

## Impact of Tumor Cell VEGF Expression on the *In Vivo* Efficacy of Vandetanib (ZACTIMA™; ZD6474)

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**Abstract.** VEGF is the key player in tumor angiogenesis. In the current study, the impact of VEGF expression on the response of tumors to the VEGFR2 associated tyrosine kinase inhibitor vandetanib was evaluated. *Materials and Methods:* Human colon carcinoma (HT29) and murine squamous carcinoma (SCCVII) clonal cell lines expressing varying levels of VEGF were established and their response to vandetanib was assessed in tissue culture and as solid tumors. *Results:* Vandetanib treatment had no effect on tumor cell clonogenic cell survival in vitro but doses  $\geq 10$  nM significantly reduced endothelial cell migration. In vivo, tumors derived from cell clones expressing high levels of VEGF displayed significantly enhanced angiogenesis and more aggressive growth. An intradermal angiogenesis assay was used to demonstrate that a 4-day treatment with vandetanib (50 mg/kg/day) was able to significantly inhibit blood vessel growth induced by both parental and high VEGF-expressing tumor cell clones. In the HT29 tumor model, treatment response to vandetanib (50 mg/kg/day, Monday-Friday for 2 weeks) was greatest in xenografts derived from the highest VEGF-expressing cell clones. A similar trend was noted in the SCCVII tumor model. The present findings indicate that vandetanib therapy effectively counteracted the aggressive feature of tumor growth resulting from VEGF over-expressing tumor cells and suggest that such tumors may be particularly well suited for anti-VEGF interventions.

It is widely known that angiogenesis plays a key role in the progressive growth of solid tumors and the metastatic

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*Key Words:* VEGF, tumor vascularity, angiogenesis, anti-angiogenic therapy, vandetanib, ZACTIMA™, ZD6474.

cascade of cancer cells (1, 2). Vascular endothelial growth factor (VEGF) is the key mediator of angiogenesis (3). It is the only growth factor shown to be a specific and critical mitogen for endothelial cells while devoid of consistent and significant mitogenic effects in other cell types (4, 5). VEGF induces endothelial cell migration and tube formation (6), functions as a survival factor for endothelial cells (7) and increases vascular permeability which allows leakage of plasma proteins and the development of an extravascular matrix, thus further enhancing the niches for subsequent endothelial cell growth (8). VEGF also induces chemotaxis (9) and the expression of other pro-angiogenic factors (10) such as plasminogen activators (11) and collagenases (5, 12). Its expression is up-regulated in human cancer as a consequence of oncogene activation, loss of tumor suppressor function or hypoxia resulting from suboptimal perfusion (13, 14). Clinically, high levels of plasma VEGF have been associated with poor prognosis in patients with a variety of tumors (15-18).

Recognition of the central role of VEGF in tumor angiogenesis has led to the development of various strategies to inhibit VEGF signaling. Such strategies include antisense oligonucleotides, antibodies or receptor tyrosine kinase inhibitors (19-22). Of these approaches, the small molecule receptor tyrosine kinase inhibitors are particularly attractive. They are orally active and thus easily compatible with long term use. Vandetanib (ZACTIMA™; ZD6474) is a lead compound in this class. It is a once-daily oral anticancer drug that inhibits VEGF, epidermal growth factor (EGF) and RETS (rearranged during transfection) receptor signaling (23). *In vivo*, vandetanib antitumor effects have been demonstrated in a wide range of preclinical models including lung, prostate, breast, ovarian or vulval tumors (23-26). Moreover, vandetanib also has the potential to inhibit metastasis by prevention of primary tumor dissemination, as well as by inhibiting the growth of metastatic foci (27, 28). Currently, vandetanib is being evaluated in over 50 clinical trials of different tumor types ([www.clinicaltrial.gov](http://www.clinicaltrial.gov)).

Though evidence exists to suggest that increased VEGF expression levels may be related to more aggressive tumor behavior and poor treatment outcome (15-18), little is known about whether VEGF levels affect VEGF-directed anticancer therapy. Insights into this question are difficult to obtain from clinical trial data due to the significant inter-patient variations, heterogeneous tumor populations and differences in patient treatments associated with such studies. Preclinical investigations employing different tumor cell lines or models with varying levels of endogenous VEGF expression also cannot address this issue adequately because other differences between the models including genetic backgrounds, intrinsic drug sensitivity, tumor-host interactions or tumor immunogenicities, may be involved. To address this topic directly human colorectal tumor (HT29) and murine squamous cell carcinoma (SCCVII) models from clonal cell lines with altered VEGF expression levels were established and characterized (29). The clones were established by clonal selection from tumor cells infected with a recombinant adeno-associated virus (rAAV) vector containing the *VEGF*<sub>165</sub> gene. Consequently the lines carry the same genetic background in the presence of varying levels of stable VEGF expression. *In vitro* the clonal cell lines have similar growth characteristics and expression profiles of other key pro-angiogenic factors, but *in vivo*, tumors from high VEGF expressing clones grow faster, have increased blood vessel densities and are better oxygenated (29). The goal of the present study was to examine the therapeutic impact of targeting VEGF signaling using the small molecule VEGFR2 associated tyrosine kinase inhibitor vandetanib in tumors derived from cancer cell lines of varying VEGF expression.

## Materials and Methods

**Cell lines.** Human colon carcinoma (HT29) and murine squamous cell carcinoma (SCCVII) cells were obtained from the American Type Culture Collection (Manassas, VA, USA) and Stanford University (Palo Alto, CA, USA), respectively. Cells were maintained in Dulbecco's modified Eagle's medium (HT29) and alpha-minimal essential medium (SCCVII) supplemented with 10% fetal calf serum (FBS) and 0.01% (v/v) L glutamine and penicillin/streptomycin (Invitrogen, Grand Island, NY, USA). Human microvascular endothelial cells of the lung (HMVEC L) were obtained from Cambrex Corporation (Baltimore, MD, USA) and maintained in EGM 2 medium (Clonetics, Inc., San Diego, CA, USA) supplemented with 2 ng/mL VEGF. All cells were maintained at 37°C in a 5% CO<sub>2</sub> incubator.

**Generation of stable tumor cell lines by rAAV transfection.** HT29 or SCCVII clonal cell lines expressing various levels of VEGF were derived as previously described (29). Briefly, 1×10<sup>4</sup> HT29 or SCCVII cells were suspended in 50 µL of serum- and antibiotic-free medium. Recombinant adeno-associated virus (rAAV) vectors containing the cassette for human *VEGF*<sub>165</sub> and a neomycin resistance gene were then added at a multiplicity of infection (MOI) of 10,000. The cells were incubated for 3 h before changing to

selection media that contained 1 mg/mL geneticin for another 48 h at 37°C. Surviving cells were plated at a low density in 60-mm dishes to obtain clones. Stable cell lines were maintained in the appropriate cell media with the addition of 500 µg/mL geneticin.

**Animals and tumor models.** Either 1×10<sup>6</sup> HT29 tumor cells or 1×10<sup>5</sup> SCCVII tumor cells (in a volume of 0.02 mL phosphate-buffered saline) were injected intramuscularly into a single hind limb of 6- to 8-week-old female NCR nu/nu or C3H/HeJ mice (Frederick Cancer Research Facility, MD, USA), respectively. The mice were maintained under specific pathogen-free conditions (University of Florida Health Science Center Vivarium) with food and water provided *ad libitum*.

**Drug preparation.** Vandetanib was a gift from AstraZeneca Pharmaceuticals (Macclesfield, UK). Stock solutions of drug (10 mM) were prepared by suspending the agent in 10% (v/v) Tween 80 and N-2-hydroxyethylpiperazine-N'-2-ethane-sulfonic acid (HEPES). Working dilutions were made by serial dilution of the stock solution in sterile saline. All drug preparations were kept refrigerated in the dark and used within 1 week of preparation.

**Clonogenic cell survival.** HT29 or SCCVII cells were seeded into 60 mm tissue culture dishes and allowed to attach overnight. Vandetanib was added at doses ranging from 0-100 nM/mL for a period of 24 h. After treatment, the cells were trypsinized, counted and plated at three different concentrations in 60-mm petri dishes. Ten to fourteen days later, plates were stained with crystal violet and colonies of greater than 50 cells were counted with an aid of a dissecting microscope.

**Endothelial cell migration.** To assess endothelial cell migration *in vitro*, 1×10<sup>5</sup> HMVEC-L cells were seeded in modified Boyden chambers and treated with 5 to 500 nM vandetanib or vehicle control. After 48 h of treatment, the number of cells migrating through the 8 µm porous membrane was counted using a microscope at a ×20 magnification in 10 randomly selected fields on each filter.

**Intradermal tumor angiogenesis.** The extent of induction of angiogenesis by HT29 or SCCVII tumor cells and the degree of inhibition of this process by vandetanib treatment *in vivo* were determined using an intradermal tumor cell angiogenesis assay as described previously (30, 31). Briefly, 1×10<sup>5</sup> HT29 or SCCVII cells suspended in 10 µL of phosphate-buffered saline were injected intradermally at four sites on the ventral surface of a female nude mouse. One drop of 0.4% trypan blue solution was added to the injected mixture to aid visualization of the injection sites. Vandetanib was administered daily by oral gavage (50 mg/kg) for 4 days commencing on the day before tumor cell inoculation. Control mice received the blank vandetanib carrier. Mice were euthanized 3 days after tumor cell injection. The skin was carefully separated from the underlying muscle and the number of vessels counted using a dissecting microscope. Scoring of all of the reaction areas was carried out at the same magnification (×5) and only vessels readily detected at this magnification were counted. The sites of injection, recognized by local swelling and blue staining, were exposed by carefully removing fat or other tissue covering the area. All vessels that touched the edge of the tumor inoculates were counted. All the animals in the experiments were pre-coded and vessel counts in each animal were scored twice. The resultant data points for each treatment group were pooled for statistical analysis (Wilcoxon rank sum test).

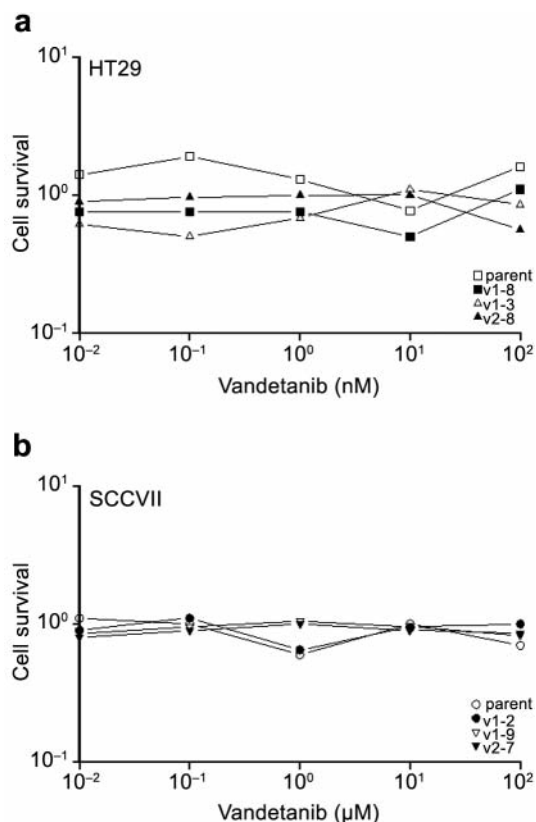


Figure 1. Clonogenic cell survival of HT29 (a) and SCCVII (b) parental and clonal cell lines assessed 24 h after treatment with a range of doses of vandetanib. Data shown are the average of 2 independent experiments.

**Assessment of tumor response to vandetanib.** When tumors reached a size of  $\sim 200$  mm<sup>3</sup>, mice were randomly assigned to control or treatment groups. Vandetanib was administered daily by oral gavage (0.1 mL/10 g body weight) for a period of 2 weeks (Monday to Friday). Control mice received vandetanib blank carrier by oral gavage. Following treatment, tumor size was measured by passing the tumor-bearing leg through a series of increasing diameter holes in an acrylic plate. The diameter of the smallest hole that the tumor bearing leg could pass through was recorded. The diameter was converted to a tumor volume using the following formula: tumor volume =  $1/6(\pi d^3) - 100$ , where  $d$  is the hole diameter and 100 represents a volume correction factor determined for a mouse leg without a tumor (32). The time for the tumors in the various treatment groups to grow to 5 times the starting size were recorded and compared (Wilcoxon rank-sum test).

## Results

Clonal cell lines expressing various levels of VEGF were established from both HT29 and SCCVII parental lines by viral infection as previously described (29). Three clones were ultimately chosen for each tumor model: the first

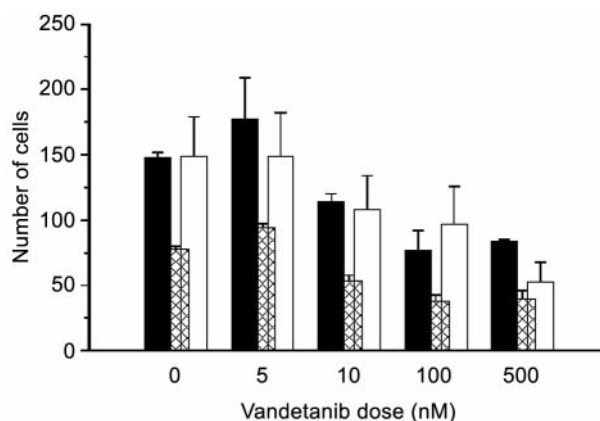


Figure 2. The effects of vandetanib treatment on endothelial cell migration through a modified Boyden chamber after 24-hour exposure to 5-500 nM vandetanib. Results for 3 independent experiments are indicated by the different shaded bars. In all cases, data obtained for vandetanib doses  $\geq 10$  nM were significantly different ( $p < 0.01$ ) versus controls (Student's *t*-test).

expressed VEGF at a level comparable to that of the parental cell line, the second expressed VEGF at an intermediate level, and the third expressed VEGF at a high level (29). In the HT29 model, the VEGF expression level of parental cells was 0.25  $\mu$ g/mL; clones selected for study expressed 0.3, 6.1 and 18  $\mu$ g/mL VEGF, respectively. These clones were named v1-8, v1-3 and v2-8 on the basis of the numbering system used during the original clone selection process (29). A similar nomenclature also was applied in the SCCVII model. For that cell line, neither the parental cells nor the v1-2 clone expressed detectable levels of human VEGF; clones v1-9 and v2-7 expressed 6.9 and 13.8  $\mu$ g/mL, respectively. Importantly, manipulation of the VEGF expression levels did not significantly affect the *in vitro* growth rate of these lines nor the expression levels of other pro-angiogenic growth factors including bFGF, PDGF, Ang 1 and Ang 2 (29). In addition, RT-PCR confirmed the lack of expression of VEGFR1 and VEGFR2 on these tumor cell lines (29).

Initial studies examined the inherent *in vitro* sensitivities of the parental and clonal cell lines to a range of doses of vandetanib. The results showed that doses as high as 100 nM (HT29) or 100  $\mu$ M (SCCVII) did not affect tumor cell survival in either the parental lines or any of the clonal cell lines (Figure 1). Endothelial cell (HMVEC-L) survival was also not affected by vandetanib doses up to 100 nM (data not shown). However, unlike tumor cells, vandetanib doses  $\geq 50$  nM did inhibit endothelial cell growth (32) and endothelial cell migration through a porous membrane was impaired at doses  $\geq 10$  nM (Figure 2).

To determine the effect of vandetanib treatment on HT29 or SCCVII tumor cell-induced angiogenesis *in vivo*, an

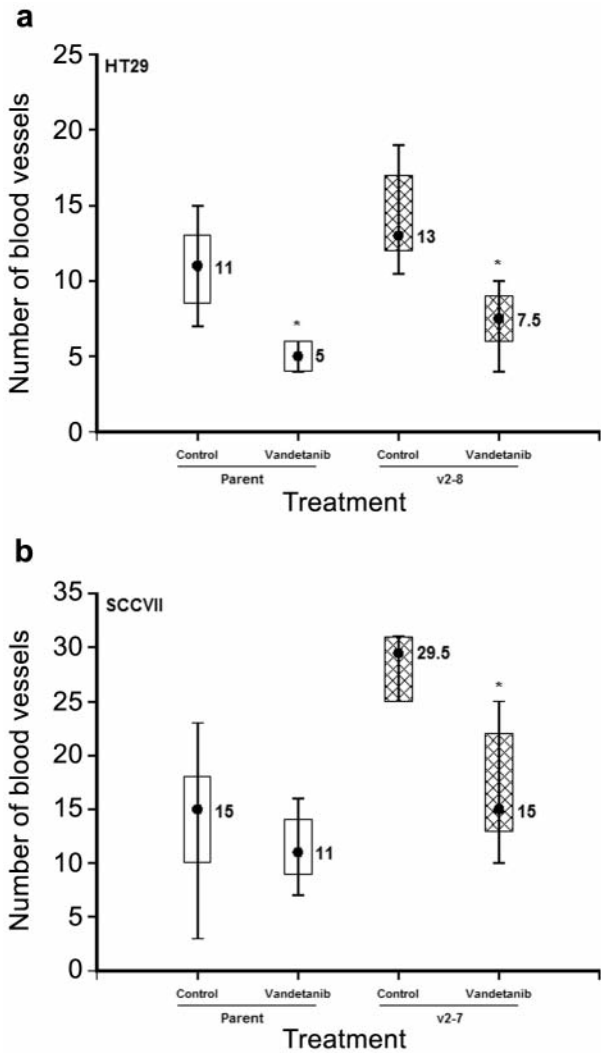


Figure 3. Nude mice (5/group) were injected intradermally with  $1 \times 10^5$  HT29 (a) or SCCVII (b) tumor cells at 4 sites per mouse. Three days later, the numbers of blood vessels induced at each inoculate were counted. In groups receiving vandetanib (50 mg/kg), the drug was given daily commencing on the day before tumor cell inoculation. Data shown are the median and 75th and 90th percentiles. Asterisks indicate statistical significance ( $p < 0.05$ , Wilcoxon rank-sum test) compared with the carrier-treated control groups.

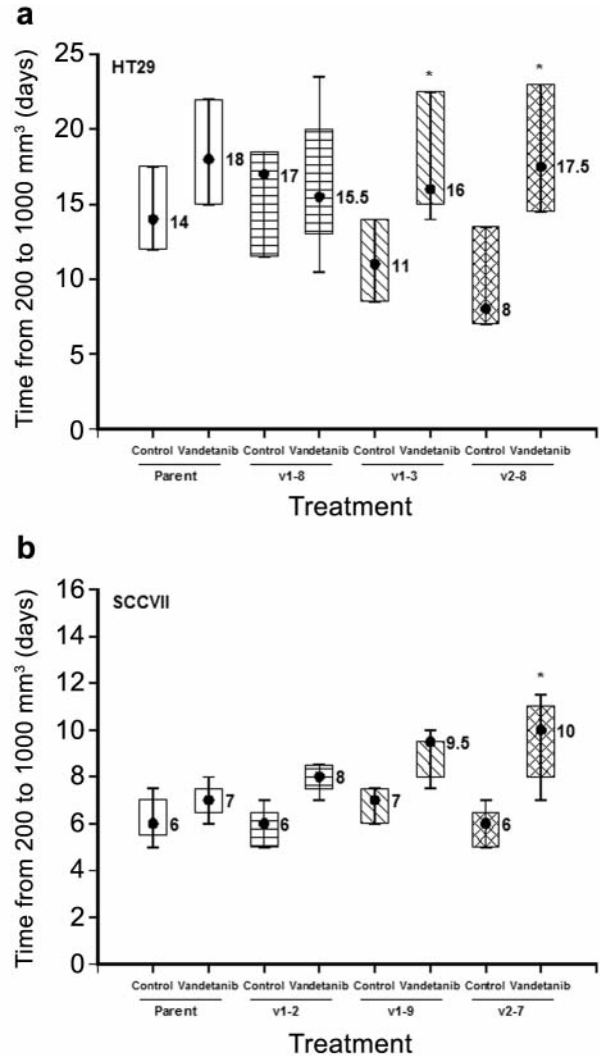


Figure 4. Response of tumors from HT29 (a) and SCCVII (b) parental and clonal cell lines to vandetanib treatment. Vandetanib (50 mg/kg) was administered on a daily basis (Monday-Friday) for a 2-week period commencing when the tumors reached a size of approximately  $200 \text{ mm}^3$ . Asterisks indicate a p-value of less than 0.05 when the growth delay of each group was compared via the Wilcoxon rank-sum test to that of the parental group.

intradermal assay (30, 31) was used. Mice received an intradermal inoculation of parental or high-expressing VEGF<sub>165</sub> (v2-8, HT29; v2-7, SCCVII) clonal tumor cells ( $1 \times 10^5$ ). Vandetanib (50 mg/kg/day) or its carrier was administered by oral gavage for 4 days commencing on the day prior to tumor cell inoculation. Three days later, all mice were euthanized and the skin flap containing the inoculation site was excised. The number of blood vessels intersecting each inoculate was counted. The results (Figure 3) showed that even this short-term treatment of a few days significantly decreased

the numbers of vessels induced by the tumor cell inoculates. Both the parental and high-expressing VEGF variants of the HT29 tumor model showed an approximately 2-fold reduction in the number of blood vessels induced (Figure 3a). A similar reduction in new blood vessel growth also was observed when mice injected with the high VEGF expressing SCCVII clone (v2-7) were treated with vandetanib (Figure 3b). These findings indicate that vandetanib treatment could be equally effective in blocking VEGF signaling in both low and high VEGF-expressing tumors.

It has previously been shown that vandetanib treatment can increase both the growth delay and the efficacy of ionizing radiation in HT29 xenografts (32). In the present study, the response of tumors established from parental HT29 cells was compared to that of tumors derived from HT29 cell clones with increased VEGF expression. Once established (~200 mm<sup>3</sup>), tumors were treated with 50 mg/kg oral doses of vandetanib daily for a two week period (days 1-5 and 8-12). In the HT29 tumor model the anti-VEGF therapy resulted in tumor growth delays which increased with increasing VEGF expression (4 days for the parental cell line; 10 days for the high-expressing VEGF clone) (Figure 4a). Mice bearing SCCVII tumors were only treated with vandetanib for a period of 5 days because the rapid growth rate of this tumor model prevented the intended 2-week course of vandetanib from being given. Even so, the 5-day vandetanib treatment did lead to the greatest responses in tumors derived from the higher VEGF expressing SCCVII clones (Figure 4b).

## Discussion

The central role of VEGF in tumor angiogenesis and disease progression has been well established (2, 3). VEGF impacts multiple steps in the angiogenic process including endothelial proliferation, migration, survival, and chemotaxis (3, 33). VEGF mRNA overexpression has been associated with an array of cancer types (34) and evidence exists to implicate VEGF as an indicator of poor prognosis in human cancer (15, 18, 35, 36). To ascertain whether VEGF expression influences the response of tumors to anti-VEGF therapy, clonal cell lines expressing varying VEGF levels (29) were grown as solid tumors which display a difference in tumor vascularity and hypoxia (29). The response of these tumor models to the small molecule VEGFR2 signaling inhibitor vandetanib was then assessed in the absence of confounding interpretational difficulties associated with inter-tumor model comparisons.

In *in vitro* assays, vandetanib had little effect on HT29 and SCCVII tumor cell survival (Figure 1) but did inhibit endothelial cell function at low doses (Figure 2), supporting the notion that the *in vivo* effects of this agent reflect its anti-angiogenic properties.

The present results (Figure 3) confirmed previous observations (29) that higher VEGF, expressing tumor cells (both HT29 and SCCVII clones) induced significantly more blood vessels when assessed in an intradermal angiogenesis assay (29). Vandetanib treatment reduced the tumor cell induced angiogenesis in both tumor cell types irrespective of their VEGF expression (Figure 3). For the HT29 model, inhibition of angiogenesis was observed for both high VEGF-expressing cells and parental tumor cells (Figure 3a). Inhibition of angiogenesis by vandetanib treatment was evident for both the high VEGF-expressing SCCVII clone

and the parental tumor cells (Figure 3b), although the suppression was significant only for the high VEGF-expressing cells (Figure 3b). These findings suggest vandetanib treatment can efficiently counteract the pro-angiogenic effect of VEGF and impair tumor-induced angiogenesis regardless of the level of VEGF expression.

Xenograft tumors grown from HT29 cells expressing higher levels of VEGF were increasingly more aggressive (grew more rapidly from 200 to 1000 mm<sup>3</sup>, Figure 4a) than those from parental HT29 cells, as has been reported previously (29). When such tumors were treated with vandetanib, the greatest tumor growth delays were observed in xenograft tumors derived from clonal cells expressing the highest levels of VEGF (Figure 4a). Since vandetanib treatment led to 16-18 day response in all HT29 tumor groups (parent, v1-8, v1-3, v2-8), this implies that at least in this model, this agent effectively counteracted the aggressive feature of tumor growth resulting from VEGF over-expression. Increasing tumor responses (longer growth delays) in tumors derived from the highest VEGF-expressing tumor clone (v2-7) treated with vandetanib were also observed in the SCCVII tumor model (Figure 4b).

The results of this study indicate that vandetanib treatment may overcome the aggressive tumor behavior associated with overexpression of VEGF. In both the human colorectal model HT29 and the rodent carcinoma model SCCVII, greater *in vivo* response to vandetanib was observed with increasing levels of VEGF expression. Taken together these findings support the use of VEGFR2-associated VEGF signaling inhibition with small molecules such as vandetanib in cancer patients with high levels of VEGF expression.

## Acknowledgements

This work was supported by U.S. Public Health Service grant CA089655. The authors thank Chris Pampo and Sharon Lepler for excellent technical assistance.

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Received October 31, 2008  
 Revised December 12, 2008  
 Accepted March 11, 2009