InChI Applications in ChEMBL

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Virtual Workshop on International Chemical Identifiers (InChl) 24-03-2021



Outline

- ChEMBL Database overview and content
- InChI uses in ChEMBL and UniChem
 - Compound Registration
 - ChEMBL Structure Pipeline
 - Cross-References



ChEMBL Database

- Open access database for drug discovery
- Core bioactivity data from:
 - Key MedChem journals
 - Public databases
 - Review articles, book chapters, etc
 - Contributed datasets
- Integrated with data on:
 - Clinical development and marketed drugs (from ClinicalTrials.gov, FDA, USANs, INNs)

(୦)

14,347

Targets

• Latest release (version 28) statistics:



InChl Applications in ChEMBL





InChl Applications in ChEMBL – Registration





ChEMBL Compound Registration

- Standardised V2000 molfile as the primary chemical structure representation
- Standard InChI and corresponding hashed InChIKey are calculated from the molfile using software provided by InChI Trust (current version 1.05)
- Standard InChI and InChIKey used as a measure of uniqueness for a chemical structure in ChEMBL
- Compounds from different sources with the same Standard InChI and InChIKey are considered to be the same compound and are assigned the same ChEMBL identifier (CHEMBL_ID)
- However, V2000 molfile and Standard InChI have a few limitations, *e.g.*,
 - Do not support the use of relative stereochemistry, only absolute or no stereochemistry



ChEMBL Compound Registration

- Several drugs are formulated as racemic mixtures (e.g., Orphenadrine, Milnacipran) or as unequal mixture of both enantiomers (e.g., Amphetamine)
- Chemical representation of these mixtures is a challenge and different data providers may have different registration rules

Example of structure representations from various sources for Milnacipran





ChEMBL Compound Registration

- Relative stereochemistry can be annotated in V3000 molfiles
- ChEMBL has started to store it internally for manually annotated compounds

Structure representation of different molfile versions (and corresponding Standard InChIKey) for Milnacipran



• V3000 molfile not fully represented by Standard InChl



InChl Applications in ChEMBL – Structure Pipeline





ChEMBL Structure Pipeline

- ChEMBL Structure Pipeline is comprised of three processes:
 - *Checker*: identifies and validates structures and identifies problems before structures are loaded into the database
 - Standardizer: processes (standardises) chemical structures according to a set of predefined rules
 - GetParent: generates parent structures based a set of rules and defined list of salts and solvents
- Standardizer and GetParent have been rewritten and adapted from rules originally implemented using a commercial software toolkit
- Checker was developed more recently in an attempt to identify problem structures



ChEMBL Structure Pipeline – *Checker*

- *Checker* process validates structures prior to loading compounds into ChEMBL
- If an error or problem is detected in the structure, a score is assigned
- The higher the score, more serious is the problem

Penalty Score	Penalty Explanation	Database Loader Actions
7	Error -9986 (Cannot process aromatic bonds) Illegal input InChI: Unknown element(s)	a fatal error and no data is loaded into ChEMBL
6	all atoms have zero coordinates InChI: Accepted unusual valence(s) InChI: Empty structure molecule has 3D coordinates molecule has a radical that is not found in the known list molecule has six (or more) atoms with exactly the same coordinates number of atoms less than 1 polymer information in mol file V3000 mol file	data is loaded into the database but without a molfile
5	InChI_RDKit/Mol stereo mismatch Mol/Inchi/RDKit stereo mismatch RDKit_Mol/InChI stereo mismatch molecule has a bond with an illegal stereo flag molecule has a bond with an illegal type molecule has a crossed bond in a ring molecule has two (or more) atoms with exactly the same coordinates	all data is loaded
2	InChI_Mol/RDKit stereo mismatch molecule has a stereo bond in a ring molecule has an atom with multiple stereo bonds molecule has a stereo bond to a stereocenter molecule has the 3D flag set for a 2D conformer Other InChI Warnings	

 InChI warnings are a component of the *Checker*



ChEMBL Structure Pipeline Publication

Bento et al. J Cheminform (2020) 12:51 https://doi.org/10.1186/s13321-020-00456-1 Journal of Cheminformatics

METHODOLOGY

Open Access

Check for

An open source chemical structure curation pipeline using RDKit

A. Patrícia Bento¹, Anne Hersey¹, Eloy Félix¹, Greg Landrum², Anna Gaulton¹, Francis Atkinson^{1,3}, Louisa J. Bellis^{1,4}, Marleen De Veij¹ and Andrew R. Leach^{1*}



ChEMBL Structure Pipeline Updates – Standardizer

- Generate canonical tautomer
 - Standard InChI is independent of the tautomeric form of a compound and hence different tautomers of a compound will have the same Standard InChI. Therefore, in ChEMBL they are considered as being the same compound
 - However, Standard InChI has a few limitations, e.g.,
 - unable to recognise some keto-enol tautomers as being the same compound





ChEMBL Structure Pipeline Updates – *Standardizer*

Generate canonical tautomer

• Update





InChl Applications in ChEMBL – Cross References





UniChem

- Cross-references to compounds in other sources provided by UniChem
- UniChem is a compound identifier mapping service developed at EMBL-EBI that provides structure-based hyperlinks to chemical structures in other databases
- Historically, each EMBL-EBI resource would individually manage its own links to all other resources
- In UniChem, mappings are maintained centrally, allowing for the simple inclusion of additional resources





UniChem

- Standard InChI and InChIKey are the chemical structure normalizing key for UniChem
- For UniChem, each new structure (as defined by the Standard InChI) is assigned a UniChem identifier (UCI) within the system and this association between the structure and its UCI is never updated or deleted. The only updates permitted are to indicate whether the assignment is 'current' or 'obsolete'.
- All structures and assignments of whatever historical status are captured
- UniChem Stats

Release Date	04-MAR-2021
Total number of Structures	176,340,296
Total number Assignments*	285,463,117
Number Current Assignments	214,367,308
Number Obsolete Assignments	71,095,809
Number of Sources	40



• UniChem can be used to:

www.ebi.ac.uk/unichem/

 Search for identical small molecule data across a wide variety of chemically aware resources, using either src_compound_ids, InChI or InChIKeys

				Show 50	o 😑 entries					Apply filter:
	IniChe	m		src_id	Source Name	src_compound_id	Currently Assigned	LR *	UCI **	Standard InChIKey
				1	chembl	CHEMBL99946	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Home / Search	EBI > Databases > Small Mo	lecules → UniChem		7	chebi	<u>136040</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Web Services				15	surechembl	SCHEMBL1414867	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Connectivity Search	Home / Search			2	drugbank	<u>DB08918</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Sources	UniChem efficiently produce	s cross-references between chemical structure identifiers from different databases (more backgr	ound)	4	gtopdb	7435	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Seneral Into	chieffen endernig preduced		ound).	9	zinc	ZINC00000000506	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Getting in touch		Read More		11	ibm	BCAB701D6C6F434F6E3261DFA5BDCE4A	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
FAQ				14	fdasrs	UGM0326TXX	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Downloads				21	pubchem_tpharma	<u>16010452</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Connectivity Info	Query UniChem			22	pubchem	<u>6917779</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
+ Other				29	nikkaji	<u>J259.860D</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Analysis.		GJJFMKBJSRMPLA-DZGCQCFKSA-N		31	bindingdb	<u>50032379</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Top Level Stats	Query term(s):			34	drugcentral	4864	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
Structures by Source				39	chemicalbook	<u>CB13133687</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
FULIK		src compound id InChI InChIKev		40	dailymed_old	levomilnacipran	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
FIKHB				45	dailymed	LEVOMILNACIPRAN	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
SCFIB		Submit Query		46	clinicaltrials	<u>F 2695</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
				46	clinicaltrials	<u>F-2695</u>	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
				46	clinicaltrials	F2695	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
				46	clinicaltrials	FETZIMA	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
		Example Queries		46	clinicaltrials	LEVOMILNACIPRAN	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
				47	rxnorm	FETZIMA	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
				47	rxnorm	LEVOMILNACIPRAN	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
				47	rxnorm	LEVOMILNACIPRAN HYDROCHLORIDE	Yes		10620	GJJFMKBJSRMPLA-DZGCQCFKSA-N
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	Bio	MedBridges chemical keys for the slocks		Showing 1	to 24 of 24 entries					

EMBL-EBI

- UniChem can be used to:
 - Search for molecules with the same atom connectivity but different stereochemistry or isotopic composition or existing in a different salt form, using either src_compound_ids or InChIKeys



Different stereochemistry and isotopes



- UniChem can be used to:
 - Search for molecules with the same atom connectivity but different stereochemistry or isotopic composition or existing in a different salt form, using either src_compound_ids or InChIKeys



EMBL-EB

- UniChem can be used to:
 - Search for molecules with the same atom connectivity but different stereochemistry or isotopic composition or existing in a different salt form, using either src_compound_ids or InChIKeys



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Query term:	GJJFMKBJSRMPLA-D	ZGCQCFKSA-N		
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	/p	Protons. The number	er of protons that must be added to o	r removed from the form	ula to give th	e input c	omposi	tion. (se	e * belo	w)				
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matches	1	ChEMBL	CHEMBL471001	1		0	0	1	0	0	0		GJJFMKBJSRMPLA- ZFWWWQNUSA-N	1
matches	1	ChEMBL	CHEMBL99946	1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
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matches	2	DrugBank		1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
matches	4	Guide to Pharmacology	7435	1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
matches	4	Guide to Pharmacology	<u>7436</u>	1		0	0	1	1	1	0		GJJFMKBJSRMPLA- UHFFFAOYSA-N	1
matches cpts of	5	PubChem: Drugs of the Future	<u>12013109</u>	1	HCI		0	0	0	0	0		XNCDYJFPRPDERF- NQQJLSKUSA-N	1
matches	7	ChEBI	135005	1		0	0	1	0	1	0		GJJFMKBJSRMPLA- UHFFFAOYSA-N	1
matches	7	ChEBI	<u>136040</u>	1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
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matches	1	ChEMBL	CHEMBL252923	1		0	0	0	1	0	0	1	GJJFMKBJSRMPLA- HIFRSBDPSA-N	1
matches	1	ChEMBL	CHEMBL259209	1		0	0	1	1	1	0		<mark>GJJFMKBJSRMPLA</mark> - UHFFFAOYSA-N	1
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matches	1	ChEMBL	CHEMBL99946	1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
matches	2	DrugBank		1		0	0	1	1	1	0		GJJFMKBJSRMPLA- UHFFFAOYSA-N	1
matches	2	DrugBank		1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
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matches	4	Guide to Pharmacology	<u>7436</u>	1		0	0	1	1	1	0		GJJFMKBJSRMPLA- UHFFFAOYSA-N	1
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matches	7	ChEBI	<u>135005</u>	1		0	0	1	1	1	0		GJJFMKBJSRMPLA- UHFFFAOYSA-N	1
matches	7	ChEBI	<u>136040</u>	1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
matches cpts of	7	ChEBI	<u>31854</u>	1	HCI		0	0	0	0	0	1	XNCDYJFPRPDERF- PBCQUBLHSA-N	1
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	/р	Protons. The number	er of protons that must be added to o	r removed from the forn	nula to give th	ne input c	omposi	tion. (se	e * belc	ow)				
-	/b	Double bond stereo												
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matches cpts of	1	ChEMBL	CHEMBL2094056	1	HCI		0	1	0	0	0		XNCDYJFPRPDERF- SLHAJLBXSA-N	1
matches cpts of	1	ChEMBL	CHEMBL2105732	1	HCI		0	0	0	0	0		XNCDYJFPRPDERF- NQQJLSKUSA-N	1
matches	1	ChEMBL		1		0	0	0	1	0	0	1	GJJFMKBJSRMPLA- HIFRSBDPSA-N	1
matches	1	ChEMBL	CHEMBL259209	1		0	0	1	1	1	0		GJJFMKBJSRMPLA- UHFFFAOYSA-N	1
matches cpts of	1	ChEMBL	CHEMBL3527085	1	HCI		0	1	1	1	0		XNCDYJFPRPDERF- ZEVXLFMTSA-N	1
matches cpts of	1	ChEMBL	CHEMBL4297064	1	HCI		0	1	1	1	0		XNCDYJFPRPDERF- UHFFFAOYSA-N	1
matches	1	ChEMBL		1		0	0	1	0	0	0		GJJFMKBJSRMPLA- ZFWWWQNUSA-N	1
matches	1	ChEMBL		1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
matches	2	DrugBank		1		0	0	1	1	1	0		GJJFMKBJSRMPLA- UHFFFAOYSA-N	1
matches	2	DrugBank		1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
matches	4	Guide to Pharmacology	7435	1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
matches	4	Guide to Pharmacology	<u>7436</u>	1		0	0	1	1	1	0		GJJFMKBJSRMPLA- UHFFFAOYSA-N	1
matches cpts of	5	PubChem: Drugs of the Future	<u>12013109</u>	1	HCI		0	0	0	0	0		XNCDYJFPRPDERF- NQQJLSKUSA-N	1
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matches	7	ChEBI	<u>136040</u>	1		0	0	0	0	0	0		GJJFMKBJSRMPLA- DZGCQCFKSA-N	1
matches cpts of	7	ChEBI	<u>31854</u>	1	HCI		0	0	0	0	0	1	XNCDYJFPRPDERF- PBCQUBLHSA-N	1
matches	7	ChEBI		1		0	0	0	1	0	0	1	GJJFMKBJSRMPLA- HIFRSBDPSA-N	1
Showing 1 to 15	5 of 238 e	ntries									Fi	rst Pre	vious 1 2 3 4 5 Nex	t Last



- UniChem can be used to:
 - Automatically generate hyperlinks from within a resource to other resources

Compound Report Card

Name And Classification		- UniChem	Cross References
Image: state stat	ID: CHEMBL99946 Name: LEVOMILINACIPRAN Max Phase: Improved Improve	Image: Stress of the stress of th	i S0032379 iiiiiiiiiiiiiiiiiiiiiiiiiiiiiiiiiiii



- UniChem can be used to:
 - Automatically generate hyperlinks from within a resource to other resources

Compound Report Card

ame And Classification			- Unicher	n Connectivity Layer Cro	SS References								
NH ₂ (c,	ID: Name: Max Phase: Molecular Formula: Molecular Weight: ChEMBL Synonyms:	CHEMBL99946 LEVOMILNACIPRAN Approved C15H22NZO 246.35 F2695 F 2695 F-2695 LEVOMILNACIPRAN	C UniChem GJJFMKBJSF Full InCh	UniChem Connectivity Layer Cross References for CHEMBL99946									
	Synonyms From Alternate Forms:	F 2695 HYDROCHLORIDE F-2695 HYDROCHLORIDE F2695 HYDROCHLORIDE LEVOMILNACIPRAN HYDROCHLORIDE	InChI=1S/0	15H22N20/c1-3-17(4-2)14(18)15(10-13(15)11-16)12-8-6-5-7-9-12/h5-9,13H,3-4,10-11,16H2,1-2H3, S	/t13-,15+/m0/s1 I P	SI	IP SP	SIP				
	Trade Names: Molecule Type:	FETZIMA Small molecule	ChEMBL	CHEMBL99946 CHEMBL2105732	CHEMBL252923 CHEMBL4297064 CHEMBL471001 CHEMBL2094056 CHEMBL259209 CHEMBL1237129		CHEMBL3527085						
Structure Search			Guide to Pharmaco	logy 7435	7436								
			PubChem Drugs of Future	: he 12013109									
			ChEBI NIH Clinic	136040 al	94468 135005 31854 SAM001247035								
			ZINC	ZINC00000000506	ZINC00000014037 ZINC000005649654 ZINC00001846944								
			eMolecule	s 31706396 30100405	36754037 36500460 36500460 30069726 29543180		36754038						
			IBM Pate System	BCAB701D6C6F434F6E3261DFA5BDCE4	7B5D61094EBF88BDDBD5CAC963214F7A AACD9B490ECD4EA5F9FD06A92ED88D94 9ED58872F7C7F5510B85B6D53849CB 0E9CA3DADB8667A092086134ABFAEF8 DCFA8846FBE580E408339B6DC13975EC 0EC7C98A888839109AE7F8FC4759DE34								
			FDA SRS	UGM0326TXX 371U2ZK31U	ES1038J3C4								



UniChem Publications

Chambers et al. Journal of Cheminformatics 2013, 5:3 http://www.jcheminf.com/content/5/1/3

DATABASE





UniChem: a unified chemical structure cross-referencing and identifier tracking system

Jon Chambers^{1*}, Mark Davies¹, Anna Gaulton¹, Anne Hersey¹, Sameer Velankar², Robert Petryszak³, Janna Hastings⁴, Louisa Bellis¹, Shaun McGlinchey¹ and John P Overington¹ Chambers et al. Journal of

Chambers *et al. Journal of Cheminformatics* 2014, **6**:43 http://www.jcheminf.com/content/6/1/43



DATABASE

Open Access

UniChem: extension of InChl-based compound mapping to salt, connectivity and stereochemistry layers

Jon Chambers^{*}, Mark Davies, Anna Gaulton, George Papadatos, Anne Hersey and John P Overington



UniChem Update Coming Soon

- New interface
- New functionalities

Unich	nem			
SEARCHES	SOURCES	ABOUT	API	
	Unichem Compound search			
	Compound Source Find sources for a given comp	es ound	Connectivity Connectivity search	



Acknowledgements

ChEMBL Structure Pipeline

ChEMBL



Greg Landrum

wellcome^{trust}

Strategic Award

RDKit Support and Chemical Data Science - T5 Informatics

Anne Hersey Patrícia Bento Eloy Félix Anna Gaulton Francis Atkinson Louisa Bellis Marleen de Veij Andrew Leach UniChem

ChEMBL

Jon Chambers Anne Hersey Anna Gaulton Mark Davies George Papadatos Shaun McGlinchey Louisa Bellis James Blackshaw John Overington



Chris Steinbeck Janna Hastings



Sameer Velankar



Robert Petryszak

Funding:



BioMedBridges EU-OPENSCREEN:



