

Ungulate effects on root fungi community at Mediterranean montane areas

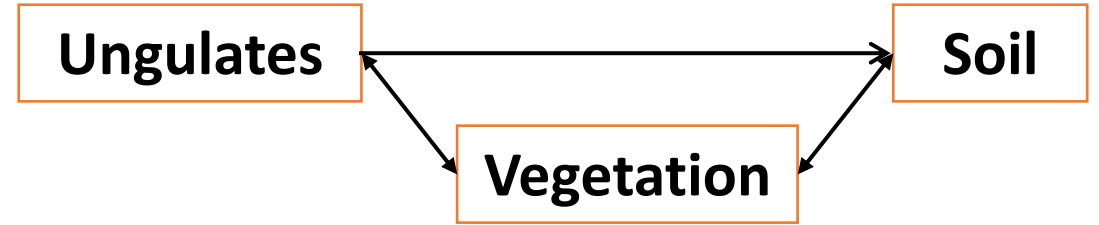
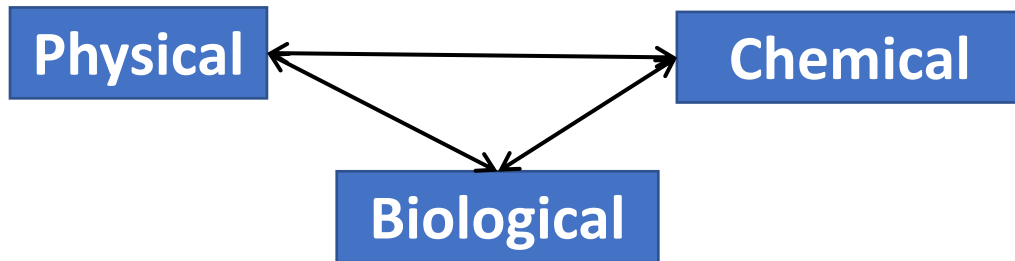
Dra. Elena Baraza; Dra. Jennifer Adams Krumins; Daniel Gamba; INCREMENTO-
Consortium; Dr. Ramon Perea

The Incremento Consortium

Carlos Alonso¹, Elena Baraza², Matthew Brolly³, Niall Burnside³, Juan Antonio Calleja⁴, Mónica Candel⁵, Jesús Cardells⁶, Joao Carvalho⁷, Marcus Clauss⁸, Rafaela Cuenca⁹, Pedro Fernández-Llario¹⁰, Iolanda Filella¹¹, Carlos Fonseca⁷, Daniel Gamba¹, Carlos Hernández^{1,11}, Miguel Ibañez-Alvarez², Anna Jolles¹², Jennifer Krumins¹³, Santiago Lavín⁹, Víctor Lizana⁶, María Martínez-Jauregui¹⁴, Gregorio Mentaberre¹⁵, Marta Peláez¹, Josep Peñuelas¹¹, Maurizio Ramanzin¹⁶, David Risco^{10,17}, Alfonso San Miguel¹, Rita Tinoco⁷, Rafael Villafuerte-Jordán¹⁴.



Soil Characteristics



<https://alternativa-verde.com/2017/04/02/pastizales-y-desertificacion-la-salvacion-esta-en-la-ganaderia/>

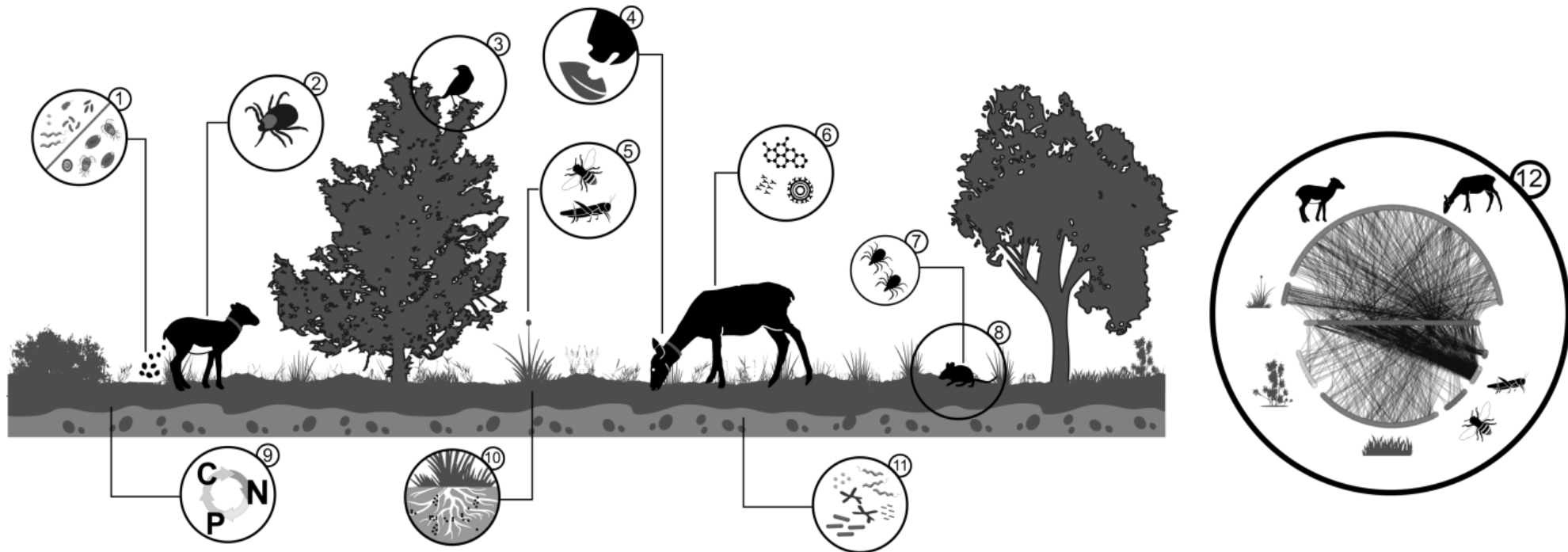
Variable	Decrease	Same	Increase	Variable	Total
Apparent density	70%	19%	0	11%	27
C/N ratio	13%	33%	20%	33%	15
% OM o SOC	9%	30%	51%	11%	47
% N total	12%	28%	4%	2%	25
Microbial activity	30%	15%	33%	22%	27
Ph	36%	36%	12%	16%	25

TFG Universitat de les Illes Balears: Abellán Pérez, Rubén

Systematic revision of exclosure experiments to investigate herbivore impacts found strong variability in direction, magnitude, and modifiers of ungulates effects

Global Project

Main Hypothesis: Changes in wild ungulate populations cause significant changes in the structure, composition and functioning of ecosystems, and therefore, in the provision of ecosystem services.



1= Microbes in faeces 2= Ticks 3=Vertebrate diversity and seed dispersal; 4= Plant preferences, herbivory and plant regeneration; 5=Invertebrate diversity, pollination and pests; 6= Haematology, nutritional condition; 7-8= small mammal abundance and disease risk; 9=Biogeochemical cycling; 10= mycorrhizal functioning; 11= microbial diversity; 12=Mutualistic and multi-trophic networks

Specific Objective:

Does the community of fungi associated with the roots of plants that grow in areas with the presence of ungulate herbivores change compared to those that grow without them?



Two areas of Spain
Quintos (Toledo)
Muela (Valencia)
with a drier climate



QUINTOS DE MORA

Long time exclusions since 2007

We collected five samples of *Agrostis sp* and *Rubus sp* roots per plot



QUINTOS DE MORA

Five samples of *Cistus ladanifer* and *Thymus communis* roots per plot

Non-ungulates

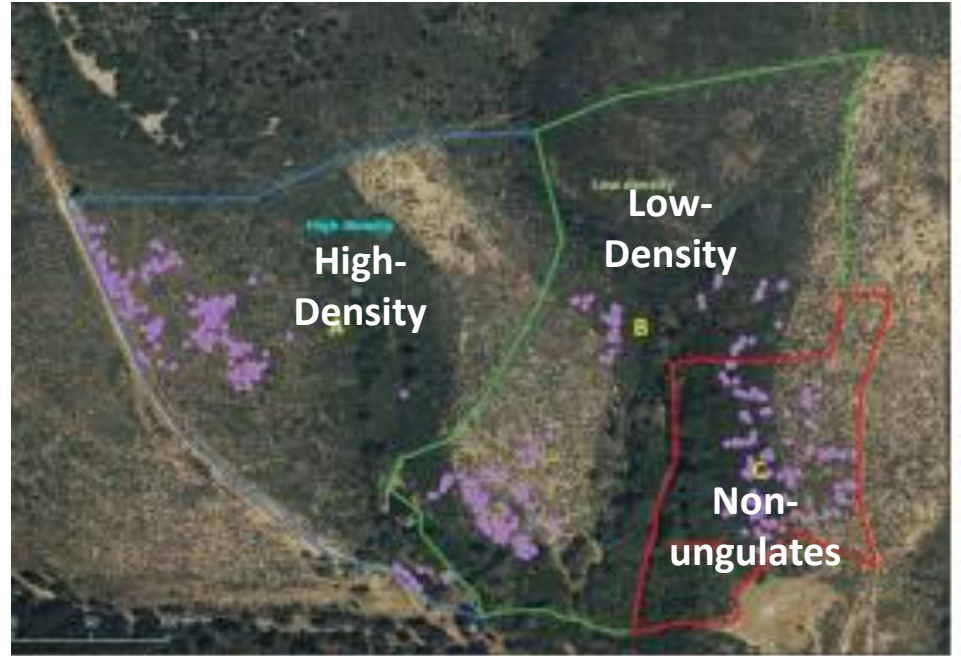
Only during 3 months



Low Density
45 ind/km²

High density
90 ind/km²

small female
deer



MUELA DE CORTES

Five samples of *Cistus albidus* and *Thymus mastichina* roots per plot.

Non-ungulates



Low Density
45 ind/km²

High density
90 ind/km²

small female
deer

Low-Density

Non-ungulates

High-Density

Non-ungulates



DNA was extracted from roots and analyzed via Illumina sequencing of the Internal Transcribed Spacer (ITS)

The primers ITS4Rev/ITS2F-ITS7Fw were used.

DADA2 pipeline in QUIMEII

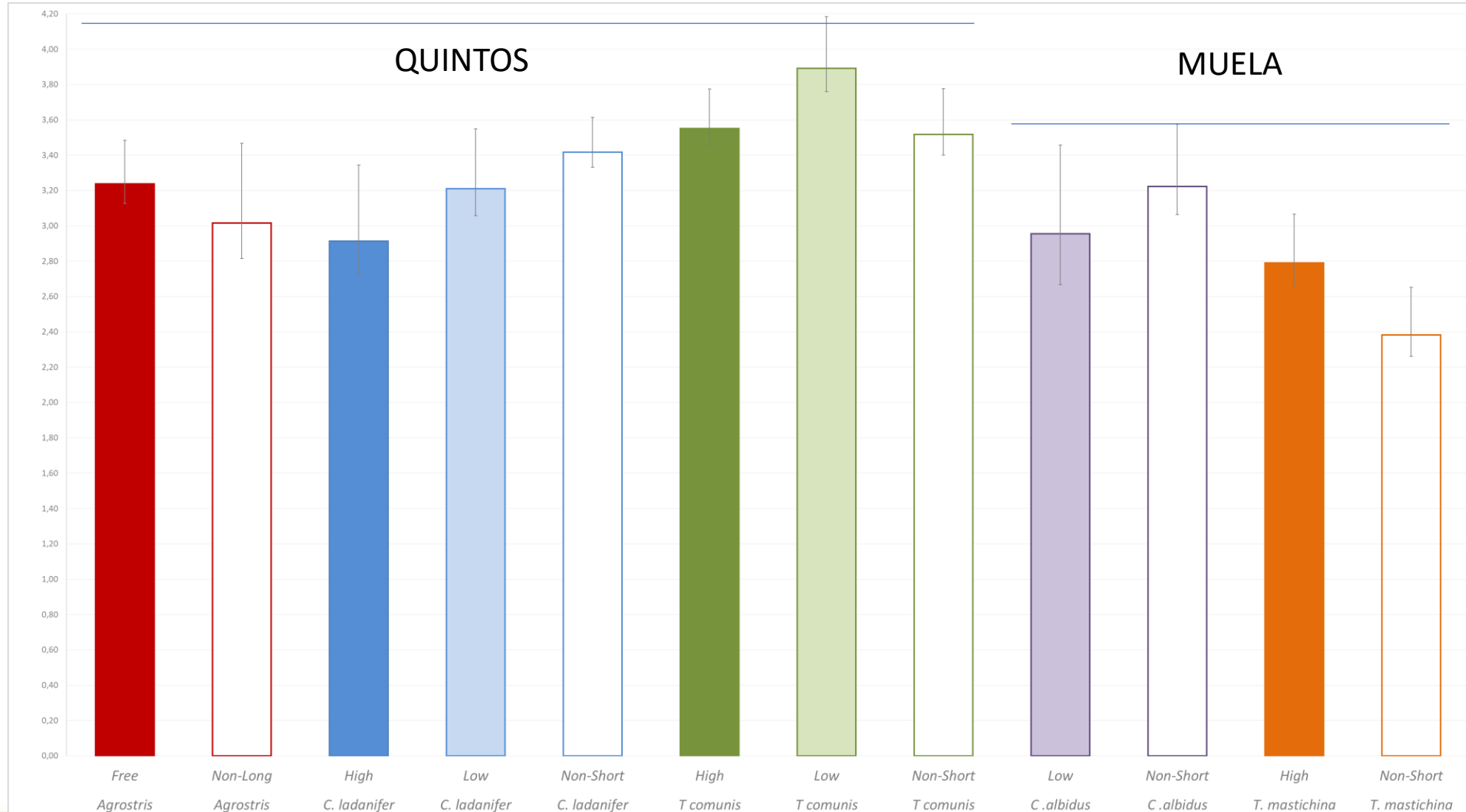
5105 ASV classify as Fungi

ASVs with less than 25 reads were eliminate

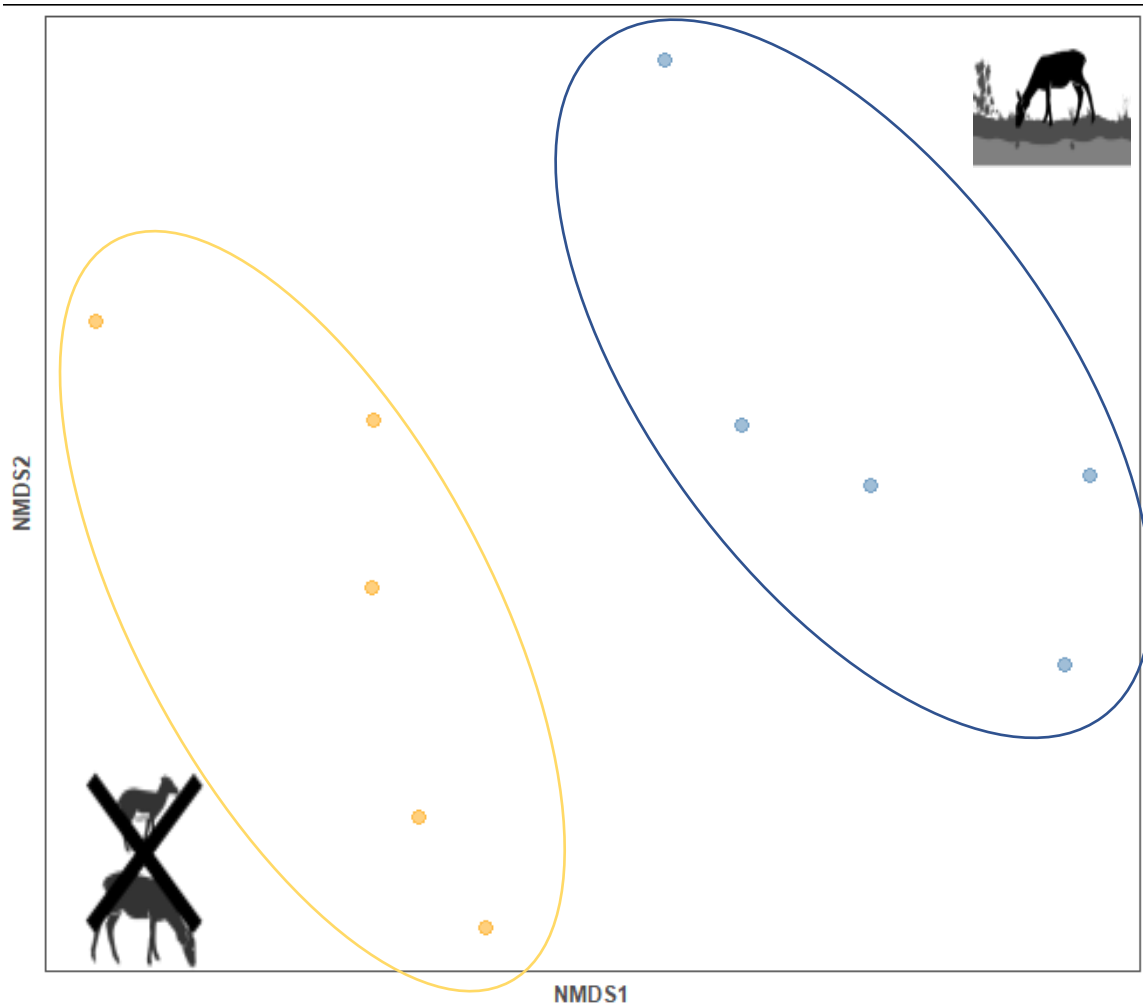
2560 ASVs considered

General Shannon Index diversity of ASVs

Not statistically significant effects



After 14 years of ungulate exclusion fungi community associate to roots of *Agrostis sp* is significantly different.



Ungulates exclusion explanatory value for the fungal communities was fitted into multidimensional scaling ordination (beta diversity was calculated using Bray-Curtis dissimilarity indices) and tested using permutation test using “envfit” function in vegan package.

Ungulates exclusion Goodness of fit:
 $R^2 = 0.5576$ $P = 0.007$

No *Rubus* root samples amplify

Quintos after 3 month of Ungulate presence



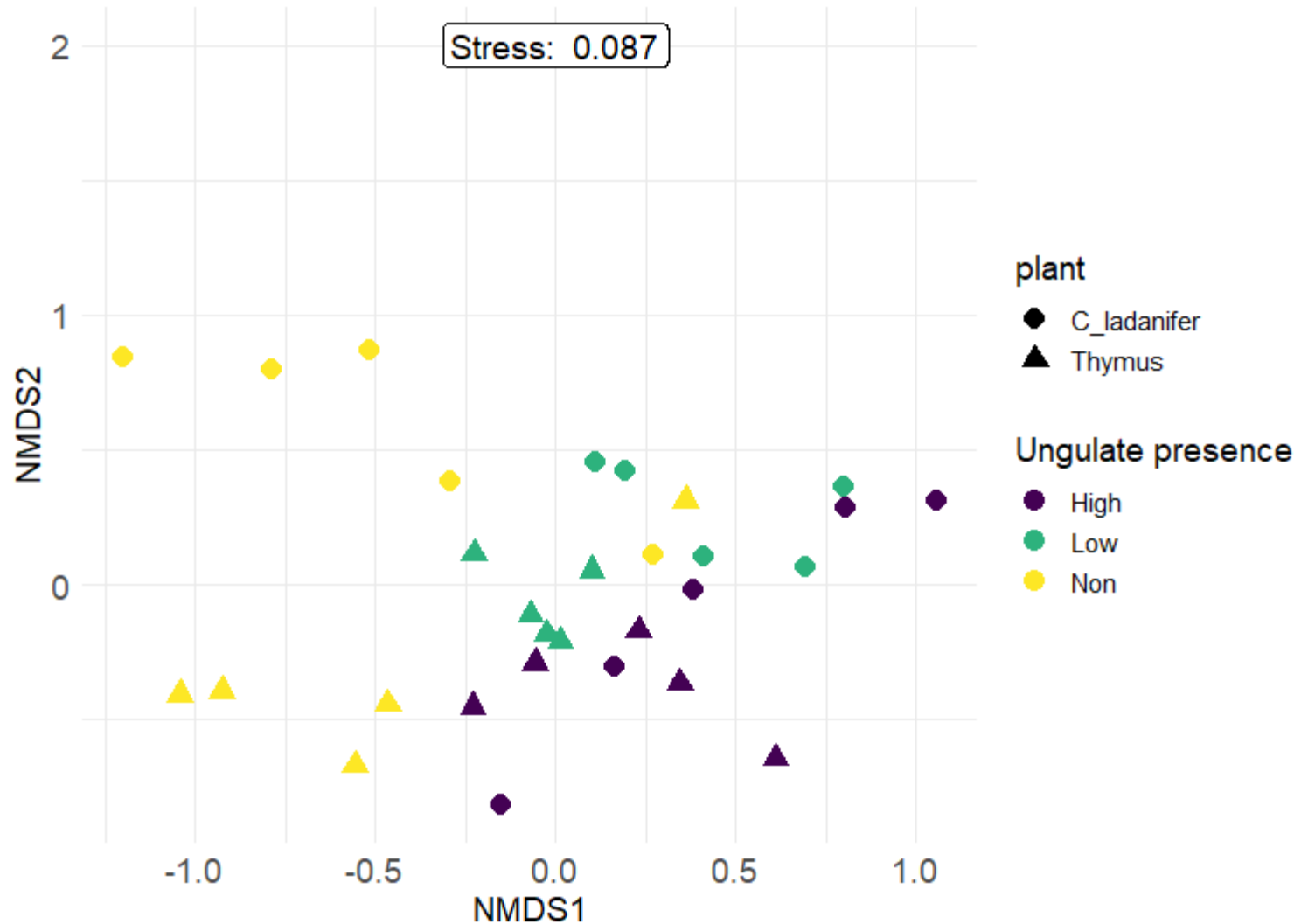
Quintos after 3 month of Ungulate presence

Ungulates density and plant species explanatory value for the fungal communities was fitted into multidimensional scaling ordination (beta diversity was calculated using Bray-Curtis dissimilarity indices) and tested using permutation test using “envfit” function in vegan package.

Goodness of fit:

Ungulate density $R^2=0.3356$ $P= 0.0002$

Plant species $R^2=0.1681$ $P= 0.005$



Muela after 3 month of Ungulate presence



Some samples did not amplify, losing important informationj

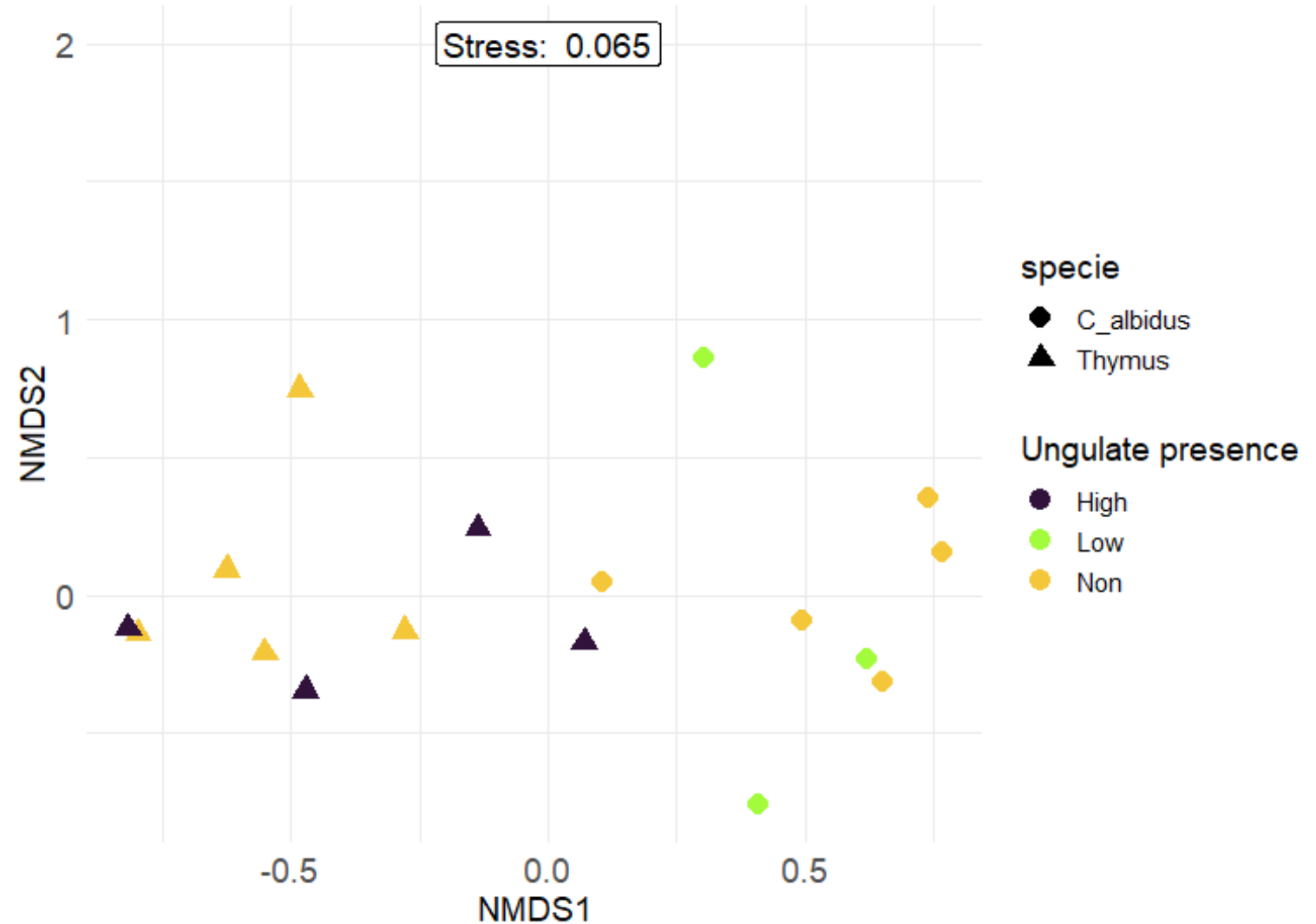
Ungulates density and plant species explanatory value for the fungal communities was fitted into multidimensional scaling ordination (beta diversity was calculated using Bray-Curtis dissimilarity indices) and tested using permutation test using “envfit” function in vegan package.

Goodness of fit:

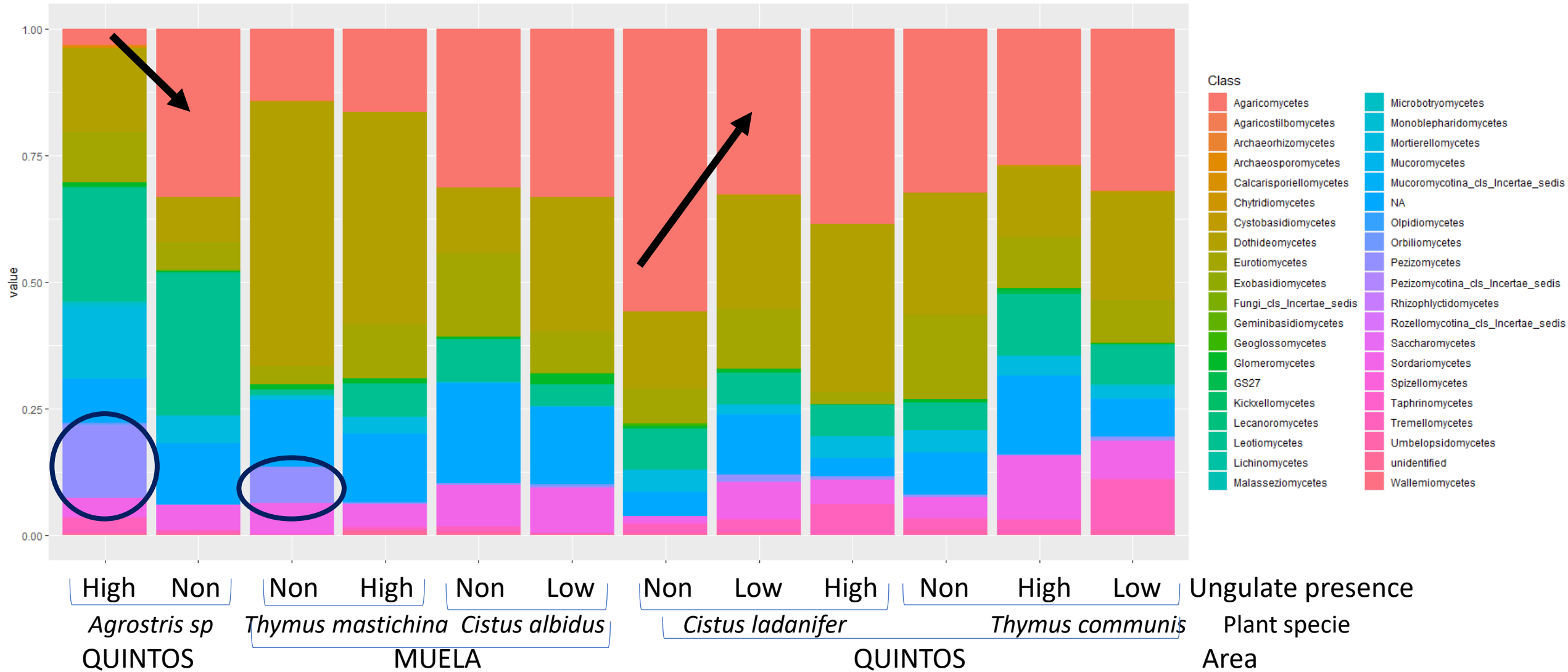
Ungulate density R^2
=0.149 P= 0.3144

Plant species R^2 =0.526 P< 0.0001

Muela after 3 month of Ungulate presence



Fungi class relative abundance per area, plant specie and ungulate density.



Conclusions:

- Fungi community associate to root change with ungulate density.
- This differentiation between fungi community of roots depended on time of exclusion and environmental conditions.
- The differentiation between grazing plots and ungrazing plots, of fungi class composition community associated to roots, is different depending on the species of plant analyzed.
- The observation of the variation in the community of fungi associated with the roots of shrubs that grow in areas with and without the presence of ungulates may be the result of a direct effect of herbivores on the plant, which varies its physiology by altering its microbiome, or it could be an indirect effect resulting from changes in the soil fungal community.

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Thank you



RESERVA VALENCIANA DE CAZA DE MUELA DE CORTES



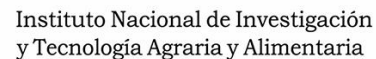
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Project



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The Incremento Consortium

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