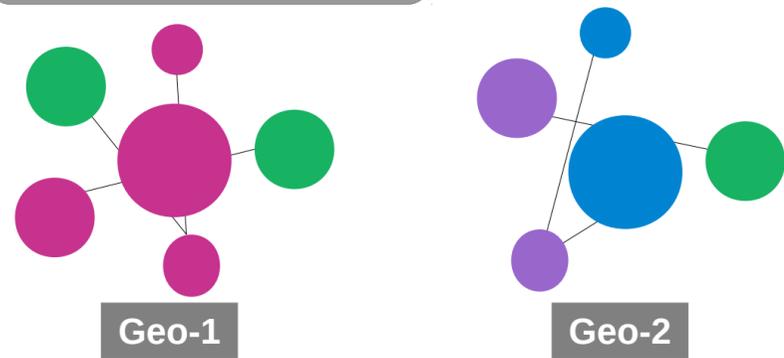
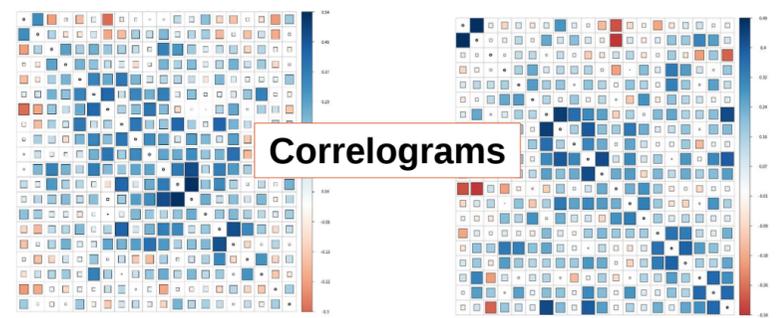


CATEGORICAL NETWORKS

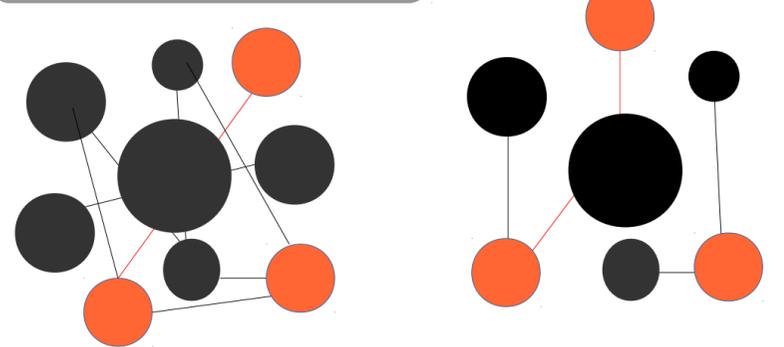


Node metadata and centrality mapped categorical networks



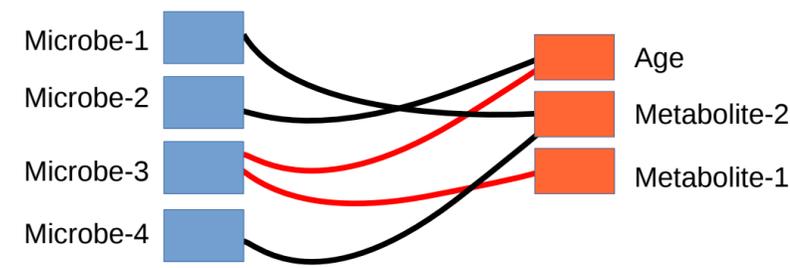
Correlograms

INTEGRATED NETWORKS



COMPLETE

Bi-partite



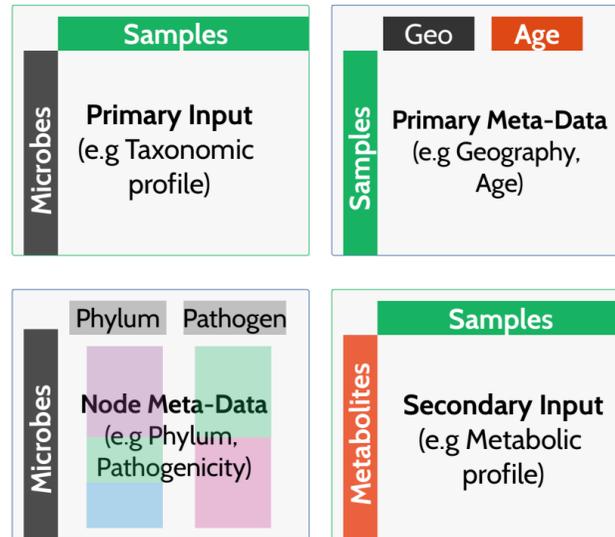
Bi-partite Sankey

Data Filtration

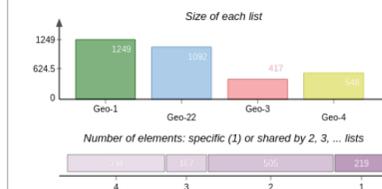
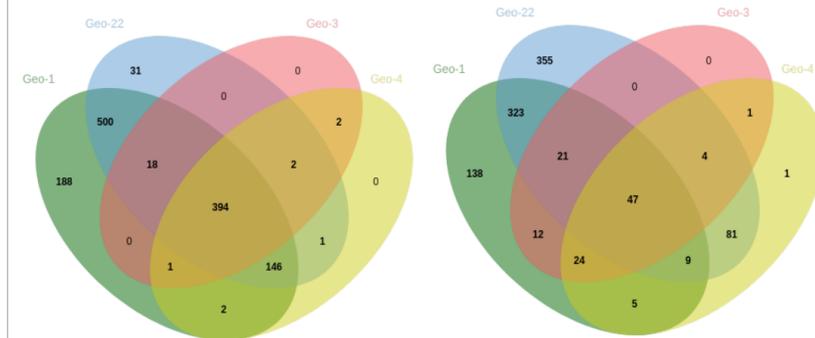
Data Normalization  
Transformation

Network inference  
Algorithms  
(e.g CCREPE, SPARCC)

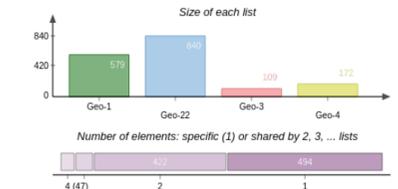
INPUT DATA



COMPOSITIONAL VENNS

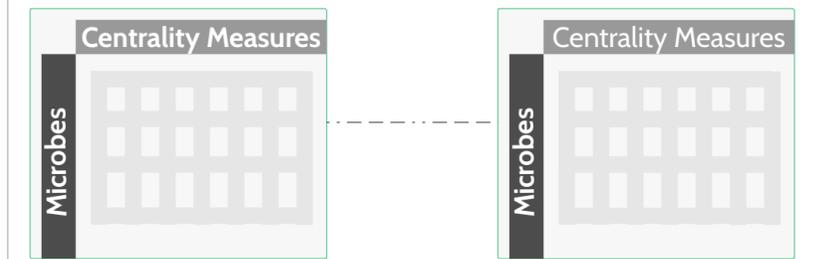


Node composition  
Venn diagram

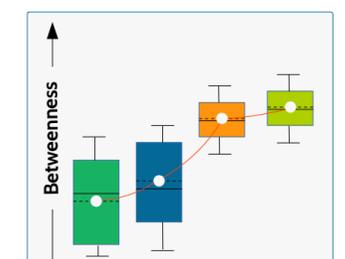
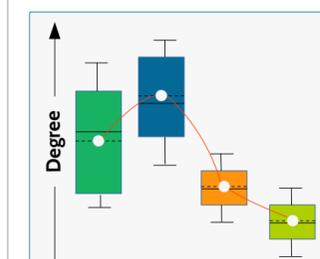


Edge composition  
Venn diagram

NETWORK PROPERTIES



Property tables for all Categories



Grouped box-plots for all properties

MetaN&ts

Inference and meta-insights for microbial correlation networks

## Data Structure and Management

### Input Data

#### Primary Abundance Data

Data	SMP1	SMP2	SMP3	SMP4	SMP5	SMP6
Node1	20	32	12	12	12	12
Node2	32	24	45	45	45	45
Node3	1	4	0	0	0	0
Node4	1	9	0	0	0	0
Node5	34	23	45	45	45	45
Node6	0	0	2	2	2	2
Node7	45	54	34	34	34	34
Node8	78	67	60	60	60	60
Node9	121	232	212	212	212	212

e.g Taxonomic abundance data

#### Sample MetaData

Smps	CatM1	CatM2	ConM1
SMP1	Grp1	GrpA	23
SMP2	Grp1	GrpB	34
SMP3	Grp1	GrpB	21
SMP4	Grp2	GrpA	54
SMP5	Grp2	GrpA	78
SMP6	Grp2	GrpB	17

required

#### Node MetaData

Node	NM1	NM2	NM3
Node1	Cat1	Catx	Cata
Node2	Cat1	Catx	Cata
Node3	Cat1	Catx	Cata
Node4	Cat2	Caty	Catb
Node5	Cat2	Caty	Catb
Node6	Cat2	Catz	Catb
Node7	Cat2	Catz	Catb
Node8	Cat1	Catz	Catb
Node9	Cat1	Catz	Catb

optional

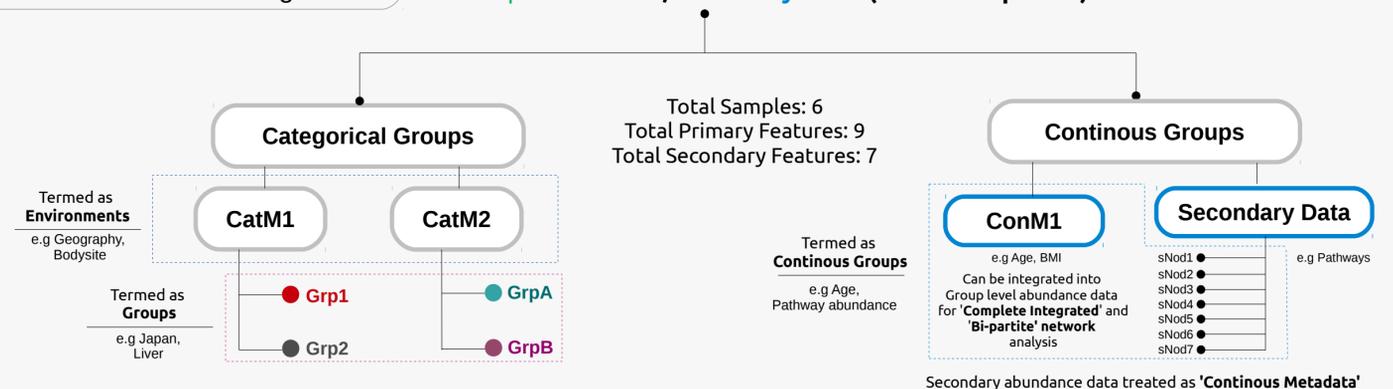
#### Secondary Abundance Data

Data	SMP1	SMP2	SMP3	SMP4	SMP5	SMP6
sNod1	0	2	2	2	1	21
sNod2	3	2	5	5	14	41
sNod3	4	4	0	1	10	10
sNod4	1	0	10	0	01	01
sNod5	3	2	5	4	4	47
sNod6	0	0	2	2	21	21
sNod7	4	4	14	3	3	30

e.g. Pathway abundance data, Gene expression data etc

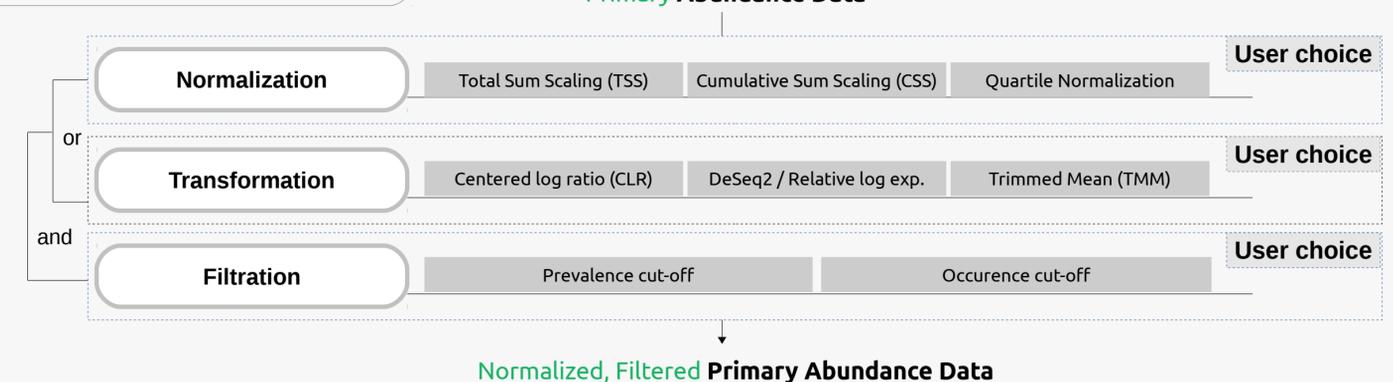
### Meta Data Processing

#### Sample MetaData, Secondary Data (inter-omic profile)



### Primary Data Processing

#### Primary Abundance Data



## Network Methods, Dashboard, Visualizations

### Network Inference Methods

Reboot/CCREPE Sparcc NAMAP Classical (Pearson, Spearman)

### User choice

### Categorical Networks

Comprehensive analysis and visualization of all Categorical networks identified through metadata

### Multi-modular

Multiple analysis And visualization Modules in the personal dashboard

### Customizations

- Layouts
- Property mapping
- Plot Downloads
- Cytoscape and Gephi compatible files

### Compositional Comparisons

Compare networks of various groups within a Category of Metadata (e.g networks of all states in a given Geography), in terms of their Node and Edge composition

## Network Comparison, Properties and Visualizations

Dynamically generated Venn Diagrams for Node as well as Edge composition of various network groups in an environment

### Continuous Metadata

Integrate continous meta-data with primary data for finding correlating taxa against Continous factors

### Searchable inter-omic data

Search and specify inter-omic features of interest for integrating into network inference

### Bi-partite plots

Edges exclusively between the inter-omic data and continous metadata groups can be visualized through highly intuitive Bi-partite plots

### Network Properties

Interactive, searchable and exportable network property tables for each group of an environment

### Network Properties Box Plots

Interactive,downloadable, trend enabled, boxplots (and variants) for each network property, for each group of an environment