National Library of Medicine

Data driven cure:
How the NLM inspires
HIV discoveries through
basic science

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Provided by: Nicole Scott (nicole.scott@nih.gov)

HIV/AIDS Literature



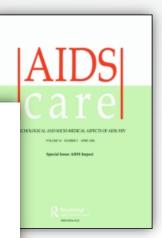
JOINT COLLECTION DEVELOPMENT POLICY STATEMENT

OF

THE NATIONAL LIBRARY OF MEDICINE

And

THE LIBRARY OF CONGRESS

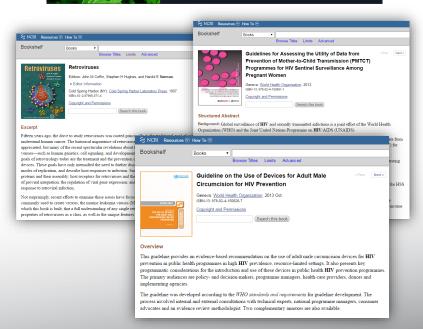




- Free full-text archive of biomedical and life sciences literature
- Includes full text of >300,000 HIV/AIDS articles cited in PubMed

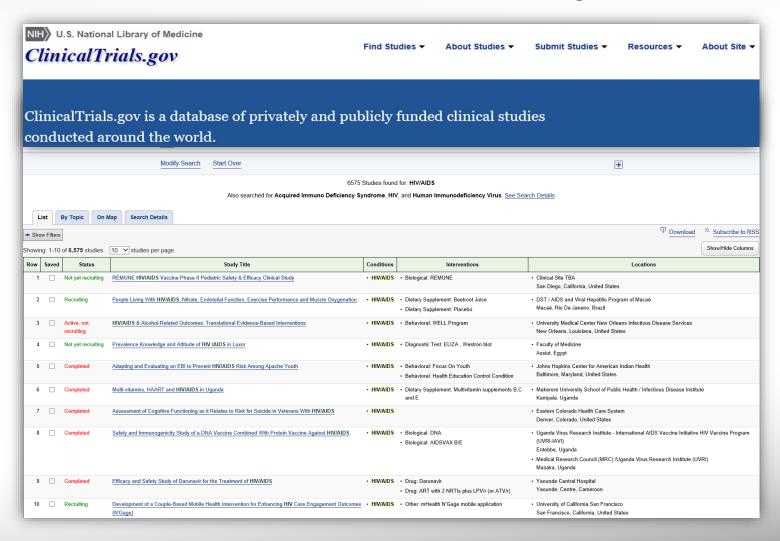






ClinicalTrials.gov

Information on > 6,500 HIV/AIDS studies from >1,000 organizations

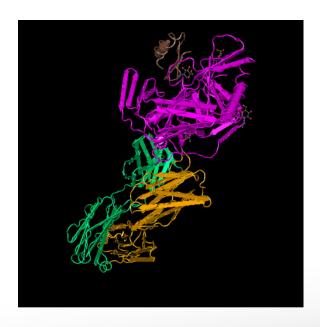


NCBI's Data & Computational Resources

HIV Protein Structures

NCBI's collection of HIV protein structures numbers almost 1,600

Crystal structure of HIV Gp120 protein with bound antibody





NCBI HIV-1: human interaction database

2,500 manually-curated interactions

14,000+ links to PubMed records

Vpu inhibits NFKB1

Protein: NP_003989.2, GeneRIF: Most HIV-1 and primate lentiviral Vpu proteins inhibit IKK-beta-induced NF-kappaB activation except HIV-1 group N Vpu proteins . Pubmed ID: 25620704

See: https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4383939/



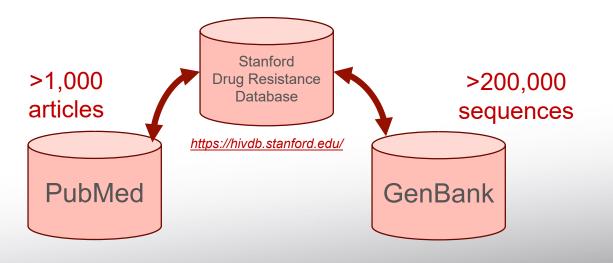
Surveillance of HIV mutations in individuals

HIV has a high mutation rate

Sequence is needed for evaluation & comparison

Combination antiretroviral therapies effective against resistance

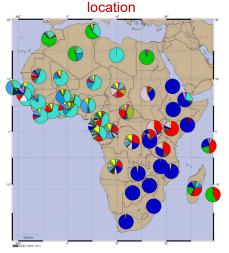
Patient virus population informs treatment strategies





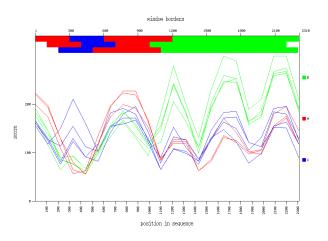
Population detection/surveillance

Los Alamos National Laboratory HIV sequences based on geographical



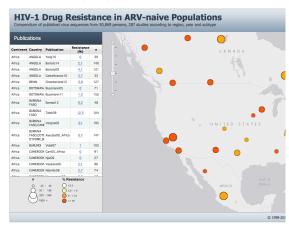
Derived in part from >3,500 GenBank records

NCBI Virus Genotyping Tool



Uses BLAST to identify the genotype (or subtype) of HIV-1 sequence

Stanford University HIV drug resistance based on geographical location

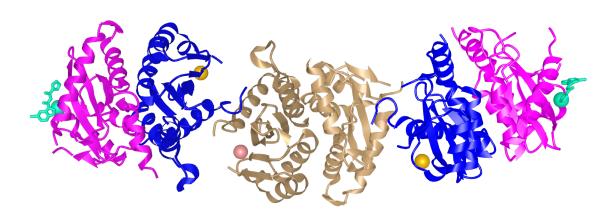


Primary data mostly found in GenBank





Vaccine and targeted drug discovery



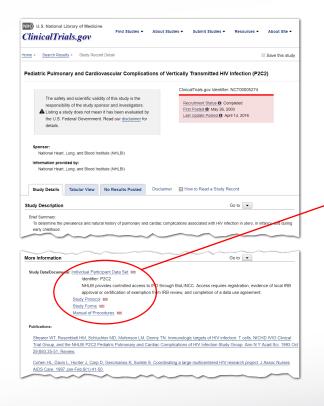
PDB accession code: 1QS4. From: "In-Silico docking of HIV-1 integrase inhibitors reveals a novel drug type acting on an enzyme/DNA reaction intermediate" Retrovirology vol. 4 21. 20 Mar. 2007, doi:10.1186/1742-4690-4-21

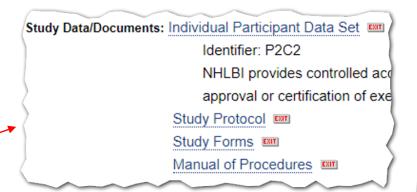


Lister Hill Center Computational Clincal Support

Identification of Research Common Data Elements in HIV/AIDS Using Data Science Methods

ClinicalTrials.gov example of data sharing (HIV trial)





Automatic Detection of Pulmonary Disease in Chest X-rays to Combat HIV/AIDS

Workflow Schematic

