

Acesse meu site e entre em contato:

https://endriowebers.github.io/

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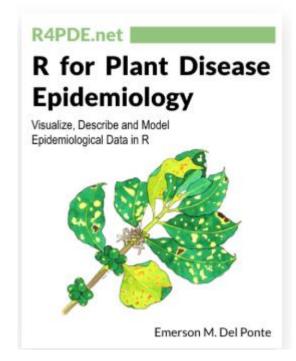
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• Romero, D., De Vicente, A., Rakotoaly, R. H., Dufour, S. E., Veening, J. W., Arrebola, E., ... & Pérez-García, A. (2007). The iturin and fengycin families of lipopeptides are key factors in antagonism of Bacillus subtilis toward Podosphaera fusca. *Molecular Plant-Microbe Interactions*, 20(4), 430-440.

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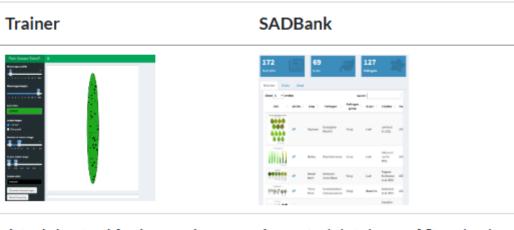
Links uteis:

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Links uteis

• https://emersondelponte.netlify.app/apps



A training tool for increasing the accuracy of visual assessments of plant disease severity (percent diseased area). A curated database of Standard Area Diagrams (SADs) for aiding visual assessments of disease severity.

Links uteis

https://tiagoolivoto.github.io/pliman/

pliman 2.1.0 ★ Home Examples ▼ Functions News ▼ More ▼ Search for

pliman



The pliman (**pl**ant **im**age **an**alysis) package is designed to analyze plant images, particularly for leaf and seed analysis. It offers a range of functionalities to assist with various tasks such as measuring disease severity, counting lesions, obtaining lesion shapes, counting objects in an image, extracting object characteristics, performing Fourier Analysis, obtaining RGB values, extracting object coordinates and outlines, isolating objects, and plotting object measurements.

pliman also provides useful functions for image transformation, binarization, segmentation, and resolution. Please visit the Examples page on the pliman website for detailed documentation of each function.