

#### ос Источники данных

ChEMBL is a large, open-access drug discovery database that aims to capture Medicinal Chemistry data and knowledge across the pharmaceutical research and development process

https://academic.oup.com/nar/article/47/D1/D930/5162468

#### ос Источники данных

- JMedChemComm
- Journal of Medicinal Chemistry
- ACS Medicinal Chemistry Letters
- European Journal of Medicinal Chemistry
- Bioorganic & Medicinal Chemistry
- Bioorganic & Medicinal Chemistry Letters
- Journal of Natural Products

For these journals, every article in each new issue is screened for the presence of quantitative small molecule (or peptide) bioactivity data.

CC ИСТОЧНИКИ ДАННЫХ  Top 15 journals covered by ChEMBL (release 24), according to numbers of articles extracted					
Journal	Number of documents				
Bioorg. Med. Chem. Lett	. 21 197				
J. Med. Chem.	21 032				
Bioorg, Med. Chem.	6996				
J. Nat. Prod.	6701				
Eur. J. Med. Chem.	5514				
Antimicrob. Agents Cher	mother. 2121				
ACS Med. Chem. Lett.	1378				
Med. Chem. Res.	1309				
MedChemComm	892				
J. Agric. Food Chem.	422				
Drug Metab. Dispos.	272				
J. Pesticide Sci.	245				
Nat. Chem. Biol.	170				
Crop Protection	129				
Pest. Manag. Sci.	126				
Others	570				

Data are also sometimes extracted from other journals and articles, when relevant to an area of particular interest

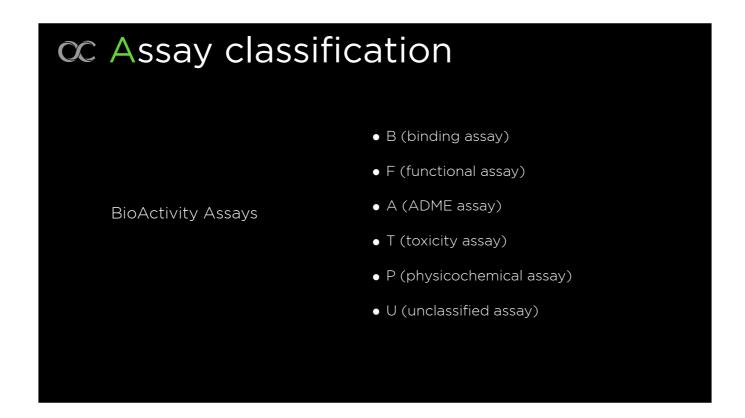
# • Patents • deposited data sets (often in collaboration with the depositor) • data from PubChem • data from BindingDB

In addition to data from the peer-reviewed literature, ChEMBL now incorporates some data extracted from patent documents. This data extraction effort is focused on patents containing compounds and bioactivity data for targets that are not currently well represented in ChEMBL (and in particular, those of interest to the NIH Illuminating the Druggable Genome project: https://commonfund.nih.gov/IDG/understudiedproteins (7)). To date, 74 050 activity measurements have been extracted from 241 patents, and these are available in ChEMBL as source 38. Additional, focused sets may also be included in future releases, where these add value to the existing ChEMBL data. A larger set of bioactivity data extracted from granted US patents by BindingDB (described previously) is also available in ChEMBL as source 37 (3).

#### ∞ Типы данных

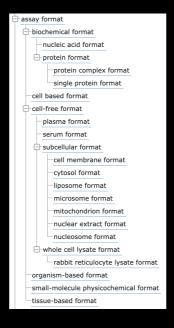
- Compounds tested
- Assays performed (type of assay, any cell-line, tissue or organism used; the property being assessed and the target (where applicable))
- Endpoints measured (IC50, Ki, CC50, etc)
- Relevant target information

From each article, details of compounds tested, assays performed, endpoints measured and relevant target information, are extracted. Compound structures are drawn in full (including any salt present) and saved in V2000 Mol file format. A brief description of the assay is abstracted; this typically includes the type of assay being performed; any cell-line, tissue or organism used; the property being assessed and the target (where applicable). Biological activity data is recorded; this includes (but is not limited to) binding measurements, efficacy in functional assays, pharmacokinetic data and toxicity endpoints.



Each assay is classified into one of the following categories: B (binding assay), F (functional assay), A (ADME assay), T (toxicity assay), P (physicochemical assay) or U (unclassified assay). Where the assay describes the interaction with a molecular target, this target is also recorded in the form of a UniProt (15) accession, or list of accessions. Quantitative and qualitative activity measurements are extracted in the form reported in the publication, with their respective activity types, units and qualifiers.

#### ∞ BAO format



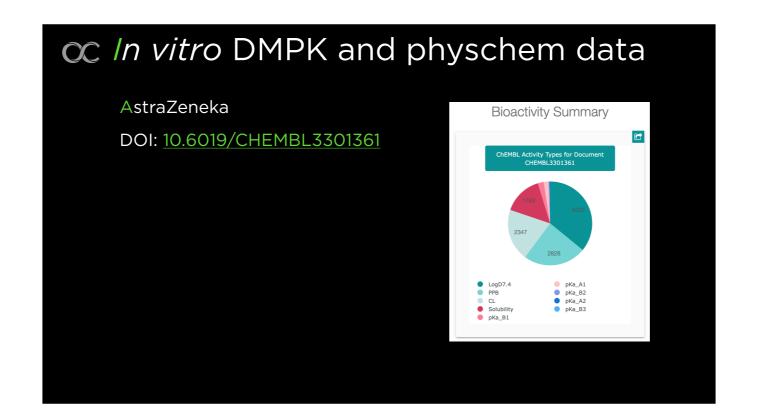
https://www.ebi.ac.uk/ols/ontologies/bao



Following initial data extraction, an extensive data curation and standardization process is applied before incorporating the information into ChEMBL. In brief, compound structures are standardized, salt-stripped and assigned identifiers based on Standard InChI (17); assay descriptions are mapped to controlled vocabularies such as the Cell Line Ontology (18), Uberon (19) and BioAssay Ontology (20); activity measurements and units are converted to a standard form; and multi-protein targets are created where required. This same curation process is also applied to other data sources within ChEMBL such as data extracted from patents, deposited data sets (often in collaboration with the depositor) and data from PubChem BioAssay and BindingDB.



ftp://ftp.ebi.ac.uk/pub/databases/chembl/ChEMBLdb/latest/



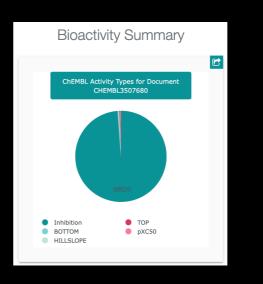
Experimental data determined at AstraZeneca on a set of compounds in the following assays: pKa, lipophilicity (LogD7.4), aqueous solubility, plasma protein binding (human, rat, dog, mouse and guinea pig), intrinsic clearance (human liver microsomes, human and rat hepatocytes). The references provided for the assays exemplify the experimental procedures used in generating the data.

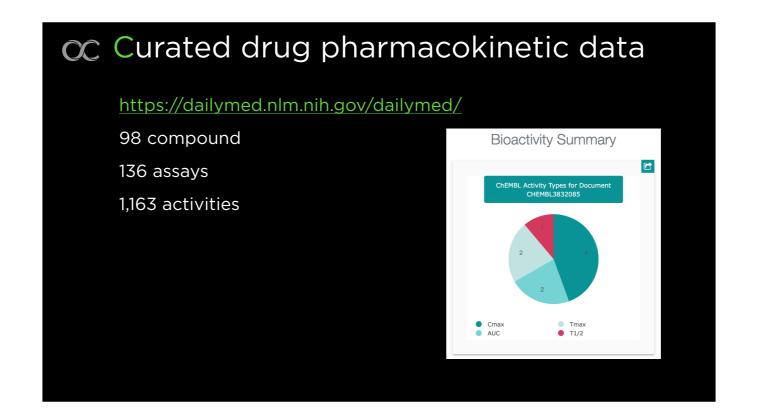
# oc Toxicity

Gates Library compound collection

DOI:10.6019/CHEMBL3507680

Human HepG2 cell viability assay in 384 well format





Knowledge of the pharmacokinetic properties of drugs is critical in understanding their safety and efficacy profiles, yet such data are often not readily available in a structured form. In order to begin to address this, pharmacokinetic measurements for 85 drugs have been extracted from reference books (21) and drug prescribing information (https://dailymed.nlm.nih.gov/dailymed/) and incorporated into ChEMBL as source 39



#### ∞ PARP1 case

Molecule ChEMBL \$ ID	Compound $_{\stackrel{\triangle}{\phi}}$ Key	Standard 🕏	Standard Relation	Standard 💂	Standard Units	pChEMBL $_{\oplus}$ Value	Comment ⊕	Assay ChEMBL \$ ID	Assa Desc
CHEMBL1949855	SID103905306	IC50	=	5650.0	nM	5.25	No Data	CHEMBL1953234	Inhibit PARP1 TACS- substr mins t colorir
CHEMBL1949862	26	IC50	=	1970.0	nM	5.71	No Data	CHEMBL1953234	Inhibit PARP1 TACS- substr mins t colorir
CHEMBL151549	32	IC50	=	490.0	nM	6.31	No Data	CHEMBL1953234	Inhibit PARP1 TACS- substr mins t colorir
work.	10i	PF50	=	10.1	No Data	No Data	No Data	CHEMBL2412859	Chemo factor, EC50 I humar to EC5 in hum



			count
BAO_Label	Standard_Type	Standard_Relation	
assay format	IC50	'<'	21.0
		9	246.0
		'>'	22.0
cell-based format	Activity	'='	8.0
	EC50	'='	281.0
		'>'	8.0
	ED50	<b>'</b>	34.0
		'>'	9.0
	IC50	'<'	1.0
		~	174.0
		'>'	15.0
single protein format	EC50	~	47.0
		'>'	2.0
	IC50	'<'	2.0
		9	1640.0
		'>'	170.0
	Kd	'<'	38.0
		9	107.0
		'>'	2.0
	Ki	'<'	4.0
		'='	1165.0
		'>'	3.0



• Биохимическая активность (на белке)

Активно: IC<sub>50</sub> <= 50 нМ</li>

• Неактивно: IC<sub>50</sub> > 50 нМ

• Собрать как можно больше данных

#### oc BAO Label & Assay format

			count
BAO_Label	Standard_Type	Standard_Relation	
assay format	IC50	'<'	21.0
		~	246.0
		'>'	22.0
cell-based format	Activity	~	8.0
	EC50	~	281.0
		'>'	8.0
	ED50	~	34.0
		'>'	9.0
	IC50	'<'	1.0
		9	174.0
		'>'	15.0
single protein format	EC50	~	47.0
		'>'	2.0
	IC50	'<'	2.0
		9	1640.0
		'>'	170.0
	Kd	'<'	38.0
		9	107.0
		'>'	2.0
	Ki	'<'	4.0
		9	1165.0
		'>'	3.0

# oc BAO Label & Assay format

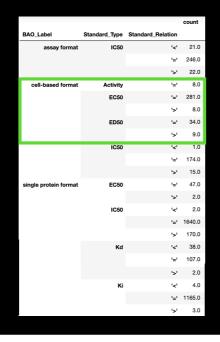
			count
BAO_Label	Standard_Type	Standard_Relation	
assay format	IC50	'<'	21.0
		9	246.0
		'>'	22.0
cell-based format	Activity	9	8.0
	EC50	9	281.0
		'>'	8.0
	ED50	9	34.0
		'>'	9.0
	IC50	'<'	1.0
		~	174.0
		'>'	15.0
single protein format	EC50	느	47.0
		'>'	2.0
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		9	1640.0
		'>'	170.0
	Kd	'<'	38.0
		9	107.0
		'>'	2.0
	Ki	'<'	4.0
		9	1165.0
		'>'	3.0

			count
BAO_Label	Standard_Type	Standard_Relation	
assay format	IC50	'<'	21.0
		~	246.0
		'>'	22.0
cell-based format	Activity	9	8.0
	EC50	9	281.0
		'>'	8.0
	ED50	9	34.0
		'>'	9.0
	IC50	'<'	1.0
		9	174.0
		'>'	15.0
single protein format	EC50	9	47.0
IC		'>'	2.0
	IC50	'<'	2.0
		9	1640.0
		'>'	170.0
	Kd	'<'	38.0
		9	107.0
		'>'	2.0
	Ki	'<'	4.0
		9	1165.0
		'>'	3.0

$$K_i = \frac{IC_{50}}{1 + [L]/K_D}$$

• Можно извлечь только неактивные

		•	count
BAO_Label	Standard_Type	Standard_Relation	
assay format	IC50	'<'	21.0
		~	246.0
		'>'	22.0
cell-based format	Activity	'='	8.0
	EC50	9	281.0
		'>'	8.0
	ED50	~	34.0
		'>'	9.0
	IC50	'<'	1.0
		'='	174.0
		'>'	15.0
single protein format	EC50	'='	47.0
		'>'	2.0
	IC50	'<'	2.0
		'='	1640.0
		'>'	170.0
	Kd	'<'	38.0
	Ki	'='	107.0
		'>'	2.0
		'<'	4.0
		-	1165.0
		'>'	3.0



- Исключим cell death assays
- Исключим cell proliferation assay
- Учтем специфичные к PARP1 методики
- Учтем только активные соединения

# cc Duplicates

	IDs	value
		count
0	CHEMBL1086580	15.0
1	CHEMBL1094636	10.0
2	CHEMBL1173055	9.0
3	CHEMBL190434, CHEMBL426270	4.0
4	CHEMBL194482	2.0
5	CHEMBL249813	3.0
6	CHEMBL251027	3.0
7	CHEMBL338790	2.0
8	CHEMBL339695	2.0
9	CHEMBL372303	12.0
10	CHEMBL3764816	3.0
11	CHEMBL3912508	2.0
12	CHEMBL3960883	2.0
13	CHEMBL506871	15.0
14	CHEMBL521686	27.0

# cc 'CHEMBL521686'

	Assay_ChEMBL_ID	value	Standard_Relation
2275	CHEMBL3887957	0.90	~
2266	CHEMBL3743825	1.00	~
2254	CHEMBL3995357	1.38	~
2270	CHEMBL3995357	1.40	~
2252	CHEMBL3767923	1.94	~
2279	CHEMBL4029615	2.09	~
2288	CHEMBL4055669	3.59	~
2261	CHEMBL2330729	4.00	~
2258	CHEMBL3107467	4.50	9
2284	CHEMBL982809	5.00	~
2262	CHEMBL3428885	5.00	~
2272	CHEMBL3295803	6.00	~
2253	CHEMBL2412862	6.00	~
2276	CHEMBL3293313	7.00	~
2267	CHEMBL4055845	8.10	~
2289	CHEMBL3995359	10.00	~
2283	CHEMBL3107875	10.00	~
2255	CHEMBL3995359	10.23	~
2257	CHEMBL3738635	11.90	9
2282	CHEMBL3887586	12.00	~
2251	CHEMBL3241058	15.50	¥
2287	CHEMBL3995358	18.90	~
2286	CHEMBL3887955	43.00	~
2290	CHEMBL3887955	50.00	'<'
2269	CHEMBL3995358	69.18	9
2259	CHEMBL3371141	84.96	~
2273	CHEMBL3887956	86.32	~