I am honored and humbled to be one of the awardees of the 2014 A. Champalimaud Vision Award. I offer my heartfelt thanks to the Champalimaud Foundation President, Leonor Beleza, and to the Award Committee Members for this wonderful recognition.

I feel especially fortunate to have had the opportunity to witness my scientific discoveries move from the bench to the clinic. Scientific discovery is hugely exciting, but the ability to translate that work into potentially helping someone lead a better life is even more fulfilling. This Award is dedicated to the patients.

Introduction

The existence of factors capable of inducing growth of cells and tissues was hypothesized already at the beginning of the last century. In 1913, Carrel described the ability of tissue extracts to stimulate cell proliferation in cultured tissue explants. Early investigators also speculated that biochemical mediators are responsible for the growth of blood vessels associated with tumorigenesis and other pathological conditions (reviewed previously). In 1939, the observation that tumors transplanted in transparent chambers inserted in the rabbit ear induce rapid and extensive neovascular growth, led Ide et al. to postulate the existence of a tumor-derived “blood vessel growth stimulating factor.” In 1945, Algire et al. announced the seminal hypothesis that “the rapid growth of tumor transplants is dependent upon the development of a rich vascular supply.” In the late 1940s and in the 1950s, other investigators postulated the existence of a diffusible angiogenic factor (“Factor X”), produced in the ischemic retina. This hypothetical molecule was thought to be responsible for neovascularization associated with diabetic retinopathy and other retinal disorders. Notwithstanding these seminal studies, very little progress was possible at that time, given the daunting challenge of isolating growth factors, which typically are active at very low concentrations. Beginning in the 1970s, the availability of powerful protein purification techniques, combined with the development of cDNA cloning methodologies, enabled major advances. The greatest challenge at that time was purifying the proteins to homogeneity to obtain a partial amino acid sequence, which could be used to design probes suitable for cDNA cloning, thus, dramatically expanding the possibilities of investigating the molecules of interest.

In 1971, Folkman published an elegant synthesis of the aforementioned early studies and hypotheses, and also proposed that antiangiogenesis could be a novel strategy to inhibit tumor growth. This key hypothesis stimulated the search for regulators of angiogenesis. By the mid 1980s, several proangiogenic molecules had been identified and characterized, including epidermal growth factor (EGF), tumor growth factor (TGF–α, TGF–β), a-fibroblast growth factor (aFGF), bFGF, and angiogenin (reviewed previously). However, while these factors were able to promote angiogenesis in various bioassays, their role as endogenous mediators of angiogenesis remained uncertain, suggesting that in all likelihood some key molecules remained to be discovered (reviewed previously).
The Discovery of Vascular Endothelial Growth Factor (VEGF)

Independent efforts contributed to the discovery of VEGF. In 1983, Senger et al. reported an initial biochemical characterization of vascular permeability factor (VPF), a permeability-enhancing protein identified in the conditioned media of a guinea pig tumor cell line. However, the lack of amino acid sequence data precluded molecular cloning and establishing whether VPF was distinct from the known mediators of vascular permeability or from other proteins. Therefore, it is not surprising that limited progress in elucidating the significance and function of VPF took place during the next several years. Senger et al. reported the full purification of guinea pig VPF in 1990.

In 1989, we reported the isolation and cloning of a heparin-binding endothelial cell mitogen. This project began while I was a postdoctoral fellow at the University of California, San Francisco (UCSF) in the mid 1980s. At that time, I was able to isolate and culture a population of nonhormone-secreting cells from bovine pituitary, termed “follicular” or “folliculo-stellate” cells. Earlier investigators noted that they establish intimate contacts with the pituitary perivascular spaces, suggesting a role in the development or maintenance of the pituitary vasculature. In the course of these studies, I discovered that follicular cells release in their culture supernatants an endothelial cell mitogen. In 1988, I joined Genentech, where I had the opportunity to pursue the isolation of this mitogen. By early 1989, we were able to determine the amino terminal amino acid sequence of the purified protein. We found that this sequence was unique, since it had no match with known sequences in available databases. Because this molecule appeared to have growth-promoting activity selectively for vascular endothelial cells, we proposed the name “vascular endothelial growth factor” (VEGF). We then isolated bovine and human clones encoding multiple molecular species (isoforms) of VEGF, due to alternative mRNA splicing. In this early study, we identified three VEGF isoforms: VEGF_{121}, VEGF_{165}, and VEGF_{189}. Subsequent studies revealed the existence of additional VEGF isoforms (reviewed previously).

After our cloning paper was accepted for publication, we learned that a group at the Monsanto Company had submitted at approximately the same time a manuscript reporting on the cloning of VPF. These investigators described a human clone that encoded a protein identical to VEGF_{189}. This group followed up on the earlier work by Senger et al. and was able to isolate and sequence VPF. Therefore, it appeared that the same molecule possesses mitogenic and permeability-enhancing activities.

VEGF as a Key Regulator of Normal and Tumor Angiogenesis

The cloning of VEGF (today also known as VEGF-A following the discovery of several related molecules, VEGF-B, VEGF-C, VEGF-D, and PIGF) generated significant interest in the angiogenesis field, but it took several years before we could establish that VEGF was truly a pathophysiologically relevant mediator. It became clear that the VEGF isoforms are well suited to generate biochemical gradients, a requirement for angiogenesis in vivo, due to their differential diffusibility, which depends on their affinity for heparan-sulfate proteoglycans. A key question was whether VEGF has a role as an angiogenic factor in vivo. The earliest evidence that VEGF expression is temporally and spatially correlated with neovascularization was from a study published in 1990, where we examined the expression of VEGF mRNA in the rat ovary by in situ hybridization. We reported that the expression was low in the avascular granulosa cells, but was strongly upregulated in the highly vascularized corpus luteum. Furthermore, in 1992 we reported that the high affinity binding sites for VEGF are selectively expressed in endothelial cells in vivo.

The identification of the VEGF tyrosine kinase receptors represented a major milestone in the quest to understand VEGF action. In 1992, in collaboration with Lewis (Rusty) Williams at UCSF, we identified the \textit{fms}-like tyrosine kinase (presently known as VEGFR-1) as a high-affinity VEGF receptor. In the same year, Terman et al. identified a highly homologous tyrosine kinase receptor, known as KDR or VEGFR-2. It now is well established that VEGFR-2 is the main signaling VEGF receptor. Figure 1 illustrates the current view of the roles of the VEGF receptors and signaling pathways.

To elucidate the role of VEGF in vivo, we employed multiple strategies to inhibit its function. In 1993, we reported that administration of an anti-VEGF monoclonal antibody substantially reduced
the growth of several human tumor cell lines implanted in immunodeficient mice. These findings were unexpected at that time, as it was widely believed that tumor angiogenesis is multifactorial and, therefore, reflects the contribution of numerous mediators. They paved the way for subsequent clinical development of VEGF inhibitors as cancer therapeutics, including a humanized variant of this anti-VEGF...
antibody (bevacizumab), which has been approved for therapy of multiple tumor types. The impact of
these findings, however, went beyond the tumor angiogenesis field. They rapidly stimulated studies
aimed at directly probing the role of VEGF in other biological contexts, including intraocular neovascular-
ization.

Inactivation of the vegf gene in mice provided evidence for the crucial role of this molecule in the
early development of the vasculature. These findings, reported in 1996 by our group and by Carmeliet et
al., were nothing short of striking. Loss of even a single vegf allele resulted in defective vascular
development and early embryonic lethality. This phenotype was even more dramatic than that of
VEGFR2 knockout, which requires inactivation of both alleles to elicit a lethal phenotype. A few years
later, we developed VEGF loxP mice. Crossing these mice with various Cre-transgenic lines enables
conditional VEGF inactivation in specific cell types or tissues. These studies reinforced the notion that
VEGF is required for angiogenesis in many tissues and organs (reviewed previously). In parallel with
the genetic reagents, we developed soluble chimeric VEGF receptors (or “VEGF-traps”), which, unlike
many monoclonal antibodies, can block VEGF across species. Also, structure–function studies of
VEGFR-1 led to the discovery that of the seven extracellular Ig-like domains, domain two is the
critical element for high-affinity VEGF binding, enabling the design of smaller and more stable
soluble receptors. The availability of these tools allowed us to establish the role of VEGF in neovascularization associated not only with such essential physiological processes as organ and skeletal growth, or cyclical growth of the ovarian corpus luteum, but also with pathological retinal neovascularization.

Looking back at that period, it is almost impossible not to recall a sense of excitement permeating
through the angiogenesis field. After decades of largely descriptive work, it finally was possible to
unravel some of the secrets of this process and provide a molecular explanation for a variety of fundamental
pathophysiological processes. A commentary by Klagsbrun and Soker, published in 1993, reflects
this excitement. According to the authors, “…VEGF/VPF may be the best candidate for the principle
regulator of normal and tumor angiogenesis.” I feel extremely fortunate that my lab was at the forefront
of this revolution.

As pointed out above, by the early 1990s it was apparent that VEGF was implicated in normal as well
as in pathologic angiogenesis. Vascular endothelial growth factor also had several features consistent with
“Factor X,” being diffusible and selective for vascular endothelial cells. Also, in 1992 two studies
reported that VEGF mRNA expression is induced by hypoxia. Therefore, it is not surprising that
VEGF became the top candidate as a mediator of retinal ischemia-related neovascularization. In 1994,
in a collaborative study with Lloyd Aiello and George King at the Joslin Diabetes Center in Boston, we
tested this hypothesis. Taking advantage of sensitive assays newly developed in our group, we measured
the VEGF levels in the eye fluids from 164 patients. We found a striking correlation between VEGF
concentrations and active proliferative retinopathy associated with diabetes, occlusion of central retinal
vein, or prematurity. At approximately the same time, a French group also reported similar
findings.

Subsequent studies revealed that VEGF upregulation in the eye is not limited to ischemic retinal
disorders. In 1996, two groups reported the immunohistochemical localization of VEGF in choroidal
neovascular membranes from patients with wet age-related macular degeneration (AMD), the leading
cause of irreversible severe vision loss in the adult population.

Proof-of-concept studies supported the hypothesis that VEGF is, indeed, a major mediator of
intraocular neovascularization. As already mentioned, administration of chimeric soluble VEGF
receptors resulted in a marked reduction of retinal neovascularization in a mouse model of retinopathy
of prematurity. Also, in collaboration with Tony Adamis and Joan Miller, we tested the effects of the
anti-VEGF monoclonal antibody used in the cancer studies in a primate model of iris neovascularization
induced by central retinal vein occlusion. Similar to the tumor models, we observed a substantial inhibition of blood vessel growth following administration of the antibody. These effects were not limited to models of retinal ischemia. As described in the next section, wet (neovascular)
AMD became the primary clinical target of our anti-VEGF efforts in the eye. To this end, we engineered an affinity-matured antibody fragment (Fab) derived from the murine antibody parent of bevacizumab. Krzystolik et al. kindly agreed to support these efforts by testing this Fab, subsequently known as ranibizumab, in a primate model of choroidal neovascularization. These studies showed a dramatic inhibition of neovascularization and leakage following intravitreal administration of ranibizumab.

An Anti-VEGF Therapy for the Eye

The development path of anti-VEGF agents for wet AMD and other intraocular neovascular disorders has been described previously. We initially considered testing the intravenous administration of bevacizumab, but the possibility of cardiovascular adverse events in elderly patients led us to discard this possibility in favor of the intraocular route of administration. However, one could not rule out that long-term injection of full-length antibodies in human eyes might result in complement-mediated or cell-dependent cytotoxicity that might be triggered by interaction of the antibody Fc portion with receptors in inflammatory or immune cells. We felt that removing the Fc would be prudent. As already noted, we created an affinity-matured Fab variant of bevacizumab to further enhance its binding affinity. Genentech initiated the first clinical trial in subjects with wet AMD in February 2000. After encouraging data from phase I and phase II studies, ranibizumab was tested in pivotal phase III trials. Examining in detail the phase III studies with ranibizumab and other VEGF inhibitors (bevacizumab and aflibercept) is beyond the scope of this article, which mainly focuses on the discovery and science of VEGF. A very recent review summarizes such clinical trials and discusses a decade of clinical experience with VEGF inhibitors. Therefore, the cost and burden of chronic therapy in some cases limits benefit of anti-VEGF treatment. It is hoped that long-acting delivery technologies will address the gap in visual outcomes between clinical trials and “real life” clinical practice.

Conclusions

I am gratified and humbled that work that I initiated almost 30 years ago during my years as a postdoctoral fellow eventually resulted in a therapy for wet AMD and other intraocular neovascular disorders. The magnitude of the benefit, particularly the visual acuity gains, vastly exceeded my expectations, considering that previous treatments only slowed down the rate of vision loss.

Numerous trials currently are exploring a variety of novel therapeutic agents. These efforts give hope that combining VEGF inhibitors with agents that target additional pathways may go beyond the benefits achieved so far from targeting VEGF alone.

Acknowledgments

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References

28. Kim KJ, Li B, Winer J, et al. Inhibition of vascular endothelial growth factor-induced an-


