
Supplementary information

Perspectives on ENCODE

In the format provided by the authors and unedited

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- 111) Center for Molecular Medicine and Department of Biochemistry and Molecular Biology, University of Georgia, Athens, GA 30602, USA
- 112) Gift of Life Donor Program, 401 North 3rd Street, Philadelphia, PA 19123
- 113) American Society for Radiation Oncology, 251 18th Street South, 8th Floor, Arlington, VA 22202
- 114) National Cancer Institute, National Institutes of Health, 9609 Medical Center Drive, Bethesda, MD 20892
- 115) Leidos Biomedical, Inc., 8560 Progress Drive, Frederick, MD 21701
- 116) National Disease Research Interchange (NDRI), 8 Penn Center, 15th Floor, 1628 JFK Boulevard, Philadelphia, PA 19103
- 117) 4407 Puller Dr, Kensington, MD 20895

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Supplementary Note 1 Useful URLs

ENCODE Portal

- a) ENCODE Portal: <https://www.encodeproject.org>

Using ENCODE Data

- b) ENCODE Data use policy: <https://www.encodeproject.org/about/data-use-policy/>
- c) ENCODE Tutorials and handouts: <https://www.encodeproject.org/tutorials/>

Accessing ENCODE data

- d) ENCODE encyclopedia: <https://www.encodeproject.org/data/annotations/>
- e) ENCODE SCREEN: <http://screen.encodeproject.org>
- f) ENCODE Registry of candidate regulatory elements:
https://www.encodeproject.org/matrix/?type=Annotation&annotation_type=candidate+regulatory+elements&files.file_type=bed+bed3%2B
- g) Factorbook of transcription factors assayed by ENCODE: <http://factorbook.org>
- h) REST API: <https://www.encodeproject.org/help/rest-api/>
- i) All ENCODE data:
<https://www.encodeproject.org/matrix/?type=Experiment&award.project=ENCODE>
 - 1. all ENCODE mouse data:
https://www.encodeproject.org/matrix/?type=Experiment&award.project=ENCODE&replicates.library.biosample.donor.organism.scientific_name=Mus+musculus
 - 2. all ENCODE human data:
https://www.encodeproject.org/matrix/?type=Experiment&award.project=ENCODE&replicates.library.biosample.donor.organism.scientific_name=Homo+sapiens
 - 3. all modENCODE/modERN data:
<https://www.encodeproject.org/matrix/?type=Experiment&award.project=modENCODE&award.project=modERN>
 - i. all worm data:
https://www.encodeproject.org/matrix/?type=Experiment&award.project=modENCODE&award.project=modERN&replicates.library.biosample.donor.organism.scientific_name=Caenorhabditis+elegans
 - ii. all fly data:
<https://www.encodeproject.org/matrix/?type=Experiment&award.project=modENCODE&award.project=modERN>
- 4. ENCODE EN-TEEx samples:
https://www.encodeproject.org/matrix/?searchTerm=gtex&type=Experiment&status=released&replicates.library.biosample.biosample_type=tissue&y.limit=
- 5. ENCODE 3 mouse matrix samples:
https://www.encodeproject.org/matrix/?searchTerm=ENCODE+3&type=Experiment&award.rfa=ENCODE+3&replicates.library.biosample.donor.organism.scientific_name=Mus+musculus&biosample_type=tissue&organ_slims=brain&organ_slims=heart&organ_slims=liver&organ_slims=limb&organ_slims=epithelium&organ_slims=lung&organ_slims=stomach&organ_slims=kidney&organ_slims=intestine&organ_slims=embryo

Examples of how ENCODE data are used

- j) Community and ENCODE consortium publications: <https://www.encodeproject.org/publications/>

How ENCODE data are generated and processed

- k) ENCODE Experimental guidelines/protocols and data standards/quality metrics: <https://www.encodeproject.org/data-standards/>
- l) ENCODE data processing pipelines: <https://www.encodeproject.org/pipelines/>
- m) Software tools: <https://www.encodeproject.org/software/>
- n) Ontologies used by ENCODE: <https://www.encodeproject.org/help/getting-started/#Ontologies>
 1. experimental assay types (OBI: http://obi-ontology.org/page/Main_Page)
 2. cell lines (EFO: <http://www.ebi.ac.uk/efo/>)
 3. primary cells (CL: <http://cellontology.org/>)
 4. tissues (UBERON: <http://uberon.org/>)
- o) Antibody characterization: <https://www.encodeproject.org/search/?type=AntibodyLot>
- p) ENCODE Help Desk: encode-help@lists.stanford.edu

ENCODE Participants

- q) ENCODE 3 participants: <https://www.genome.gov/27568353/encode-phase-iii-participants-and-projects/>
- r) ENCODE 4 participants: <https://www.genome.gov/26525220/encode-participants-and-projects/>

ENCODE is a member of these organizations

- s) International Human Epigenome Consortium (IHEC): <http://ihec-epigenomes.org/welcome/>
- t) Global Alliance for Genomics and Health (GA4GH): <https://www.ga4gh.org>

ENCODE Tissues and Cell Lines: Current and Planned

- u) <https://www.encodeproject.org/proposed-biosamples/>

Related Projects

- v) NIH Roadmap Epigenomics Program (REMC): <https://commonfund.nih.gov/epigenomics/index>
- w) International Human Epigenome Consortium (IHEC): <http://ihec-epigenomes.org>
- x) Canadian Epigenetics, Environment and Health Research Consortium (CEEHRC) Network: <http://www.epigenomes.ca>
- y) Blueprint: <http://www.blueprint-epigenome.eu>
- z) 4D Nucleome Program (4DN): <https://commonfund.nih.gov/4Dnucleome/index>
- aa) The Cancer Genome Atlas (TCGA): <https://cancergenome.nih.gov>
- bb) Genotype and Tissue Expression Project (GTEx): <https://commonfund.nih.gov/GTEx/index>
- cc) PsychENCODE: https://www.nimhgenetics.org/available_data/psychencode/
- dd) Functional Annotation of the Mammalian Genomes (FANTOM) <https://fantom.gsc.riken.jp/>
- ee) The Human Cell Atlas (HCS) <https://www.broadinstitute.org/research-highlights-human-cell-atlas>
- ff) The Human Biomolecular Atlas Program (HuBMAP) <https://commonfund.nih.gov/hubmap>
- gg) Functional Annotation of Animal Genomes (FAANG): <http://www.faang.org>