppies. They found no significant difference in stenosis rates whether a TAT was used or not. Interestingly, they observed a shelf of stenotic tissue on the posterior wall of the esophagus at the site where the intraluminal silk knots had been tied, compared to a thin linear scar on the anterior wall, where knots had been tied extraluminally.

More recently, the use of a tubed latissimus dorsi musculocutaneous flap (LDMCF) for esophageal replacement has been studied in canine puppies [54, 55]. Long-term follow-up of these animals showed good survival rates with only occasional vomiting. Although contrast studies showed no evidence of stenosis, endoscopy demonstrated hair re-growth intraluminally, with no evidence of abnormal histological changes on sectioning of the LDMCF.

Most recently Tannuri et al. [56, 57] have studied the effects of circular myotomy (Livaditis) on anastomotic leak rates and anastomotic healing in dogs subjected to esophageal resection and an anastomosis under tension. They found an equal leak rate between groups with an anastomosis under tension and those with an additional circular myotomy. Morphometric and biochemical analysis showed reduced numbers of small vessels and collagen type V in the fibrous scars of the myotomy animals. The authors concluded that circular myotomy does not reduce the possibility of anastomotic leaks but

does promote deleterious changes in anastomotic healing.

Porcine (piglet) models of EA have been very useful for pediatric surgical research purposes because of their size and anatomical similarity to human neonates. Initial studies involved the evaluation of surgical techniques for esophageal replacement comparing gastric tube interposition with gastric tube in continuity and gastric transposition [58]. Gastric transposition gave very poor results with 100% mortality within 96 h. Gastric tube interposition was associated with improved growth, fewer clinical complications (such as vomiting, minor leak and stricture) and fewer histological changes (such as esophagitis or submucosal fibrosis) than either gastric tube in continuity (with or without posterior fundoplication) or gastric transposition.

The effects of a circular myotomy (to achieve esophageal elongation in long gap atresia) have sometimes been detrimental to patients, with muscular defects allowing ballooning of the esophagus post-operatively [59]. To try and prevent this adverse occurrence, Komuro et al. [60] applied collagen sponge scaffolds to the muscular defect in myotomized piglet esophagus. Although the loose connective tissue covering the defect did not contain muscle, when it was examined histologically, it did prevent distension of the esophagus at the point of myotomy.

The advent of robotic-assisted surgery has prompted a great volume of research in animal models of surgical disease. The advantages are many, but the techniques are not without their drawbacks. Although robots can filter out the fine tremor of a surgeon's hand and add an extra element of flexibility with intracorporeal instrument angulation and higher magnification, the prolonged set up time and added expense can detract from the overall usefulness of certain robots. Thoracoscopic repair of EA [61] has become a recognized method for dealing with this complex surgical problem and studies involving robotic-assisted thoracoscopic repair of EA in piglets have shown that the technique is both technically feasible and that it may facilitate a minimal approach in the small chest of a newborn [62, 63]. A recent interesting study looked at a novel approach to thoracoscopic esophageal anastomosis using a U-clip suturing technique [64]. The self-ligating metal U-clips were easily placed and appeared to be a feasible alternative to sutures but the study only involved a small number of piglets and therefore warrants further research to assess the technique more comprehensively.

A surgical rat model has recently been developed to examine the possibility of esophageal transplantation for use as an esophageal replacement. Initial studies by Yamataka et al. [65] looked at syngeneic esophageal transplantation into an omental pouch and found the transplants to be viable and well vascularized after a period of 10 days. They also found that the grafts could be easily mobilized into the thoracic cavity on an omental pedicle without tension. Following this study they employed allogeneic esophageal transplantation from newborn Brown-Norway rats into 5-week-old Lewis rats and studied the effects on the neo-esophagus with and without immunosuppression using FK-506 [66]. Syngeneic transplants were used as controls and showed no evidence of rejection with 100% graft survival. Grafts that received between 0.6 and 1.2 mg/kg of FK-506 for the duration of the study period showed minimal or no rejection. However, grafts receiving no immunosuppression, low dose FK-506 or immunosuppression for a shorter period showed definite histological evidence of rejection.

Knockout models of VATER

Faulty genetic signaling has often been implicated in the etiology of the VATER association. The hedgehog (Hh) family of signaling molecules mediates many inductive processes during invertebrate and vertebrate development. As previously mentioned, Sonic hedgehog (Shh) is a genetic signaling molecule crucial to normal embryonic development, which is released from the notochord and acts in a concentration dependant manner [67]. Shh has been shown to be involved in the control of cell fate determination in the central nervous system, dorsoventral patterning of somites, anteroposterior polarity of limbs and left-right asymmetry. In the developing embryo, the *Shh* pathway is essential for correct patterning of the various organ systems [68-70], including hindgut patterning [71] and lung growth [72], and is responsible for radial patterning of the gut wall by inducing inner components and by inhibiting outer components, such as smooth muscle and neurons [73]. In mutant mice with deletions of the Shh gene, VATER anomalies are commonly found [67, 74]. Homozygous Shh-null mutant mice demonstrate EA/stenosis, TEF and tracheal and lung abnormalities, similar to those seen in human neonates with foregut defects [75]. Anorectal malformations have also been shown in Shh-null mice with mutants displaying a persistant cloaca, with the distal urinary tract and anorectum sharing a common outlet [76].

Other factors implicated in *Shh*'s patterning of the embryo include the *Shh* membrane-bound protein receptors Patched (*Ptc*), Smoothened (*Smo*) and the zinc-finger transcription factors *Gli1*, *Gli2*, and *Gli3*, which are also released from the embryonic notochord and act as mediators in the nuclear transcriptional activity of *Shh* signaling [77]. These genes are known to operate in a downstream manner to *Shh* and have been shown to have an essential function in the formation of lung, trachea, esophagus and ano-rectum in mice [76].

Mutant mice lacking *Gli2* function exhibit the full spectrum of VACTERL anomalies, except renal anomalies. These include foregut defects, such as esophageal and tracheal stenosis as well as lung hypoplasia [74]. All *Gli2* null-mutants displayed an imperforate anus with recto-urethral or recto-vaginal fistula [78]. *Gli3*

null-mutants have been shown to develop vertebral, anal and limb anomalies. However, Gli3 null-mutant mice with a 50% reduction in the gene dosage of Gli2 developed EA and TEF with very poor lung development. Mice lacking both Gli2 and Gli3 did not develop esophagus, trachea or lung [76, 79].

NKX2.1 (otherwise known as *TTF-1*) is a homeodomain transcriptional factor expressed in thyroid, lungs and parts of the brain. Mouse embryos carrying a homozygous targeted disruption of the *Nkx2.1* locus demonstrate failure of septation of the anterior foregut along the dorsoventral axis, into distinct tracheal and esophageal structures [80]. This failure of septation results in a common lumen connecting the pharynx to the stomach, similar in phenotype to the human pathological condition of TEF. Further abnormalities in this knockout model include severe lung hypoplasia, with associated poor gas exchange, resulting in early postnatal death.

Naturally occurring models of anorectal malformations

Two naturally occurring animal models have long been used to study anorectal malformations, such as those seen in the VATER association. The Sd mouse, first bred by Danforth in 1930 [81], is a mutant of the normal house mouse and was found to develop urogenital and anorectal abnormalities [82]. Also known as "Danforth's short tail mouse", this model has a number of advantages because of the high percentage of abnormal animals per litter, the relative inexpense of a mouse model and the ease with which Sd mice can be bred. These mice have an autosomal-dominant trait for imperforate anus and pass on this semi-dominant trait with high penetrance, where all of the homozygotes and most of the heterozygotes are affected. Kluth et al. [83] looked at the spectrum of anorectal anomalies in this model and found that the pathological anatomical malformations in the heterozygous Sd mouse group were very similar to those seen in human neonates with anorectal anomalies as well as previous porcine models, which have been described. To date, the gene defective in Sd mice has yet to be identified.

Domestic pigs have a naturally high incidence of anorectal malformations and can be bred to generate an animal model to study the anatomy and embryology of anorectal anomalies [84]. Recent studies have developed a pig pedigree over 15 years, resulting in the development of non-syndromal anal atresia in up to 60% of newborn piglets [85]. This resource was generated by surgically treating affected piglets soon after birth and later breeding them for a number of generations. Utilizing marker analysis, the authors attempted to map susceptibility loci and found a suggestive locus on pig chromosome 15 [86]. Previous morphological studies on the internal sphincter in neonatal piglets with anorectal malformations showed that all animals had a normal internal sphincter surrounding the proximal fistula [84]. They, therefore, suggested that the fistula should be designated as an ectopic anal canal and that the incorporation of the fistula into the surgical correction of anorectal malformations may yield better results from a continence point of view.

Animal models of Hirschsprung's disease

Hirschsprung's disease (HD) is a common cause of intestinal obstruction in the newborn with an incidence estimated to be 1 in 5,000 live births. Animal models of HD have been pivotal in the understanding of the anatomy and pathophysiology of the disease and in the discovery of the genes involved in HD. Four types of HD animal models can be distinguished: natural mutants, knockout models, surgical and chemical models.

Natural mutants

Aganglionic megacolon not only occurs in humans but also affects several other species such as mice, rats and horses [87–90]. All of these mutations not only involve the intestine but another neural crest derivative, the melanocytes, since all have major areas of unpigmented skin and hair. In 1966 Lane described two strains of mice, which developed aganglionosis as an autosomal recessive condition [87]. The lethal spotting (ls) mice have approximately 2 mm of aganglionosis with a patched coat. Piebald lethal (s^{I}) mice have approximately 10 mm of aganglionosis. Later studies linked the defect to chromosome 2 and 14, respectively. In 1979 Ikadai et al. [89] described aganglionosis in rats and 80–90% of the rats had a total colonic aganglionosis. The animals had a high mortality rate and were only able to survive 3–4 weeks of age due to a severe bowel obstruction and enterocolitis. These animals showed autosomal recessive inheritance and had a white colored coat. A fourth rodent model is the dominant megacolon (Dom) mouse that was described by Lane and Liu in 1984. As the name suggests the inheritance is dominant [88]. These mice are characterized by distal colonic aganglionosis and a long hypoganglionic transition zone, but occasionally the entire large intestine is aganglionic. The allele for this mutation has been mapped on chromosome 15. Histological studies in all these rodent models using acetylcholinesterase whole-mounts demonstrated that they have a histological picture identical to the human. There is distal aganglionosis with increased nerve trunks, followed by a transition zone, which is often asymmetrical and variable in length and more proximal the plexus is near normal [91]. Many electrophysiological studies have been done in rodent models of aganglionosis. It has been demonstrated that there are abnormal discharges of myogenic action potentials in the aganglionic bowel associated with tonic constriction and a reduction in the luminal diameter [92] and that the aganglionic bowel has no inhibitory junction potentials,

which leads it to writhe in an uncontrolled manner tending to contraction. Therefore, the rodent models have helped us to clarify that the simple lack of nerve fibers is sufficient explanation for the functional obstruction seen in patients with HD and that there is no need to invoke selective disorders of various components of the autonomic nervous system to account for the clinical variability [91]. The rodent models of aganglionosis have also contributed to the understanding of the embryology of HD. Cass et al. [93] studied the migration of the neural crest cells to the gut in the lethal spotting mouse, the piebald mouse and the spotting lethal rat. They concluded that there was a slowing of migration in those neural crest cells derived from the vagal neural crest and that there were small numbers of enteric precursors in the aganglionic distal hindgut derived from the sacral neural crest that did not differ in number compared to normal embryos. Although they demonstrated that there is sacral neural crest contribution into the enteric nervous system, this contribution was shown to be functionally insignificant.

The mutations which give rise to these natural mutant mouse strains have now been identified as being the endothelin receptor-B gene (ENDR-B) for the piebald lethal mouse, the gene for endothelin-3 (ET-3) in the lethal spotting mouse and the Sox10 gene in the *Dom* mouse [94–97].

Knockout models for HD

Transgenic technology, particularly in the mouse, provides a powerful method with which to produce animal models to determine the function of genes associated with human inherited disease. Several genes have been disrupted in mice, producing phenotypes remarkably similar to HD in humans. In addition, three well-studied mutant mouse models of HD have been found to be due to disruptions of specific genes, as stated before. The Ret gene encodes a receptor tyrosine kinase, which has four ligands: glial cell line derived growth factor (GDNF), neurturin (NTN), artemin (ATM) and persephin (PSP) [98]. The complete receptor complex includes the *Ret* receptor tyrosine kinase and a glycosylphosphatidylinositol-anchored binding component ($gfr\alpha 1$, $gfr\alpha 2$, $gfr\alpha 3$) or $gfr\alpha 4$). This receptor has been suggested to function as an adhesion molecule, which is required for neural crest migration and that could also play a role in either differentiation or survival of the neural crest cells which have stopped migrating [99, 100]. Ret (-/-) transgenic mice have a homozygous, targeted mutation of the tyrosine kinase receptor resulting in a loss of its function. These mice exhibit total intestinal aganglionosis and renal agenesis [101]. The *Ret* gene has been demonstrated to be a major gene causing HD in humans. Mutations of Ret account for 50% of familial and 15-20% of sporadic cases of HD [102–105]. GDNF, one of the *Ret* receptor ligands, stimulates the proliferation and survival of neural crest derived precursor cells in the embryonic gut [106, 107]. Mice homozygous for null mutation in *Ret*, *GDNF* and *gfra1* have almost identical phenotypes characterized by failure of enteric nervous system development distal to the esophagus and absent kidneys [101, 108–112]. Although a causative role for *GDNF* mutations in some patients with HD has been suggested, the occurrence of such cases is uncommon and it is more likely that the *GDNF* mutations are involved via its interaction with the *Ret* receptor [113, 114]. No *gfra1* mutations have been identified in patients with HD [115].

The endothelins are intercellular local messengers that comprise four members to date: ET-1, ET-2, ET-3 and VIP. They transduce a signal via two cell surface transmembrane receptors: ENDR-A and ENDR-B [98]. In human fetuses, both ENDR-B and ET-3 have been demonstrated on enteric neurones and gut mesenchyme cells, suggesting that their function may be to regulate interactions between neural crest cells and gut mesenchyme cells, necessary for normal migration [94, 95]. Both ET-3 and ENDR-B genes have been disrupted and have been identified as the cause for the natural mutants lethal spotting mice and piebald lethal mice, respectively [94, 95]. Moreover, a transgenic mouse ENDR-B null, with a mutation targeted to the ENDR-B gene, has a phenotype identical to the piebald lethal mouse [116]. In the same year as the connection between mutations in the *Ret* receptor and familial HD was established, *ET-3* and ENDR-B mutations were also implicated in the disease [94, 95, 117]. However, these mutations have been demonstrated in less than 10% of the cases of HD in humans and particularly in Waardenburg syndrome [118–121]. Endothelins are initially produced as an inactive proendothelin that has to be activated by a specific enzyme, the endothelin-converting enzyme (ECE). Two ECE genes have been described, ECE-1 and ECE-2 [98]. ECE-1 knockout mice show craniofacial and cardiac abnormalities in addition to colonic aganglionosis [122]. A heterozygous ECE1 mutation has been identified in a patient with HD who also had craniofacial and cardiac defects [123].

Sox10 is a member of the SRY-related family of transcription factors and is expressed by enteric nervous system precursors before and throughout colonization of the gut mesenchyme [98]. Disruption of the Sox10 gene has been demonstrated to be the cause of the Dom mouse natural mutant [96, 97]. Interestingly, both homozygous and heterozygous animals produce a lethal HD-like phenotype [124]. Mutations in Sox10 have been identified in Waardenburg syndrome associated with HD [125].

Phox2B is a transcription factor that is essential for the development of the neural crest derivates [126] as it regulates the *Ret* expression in enteric nervous system precursors [127]. Targeted *Phox2B* gene disruption leads to a complete absence of enteric nervous system in the mice, a phenotype that is very similar to that of the *Ret* knockout mouse [127]. Recently, Garcia-Barcelo et al. [128] reported that *Phox2B* deficiency might predispose to HD in humans. *Pax3* is a member of the paired-boxcontaining family of nuclear transcription factors that is expressed in neural cell precursors giving rise to enteric ganglia and synergizes with *Sox10* to activate an enhancer in the *Ret* gene [129]. In the mouse, *Pax3* mutations result in a phenotype characterized by a white belly spot and deficient enteric ganglia in the heterozygous state. Homozygous deficient embryos die during midgestation with neural tube defects, cardiac defects and absence of enteric ganglia [129]. No *Pax3* mutations have been identified in patients with HD.

Surgically created HD

Most surgically generated experimental models of HD have been made in chick embryos because they are easily accessible and the development of their enteric nervous system has been well studied. Aganglionosis can be induced by surgical ablation of the premigratory neural crest [130]. Even though this procedure does not tell us about the mechanism of HD, it can be used to investigate possible treatment strategies. It has been possible to recolonize aganglionic bowel with neural crest cells by transplanting tissue obtained from the dorsal neural tube [131–133]. It has also been shown that neurons from more proximal regions of bowel, if taken at early stages after their differentiation, are capable of recolonizing distal bowel and forming enteric ganglia [133, 134].

Chemical models of HD

Sato et al. [135] first described a chemical model of HD in 1978. They created a segmental aganglionosis by topical application of benzalkonium chloride to the colon and rectum in rats. This model has also been successfully reproduced in mice and guinea pigs [136, 137] and has been tested not only in the intestine, as a model of HD, but also in the distal esophagus as a model of achalasia [138]. Benzalkonium chloride is a cationic surfactant agent that attaches to the cell membrane causing irreversible depolarization, which injures the cell membrane resulting in severe cell damage and cell death. Since neurons have a higher cell membrane negative charge, the depolarizing effect is more marked on them; thus, benzalkonium chloride induces a selective neuronal ablation in the intestinal wall [136]. This treatment eliminates almost all myenteric neurons and glia in treated segments. Although the aganglionic bowel does not show hypertrophic nerve bundles and the chemical does not affect the number of submucosal neurons, the treated part results in a narrow segment and the rectoanal reflex is abolished [135]. Compared to the other models of HD, this technique is cheap, easy to perform and the animals can survive longer. This chemical model of HD has been used to study functional and structural changes in the bowel resulting from loss of these neural elements and could be used to study the chronic changes caused by the aganglionic segment and the long-term effects of different surgical treatments [139–146].

Animal models of gastroschisis

Gastroschisis (GS) is a congenital anomaly, which occurs with an approximate incidence of 1 in 4,000 births. Based on embryologic studies, there are various theories on the pathogenesis of GS. The three main theories are somatopleure differentiation deficiency in embryonic mesenchyme [147], defects of the omphalomesenteric artery in utero [148], and premature atrophy or abnormal persistence of the right umbilical vein [149]. Although it is thought to be a rare condition, the incidence of GS appears to be increasing dramatically in recent years, possibly as a result of lower maternal age, alcohol consumption, cigarette smoking and drug abuse [150, 151]. Infants born with this condition have an abdominal wall defect, usually to the right of the umbilicus, which exposes the bowel to amniotic fluid during fetal life. Although bowel is only rarely resected in GS, histologic findings demonstrate thickened intestinal muscle layers, blunting of epithelial villi and the presence of a serosal fibrous peel [152]. Despite the fact that 90% of infants survive, children born with GS face potential prolonged difficulties with nutrient absorption and intestinal dysmotility. Although recent advances in antenatal diagnosis, surgical technique and neonatal intensive care have improved outcomes dramatically, a level of morbidity and mortality persists for neonates with GS. The etiology of intestinal damage in infants with GS is uncertain and is probably multifactorial. Studies in chick embryos, fetal mice, rats, rabbits and lambs have demonstrated similar changes in the affected intestine to those seen in human neonates with GS. These studies have suggested that both exposure to amniotic fluid and possible constriction of the bowel at the level of the abdominal defect play a role in the changes affecting the bowel.

Three types of GS animal models have been developed over the years: surgically created, teratogen induced and transgenic.

Surgically created gastroschisis models

The surgical creation of an abdominal defect in fetal rabbits and lambs in the 1970s allowed early researchers to study the pathophysiology of GS [153, 154]. Chick embryo models of GS followed in the 1980s, with the utilization of one of the most readily available and versatile animal models, which could be easily modified to avoid exposure of the bowel to fetal urine [155]. Most recently, the rat model of GS was developed to take advantage of its large litter size, short gestation and the ready availability of molecular probes and reagents [156].

With the exception of the chick embryo, the surgically created models involve a laparotomy, followed by a hysterotomy to expose the experimental fetus. This is generally performed under general anesthetic with tocolytic agents to prevent premature labor. A small fetal para-umbilical incision is then made to allow extrusion of bowel loops out of the abdomen, with care taken not to injure the umbilical vessels. The fetus is then replaced back into the uterine cavity to simulate human GS, where the bowel is bathed in amniotic fluid. Alternatively, the fetus can be partially replaced into the uterus with the bowel remaining outside the uterine cavity to prevent amniotic fluid from coming into contact with the fetal intestines.

In 1991, Philips modified the fetal rabbit model to yield higher survival rates (up to 82% in the GS group), resulting in significantly shorter and thicker bowel in experimental animals compared to controls [157]. Animals were generally operated on day 23-24 (Full term = 30-31 days) with specimens being harvested shortly before term. Further studies examining the effects of amniotic fluid on the exposed bowel were performed by creating GS either intra-amniotically or extra-amniotically [158]. This resulted in significantly shorter bowel in fetal rabbits, where the bowel was exposed to amniotic fluid compared to bowel that was maintained extra-amniotically and control bowel. However, there was no significant difference in intestinal weight or bowel wall thickness. Surprisingly, there was no evidence of serositis or adhesions in experimental bowel and all GS specimens were histologically normal, in contrast to bowel from human newborns with GS.

Nutrient uptake by GS intestine has also been studied using the rabbit model. Shaw employed an everted mucosal sleeve technique to determine the uptake of an amino acid (proline), and a sugar (glucose) in the small intestine (SI) of GS-affected fetal rabbits [159]. He found that the uptake of proline per milligram and centimeter of SI, and the uptake of glucose per milligram of SI was significantly impaired in GS fetuses compared with controls.

The role of nitric oxide (NO) in GS-induced intestinal damage has been studied in the fetal rabbit model of GS. Bealer measured NO synthase activity in the SI of fullterm fetal rabbits with and without GS [160]. He found that the SI of GS fetal rabbits showed a significant increase in total NO synthase activity 2.5 times greater than that of control littermates without GS. He attributed this finding to accelerated enzyme kinetics.

The effects of prokinetic agents have also been studied to determine their potential effects on intestinal hypomotility in fetal rabbit GS [161]. Cisapride was the only agent found to improve bowel contractility in newborns with GS. Erythromycin, octreotide and metoclopramide did not improve in vitro contractility of GS bowel in fetal rabbits. Most recently, gastrointestinal motility has been determined in preterm fetal GS rabbits with the administration of intragastric fluorescein [162]. This study showed that bowel exposure to amniotic fluid reduces intestinal motility and gastric contractility in the preterm rabbit fetus. The fetal rabbit model of GS has recently been used to demonstrate the feasibility of intrauterine repair of GS for the first time [163]. This short study only assessed fetal survival for 5 h after initial GS creation and therefore has not addressed the issue of long-term survival following intrauterine GS repair or if there is any reduction in intestinal damage following the procedure.

The lamb model of GS has allowed more invasive therapeutic techniques to be studied by virtue of the size of the study animal. Fetendoscopy has been employed not only to create GS surgically but also to place an amnioinfusion catheter into the amniotic cavity of the experimental fetus [164]. This fetendoscopic technique resulted in survival rates of 46.7% in the GS group. Esophageal ligation has recently been employed in the fetal lamb model of GS in an attempt to interrupt the passage of amniotic fluid through the digestive tract [165]. However, this intervention did not have a beneficial effect on the inflammatory process affecting GS bowel in this model. In the lamb model morphometric analysis of GS bowel revealed thickening of the muscularis propria with evidence of both cellular hyperplasia and hypertrophy [166]. These changes in smooth muscle were associated with a thickened submucosa with an increase in collagen content.

The chick embryo model of GS has allowed many studies on the effects of amnio-allantoic fluid (AAF) exchange on GS bowel. The amniotic cavity of chick embryos is readily accessible once an eggshell window is created. The allantoic and amniotic membranes can then be opened to create a common cavity, which mimics the amniotic cavity in humans. The characteristic picture of GS evolved only in chick embryos where herniated bowel was exposed to urine components in the allantoic fluid [167]. The use of AAF exchange in GS chick embryos has demonstrated a reduction in macroscopic and microscopic intestinal damage [168, 169], and prevented a decrease in bowel contractility [170]. Aktug et al. [171] found that AAF exchange resulted in a reduction in the levels of gastrointestinal waste products (bile acids and bilirubin) in embryonic AAF; however, it did not alter the levels of urinary waste products, such as urea and creatinine. This dilution in the concentration of gastrointestinal waste products vielded a reduction in bowel damage seen in GS chick embryos. Studies on the effects of AAF pH on GS bowel have shown that pre-treatment with alkalinization of AAF can also reduce intestinal damage [172].

Research involving the fetal rat model of GS has centered around neuronal differentiation of the bowel, as it has been proposed that delayed maturation of intestinal smooth muscle cells might account for the dysmotility seen in human neonates with GS. Survival rates of 78–91% have been achieved in this model. In the GS rat model fetus, the myenteric plexus was noted to be disorganized with absent or scattered neuronal cells suggesting a degree of neuronal immaturity [173]. Intestinal neurofilament formation also appeared immature, demonstrating delayed cytoskeletal organization and reduced synaptic activity in the myenteric neurons in the rat model of GS [174]. Immunohistochemical analysis of GS intestine showed delayed differentiation of smooth muscle cells and interstitial cells of Cajal, with the most severely affected bowel loops demonstrating the most significant aberration in pacemaker cell morphology [175].

Although fetal urine has been implicated in the etiology of intestinal damage in GS, the role of meconium has also been investigated [176]. Correia-Pinto looked at different grades of meconium contamination in the fetal rat model of GS and found that all parameters of bowel damage were adversely affected by meconium contamination of the amniotic fluid (MCAF) [177]. Intestinal length was decreased in direct correlation with the grade of MCAF. In contrast, intestinal weight, degree of fibrous peel coverage and bowel adherence was significantly increased in experimental fetuses supporting the hypothesis that meconium is at least in part responsible for bowel damage in GS.

Transgenic gastroschisis models

Transgenic mice have played a relatively minor role in animal research of GS. The Heiligenberger (HLG) inbred mouse strain was established as an inbred strain in Freiburg (Germany) in 1939 and was subsequently colony-bred from 1975 to 1992 in Essen [178]. By 1994 more than 20 generations of inbreeding had taken place and the strain was registered (HLG/Zte). In this model GS occurs spontaneously with a frequency of about 3%. This strain demonstrates a higher incidence (up to 11%) of GS when embryos at the one-cell stage are exposed to the teratogen, X-irradiation. The pattern of inheritance in this model follows a recessive mode for the HLG susceptibility alleles. Utilizing a backcross strategy and genome-wide micro-satellite typing, this trait has been mapped chromosomally. Hillebrandt found a suggestive linkage for a locus responsible for radiation-induced GS (Rigs1) in a region of mouse Chromosome 7 [179]. Additional quantitative trait loci were identified on Chromosome 11 and Chromosome 13. Pils found that an increased incidence of GS (6.5 vs. 3.5% in controls) persisted in the progeny of a second generation of HLG/ Zte mice even without the pregnant females receiving a dose of radiation [180]. These results support the hypothesis that genomic instability may play a role after radiation exposure of the zygote stage of HLG mice.

Teratogen-induced gastroschisis models

The investigation of teratogens in animal models has been of great benefit in determining the deleterious effects of compounds prior to their use in humans. To date there is no single teratogenic agent in use in pediatric surgical research, which yields a high incidence of GS when administered to chick embryos, mice, rats or rabbits with good reproducibility. Agents, such as doxorubicin hydrochloride [181], ethanol [182, 183], nitrous oxide [184], ethylene glycol [185], scopolamine hydrobromide [186], acetazolamide [187], and cyclooxygenase inhibitors [188, 189] have all been investigated and have resulted in embryonic malformations, including GS. However, the teratogenic rate in these studies is relatively low with GS affecting embryos in only 3.7– 19.8% of cases.

The injection of doxorubicin hydrochloride/saline into the albumin of chick eggs resulted in a GS rate of 19.8% [181]. However, the development of ventral defects, such as GS, in the chick embryo seems to be a perturbation of embryogenesis at a crucial early stage rather than a direct teratogenic effect of doxorubicin itself, as controls receiving the saline vector commonly developed the malformation (15.7%).

Due to the high incidence of GS in offspring of young mothers who often smoke during pregnancy, Singh studied the effects of carbon monoxide (CO) and altered protein/zinc diet on pregnant mice of the CD-1 strain [190]. He found that the incidence of GS was 47% in the low protein/zinc deficient group, which had been exposed to CO. In contrast, no fetal mice developed GS in the other groups with either low protein/zinc diet/air or CO alone. These findings contribute to the thinking of a multifactorial etiology of GS.

Alcohol has been widely studied for its teratogenic effects and in humans results in a pattern of malformations in what is known as the fetal alcohol syndrome (FAS). The main methods of administration have included intraperitoneal injection of an alcohol solution to pregnant rats [182] and oral administration of ethanol-derived diet to pregnant C57BL/6J isogenic mice [183]. Although experimental fetuses in both rat and mouse studies developed multiple anomalies, the incidence of GS was generally low with the doses given.

An interesting chemical, which has been extensively studied in the toxicology and oncology literature, is the polar solvent N-methylformamide (NMF). This compound is a metabolite of dimethylformamide, a solvent with wide applications in the chemical industry. Its teratogenic effects were noted in rabbits as far back as 1980 when oral administration resulted in a number of developmental anomalies [191]. At this time rabbit embryos appeared to be more sensitive to these chemicals than other animals and more recent studies have demonstrated similar findings. It has also been evaluated as an antineoplastic agent because it has been shown to induce tumor inhibition in human colon carcinoma cell lines when used in combination with 5-fluorouracil [192]. In a study in 1995, Kelich [193] utilized Crl:CD (SD) COBS rats and New Zealand white rabbits [Hra:(NZW)] and administered daily NMF by gavage at doses up to 75 mg/kg from GD 6–15 in rats and 50 mg/kg from GD 6-18 in rabbits. In the rat group, only one fetus developed GS out of a total of 269 (0.4%). Interestingly, in

the rabbit group 84 out of 85 (99%) fetuses developed GS, with skull and sternal anomalies being the other major malformations. With this high incidence of fetal GS in NMF-treated rabbits the potential for future development of an animal model in pediatric surgical research is significant. Further studies of this model could yield valuable results for pediatric surgical researchers worldwide.

Animal models of congenital diaphragmatic hernia

Congenital diaphragmatic hernia is a common congenital malformation with an incidence of 1 in 3,000 births. Despite progress in neonatal and intensive care units, the outcome of these patients depends mainly on the pulmonary hypoplasia that remains the principal contributor to the high mortality of CDH. Much of the current understanding of the pathophysiology of CDH originates from experimental studies.

Four types of CDH animal models have been developed over the years: naturally occurring, surgically created, teratogen induced and transgenic.

Naturally occurring CDH

A CDH pig model was described in a herd that was originally bred to produce pigs with anorectal malformations [194]. Among 93 piglets, 13 with CDH were identified, which presented with symptoms from 1 to 4 months after birth. As with humans with CDH, herniated intra-abdominal organs, stomach and intestine were found within the thoracic cavity; but crucially, these pigs lacked the pulmonary hypoplasia that is characteristic of human CDH. This pig model represents a true natural animal model of CDH, but the main disadvantage is that it requires a large herd of animals to ensure a regular production of animals with CDH.

Surgically created CDH

De Lorimier [195] was the first to create an animal model of CDH, using third trimester fetal lambs, based upon the traditional view that lung hypoplasia in CDH results from compression by abdominal viscera. This led to hypoplastic lungs but similar pressure volume curves compared to sham-operated controls. Subsequent studies attempted to simulate CDH by placing a balloon in the left chest in fetal lambs [196, 197]. Newborns had significantly reduced lung weights and air capacities, decreased pulmonary vascular bed and decreased pulmonary compliance compared with control animals. Following this it was shown that in utero deflation of the balloon improved the pathophysiologic consequences of CDH and improved survival [197]; but none of these models accurately simulated the pulmonary vascular morphological changes seen in humans with CDH. After

that, the age in which the defect was created was decreased to the second trimester in the hope that insulting the developing lung earlier in gestation would more closely simulate the human condition [198]. In this model, time-dated pregnant ewes were anesthetized and following a midline abdominal incision, the fetus was localized by palpation of the uterus. Then a hysterotomy was performed and the upper torso of the fetus delivered. Next, a posterolateral left thoracotomy was made and the diaphragm was incised widely over the superior edge of the spleen and the fundus of the stomach; the stomach, omentum and small intestine were gently pulled into the chest. The fetal thoracotomy and maternal hysterotomy were closed and pregnancy prolonged using tocolytics. This model has also been tested in rabbits [199-201]. Resulting CDH lambs have a number of similarities to the human condition: the lungs are hypoplastic and have reduced number of airway branches, diminished alveolar number, decreased size of the pulmonary vascular bed and an increase in the muscularization of the pulmonary arterial tree when compared with controls [202-204]. The prenatal repair of this surgically created diaphragmatic defect showed a reversal of pulmonary hypoplasia; however, a prospective randomized trial in humans did not show any clear benefit over standard postnatal management [205]. This redirected the efforts to approach the disease on reversing the pulmonary hypoplasia rather than focusing on the diaphragm. Extrapolating the findings from the congenital high airway obstruction syndrome in which hyperplastic lungs are found, an innovative animal model was created utilizing temporary tracheal occlusion to induce lung hyperplasia [206]. Repair of the actual diaphragmatic defect occurred postnatally. Initially, a clip was placed in the trachea through a cervical dissection. The clips, however, required surgical removal prior to birth leading to associated tracheal complications. Modification led to the current technique of endoscopic placement of a detachable balloon directly into the fetal trachea [207]. The results of this technique, in a randomized controlled trial in humans, have recently been reported [208]. The historical analysis of how pioneering fetal surgery developed for CDH reflects the importance of animal models in surgical diseases. However, since this animal CDH model is surgically created during fetal life, it is unable to yield any insights into the embryogenesis of CDH.

Teratogen-induced CDH

Although the teratological effects of the herbicide nitrofen (2,4-dichlorophenyl-*p*-nitrophenyl ether) in rodents were described a long time ago [209], they had been overlooked by pediatric surgeons until a rediscovery of this model led to an explosion in research [210-212]. Since nitrofen is administered during early organogenesis, the advantage of this model over the surgically created ones is that it gives the opportunity to examine the embryogenesis of CDH. The exact time of administration and the right dosage of nitrofen needed to create CDH have been studied over the years. In the mouse model, time-mated CD-1 or Swiss-Webster mice have to be gavage fed with 25 mg of nitrofen dissolved in 0.5 ml of olive oil on GD 8 to produce severe pulmonary hypoplasia in the whole litter and CDH in 30% of them [213]. In the rat model, time-mated Sprague-Dawley rats are gavage fed with 100 mg of nitrofen dissolved in 1 ml of olive oil on GD 9.5 in order to obtain a 60% rate of CDH, most of them being left sided. Interestingly, if rats are given nitrofen on day 11 of gestation, they mainly develop right-sided CDH [211].

In both mouse and rat models, the CDH animals have predominantly large-sized hernial defects containing liver and stomach and a marked bilateral pulmonary hypoplasia. Surprisingly, nitrofen-treated lungs without CDH are also hypoplastic [211, 212, 214, 215]. The associated malformations found in the CDH animals have been studied and they closely mimic the incidence and spectrum of those seen in humans [216–220]. Since the defect is induced at the stage when the foregut has just separated into the esophagus and the trachea, this model of CDH has given the opportunity of carefully studying the developmental anatomy of the lungs and diaphragm in CDH [212, 221-224]. Recent studies have shown that the pulmonary hypoplasia precedes the diaphragmatic defect in this model and that there could be a common mechanism underlying the etiology of CDH or that the pulmonary hypoplasia might even be the cause of the diaphragmatic defect instead of the result [225, 226]. Although all these studies provide some information about the pathogenesis of CDH, research into the etiology of CDH had been lacking. Recently, Greer et al. [227] have provided evidence suggesting that the retinoid signaling pathway may contribute to the etiology of CDH.

Even though the nitrofen model of CDH has been widely used, it is unclear whether pulmonary hypoplasia (particularly in those nitrofen-treated animals without CDH) may arise due to a direct toxic effect of nitrofen rather than being secondary to the CDH and hence has little to tell us of hypoplastic lung development in human CDH. Moreover, since there is a liver ingrowth into the chest in most of the animals with CDH and this is rarely seen in human CDH, the nitrofen CDH model has recently been questioned as a good model of CDH [228]. Nevertheless, the nitrofen model remains one of the most popular because it is cheap, technically simple and because it provides a singular opportunity for embryological studies.

Transgenic models of CDH

Ideally, the best animal model of CDH would be a gene knockout model that could accurately reproduce the malformation seen in humans. So far, several mutant phenotypes have been described in which

diaphragmatic hernia may feature but they have a range of often far more frequent and significant malformations that make them phenotypically different from human CDH. Deletion of both murine RARs is associated with a high incidence of cranial, vertebral, limb, cardiac, foregut and pulmonary malformations whilst manifesting a low frequency of CDH [229, 230]. Mice with mutations for the homeobox Hlx gene also have CDH; however, more remarkable in these mice are their large lungs and small liver and intestine [231]. Homozygous inactivation of WT-1 results in CDH but they are also severely deformed and have serious defects in the urogenital system [232]. Recently, knockout mice homozygous for Slit3 deficiency were constructed. In these mice, the most obvious phenotype is CDH, but the hernia is located on the ventral midline portion of the central tendon rather than in the posterolateral diaphragm and all hernias include the liver and the gallbladder. Moreover, 20-40% of the mice have renal and ureteral agenesis [233].

Heterogeneity between human CDH and described knockout mice restrains against the adoption of transgenic CDH models at present.

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