

# 3-Day Workshop cum Faculty Development Program on ‘Understanding Health and Wellness Through Microbiome Research’

## Day3:

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## Lecture1: (Databases)

### Reference Databases:

#### 1) 16S and other amplicon data

- a) Ribosomal Database Project (RDP)

Weblink: <http://rdp.cme.msu.edu/>

Downloading link: <http://rdp.cme.msu.edu/misc/resources.jsp>

- b) Greengenes Database

Weblink: <https://greengenes.secondgenome.com/>

- c) SILVA Database

Weblink: <https://www.arb-silva.de/>

Downloading link: <https://www.arb-silva.de/documentation/release-138/>

- d) UNITE Database

Weblink: <https://unite.ut.ee/#main>

Downloading link: <https://unite.ut.ee/repository.php>

#### 2) Whole Genome Sequences

- a) NCBI Genome: (RefSeq and GeneBank)

Weblink: <https://www.ncbi.nlm.nih.gov/datasets/genome/>

- b) NCBI-nr

Weblink: <https://www.ncbi.nlm.nih.gov/refseq/about/nonredundantproteins/>

- c) NCBI-nt

Weblink: <https://www.ncbi.nlm.nih.gov/nucleotide/>

- d) JGI (Joint Genome Institute)

Weblink: <https://data.jgi.doe.gov/>

- e) Genome Taxonomy Database (GTDB)  
Weblink: <https://gtdb.ecogenomic.org/>
- f) Metaphlan/Chocophlan
  - Clade-Specific Marker Gene Database  
Database Link:  
[http://cmprod1.cibio.unitn.it/biobakery4/metaphlan\\_databases/mpa\\_vJan21\\_CHOCOPhlanSGB\\_202103\\_marker\\_info.txt.bz2](http://cmprod1.cibio.unitn.it/biobakery4/metaphlan_databases/mpa_vJan21_CHOCOPhlanSGB_202103_marker_info.txt.bz2)  
GITHUB: <https://github.com/biobakery/MetaPhlAn/wiki/MetaPhlAn-4>
  - ChocoPhlAn Database  
Paper: <https://elifesciences.org/articles/65088>  
Github: <https://github.com/biobakery/MetaPhlAn/wiki/HUMAN-N-Workshop-on-Genomics-2023>

### 3) Functional Databases

- a) MetaCyc:  
<https://metacyc.org/>
- b) KEGG:  
<https://www.kegg.jp/>
- c) COG:  
<https://www.ncbi.nlm.nih.gov/research/cog-project/>
- d) Pfam:  
<http://pfam.xfam.org/>
- e) CAZy:  
<http://www.cazy.org/>
- f) CARD:  
<https://ardb.cbc.umd.edu/>  
<https://card.mcmaster.ca/>
- g) UNIPROT:  
<https://www.uniprot.org/>

## Microbiome Databases:

### 1) General Sequence Databases:

- a) ENA (European Nucleotide Archive)  
<https://www.ebi.ac.uk/ena/browser/home>
- b) NCBI (National Center for Biotechnology Information)

<https://www.ncbi.nlm.nih.gov/>

- c) DDBJ (DNA Data Bank of Japan)  
<https://www.ddbj.nig.ac.jp/index-e.html>
- d) China National GeneBank Database  
<https://db.cngb.org/>

## 2) Customized Microbiome Databases

- a) curatedMetagenomicData:  
Weblink: <https://waldronlab.io/curatedMetagenomicData/>  
Paper: <https://www.nature.com/articles/nmeth.4468>  
Github: <https://github.com/waldronlab/curatedMetagenomicData>
- b) MicrobiomeDB:  
Weblink: <https://microbiomedb.org/mbio/app>  
Paper: <https://academic.oup.com/nar/article/46/D1/D684/4584629?login=false>
- c) MicrobiomeHD:  
Repo-link: <https://zenodo.org/records/569601>  
Paper: <https://www.nature.com/articles/s41467-017-01973-8>
- d) QIITA:  
Weblink: <https://qiita.ucsd.edu/>  
Documentation: <https://qiita.ucsd.edu/static/doc/html/index.html>
- e) GMRepo:  
Paper: <https://pubmed.ncbi.nlm.nih.gov/34788838/>  
Weblink: <https://gmrepo.humangut.info/home>
- f) Human Microbiome Compendium:  
Weblink: <https://microbiomap.org/>  
R Package to download: <https://blekmanlab.github.io/MicroBioMap/>  
Paper: [https://www.cell.com/cell/fulltext/S0092-8674\(24\)01430-2?\\_returnURL=https%3A%2F%2Flinkinghub.elsevier.com%2Fretrieve%2Fpii%2FS0092867424014302%3Fshowall%3Dtrue](https://www.cell.com/cell/fulltext/S0092-8674(24)01430-2?_returnURL=https%3A%2F%2Flinkinghub.elsevier.com%2Fretrieve%2Fpii%2FS0092867424014302%3Fshowall%3Dtrue)
- g) HACK Profiles Repository (Our Own Creation):  
Paper: [https://www.cell.com/cell-reports/fulltext/S2211-1247\(25\)00149-4](https://www.cell.com/cell-reports/fulltext/S2211-1247(25)00149-4)  
Github: [https://github.com/tsg-microbiome/HACK\\_index\\_2024](https://github.com/tsg-microbiome/HACK_index_2024)

## 3) Metagenome Assembled Genomes

- a) UNINA MAGs Database:  
Weblink: [http://segatalab.cibio.unitn.it/data/Pasolli\\_et\\_al.html](http://segatalab.cibio.unitn.it/data/Pasolli_et_al.html)  
Paper: [https://www.cell.com/cell/fulltext/S0092-8674\(19\)30001-7?\\_returnURL=https%3A%2F%2Flinkinghub.elsevier.com%2Fretrieve%2Fpii%2FS0092867419300017%3Fshowall%3Dtrue](https://www.cell.com/cell/fulltext/S0092-8674(19)30001-7?_returnURL=https%3A%2F%2Flinkinghub.elsevier.com%2Fretrieve%2Fpii%2FS0092867419300017%3Fshowall%3Dtrue)

b) UHGG/UHGP (MGnify)

Paper: <https://www.nature.com/articles/s41587-020-0603-3>

Weblink: [http://ftp.ebi.ac.uk/pub/databases/metagenomics/mgnify\\_genomes/](http://ftp.ebi.ac.uk/pub/databases/metagenomics/mgnify_genomes/)

## Links for Sample Collection:

### For Fecal sample:

1) IHMS (International Human Microbiome Standards:

<https://human-microbiome.org/index.php#SOPS>

### For Oral sample:

1) Molecular Medicine Ireland:

<https://brd.nci.nih.gov/brd/sop/download-pdf/147>

2) CD-Genomics:

<https://www.cd-genomics.com/microbioseq/saliva-sample-collection-and-handling-instructions.html>

## Link for video for uploading samples to ENA:

<https://www.youtube.com/watch?v=n2kUjY25dPY>

(start from 15:30 mins)