

UNIVERSITÀ DEGLI STUDI DI TORINO



*Analysis of the spatial interdependence of nuclear size in confocal microscopy images of plant roots*

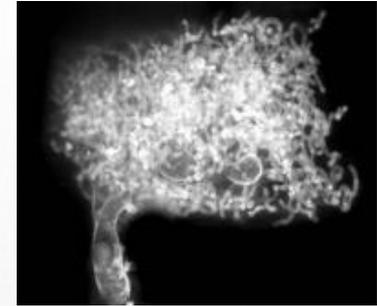
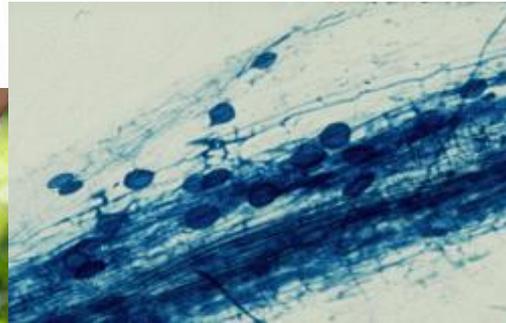
*Analisi della interdipendenza spaziale della dimensione dei nuclei in radici vegetali osservate in microscopia confocale*

Ivan Sciascia<sup>1</sup>, Andrea Crosino<sup>1</sup>, Gennaro Carotenuto<sup>1</sup>, Andrea Genre<sup>1</sup>

Keywords: mycorrhized roots, estimated variogram, nuclei size

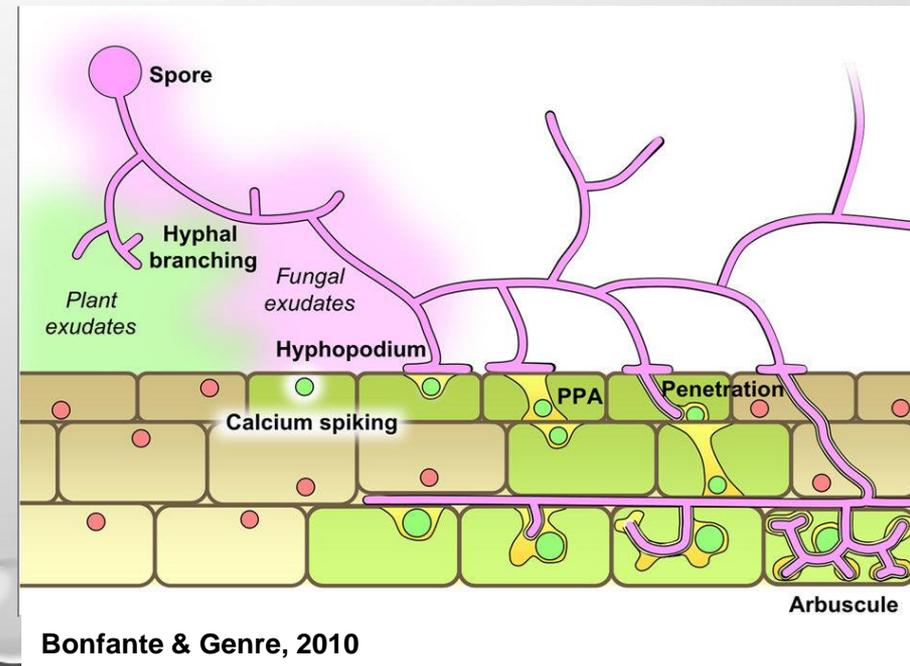
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# AM symbiosis



AM Symbiosis, beneficial interaction

Arbuscules, the structures devoted to the exchange of mineral nutrients for sugars and lipids



# AM symbiosis

The symbiosis between fungi and plants leads to a mutual exchange of nutrients which brings advantages to the plant in terms of greater growth, resistance to stress and pathogens.



# What we already know about endoreduplication and nuclear size



## Mycorrhizal symbiosis stimulates endoreduplication in angiosperms

L. D. BAINARD<sup>1\*</sup>, J. D. BAINARD<sup>1\*</sup>, S. G. NEWMAS<sup>1</sup> & J. N. KLIRONOMOS<sup>2</sup>

*Plant and Soil* 226: 37–44, 2000.  
© 2000 Kluwer Academic Publishers. Printed in the Netherlands.

## Ploidy in tomato roots as affected by arbuscular mycorrhizal colonization

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*Eur. J. Histochem.*  
45: 9-20, 2001  
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## REVIEW

## The nucleus of differentiated root plant cells: modifications induced by arbuscular mycorrhizal fungi

G. Lingua<sup>1</sup>, A. Fusconi<sup>2</sup> and G. Berta<sup>1</sup>

## METHODOLOGY ARTICLE

Carotenuto et al. *BMC Plant Biology* (2019) 19:180  
<https://doi.org/10.1186/s12870-019-1791-1>

## Size matters: three methods for estimating nuclear size in mycorrhizal roots of *Medicago truncatula* by image analysis

Gennaro Carotenuto, Ivan Sciascia, Ludovica Oddi, Veronica Volpe and Andrea Genre<sup>\*</sup>

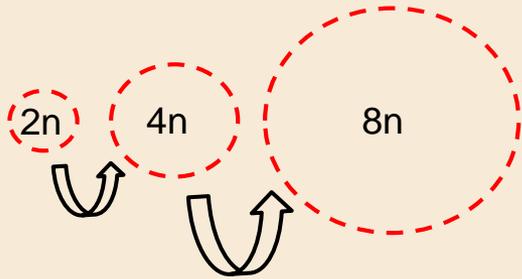
BMC Plant Biology

Open Access



Previous studies have highlighted endoreduplication in several plant-microbe interaction

Nuclei size and endoreduplication:



Double DNA content without dividing

Endoreduplication events have been analyzed through:

Use of flow cytometry to analyze the occurrence of endoreduplication in *M. truncatula* roots.

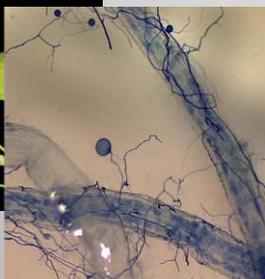
Gene expression analyses of endocycle initiation markers

Image post-processing from confocal Z-stacks

Investigation of ploidy changes in wild-type *Medicago truncatula* roots

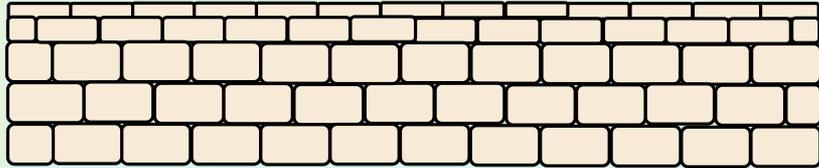
This presentation focuses on post-processing set-up

*Medicago truncatula*



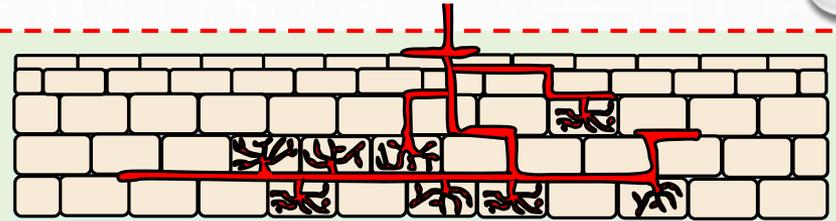
*Gigaspora margarita*

# Workflow:

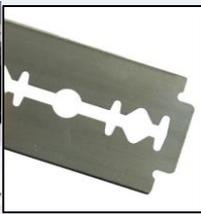


1cm-long Control root segments

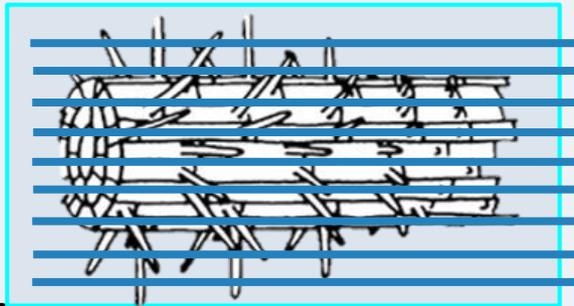
Excision



1cm-long Mycorrhizal root segments



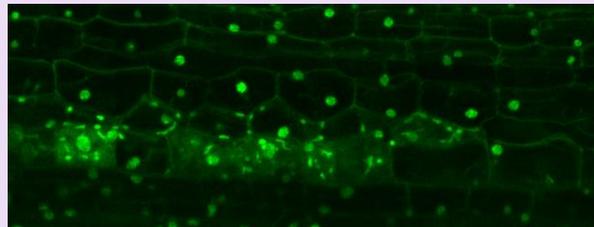
Vibratome sectioning



100 $\mu$ m-thick slices

We obtained 100 micrometer thick slices which were prepared with fluorescent dyes for DNA, that label the nuclei

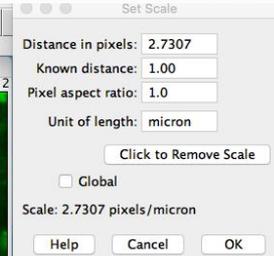
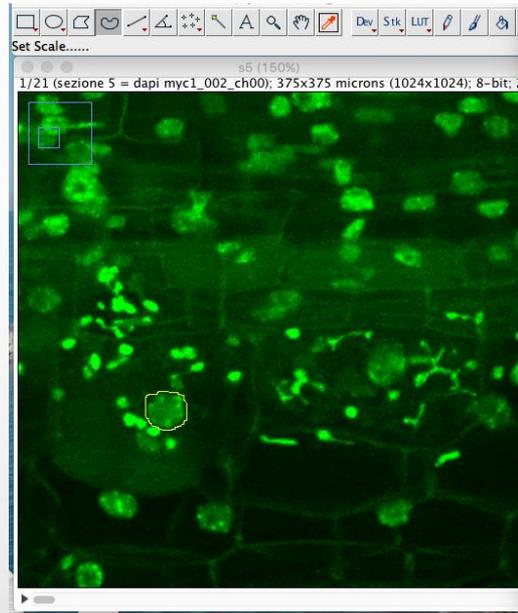
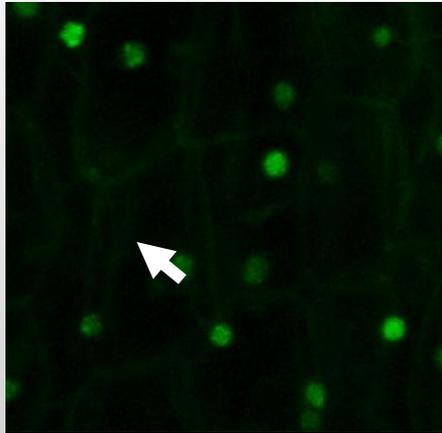
These samples were then imaged on the confocal microscope



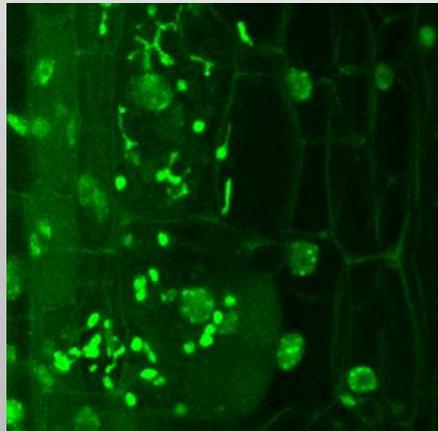
Z-series, taken with 1.5 micrometer Z-step

We observed and measured the size of hundreds of nuclei for every section

# Manual measurements with ImageJ:



Results								
	Area	Perim.	Circ.	IntDen	RawIntDen	AR	Round	Solidity
1	84.355	33.772	0.929	9435.475	70356	1.110	0.901	0.964



# Spatial statistical analysis: mark variogram

Analysis techniques derived from geostatistics dealing with spatially correlated data

Analysis of the correlation between nuclear size and distances

Mathematical models that describe variability as a function of distance

# Mark variogram: areas and distances

To calculate the semivariance as a function of the distance between the nuclei we first calculate the Euclidean distances and then the squared differences between the dimensions of the areas of the nuclei

	Area	BX	BY
1	14,484	76,904	5,127
2	13,947	37,354	31,128
3	34,735	121,216	56,396
4	8,583	58,228	59,692
5	3,219	62,256	80,566

Spatial coordinates

Euclidean distances, above the diagonal of the matrix



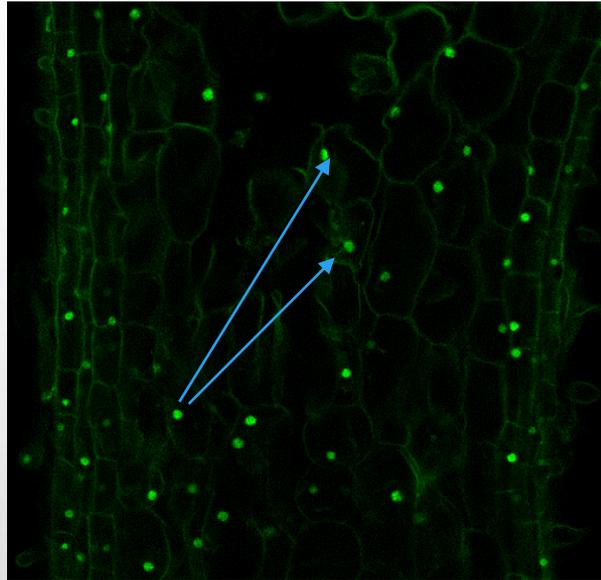
	Area	1	2	3	4	5
1	14,484		47,33133	67,76477	57,67263	76,84794
2	13,947	0,288369		87,586	35,37833	55,35545
3	34,735	410,103	432,1409		63,07418	63,72182
4	8,583	34,8218	28,7725	683,9271		21,25908
5	3,219	126,9002	115,09	993,2583	28,7725	



Squared area differences, below the diagonal of the matrix

Semivariance is a function of distance between pairs of objects and the plot experimental function is called variogram:

$$\gamma(h) = \frac{1}{2N(h)} \sum_{(p_i, p_{(i+h)}) \in S} [y(p_i) - y(p_{(i+h)})]^2$$



If we consider digital images from confocal microscopy with resolution  $\delta = 2,7307 \frac{\text{pixel}}{\mu\text{m}}$  as in Figure 1 the elements of the

formula (1) are:

$\gamma(h)$  is the semivariance in function of the distance  $h$

$N(h)$  are the root nuclei pairs at distance  $h$

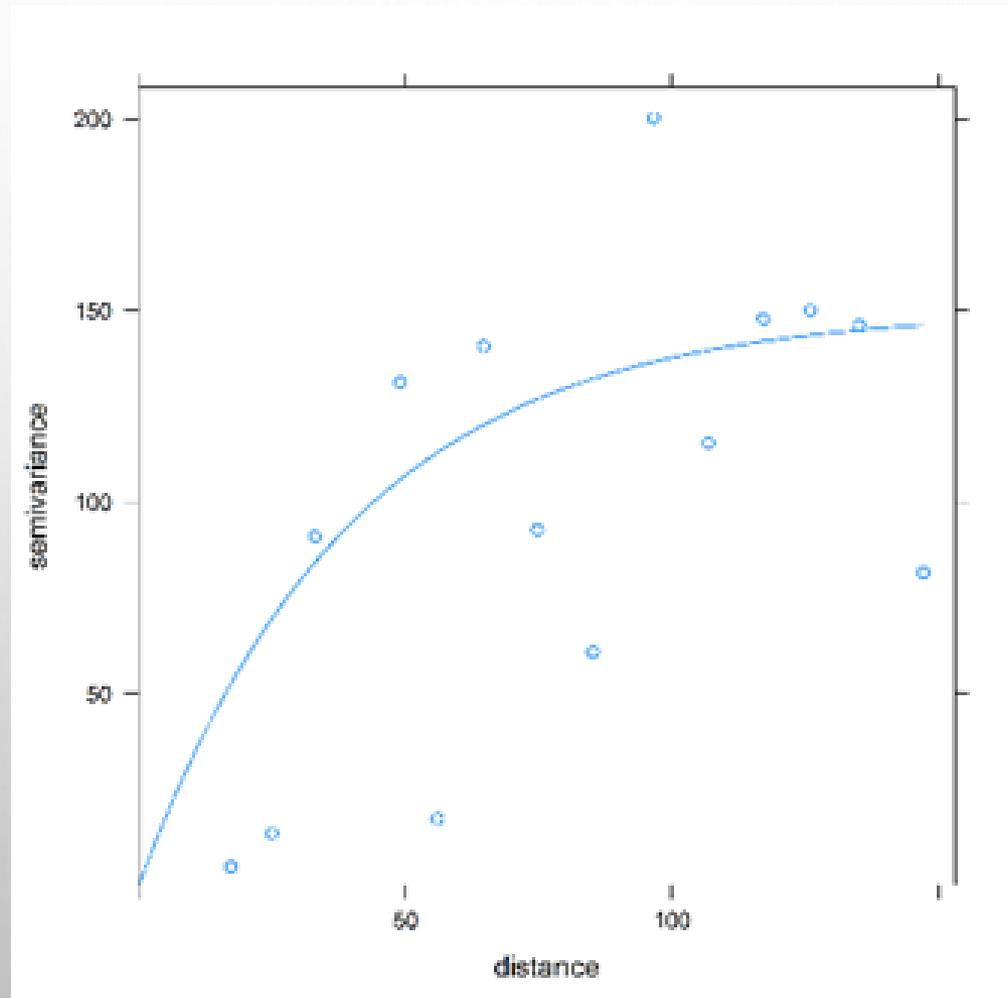
$S$  is the frame sample

$y(p_i)$  is the nucleus area in  $\mu\text{m}^2$  at the coordinates  $p_i$

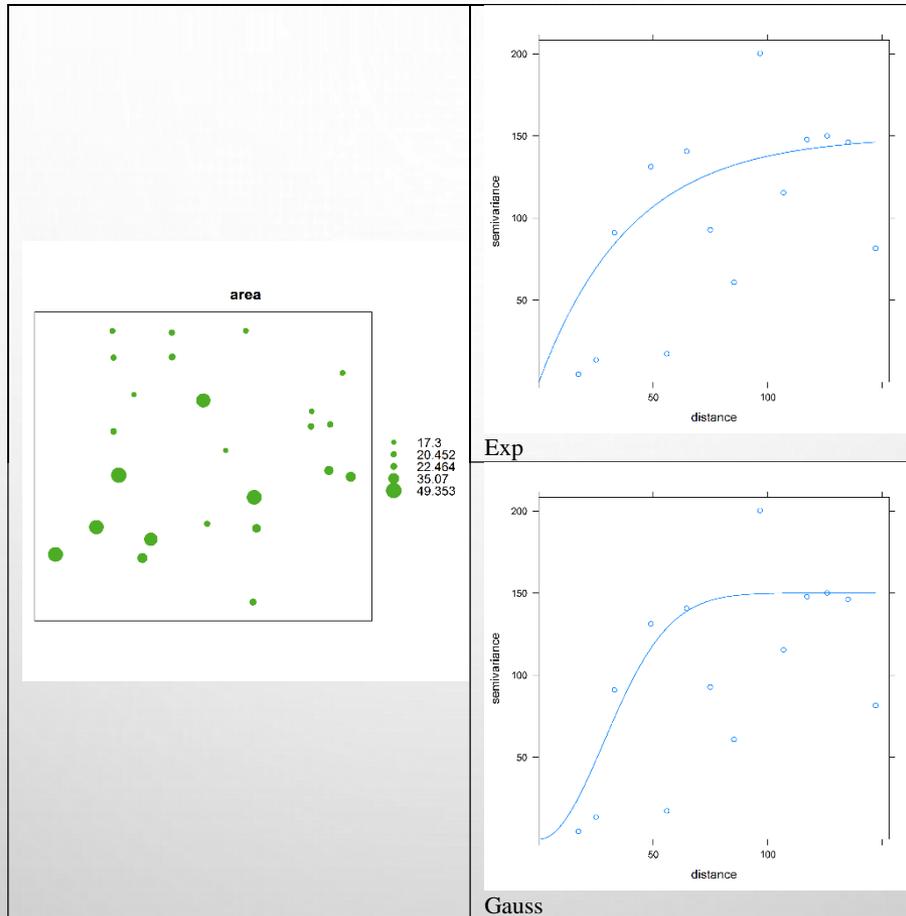
$y(p_{(i+h)})$  is the nucleus area in  $\mu\text{m}^2$  at the coordinates  $p_{(i+h)}$

# Calculation of experimental and theoretical variograms

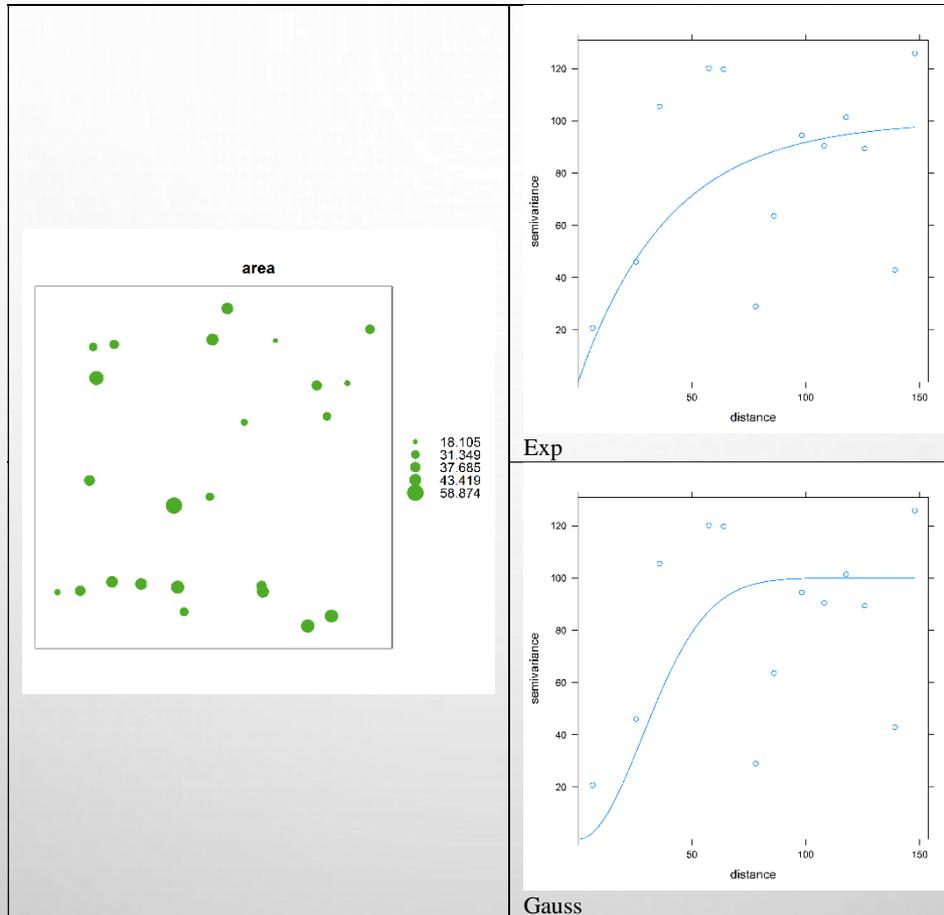
The experimental variogram consists of a cloud of points where the semivariance is a function of the distance. The theoretical variograms are point cloud adaptation curves, we use the exponential and Gaussian models.



Theoretical models have characteristic parameters, the sill, the range and the nugget. The value of the sill of the theoretical graph was set on the basis of the variance of the sample measurements while the range was visually evaluated and set equal for the two data series. The nugget was conventionally set to zero.



Area plot of control nuclei and estimated exponential and Gaussian variograms



Area plot of mycorrhized nuclei and estimated exponential and Gaussian variograms

## Performance indexes for the estimation models

<i>Model</i>	<i>MAD</i>	<i>RMSE</i>
Control Exponential	3,784341	4,735795
Control Gaussian	3,616071	4,948705
Mycorrhized Exponential	2,459242	3,201468
Mycorrhized Gaussian	2,702641	3,368925

From this first study using theoretical semivariance estimation models we observed that the estimation models perform better for the frame sample of mycorrhized roots. This may suggest minor variability in the distribution of nuclei sizes, to be investigated with other research in confocal microscopy.

Thank you for your attention!

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