

1 Assessing branching structure for biomass and wood quality estimation 2 using terrestrial laser scanning point clouds

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12 Abstract

13 Terrestrial laser scanning (TLS) accompanied by quantitative tree-modeling algorithms can
14 potentially acquire branching data non-destructively from a forest environment and aid the
15 development and calibration of allometric crown biomass and wood quality equations for species and
16 geographical regions with inadequate models. However, TLS's coverage in capturing individual
17 branches still lacks evaluation. We acquired TLS data from 158 Scots pine (*Pinus sylvestris* L.) trees
18 and investigated the performance of a quantitative branch detection and modeling approach for
19 extracting key branching parameters, namely the number of branches, branch diameter (b_d) and
20 branch insertion angle (b_a) in various crown sections. We used manual point cloud measurements as
21 references. The accuracy of quantitative branch detections decreased significantly above the live
22 crown base height, principally due to the increasing scanner distance as opposed to occlusion effects
23 caused by the foliage. b_d was generally underestimated, when comparing to the manual reference,
24 while b_a was estimated accurately: tree-specific biases were 0.89 cm and 1.98°, respectively. Our
25 results indicate that full branching structure remains challenging to capture by TLS alone.
26 Nevertheless, the retrievable branching parameters are potential inputs into allometric biomass and
27 wood quality equations.

28 **Keywords:** Forestry, LiDAR, Modeling, Point clouds, Scots pine

29 Introduction

30 Size and shape of a tree crown and branches reflect how changes in climate or silviculture affect tree
31 growth (Rubio-Cuadrado et al. 2018) and wood formation (Vanninen et al. 2000). Consequently,
32 branching structures have implications on biomass accumulation (Helmisaari et al. 2002, Ogaya et al.
33 2007) and wood quality (Huuskonen et al. 2014, Kuprevicius et al. 2013, Mäkinen 1999). Tree

34 biomass in particular is one of the most influential attributes required in terrestrial carbon cycle
35 models. Terrestrial carbon in the forest is usually estimated by multiplying forest biomass by a carbon
36 content factor (Penman et al. 2003). Wood quality estimations are considered essential to accurately
37 target harvesting operations and to optimize wood procurement for more sustainable forest resource
38 usage.

39 Allometric equations that utilize forest inventory attributes (e.g., species, diameter-at-breast height
40 (DBH), and tree height (H)) are available for biomass components such as the stem and crown
41 (Jenkins et al. 2003, Liepiņš et al. 2018, Picard et al. 2012, Repola 2009, Zianis et al. 2005). However,
42 data with which to build the models are only available for a limited number of tree species and
43 geographical regions, and they often lack explanatory variables related to crown size and structure,
44 which makes them poorly transferrable across regions (Duncanson et al. 2015, Temesgen et al. 2015).
45 The wood quality of standing timber can additionally be determined using branching parameters
46 (Benjamin et al. 2007, Lyhykäinen et al. 2009, Uusitalo 1997). Branches have a direct, mechanical
47 influence on wood quality in sawn timber, because they distort the stem wood grain orientation and
48 decrease wood stiffness and strength (Samson 2007). Branch knots are often the main defect
49 measured in the visual grading of sawn goods (STMY 2016). Potential variables to more accurately
50 explain crown biomass and wood quality may thus include geometric features of the crown and
51 branches (e.g. crown length, width, and volume, number of branches, variation in branch size, whorl-
52 to-whorl distances, height of the lowest dead branch (H_{db}), and height of the live crown base (H_{lc})).

53 Acquiring detailed data on branching structure for crown biomass and wood quality studies is
54 notoriously difficult. Destructive measurements are required for developing new models or equations
55 for species and geographical areas without existing models, and for calibrating existing models. These
56 measurements are laborious, time consuming, and often not feasible in locations such as permanent
57 sample plots or in urban or conservation areas. Terrestrial laser scanning (TLS) provides a three-
58 dimensional (3D) point-based representation of forest canopies and can be used to characterize both
59 tree- and plot-level structural information through the analysis of spatial point distribution or gap
60 probability, or by modeling individual trees using geometrical primitives (Jupp et al. 2007, Liang et
61 al. 2016, Newnham et al. 2015). The latter approaches have gained increasing interest in the literature,
62 as they have been shown to hold potential for representing the scanned tree structures with the most
63 detail, including a geometrical presentation of individual branches and twigs (Bournez et al. 2017).
64 Most geometrical tree-modeling approaches can be roughly broken down into three main steps
65 (Bucksch et al. 2008, Côté et al. 2011, Gorte et al. 2004, Liang et al. 2012, Pfeifer et al. 2004, Pyörälä
66 et al. 2018b, Raumonon et al. 2013, Xia et al. 2015, Zhong et al. 2017): First, the context of points is
67 characterized, and those belonging to the stems and branches are identified by applying point
68 classification techniques. Second, the points identified as woody components are segmented and
69 organized into a hierarchical structure resembling the tree phenology, based on point connectivity or

70 pattern recognition analyses. Third, the stems, branches, and twigs are modeled by fitting geometrical
71 primitives, most often circles or cylinders, to the points.

72 Studies applying TLS point clouds to estimate the biomass of woody tree crown components have so
73 far concurred that crown-related biomass components can be more accurately estimated using TLS
74 point cloud features than with existing allometric models (Calders et al. 2015, Hackenberg et al. 2015,
75 Hauglin et al. 2013, Kankare et al. 2013, Stovall et al. 2018, Temesgen et al. 2016). Consequently,
76 studies are increasingly using TLS data to infer the stem and branch biomass directly from the
77 geometrical tree models, although how the methodologies perform across various vertical stem
78 sections with varying branch properties remains unclear. The omission of structures along the length
79 of tree stems has a crucial effect on the performance of geometrical tree-modeling methods in the
80 extraction of branching structures and, consequently, their applicability into acquiring input for
81 biomass and wood quality equations (Pyörälä et al. 2018b).

82 In this study, we analyzed the performance of a geometrical tree-modeling method (Pyörälä et al.
83 2018b) in the detection and modeling of individual branch bases across a range of vertical locations
84 and crown conditions. Our objectives were to explicitly quantify the various factors that affected the
85 parameter extraction, and to analyze the implications of our results for the use of geometrical TLS tree
86 models in the development and calibration of biomass and wood quality models.

87 **Materials**

88 The data consisted of 158 Scot pines from six two-hectare forest stands. The stands were located in
89 southern Finland, with four stands in Evo (61.19 °N, 25.11 °E) and two in Orimattila (60.80 °N, E
90 25.73 °E). Based on stand-wise forest inventory data from 2013, the tree species composition varied
91 from nearly pure Scots pine forests to mixed Scots pine and Norway spruce (*Picea abies* H. Karst)
92 forests (Table 1). The sample trees were located in groups of three to six trees and were evenly
93 distributed around each stand. From each sample tree, *DBH* was measured with calipers on two
94 perpendicular planes. A Vertex III hypsometer (Haglöf, Sweden) was used to determine *H*, *H_{lc}*, and
95 *H_{db}*. *H* was the height from the ground to the tree top, *H_{lc}* the height from the ground to the lowest
96 living branch that was separated from the live crown by a maximum of two dead whorls, and *H_{db}* the
97 height from the ground to the lowest dead branch that was approximately over 15 mm in diameter. All
98 Vertex measurements averaged three repetitions. Table 2 shows descriptive statistics of the sample
99 trees at each stand, as measured in the field in August 2014.

100 TLS data were collected using a Faro Focus^{3D} X 330 phase-shift scanner (Faro Technologies Inc., FL,
101 USA) during August and September 2014. Each tree group was scanned from 5–10 locations to
102 ensure that each tree was recorded from all sides, resulting in 197 scans. The mean horizontal
103 scanner-to-tree distance at breast height was 9.8 m (Table 1). With the scanner settings used, point-to-

104 point sampling distance was 6.3 mm at a 10-meter distance. Six spherical targets were used within
105 each of the five tree groups to allow co-registration of the point clouds into a single combined point
106 cloud using Faro Scene 5.2.1 software. Table 1 shows the registration errors given by the software.

107 **Methods**

108 *Data sampling and manual measurements*

109 As manually identifying and measuring each individual branch from all sample trees is impractical,
110 we used a sampling scheme to divide each sample tree into stem section strata using H_{db} , H_{lc} , and H
111 derived from the field measurements. Dead crown (between H_{db} and H_{lc}) and live crown (between H_{lc}
112 and H) were divided into two strata of equal length (upper half and lower half). In other words, we
113 measured five vertical stem sections: the stem below H_{db} , the lower half of the dead crown, the upper
114 half of the dead crown, the lower half of the live crown, and the upper half of the live crown (Figure
115 1a).

116 We manually extracted the point clouds of individual sample trees from the merged point clouds. For
117 every tree, a sample height was randomly selected from each of the five stem sections. All visually
118 identifiable first-order branch bases (i.e. the branch bases diverting from the stem) 0–50 cm below and
119 above (i.e. 1 m in total) the selected height were included in the sample. We extracted points
120 belonging to each visually identified branch base and defined a tight circle that encompassed the
121 extracted points perpendicular to the longitudinal axis of the identified branch. Three geometrical
122 branch attributes were measured as illustrated in Figure 1c: branch diameter (b_d), branch height (b_h),
123 and branch insertion angle (b_a). b_d was the diameter of the manually fitted circle, b_h the difference
124 between the circle center (b_c) z-coordinate and the visually estimated root collar height, and b_a was the
125 angle between a vertical Z-axis $n_z = [0,0,1]$ and the normal $e_0 = [e_1, e_2, e_3]$ of the fitted circle, similarly
126 to the literature (Samson 1993).

127 *The quantitative branch detection and modeling -method*

128 We produced geometrical tree models for each tree that described the tree stems and the first-order
129 branching structure along the full range that was visible in the point clouds. The tree stems were
130 modeled following the approaches of Liang et al. (2012). Points associated with tree stems were
131 identified using principal component analysis (PCA), to estimate the direction vector in which the
132 points exhibit most variance. We assumed that points exhibiting most variance along a near-vertical
133 direction originated from the tree stems. Cylinders were then fitted to the stem points using a
134 weighted least-squares optimization to minimize the distance of the points to the cylinder surface.
135 After stem modeling, points within a horizontal distance of 50 cm from the modeled stem surface
136 were selected and divided into 15-cm segments, with adjacent segments overlapping by 10 cm. Points
137 around the stem model in each segment were analyzed using distributions of point density and the

138 mean distance of the points as per 360 degrees around the stem model (Figure 1b). Two point
139 distributions were used to separate branch points from noisy points, e.g. from branch bumps or other
140 stem deformations that should not have a peak in the distribution of mean distance. The distributions
141 were smoothed by a convolution with a Gaussian window function by means of the Fast Fourier
142 Transform (Cooley et al. 1965). The Continuous Wavelet Transform (CWT) function (Du et al. 2006),
143 an iterative pattern-matching algorithm, was utilized in identifying the peaks in the smoothed
144 distributions. Positions exhibiting peaks from 5 to 45° in width for the point density function and from
145 20 to 75° in width for the mean distance function were defined as branch positions. Points falling
146 within each peak were labeled as belonging to a single branch. Branch points were then projected onto
147 a horizontal plane perpendicular to the longitudinal axis of the branch and modeled as a circle using
148 the Random sample consensus (RANSAC) algorithm (Fischler et al. 1981). Circle diameter was
149 considered to represent b_d , and b_a was solved from the longitudinal axis direction of the points given
150 by PCA. Branches with b_d less than 7 mm and more than 100 mm, and with b_a less than 20° and more
151 than 120° were filtered out, as they were assumed to be false positives resulting from noise or stem
152 deformations. The method pipeline is illustrated in Figure 1 and described in more detail in Pyörälä et
153 al. (2018b).

154 *Evaluation of the quantitative method*

155 For evaluation of the quantitative method, we selected a sample of the quantitatively detected
156 branches using same sample heights as for the manual references. The sample of quantitatively
157 detected branches was compared to the reference branches using all samples from each tree (one
158 sample includes branches along one 1-m section of stem from a given stratum, i.e., five samples from
159 each tree), and the samples in each stem section strata separately (i.e., one sample from each tree).
160 The number of branches detected by the quantitative method (n_q) and the number of branches
161 identified visually (n_m) were calculated separately for each sample. Then, for each sample, we
162 calculated the difference between n_q and n_m . The number of false positives or commission errors (n_c)
163 and false negatives or omission errors (n_o) made by the quantitative method were assessed such that a
164 negative value was considered to indicate branches being omitted, and a positive value to indicate
165 branches being falsely detected (commission error). The accuracy of quantitative branch detection
166 was defined for the stem section strata as well as for entire trees (i.e., combining all samples from a
167 tree) as in Equation 1.

$$Accuracy (\%) = \frac{n_q}{n_q + |n_o| + n_c} * 100 \quad (1)$$

168 Estimates for b_d and b_a derived from the RANSAC-circle-fitting method and PCA, respectively, were
169 compared to the manual measurements. Accuracies of the b_d and b_a estimates were evaluated by
170 comparing the minimum, mean, and maximum branch parameter values between the manual

171 references and the quantitative data in each sample. We reported the root-mean-squared error (RMSE)
172 and the simple regression model R^2 between the data sets separately in each stratum and when
173 considering all samples. b_α accuracy is likely to affect b_d accuracy, as it determines the axis used to
174 project the points onto a horizontal plane for circle fitting. Therefore, we also inspected whether the
175 sample-specific mean estimation error in b_d and b_α had an impact on the other parameter's estimation
176 errors.

177 Previous research has reported that the quality and completeness of the point cloud is highly
178 dependent on scanner distance and occlusion (Abegg et al. 2017, Pyörälä et al. 2018b). Here, we
179 examined to what degree scanner distance and self-occlusion affected branch detection. The height
180 above H_{lc} was considered to represent the magnitude of foliage occlusion, because the quantity of
181 foliage borne by the live crown, and thus the occlusion, was expected to increase cumulatively above
182 H_{lc} (Figure 2). Below H_{lc} , the dead crown was supposed to bear no foliage and to cause no occlusion.
183 We evaluated the quantitative branch detection accuracy within the live crown strata in relation to the
184 3D scanner distance and height above H_{lc} using a multiple linear regression model. We reported the
185 model R^2 and inspected the magnitude of the parameter estimate values for either explanatory
186 variable: the parameter estimate value (or slope) gave the rate at which the detection accuracy was
187 estimated to decline when the scanner distance or height above H_{lc} increased by 1 m. In addition, to
188 analyze the statistical significance of the relationships, we reported the standard error (SE), t-statistic
189 and p -value of the parameters based on the Student's t-test.

190 We expected the quantitative branch detection method to be sensitive to branch size. A simple
191 regression model was used to define the impact of sample-specific mean b_d to quantitative branch
192 detection accuracy in each stem section strata and when considering all samples from a tree. We
193 reported parameter estimates, SE, t-statistic, and p -value.

194 **Results**

195 Compared to our manual measurements, the quantitative branch detection method had an overall
196 accuracy of 68.6%, and was at its highest (81.0–82.6%) in the dead parts of the crown (Figure 3). The
197 number of manually detectable branches increased with height in the dead crown, but decreased
198 significantly above H_{lc} (Table 3, Figure 3). The proportions of omission and commission errors to the
199 number of manually detected branches were 34.4% and 3.0%, respectively. The method made 30
200 commission errors in the stem section stratum below H_{db} . Based on visual inspections, the false
201 positives were due to noisy points, loose bark, stem deformations, and in one case a broken branch
202 that was detected twice by the quantitative method. In addition to causing commission errors, similar
203 factors were also found to cause some of the omission errors of branches that intuitively should be
204 detectable, if not for the aforementioned factors that introduced noise to the point distribution and
205 caused the CWT peak detection to fail (Figure 4).

206 Mean and maximum b_d were larger in the live than in the dead crown, while mean b_a was lower in the
207 live than in the dead crown (Table 3). In all, sample-specific mean b_d were estimated with an RMSE
208 of 0.94 cm (bias -0.89 cm), and mean b_a with a 7.76° RMSE and 1.98° bias (Figure 5). Estimate errors
209 of b_d and b_a were independent of the magnitude of the estimated parameter, and of the error in the
210 other parameter (Figure 6).

211 The correlation coefficient (r) between scanner distance and height above H_{lc} was 0.27. Based on the
212 multiple regression analysis, scanner distance was the main factor contributing to the diminishing
213 branch detection accuracy in the live crown (Table 4). Model RMSE and R^2 were 0.26 and 0.35,
214 respectively. The number of detected branches decreased gradually in both data sets as the distance
215 from the scanner increased, while the occlusion effect was notable only in the reference data (Figure
216 7).

217 Lastly, the simple regression analysis showed that the effect of sample-specific, manually measured
218 mean b_d to branch detection accuracy was not statistically significant, except in the lower half of the
219 dead crown (Table 5).

220 Discussion

221 We analyzed the performance of a geometrical tree-modeling method in quantitative branch detection,
222 and examined the capacity of TLS to provide input data for calibration of crown biomass and wood
223 quality equations across different parts of the stem. The method used in our study represented an
224 example of the geometrical tree-modeling methods that have gained increasing attention in recent
225 forestry-related TLS approaches. However, the methods still largely lack evaluation of their accuracy
226 in branch structure extraction with respect to limiting factors such as the decreasing point cloud
227 quality in tree crowns. Results in our current study entailed a comparison of the automated parameter
228 extraction to manual point cloud measurement data acquired using a sampling procedure from the
229 original TLS point cloud. The sampling approach may have introduced biases and accurately
230 represented the actual number of branches only in the dead parts of the crown closer to the stem base.
231 A diminishing number of manually detectable branches was observable for the sample higher up the
232 live crown (Table 3, Figure 3). Figure 7 suggested that the completeness of the manual reference data
233 above H_{lc} was affected partly by both increasing scanner distance and increasing occlusion effect
234 (Figure 2). The estimated reference branch parameters behaved as expected based on the literature: b_d
235 increased over the dead crown (Table 3) (Maguire et al. 1999, Mäkinen et al. 1998), and b_a decreased
236 with branch height (Table 3) (Mäkinen et al. 1998, Osborne et al. 2015).

237 The quantitative branch detection accuracy compared to manual measurements (64.8%) was lower in
238 our current study than in Pyörälä et al. (2018b) (69.9%), where only the log-section (stem diameter >
239 15 cm) and largest branches in each whorl were considered. The results implied that the lower parts of

240 the tree were better covered by TLS and larger branches were more easily detected. Here, the first
241 assumption was supported by the fact that the quantitative branch detection accuracy and the number
242 of reference branches decreased rapidly above H_{lc} in our data (Figure 3). This effect has been noted in
243 previous research (Boudon et al. 2014, Eysn et al. 2013) and, in this work, we aimed to quantify the
244 factors contributing to the effect. When we analyzed the interaction between the distance from the
245 scanner and the occlusion effect caused by the tree crown above H_{lc} (Figure 2, Figure 7), the scanner
246 distance affected the branch detection accuracy to such a degree that the occlusion effect was not
247 significantly present in the data (Table 4). In addition, the viewing angle between the scanner and the
248 measured point has an additional effect on the magnitude of self-occlusion, which was not considered.
249 The second assumption was not supported by our data, as in this study mean branch size did not have
250 a statistically significant effect on branch detection accuracy in most parts of the tree (Table 5), except
251 for the lower part of the dead crown, where mean b_d was at its lowest (Table 3). In general, it is worth
252 noting that there was relatively little variance among b_d in the data (Table 3).

253 Stand structure and scanning setup affect the uniformity of point cloud density and the magnitude of
254 occlusion. The theoretical probability of a laser beam hitting an object largely depended on object
255 size, the scanner angular resolution, and the evenness of the scanning location distribution, which can
256 be used to minimize the distance from the scanner to each tree and avoid occlusion (Abegg et al.
257 2017). Wilkes et al. (2017) proposed that to capture upper canopy structures, the scanning locations
258 should be sampled using a 10-by-10-m grid. In compliance with the results of our current study, we
259 also argue that to capture branches above H_{lc} , paying attention to scanner-to-tree distances and
260 covering all sides of a tree is crucial. On the other hand, shortening scanner-to-tree distances may
261 restrict the spatial coverage. The multi-scan setup used in our study ensured complete stems were
262 scanned, and had an average distance of 9.8 m between a tree and a scanner. It could be beneficial to
263 implement scanner settings that result in a smaller point-to-point sampling distance than in our study
264 (6.3 mm at a 10-m distance), to avoid compromising the size of the area covered. Further
265 improvements in data acquisition technology could include use of pulse-based laser scanners, or
266 scanners that record multiple returns or the full waveform. Nevertheless, diminishing point density
267 will always occur higher in the tree with any TLS scanning setup or technique. If more commonly
268 available in the future, in- or above-canopy measurements using laser scanners mounted on unmanned
269 aerial vehicles could enable better coverage of upper parts of the tree crowns from closer distance and
270 thus with higher point density (Jaakkola et al. 2017, Wallace et al. 2012).

271 The chosen branch detection and modeling methods may also affect the expected completeness and
272 accuracy of a retrieved branching structure. In manual inspections, the existence of a branch can
273 visually be interpreted even from a sparse point cloud using multiple viewing angles to distinguish a
274 branch-like shape from the surrounding points. The automated method used in our study was based on
275 pattern matching (Figure 1b). Noise, stem deformations, and underestimated stem diameters may

276 result in branches being omitted even if they are visible in the point cloud, as illustrated in Figure 4,
277 due to the threshold values regarding the smoothing convolution with a Gaussian window function,
278 allowed peak width in CWT, and the maximum b_a . These particular omission errors could partly be
279 avoided if the information of tree trunk points as defined in the stem-modeling phase (Liang et al.
280 2012) was used to exclude these points from the branch detection phase. We will implement this last
281 improvement in future applications of the method.

282 Most TLS point cloud based geometric tree-modeling methods found in the literature base the branch
283 detection on point connectivity analyses in the point neighborhood, which is a different approach than
284 in our present study (Bucksch et al. 2008, Gorte et al. 2004, Raunonen et al. 2013). For example, the
285 results in Boudon et al. (2014) and Bournez et al. (2017) implied that connectivity analyses also face
286 challenges higher in the tree crown given the decreasing completeness of the point cloud. One
287 possible solution to substitute for omitted branches could be an approach that combined quantitative
288 point cloud processing with a process-based tree growth model to overcome the data gaps in the point
289 cloud (Côté et al. 2011, Côté et al. 2012). However, such an approach requires existing equations
290 suited for particular species and geographical regions. One of the motivations in developing TLS
291 point cloud -based tree-modeling methods is to enable data collection for the development of new
292 equations as well as for the calibration of existing allometric models, be they process-based or
293 empirical. In such case, data gaps cannot be overcome by modeling.

294 Our results showed that b_d was generally underestimated by the quantitative method compared to the
295 manual method (Figure 5). The effect of the measurer plays a role in manual circle-fitting. A circle
296 can be fitted to the projection of branch points in various ways. In this study, the fitting aimed to
297 exclude noise and the elliptical shape of a branch bottom (Figure 1c), in contrast to our previous study
298 Pyörälä et al. (2018b), where the manual measurements did not account for noise or elliptical shape,
299 but the circle-fitting utilized all points that appeared to belong to a branch. We changed the
300 measurement principle in this study, because we considered the current approach to more realistically
301 represent the size of the branch. Despite the different measurement principles between the two studies
302 as described, the quantitative method underestimated b_d in both Pyörälä et al. (2018b) (whorl-specific
303 maximum branch diameters underestimated on average by -0.34 cm), and the current study (tree-
304 specific mean branch diameters underestimated on average by -0.89 cm) (Figure 5, Figure 6).

305 The b_a estimates from PCA did not differ statistically from the manual measurements when tree-
306 specific mean and maximum values were compared (Figure 5). As PCA is used to distinguish the
307 direction in which the points exhibit most variation, the results indicated that the longitudinal axis of
308 the branch was more completely recorded than the horizontal axis. Figure 4 showed that the insertion
309 angle estimation errors are mostly unbiased in respect to the magnitude of the manually measured b_a .
310 The sharp trend in the maximum error values for b_a resulted from the maximum value threshold that

311 was set at 120° to filter out likely false positives. In addition, the errors in b_a estimates did not have a
312 clear effect on b_d estimate accuracy (Figure 6c), although a slight trend was observable.

313 Previous studies on the accuracy of individual branch modeling are sparse. Dassot et al. (2012)
314 excluded branches below 7 cm in diameter, but reported $\pm 30\%$ differences between tree-specific
315 branch volume estimates and destructive measurements. Côté et al. (2013) combined quantitative
316 point cloud processing to process-based tree growth modeling and compared the results to destructive
317 measurements from six trees. The group reported relative RMSEs of 20% and 25% for b_d and b_a ,
318 respectively. Hackenberg et al. (2015) reported that compared to destructive biomass measurements,
319 branch modeling by means of cylinder-fitting was more accurate for branches with diameters above 7
320 cm. Lau et al. (2018) also reported accurate results in b_d estimation, but excluded branches below 10
321 cm in diameter. Based on our results and the literature, individual branches remain challenging to
322 retrieve accurately from TLS point clouds, especially for coniferous species where most branches are
323 small in diameter.

324 Our results and previous research implied that unbiased crown biomass estimates in a forest
325 environment are probably not achievable directly from branch-to-branch measurements using TLS
326 point clouds. Instead, select branching features that are extractable from TLS point clouds could be
327 used as additional explanatory variables in allometric biomass models to account for local variation in
328 crown structure (Kankare et al. 2013, Stovall et al. 2018, Temesgen et al. 2016). On the other hand,
329 based on the current results and our previous studies, it is apparently possible to derive branching
330 parameters from TLS point clouds that are applicable to wood quality estimations (Kankare et al.
331 2014, Pyörälä et al. 2018a, Pyörälä et al. 2018b). For example, H_{db} is one of the most commonly used
332 variable in estimating wood quality in standing timber, because it can be used to estimate the yield of
333 the branchless bottom logs, the most valuable log product (Lyhykäinen et al. 2009, Uusitalo 1997).
334 Our current study had false positive branch detections below H_{db} , which could lead to underestimating
335 the bottom log yield. Furthermore, a comparison to X-ray scanning data of logs in Pyörälä et al.
336 (2018a) revealed a dependency between the maximum interior knot diameter within a log measured
337 from X-ray scanning images and the maximum b_d measured manually from the TLS point cloud. In
338 Pyörälä et al. (2018b) and in our current study, the quantitative method underestimated b_d , which may
339 mean that the maximum knot size within a tree was also underestimated.

340 **Conclusion**

341 Tree crown and branching parameters are closely related to tree growth and wood formation. Due to
342 the difficulty in measuring these metrics directly by conventional, destructive means, a wide field of
343 applications relies on established relationships between crown biomass, wood quality, and more
344 readily measurable tree characteristics such as DBH , H , and H_{db} . However, laborious local reference
345 measurements are required to build such equations for new species and regions. Furthermore, in the

346 case of biomass modeling, the accuracy of allometric models is lower at predicting the biomass of the
347 crown components than of the stem. Using TLS point cloud -based geometrical tree-modeling
348 approaches to measure variables describing the branching structure could reduce the need for
349 destructive measurements and improve the accuracy of the equations in estimating the crown biomass
350 and the expected wood quality, even if full branching structures are not captured. The initial equation
351 development would also require species-specific destructive measurements of tree biomass, but the
352 extractable branching parameters describe tree-specific differences in growth and wood formation
353 across various regions in more detail than the currently applied equations.

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539 **Table 1.** Stand information from a stand-wise forest inventory in 2013: Site location, site type (VT =
540 Sub-xeric heath forest i.e. *Vaccinium* type, MT = Mesic heath forest, i.e. *Myrtillus* type, OMT = Herb-
541 rich heath forest, i.e. *Oxalis-Myrtillus* type) and volume of pine (V_{Pine}) and other species (V_{other}) per
542 hectare. Scanning information: Mean 2D scanner-to-tree distance, the standard deviation (\pm SD), and
543 mean registration error (\pm SD).

Stand	Location	Site type	V_{Pine} (m ³ *ha ⁻¹)	V_{other} (m ³ *ha ⁻¹)	Distance to the scanner (m)		Registration error (mm)	
					Mean	\pm SD	Mean	\pm SD
1	Evo	VT	220	10	8.9	4.18	1.2	3.5
2	Evo	VT	250	20	10.85	4.81	1	2.2
3	Evo	OMT	200	20	9.47	4.25	0.7	1.7
4	Evo	MT	140	260	10.15	5.31	1.5	5.6
5	Orimattila	MT	80	170	11.93	4.64	1.4	2.4
6	Orimattila	VT	170	80	12.18	5.11	1.7	3.3
<i>Mean</i>			<i>176.7</i>	<i>93.3</i>	<i>9.79</i>	<i>4.83</i>	<i>1.3</i>	<i>5.6</i>

544

545 **Table 2.** Sample tree field measurement statistics. Mean and standard deviation (\pm SD) values of
546 diameter at breast height (DBH), tree height (H), the height of the lowest dead branch (H_{db}), and base
547 height of the live crown (H_{lc}) of the sample trees.

Stand	Number of trees	DBH (cm)		H (m)		H_{db} (m)		H_{lc} (m)		
		Mean	\pm SD	Mean	\pm SD	Mean	\pm SD	Mean	\pm SD	
1	30	28.9	3.4	22.6	1.2	4.8	2	14.4	1.7	
2	20	32.8	2.5	27	1.5	8.7	1.5	17.1	1.8	
3	30	28.6	4.2	22.9	1.8	4.4	2.3	13.9	1.9	
4	24	35.8	5.0	29	1.4	8.5	1.6	20.9	1.8	
5	28	34.7	4.2	27.7	1.8	10	2.9	18.8	1.6	
6	26	32.1	6.0	26	2.4	7.6	2.2	15.4	1.7	
<i>All trees</i>		<i>158</i>	<i>31.8</i>	<i>5.2</i>	<i>25.5</i>	<i>3</i>	<i>7</i>	<i>3.1</i>	<i>16.6</i>	<i>3.0</i>

548

549 **Table 3.** Number of manually measured branches (N), minimum (Min), mean, maximum (Max), and
550 standard deviation (\pm SD) of their height (b_h), diameter (b_d), and insertion angle (b_a) in different stem
551 sections. H_{db} = Height of the lowest dead branch.

	N	b_h (m)				b_d (cm)				b_a (°)			
		Min	Mean	Max	\pm SD	Min	Mean	Max	\pm SD	Min	Mean	Max	\pm SD
Below H_{db}	22	0.99	5.30	9.31	2.83	1.02	1.58	2.22	0.37	53.46	69.51	84.08	9.25
Dead crown, lower half	702	1.94	9.42	16.36	3.17	0.77	2.06	5.72	0.65	36.21	67.23	99.63	11.36
Dead crown, upper half	1075	7.66	13.33	20.51	2.71	0.78	2.25	5.56	0.66	30.87	64.37	98.39	11.66
Live crown, lower half	720	12.52	17.35	24.54	2.64	1.08	2.35	5.99	0.67	28.82	61.95	94.43	11.38
Live crown, upper half	42	15.85	19.92	25.61	2.72	1.39	2.44	4.71	0.78	20.80	63.37	78.95	13.60
Total	2561	0.99	13.43	25.61	4.24	0.77	2.21	5.99	0.67	20.80	64.75	99.63	11.67

552

553 **Table 4.** Results of the multiple regression analysis on the effect of scanner distance and occlusion
554 (Height above the live crown base (H_{lc})) on the quantitative branch detection accuracy. SE =
555 parameter estimate standard error, t = Student's t-test statistic of the parameter estimate, p =
556 probability value of the t-test. Statistically significant parameter estimates ($p < 0.05$) are marked with
557 an asterisk (*).

Quantitative branch detection accuracy above H_{lc}	Estimate	SE	t	p
Intercept	1.564	0.165	9.494	<0.05*
Distance from the scanner	-0.062	0.008	-7.970	<0.05*
Height above H_{lc}	-0.003	0.016	-0.170	0.87

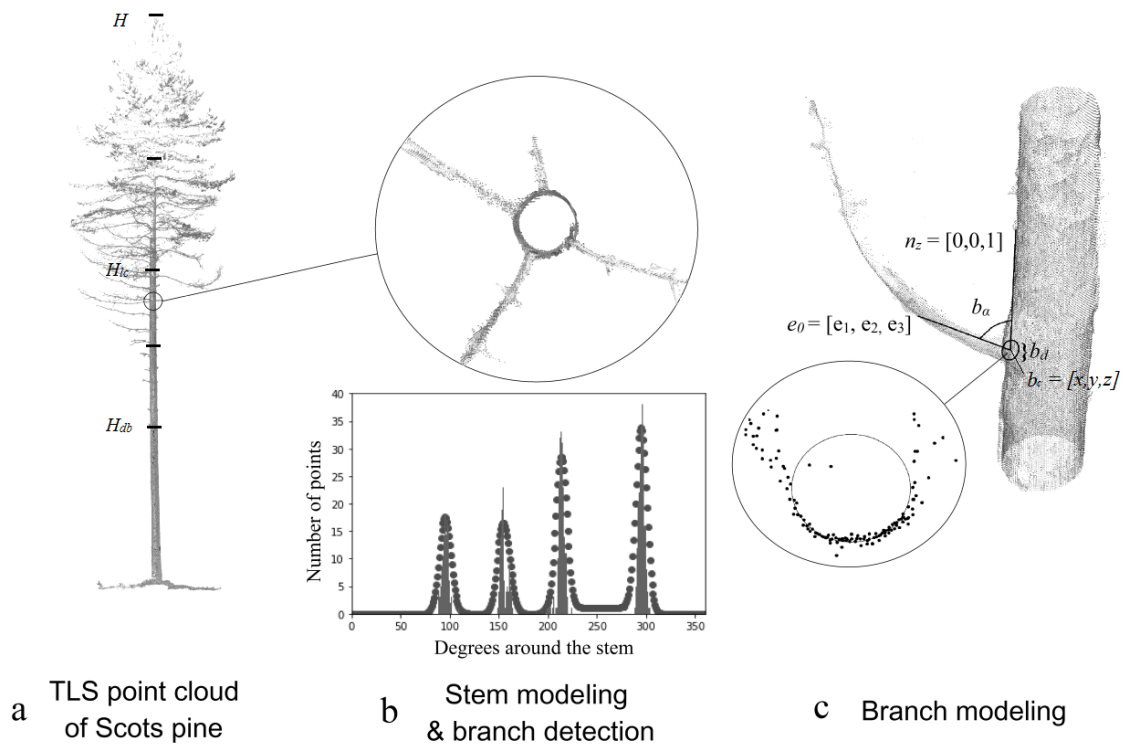
558

559 **Table 5.** Results of the simple regression analysis on the effect of the sample-specific mean branch
560 diameter on the quantitative branch detection accuracy. SE = parameter estimate standard error, t =
561 Student's t-test statistic of the parameter estimate, p = probability value of the t-test. Statistically
562 significant parameter estimates ($p < 0.05$) are marked with an asterisk (*).

Quantitative branch detection accuracy	Estimate	SE	t	p
<i>Below Hdb</i>				
Intercept	-0.102	0.616	-0.166	0.875
Mean bd	0.390	0.402	0.970	0.377
<i>Dead crown, lower half</i>				
Intercept	0.401	0.097	4.131	<0.05*
Mean bd	0.110	0.044	2.493	<0.05*
<i>Dead crown, upper half</i>				
Intercept	0.598	0.119	5.037	<0.05*
Mean bd	0.019	0.050	0.378	0.706
<i>Live crown, lower half</i>				
Intercept	0.229	0.156	1.469	0.145
Mean bd	0.028	0.064	0.441	0.660
<i>Live crown, upper half</i>				
Intercept	-0.205	0.245	-0.836	0.431
Mean bd	0.113	0.098	1.154	0.286
<i>Full tree</i>				
Intercept	0.476	0.109	4.374	<0.05*
Mean bd	0.064	0.047	1.353	0.178

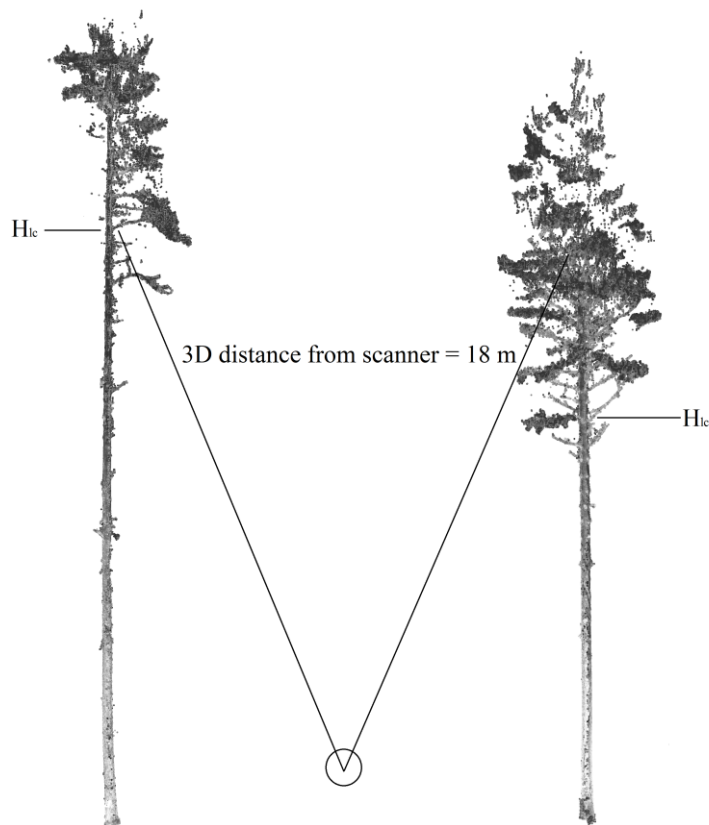
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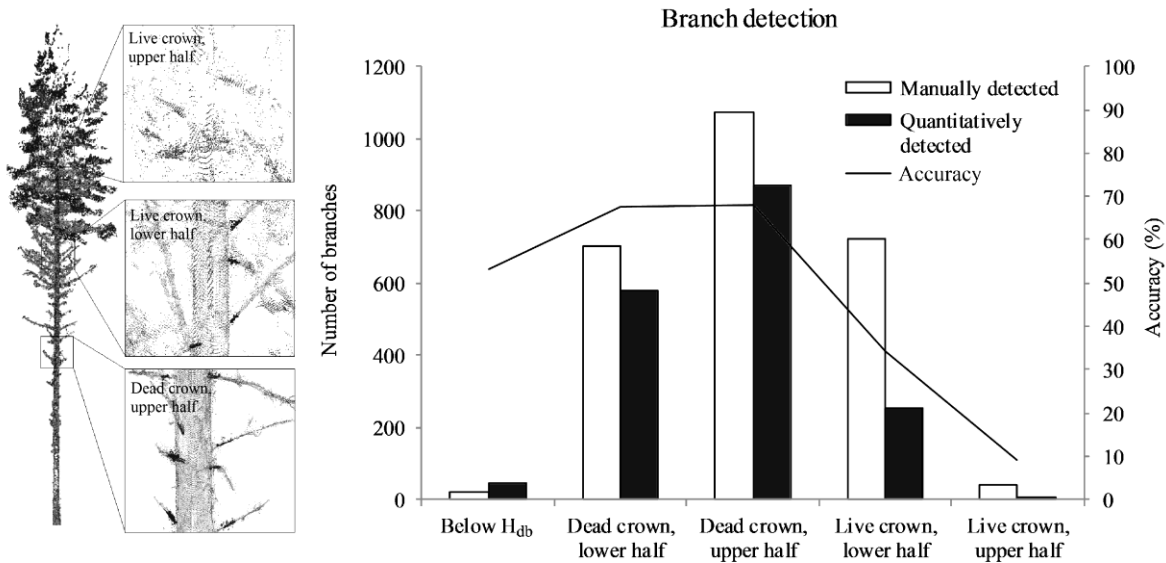
565

566 **Figure 1.** The sampling scheme and principle of the quantitative method applied for detecting and
 567 modeling branches. a) H_{db} , H_{lc} and H show the heights of the lowest dead branch, live crown base,
 568 and tree top, respectively. Black lines show the limits between the sampled strata. b) The tree point
 569 cloud was segmented vertically into slices of 15 cm thickness and 50 cm radius. For each segment, the
 570 stem was modeled as a cylinder and the point distribution around the stem model was analyzed.
 571 Locations that exhibit peaks in the point distribution were considered potential branch locations and
 572 points from these locations were extracted for the modeling. c) The branch insertion angle (b_a) was
 573 solved from the eigenvector e_0 of the branches and Z-axis n_z using principal component analysis, and
 574 the branch diameter (b_d) and branch location (b_c) were modeled by means of circle-fitting.

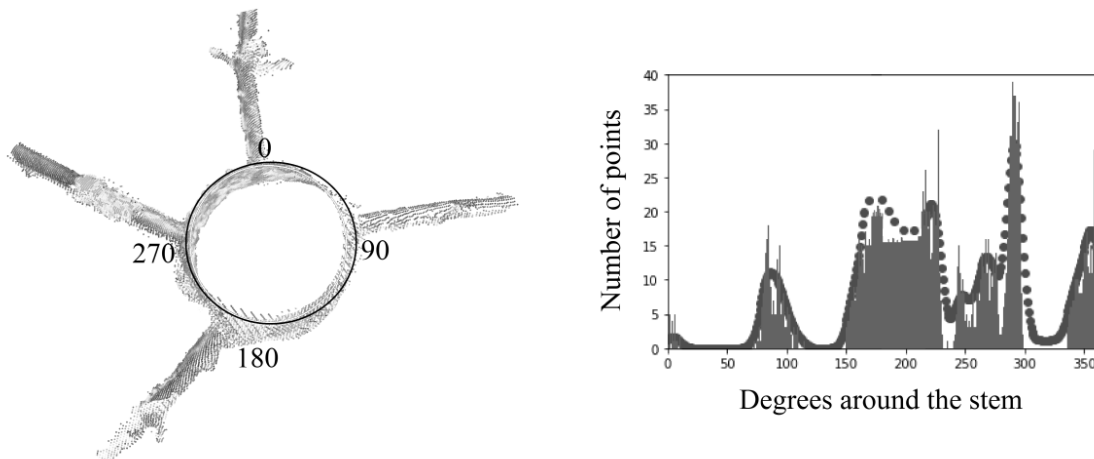


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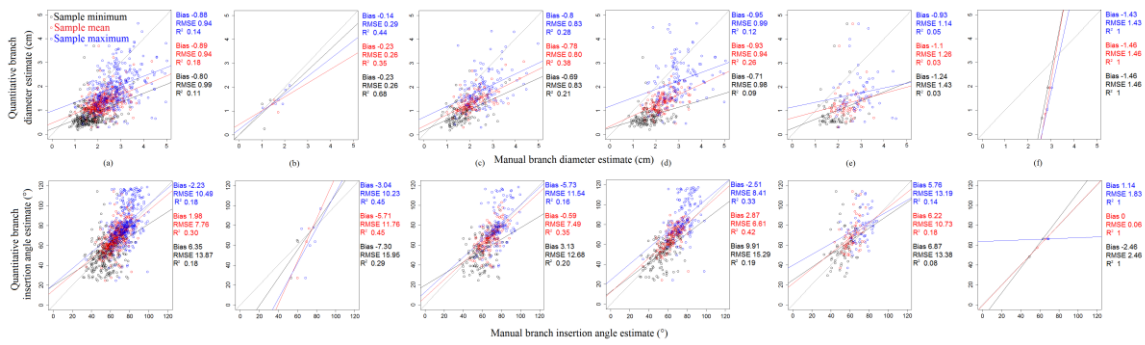
576 **Figure 2.** The simplified concept of the occlusion effect as indicated by the height above the base
 577 height of the live crown (H_{lc}). The figure illustrates two sample trees with different live crown base
 578 heights. The scanner was situated at an equal distance from either tree. On the left-hand side, the line
 579 shows the 3D distance from the scanner (18 m) to a branch 0 m above H_{lc} . On the right-hand side, the
 580 line shows an equal 3D distance from the scanner to a branch approximately 6 m above H_{lc} . Due to
 581 the unequal quantity of accumulated foliage between the scanner and each branch, the right-hand-side
 582 branch is supposed less likely visible than the left-hand-side branch. Therefore, the height above H_{lc}
 583 was considered to represent the magnitude of occlusion in this study.



584
 585 **Figure 3.** Branch detection performance in different stem sections. Left: illustrations of branch
 586 detection performance in the dead crown and lower and upper half of the live crown; branches
 587 detected quantitatively are highlighted in distinctive dark gray. Right: White and gray bars represent
 588 the number of branches detected manually and quantitatively, respectively. The line shows the
 589 accuracy (%) of the quantitative method compared to the manual method. H_{db} is the height of the
 590 lowest dead branch measured in the field.



591
 592 **Figure 4.** The effect of stem model to branch detection: forcing a cylindrical shape to the stem points
 593 excludes stem deformations, branch bumps, or other anomalies from the stem model. In our branch
 594 detection method, such occurrences distort the analyzed point densities around the stem model: in the
 595 illustrated whorl, the branch at approximately 220 degrees was omitted by the Continuous Wavelet
 596 Transform peak detection due to stem deformation between 150 and 270 degrees that distorts the
 597 baseline.



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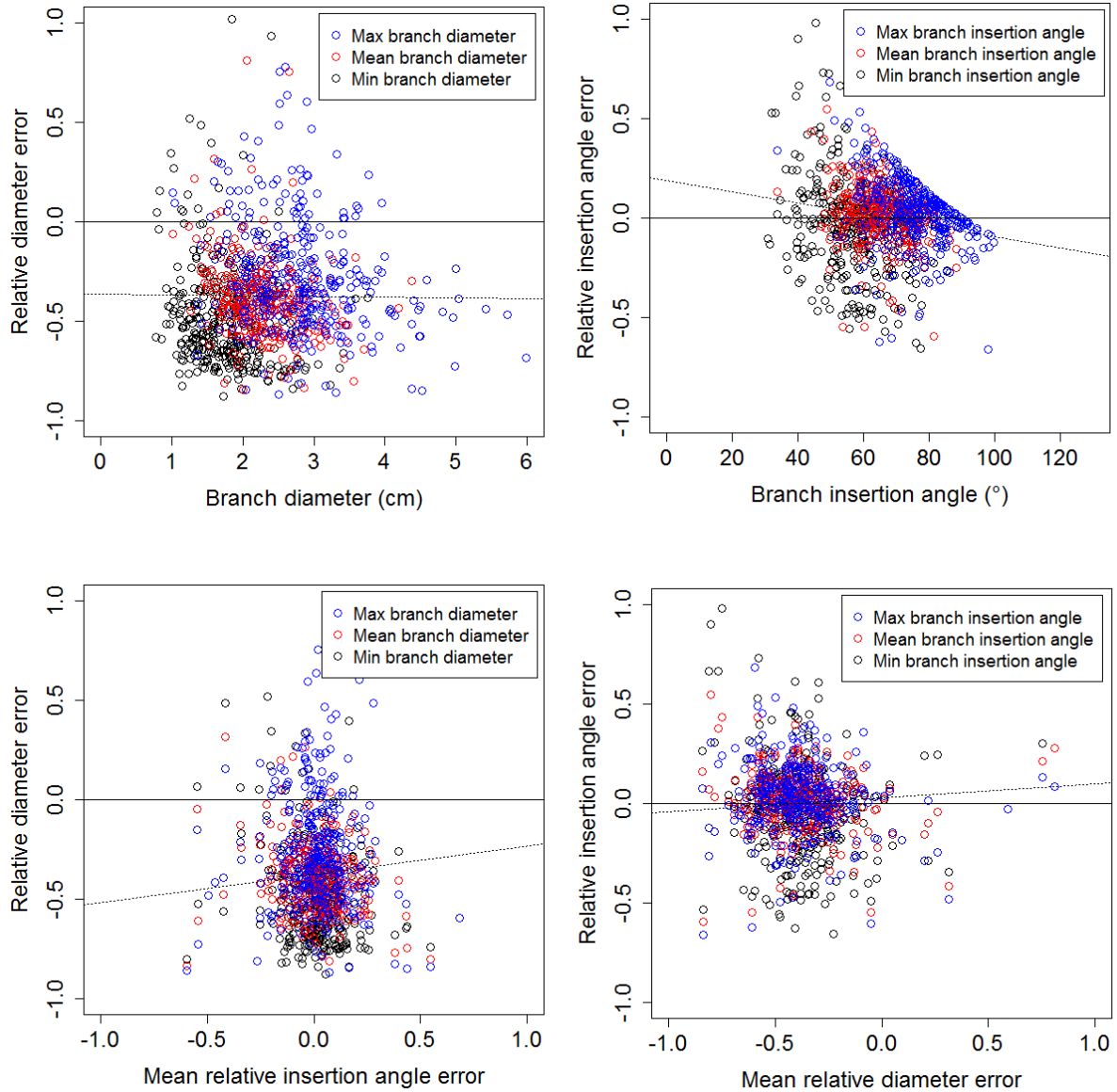
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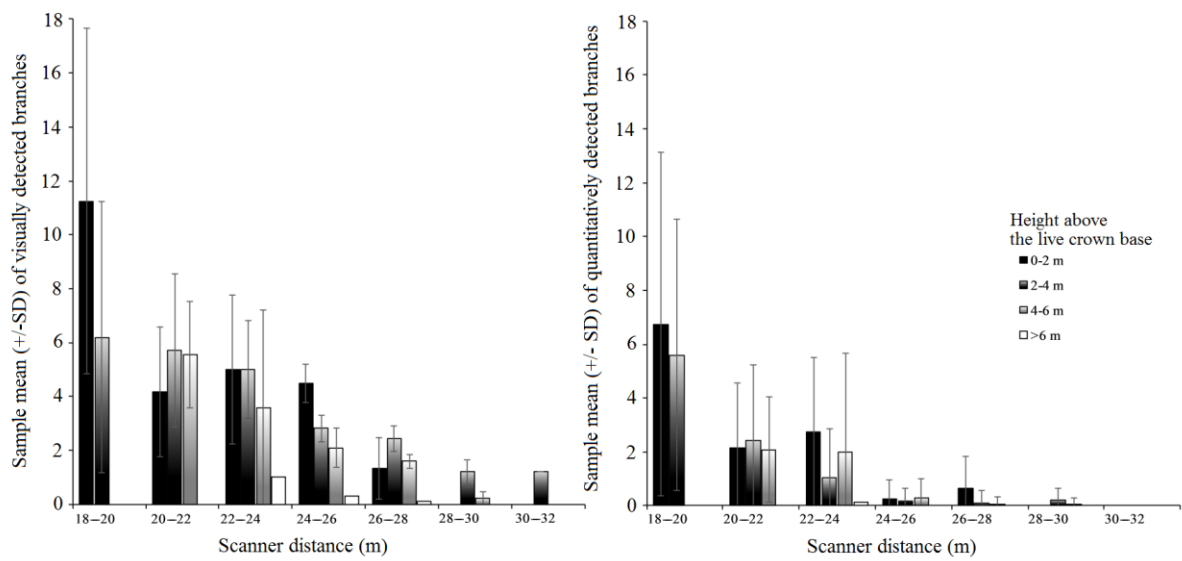
Figure 5. Comparisons between manually measured branch parameter references and quantitatively estimated branch parameters using samples from (a) all five strata in each tree, (b) below the lowest dead branch, (c) the lower half of the dead crown, (d) the upper half of the dead crown, (e) the lower half of the live crown, and (f) the upper half of the live crown. The upper row gives the results of the branch diameter comparisons between the data sets and the bottom row those of the branch insertion angle. Blue indicates the sample-specific maximum value of the branch parameter, red the mean, and black the minimum. The bias and root-mean-squared error (RMSE) give the error of the quantitative minimum, mean, and maximum estimates compared to the manual minimum, mean, and maximum observations. R^2 indicates the regression model fit (solid line) between the two.



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609 **Figure 6.** Sample-specific relative errors of b_d and b_a in respect to the reference measurement value,
 610 and the mean relative estimation error of the other branch parameter. Each observation refers to the
 611 minimum (black), mean (red), or maximum (blue) values of the branch parameter in a sample (i.e.,
 612 branches along one 1-m stem section). A negative value on the y-axis refers to the quantitative model
 613 underestimating the parameter, and vice versa. The dotted line shows the best linear fit between the
 614 variables.

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 617 **Figure 7.** The effect of 3D distance from the scanner and the self-occlusion as indicated by the height
 618 above the base height of the live crown (H_{lc}), (color-coded bars) on the sample-specific mean number
 619 (\pm standard deviation) of manually detected branches (left), and quantitatively detected branches
 620 (right).