

Quantification of Angiogenesis in Breast Cancer by Automated Vessel Identification in CD34 Immunohistochemical Sections

LARS TORE GYLAND MIKALSEN¹, HARI PRASAD DHAKAL²,
ØYVIND S. BRULAND^{3,4}, JAHN M. NESLAND² and DAG RUNE OLSEN⁵

¹*Department of Physics, University of Oslo, Oslo, Norway;*

²*Department of Pathology, Faculty Division, The Norwegian Radium Hospital,
Medical Faculty, University of Oslo, Oslo, Norway;*

³*Department of Oncology, The Norwegian Radium Hospital, Oslo University Hospital, Oslo, Norway;*

⁴*Institute for Clinical Medicine, University of Oslo, Oslo, Norway;*

⁵*Faculty of Mathematics and Natural Sciences, University of Bergen, Bergen, Norway*

Abstract. *Background/Aim: Tumor growth is dependent upon angiogenesis. Tumor vascularity, as measured by microvessel density or Chalkley counts, has been shown to predict treatment outcome. However, many issues related to reproducibility and methodology have prevented its clinical application. We present a method of automatic vessel identification applied to CD34 immunohistochemical sections which facilitates increased reproducibility. Materials and Methods: Pixel colour information was used to identify CD34 stain. In order to reduce the effects of noise and background, stained areas smaller than 3.5 µm were ignored. Results: Comparing automatic and manual vessel counts in 50 randomly selected breast cancer cases, the method achieved an intraclass correlation coefficient of $r_a^2=0.96$ and a 95% confidence interval for the percentage difference between the counts from -26.1% to 10.8%. The method was also found to have a sensitivity approaching 100%. Conclusion: The method can reliably be used on colour photographs of staining for CD34 to quantify angiogenesis.*

Formation of new vasculature is required for tumor growth beyond the size of 1-2 mm (1). The clinical importance of angiogenesis has been documented in various types of carcinomas (2, 3), including breast cancer (4-11). Breast carcinomas with high vascularity have been shown to be more aggressive and more likely to form metastases (8).

Correspondence to: Lars Tore Gyland Mikalsen, Department of Physics, University of Oslo, Postboks 1048 Blindern, N-0316, Oslo, Norway. Tel: +47 22857375, Fax: +47 22855671, e-mail: l.t.g.mikalsen@fys.uio.no

Key Words: Angiogenesis, breast cancer, image analysis, CD34, MVD, vessel identification.

Methodological differences, observer variation and conflicting results have, however, kept angiogenesis quantification from being useful in routine clinical management (7, 12, 13). In addition to challenges with quality control and reproducibility, the features of the vascular system related to prognosis are still not well defined.

Microvessel density (MVD) was the first surrogate marker for angiogenesis (8, 14) and the most commonly used (7, 15). The second international consensus report on angiogenesis quantification in solid tumors (12), however, recommends the use of the Chalkley count, a relative area estimate (16). The two methods measure different aspects of the vasculature and do not provide the same clinical information. MVD, Chalkley count, or both have been found to be significantly related to survival of patients for several types of carcinomas (2). However, in breast cancer studies, only Chalkley count predicted survival when both methods were applied to the same patient cohorts (2, 10, 17). Thus, in addition to methodological issues, the parameter selection is clinically relevant.

Many reports deal with vascular density (3), whereas features related to vessel morphology and growth patterns have received far less attention. Although some have been used qualitatively (18, 19), many of the contextual features are tedious, difficult or impossible to quantify manually, but are obtainable through the use of image analysis (16, 20-24). Weyn *et al.* investigated a large panel of parameters and reported that for all investigated tumors, the prognosis was more correlated to contextual parameters than to MVD (22). Image analysis is an important tool for investigation of both conventional, shape and contextual quantifiers. Prior to parameter generation, however, vessels must be identified and each pixel labelled accordingly. Although it is possible to do this manually, an automatic method is a near-requirement for larger studies of clinical material. The accuracy of the method will depend on the material in question, and needs to be explored.

We present a conceptually simple automatic segmentation method, relying on colour hue and blue channel intensity, along with accuracy tests that do not require manual delineation. The results show that the fully automated counts are highly consistent with manual counting.

Materials and Methods

Patients and tumors. We have examined photographs of CD34-stained sections from 420 primary invasive breast carcinomas from the 920 patients enrolled in the Oslo Breast Cancer Micrometastasis Project from 1995 to 1998. The study was approved by the Regional Ethical Committee and written consent was obtained from all patients. The clinical material has previously been reported on (25-27), with stains studied (4, 17).

Immunohistochemistry. Four-micrometer-thick sections from paraffin-embedded blocks with representative tumor tissue were prepared as previously described (4). Briefly, deparaffinised sections were microwaved in Tris/EDTA (pH 9.0), followed by 5 min treatment with 0.03% hydrogen peroxidase. The sections were incubated with monoclonal murine antibody (IgG1) QBEND-10 (Monosan, the Netherlands) against CD34 at room temperature for 30 min, then with peroxidase-labelled polymer conjugated to goat antimouse antibody for 30 min, and finally with 3-3'-diaminobenzidine tetrahydrochloride for 10 min [Dako EnVision™ + System Peroxidase (DAB) (K4007; DakoCytomation, CA, USA)]. Counterstaining was performed using haematoxylin. Appropriate negative and positive controls were included. The immunostaining was performed with a Dako Autostainer.

Microscopy. A careful scan of the tumor with a light microscope (Axiphote microscope; Zeiss, Germany; with a Plan-neofluar 10/0.30 objective lens) at low magnification was used to identify the three most vascular regions in the tumor, disregarding any pre-existing mature vessels. These areas were photographed at $\times 100$ magnification with a cropped square field size of $532.6 \times 717.4 \mu\text{m}$, corresponding to the circular area of a $\times 310$ magnification field in the ocular. For some of the smallest tumors only one or two fields were selected. The photographs were taken at a resolution of 1550×2080 pixels on a Leica DFC320 digital camera, using automatic white balance and exposure. The pixel size was measured to be $0.3436 \mu\text{m}$ using a stage micrometer.

The most vascular of the three fields was then determined manually by visual comparison of the images, where the number of vessels was considered more important than the area in fields with similar vascularity. In total 54 out of 445 cases (10.5%) contained prominent areas of positively stained non-endothelial cells (Figure 1F), e.g. fibroblasts, and were labelled as 'background' in this context. The remaining cases were labelled as 'clean'. The manual selection of the field with highest vascularity was carried out to avoid removal of all cases with fields containing background, even when it was not the most vascular. Case diversity and image quality is exemplified in Figure 1.

Automatic vessel identification. The CD34-positive cells in the images were identified using a segmentation algorithm developed specifically for this material in the Python programming language (28), using the *SciPy* extension (29).

Uneven illumination was corrected for using a 2-degree polynomial fit of the mean background pixel intensity of all images in the set. White balance was adjusted by balancing the RGB colour channels so that the average colour became neutral in light non-overexposed pixels.

The blue colour channel was chosen as an intensity map of the image in order to maximize the contrast of the vessels (orange-red) while minimizing that of the most prominent non-CD34 feature, blue cell nuclei. A combination of two different histogram thresholding methods was used to set an intensity cut-off value: Otsu's method (30), which maximizes between-class variance, and Kittler and Illingworth's method (31) which minimizes the average pixel classification error rate of fitted normal distributions. The smaller of the two obtained threshold values, th_{min} , was compared to a second value, obtained by first selecting the maximum value, th_{max} , and then recalculating the two thresholds using only the portion of the histogram lower than th_{max} . The second value was set to the maximum value of that result, $th_{max, max}$. The final intensity threshold was set to the smallest of the two values, either th_{min} or $th_{max, max}$. All pixels with higher intensity than the selected cut-off were marked as non-endothelial.

In order to differentiate between densely coloured blue cell nuclei (Figure 1D) and CD34 stains, a map of the image's hue values was calculated. Each pixel with a hue located between 150° (mint) and 270° (indigo) were marked as non-endothelial. Pixels of neutral colour were not altered. The remaining pixels, which were both dark and non-blue, were considered CD34-positive.

Using mathematical morphology, the following post-processing steps were carried out: i) Object contours were smoothed using a diameter of 2.5 pixels; ii) CD34 objects were expanded by 3 pixels into adjacent underexposed pixels; iii) any contiguous underexposed region completely enclosed by CD34 staining was reclassified as CD34; iv) Lumens (gaps) thinner than $1.0 \mu\text{m}$ and CD34 objects (vessels) thinner than $w_{min}=3.5 \mu\text{m}$ (half the width of an erythrocyte) at their widest were removed; v) CD34 regions without at least one pixel darker than half the intensity threshold were removed.

From this a complete map of the image was obtained, containing information about the class of each pixel, whether non-endothelial, endothelial or the endothelial region has been removed in post processing due to size requirements (Figure 2).

Vessel count comparison. As a quantitative measure of the program's ability to recognize vessels, the number of vessels in a section, i.e. MVD, was automatically obtained by counting each contiguous foreground object in the image vessel map. The automatic vessel count was compared to a manual count made by an experienced pathologist. The manual counts were carried out on the images according to Weidner *et al.*'s criteria (8, 14), without knowledge of the program results. Microscopy comparisons were not made due to the differences in field size and shape.

Fifty-one out of the 391 clean cases were manually counted and used as training material in the development of the procedure. The cases were recounted 10 months later by the same observer to establish intra-observer variation in manual counts. Fifty test cases were randomly sampled from the remaining material and used to evaluate the program's performance. Finally, counts were made for 25 cases from the background material to investigate the need for a manual selection of cases.

To show the degree of equivalence between the two methods we provide the 95% limits of agreement according to (32, 33), i.e. the

interval of the differences will be located within with 95% probability. Percentage differences were used, and assumed to be normally distributed and independent of the counts. The limits were calculated using ± 1.96 unbiased standard errors. Values are reported with 95% confidence intervals (CI).

An alternate way of comparing the counts is through the use of correlation coefficients (ICC) (34): The ICC(2,1), which treats observer variance as random effects and measures the method agreement (r_a^2); and the ICC(3,1), which treats observer variance as fixed effects and measures the method consistency (r_c^2). While inter-observer variation in the manual method is a random effect, the defined methodological differences of the automatic method are fixed effects. Even though the fixed effects are likely larger, we have chosen to use the random effect model as it will, in most cases, result in a lower value (34), thus preventing an overestimation of the reliability.

The calculations were performed in MatLab® (version (R2010a); The MathWorks, Inc., Natick, MA, USA).

Qualitative evaluation of the segmentation. A qualitative evaluation of how well the program recognized CD34 stains was made for each case in the clean set. The calculated vessel maps were compared to the images and the cases were categorised into one of three qualitatively defined categories. A segmentation result was considered 'good' if it was highly unlikely that the total number of errors made constituted more than 5% of either MVD or total endothelial area; 'medium' if outside this tolerance, but still providing a decent representation, *i.e.* parameters generated from the map should be similar to their true values; and 'poor' if it was not immediately clear that this was the case.

Vessels were allowed to be fragmented, merged or ignored if this reflected the stains. Objects removed due to the size requirements were not considered errors.

The evaluation was performed in a custom program written in *Python* that allowed the user to overlay calculated maps of both objects identified as vessels and objects that would have been identified as vessels were it not for the size requirements. The latter is important to distinguish between vessels omitted due to size alone and vessels omitted due to errors made by the program.

Results

Comparison of automatic and manual counts. The manual and automatic vessel counts are shown in Figure 3A. The method agreement between the two methods was $r_a^2=0.96$, comparatively, the Pearson's correlation was $r_p^2=0.97$. Testing method equivalence, the mean relative difference (measurement bias) was $d=-7.63\pm 2.61\%$ and the standard error $SE=9.41\%$. This gives a lower boundary of -26.08% and an upper boundary of 10.82% ($CI=\pm 4.52\%$) for the 95% limits of agreement (Figure 4A). A frequency histogram of the percentage differences was made to ensure that they were approximately normal in accordance with the applied statistics (Figure 4B). The background cases had a very poor agreement (Figure 3A) with $r_a^2=0.03$. The 95% limits of agreement were -22.80 to 109.79% ($CI=\pm 22.97\%$).

There is no intra-method variation in the automatic method. The two manual counts were highly consistent (Figure 3B), with $d_{m,m}=0.82\pm 1.14\%$ and $SE_{m,m}=4.15\pm 1.97\%$.

The manual intra-method variance contributes $\frac{1}{4}(SE_{m,m})^2$ to the measured inter-method variance, corresponding to 2.52% of the measured standard error. The intra-rater agreement was $r_a^2=0.99$.

The automatic method's vessel criteria differ from Weidner *et al.*'s criteria by introducing a size requirement for the vessels. Figure 5 shows the effect of w_{min} on both correlation coefficients and the 95% limits of agreement. The methods were found to provide strong correlations and a narrow region for the limits of agreement for values between roughly 2.5 and 4.5 μm .

Qualitative evaluation of the automatic vessel identification.

In the qualitative test, 99.5% of the cases were placed in either the 'good' or the 'medium' group. Both of these groups are considered to map the stains in an adequate way (Table I); 0.5% of the cases were deemed to contain too many errors and were placed in the 'poor' category. Table I also shows the largest cause of errors in the 'medium' and 'poor' images.

During the qualitative evaluation, independent vessels that were thicker than w_{min} , clearly defined and readily identifiable by manual inspection, but not found by the program, were counted: 21 such vessels were found, while the total number of automatically identified objects was 29,753, giving a sensitivity estimate of 99.93%.

Discussion

Tumour angiogenesis is important in the metastatic process (8, 35) and of clinical value, both as a prognostic marker (2, 14) and as a predictive marker for targeted therapy (3, 18, 36). An international consensus report on angiogenesis quantification concluded that more reliable parameters were urgently needed (12). Many of the challenges listed, however, are still unresolved. They include methodological difficulties such as reproducibility of vessel identification and field selection. The latter has been proposed as being solved through automatic quantification of the entire section prior to field selection (24, 37); however, it is unclear which parameter it is that best quantifies the degree of vascularity and at what field size it should be measured. The potential role of contextual parameters as prognostic markers complicates this further. Additionally, the choice of endothelial marker best suited for these tasks, and the selection of cut-off values for patient group stratification are not established. (7, 13, 38).

Image analysis has the potential to improve upon several of these areas. The main advantage is the additional morphometric parameters that can be calculated (16, 38, 39). A more complete description of the vascular geometry can thus be investigated with respect to predictive abilities. Furthermore, parameter reproducibility is increased through

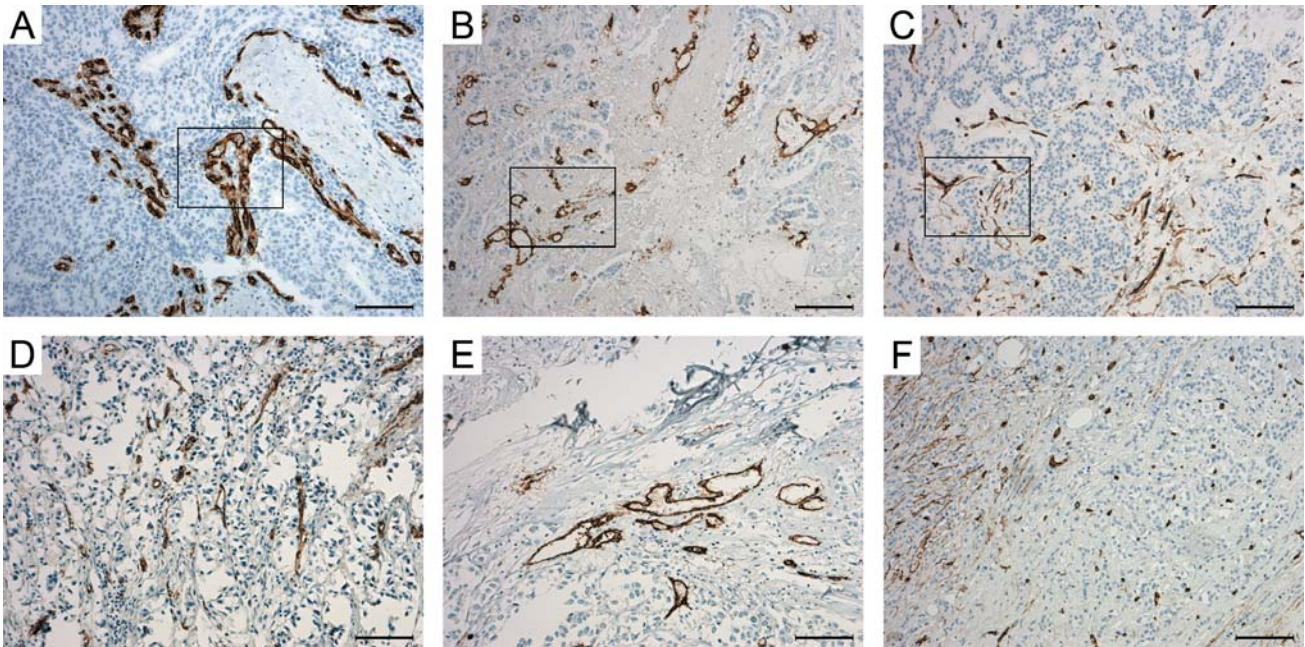


Figure 1. Images from the material exemplifying case diversity. There are large variations in: vessel counts and endothelial growth patterns (A-C), tissue patterns (A, B, and D), cell nuclei intensity (D), and, in rare cases, artifacts (E). In some cases (~10%), fibroblasts were positively stained (left side of F), making them unsuited for automatic segmentation. Scale bars=100 μ m. Insets are further described in Figure 2.

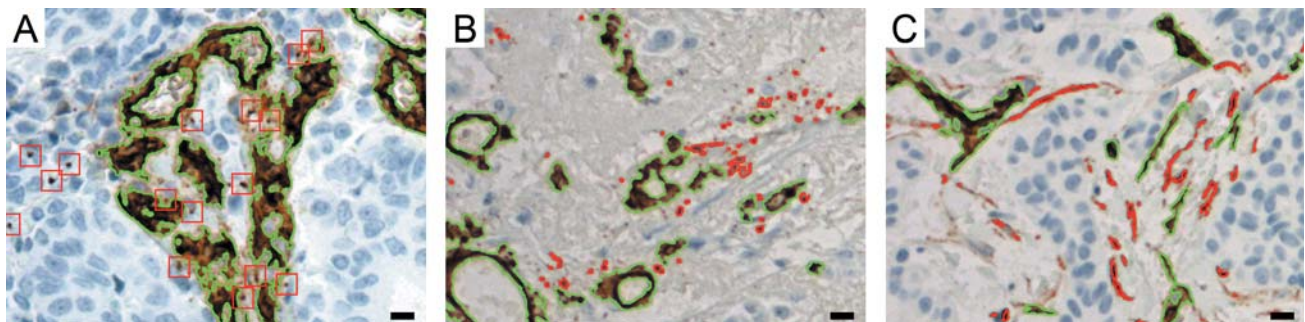


Figure 2. Enlarged sections from the inset of Figure 1 (A-C) respectively. Stains that were thinner than 3.5 μ m were removed in post processing. Green outlines show the final results. A: Red squares show the largest removed objects. The stains are similar to those of the remaining vessels, but are much smaller. B: There are numerous removed objects (red outlines), most of which are very small and have a slightly different colour hue. C: A rare example where long thin structures like vessels are removed (red outlines). The structures have a width close to w_{min} , resulting in a seemingly inconsistent removal of structures with similar morphology. Scale bars=10 μ m.

mathematical standardisation of the procedure, thus ensuring objectivity by eliminating intra- and inter-observer variation (39, 40). Finally, field size effects may be investigated through image cropping.

Parameter calculation requires the identification of all endothelial pixels in the images. Due to the laborious nature of manual delineation, it is highly desirable to use an automatic method. Biological differences, immunohistochemical protocol, or image acquisition setup can have large impacts on

the identification accuracy. For this reason the performance of the method needs to be carefully evaluated.

Although some studies have previously been carried out using automatic vessel identification (22, 23, 37, 41-49), none were found suitable for our purposes. They were either applied to qualitatively different images, insufficiently documented, insufficiently investigated for accuracy or of lower than desired accuracy. We have implemented a conceptually simple algorithm, identifying

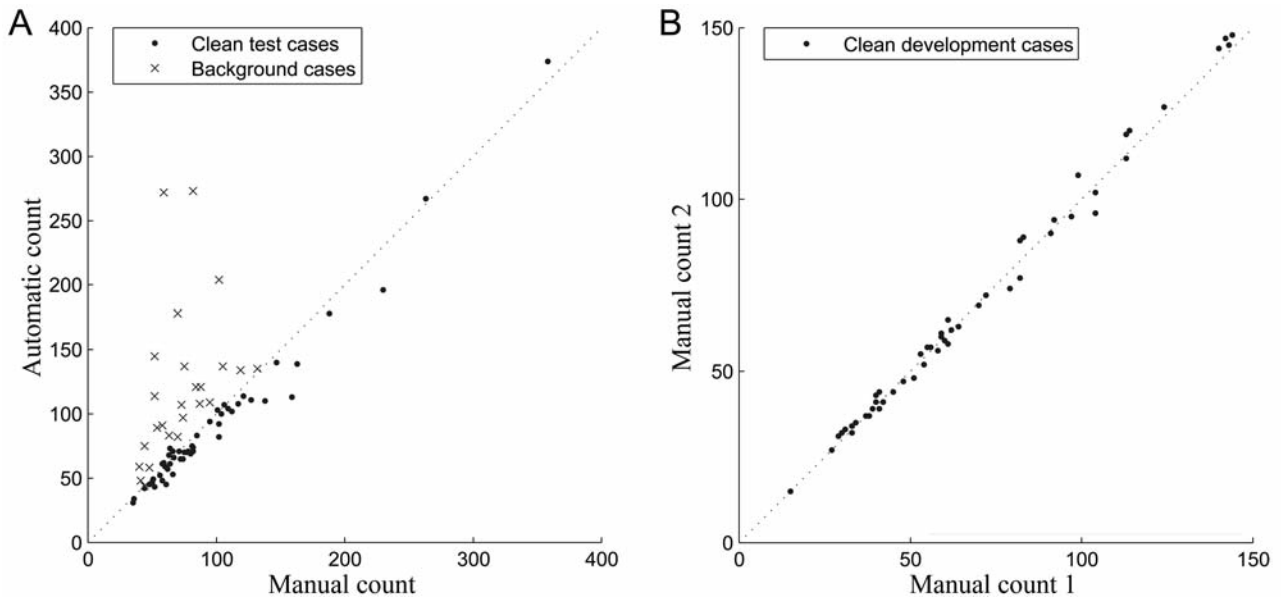


Figure 3. A: Comparison of automatic and manual counts for 50 clean test cases and 25 cases with various degrees of background staining. B: A manual count-recount comparison of the 51 clean training cases.

vessels from colour information and blue channel intensity, with only a few steps of post-processing. This method provided excellent results in the quantitative method validation.

The direct count comparison performed has the advantage of providing quantitative accuracy estimates without requiring manual delineation. There are, however, several potential weaknesses. False-positive and -negative errors cancel each other out; furthermore, no emphasis is placed on how well the vessels are delineated, possibly allowing significant errors in contextual, morphometric and area measurements without affecting counts. For this reason a qualitative evaluation was performed. It showed that vessels were correctly identified and delineated to a high precision in the vast majority of cases. The sensitivity was found to be near 100%; consequently, the observed negative bias in the automatic counts was primarily caused by the removal of small stains, a defined methodological difference.

The size filter was necessary to achieve reliable results (Figure 5) through the removal of noise and of tiny stain fragments, effectively also causing small vessels to be removed (Figure 2). This constitutes an important divergence from Weidner *et al.*'s criteria, where all vessels are counted. Although the clinical significance of these stains is unknown, the strong count comparison shows that this approach is compatible with Weidner *et al.*'s method.

CD34 is known to be a reliable marker for endothelial identification. However, in a low percentage of cases (fewer than 10%), stromal elements express CD34 (Figure 1F),

Table I. Qualitative method evaluation. Each case in the clean material was automatically segmented by the program, and manually assigned to one of three groups depending on a subjective evaluation of the result (see Materials and Methods). 'Medium' and 'poor' cases were additionally categorized by the primary cause of segmentation errors.

	Good	Medium	Poor
No issues	329 (87.5%)	Ø	Ø
Artefact or non-CD34 issue	Ø	10 (2.7%)	0
Background stains	Ø	17 (4.5%)	2 (0.5%)
Vessel colour or intensity	Ø	6 (1.6%)	0
Vessel morphology	Ø	12 (3.2%)	0
Sum	329 (87.5%)	45 (12.0%)	2 (0.5%)

making it difficult to use these cases for automatic analysis (Figure 3 and Figure 5).

Conclusion

We found that CD34 stains can be accurately identified using the presented automatic identification method, provided that a manual selection of cases is made. The quantitative test demonstrated the very strong relationship between manual and automatic counts and the qualitative test verified the integrity of these results. The method can thus be used in the calculation of a wide range of angiogenesis parameters, and provide important clinical information.

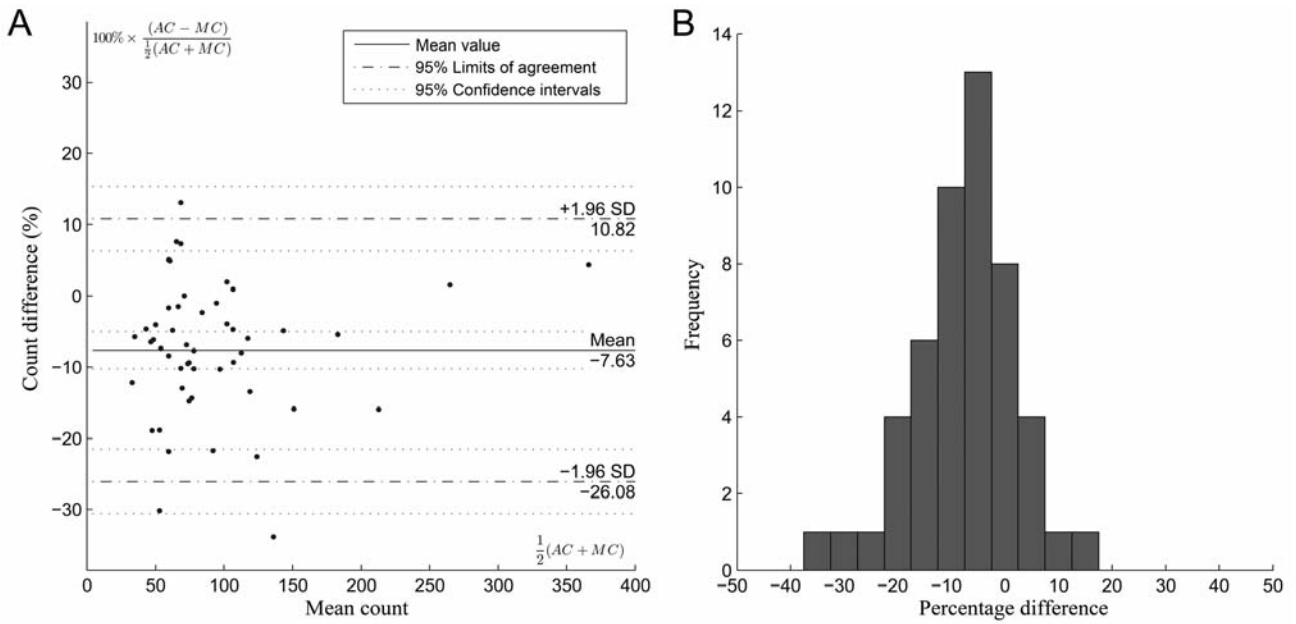


Figure 4. The agreement between automatic (AC) and manual vessel counts (MC) is measured as the region in which the percentage differences will fall within with 95% probability. A: Bland-Altman plot showing the percentage difference between the counts plotted against the average count, with the 95% limits of agreement imposed. B: A frequency histogram showing that the percentage difference distribution is approximately normal, a prerequisite for the limits calculation.

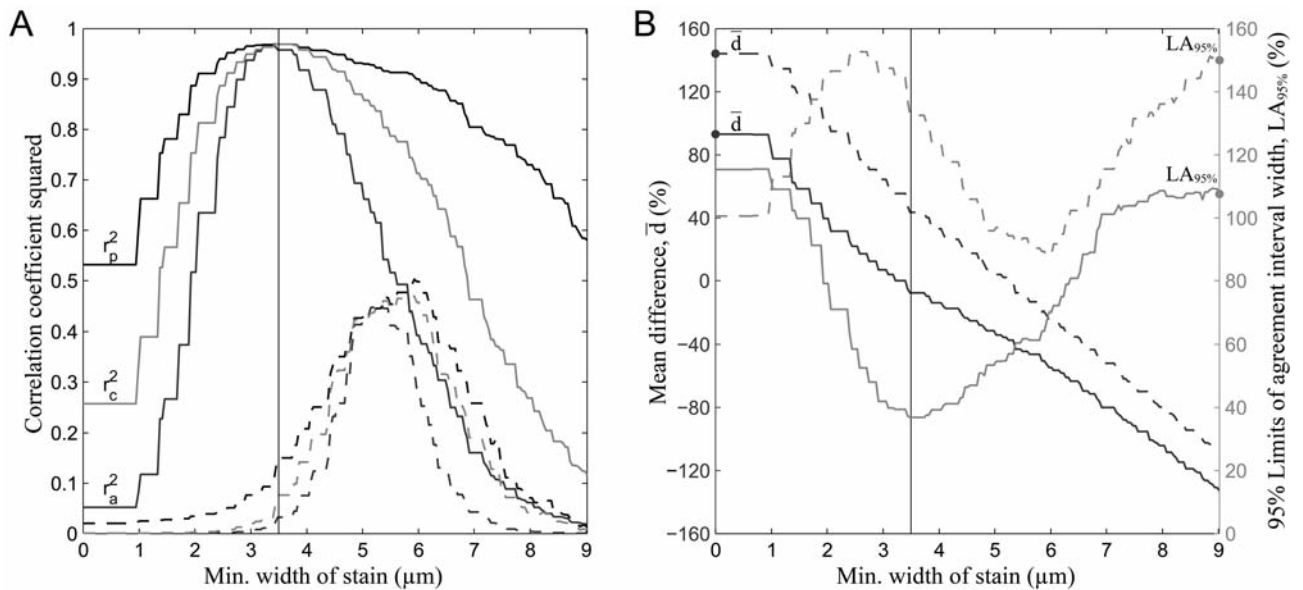


Figure 5. The effects of the minimum vessel width parameter (w_{min}) on the vessel count comparison. Solid lines show the clean cases and dashed lines background cases. A: Pearson's (r_p), intraclass consistency (r_c) and intraclass agreement (r_d) correlation coefficients squared. B: The measurement bias (\bar{d}) and variability ($LA_{95\%}$) as measured by the 95% limits of agreement.

Author Contribution Statement

DRO, JN, HD, LTM and ØB conceived and designed the research. HD and LTM acquired the data; LTM developed the automatic method and analysed the data; LTM wrote the manuscript; HD, JN, ØB and DRO revised the manuscript; all Authors gave final approval of the submitted version.

Acknowledgements

We are grateful to Ellen Hellesylt and Mette Førsund for the high-quality immunohistochemistry, and the Oslo Breast Cancer Micrometastasis Project for permitting the use of data.

References

- Folkman J: What is the evidence that tumors are angiogenesis dependent? *JNCI* 82: 4-6, 1990.
- Offersen BV, Borre M and Overgaard J: Quantification of angiogenesis as a prognostic marker in human carcinomas: a critical evaluation of histopathological methods for estimation of vascular density. *Eur J Cancer* 39: 881-890, 2003.
- Hlatky L, Hahnfeldt P and Folkman J: Clinical application of antiangiogenic therapy: microvessel density, what it does and doesn't tell us. *JNCI* 94: 883-893, 2002.
- Dhakal HP, Naume B, Synnestvedt M, Borgen E, Kaaresen R, Schlichting E, Wiedswang G, Bassarova A, Giercksky KE and Nesland JM: Vascularization in primary breast carcinomas: its prognostic significance and relationship with tumor cell dissemination. *Clin Cancer Res* 14: 2341-2350, 2008.
- Hansen S, Grabau DA, Sorensen FB, Bak M, Vach W and Rose C: Vascular grading of angiogenesis: prognostic significance in breast cancer. *Br J Cancer* 82: 339-347, 2000.
- Nieto Y, Woods J, Nawaz F, Baron A, Jones RB, Shpall EJ and Nawaz S: Prognostic analysis of tumour angiogenesis, determined by microvessel density and expression of vascular endothelial growth factor, in high-risk primary breast cancer patients treated with high-dose chemotherapy. *Br J Cancer* 97: 391-397, 2007.
- Uzzan B, Nicolas P, Cucherat M and Perret GY: Microvessel density as a prognostic factor in women with breast cancer: a systematic review of the literature and meta-analysis. *Cancer Res* 64: 2941-2955, 2004.
- Weidner N, Semple JP, Welch WR and Folkman J: Tumor angiogenesis and metastasis – correlation in invasive breast carcinoma. *N Engl J Med* 324: 1-8, 1991.
- Weidner N: Tumoural vascularity as a prognostic factor in cancer patients: The evidence continues to grow. *J Pathol* 184: 119-122, 1998.
- Hansen S, Sorensen FB, Vach W, Grabau DA, Bak M and Rose C: Microvessel density compared with the Chalkley count in a prognostic study of angiogenesis in breast cancer patients. *Histopathology* 44: 428-436, 2004.
- Offersen BV, Sorensen FB, Yilmaz M, Knoop A and Overgaard J: Chalkley estimates of angiogenesis in early breast cancer – relevance to prognosis. *Acta Oncol* 41: 695-703, 2002.
- Vermeulen PB, Gasparini G, Fox SB, Colpaert C, Marson LP, Gion M, Belien JA, de Waal RM, Van Marck E, Magnani E, Weidner N, Harris AL and Dirix LY: Second international consensus on the methodology and criteria of evaluation of angiogenesis quantification in solid human tumours. *Eur J Cancer* 38: 1564-1579, 2002.
- Fitzgibbons PL, Page DL, Weaver D, Thor AD, Allred DC, Clark GM, Ruby SG, O'Malley F, Simpson JF, Connolly JL, Hayes DF, Edge SB, Lichter A and Schnitt SJ: Prognostic factors in breast cancer. College of American Pathologists Consensus Statement 1999. *Arch Pathol Lab Med* 124: 966-978, 2000.
- Weidner N, Folkman J, Pozza F, Bevilacqua P, Allred EN, Moore DH, Meli S and Gasparini G: Tumor angiogenesis - a new significant and independent prognostic indicator in early-stage breast-carcinoma. *JNCI* 84: 1875-1887, 1992.
- Gasparini G: Clinical significance of determination of surrogate markers of angiogenesis in breast cancer. *Crit Rev Oncol Hematol* 37: 97-114, 2001.
- Fox SB, Leek RD, Weekes MP, Whitehouse RM, Gatter KC and Harris AL: Quantitation and prognostic value of breast cancer angiogenesis: comparison of microvessel density, Chalkley count, and computer image analysis. *J Pathol* 177: 275-283, 1995.
- Dhakal HP, Bassarova A, Naume B, Synnestvedt M, Borgen E, Kaaresen R, Schlichting E, Wiedswang G, Giercksky KE and Nesland JM: Breast carcinoma vascularity: a comparison of manual microvessel count and Chalkley count. *Histol Histopathol* 24: 1049-1059, 2009.
- Sharma S, Sharma MC and Sarkar C: Morphology of angiogenesis in human cancer: a conceptual overview, histoprognostic perspective and significance of neoangiogenesis. *Histopathology* 46: 481-489, 2005.
- Safali M, Karslioglu Y, Arpacı F, Kurt B and Gunhan O: A distinct microvascular pattern accompanied by aggressive clinical course in breast carcinomas: a fact or a coincidence? *Pathol Res Pract* 206: 93-97, 2010.
- Korkolopoulou P, Konstantinidou AE, Kavantzias N, Patsouris E, Pavlopoulos PM, Christodoulou P, Thomas-Tsagli E and Davaris P: Morphometric microvascular characteristics predict prognosis in superficial and invasive bladder cancer. *Virchows Arch* 438: 603-611, 2001.
- Weidner N: Chapter 14. Measuring intratumoral microvessel density. *Methods Enzymol* 444: 305-323, 2008.
- Weyn B, Tjalma WA, Vermeylen P, van Daele A, Van Marck E and Jacob W: Determination of tumour prognosis based on angiogenesis-related vascular patterns measured by fractal and syntactic structure analysis. *Clin Oncol (R Coll Radiol)* 16: 307-316, 2004.
- Goutzani LP, Papadogeorgakis N, Pavlopoulos PM, Petsinis V, Plochoras I, Eleftheriadis E, Pantelidaki A, Patsouris E and Alexandridis C: Vascular fractal dimension and total vascular area in the study of oral cancer. *Head Neck* 31: 298-307, 2009.
- van Niekerk CG, van der Laak JA, Borger ME, Huisman HJ, Witjes JA, Barentsz JO and Hulsbergen-van de Kaa CA: Computerized whole-slide quantification shows increased microvascular density in pT2 prostate cancer as compared to normal prostate tissue. *Prostate* 69: 62-69, 2009.
- Naume B, Borgen E, Kvalheim G, Kaarsen R, Qvist H, Sauer T, Kumar T and Nesland JM: Detection of isolated tumor cells in bone marrow in early-stage breast carcinoma patients: comparison with preoperative clinical parameters and primary tumor characteristics. *Clin Cancer Res* 7: 4122-4129, 2001.

- 26 Naume B, Wiedswang G, Borgen E, Kvalheim G, Karesen R, Qvist H, Janbu J, Harbitz T and Nesland JM: The prognostic value of isolated tumor cells in bone marrow in breast cancer patients: evaluation of morphological categories and the number of clinically significant cells. *Clin Cancer Res* 10: 3091-3097, 2004.
- 27 Wiedswang G, Borgen E, Karesen R, Kvalheim G, Nesland JM, Qvist H, Schlichting E, Sauer T, Janbu J, Harbitz T and Naume B: Detection of isolated tumor cells in bone marrow is an independent prognostic factor in breast cancer. *J Clin Oncol* 21: 3469-3478, 2003.
- 28 Python programming language version (2.6.6); Python Software Foundation. [Online]. URL: <http://www.python.org/> last accessed on October 28, 2011.
- 29 Jones E, Oliphant T and Peterson P: SciPy version (0.8.0): Open Source Scientific Tools for Python. URL: <http://www.scipy.org/> last accessed on October 28, 2011.
- 30 Otsu N: Threshold selection method from gray-level histograms. *IEEE Trans Systems Man Cybernetics* 9: 62-66, 1979.
- 31 Kittler J and Illingworth J: Minimum error thresholding. *Pattern Recog* 19: 41-47, 1986.
- 32 Bland JM and Altman DG: Statistical methods for assessing agreement between two methods of clinical measurement. *Lancet* 1: 307-310, 1986.
- 33 Bland JM and Altman DG: Applying the right statistics: analyses of measurement studies. *Ultrasound Obstet Gynecol* 22: 85-93, 2003.
- 34 Shrout PE and Fleiss JL: Intraclass correlations: uses in assessing rater reliability. *Psychol Bull* 86: 420-428, 1979.
- 35 Fox SB, Generali DG and Harris AL: Breast tumour angiogenesis. *Breast Cancer Research* 9: 216, 2007.
- 36 Kerbel RS: Tumor angiogenesis. *N Engl J Med* 358: 2039-2049, 2008.
- 37 Belien JA, Somi S, de Jong JS, van Diest PJ and Baak JP: Fully automated microvessel counting and hot spot selection by image processing of whole tumour sections in invasive breast cancer. *J Clin Pathol* 52: 184-192, 1999.
- 38 Fox SB and Harris AL: Histological quantitation of tumour angiogenesis. *APMIS* 112: 413-430, 2004.
- 39 Vermeulen PB, Gasparini G, Fox SB, Toi M, Martin L, McCulloch P, Pezzella F, Viale G, Weidner N, Harris AL and Dirix LY: Quantification of angiogenesis in solid human tumours: an international consensus on the methodology and criteria of evaluation. *Eur J Cancer* 32A: 2474-2484, 1996.
- 40 Wild R, Ramakrishnan S, Sedgewick J and Griffioen AW: Quantitative assessment of angiogenesis and tumor vessel architecture by computer-assisted digital image analysis: effects of VEGF-toxin conjugate on tumor microvessel density. *Microvasc Res* 59: 368-376, 2000.
- 41 Ranefall P, Wester K, Busch C, Malmstrom PU and Bengtsson E: Automatic quantification of microvessels using unsupervised image analysis. *Analyt Cell Pathol* 17: 83-92, 1998.
- 42 Goddard JC, Sutton CD, Furness PN, Kockelbergh RC and O'Byrne KJ: A computer image analysis system for microvessel density measurement in solid tumours. *Angiogenesis* 5: 15-20, 2002.
- 43 Kim NT, Elie N, Plancoulaine B, Herlin P and Coster M: An original approach for quantification of blood vessels on the whole tumour section. *Analyt Cell Pathol* 25: 63-75, 2003.
- 44 Mertz KD, Demichelis F, Kim R, Schraml P, Storz M, Diener PA, Moch H and Rubin MA: Automated immunofluorescence analysis defines microvessel area as a prognostic parameter in clear cell renal cell cancer. *Hum Pathol* 38: 1454-1462, 2007.
- 45 Righi M, Giacomini A, Lavazza C, Sia D, Carlo-Stella C and Gianni AM: A computational approach to compare microvessel distributions in tumors following antiangiogenic treatments. *Lab Invest* 89: 1063-1070, 2009.
- 46 Brawer MK, Deering RE, Brown M, Preston SD and Bigler SA: Predictors of pathologic stage in prostatic carcinoma. The role of neovascularity. *Cancer* 73: 678-687, 1994.
- 47 Simpson JF, Ahn C, Battifora H and Esteban JM: Endothelial area as a prognostic indicator for invasive breast carcinoma. *Cancer* 77: 2077-2085, 1996.
- 48 van der Laak JA, Westphal JR, Schalkwijk LJ, Pahlplatz MM, Ruiters DJ, de Waal RM and de Wilde PC: An improved procedure to quantify tumour vascularity using true colour image analysis. Comparison with the manual hot-spot procedure in a human melanoma xenograft model. *J Pathol* 184: 136-143, 1998.
- 49 van Der Laak JA, Pahlplatz MM, Hanselaar AG and de Wilde PC: Hue-saturation-density (HSD) model for stain recognition in digital images from transmitted light microscopy. *Cytometry* 39: 275-284, 2000.

Received September 14, 2011

Revised November 4, 2011

Accepted November 7, 2011