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Evaluation of growth hormone (GH) action in mice: discovery of GH receptor antagonists and clinical indications

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Abstract

The discovery of a growth hormone receptor antagonist (GHA) was initially established via expression of mutated GH genes in transgenic mice. Following this discovery, development of the compound resulted in a drug termed pegvisomant, which has been approved for use in patients with acromegaly. Pegvisomant treatment in a dose dependent manner results in normalization of IGF-1 levels in most patients. Thus, it is a very efficacious and safe drug. Since the GH/IGF-1 axis has been implicated in the progression of several types of cancers, many have suggested the use of pegvisomant as an anti-cancer therapeutic. In this manuscript, we will review the use of mouse strains that possess elevated or depressed levels of GH action for unraveling many of GH actions. Additionally, we will describe experiments in which the GHA was discovered, review results of pegvisomant's preclinical and clinical trials, and provide data suggesting pegvisomant's therapeutic value in selected types of cancer.

Keywords

Growth hormone receptor antagonist (GHA); GHA; Somavert; pegvisomant; GHR^{-/-} mice; FaGHRKO; cancer

1. Introduction

Although definitions differ, translational research typically refers to basic science research that is effectively “translated” into clinical benefits. This review describes several decades of

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research at the “bench” in the area of growth hormone (GH). Much of the bench discoveries originated through the study of several strains of genetically engineered mice with altered GH induced intracellular signaling. Analysis of these mice has led to a better understanding of the basic function of GH as well as its fundamental and tissue-specific role in various physiological processes. The discovery of the GH receptor antagonist (GHA) originated in mice, which resulted in a drug Somavert® (pegvisomant for injection) for the treatment of patients with acromegaly. In this review, we will summarize the general phenotypes of select mouse lines that have extreme alterations in GH action and will highlight data referring to cancer using these mice. An overview of the basic phenotypes of these mice is provided in Table 1 and pictures of the mice are shown in Figures 1 and 2. We will also compare the general features of these mice to cognate clinical populations. Finally, we will conclude by describing how pegvisomant may have applications directed toward several types of cancer.

2. GH excess

2.1. Genetically engineered mice with excess GH

Over thirty years ago, a transgenic mouse that expressed rat GH was first described (Palmiter et al., 1982). In subsequent years, multiple research groups have generated and characterized various species of GH transgenes in mice (e.g. ovine GH, human GH, bovine GH). While GH transgene expression results in a fairly uniform phenotype, there are some differences among the various transgenic lines that are likely due to diverse levels of GH expression as well as the species of the transgene. Regarding the latter point, human GH can bind to both GH and prolactin receptors whereas rat, bovine and ovine GH bind exclusively to the GH receptor. Thus, overexpression of human GH in mice results in a physiological state in which both GH receptors and prolactin receptors are simultaneously activated whereas rat, bovine, or ovine GH transgene expression elucidate the specific role of GH action.

GH transgenic mice have a phenotype that resembles the conditions found in acromegaly and will be more fully outlined in the subsequent section. As expected GH transgenic mice have elevated plasma levels of GH and insulin like growth factor-1 (IGF-1) (Kopchick et al., 1999). These mice also exhibit accelerated somatic growth starting around three weeks of age and remain larger than littermate controls through their adult life (Kaps et al., 1999; Knapp et al., 1994; Palmer et al., 2009; Palmiter et al., 1982) (Figure 1) The extreme elevation in circulating GH also results in hyperinsulinemia despite euglycemia (Kopchick et al., 1999; Palmer et al., 2009) that is likely due to GH’s diabetogenic effect. Reports detailing body composition of GH transgenic mice are inconsistent, but the discrepancy appears to be primarily due to the age of the mice when examined. Data in young GH transgenic mice are the most conflicting with male (3 months of age) and female mice (6 months of age) showing increases in fat mass in some studies (Eckstein et al., 2002; Kaps et al., 1999; Palmer et al., 2009; Pomp et al., 1996) and decreases in fat mass in others (Pomp et al., 1996). Again, the level or species of transgene expression may contribute to these discordant findings. Regardless, in adulthood (older than 4–6 months of age in males and > 6 months of age in females), GH transgenic mice are consistently reported to be lean with a significant reduction in the mass of all adipose depots (Berryman et al., 2004; Palmer et al., 2009). Transgenic mice also have altered adipokine and cytokine levels with reduced leptin and adiponectin (total as well as high molecular weight forms) levels as well as increased resistin in the circulation and TNF- α , and IL-6 levels in adipose tissue as compared to control mice (Berryman et al., 2004; Lubbers et al., 2012; Wang et al., 2007). The importance of GH in nutrient partitioning is highlighted by studies in which diet is altered in these mice. That is, GH transgenic mice fed a high fat diet are resistant to diet-induced obesity and gain substantial lean mass with the additional calorie intake (Berryman et al.,

2006; Olsson et al., 2005). In contrast, calorie restriction improves their insulin levels and alters lipid metabolism (Wang et al., 2007). Importantly, many of the reports, such as data generated for body composition (Palmer et al., 2009), show gender differences corresponding to transgene expression with females having a less dramatic phenotype than male transgenic animals.

GH transgenic mice show signs of premature aging and have drastically reduced (~50%) lifespans compared to littermate controls (Bartke, 2003; Wolf et al., 1993). The cause of their premature death is multifactorial. Tissue-specific pathological organ damage has been noted as a result of elevated GH levels. Examples include glomerulosclerosis and triglyceride deposition in the kidney, resulting in marked kidney damage (Doi et al., 1991; Kumar et al., 2011). There is also evidence of impaired cardiac and vascular function (Bohlooly et al., 2005; Bollano et al., 2000; Izzard et al., 2009; Miquet et al., 2011) as well as lipid abnormalities (Frick et al., 2001). Moreover, most GH sensitive tissues are enlarged in proportion to the larger size of the mouse, but several tissues, such as liver and spleen, experience disproportionate increases in size, which likely influences organ function (Liu et al., 1995; Machado et al., 2005; Orian et al., 1989; Quaife et al., 1989; Shea et al., 1987; Yang et al., 1993a; Yang et al., 1993b). Finally, GH transgenic mice have greater incidence of tumors. For example, hGH transgenic mice have early onset of mammary tumors (Cecim et al., 1994) although the role of prolactin versus GH signaling in these mice has not been resolved. Further, the hepatomegaly in these transgenic mice is linked to hypertrophy and hyperplasia of hepatocytes, which appears to contribute to an increase the frequency of tumors and dysregulation of several oncogenic pathways in this tissue (Miquet et al., 2013; Quaife et al., 1989). The chronic expression of IGF-1 results in a different phenotype, suggesting that the elevation of GH has a direct impact on several of the pathologies seen in GH transgenic mice (Quaife et al., 1989).

2.2. Excess GH in humans (acromegaly/gigantism)

Acromegaly is a hormonal disorder characterized by chronically elevated levels of circulating GH resulting in subsequent elevated levels of IGF-1. Typically, the condition is caused by a GH-secreting pituitary adenoma. As such, the treatment of acromegaly usually aims at removing the tumor with subsequent control of tumor growth as well as normalizing GH and IGF-1 levels. If GH and IGF-1 are in excess during childhood prior to closure of the epiphyseal growth plate and left untreated, then excessive growth or gigantism results. GH excess occurring in adulthood results in enlargement of extremities along with thickening of the skin, insulin resistance, sleep apnea, and cardiac abnormalities. The condition results in significant morbidity and increased mortality, as reviewed (Melmed, 2009; Melmed et al., 2009; Sherlock et al., 2011). The clinical presentation of these conditions share many characteristics with GH transgenic mice, including elevations in the activity of the GH/IGF-1 axis, increased lean mass and reduced fat mass, impaired glucose homeostasis and increased incidence of select cancers. However, there are important distinctions. First, the cause of chronic excess of GH in acromegaly, as stated above, is usually due to autonomous GH secretion from a pituitary adenoma whereas, in GH transgenic mice, the GH excess is due to transgene insertion resulting in ectopic GH secretion. Second, the level of circulating GH is often markedly greater in transgenic mice than what is observed in acromegaly. Finally, the transgenic mice have excessive GH gene expression for most of their postnatal life and, thus, resemble the excessive somatic growth associated with acromegalic-induced gigantism. Regardless, GH transgenic mice provide a valuable means to evaluate the impact of GH excess on specific tissues and organ systems.

3. GH resistance

3.1. Genetically engineered mice with disruption of the GHR gene

In attempts to better understand the numerous *in vivo* actions of GH, our laboratory generated the GH receptor gene disrupted mouse (GHR^{-/-}) in the mid 1990's (Zhou et al., 1997) (Figure 1). They also are termed global GHR^{-/-} mice. Since these mice are completely resistant to GH action, the unique phenotype of these mice provides key evidence as to the actions of GH. Thus, they have been and continue to be used to study the various activities of GH. In fact, since their creation, this mouse has been used in over 100 published studies (List et al., 2011).

As might be expected, GHR^{-/-} mice have an overall phenotype that is opposite to that of the GH transgenic mice previously described. They have a pronounced decrease in body size with extremely low circulating IGF-1 despite elevated GH levels (Zhou et al., 1997). Organ weights are proportionally decreased with the exception of liver and kidney, which are disproportionately smaller, and brain and select adipose depots, which are disproportionately larger (Berryman et al., 2010; Coschigano et al., 2003a). In addition to the obvious effects on body size, GHR^{-/-} mice have many other unique attributes that highlight the numerous roles of GH on metabolism (List et al., 2011). For example, GHR^{-/-} mice are exceptionally insulin sensitive with low levels of circulating insulin (Al-Regaiey et al., 2005; Bartke et al., 2004; Berryman et al., 2006; Bonkowski et al., 2009; Coschigano et al., 2003a; Dominici et al., 2000; Egecioglu et al., 2006; Hauck et al., 2001; Liu et al., 2004; Panici et al., 2009). Fasting blood glucose levels are decreased at younger ages in GHR^{-/-} mice compared to controls but tend to normalize at advanced ages (Al-Regaiey et al., 2005; Liu et al., 2004). Also, GHR^{-/-} mice are obese with a lifelong increase in percent body fat (Berryman et al., 2004; Berryman et al., 2006; Berryman et al., 2010). Consistent with their increased adiposity, GHR^{-/-} mice have elevated levels of leptin (Al-Regaiey et al., 2005; Berryman et al., 2004; Egecioglu et al., 2006). Adiponectin levels (both total and high molecular weight forms) also are increased in GHR^{-/-} mice, which is counterintuitive since adiponectin levels are usually negatively correlated with increased adiposity (Al-Regaiey et al., 2005; Berryman et al., 2004; Lubbers et al., 2012; Nilsson et al., 2005). However, since adiponectin is also positively associated with improved insulin sensitivity, this adipokine appears to be more highly correlated with insulin sensitivity in a GH resistant state. Another trait of the GHR^{-/-} mice that may in part explain the "healthy obesity" is that the adipose tissue is not uniformly increased, but rather almost exclusively increased in the subcutaneous white adipose tissue (WAT) depot (Berryman et al., 2004; Berryman et al., 2010). The concept that subcutaneous WAT might be healthier than other WAT depots has been supported by a number of studies and has been recently reviewed (Lee et al., 2013). Perhaps, the most striking observation from the GHR^{-/-} mice is that they are extremely long-lived (Coschigano et al., 2003a). In fact, GHR^{-/-} mice are officially recognized as the world's longest-lived laboratory mouse (<http://www.methuselahfoundation.org/>). The precise mechanisms involved in lifespan extension seen in GHR^{-/-} mice are the focus of many laboratories (including our own) in the aging field. However, reduction in neoplastic diseases are a major factor as 83% of WT mice die from neoplastic disease compared to 42% (49% reduction) of the GHR^{-/-} mice (Ikeno et al., 2009), as more fully described in the subsequent section on pegvisomant and cancer. In addition to decreased cancer, GHR^{-/-} mice also appear to be protected from diabetes (type 1 and 2). For example, streptozotocin-induced type 1 diabetes induction results in glomerulosclerosis, increases in glomerular volume, and increases in the ratio of mesangial area to total glomerular area in wild-type mice while GHR^{-/-} mice show none of these pathological changes (Bellush et al., 2000). Furthermore, GHR^{-/-} mice on a high-fat diet have lower levels of insulin and glucose

compared to WT mice on a high-fat diet despite a larger increase in adiposity (Berryman et al., 2006).

In attempts to further understand the health benefits of the GHR^{-/-} mice, our laboratory and others have recently generated and characterized various tissue-specific GHR gene disrupted mice using the Cre-LoxP system. Deletion of GHR in liver results in decreased circulating IGF-1, elevated GH, insulin resistance and hepatic steatosis (Fan et al., 2009). Two separate groups have deleted GHR in skeletal muscle and have reported different results. Using the Mef-2c promoter/enhancer to drive Cre recombinase expression, Mavalli and colleagues (Mavalli et al., 2010) observed peripheral adiposity and impaired glucose metabolism. However, when the muscle creatine kinase (MCK) promoter/enhancer is used, Vijayakumar and colleagues (Vijayakumar et al., 2012) reported reduced adiposity and improved glucose metabolism. Tissue specific disruption of GHR in pancreatic β -cell results in impaired insulin secretion (Wu et al., 2011). In our laboratory, we generated fat specific GHR disruption (FaGHRKO mice), which have a unique phenotype (List et al., 2013) (Figure 2). They have a normal GH/IGF-1 axis and glucose homeostasis, but are severely obese. Furthermore, FaGHRKO mice do not have increased adiponectin levels which are seen in global GHR^{-/-} mice, suggesting that GH does not act directly on adipose tissue to regulate adiponectin as previously suggested (List et al., 2013). Thus, the removal of GHR in specific tissues allows researchers to better evaluate organ-specific actions of GH. Collectively, these results demonstrate that disruption of GHR in specific tissues can dramatically influence glucose homeostasis and adiposity. Nonetheless, no single tissue specific deletion of the GHR results in an overall phenotype that resembles those of GHR^{-/-} mice and implies some level of important tissue crosstalk for the unique phenotype observed in these global GHR^{-/-} mice. Finally, the incidence and progression of neoplastic disease has not been reported in any of these GHR tissue specific gene disrupted mice.

3.2. Disruption of GHR in humans (Laron Syndrome)

Laron syndrome (LS; also known as GH insensitivity or resistance syndrome) is a genetic disorder most commonly caused by mutations in the GHR gene; however, mutations in other genes in the GH/GHR induced signaling pathway have also been described (Hwa et al., 2007; Hwa et al., 2011; Laron et al., 1993; Rosenfeld et al., 2007). Thus, the GHR^{-/-} mouse generated in our laboratory serves as an excellent model of LS in humans and a thorough comparison has been the subject of a recent book (Laron and Kopchick, 2011). GHR^{-/-} mice possess many similar traits to the human condition. These similarities include dwarfism, decreased IGF-1, increased GH, GH resistance, delayed sexual development, decreased muscle mass, decreased bone mineral density, decreased cranial bone size, hypoglycemia at younger ages, increased adiposity with greater increases in subcutaneous WAT, increased leptin and adiponectin levels and astonishingly, a resistance to diabetes and cancer (Bellush et al., 2000; Guevara-Aguirre et al., 2011; Ikeno et al., 2009; Laron and Kopchick, 2011).

There are a few potential differences between individuals with LS and GHR^{-/-} mice regarding glucose metabolism and longevity. These differences appear to depend on the particular population of LS individuals evaluated. Laron and colleagues suggest that LS individuals tend to develop hyperinsulinemia at advanced ages (Laron and Kopchick, 2011) while GHR^{-/-} mice have low levels of insulin throughout life. This difference may partially explain the discrepancy in lifespan. As stated earlier, GHR^{-/-} mice are extremely long-lived, while individuals with LS appear to have normal lifespans (Kopchick and Laron, 2011). Unfortunately, longevity studies in individuals with LS have limitations as this syndrome is extremely rare, making it difficult to study, and individuals with LS have a higher incidence of accidental and alcohol related deaths (Guevara-Aguirre et al., 2011). However, a recent study, which reports on a group of LS individuals from Ecuador, which

have been studied for over 22 years, shows that individuals with LS have dramatic reduction in insulin levels, improved insulin sensitivity, with significantly decreased HOMA-IR scores, and, as stated above, have no cases of cancer or diabetes (Guevara-Aguirre et al., 2011). Thus, it appears that humans with LS actually are indeed resistant to diabetes and cancer similar to the GHR^{-/-} mice.

4. GH Receptor Antagonist (GHA)

4.1. Development of the GHA

In 1991, our laboratory first described mice that express a GHA transgene (Chen et al., 1991a; Chen et al., 1991b; Chen et al., 1991c). The GHA transgene was a mutated bovine (b) GH gene in which the codon for glycine at position 119 was replaced with a larger amino acid, namely arginine, as described in detail below. Expression of the mutated GH transgene decreased GH action, resulting in dwarf mice (Figure 1), and was subsequently determined to be a classic receptor antagonist (Okada et al., 1992).

The fortuitous discovery of GHA is worthy of discussion as it not only has provided a better understanding of how GH interacts with its receptor but also has provided a novel mouse strain to assess the impact of a reduction in GH action in an animal system. However, most importantly, the discovery and characterization of GHA resulted in the development of an innovative drug, pegvisomant, used for the treatment of patients with acromegaly. This subject will be more thoroughly described later in this review. The GHA and resulting mice helped unravel several basic features of the GH/GHR interaction. For example, GHA helped to establish that the stoichiometry of GH:GHR is 1:2 (Cunningham et al., 1991; de Vos et al., 1992) and that two distinct regions of a single GH molecule interact with one receptor homodimer. At the time, one of the regions in GH responsible for binding to the GHR, referred to as site 1, had already been identified and shown to have a high affinity for the GHR (Bass et al., 1991; Cunningham et al., 1989; Cunningham et al., 1991; Cunningham and Wells, 1989; Fuh et al., 1992; Pearce et al., 1996). However, the second GH site remained obscure despite some hint that the region responsible for binding at Site 2 might be located in helix 3 since this region had previously been implicated in bone growth (Hara et al., 1978). A thorough assessment of the amino acids within helix 3 revealed that this helix had an 'imperfect' amphipathic configuration due to three amino acids. We hypothesized that engineering a perfect amphipathic configuration for helix 3 might result in greater growth promoting activity. Thus, we engineered a mutated bGH gene in which the three amino acid were converted to amino acids that would result in a perfect amphipathic helix (Q117L, G119R and A122D). Surprisingly, instead of augmenting GH's growth-promoting activity, the perfect amphipathic amino acid substitutions in helix 3 antagonized the function of GH *in vitro* and *in vivo* (Chen et al., 1991a; Chen et al., 1991b; Chen et al., 1990). Subsequently, only the G119R substitution was shown to be responsible for the GH antagonistic effect (Chen et al., 1991c). The mutation of the glycine codon in this helix allows the GHA site 1 to bind properly to the preformed GHR dimer on the surface of target tissues, but prevents proper or functional binding at GH's Site 2. This improper interaction fails to induce intracellular GHR signaling (Chen et al., 1991a; Chen et al., 1991c; Chen et al., 1990). Thus, GHA acts as a competitive inhibitor of endogenous GH or a classic receptor antagonist. For human GH, additional work revealed that similar antagonist effects could be achieved by substituting the corresponding glycine with lysine at position 120 (Chen et al., 1994). The study of GHA transgene expression in mice (GHA mice) has been helpful in understanding not only the potential impact of GH deficiency but also provides the foundational work for developing the GHA into a pharmaceutical agent.

4.2. Genetically engineered mice expressing the GHA transgene

The phenotype of GHA mice is generally intermediate between that of WT and GHR^{-/-} mice. For example, GHA mice are dwarf (Chen et al., 1991a; Chen et al., 1991c; Chen et al., 1990), but not as dramatic as seen in GHR^{-/-} mice (Coschigano et al., 2003a). Interestingly, while notably dwarf in early life, the body weight of male GHA mice gradually catches up to that of control mice (Berryman et al., 2013; Coschigano et al., 2003a). However, the increase in body weight in later adult life is due to marked increases in adiposity and not due to an increase in body length or lean tissue (Berryman et al., 2013). Like GHR^{-/-} mice, GHA mice exhibit a disproportionate increase in the subcutaneous fat depot (Berryman et al., 2004; Yakar et al., 2004) although other fat pads are also significantly increased in older GHA mice as compared to controls (Berryman et al., 2013).

Circulating levels of several factors have been reported in these mice. Serum IGF-1 and IGFBP-3 are reduced by ~80% and 30%, respectively compared to controls (Coschigano et al., 2003a). Glucose homeostasis is moderately improved in young mice with low to normal plasma levels of glucose and insulin (Coschigano et al., 2003a; Yakar et al., 2004). However, there is evidence of increasing insulin levels with advancing age in male GHA mice (Berryman et al., 2013; Coschigano et al., 2003a). Finally, adipokine levels are dramatically altered in GHA mice. Like GHR^{-/-} mice, total and high molecular weight adiponectin levels as well as leptin levels are significantly increased (Berryman et al., 2004; Berryman et al., 2013; Lubbers et al., 2013; Yakar et al., 2004). Notably, the leptin levels surge in later life (by 18 months) in GHA mice reaching levels ~5–7 fold higher than littermate controls (Berryman et al., 2013), which does not appear to occur in GHR^{-/-} mice (unpublished results). The relevance of this increase in the context of relatively consistent production of other adipokines is a subject of current investigation. Regardless, these collective data reveal an important effect of age on the overall phenotype of GHA mice.

Only one study has reported lifespan data for GHA mice (Coschigano et al., 2003a). Although this study reported a propensity for GHA mice to live longer than littermate controls, especially for females, the difference was not significant. Noteworthy is the fact that GHA mice are an exception to other mouse strains with decreases in GH action that experience major improvements in lifespan, such as the aforementioned GHR^{-/-} mice. However, it should also be mentioned that GHA mice, with their severe obesity especially with advancing age (Berryman et al., 2013), evidently do not live shorter than littermate controls. Thus, expression of the antagonist appears to confer some resiliency to the morbidities that are commonly associated with an obese state. Regardless, GHA mice are valuable for lifespan studies as they offer both an exception to the dogma that decreases in GH signaling increases lifespan and they defy the common pathologies that accompany obesity. Moreover, GHA mice could be considered a more clinically relevant mouse line than GHR^{-/-} mice; after all, repression of GH action is pharmacologically achievable with the use of a GHA (Somavert®, pegvisomant for injection) whereas total repression of GH action, as in GHR^{-/-} mice, is not and would not be clinically desirable.

5. Pegvisomant

5.1. Development of pegvisomant

The discovery of the GHA via the generation of the GHA transgenic mouse and confirmation *in vitro* led to the exciting speculation as to whether GHA could be used in clinical settings where a decrease in GH action was desirable. The obvious initial indication would be one where there are elevated serum GH levels, such as that found in patients with acromegaly. For a GHA to be effective, it would have to be present in serum at much higher levels than endogenous GH in order to effectively block the GHR. This posed a major

problem in that the serum half-life of GH and the GHA is approximately 15 minutes (Veldhuis et al., 2002), and endogenous GH is released in a pulsatile manner throughout the day. Therefore, the half-life of GHA would have to be greatly extended to continually block endogenous GH signaling. Since the serum half-life of GH was known to increase via conjugation of 5-kDa polyethylene glycol (pegylation, PEG) moieties (Clark et al., 1996), the same modification was investigated as a method to improve the pharmacokinetics of GHA. PEG binds to the primary amine groups in lysine residues as well as the N-terminus of proteins, thus increasing the size of the protein and reducing its clearance from serum. However, it was critical to avoid steric hindrance that could be caused by the addition of PEG molecules to lysine residues in regions of the protein that are important for GHR binding. In the case of GHA, there were two lysine residues at positions 168 and 172 that are critical for binding of site 1 to the GHR. In order to preserve binding affinity of PEG-G120K, these residues were mutated to K168A and K172R so as to prevent pegylation at these sites. Other modifications were made to increase PEG-G120K's binding affinity and to increase the potency of the drug. Six mutations resulting in the following amino acid substitutions, H18D, H21N, R167N, D171S, E174S and I179T, were introduced into that molecule and had previously been shown to increase the binding affinity at site 1 (Cunningham et al., 1991; Pradhananga et al., 2002). These site 1 amino acid substitutions also abolish the ability of the molecule to bind to the prolactin receptor (Goffin et al., 1999), thus reducing off-target pharmacologic effects.

The resulting recombinant protein, pegvisomant (Somavert® pegvisomant for injection), consists of a G120K molecule that is conjugated to PEG- at 4–5 sites and contains the above-mentioned 8 amino acid changes (Figure 3). Pegylation increases the serum half-life to approximately 100 hours (Muller et al., 2004) and acts as an effective GHA despite a 20-fold reduced affinity for the GH receptor due to the pegylation (Pradhananga et al., 2002).

5.2. Characterization of pegvisomant in vivo

In mice, very high doses of pegvisomant (10–20 times that normally given to treat acromegaly in humans) can reduce IGF-1 levels up to 70% (Kopchick et al., 2002). Also, GHA (as noted above) expressed in transgenic mice is effective at inhibiting GHR activity. However, human (h) GHA fails to antagonize GHR activity in GH-deficient dwarf rats when co-infused with GH (Mode et al., 1996). We are still puzzled by this observation and do not have an explanation.

Two studies designed to demonstrate the safety and effectiveness of pegvisomant in primates were conducted in rhesus monkeys. In the first study, a weekly bolus of 1 mg/kg of pegvisomant was injected subcutaneously into ovariectomized monkeys (Wilson, 1998). Serum IGF-1 levels were 61% lower in the pegvisomant injected animals compared to controls after 3 days and demonstrated maximal depression 10 days post injection. IGF-1 levels normalized to that of controls 14 days post injection. In the follow-up study, five ovariectomized monkeys were given the same dose of pegvisomant (Wilson, 2000). Within 24 hours of pegvisomant administration, IGF-1 levels began to fall, continued to decline through day 5, resulting in an overall 78% decline. These studies demonstrated that pegvisomant was well tolerated and effective in inhibiting endogenous GH action in primates.

5.3. Clinical use of pegvisomant/Somavert

5.3.1. Phase I-III clinical trials—Once pegvisomant was shown to be safe in primates and an effective suppressor of IGF-1, clinical trials in humans commenced. In a phase I, placebo-controlled study, 36 healthy young men received single subcutaneous dose of pegvisomant at 0.03, 0.1, 0.3 or 1 mg/kg (Thorner et al., 1999). Pegvisomant caused no

serious adverse events at any of the above doses. IGF-1 levels were decreased in a dose dependent manner for the 0.3 and 1.0 mg/kg doses. Maximum suppression for the 1 mg/kg pegvisomant dosage was observed 5 days post-injection with IGF-1 being reduced to nearly 50% of baseline levels. There were no substantial changes in circulating GH levels at any of the doses given. However, there was an extremely small, yet significant rise, in mean GH levels for the 1 mg/kg dose after 48hr in one of two GH quantification assays utilized, which may indicate reduced negative IGF-1 feedback on the pituitary.

The first study reporting treatment of acromegalic patients with pegvisomant was presented at The Endocrine Society's Annual Meeting in 1998 (van der Lely et al., 1998). Approximately two thirds of the patients had previously received surgery, radiotherapy and octreotide treatment. Medical treatment was withdrawn for three weeks from all patients prior to receiving pegvisomant. In this double blind, placebo-controlled phase II study, 46 patients with acromegaly were given placebo or a 30mg/week or 80mg/week dose of pegvisomant. The drug was well tolerated, and no patients were dropped due to adverse events. Total IGF-1 levels were reduced by 16% in patients receiving 30mg/week while total IGF-1 levels were reduced by 31% in those receiving 80mg/week after six weeks of treatment. In addition, free IGF-1 levels in patients receiving the 80mg/week dose was reduced by 47% after 6 weeks of treatment.

A phase III clinical trial, treating patients with acromegaly established the short term safety and efficacy of pegvisomant (Trainer et al., 2000). In this randomized, double-blind, placebo-controlled study, 112 patients with acromegaly received daily subcutaneous doses of pegvisomant for 12 weeks. Patients were selected whose serum IGF-1 level were greater than 1.3 times the upper level of the normal age-adjusted range and who had withdrawn from the use of other drugs. Patients were randomly selected to receive a daily dose of 10, 15 or 20 mg of pegvisomant or placebo. The drug was well tolerated, with only one patient classified as experiencing a serious adverse reaction. This patient had elevated levels of alanine and aspartate aminotransferases in the serum that returned to normal within eight weeks of discontinuation of treatment. All three treatment groups demonstrated a dose dependent decrease in serum IGF-1, IGFBP-3 and acid-labile subunit levels. Normalization of serum IGF-1 levels was achieved in 38% of patients receiving 10 mg/day and 82% of those receiving 20mg/day. In addition, the IGF-1 levels were normalized rapidly, with 75% of the maximal reductions occurring within the first two weeks of treatment. While three patients at the 20 mg/day dose did not achieve normal IGF-1 levels, they all showed substantial decreases in IGF-1 levels. Also, there was a dose-dependent increase in serum GH levels, coinciding with the decrease in serum IGF-1 levels. This reflects the negative feedback loop that helps regulate GH secretion by the somatotrophs of the pituitary. Evaluation of typical symptoms associated with acromegaly revealed improvement in the treatment groups with regards to decreases in soft tissue swelling, excessive perspiration and fatigue.

5.3.2. Initial long-term trials—While the phase III trial demonstrated the short term safety and efficacy of pegvisomant, questions regarding its long-term use remained. To address this issue, a new study enlisted a cohort of 160 patients for pegvisomant treatment of an average of 425 days (van der Lely et al., 2001). As in the above study, other medical interventions were discontinued and patients had serum IGF-1 level at least 1.3 times the upper limit of the normal range. Patients received initial doses of 10 mg/day of pegvisomant and that dose was increased or decreased in increments of 5 mg/day (with a maximum dose of 40 mg/day) until IGF-1 level normalization was achieved. For data analysis, patients were divided into groups receiving pegvisomant for 6 months, 12 months, or 18 months. Patients in the 12 month group were also included in the 6 month group while patients in the 18 month group were included in both the 6 and 12 month groups. Remarkably, serum IGF-1

levels were normalized in 97% of patients (n=90) treated with pegvisomant for 12 months or more. Similar to results in the short-term Phase III clinical trial study, the negative feedback loop that helps regulate GH secretion from the pituitary resulted in an increase in serum GH levels in all patient groups, rising by 12.5 ug/L in the 6 and 12 month group and by 14.2 ug/L in the 18 month group. GH levels returned to baseline levels within 30 days in patients withdrawn from pegvisomant and not placed on alternative medical therapy, demonstrating that pegvisomant treatment was responsible for the elevated GH levels. Antibodies to pegvisomant were detected in roughly 17% of patients, although none showed signs of tachyphylaxis.

Serum insulin and glucose concentrations decreased in all three treatment groups demonstrating an improvement in parameters of insulin resistance. This is extremely important since nearly one third of untreated acromegalics develop type 2 diabetes. The improvement of these important metabolic parameters via pegvisomant treatment of acromegalic individuals are benefits not observed when a somatostatin analog is used (Barkan et al., 2005). The effect of pegvisomant treatment on pituitary tumor growth was assessed by magnetic resonance imaging. During the early human clinical trials using pegvisomant, the fear of increases in pituitary tumor size was apparent since similar scenarios occurred when dealing with Nelson's syndrome following bilateral adrenalectomy (Nelson et al., 1960). It should be noted that bilateral adrenalectomy completely ablates cortisol from the individual while pegvisomant merely normalizes IGF-1 levels. The mean tumor volume was unchanged for the 131 patients treated with pegvisomant for which scans were available. Subsequent studies with more patients and longer-term follow up have demonstrated that pegvisomant treatment is not associated with tumor progression.

5.3.3. The long term trial: ACROSTUDY—Due to the new mechanism of action for pegvisomant, the Food and Drug Administration (FDA) and European Medicines Agency (EMA) required an ongoing observational survey to determine pegvisomant's long-term efficacy and safety. This study, carried out by Pfizer Inc, was started in 2004 under the name ACROSTUDY (Trainer, 2009).

As of February 2009, more than 792 patients had been included in the study with a mean of 3.3 years of pegvisomant treatment. The majority of patients were receiving daily injections of pegvisomant, and 67% were not undergoing additional treatment for acromegaly. After 12 months of treatment, average serum IGF-1 levels dropped from 518 ng/ml to 277 ng/ml. Normalization of serum IGF-1 levels was relatively constant at 62% regardless of the length of treatment. The 62% normalization rate was unexpectedly low given that the results from the phase III study demonstrated a 97% normalization rate (van der Lely et al., 2001). The reason for this discrepancy is believed to be due to a failure of physicians to increase the dose of pegvisomant in patients who do not show IGF-1 normalization (Trainer, 2009). This is evidenced by the fact that four out of five patients with higher than normal IGF-1 levels after three years of treatment remained on a dose of 20 mg/day or lower. Therefore, it is believed that IGF-1 normalization in the vast majority of patients can be achieved by vigilant pegvisomant dose titration. In addition, those patients taking other medical therapies in combination with pegvisomant did not exhibit increased IGF-1 normalization rates compared to those on IGF-1 monotherapy.

Of the more than 792 patients, adverse events were reported in 142 patients and 56 of those were attributed to pegvisomant therapy. The most common adverse event, which occurred in 29 patients (3.7%), was abnormal elevation of liver enzymes. Liver enzyme levels were returned to normal in 10 of these patients, half of which remained on pegvisomant and several of whom stopped treatment and subsequently restarted on pegvisomant. The 56 reported serious adverse events occurred in a total of 46 patients. Thirteen of these were

attributed to pegvisomant, with 9 due to elevated liver enzymes. Importantly, no patient treated with pegvisomant showed any evidence of sustained liver damage.

Pituitary tumor growth was reported as a serious adverse event in three patients. 411 patients underwent pre-treatment and post-treatment MRI imaging to investigate what effect, if any, pegvisomant has on pituitary tumor volume. Of these patients, 70 patients had a change in tumor size during treatment, with 22 showing an increase and 31 showing a decrease. Of those with lower tumor volume, 26 had been either previously treated with radiotherapy or were on combination therapy (either somatostatin analogue, dopamine agonist or both). It is believed that these additional treatments are the likely cause of tumor shrinkage. Of those with an increase in tumor volume, 6 had previously received radiation therapy and two had discontinued somatostatin analogue treatment. Tumor growth was observed before the initiation of pegvisomant treatment in two of the 22 patients. In 11 patients, additional review was unable to confirm a suspected increase in tumor size by the investigator.

6. GH/IGF-1 association with cancer

Decades of research have clearly established a central role of the GH/IGF-1 axis in postnatal growth. More recently, considering the ability of GH to promote cell proliferation and angiogenesis and inhibit apoptosis, many have turned their attention of the ability of GH/IGF-1 to be involved in several types of cancers. In that regard, a recent pathway-based approach of genome-wide association studies (GWAS) identified GH induced signaling pathways as the third highest pathway associated with breast cancer susceptibility (Menashe et al., 2010). Therefore, it is possible that inhibitors of this pathway, such as GHA, may prove to be effective therapeutics to inhibit the growth and/or progression of numerous cancers.

6.1. Human epidemiological studies

Numerous studies in humans have provided evidence that the GH/IGF-1 axis plays a role in the development and/or progression of human cancers. This literature is considerable and will not be reviewed here. However, we will point out three examples. A study depicting the association of height and cancer in over 300 individuals found a significant increase in the incidence of colon, breast and prostate cancer in taller individuals (Gunnell et al., 2001). Similarly, studies of individuals with elevated GH levels due to acromegaly indicated that there is a moderate risk in developing colorectal cancer (Melmed, 2009; Renehan and Brennan, 2008). As previously stated, Laron individuals have an inactivating mutation within their GHR gene, and as such, are insensitive to GH action. Data from 2 cohorts of Laron patients strongly implicate GH action in human cancer. In one cohort of 230 Laron patients, none of the individuals developed cancer (Steerman et al., 2011). In the second cohort located in Ecuador, out of a population of 99 Laron patients, only 1 patient exhibited a malignancy that turned out to be non-lethal in contrast to a 17% cancer incidence in control individuals (Guevara-Aguirre et al., 2011). Thus, there is an extraordinary and exciting reduction in the incidence of cancer in Laron Syndrome patients.

6.2. Evidence from animal studies

As discussed previously, a study by Ikeno (Ikeno et al., 2009) detailed the overall reduced occurrence of neoplasms in GHR^{-/-} mice throughout their lifespan and complements that seen in Laron patients. A longevity study was performed on GHR^{-/-} mice and their WT littermates. Histopathological examination was performed on the mice following their death. While neoplastic lesions are found in more than 90% of WT mice, such lesions are found in only 68% of the GHR^{-/-} mice. In addition, the tumor burden (number of different types of tumors) in the GHR^{-/-} mice is reduced by 47% compared to WT mice and shows that the

overall tumor incidence is reduced in GHR^{-/-} mice. As for the cause of death, 83% of WT mice die from neoplastic disease while only 42% (49% reduction) of the GHR^{-/-} mice die from such lesions.

There is mounting evidence from animal studies that implicate GH in oncogenesis, as recently reviewed by Chhabra, Waters and Brooks (Chhabra et al., 2011). However, mammary cancers are perhaps the best studied (Perry et al., 2008). A study utilizing the GHA mouse model demonstrated that inhibition of the GH action could provide striking protection from breast cancer *in vivo* (Pollak et al., 2001). In this study, 8week-old GHA and control mice were given a weekly gavage of the carcinogen, dimethylbenz(a)anthracene (DMBA) and monitored for tumor incidence. At the end of the 39 week study period, 68% of GHA animals remained without tumors, while only 32% of control animals were tumor free.

Another study utilizing the Spontaneous Dwarf Rat (SDR), which has an inactivating mutation in the GH gene Gh^{dr/dr}, was conducted to assess GH signaling in the progression of mammary carcinogenesis (Swanson and Unterman, 2002). Female SDR and control rats were subjected to chemical carcinogenesis with N-methyl-N-nitrosourea (MNU) or DMBA and observed for 180 days. All of the wild-type rats ($n = 10$) developed multiple mammary cancers (5.3/rat) while only 3 SDR rats ($n = 15$) developed cancers (0.2/rat). A subsequent study examined the effect of GH administration to SDR along with MNU treatment (Shen et al., 2007). In this study, SDR and control rats were injected with GH or vehicle following MNU exposure. Tumor occurrence did not occur in SDR while 80% of control animals and the GH-treated SDR group developed tumor at two months following MNU exposure. Importantly, cessation of GH injections from SDR rats resulted in the rapid regression of all tumors and subsequent resumption of GH injections resulted in a rapid resurgence of tumors at the original tumor sites. This work demonstrated that SDR rats treated with GH become vulnerable to mammary carcinogenesis and that these advanced rat mammary cancers are dependent on GH for their survival. In a similar study, SDR rats were injected with MNU and treated with GH with or without estradiol (E2) and progesterone (P4), with E2 and P4 alone, or with IGF-1 (Thordarson et al., 2004). As demonstrated above, the GH-treated SDR rats exhibited a 100% tumor incidence by 10 weeks of treatment, while only 50% of the IGF-1 treated SDR rats developed tumors by week 14. As expected, SDR animals treated with E2 + P4 had no tumors, while, paradoxically, GH + E2 + P4 treated animals also showed almost no tumor incidence. This suggests that steroid treatment blocked the action of GH possibly by reducing serum IGF-1 levels.

Our GHR^{-/-} mouse model has also been used to examine the role of the GH/IGF-1 axis in mammary carcinogenesis. GHR^{-/-} animals were crossed to C3(1)/TAg mice, a mouse line that develops estrogen receptor α negative mammary cancers (Zhang et al., 2007). While TAg/GHR^{+/+} animals developed an average of nearly 10 tumors per animal, TAg/GHR^{-/-} animals only showed ~3 tumors per animal. Even more striking than the number of tumors was the tumor size. TAg/GHR^{+/+} tumors grew to more than 10 times the volume of those found in GHR disrupted animals. These experiments demonstrate that the disruption of the GH/IGF-1 axis significantly retards TAg-driven mammary carcinogenesis, and once again, suggest that antagonists of GH induced signaling may be an effective strategy to inhibit the progression of estrogen-independent breast cancer. A similar study was used to examine the role of the GH/IGF in prostate carcinogenesis (Wang et al., 2005). In this study, GHR^{-/-} mice were again crossed with C3(1)/TAg mice. The resulting mice develop prostate neoplasia as a result of large T antigen expression that progresses to invasive prostate carcinoma. TAg/GHR^{+/+} and TAg/GHR^{-/-} mice were generated and sacrificed at 9 months of age. While seven of eight TAg/GHR^{+/+} mice developed prostate neoplasias, only one of

eight Tag/GHR^{-/-} mice developed such lesions suggesting that disruption of GH signaling confers resistance to prostate carcinogenesis.

6.3. Use of pegvisomant in cancer xenographs

The above studies as well as many confirming *in vitro* studies argue for the use of pegvisomant for selected cancers. In an early study, pegvisomant was examined for its impact on cancer cell xenographs in athymic nude mice (Divisova et al., 2006). Treatment of WT FVB/N mice with pegvisomant severely reduced mammary development and reduced mammary gland GH and IGF-1 signaling. In this study, human breast cancer cells (MCF-7) were grown as subcutaneous xenographs in the thoracic mammary fat pad until they reached a volume of approximately 100–200mm³. The mice were then randomized and injected daily with pegvisomant (250mg/kg) or vehicle. MCF-7 xenographs stopped growing upon pegvisomant treatment while MCF-7 control animals showed a steady increase in tumor size. In another study, pegvisomant was analyzed for its ability to inhibit the growth of human colorectal carcinoma cell lines in nude mice (Dagnaes-Hansen et al., 2004). In this study, Colo205 and HT-29 cell xenographs were grown in nude mice until they reached a volume of 90–105mm³ and 60–70mm³, respectively. The mice were then separated into 2 groups and injected every other day with either saline or pegvisomant at 60mg/kg. After 16 days of injections, pegvisomant treatment caused a 39% reduction in tumor volume and a 44% reduction in tumor weight in the nude mice with the COLO 205 colorectal cancer. However, pegvisomant had no effect on the HT-29 colorectal cancer and suggests the inhibition of GH/IGF-1 may represent a treatment for some colorectal cancers. In a third study, pegvisomant was assessed for antitumor activity against human meningiomas in nude mice (McCutcheon et al., 2001). Primary cultures of human meningioma tumors were xenografted into athymic mice. Mice were then injected with pegvisomant or vehicle for 8 weeks. Following treatment tumor growth was assessed. The mean tumor size in the vehicle injected cohort increased from 284mm³ to 350mm³ while those in the pegvisomant treated group decreased from 291mm³ to 198mm³. This study complemented their earlier work which demonstrated that the growth rate of GHR expressing primary meningioma cultures was decreased via the blockade of the GH/IGF-1 axis by B2036 (a nonpegylated version of pegvisomant) and suggested that the down-regulation of this axis may represent a potent therapeutical approach for this disease (Friend et al., 1999).

6.4. Additional human cancer studies

The role of the GH/IGF-1 axis has been studied on glioma cell lines (Friend et al., 2001). In one study both the GH and IGF-1 receptors were found to be expressed in 5 immortalized glioma cell lines. Treatment of two of these lines (U251 and U373) with IGF-1 resulted in increased mitogenesis as assessed by thymidine uptake. U251 cells were then xenografted into immunocompromised (*scid*) mice that had been crossed with the *lit* mouse strain. 10 of *scid/lit* heterozygous mice and 10 *scid/lit* homozygous were injected with 10⁶ U251 and 20 more injected with U373 cells. Five mice developed tumors in each group. The initial tumor volume in the U251 *lit/+* and *lit/lit* mice was 269 and 215 mm³, and volumes after 8 weeks were 2,589 and 1,819 mm³, respectively. Initial tumor volumes for U373 cells were 324 and 266mm³ in the *lit/+* and *lit/lit* mice, respectively, and after 8 weeks, the volumes were 526 and 599 mm³, respectively. These results were not statistically significant and suggest that a low GH/IGF-1 environment does not inhibit the growth of these tumors *in vivo*.

The possibility of the GH acting to promote growth in an autocrine/paracrine manner has also been reported by Lobie and colleagues. In one study, human endometrial carcinoma cell lines, RL95-2 and AN3, were stably transfected with a plamid designed to express human (h) GH (Pandey et al., 2008). Both AN3-vector (control cells) and AN3-hGH cell lines expressed equivalent levels of the GHR while AN3-hGH secreted detectable levels of hGH

into the culture medium. The AN3-hGH cells were found to grow significantly faster over a 14 day time period compared to control AN3-vector containing cells. The AN3-hGH cells also demonstrated enhanced anchorage-independent cell growth as indicated by colony formation in soft agar. Similar results were found with the RL95-2 cell line. Both control and RL95-2-hGH expressed the GHR, and both cell lines express endogenous hGH. However, RL95-2-hGH produced and secreted increased levels of hGH into the culture medium. Once again, the total cell number of RL95-2-hGH cells increased much faster than control cells over a 14 day period and anchorage-independent growth was also increased. In addition, antagonism of GH signaling by B2036 resulted in a reduction in oncogenic cell characteristics. When xenografted into athymic mice, RL95-2-hGH cells tumors grew significantly faster than RL95-2-vector tumors. In a second study, the autocrine effects of hGH on primary human mammary carcinoma cultures was examined. Primary cultures that were negative for hGH mRNA were highly sensitive to exogenous hGH stimulation. As expected, since there is no autocrine hGH stimulation in these cells, they were not affected by B2036. In contrast, B2036 significantly inhibited the proliferation of primary cultures that were positive for hGH mRNA. These cells theoretically would produce autocrine hGH.

7. Conclusion

The mouse strains generated in our laboratory with either enhanced or repressed GH action have elucidated much about the *in vivo* functions of GH including metabolic and adipogenic effects and possible contributions to oncogenesis. The generation and characterization of the GHA mouse strain resulted in the discovery of a new class of drugs, namely the GHA, of which Somavert® (pegvisomant for injection) was developed. It is currently used in the treatment of patients with acromegaly. Considering the body of evidence from *in vitro* and *in vivo* studies and human epidemiological reports, there is a growing belief that the GH/IGF-1 signaling pathway is capable of promoting growth of selected types of cancer. As such, inhibitors of this pathway using agents such as pegvisomant may prove to be an effective therapeutic intervention for certain cancer indications. Hopefully, future clinical trials will determine whether pegvisomant is of value in the treatment of these cancers.

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Abbreviations

GH	growth hormone
GHA	growth hormone receptor antagonist
TNF-α	tumor necrosis factor alpha
IL-6	interleukin-6
IGF-1	insulin-like growth factor-1
GHR$-/-$	growth hormone receptor gene disrupted mice
MCK	muscle creatine kinase
FaGHRKO	Fat specific growth hormone receptor knockout mice
WAT	white adipose tissue

LS	Laron Syndrome
bGH	bovine growth hormone
PEG	polyethylene glycol
DMBA	dimethylbenz (a) anthracene
SDR	spontaneous dwarf rat
MNU	N-methyl-N-nitrosourea
E2	estradiol
P4	progesterone

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Highlights

- Excess GH leads to poor health outcomes
- Removal of GH action is associated with resistance to diabetes and cancer
- The GHA was discovered using transgenic mice that express mutated GH genes
- Somavert® (pegvisomant for injection) is an approved drug for acromegaly
- GHA may have applications for selected cancer indications



Fig. 1. Photo of mice with altered GH action in the same genetic background (C57BL/6J)
A giant bGH transgenic mouse (mouse on left), wild-type mouse (2nd mouse from the left), a dwarf GHA transgenic mouse (3rd mouse from the left) and a dwarf GHR^{-/-} mouse (4th mouse from the left). These mice represent elevated, normal, decreased and absent levels of GH action, respectively.



Fig. 2. Photo of mice with removal of GHR specifically in adipose tissue
A wild-type mouse is shown on the left and a FaGHRKO mouse is shown on the right. FaGHRKO mice are obese demonstrating a direct action of GH on adipose tissue (List et al., 2013).

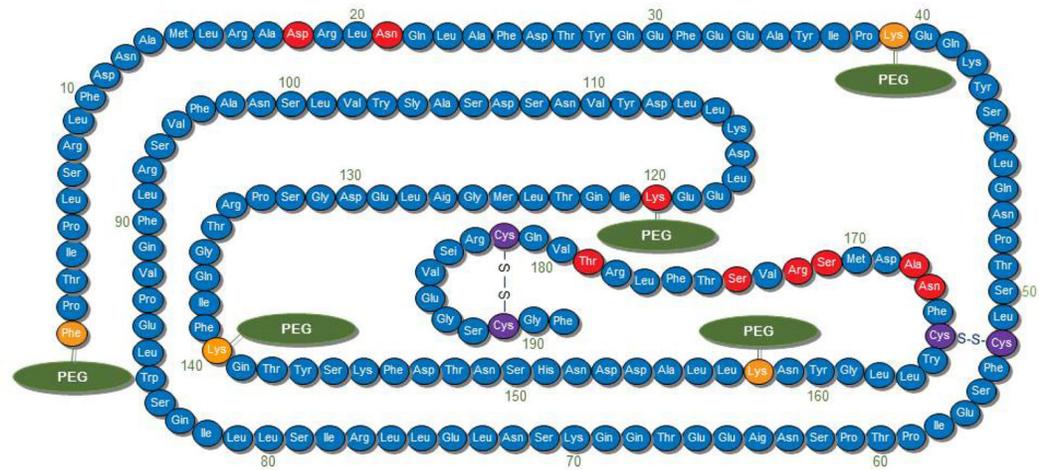


Fig. 3. Pegvisomant – the amino acid sequence

Five pegylation sites are indicated with a green PEG notation. The corresponding Lys residues are depicted in orange (except the Lys at position 120 which is shown in red). Other amino acid changes found in pegvisomant that differ from wild type hGH are displayed in red. Importantly, Lys at position 120 replaces Gly. It is this crucial amino acid substitution that generates the GHA.

Table 1

Phenotypic comparison of genetically modified mice produced in our laboratory with altered activity of GH

Mouse Model	bGH ^a	GHR-/- ^b	GHA ^c	FaGHRKO ^d
Level of GH signaling	↑↑	none	↓	↓fat only
Growth/Body Weight	↑↑	↓↓	↓	↑↑
Body Composition (Adult)				
% Lean Mass	↑↑	↓↓	↓	↔
% Fat Mass	↓↓	↑	↑↑	↑↑
GH/IGF-1 Axis				
GH	↑↑	↑	↑	↔
IGF-1	↑↑	↓↓	↓	↑/↔
IGFBP-3	↑	↓	↓	↔
Plasma levels				
Glucose	↑/↔	↓	↔	↔
Insulin	↑	↓↓	↔	↔
Leptin	↓	↑	↑↑	↔/↑
Adiponectin	↓	↑↑	↑	↓/↔
Reproduction				
Sexual Maturity (Time)	↓↓	↑	n.d.	n.d.
Fertility		↓	n.d.	n.d.
Morbidity				
Tumor Incidence	↑	↓↓	n.d.	n.d.
Cardiac/Vascular	↑	↔	n.d.	n.d.
Deficits				
Kidney Impairments	↑	↓	↓	n.d.
Lifespan	↓	↑	↔	n.d.

n.d. = not determined

^a(Bartke et al., 1998; Berryman et al., 2004; Berryman et al., 2006; Cecim et al., 1995; Lubbers et al., 2013; Olsson et al., 2003; Olsson et al., 2005; Palmer et al., 2009; Palmiter et al., 1982; Palmiter et al., 1983)

^bas reviewed in (List et al., 2011)

^c(Berryman et al., 2004; Chen et al., 1996; Chen et al., 1991a; Chen et al., 1991c; Coschigano et al., 2003b; Flyvbjerg et al., 1999; Lubbers et al., 2013; Berryman, 2013 #5358; Yakar et al., 2004)

^d(List et al., 2013)