Actinomycete BGC ? Metabolomics Mini-Pipeline Report

Run Metadata

- Generated at: 2025-10-03T19:00:27.095345 UTC

Methods

Methods Documentation / ????

> ???????????????????????????????????

> English: This document will be referenced in the final report, detailing data sources, processing steps, scoring rules, and limitations.

Data Sources / ????

- TODO: antiSMASH / DeepBGC / PRISM ???????

- TODO: LC-MS/MS ???????

- TODO: NPAtlas/MIBiG ?????

Processing Steps / ????

- TODO: BGC ??????

- TODO: LC-MS/MS ????

- TODO: ????????

- TODO: ????????

- TODO: ????????

- TODO: ADMET ????????

Parameters & Versions / ?????

- TODO: ??????????????????

Limitations & Future Work / ???????

- TODO: ?????????????????

Top Candidates

Top Candidates

- **Rank 1** ? Compound CMP001 | Score: 0.760 | ADMET: Pass | Cluster: CLUSTER_001
- **Rank 2** ? Compound CMP002 | Score: 0.760 | ADMET: Pass | Cluster: CLUSTER_002
- **Rank 3** ? Compound CMP003 | Score: 0.760 | ADMET: Pass | Cluster: CLUSTER_003
- **Rank 4** ? Compound <NA> | Score: 0.400 | ADMET: Pass | Cluster: nan

Figures

- cluster_sizes.csv
- cluster_summary.txt
- top_scores.png

NaN

1.0

0.0

Ranking Snapshot

CompoundID EvidenceScore EvidenceCo	ount BGCUIDs
FeatureIDs logP TPSA HBD HBA MV	W ToxicityFlag RuleOfFivePass ClusterID ClusterSize
Novelty ADMETScore AggregateScore R	Rank EvidenceSummary
CMP001 0.6 1	SampleA_BGCUID_001
F001 F002 F003 1.00 48.0 4.0 8.0 166.0	Low True CLUSTER_001 1.0 1.0
1.0 0.76 1 1 evidence links	
CMP002 0.6 2	SampleA_BGCUID_001 SampleA_BGCUID_002
F001 F002 F003 1.25 36.0 3.0 6.0 1	90.0 Low True CLUSTER_002 1.0
1.0 1.0 0.76 2 2 evidence lin	ks
CMP003 0.6 1	SampleA_BGCUID_001
F001 F002 F003 1.25 60.0 5.0 10.0 206.0	0 Low True CLUSTER_003 1.0 1.0
1.0 0.76 3 1 evidence links	
NaN	0.5
SampleA_BGCUID_001 SampleA_BGCUID	D_002 SampleB_BGCUID_001

4 9 evidence links

NaN

NaN

NaN

F001|F002|F003|F004|F005|F006 NaN NaN NaN NaN NaN

0.40