4 Animal Models of Aganglionosis

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4.1 Introduction

Humans are not the only mammals to suffer from aganglionosis. Aganglionosis has also been described in mice, rats, horses, cats and dogs. Rodent animal models have contributed significantly to our understanding of Hirschsprung's disease (HSCR). Over the last decade, the understanding of the genetics and cell biology of the development of the enteric nervous system (ENS) has made great progress. Rodent animal models have shown many points of correlation with humans in regard to ENS development, both normal and abnormal. Nevertheless, the link between the genotype and the phenotype is often indirect, and so many questions have yet to be answered. This chapter deals with the characteristics of aganglionosis in rodents with emphasis on how knowledge of the animal models has contributed to our understanding of the genetics and pathogenesis of HSCR and allied disorders.

4.2 History

4.2.1 Rodents

The first description of aganglionosis in mice was by Derrick and St George-Grambauer in 1957 [1]. They found approximately 3.2 per 1000 of their colony developed aganglionosis. The average length of aganglionosis was 15–20 mm with colonic distension extending into the first few millimeters of aganglionic colon. There was no association with a white or patched color coat. The next description was by Bielschowsky and Schofield in 1962 [2]. In this colony 10% of the offspring were affected and there was an association with a white colored coat. These mice also had a high incidence of mammary cancer and pituitary adenomas. Outbreeding experiments suggested an autosomal recessive trait with modification of the trait by other genetic factors.

In 1966, Lane described two strains of mice which developed aganglionosis as an autosomal recessive condition [3]. The lethal spotting (ls) mice have approximately 2 mm of aganglionosis with a patched coat, and later studies linked the defect to chromosome 2. Piebald lethal $(s¹)$ mice had approximately 10 mm of aganglionosis and linkage studies suggested that the defect was on chromosome 14. In 1979, Ikadai et al. [4] described aganglionosis in spotting lethal (sl) rats. The animals had two lengths of aganglionosis: total colonic aganglionosis (TCA) and mid colon. These animals again showed autosomal recessive inheritance and had a white colored coat. A fourth rodent model is the Dominant megacolon (Dom) mouse

in which the aganglionic colon has a long hypoganglionic transition zone (Table 4.1) [5].

4.2.2 Other Mammals

There have been isolated reports of aganglionosis in a range of animals including cats [6, 7], horses [8–11] and pigs [12, 13].

4.3 Histologic Anatomy

Are rodents good histologic models of HSCR? The first histologic studies were performed by Lane [3]. These were restricted to showing there was aganglionosis in the terminal bowel, and in documenting the length of aganglionosis.

Bolande and Towler [14] and Bolande [15] investigated the lethal spotting mouse using histochemical and ultrastructural studies. Histology showed hypoganglionosis in the distal bowel but there was no dense ingrowth of nerve fibers. Boley [16] suggested that these findings of no hypertrophied nerve trunks indicated that these mice were not a good model of human disease. In the distal narrowed segment there was a reduction in adrenergic and cholinergic fibers. In the dilated part of the bowel there was an increase in adrenergic fibers. The ultrastructural studies showed that just above the transition zone, there were secondary degenerative changes in the ganglion cells, which increased with age, resulting in what appeared to be secondary cell death and abiotrophy. Webster [17, 18] performed detailed studies in lethal spotting and piebald lethal mice using cholinesterase stains and fluorescence to delineate the adrenergic nerves.In postnatal mice of both strains he demonstrated normal innervation in the proximal bowel followed by a transition zone and then an aganglionic zone with increased nerve trunks and a decrease in the innervation of the circular muscle fibers. In the most distal aganglionic colon, just above the anal sphincters, there appeared to be a variable, but denser, innervation to the circular muscle involving cholinergic nerves. Bu'Lock et al. [19] found a selective depletion of substance P in the transitional zone in piebald lethal mice. However, the change was from 10% in normal mice to 5% in mutant mice, and the study failed to confirm a previous report from the same laboratory of a decrease in substance P in the mutant ileum, indicating that the variability between animals and sensitivity of the techniques can make conclusions difficult.

In the spotting lethal (sl) rat model Ikadai et al. [4, 20] and Horie et al. [21] studied the length of aganglionosis and found that there were two subgroups, one in which there was TCA and a second, less numerous group, in which ganglion cells extended to the proximal half of the colon. The visible cone was often distal to the commencement of aganglionosis. In a histologic study of sl rats using whole-mounts and AchE, tyrosine hydroxylase and substance P, Nagahama et al. [22] showed aganglionosis of the colon along with increased nerve trunks. These changes are similar to those seen in humans with TCA. However, even in the proximal ganglionated duodenum there were changes in the two dimensional structure of the enteric plexus, with the lattice pattern being irregular. This raises the question as to whether the proximal gut is entirely normal, and if there is histologic abnormality in some of the subtle architecture, does this mean diminished function? The bowel also has many functions, so it may be possible for example that water absorption is affected while propulsive activity is normal.

An ultrastructural study in the sl rat model confirmed that almost no nerve terminals were present in the circular muscle layer of any regions of the constricted intestine, but some terminals were observed in the longitudinal muscle layer of that segment. The authors concluded that the denervated circular muscle layer is related to the production of a constricted segment, irrespective of the presence or absence of nerve terminals in the longitudinal muscle layer [23].

Table 4.1 Naturally occurring rodent animal models of HSCR

Studies in our laboratory using AchE whole-mounts have demonstrated that the three rodent animal models have a histologic picture identical to that of humans. There was distal aganglionosis with increased nerve trunks, followed by a transition zone which was often asymmetric and variable in length, and more proximally the plexus was near normal in two-dimensional architecture. In spotting lethal rats, particularly those with TCA, the proximal small bowel had an abnormal architecture, but inconsistently so. We could not demonstrate as clearly as Nagahama et al. [22] that the duodenal architecture was abnormal. In our experiments the normal two-dimensional architecture of the enteric plexus was itself variable with some areas looking open and other areas with a tight regular lattice-like structure.

In the lethal spotting mice we were able to identify increased nerve trunks, in contrast to Bolande [15] who could not detect them. The animals we investigated had a longer length of aganglionosis (about 4 mm) and it may be that the genetic background of the animals used by Bolande was such that they had a very short length of aganglionosis, and were more a hypoganglionic model.

The published literature suggests that the length of aganglionosis in each animal model is relatively consistent within each strain [3]. However, in reality there is a considerable variability and while in the majority of cases the length of aganglionosis in piebald lethal is greater than in lethal spotting mice, there is overlap. We have noticed that if heterozygote animals with the least pigment are chosen as mating pairs then the length of aganglionosis tends to be longer. Alternatively, if the mutant animals are back-crossed with C57 or Castaneus stock, the length of aganglionosis reduces. Some F2 offspring from a lethal spotting mouse crossed with Castaneus apparently had anatomically normal distal colon although the color of the coat was that of a mutant animal. The most dramatic alteration in length of aganglionosis occurs when spotting lethal rats are crossed with DA rats: the predominant ileal aganglionosis changes to distal colonic aganglionosis. The change is a dramatic step-like decrease in the length of aganglionosis, suggesting that an extra quantum of enteric neuronal precursors has been created or there are regional differentiations in the bowel such that a region is either filled or remains aganglionic. It is of interest that in humans there also appear to be two common points of cessation of enteric innervation, either the sigmoid colon (in 80% of patients) or about 10 cm proximal to the ileocecal valve (in 10% of patients).

The sl rat has also been instructive in further elucidating the distal aganglionic bowel, namely that the gut is not completely aganglionic. There are groups of ganglion cells (seen best with NPDH-diaphorase staining) associated with the hypertrophied nerve fibers. We postulate that these can only have arisen from the sacral neural crest (there is a gap of many centimeters before ganglion cells are seen in the small bowel). Furthermore, we pos-

tulate that the presence of occasional clusters of ganglion cells in the "aganglionic" distal bowel in HSCR patients, if sampled on rectal biopsy, may cause a temporary diagnostic error. This may be the basis of the rarely described patients with "acquired aganglionosis", where the patient clinically has HSCR but the first biopsy suggests the presence of ganglion cells.

Morphologic studies in mice have shown that the density and distribution of the interstitial cells of Cajal (ICC) in the aganglionic region of the colons are similar to those of ICC at the same level of the colon in agematched wild-type controls [24, 25]. Therefore, it appears that the enteric neurons are not necessary for the development of ICC. Data from humans are inconsistent which could be attributed to the regional differences in the density of ICC in the colon [26–28]. More recently Taniguchi et al. [29] have shown that the aganglionic intestine of ls/ls mice induces secondary disturbances during the normal development of ICC, in the form of fewer cytoplasmic processes and lack of attachment to the intermuscular nerves.

4.4 Physiology

The principle work, which has been on the piebald lethal mouse, is that of Wood et al. [30–32]. In a series of experiments the colon was inspected under video cameras and at the level of aganglionosis there was a functional obstruction. Pellets would move down the bowel and as soon as they reached the aganglionic zone they would stop and at that stage reverse peristalsis would commence [31].

Electrophysiologic studies in piebald lethal mice have shown that there are abnormal discharges of myogenic action potentials in the aganglionic bowel associated with tonic constriction and a reduction in the luminal diameter [30]. Furthermore, the proliferated extrinsic cholinergic nerve fibers appear to be not related to the narrowing of the aganglionic colon [33].

Electrophysiologic experiments in our laboratory have demonstrated that the aganglionic bowel in mouse models has no inhibitory neuromuscular junction potentials (IJP) and only occasional excitatory junction potentials which on repeated stimulation fatigue quickly. The aganglionic circular muscle, lacking inhibition, writhe in an uncontrolled manner and tend to contract [34]. The visual and tension studies confirm the impression that the aganglionic smooth muscle lacks the stabilizing influence of the nonadrenergic noncholinergic (NANC) inhibitory nerves. The smooth muscle appears hyperexcitable and in constant motion. This would confirm the hypothesis proposed by Alvarez [35] that the simplest explanation for the finding in HSCR is that nerves to smooth muscle normally function to keep the muscle from contracting into a knot. This is also the conclusion of Richardson

[36] who performed a pharmacologic study on the lethal spotting mouse.

The simple lack of nerve fibers is sufficient explanation for the functional obstruction seen in the rodent models and in patients with HSCR. There is no need to invoke selective and subtle disorders of various components of the autonomic nervous system to account for the clinical variability.

Thorough studies on the electrophysiology of the mutant rat colon have shown an absence of IJPs except at the sphincter where there is an evoked inhibitory response (in addition to an excitatory response) [37–40]. Another study in piebald lethal mice has shown an increase in basal contractile activity and a reduction in responsiveness to vasoactive intestinal peptide [41]. This supports a generalized reduction in the function of the inhibitory innervation of the aganglionic colon.

In 1990, Bult et al. [42] provided evidence that nitric oxide (NO) is released on stimulation of the inhibitory NANC nerves of canine ileocolonic junction. Since then, substantial evidence has accumulated indicating that NO is the primary nitrergic inhibitory neurotransmitter in the gut of various species [43–46]. More recently de Lorijn et al. [47] have shown that the inhibitory innervation of the murine internal anal sphincter and the rectoanal inhibitory reflex are mediated by NO, and the rectoanal inhibitory reflex requires an intact network of ICC in the internal anal sphincter. Thus both loss of nitrergic innervation and deficiency of ICC lead to impaired anal relaxation and may play an important role in rectal evacuation disorders.

4.5 Embryologic Studies on Rodent Models of Aganglionosis

Webster [17, 18] reported studies on both the ls and $s¹$ mice in which he used a nonspecific esterase stain to follow migratory patterns of enteric neuronal precursors in embryos of mutant mice. In both cases there was slowing of migration such that the migrating vagal neural crest cells (VNCC) did not keep pace with the rapidly elongating gut, although the cells still showed signs of distal migration for several days after the usual time of cessation of migration of VNCC in the normal embryos. Webster interpreted these results as suggesting a defect in the vagal neural crest. Using the nonspecific esterase strain in the mutant and control animals there was no evidence of a sacral neural crest input. However, Rothman and Gershon found different results studying the same ls mouse strain [48, 49]. They found that the gut microenvironment in the embryo is unreceptive and cannot support enteric neuronal precursors. The principal experiments in reaching this conclusion were cocultures where isolated segments of aganglionic colon from mutant mice were placed next to sources of neural crest cells (either vagal crest or proximal gut). Aganglionic gut from the mutants

was never normally innervated whereas distal gut from the normal embryo was receptive to ingrowth of enteric neuronal precursors. Rather, enteric neuronal precursors tended to avoid aganglionic gut tissue [50]. Other studies looking at the extracellular matrix have shown increases in laminin, collagen type IV and chondroitin sulfate in the distal aganglionic gut [51, 52]. These authors concluded that it is an excess of these extracellular matrix molecules, in particular laminin, which results in a hostile local microenvironment and this causes aganglionosis. They proposed that there is no defect in the émigrés from the neural crest.

Commencing in 1984 our laboratory studied the ls, s¹ mouse and sl rat animal models, using histology, tissue culture and the kidney capsule techniques. We used three techniques in an attempt to overcome possible artifacts with any one technique, and three animal models to detect any differences between each of the animal strains. The results in all animals, and with each of the three techniques, agreed with those of Webster. In particular, there was slowing of migration in what was the one predominant source of enteric neurons, namely vagal neural crest cells. The slowing in the migration of these cells occurred well before the eventual aganglionic zone was reached and migration spluttered on for several days after it would normally have ceased. In the ls and s¹ mice the slowing commenced in the terminal small bowel whereas in the sl rats the slowing was apparent even in the proximal small bowel. The appearances were most consistent with the interpretation of a lesser population of enteric precursors in the mutants and insufficient numbers to fully colonize the embryonic gut [53].

We made one new finding which shed some light on the contentious debate as to the existence and extent of a contribution of the second sacral neural crest to the vagal neural crest cells: there were small numbers of enteric precursors in the most distal hindgut. These cells were usually in groups of two to four (maximum ten) in contrast to the vagal émigrés which proliferated into the thousands. There was no obvious difference in the numbers of these sacral neural crest cells in the mutants compared to normal embryos [34]. This result was confirmed using the three different experimental techniques. Therefore, as in most good debates, both sides are correct—there is a sacral neural crest contribution to the ENS, but this contribution is functionally insignificant. Nevertheless, failure of the vagal derived neural crest cells to colonize the hindgut is the prime cause of failure of hindgut enteric nervous development. Furthermore, the interaction between sacral and vagal enteric neural crest cells may be necessary for sacral neural crest cell contribution to the ENS [54, 55].

However, the use of immunohistochemistry, special stains and techniques such as the kidney capsule were limited in achieving a full understanding of how aganglionosis arises in the animal models. Ultimately, all the techniques are indirect, and the debate between the

various theories (see below) could not be settled. Therefore, the new techniques of molecular genetics, that is linkage studies and gene knockouts, were utilized. Initially, the first strategy was linkage studies, as there was no knowledge of which genes were likely to be involved and so knocked out. Theoretically the linkage strategy was simple but during the laborious application of these techniques the answers came from knocking out genes known to have a completely unrelated physiologic function with the unexpected finding that aganglionosis resulted.

4.6 Molecular Genetics

We will focus here on the animal work that lead to a better understanding of the molecular genetics of HSCR and allied disorders (for further details see Chapter 5).

4.6.1 Backcross and Linkage

As the rodent model animals are inbred, if polymorphisms are introduced by outbreeding (backcrossing) then a linkage strategy should allow the chromosomal area responsible for the mutation to be progressively narrowed and finally the gene isolated [56].

In several laboratories (including ours) a lethal spotting backcross strategy was used to try and localize the gene. It was already known that the chromosomal location was on mouse chromosome 2, and backcross studies narrowed the area to between GNAS and endothelin 3 (*edn3*) (Ramu E et al., unpublished work; [57]). However before further work could be done to walk into the mutation area, the answer came from knockout experiments (see below). A similar backcross strategy was used in the s¹ mouse. Work was more successful in these experiments, in that the regions of interest were localized and cloned, but it remained difficult to identify the gene involved [58]. Similarly in the sl rat, a project was initiated in our laboratory to localize the gene responsible for aganglionosis in the rat, using the mutant animals crossed with the DA rat. The DA rat was chosen because of its heavy pigmentation which allowed the wild types and heterozygotes to be clearly distinguished from mutants in the F2 offspring litter. However, while this work was progressing the answer came from knockouts of genes found initially in humans and adult animals whose full function was being explored by the use of experiments in which the gene was removed and then what happened in the offspring was observed.

4.6.2 Knockout Models

With advances in knockout and transgenic technology, many molecules and several signaling pathways have

been identified as important in the control of mammalian ENS development (Table 4.2).

4.6.2.1 *Ret/Gdnf/Gfrα1* **and** *Ret/Ntn/Gfrα2* **Knockout Mouse**

The *Ret/Gdnf/Gfrα1* signaling pathway is of importance in ENS development, having been shown to promote survival of neurons, mitosis of neuronal progenitor cells, differentiation of neurons and neurite extension [59, 60].

The first targeted gene deletion knockout model was of the *ret* gene which unexpectedly produced a phenotype similar to aganglionosis. The *ret* gene had been initially isolated in a tumor cell line [61]. Subsequent examination of these animals showed a total absence of ganglion cells throughout the gut and associated renal anomalies [62]. *Ret* is normally expressed in the embryonic gut [63]. *Ret−/−* mice exhibit a failure of neural crest colonization of the gut distal to the gastric cardia. The esophagus and gastric cardia also exhibit a reduced population of neurons and glia [64, 65].

This animal work was critically important in identifying *RET* as a firm human candidate gene in the area previously identified as deleted on chromosome 10 [66–69]. Without this vital research, *RET* would have remained only one of a dozen or so candidate genes in this deleted area of the human chromosome. It was a combination of both the human and animal works that allowed the early identification of *RET* as the first gene responsible for HSCR [70, 71].

Likewise, *gdnf−/−* and *gfrα1−/−* mice have almost identical phenotypes to *ret−/−* mice [72, 73]. To date, a few *GDNF* mutations and no *GFRα1* mutations have been identified in humans with HSCR [74–76].

Gfrα2 and *ntn* are, like *gfrα1* and *gdnf*, capable of forming a *ret*-activation complex and are thought to be required for the maintenance of a subset of enteric ganglia [77]. *Ntn−/−* and *gfrα2−/−* mice exhibit a decrease in the density of cholinergic neurons in the ENS but no renal abnormalities, and the mice survive and breed [77, 78]. It appears that *NTN* mutation alone does not result in HSCR, but could contribute to the severity of HSCR due to other mutations [79]. To our knowledge, the *GFRα2* mutation has not been identified yet in humans with HSCR.

4.6.2.2 *Ednrb/Edn3/Ece1* **Knockout Mouse**

Endothelins had been discovered while searching for contractile substances in the pig aorta [80]. Scientists interested in the biologic function of endothelin genes also produced a series of targeted gene deletion knockouts in an attempt to see how animals without the gene would function. The first of these was an endothelin-1 (*edn1*) deletion, due to disruption of the endothelin-convert-

aOf homozygous mice

ing enzyme-1 gene (*ece1*), which produced craniofacial and cardiac defects in addition to colonic aganglionosis, suggesting that the endothelins were important in neural crest development [81, 82]. In follow-on experiments, when the endothelin-3 gene (*edn3*) was made functionally null, a phenotype resulted which looked identical to the lethal spotting mouse. Similarly when the endothelin B receptor (*ednrb*) was made nonfunctional, the offspring resembled piebald lethal animals. Examination of the gut confirmed aganglionosis. Researchers then examined the lethal spotting mice and piebald lethal mice for defects in *edn3* and *ednrb*, respectively. They confirmed that *edn3* and *ednrb* have a role in the migration and development of the ENS, and defects of the endothelin signaling pathway are responsible for the phenotypes of these animal

models. Namely, in the lethal spotting mouse there was a point mutation in the proendothelial-3 gene which prevented cleavage and resulted in no *edn3*; and in the piebald lethal mouse there was a complete deletion of *ednrb* (Table 4.1) [83–86].

The shorter length of aganglionosis in the lethal spotting mouse is thought to be due to the other endothelins (*edn1* and *edn2*) partially reacting with the *ednrb* and producing a milder form of aganglionosis. This would explain why in the piebald lethal animals the length of aganglionosis is on average about 10 mm, whereas in the lethal spotting mouse the length of aganglionosis is about 2 mm.

In our laboratory we have crossed both lethal spotting and piebald lethal animals and the resultant length of aganglionosis appears to be similar or only slightly longer than that of the piebald lethal mouse. Therefore there appears to be no additive factor between the two genes and the defect produced by the piebald lethal lesion which is due to the absence of *ednrb* is the limiting factor in the length of aganglionosis. However, when the animals are backcrossed with either C57 or Castaneus animals, the innervation of the most distal colon appears to be near normal. Therefore, there are modifier genes affecting the length of aganglionosis in these mouse models.

Knowledge that *ret* and *ednrb* defects are involved in mice suggested that defects in these genes were candidates for the spotting lethal rat. First a defect in *ret* was excluded, along with *edn3*. Working initially on a cDNA from the rat *ednrb*, we noted a 250–300 bp deletion. Further experiments using both the cDNA and genomic DNA localized the defective area to the end of the first translating exon and the next 17 bp sequence of the first intron of the *ednrb*. This 301 bp deletion results in alternative splicing which results in either a stop codon or an in-frame 270 bp deletion and a protein product with an inability to insert into the cell membrane [87, 88].

A recent study has shown that abnormalities of the ENS in heterozygous *ednrb*-deficient spotting lethal rats resemble those in intestinal neuronal dysplasia B (INDB) [89].

These mouse animal experiments again led the way to the discovery of defects in the same genes in humans. Subsequently, defects in *EDNRB* [90–94] and *EDN3* [95, 96] have been found in humans.

4.6.2.3 *Sox10* **Knockout Mouse**

Sox10 was identified while doing a comparative study of human/mouse sequences [97]. It is a member of the sry-related family of transcription factors [98]. The naturally occurring Dom mouse model of HSCR was used to identify the role of *sox10* in ENS development [99]. It appears that early death of neural crest cells is responsible for the complete aganglionosis of the gut in *sox10Dom*/ *sox10Dom* mice [100]. *Sox10Dom/+* mice exhibit distal hindgut aganglionosis and pigmentation abnormalities [101]. A similar gene mutation has been identified in patients with Waardenburg-Shah syndrome [102].

4.6.2.4 *Phox2B* **Knockout Mouse**

Phox2b is a homeodomain-containing transcription factor that regulates *ret* expression and thus it is essential for ENS development [103, 104]. *Phox2b−/−* mice exhibit neural crest colonization of foregut only. Subsequently, the foregut neural crest cells undergo apoptosis. At birth, animals exhibit complete aganglionosis of the gut [104]. *PHOX2B* is proposed to be a candidate gene in patients with Haddad syndrome [104, 105].

4.6.2.5 *Pax3* **Knockout Mouse**

Pax3 is a member of the paired-box-containing family of transcription factors [106]. It appears that *pax3* is required for the formation of enteric ganglia and functions with *sox10* to modulate *ret* expression, and thus there are no enteric neurons caudal to the stomach in *pax3−/−* mice [107]. Patients with Waardenburg syndrome without HSCR usually have mutations in *PAX3* [108].

4.6.2.6 *Hox11L1* **Knockout Mouse**

Hox11L1 is a homeobox transcription factor which may play a role in neural crest cell proliferation or differentiation [109]. *Hox11L1−/−* mice develop an INDB-like condition, followed by death of some of the enteric neurons [109, 110]. The *Hox11L1−/−* mouse has been proposed as a model for INDB. Further knowledge of the regulatory genes and the transcriptional targets of *Hox11L1* may produce candidate genes for involvement in INDB, and thus a better understanding of this controversial disease entity [59, 111].

4.6.2.7 *Ihh−/−* **and** *Shh−/−* **Knockout Mouse**

Indian hedgehog (*Ihh*) and Sonic hedgehog (*Shh*) genes may influence survival and/or differentiation of neural crest cells [112]. *Ihh−/−* and *Shh−/−* mice die during early embryonic stages. Late fetal *Ihh+/−* mice exhibit a dilated region of the colon, with missing enteric neurons in some parts of the small intestine and the dilated region of the colon [113]. *Shh+/−* mice do not lack an ENS in any part of the gut, but nerve cell bodies are present within the mucosa [113]. Both *IHH* and *SHH* are possible candidate genes for ENS defects in humans [59].

4.7 Contribution of Animal Models to Theories as to the Cause of Aganglionosis

There are two broad theories as to the cause of aganglionosis (with many internal minor refinements of the theories being possible), and workers investigating the animal models have found evidence for both theories.

4.7.1 Defect of Central Vagal Neural Crest Cell Production and Migration

Webster's and our interpretation of the slowing of migration found in mutant embryos is that this suggests an early defect in the vagal neural crest in the production of sufficient neural crest cells to adequately populate the gut. This hypothesis would fit with the ablation experiments of Yutema and Hammond (see Chapter 2).

As a refinement to this theory we hypothesize that the uneven distribution we see in humans and mutant animal experiments suggests evidence of clones or discrete quanta of precursors or mother cells, perhaps as low as four to six quanta from which arise all progeny that normally populate the gut. If one quantum is missing, distal colonic aganglionosis results; if two quanta are missing, ileal aganglionosis results; if three are missing, mid-smallgut aganglionosis results; if four quanta are missing, total intestinal aganglionosis results [34]. Certainly in the mutant rats we do not see an even or random distribution of the site of the commencement of aganglionosis; rather there are three most frequent "nodal regions" where aganglionosis commences. Our hypothesis is that the quanta are generated at a vagal neural crest level over several somites and the defect in the mutants occurs initially at a premigratory stage, with the eventual extent of bowel aganglionosis being merely a later playing out of this early vagal neural crest defect. It is as if there is a "checkerboard" of potential spaces to be filled and a number of precursor families to fill the spaces. Because the spaces are always filled from a proximal direction, no matter which family is missing, the result will be distal aganglionosis. The only variable is that if there are more families missing then the length of aganglionosis will be longer (Cass, First World Workshop in Hirschsprung's Disease, Sestri Levante, 1993, unpublished data).

4.7.2 Defect in the Local Gut Microenvironment

Gershon and coworkers extensive experimental work supports defects in the mutants being in the gut microenvironment itself and being specifically related to excess laminin causing migrating enteric neuronal precursors to mature early and hence not to continue to divide or migrate [114]. Nishijima et al. [115] found migration down the mouse embryonic gut was not an even process but rather proceeded in bursts followed by a pause. In the lethal spotting mouse mutants, migration proceeded normally but then suddenly stopped at the last of these boundaries, resulting in aganglionosis. The authors interpreted this result as indicating that the gut had subtly different microenvironments, and an intrinsic defect in the last 2 mm of the mouse colon resulted in aganglionosis. Similarly Kapur et al. [116], using transgenic mice with a cell label DβH-inlacZ and chimeric animals, showed that the enteric neurons from ls/ls could populate the distal gut. The explanation was that the enteric neuronal precursor from the normal embryo contributed a factor that overcomes the microenvironmental defect [116–118]. In our experiments, we could not demonstrate defects in the extracellular matrix in early embryos [34, 53]. Rather

the increase in the extracellular matrix components appeared to be a secondary event [34, 53].

4.8 Summary

In summary, the animal models of aganglionosis have been pivotal in the discovery of the genes of HSCR. In future, animal models will continue to contribute to the understanding of how the genes interact and are modified by yet other genes. In addition, animal models of aganglionosis will continue to contribute to the anatomic, physiologic and pharmacologic understanding of aganglionosis.

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