

Defining the Chemical Space of Public Genomic Data

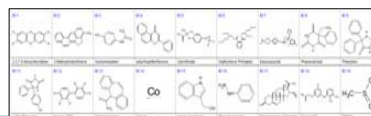
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TOXICOGENOMICS

Project Goals

- Chemically annotate the National Center for Biotechnology Information's (NCBI's) Gene Expression Omnibus (GEO)⁵, the European Bioinformatics Institute's (EBI's) ArrayExpress¹, Environment, Drug, Gene Expression (EDGE)⁶ and the National Institute of Environmental Health Sciences CEBS³ genomic data repositories.
- Create a Structure Index of the three data repositories that can integrate these data with other public data sources (e.g., PubChem⁸).
- Populate Structure Index files with historical toxicological and chemical data from DSSTox Data Sources through cross referencing.
- Explore methodologies to address concerns of public genomic data use, including: across laboratory and platform comparisons, as well as extrapolation between species, doses, chemicals, and endpoints.
- Restructure Genomic Data for insertion into a Chemogenomic Database.
- Use Chemogenomic Database in conjunction with both developed and adopted methodologies, as well as data mining tools, to guide future experiments, discern new patterns of toxicological interest, and explore new hypotheses related to toxicological potential.

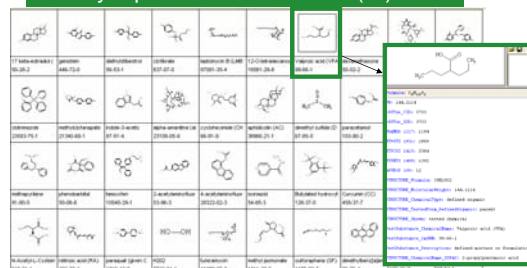
EDGE Database



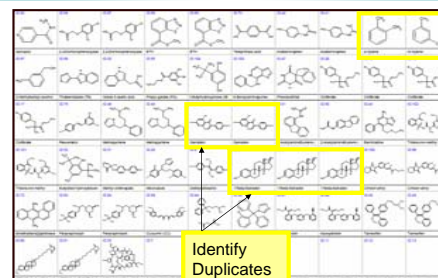
Statistics

- 25 Records
- 18 Chemical Structures
- 8 Chemical Structures overlap GEOMIS (GEO)
- 7 Chemical Structures overlap AREXCH (ArrayExpress)
- 6 Chemical Structures overlap CSTARC (CEBS)
- Chemical Structures overlap 82 Toxicological Records

ArrayExpress Structure Index (SI) File



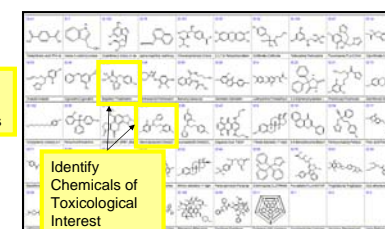
ArrayExpress Database



Statistics

- Initially 129 Chemical Exposure Records
- Currently 351 Chemical Exposure Records
- 129 Chemical Structures overlap 273 Toxicological Records

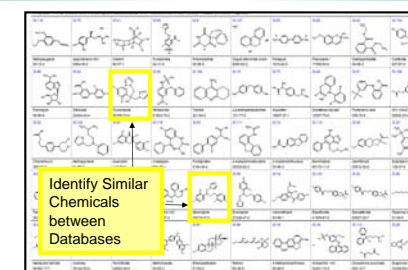
GEO Database



Statistics

- Initially 129 Chemical Exposure Records
- Currently Over 500 Chemical Exposure Records
- 129 Chemical Structures overlap 246 Toxicological Records

CEBS Database



Statistics

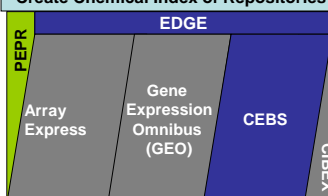
- Initially 135 Chemical Exposure Records
- Currently All Records contain Toxicology Information
- Chemical Structures overlap 538 Toxicological Records

Methods/Approaches

Identification of Genomic Repositories and Databases

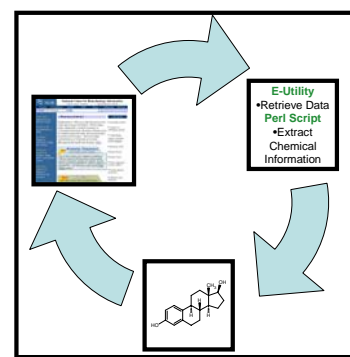
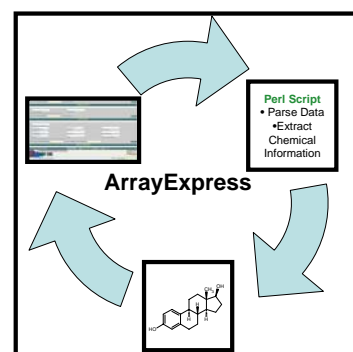
Identification of Genomic Repositories and Databases of Possible Toxicogenomic Interest

Create Chemical Index of Repositories



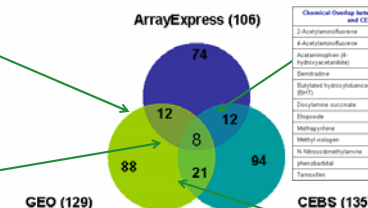
Legend:

- Indexed Databases
- MGED Databases
- Small Database



Chemical Space

Overlap between Genomic Repository Chemical Space

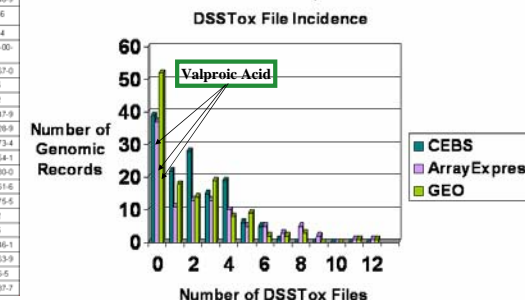


Chemical Overlap between ArrayExpress and GEO	
17beta-Estradiol	50-28-2
Genistein	446-72-0
Dexamethasone	50-02-2
Cyclohexanone	66-81-9
Salicylic Acid	30-79-4
Hydrogen peroxide	7722-84-1
Terephthalic acid (TPA)	100-21-0
Fenpropimorph	67306-03-0
5-Fluorouracil	51-21-8
Lithyminine	6890-02-3
Mitomycin C	50-07-7
Paclitaxel	33069-62-4

Chemical Overlap between ArrayExpress, GEO, and CEBS	
cis-Octachlorodioxane platinum	15663-27-1
Clotriazole	637-07-0
Clotrimazole	23693-75-1
Ethanol	64-17-5
Ethinyl estradiol	57-63-6
Lipopolysaccharide (LPS)	NOCAS
Muscadine	77016-47-0
Valproic acid	99-66-1

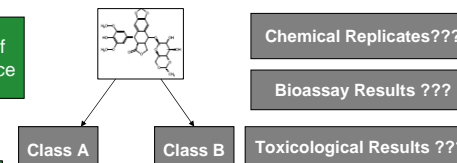
Chemical Overlap between GEO and CEBS	
1,3-Bis(phenylethyl)-1H-imidazole	154-93-8
Adamsyn, hydrochloride	25316-40-9
Allyl alcohol	107-18-6
alpha-Naphthyl isothiocyanate	551-26-4
Alorvastatin	13623-00-5
Benzaldehyde	41859-67-0
Carbon tetrachloride	56-23-5
Dimethylformamide	68-12-0
Econazole	27220-47-9
Fenflurine	49602-28-9
Fluorazone	86366-73-4
Fluvestron	93967-54-1
Gendrol	25912-30-0
Isosaccharin	84825-61-6
Levonorgestrel	75330-75-5
Methotrexate	59-05-2
N-Nitrosodimethylamine	55-18-5
Rifampin	13292-46-1
SARVASTATIN	79902-43-9
Sodium arsenite	7704-48-5
Troglitazone	97322-80-7

Overlap between Genomic Chemical Space and DSSTox Chemical Space

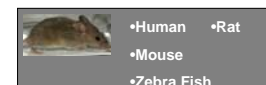


Future Analysis

1. Analysis of Chemical Space



2. Coverage of Species



3. Coverage of Platforms

References

- ArrayExpress <http://www.ebi.ac.uk/ArrayExpress>
- ArrayTrack <http://www.fda.gov/nctr/science/centers/toxicoinformatics/ArrayTrack>
- CEBS <http://cebs.niehs.nih.gov>
- Comparative Toxicogenomics Database (CTD) <http://www.ctd.mdibl.org>, Mattingly et al, **The Comparative Toxicogenomics Database (CTD): A Cross-Species Resource for Building Chemical-Gene Interaction Networks**, ToxSci. 2006, in press
- Gene Expression Omnibus (GEO) <http://www.ncbi.nlm.nih.gov/geo>
- Environment, Drug, and Gene Expression (EDGE) <http://edge.oncology.wisc.edu/>
- Iconix B. Ganter et al, **Development of a large-scale chemogenomics database to improve drug candidate selection and to understand mechanisms of chemical toxicity and action**, Journal of Biotechnology (2005) 119: 219-44.
- PubChem <http://pubchem.ncbi.nlm.nih.gov>

This work was reviewed by U.S. EPA and approved for publication but does not necessarily reflect official Agency policy.