CD133 positive endothelial progenitor cells contribute to the tumour vasculature in non-small cell lung cancer

W Hilbe, S Dirnhofer, F Oberwasserlehner, T Schmid, E Gunsilius, G Hilbe, E Wöll, C M Kähler

Aims: Recent results generated in a mouse model suggest that tumour angiogenesis/vasculogenesis can be initiated and maintained by bone marrow derived endothelial progenitor cells. This present study investigated the distribution and frequency of CD133 positive endothelial progenitor cells in patients with non-small cell lung cancer (NSCLC) (tumour tissue and tumour free lung regions) and healthy controls using fresh frozen specimens. The novel marker CD133 identifies human haemopoetic precursor cells, in addition to human endothelial progenitor cells.

Methods: Seventy nine lung cancer specimens and 66 adjacent histologically tumour free tissues of the same patient cohort were analysed; 11 postmortem specimens from control patients who did not suffer from malignant disease served as controls. Cryostat sections were stained for CD133, CD31, vascular endothelial growth factor receptor 2 (VEGFR-2; KDR), p53, and the proliferation marker Ki-67, and the correlations were analysed.

Results: Forty three of 63 evaluable tumour specimens had increased numbers of CD133 positive cells and in some cases capillary forming CD133 positive structures were detectable. In addition, 30 of 63 specimens had raised expression of KDR and 29 of 63 had increased MVD. Increased CD133 expression marginally correlated with raised KDR expression but not with p53 and Ki-67.

Conclusion: A significant increase in CD133 positive cells was documented in patients with NSCLC, suggesting an involvement of endothelial progenitor cells in tumour vasculogenesis and tumour growth in these patients.

Angiogenesis is essential for tumour growth and metastatic spread and represents an important prognostic indicator in non-small cell lung cancer (NSCLC). Because different anti-angiogenic substances are currently under development and some of them have already been tested in clinical trials, the determination of the neoangiogenic potential of malignant tumours is of pivotal interest.

The growth of mature endothelial cells (ECs) depends on different growth factors leading to the recruitment of ECs from neighbouring pre-existing capillaries. Vascular endothelial growth factor (VEGF) is one of the key stimuli of endothelial proliferation and migration, and thus plays an essential role in physiological and pathological angiogenesis, and has also been reported to correlate with the neoangiogenesis occurring during cancer. Furthermore, VEGF is one of the key molecules in recruiting and activating endothelial progenitor cells (EPCs) from the bone marrow. In humans, increased immunolocalisation of VEGF in NSCLC specimens seems to be associated with a poor outcome. However, the impact of increased VEGF expression alone on tumour angiogenesis is ambiguous, and the determination of VEGF alone seems to be an inadequate means of defining angiogenic activity in NSCLC.

Furthermore, recent evidence suggests that the recruitment of ECs from neighbouring pre-existing capillaries is not the only source of increased tumour vascularisation during cancer spread. Lyden and colleagues demonstrated in a mouse model that the recruitment of bone marrow derived endothelial precursors plays an essential role in tumour vascularisation and tumour growth. These bone marrow derived endothelial cells are thought to contribute to angiogenic processes in cancer tissue and also to have the capacity to form new vessels—a process known as vasculogenesis. EPCs represent an immature subset of CD34 positive cells expressing the 120 kDa glycosylated polypeptide CD133. In contrast, mature ECs present in pre-existing vessels are CD133 negative. Interestingly, a more recent study reported that increased numbers of circulating CD133 positive EPCs occur in patients with breast cancer. These findings support the hypothesis of a functional role of EPCs in vascularisation and growth of human solid organ tumours.

“Endothelial progenitor cells represent an immature subset of CD34 positive cells expressing the 120 kDa glycosylated polypeptide CD133”

Until now, the standard method of measuring tumour angiogenesis has been the determination of intratumorous microvessel density (MVD), which is closely correlated with tumour growth and postoperative prognosis. Thus, patients with highly vascularised tumours suffer from early metastatic disease and reduced survival.

Multiple markers have been found to be altered in patients with NSCLC and some of them are thought to influence vascularisation of the tumour. Mutated p53 has been shown to potentiate the induction of VEGF secretion and to reduce the expression of the antiangiogenic factor thrombospondin. Bcl-2 and p53 have been shown to regulate VEGF mediated angiogenesis in NSCLC.

In addition, the proliferation associated marker Ki-67 is an essential prognostic parameter for NSCLC.

Abbreviations: EC, endothelial cell; EPC, endothelial progenitor cell; MVD, microvessel density; NSCLC, non-small cell lung cancer; VEGF, vascular endothelial growth factor; VEGFR, vascular endothelial growth factor receptor.
Here, we show for the first time that increased numbers of CD133 positive EPCs are found in NSCLC specimens and that they may contribute to tumour vasculogenesis. Furthermore, the relation between this increase and VEGF receptor 2 (VEGFR-2; KDR) expression and its correlation with MVD, proliferative activity, and p53 expression is presented.

PATIENTS AND METHODS

Patients

Fresh tumour tissues from 79 patients with NSCLC were obtained for routine diagnosis. All patients were treated surgically with curative intention. One portion of the sample was fixed in 10% formalin and another part was immediately snap frozen in liquid nitrogen and stored at −80°C for immunohistology. All tumour specimens were evaluated histopathologically according to the World Health Organisation classification.

As a result of an alveolar pattern of tumour infiltration, with consecutively increased numbers of capillaries, 16 patients were excluded from further evaluation, because in these cases the determination of MVD could have produced false positive results. In the remaining 63 patients, the histological diagnosis was as follows: 25 cases of adenocarcinoma, 34 cases of squamous cell carcinoma, two cases of large cell carcinoma, and two cases of carcinoid tumour. According to UICC recommendations, 42 patients were classified as stage I, 13 patients as stage II, and eight patients as stage III. Patients in advanced clinical stages were treated with chemotherapy and/or radiotherapy.

Immunocytochemical staining

Frozen specimens were cut at 3–5 μm in a cryostat, fixed, incubated with monoclonal antibodies at appropriate dilutions, and then incubated with an alkaline phosphatase complex (D651; Dako, Glostrup, Denmark), as reported previously. All antibodies were diluted with phosphate buffered saline with 1% bovine serum albumin. The following monoclonal antibodies were used: CD133/1 (clone AC133; Miltenyi Biotec, Bergisch Gladbach, Germany); anti-VEGFR-2 (clone KDR-2; Sigma-Aldrich, St Louis, Missouri, USA); CD31 (clone WM59; Serotec, Kiddington, UK); PAb 1801 to detect p53 (clone 1801; Pharmingen, San Diego, California, USA), and Ki-67 (Dakopatts, Glostrup, Denmark).

Immunohistochemical evaluation

Slides were evaluated independently by two of the authors (WH and SD) using a semiquantitative method on a Zeiss AXIOSKOP 2 microscope. Immunopositive cells were counted from representative areas of the sections and expressed as a percentage. The intensity of immunostaining was categorised as follows: −, negative; +, low; ++, moderate; and ++++, high.

Cutoff values were defined by analysing adjacent histologically confirmed tumour free areas of 66 patients with NSCLC (mean percentage ± 2SD): CD133, 0.7%; VEGFR-2 (KDR), 12.3%; p53, 1.3%; and Ki-67, 2.7%. For VEGFR-2 expression a second cutoff was introduced, because the calculated value (12.3%, “high expression”) could not distinguish between the subgroups sufficiently. Therefore, the cutoff value was defined by using the median value of the tumour free cohort (2.5%, “low expression”). For CD31, the cutoff was defined as the median number of positive cells of the three most intense areas: 15%. Only cases with moderate (++) or high (++++) staining intensity and a percentage of immunopositive cells above the cutoff point were scored as “positive”.

Microvessel counting was performed in accordance with previous reports, with slight modifications, using CD31. EPCs localised in capillary structures were characterised further by CD133 and VEGFR-2 (KDR). For all three tested antibodies, the most intense regions were identified under low magnification (×100), after which the positive cells were...
counted under higher magnification (×400) in these regions. Ki-67 and p53 analysis was performed on the whole cell population.

Summary of the procedure for immunohistochemical analysis
(1) Counting the number of “immunopositive cells” expressed as a percentage.
(2) Assessing the intensity of immunostaining (−, negative; +, low; ++, moderate; and ++++, high).
(3) “Positive cases”: moderate (+++) or high (++++) staining intensity and a percentage of “immunopositive cells” above the cutoff points.

Statistical analysis
The Pearson χ² test was used to compare the results of two or more subgroups. Significance was determined by means of 95% confidence intervals. All statistical procedures were performed using SPSS, version 7.5, statistical software (SPSS Inc, Chicago, Illinois, USA).

RESULTS
Patients
The median age of the 63 patients evaluated was 62 (range, 37–79) years at the time of surgery. The male : female ratio was 4.2 : 1 (51 : 12). The median observation time was 17 (range, 1–95) months, the median survival was not reached, and 18 patients died.

The analysis was based on three different groups of tissues, namely:

(1) Tumour specimens of patients with NSCLC (tumour group).
(2) Tumour free specimens of patients with NSCLC (tumour free group).
(3) Tumour free specimens of non-tumour patients (healthy control group).

Immunohistochemistry
Tables 1 and 2 give the results of the immunohistochemical staining. Increased numbers of CD133 positive cells were found in 43 of 63 cases within the tumour group, and some cases revealed capillary forming structures positive for CD133 staining (figs 1–4). Overall, the median number of CD133 positive cells within the tumour specimens was 0.2% (range, 0–50%) and 43 were over the cutoff value (0.7%). In contrast, in tumour free samples (n = 66) of the same NSCLC patient cohort, only single CD133 positive cells were detected (median, 0%; range, 0–1%) and no CD133 positive capillary structures were seen. The same data were obtained in the control group (median, 0%; range, 0–1%). In the tumour group CD133 expression correlated marginally with VEGFR-2 expression (cutoff value of 2.5%; p = 0.053).

Increased numbers of VEGFR-2 (KDR) immunopositive cells were detected in 30 of 63 samples within the infiltrating tumour using a cutoff value of 2.5%. The median number of VEGFR-2 positive cells was 2% (range, 0–20%). VEGFR-2 expression (median, 2.5%; range, 0–20%) in the tumour free regions of these patients revealed similar values: 32 of 66 scored positive. Applying the calculated cutoff value (12.3%), five patients in the tumour group and four in the tumour free cohort had raised numbers of VEGFR-2 positive cells. Similar values were detected (2% VEGFR-2 positive cells, range, 0.2–20% number of positive cases, four of 11 (cutoff, 2.5%) or one of 11 (cutoff, 12.3%)) in the control group. There were no significant differences between the three cohorts analysed. When a lower cutoff value was applied, the distribution of VEGFR-2 positive cases within the three groups still did not differ (cutoff, 2.5%: 36% control group v 48% tumour free group v 48% tumour group).

Increased MVD (as assessed by CD31 expression) was detected in 29 of the 63 tumour samples. MVD correlated with tumour grade (p = 0.033). In detail, MVD positive cases revealed a higher rate of grade III tumours (15 of 29) when compared with MVD negative tumours (nine of 29).

The frequency of Ki-67 and p53 expression did not correlate with the angiogenic parameters (CD133, CD31/MVD, and VEGFR-2 (KDR)). Within the analysed clinical parameters (sex, histology, and stage) the angiogenic markers were distributed equally in the tumour cohort (table 1).

<table>
<thead>
<tr>
<th>Table 1</th>
<th>Comparison of angiogenic markers with clinical parameters</th>
</tr>
</thead>
<tbody>
<tr>
<td>No. (%)</td>
<td>MVD</td>
</tr>
<tr>
<td>No. of positive cases</td>
<td>29</td>
</tr>
<tr>
<td>Diagnosis</td>
<td></td>
</tr>
<tr>
<td>AC</td>
<td>11 (38%)</td>
</tr>
<tr>
<td>SCC</td>
<td>17 (59%)</td>
</tr>
<tr>
<td>Other</td>
<td>1 (3%)</td>
</tr>
<tr>
<td>Stage</td>
<td></td>
</tr>
<tr>
<td>Stage 1</td>
<td>17 (59%)</td>
</tr>
<tr>
<td>Stage 2</td>
<td>6 (21%)</td>
</tr>
<tr>
<td>Stage 3</td>
<td>6 (21%)</td>
</tr>
<tr>
<td>Died</td>
<td>9 (31%)</td>
</tr>
</tbody>
</table>

AC, adenocarcinoma; MVD, microvessel density; SCC, squamous cell carcinoma; VEGFR-2 (KDR), vascular endothelial growth factor receptor 2.
**Table 2** Expression profiles of the three angiogenic markers analysed

<table>
<thead>
<tr>
<th></th>
<th>Control group</th>
<th>Non-tumour specimens from NSCLC group</th>
<th>NSCLC specimens</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number CD133</td>
<td>11</td>
<td>66</td>
<td>63</td>
</tr>
<tr>
<td>Number of positive cells (median, range)</td>
<td>0% 0–1% 0% 0–1% 0.2% 0–50%</td>
<td>0% 0% 0% 0% 0% 0%</td>
<td>0% 0% 0% 0% 0% 0%</td>
</tr>
<tr>
<td>VEGFR-2</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Number of positive cells (median, range)</td>
<td>2% 0.2–20% 2.50% 0–20% 2% 0–20%</td>
<td>2% 0.2–20% 2.50% 0–20% 2% 0–20%</td>
<td>2% 0.2–20% 2.50% 0–20% 2% 0–20%</td>
</tr>
<tr>
<td>Cases above cutoff value (0.7%)</td>
<td>0% 0% 0% 0% 0% 0%</td>
<td>0% 0% 0% 0% 0% 0%</td>
<td>0% 0% 0% 0% 0% 0%</td>
</tr>
<tr>
<td>Cases above the cutoff values</td>
<td>0% 0% 0% 0% 0% 0%</td>
<td>0% 0% 0% 0% 0% 0%</td>
<td>0% 0% 0% 0% 0% 0%</td>
</tr>
<tr>
<td>High expression: cutoff 12.3% (number, %)</td>
<td>4 36% 32 48% 30 48%</td>
<td>4 36% 32 48% 30 48%</td>
<td>4 36% 32 48% 30 48%</td>
</tr>
<tr>
<td>MVD (CD31)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Number of positive cells (median, range)</td>
<td>NE NE 6% 0–32%</td>
<td>NE NE 6% 0–32%</td>
<td>NE NE 6% 0–32%</td>
</tr>
<tr>
<td>Cases above cutoff value (15) (number, %)</td>
<td>NE NE 29 46%</td>
<td>NE NE 29 46%</td>
<td>NE NE 29 46%</td>
</tr>
<tr>
<td>Cases above the cutoff values</td>
<td>NE NE 29 46%</td>
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</tr>
<tr>
<td>Low expression: cutoff 2.5% (number, %)</td>
<td>0% 0% 1% 2% 25 63%</td>
<td>0% 0% 1% 2% 25 63%</td>
<td>0% 0% 1% 2% 25 63%</td>
</tr>
<tr>
<td>Number of positive cells (median, range)</td>
<td>0% 0% 0% 0% 0% 0%</td>
<td>0% 0% 0% 0% 0% 0%</td>
<td>0% 0% 0% 0% 0% 0%</td>
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<tr>
<td>Cases above cutoff value (0.7%)</td>
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<td>0% 0% 0% 0% 0% 0%</td>
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Comparison of the three cohorts: tumour free control group ("control group"), tumour free specimens from patients with NSCLC, and tumour specimens from patients with NSCLC.

**Take home messages**

- Forty three of 63 patients with non-small cell lung cancer had increased expression of CD133 positive cells in their tumour tissue
- Thirty of 63 specimens also had raised expression of KDR and 29 of 63 had increased microvessel density
- These results suggest that endothelial progenitor cells (EPCs) are involved in tumour vasculogenesis and tumour growth in these patients
- Treatment strategies targeting the mobilisation of EPCs from the bone marrow and the homing of these cells into malignant tumours might be an alternative approach for the treatment of patients with cancer

**Survival analysis**

None of the tested angiogenetic markers was significantly related to survival in univariate analysis in the tumour cohort.

**DISCUSSION**

Tumour vascularisation seems to be essential for tumour growth and metastatic spread. In addition to angiogenesis—the migration and sprouting of ECs from neighbouring pre-existing capillaries—tumour induced vasculogenesis mediated by bone marrow derived EPCs is thought to be a possible mechanism of vascularisation. Asahara et al showed that EPCs isolated from peripheral blood are incorporated into sites of active angiogenesis, and that EPCs might help augment collateral vessel growth within ischaemic tissue. Furthermore, EPCs are thought to contribute to pathological neovascularisation—for example, in sites of myocardial infarction, and Lyden et al demonstrated in a mouse model that EPCs seem to be key cells for tumour vascularisation and growth. Until recently, no data were available on the presence of EPCs in solid organ cancer, and their possible contribution to tumour induced vasculogenesis in humans. However, a recent study has shown that increased numbers of circulating CD133 positive EPCs can be found in patients with breast cancer. Here, we show for the first time that increased numbers of CD133 positive EPCs are present in patients with NSCLC, and that they seem to contribute to capillary forming units in invading tumour formations (figs 1–4). Increased numbers of CD133 expressing cells were found in 43 of the 63 samples, correlating marginally with KDR expression. Previous phenotypic analyses revealed that most CD133 positive cells derived from CD34 positive cells display endothelial features, including the expression of VEGFR-2 (KDR). Not surprisingly, we found a correlation between CD133 and VEGFR-2 (KDR) expression in our present study. However, CD133 did not correlate with MVD, as defined by CD31 positivity. This result is not surprising because CD31 is expressed on activated endothelial cells and on non-activated ones. CD31 expression can also be detected in many other cell types present in cancer tissues.

All three evaluated cohorts (NSCLC specimens, and tumour free tissue from patients and the control group) revealed similar levels of VEGFR-2 expression. A possible explanation for this observation might be that VEGFR-2 staining is not endothelial cell specific. VEGFR-2 is expressed on endothelial cells, alveolar type II cells, and on endothelial progenitor cells. In addition, normal lung tissue has a high density of blood vessels, and consequently has high VEGFR-2 expression. Thus, VEGFR-2 expression seems to be insufficient to discriminate highly active tumours.

In conclusion, we have shown for the first time that increased numbers of CD133 positive endothelial progenitor cells can be found in non-small cell lung cancer tissue and that these cells seem to contribute to the formation of capillaries.”

**ACKNOWLEDGEMENTS**

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**Authors’ affiliations**

W Hilbe, F Oberwasserlechner, E Gunsilius, E Wöll, C M Kahler, Department of Internal Medicine, University Hospital, Anichstrasse 35, A-6020 Innsbruck, Austria.
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