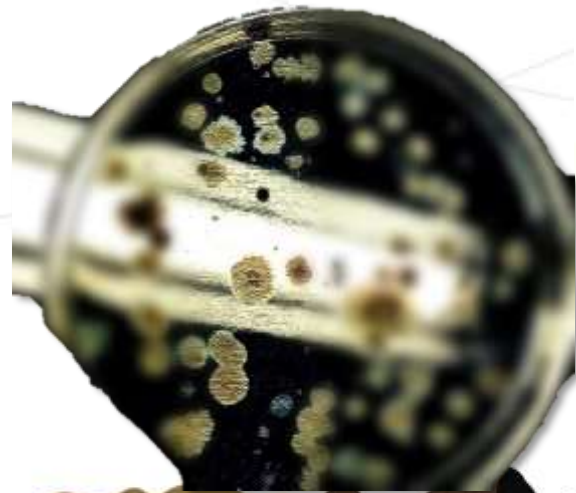




INTEGRATING MICROBIOLOGICAL QUALITY INDICATORS AND SOIL PROPERTIES THROUGH SCORE FUNCTIONS TO ASSESS LAND USES CHANGES IN COLOMBIAN *ANDISOLS*



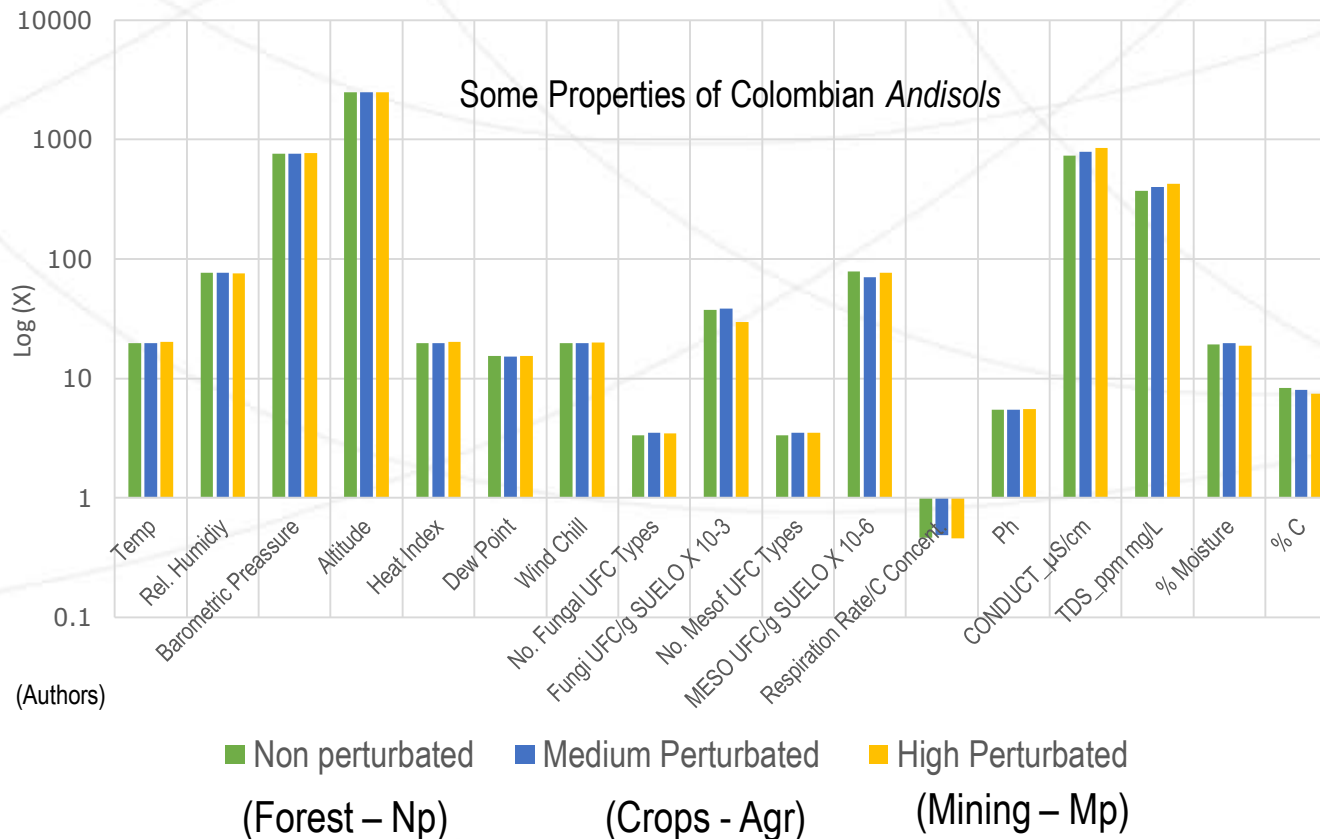
Aranguren R., Cañón J., Florez M., Lumini E.

GLOBAL SYMPOSIUM ON SOIL
BIODIVERSITY | 19-22 April 2021 GSOBIT
2021



PROBLEM STATEMENT

We need to perform **quantitative assessments of Soil Quality (SQ)** to Identify changes in Soil Properties and Functionality (Hinge, 2019)



SQ MEASURES NEED INDICATORS WITH:

INTEGRALITY
(Without loss of information)

REPRESENTABILITY
(**Specificity** - Treats – Soil Functions.)

DIFFERENTIATION

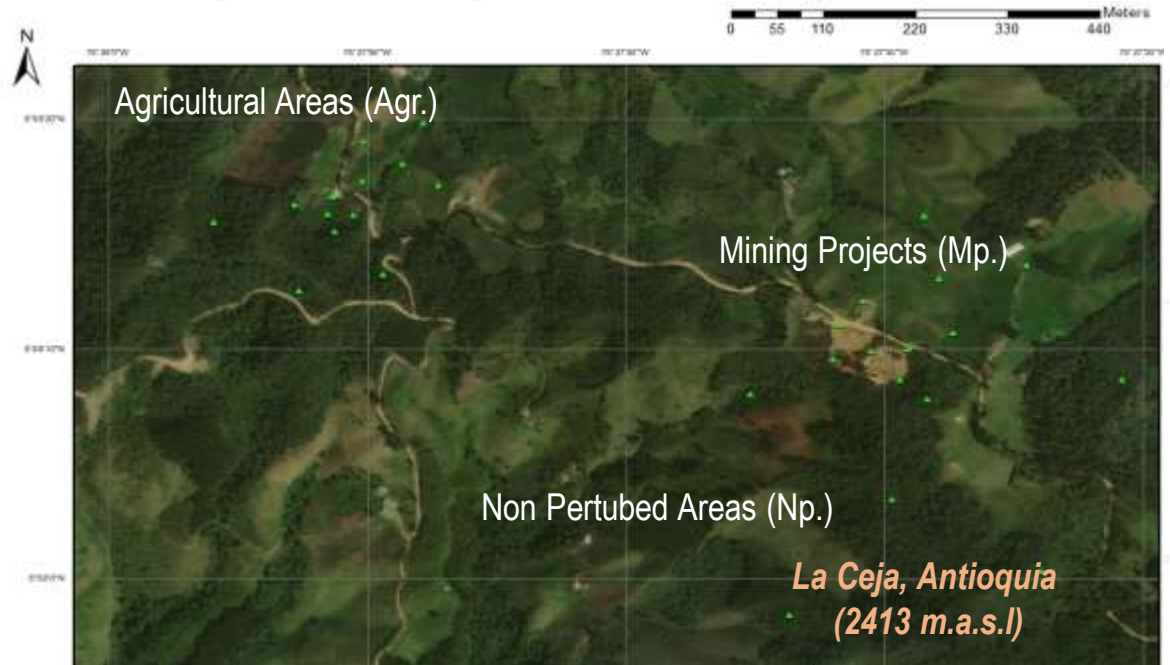
EARLY WARNINGS
(A signal of Impacts)

SIMPLE INTERPRETATION

LOW COST
(Analysis – Approach)



MATERIAL AND METHODS



Antioquia, Colombia

La Ceja , El Retiro, Envigado

Land uses: Agr, Mp., Np.

Period: September – October 2018

Sampled Points: 90

AIM: To build a Minimum Data Set (MDS) to measure SQ in Colombian *Andisols* based on **microbial indicator**, soil properties and environmental variables (with low measurement cost).

Why: 14% National Mining Projects are located in Antioquia.
Andisols of Antioquian Andean Area are the **most treated** soils.
1st producer of non metallic minerals.
(DIASAM, 2019)



MATERIAL AND METHODS 2

Treats by Mp. and Agr. Activities

SOM decline

Salinization

Soil Functions Affected

Habitat provision (Loss of Biodiversity)

OM cycling (C sequestration/Respiration)

Statistical Analysis

ANOVA/LSD test to Identify Differences between land uses.

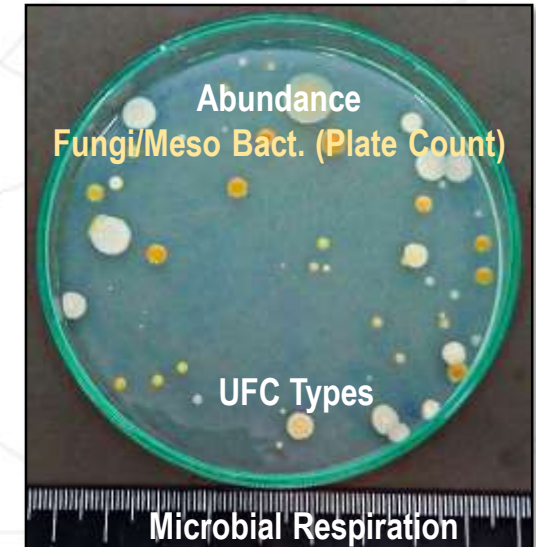
Integration of Variables by Geometric Means (GM).

Create a MDS by Factor Analysis and Correlation Factors.

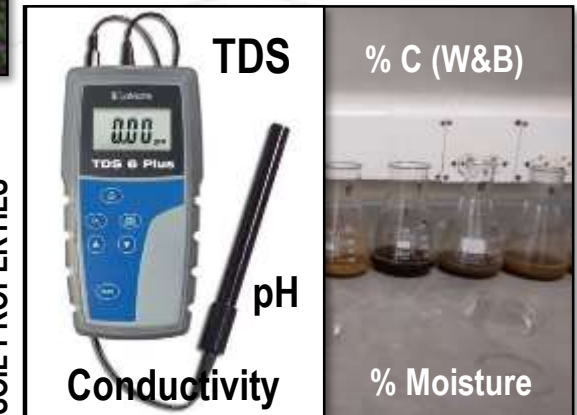
Use Scoring Function to qualify land uses according to MDS (Linear and Non-Lin.).



MICROBIAL INDICATORS



SOIL PROPERTIES



(Raiesi & Kabiri, 2016)

RESULTS AND DISCUSSIONS

1. CAPACITY OF SHOWING IMPACTS/CHANGES IN THE SOIL

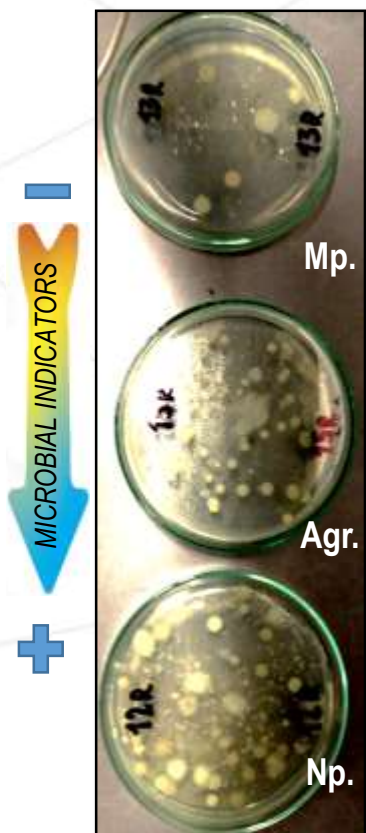
- Dif. Between each land use
- Dif. Between Np. & Perturbed Areas (Agr. & Mp.)
- Dif. Between Np. & Higher Perturbed Areas (Mp.)

Test ANOVA/LSD in a set of soil variables in Colombian Andisols using some GM

VARIABLE	LAND USES								
	Sig.	Agr. - Mp. Dif.	Limit (+/-)	Sig.	Agr. - Np. Dif.	Limit (+/-)	Sig.	Mp. - Np. Dif.	Limit (+/-)
Temperature (°C)	X	-2.2340	1.0556	-	0.4158	1.0556	X	2.6498	1.0556
Dew Point (°C)	X	-0.9960	0.7363	-	-0.6448	0.7363	-	0.3512	0.7363
GM Temps. (T, DP, WCh)	X	-0.0768	0.0371	-	0.0042	0.0371	X	0.0809	0.0371
RHA (%)	X	5.0928	4.0063	X	-5.4925	4.0063	X	-10.5853	4.0063
pH	-	-0.1280	0.1378	-	-0.1000	0.1378	-	0.0280	0.1378
Conductivity (µS/cm)	-	-95.6333	108.5890	-	49.7333	108.5890	X	145.3670	108.5890
TDS (ppm)	-	-33.5111	50.8755	-	36.6889	50.8755	X	70.2000	50.8755
Soil Moisture (%)	X	5.2579	2.2919	-	-1.4789	2.2919	X	-6.7368	2.2919
Carbon content (%)	X	2.5317	1.1309	X	-2.4369	1.1309	X	-4.9686	1.1309
GM (Moist, C)	X	0.8232	0.2896	X	-0.3499	0.2896	X	-1.1731	0.2896
Mesof. Abundance (CFU/g Soil)	X	23.8621	15.7730	-	-1.4498	15.7730	X	-25.3118	15.7730
Fungi Abundance (CFU/g Soil)	X	27.0980	8.2477	X	16.3922	8.2477	X	-10.7059	8.2477
GM Microbial Abundances (Fungi/Mesof. Ln)	X	0.2428	0.0766	X	0.0786	0.0766	X	-0.1642	0.0766
No. FCU Types Mesof.	-	-0.3000	0.4866	X	0.8000	0.4866	X	1.1000	0.4866
No. FCU Types Fungi	-	0.0000	0.4604	X	0.7667	0.4604	X	0.7667	0.4604
GM CFU Types (Fungi/Mesof. Ln)	-	-0.0726	0.1445	X	0.1613	0.1445	X	0.2339	0.1445
MRRat/C	X	-0.9143	0.3003	X	0.3847	0.3003	X	1.2991	0.3003

(Authors)

Fungi Abundance CFU/g Soil x 10⁻³



RESULTS AND DISCUSSIONS

Spearman's Correlation Coefficients

Fertilization/TDS/MI
(Traoré et al. 2007)



	RAH	GM Temp	GM Types Fung./Bact.	GM Abundance Fung./Bact.	MRRat/C	Conductivity	TDS	% Moisture	% C
RAH		0.0001	0.636	0.3061	0.0023	0.04	0.1578	0.0078	0.002
GM Temp	0.0001		0.5594	0.4913	0.0192	0.3443	0.2951	0.0103	0.0076
GM Types Fung./Bact.	0.636	0.5594		0.0124	0.0164	0.8774	0.9039	0.0615	0.0345
GM Abundance Fung./Bact.	0.3061	0.4913	0.0124		0.0065	0.612	0.7507	0.1672	0.0061
MRRat/C	0.0023	0.0192	0.0164	0.0065		0.2874	0.1954	0	0
Conductivity	0.04	0.3443	0.8774	0.612	0.2874		0.0002	0.0895	0.6701
TDS	0.1578	0.2951	0.9039	0.7507	0.1954	0.0002		0.0171	0.6576
% Moisture	0.0078	0.0103	0.0615	0.1672	0	0.0895	0.0171		0.0011
% C	0.002	0.0076	0.0345	0.0061	0	0.6701	0.6576	0.0011	

(Authors)

	Factor 1	Factor 2	Factor 3	Communality	Specific Variance
RAH(%)	0.05143	-0.10029	0.88705	0.79956	0.20043
GM Temp.	-0.05526	0.04772	-0.84359	0.71698	0.28301
GM Types Fung./Bact.	-0.44932	0.03209	0.15258	0.22620	0.77379
GM Abundance Fung./Bact.	0.73042	-0.19654	-0.14059	0.59191	0.40808
MRRat/C	-0.86696	0.10596	-0.21097	0.80735	0.19264
Conductivity(μS/cm)	-0.08549	0.95965	-0.08046	0.93471	0.06528
TDS(mg/L)	-0.11754	0.96944	-0.08060	0.96014	0.03986
%Moisture	0.70384	0.27847	0.30760	0.66756	0.33243
%C	0.80971	-0.16461	0.28574	0.76438	0.23560
Eigenvalue	3.1296	1.8449	1.4943		
Variance (%)	34.773	20.499	16.604		
Cumulative variance (%)	34.773	55.272	71.876		



Vegetal Cover Loss

(Procházka et al. 2011)

FA/CI and Eigenvectors of a set of variables to measure soil functionality in Colombian Andisols

RESULTS AND DISCUSSIONS

SCORING FUNCTION TO MEASURE CHANGES IN SOIL THROUGH MDS VARIABLES

A) $L_{SF}(Y) = x/x_{max}$

B) $NL_{SF}(Y) = \frac{a}{\left(1 + \left(\frac{x}{x_0}\right)^b\right)}$

C) $SQI = \sum_{i=1}^n w_i s_i$

(A) Linear Scoring Function Transformation.

(B) Non linear Scoring Function Transformation.

(C) Additive equation.

(Raiesi & Kabiri, 2016)

*$L_{SF}(Y)$ WAS EFFECTIVE TO SHOW
CHANGES ON OUR MDS VARIABLES*

SQI by Land Uses.
NT, non-transformed variables; (CI) calculated with CI.



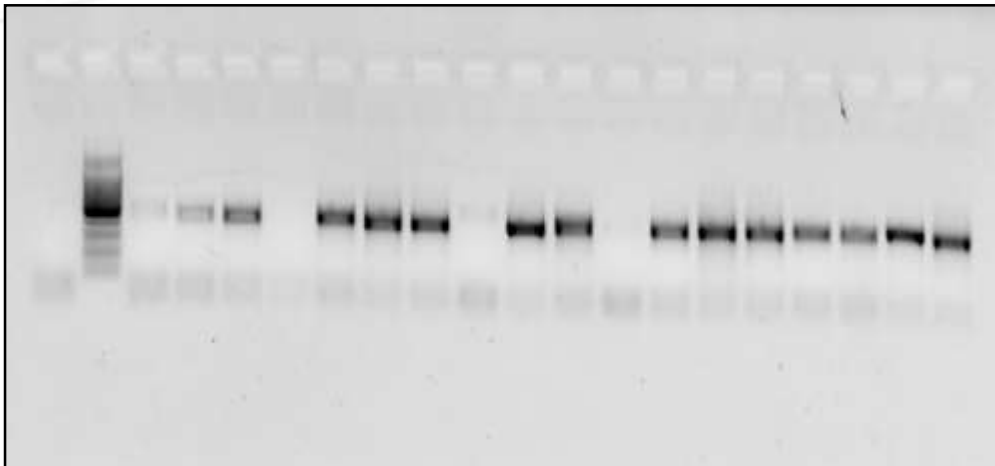
ICs increase sensitivity of the SQI to identify changes in soil functionality

(Hussain et al. 1999).



FINAL REMARKS

- **GEOMETRIC MEANS** are a good alternative for including significant variables of **SOIL FUNCTIONALITY** in an SQI.
- **SQI** calculated with IC and non-linear functions are **more sensitive to disturbances**. This sensitivity allows to discriminate between degrees of impact using variables with **low measurement cost**.
- We suggest using robust **indicators of microbial diversity and functionality** (specially on Fungal communities) in **MDS/SQI**, since these indicators are important to discriminate impacts on soil functionality.
- **Further work:** we will integrate three more factors into the SQI: 1) the structure of Fungal communities using Metabarcoding of Fungal nuclear ribosomal ITS2 region with primers fITS7/ITS4; 2) Glomaline concentration from soil samples; 3) Mycorrhizal Diversity from trap cultures.



Grazie Mille

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Acknowledgments

