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A machine learning method to assess growth patterns in plants of the family Lemnaceae.

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Abstract: Numerous new technologies have been implemented in image analysis methods that help researchers withdraw scientific conclusions from biological phenomena. Plants of the family Lemnaceae (duckweeds) are the smallest flowering plants in the world, and biometric measurements of single plants and their growth rate are highly challenging. Although the use of software for digital image analysis has changed the way scientists extract phenomenological data (also for studies on duckweeds), the procedure is often not wholly automated and sometimes relies on the intervention of a human operator. Such a constraint can limit the objectivity of the measurements and generally slows down the time required to produce scientific data. Here is the need to implement image analysis software with artificial intelligence that can substitute the human operator.

In this paper, we present a new method to study the growth rates of the plants of the Lemnaceae family based on the application of machine learning procedures to digital image analysis. The method is compared to existing analogical and computer-operated procedures. Results showed that our method drastically reduces the time consumption of the human operator while retaining a high correlation in the growth rates measured with other procedures

As expected, machine learning methods applied to digital image analysis can overtake the constraints of measuring growth rates of very small plants and might help duckweeds gain worldwide attention thanks to their great nutritional qualities and biological plasticity.

Keywords: Duckweed; Machine learning; Image analysis; Machine training; Aquatic plants; Lemnaceae; Lemna

1. Introduction

Image analysis has changed the way scientists experiment in numerous fields [1]. The image analysis approach allows scientists to frame time-specific data that can be analysed later. This methodology has been adopted in multiple plant science research fields [2]. Image analysis software is the go-to technology to correctly satisfy the needs of modern research data. In several areas of study (including genetics), the requirement of an image analysis software that could quantify tiny differences among plants phenotypes has been mandatory and has led us to enter the so-called "big data era" in plant science [3–4]. Thanks to the breach made by the genetic field, this software analysis method soon became mandatory in numerous other areas, such as botany, agronomy, and forestry [5–10].

Within the "big data era", scientists are now phasing the new challenge of extrapolating scientific-sounding data from the monstrous amount produced by image analysis [11–12]. Here is the birth of the "artificial intelligence era" [13], in which computer intelligence substitutes humans to extrapolate scientific quality data among large quantities.

The advent of artificial intelligence in plant science is already paying off [14]. In numerous fields of plant science, this technology is speeding up the process and excluding numerous errors made by human operator [11]. Although artificial intelligence interfaces are still too complicated for most plant biologists, some software leads to better use of this technology in numerous research fields [14–15]. Among others, ilastik® is a supervised machine learning software (learning from training data) that brings machine-learning-based image analysis to end-users without extensive computational expertise [16].

Ilastik® provides end-users with a supervised machine learning experience without requiring extensive training data. This is achieved thanks to the accurate machine training feature of the software that can fine-tune training via a "paint" like interface [17–18]. Ilastik® contains pre-defined workflows that can be used for image segmentation, object classification, counting and tracking [16]. Moreover, a specific setup of the machine training process can be reutilised numerous times, and applying a particular feature of the program, "batch analysis" can be performed theoretically with an infinite number of images [17].

This paper proposes the use of ilastik® in a low-cost setup aimed at getting a new standardised method to perform image analysis of the aquatic plant family Lemnaceae. These hydrophytes have been often mentioned referring to their small size and fast growth [19–21]. However, their current appreciation is moving toward these plants' exceptional nutritional qualities [21]. Additionally, Lemnaceae are gaining worldwide attention in numerous other fields, such as phytoremediation, plant science, biomonitoring, and closed bioregenerative systems [22–24]. Due to the simplicity of these biological systems, numerous scientists are evaluating the possibility of using these plants as a model [25]. Research in all these fields is constrained by the extremely small size of individuals that prevents applying the methods that are commonly used for biometric measurements and plant growth rates in all other flowering species. We suggest a new image analysis method via machine learning to boost knowledge and standardise the scientific analysis of the Lemnaceae plant's growth. This approach can increase confidence in experimental results and speed up image analysis techniques by offloading the image analysis process to a machine [26]. Due to scientists' strong interest in this family of plants, we shared the view that it is mandatory to standardise analysis methods [27] and decided to contribute to achieving such a goal.

In particular, we focused on methods able to identify fine changes in growth phenological processes more effectively than those based on the number of fronds used in the past [28]. The new method had to be applied in any growth-related tests, such as bioassay and laboratory tests, for Lemnaceae and other floating aquatic plants.

More specifically, our work aimed to validate the utilisation of Ilastik® software in monitoring the growth rate of Lemnaceae. Our approach was to highlight the possible effects of two light treatments on plant growth by applying the previously used analyses and the newly proposed method.

2. Results

To evaluate the use of the machine learning system, we cultivated *Lemna minor* (9440) under different light quality treatments. Moreover, we have studied the growth rates with different methods. More specifically, the standard gold method has been defined by the ISO 20079 protocol. This method requires counting the number of fronds over a period of time at constant time intervals. Furthermore, two digital methods were also investigated, one previously described by Haffner et al. [1] and the newly described ilastik® method. The three methods used the Naumann et al. [2] equation to calculate

growth rates. The three methods' results were compared to appreciate any existing differences. Additionally, the Ilastik® method's results were compared with those produced in Fiji.

Plants cultivated under the two different combinations of light conditions (White and White + Red) grew healthy, with no visual sign of overall differences. Images of plants at the beginning and end of the experiment were used to calculate the relative growth rates (RGR) of *Lemna minor* by applying the three different methods (the ISO 20079 frond number evaluation, the Fiji image analysis software, and the ilastik® machine learning method). The ANOVA results showed no significant difference ($P=0.985$) in the mean growth rates calculated with the three methods (Figure 1). Therefore, they are equally valid in calculating the growth rate of duckweeds.

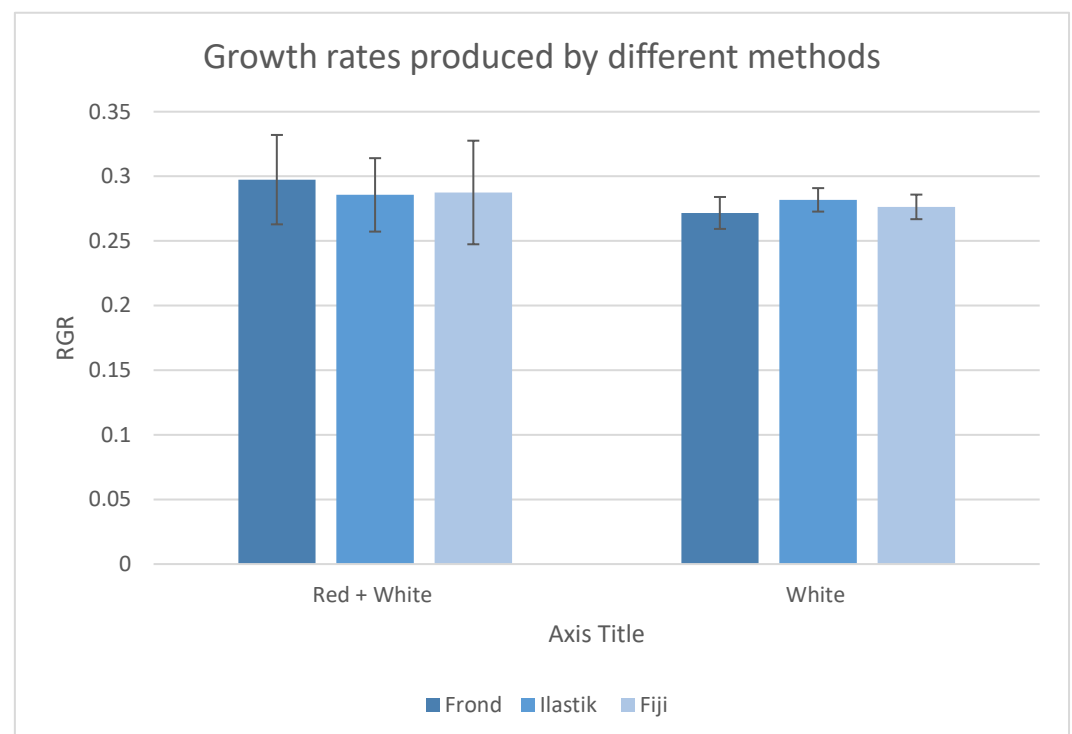


Figure 1. Graph compares RGR calculated with three methods (frond number, ilastik® and Fiji) for the two experiment setups (red light treatment and control).

Unlike the ISO 20079 method, the other two allowed us to calculate plant growth throughout the experiment by analysing a series of photos taken at regular intervals. We used these data to compare the two computerised methods further. Data showed a high correlation between the measurements by ilastik® and Fiji methods, as represented in the scatter diagram (Figure 2). The strong correlation is supported by calculated coefficients of 0.99 for the Pearson coefficient and 0.99 for the Spearman coefficient. More specifically Pearson coefficient showed an almost perfect strength of agreement among data compared. While Spearman's rho coefficient of rank correlation is 0.995. The 95% confidence interval ranges from 0.993 to 0.998. The conclusion is that there is a significant relationship between the two variables.

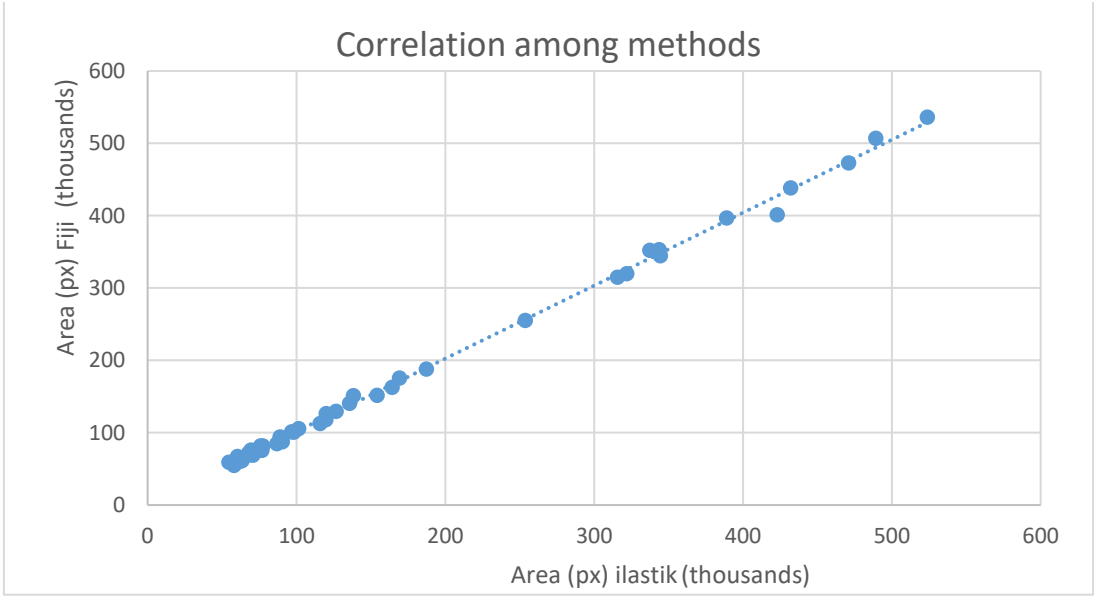


Figure 2. Graph shows plotted output data (pixel) of the same images analysed with the two methods (Fiji and ilastik®). As demonstrated by the data visualisation, a strong correlation between the two methods is present.

Due to the presence of outliers to the median line, we compared results by means of difference. The Bland-Altman (B&A) analysis is reported in Figure 3.

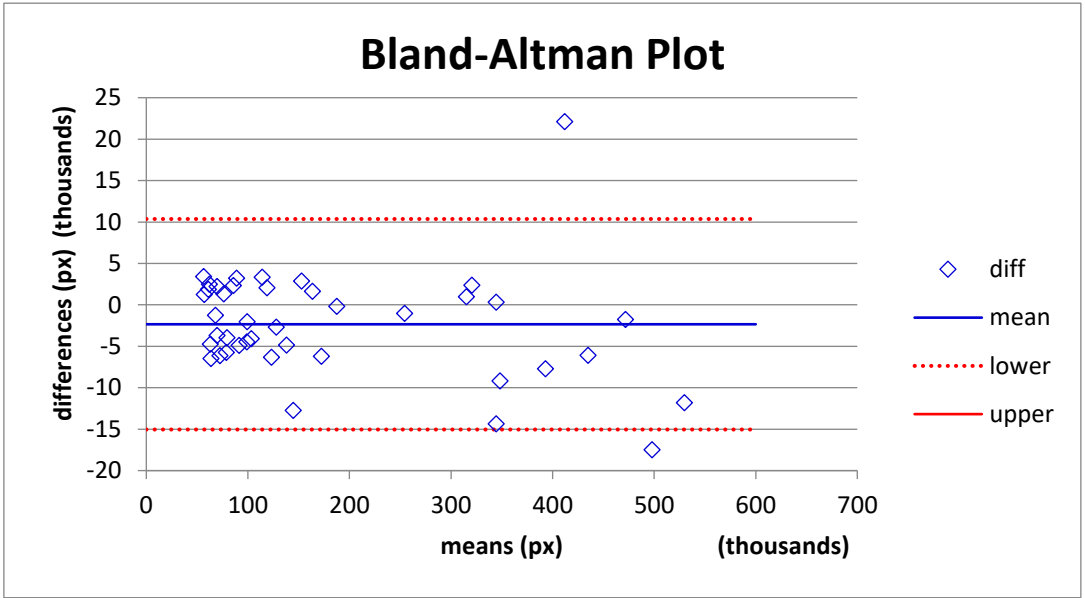


Figure 3. The B&A plot can be evaluated in good agreement according to the scatter dispersion. The scattering of points is reduced, and points lie relatively close to the line representing mean bias. It is essential to consider the big numerical difference existing among data; this difference in the two outlier cases might (outside the limits of agreement) be due to human error during the Fiji analysis [3].

Results from the Heteroskedascity test with the White method have a p-value of 0.06; we fail to reject the null hypothesis and conclude that residuals show a homoscedastic distribution.

The three statistical analyses performed show that the two analysis methods can be used interchangeably.

2.1 Time to data

The previous paragraph shows that the newly described method is perfectly coherent with the results produced with the previously described method (Fiji) regarding data outcomes. We now consider the possible benefits and advantages in terms of time to produce data.

The time required to run the analysis with both software by the same operator was 95.4 seconds for Fiji software and 300 seconds for ilastik® per picture. The main difference between the two methods was that the operator who wanted to run the additional analysis with Fiji needed to start over again. This required the same amount of time per analysis (Figure 4). Differently, the operator that trained the machine by using ilastik® to analyse the first image needed a time longer than that for one image with Fiji, but the operator could immediately run any batch analysis with no additional time required because the machine performed the same task for any number of pictures selected (Figure4).

Furthermore, in the case of ilastik®, it was possible to save the machine training parameters to be applied to possible future pictures taken under identical conditions (light, distance from the camera, camera setup etc.).

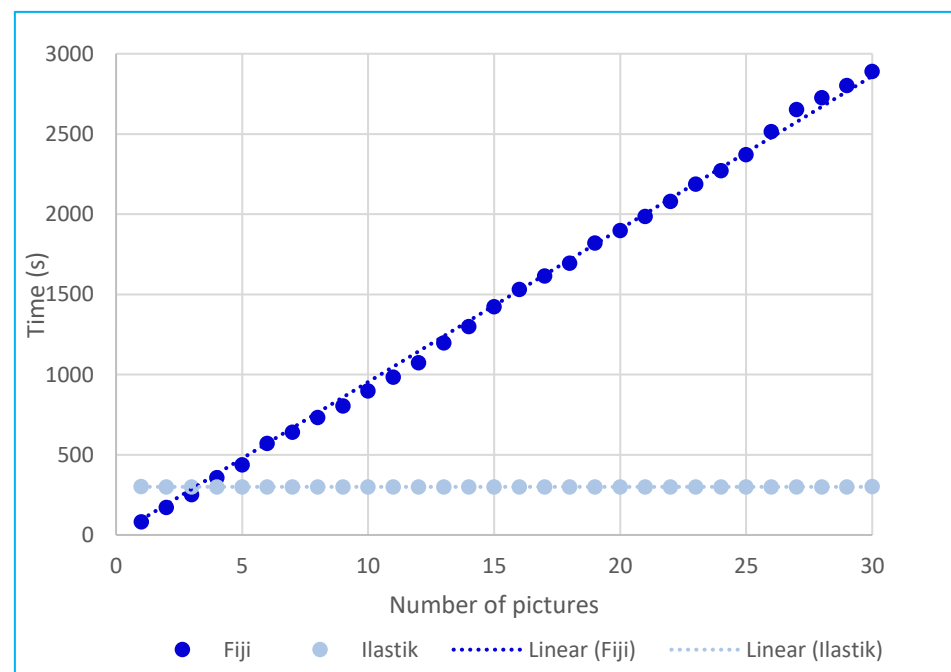


Figure 4. The time required by the operator to analyse the area occupied by fronds of Lemnaceae. The dark blu line represents the time required by the operator by employing the Fiji software; the light blue line represents the time requirements with the ilastik® software. The x-axis represents the number of pictures the y-axis represents the time requirement per picture.

According to the data recorded during the tests, the predictive analysis of time necessary to measure plant growth as a function of the number of images is in perfect accordance

with the time model described by formulas 5 and 6. Equation (5) describes plotted data from Fiji software:

$$(5) y_1 = 95.4x$$

Where y refers to the time (in seconds) required by the operator to perform the analysis and x is the number of images to be analysed. Equation (6) describes the plotted data from the analysis conducted with ilastik® software:

$$(6) y_2 = 300$$

3. Discussion

In this paper, we have demonstrated how a newly described method can be effective in calculating the biological effect of utilising machine learning during image analysis. Plants grown with different light recipes haven't shown different growing patterns by means of relative growth rates. Furthermore, data showed a strong correlation between our newly designed method with the older one (Fiji). Such a strong correlation maximises confidence in the new adoption of the method. Moreover, we have demonstrated how the presence of outliers has been warded off by testing for heteroskedasticity.

It is important to remark on the importance of RGR calculation in the field of study of the Lemnaceae plants as one of the few growth monitoring tools. Upon having compared the RGR outcomes obtained with the different methods, we can conclude that the three methods are equally valuable for studying the growth rate of duckweed plants.

Nevertheless, the methods relying on image analysis do not require destructive measurements and can facilitate other *in vivo* analyses such as genetics [2].

The newly proposed approach of using a machine learning system has numerous advantages and fewer disadvantages than previously proposed methods based on image analysis [27]. In fact, with the initial setup of a photo booth box, researchers can rely on coherent methods that discard human input in the analysis process. It is important to remark that the value of the old methods remain not lowered; however, thanks to the more substantial presence of open-source software and more available technologies, tweaking these systems to researchers' needs has become more accessible.

Results show that the newly proposed method is faster and as reliable as the other methods previously used to measure the growth rate of Lemnaceae [31–32]. Our experiment did not compare results with a fresh or dry weight of plants nevertheless, this was not the main aim of our work, and we considered reliable the correlation between weight and frond area [32]. The application of our method drastically reduces the time required by the operator to analyse the growth rate in Lemnaceae to only the time needed to train the machine. Noticeably, the latter corresponds to the time usually required to analyse a few images with the so far used image analysis methods. Moreover, by applying ilastik® procedure to different experiments designed to use the same photographic conditions and identical clones, the saved machine training can be stored by the researchers and used theoretically infinite times. In these cases, our machine learning approach simplifies the analysis methods to the "click of a button".

Overall, both our result and what has been previously reported in the literature, confirmed the urge to use computer image processing to speed up the innovation process in Lemnaceae. As previously mentioned by other authors [33], the usage of this technology

with low-cost hardware can define new qualitative standards in determining the growth rate of Lemnaceae.

4. Materials and Methods

Plants of *Lemna minor* were grown under identical environmental conditions through temperature, nutrient media, and background light conditions. The advent of Light-emitting diode (LED) technology allows scientists to provide plants with the exact amount and quality of light needed to maximise growth and efficiency. We experimented with different light recipes to validate the machine learning method. More specifically, half of the plant samples were treated with a background light (white) and the other half with the integration of red light (white+red). Details of the cultivation and experiment setup as well as of the data analyses, are reported below.

4.1 Plant cultivation

Lemna minor (9440) was cultivated for 168h in a controlled temperature chamber FOC 200IL Velp scientifica® at a constant temperature of 25±0.5 C°. Five plants with two or more fronds were cultivated in a 150ml glass baker with 100 ml of Murashige and shook growth medium (Sigma-Aldrich - Murashige and Skoog basal medium) (Ph adjusted to 5.8). The bakers were covered with a petri dish to avoid water evaporation. The growth chamber was illuminated with a background white LED light.

Pictures of the growing plants were taken every 24 h with a Sony® alpha 7 II camera equipped with a Sigma® 50mm Art F1.4, mounted on a fixed stand. Photos were shot under an illuminated photo boot with a white background to guarantee optimal sample illumination and contrast. Furthermore, camera photo parameters were kept constant throughout the experiment.

4.2 Photo booth setup

To maintain a constant photo shooting environment, we have set up a photo booth in a dark room of our laboratory. This approach guarantees stable light conditions and centring the samples to the camera frame. We have achieved so by buying a product photo boot online and a camera tripod. Both components were fixed to a table to keep camera distance and centring constant throughout the experiment.

4.3 Light quality and quantity

We have opted for a different light quality setup to stimulate differences in growth; we decided to use the following light treatment setup. Plants were exposed to the same white background light. The existing difference among samples was due to providing extra monochromatic lighting to the red treatment. More specifically, single 3w red coloured LEDs (no branded) were installed to achieve light treatment. Light quality and quantity are described in the following table. They are expressed as average among the three replicas per treatment. Light quality and quantity were measured with a spectroradiometer (SS-110, Apogee Instruments Inc.) to control the emission spectrum of each light treatment.

Table 2. Total Pothon flux density (PPFD) (μmol-s-1), Photosynthetic Photon Flux (PPF) (μmol-m-2-s-1), Yield Photon Flux (YPF) (μmol-s-1), Photosynthetic photon efficacy (PPE) (PPF umol / watts), R/FR is the red (R) light relative to the amount of far-red (FR) light.

Total PPFD	Stdev.	PPF	Stdev.	YPF	Stdev.	PPE	Stdev.	R/FR	Stdev.
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Red treat- ment	128.34	1.1	126.55	1.07	110.72	0.94	0.88	0	10.62	0.03
Control	129.69	0.98	126.53	0.61	107.06	0.66	0.83	0.02	7.08	0.01

4.4 Measuring systems

In this study, we have adopted three different methods to evaluate the relative growth rates of *Lemna minor* during the experiment. As described by the ISO 20079, we have used frond number as an evaluation method for growth during the investigation [2]. The other two approaches were achieved via computer software (Fiji and ilastik® (Figure 7)). Both methods produced quantitative information on the area occupied by the plant (in pixel).

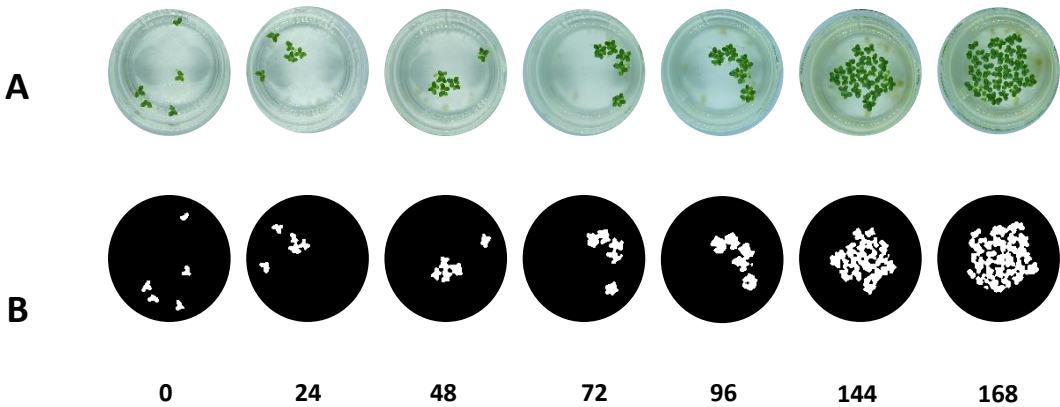


Figure 5. A) pictures of *Lemna minor* at different time intervals B) pictures processed by the ilastik® software.

A plant's growth rate can be calculated from the area or number of fronds described by Haffner et al. [1]. We employed Fiji software as defined by Haffner et al. [1]. The ilastik® software has been used following the protocol described in supplementary materials XX. When training the machine, we started from the pictures where the frond number is the highest to better train the machine in understanding the picture composition. The output file from the software and the feature selected for the first picture is saved and could be used for other analyses. The three measuring systems were used to calculate the relative growth rate (RGR) described by Naumann et al. [2].

4.5 Statistical analysis

Statistical analysis was conducted following four-step phases; first, we compared the three methods' relative growth rates with the formula described by Naumann et al. [2]. In this phase, the outcome for the six growth rates compared to performing a one-way ANOVA analysis was performed with the SPSS software (IBM inc.). The ANOVA was fundamental to confirm that the three methods' growth rates were in accordance. More specifically, the two computed methods were in accordance with the gold standard defined by the ISO 20079 protocol. Subsequently, we compared the proposed method (ilastik® software) with the previously described (Fiji). Agreement among the two computed methods (Fiji and ilastik®) has been shown by calculating correlation coefficients with Pearson and Spearman. As described by McBride, correlation can be defined as almost perfect because the Value of ρ is 0,999 in the range >0.99 [5]. Furthermore, we compared utilising differences between the two measurement techniques with the Bland-Altman plot to underline the presence of bias between the two

methods. As described by Dogan [29], Bland and Altman's limits of agreement (LoA) have conventionally been used in medical research to evaluate the agreement between two methods of measurement for quantitative variables [7]. Nevertheless, Bland and Altman's LoA method may be misleading in the presence of heteroskedastic distributions [8]. Due to the fact of outliers in the Bland and Altman graphic representation, we opted to test heteroskedasticity with the White test because it can better perform in the presence of non-linear forms of heteroskedasticity (presence of outliers).

4.6 Time analysis of the software utilisation

The following formula has been utilised to compare the time usage of the two software:

$$(1) \quad y = a + bx$$

Where y is the time required by the operator to perform analysis with the software under evaluation, a is the time to set up the analysis with the given software. The letter b indicates the time for the operator to analyse a single image, and x is the number of images analysed.

Consequently, the equation can be solved respectively for Fiji (2) and ilastik® (3).

$$(2) \quad y_1 = bx$$

$$(3) \quad y_2 = a$$

To validate what has been modelled by the mathematical equations, we have provided quantitative data about the time to analyse the two computed methods.

4.7 Relative Growth rate

Growth has been calculated following the growth rate calculation, Ziegler et al. 2014 [9].

$$(4) \quad RGR = (\ln X_{t_n} - \ln X_{t_0}) / (t_n - t_0)$$

Where X is the pixel in the area as described by Haffner et al. [1] and t_0 , t_n represents the start and the end of the test, respectively.

5. Conclusions

We have presented a novel method designed to rapidly and inexpensively quantify the Lemnaceae growth rate by tracking frond surface variance at different time intervals. We have shown that machine learning technology can substitute traditional methods and document biological phenomena if well applied to the observed system. The proposed approach helps the researcher train the machine ad-hoc to the specific requirements. This customisable method can be used across different Lemnaceae applications and other surface-floating aquatic plants.

It is important to remark on the importance of RGR calculation in the field of study of the Lemnaceae plants as one of the few monitors for growth. Upon having compared the RGR outcomes obtained with the different methods, we can conclude that the three methods are equally valuable for studying the growth rate of duckweed plants.

Nevertheless, the methods relying on image analysis do not require destructive measurements and can facilitate other *in vivo* analyses such as genetics [2].

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Supplementary Materials: The following supporting information can be downloaded at: www.mdpi.com/xxx/s1 here you will find a step by step guide to perform the RGR analysis with ilastik software.

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References

1. William, W.; Ware, A.; Basaza-Ejiri, A.H.; Obungoloch, J. A Review of Image Analysis and Machine Learning Techniques for Automated Cervical Cancer Screening from Pap-Smear Images. *Computer Methods and Programs in Biomedicine* 2018, 164, 15–22, doi:10.1016/J.CMPB.2018.05.034.
2. Lobet, G. Image Analysis in Plant Sciences: Publish Then Perish. *Trends in Plant Science* 2017, 22, 559–566, doi:10.1016/J.TPLANTS.2017.05.002.
3. Popescu, G. v.; Noutsos, C.; Popescu, S.C. Big Data in Plant Science: Resources and Data Mining Tools for Plant Genomics and Proteomics. *Methods Mol Biol* 2016, 1415, 533–547, doi:10.1007/978-1-4939-3572-7_27.
4. Berman, J.J. *Principles of Big Data*; Elsevier Inc., 2013; ISBN 9780124045767.
5. Oliveira, M.C.; Osipitan, O.A.; Begcy, K.; Werle, R. Cover Crops, Hormones and Herbicides: Priming an Integrated Weed Management Strategy. *Plant Sci* 2020, 301, doi:10.1016/J.PLANTSCI.2020.110550.
6. Chlingaryan, A.; Sukkarieh, S.; Whelan, B. Machine Learning Approaches for Crop Yield Prediction and Nitrogen Status Estimation in Precision Agriculture: A Review. *Computers and Electronics in Agriculture* 2018, 151, 61–69, doi:10.1016/J.COM-PAG.2018.05.012.
7. McCabe, M.F.; Tester, M. Digital Insights: Bridging the Phenotype-to-Genotype Divide. *Journal of Experimental Botany* 2021, 72, 2807–2810, doi:10.1093/JXB/ERAB108.
8. Zhang, D.; Tsai, J. Advances in Machine Learning Applications in Software Engineering. *Computer Science* 2007, doi:10.4018/978-1-59140-941-0.
9. Cortés, A.J.; Restrepo-Montoya, M.; Bedoya-Canas, L.E. Modern Strategies to Assess and Breed Forest Tree Adaptation to Changing Climate. *Frontiers in Plant Science* 2020, 11, doi:10.3389/FPLS.2020.583323.
10. Ma, C.; Xin, M.; Feldmann, K.A.; Wang, X. Machine Learning–Based Differential Network Analysis: A Study of Stress-Responsive Transcriptomes in Arabidopsis. *The Plant Cell* 2014, 26, 520–537, doi:10.1105/TPC.113.121913.
11. Ma, C.; Zhang, H.H.; Wang, X. Machine Learning for Big Data Analytics in Plants. *Trends in Plant Science* 2014, 19, 798–808, doi:10.1016/J.TPLANTS.2014.08.004.
12. Marx, V. The Big Challenges of Big Data. *Nature* 2013 498:7453 2013, 498, 255–260, doi:10.1038/498255a.
13. Schatz, M.C. Computational Thinking in the Era of Big Data Biology. *Genome Biol* 2012, 13, 177, doi:10.1186/GB-2012-13-11-177/METRICS.
14. Bruce Ratner. *Statistical and Machine-Learning Data Mining*; Routledge, Taylor & Francis Group, 2017; ISBN 9780367573607.
15. Bassel, G.W.; Glaab, E.; Marquez, J.; Holdsworth, M.J.; Bacardit, J. Functional Network Construction in Arabidopsis Using Rule-Based Machine Learning on Large-Scale Data Sets. *The Plant Cell* 2011, 23, 3101–3116, doi:10.1105/TPC.111.088153.
16. Berg, S.; Kutra, D.; Kroeger, T.; Straehle, C.N.; Kausler, B.X.; Haubold, C.; Schiegg, M.; Ales, J.; Beier, T.; Rudy, M.; et al. Ilastik: Interactive Machine Learning for (Bio)Image Analysis. *Nature Methods* 2019 16:12 2019, 16, 1226–1232, doi:10.1038/s41592-019-0582-9.
17. Sommer, C.; Straehle, C.; Kothe, U.; Hamprecht, F.A. Ilastik: Interactive Learning and Segmentation Toolkit. *Proceedings - International Symposium on Biomedical Imaging* 2011, 230–233, doi:10.1109/ISBI.2011.5872394.

18. Haubold, C.; Schiegg, M.; Kreshuk, A.; Berg, S.; Koethe, U.; Hamprecht, F.A. Segmenting and Tracking Multiple Dividing Targets Using *ilastik*. *Advances in Anatomy Embryology and Cell Biology* 2016, 219, 199–229, doi:10.1007/978-3-319-28549-8_8.
19. Sree, K.S.; Sudakaran, S.; Appenroth, K.J. How Fast Can Angiosperms Grow? Species and Clonal Diversity of Growth Rates in the Genus *Wolffia* (Lemnaceae). *Acta Physiologiae Plantarum* 2015, 37, doi:10.1007/S11738-015-1951-3.
20. Appenroth, K.-J.; Sree, K.S.; Bog, M.; Ecker, J.; Seeliger, C.; Böhm, V.; Lorkowski, S.; Sommer, K.; Vetter, W.; Tolzin-Banasch, K.; et al. Nutritional Value of the Duckweed Species of the Genus *Wolffia* (Lemnaceae) as Human Food. *Frontiers in Chemistry* 2018, 0, 483, doi:10.3389/FCHEM.2018.00483.
21. Appenroth, K.J.; Sree, K.S.; Böhm, V.; Hammann, S.; Vetter, W.; Leiterer, M.; Jahreis, G. Nutritional Value of Duckweeds (Lemnaceae) as Human Food. *Food Chemistry* 2017, 217, 266–273, doi:10.1016/J.FOODCHEM.2016.08.116.
22. Fujita, M.; Mori, K.; Kodera, T. Nutrient Removal and Starch Production through Cultivation of *Wolffia* *Arrhiza*. *J Biosci Bioeng* 1999, 87, 194–198, doi:10.1016/S1389-1723(99)89012-4.
23. Gupta, C.; Prakash, D. Duckweed: An Effective Tool for Phyto-Remediation. <http://dx.doi.org/10.1080/02772248.2013.879309> 2014, 95, 1256–1266, doi:10.1080/02772248.2013.879309.
24. Romano, L.E.; Aronne, G. The World Smallest Plants (*Wolffia* Sp.) as Potential Species for Bioregenerative Life Support Systems in Space. *Plants* 2021, Vol. 10, Page 1896 2021, 10, 1896, doi:10.3390/PLANTS10091896.
25. Acosta, K.; Appenroth, K.J.; Borisjuk, L.; Edelman, M.; Heinig, U.; Jansen, M.A.K.; Oyama, T.; Pasaribu, B.; Schubert, I.; Sorrels, S.; et al. Return of the Lemnaceae: Duckweed as a Model Plant System in the Genomics and Postgenomics Era. *The Plant Cell* 2021, doi:10.1093/PLCELL/KOAB189.
26. Berg, S.; Kutra, D.; Kroeger, T.; Straehle, C.N.; Kausler, B.X.; Haubold, C.; Schiegg, M.; Ales, J.; Beier, T.; Rudy, M.; et al. *Ilastik*: Interactive Machine Learning for (Bio)Image Analysis. *Nature Methods* 2019 16:12 2019, 16, 1226–1232, doi:10.1038/s41592-019-0582-9.
27. Haffner, O.; Kučera, E.; Drahoš, P.; Cigánek, J.; Kozáková, A.; Urminská, B. Lemna Minor Bioassay Evaluation Using Computer Image Analysis. *Water* 2020, Vol. 12, Page 2207 2020, 12, 2207, doi:10.3390/W12082207.
28. Naumann, B.; Eberius, M.; Appenroth, K. Growth Rate Based Dose-Response Relationships and EC-Values of Ten Heavy Metals Using the Duckweed Growth Inhibition Test (ISO 20079) with *Lemna Minor* L. Clone St. *J Plant Physiol* 2007, 164, 1656–1664, doi:10.1016/J.JPLPH.2006.10.011.
29. JS, K. Why Bland-Altman Plots Should Use X , Not $(Y+X)/2$ When X Is a Reference Method. *Stat Med* 2008, 27, 778–780, doi:10.1002/SIM.3086.
30. Hayes, A.F.; Cai, L. Using Heteroskedasticity-Consistent Standard Error Estimators in OLS Regression: An Introduction and Software Implementation. *Behavior Research Methods* 2007 39:4 2007, 39, 709–722, doi:10.3758/BF03192961.
31. Mazur, R.; Szoszkiewicz, K.; Lewicki, P.; Bedla, D. The Use of Computer Image Analysis in a *Lemna Minor* L. Bioassay. *Hydrobiologia* 2018, 812, 193–201, doi:10.1007/S10750-016-2972-7/TABLES/4.
32. Haffner, O.; Kučera, E.; Drahoš, P.; Cigánek, J.; Kozáková, A.; Urminská, B. Lemna Minor Bioassay Evaluation Using Computer Image Analysis. *Water* 2020, Vol. 12, Page 2207 2020, 12, 2207, doi:10.3390/W12082207.
33. Ziegler, P.; Adelman, K.; Zimmer, S.; Schmidt, C.; Appenroth, K. Relative in Vitro Growth Rates of Duckweeds (Lemnaceae) - the Most Rapidly Growing Higher Plants. *Plant Biol (Stuttg)* 2015, 17 Suppl 1, 33–41, doi:10.1111/PLB.12184.
34. McBride, G.B. A Proposal for Strength-of-Agreement Criteria for Lin's Concordance Correlation Coefficient.; 2005;
35. Doğan, N.Ö. Bland-Altman Analysis: A Paradigm to Understand Correlation and Agreement. *Turkish Journal of Emergency Medicine* 2018, 18, 139, doi:10.1016/J.TJEM.2018.09.001.
36. Sedgwick, P. Limits of Agreement (Bland-Altman Method). *BMJ* 2013, 346, doi:10.1136/BMJ.F1630.
37. Ludbrook, J. Confidence in Altman-Bland Plots: A Critical Review of the Method of Differences. *Clin Exp Pharmacol Physiol* 2010, 37, 143–149, doi:10.1111/J.1440-1681.2009.05288.X.