
Cambridge Cheminformatics Network Meeting

*Setting up Cheminformatics Support for
the Open Source Malaria Project*

Chris Swain

Cambridge MedChem Consulting

<http://www.cambridgemedchemconsulting.com>

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Open Source Malaria



- The Open Source Malaria project is trying a different approach to curing malaria. Guided by open source principles, everything is open and anyone can contribute.
- <http://opensourcemalaria.github.io/NewSite/>
- This Landing Page aggregates the most recent activity in Open Source Malaria. In open source research all data and ideas are freely shared, anyone may participate as an equal partner and there will be no patents - think “Linux for Malaria Research”.

Organisation of project uses GitHub

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OpenSourceMalaria / OSM_To_Do_List Watch 39 Star 15 Fork 4

Issues Pull requests Labels Milestones is:open New Issue

Clear current search query, filters, and sorts

132 Open 206 Closed Author Labels Milestones Assignee Sort

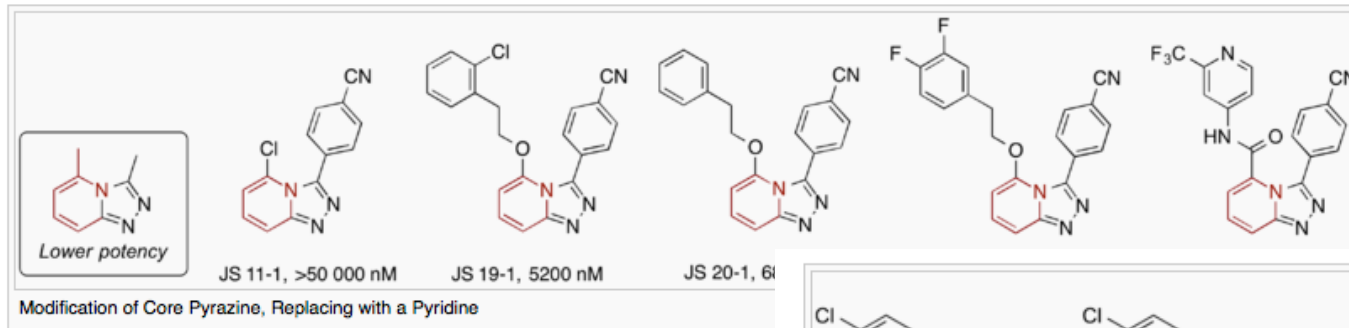
- First Year project student (TSP) targets for OSM** Being Synthesised Now High Priority Series 4 Synthetic Chemistry Needed #338 opened 2 days ago by alintheopen
- New tool for 3D conformations exploration** #337 opened 3 days ago by lpatiny
- Competition for Making a Graphical Abstract** Collaboration Request Competition High Priority Paper 1 #336 opened 11 days ago by alintheopen
- Structures missing** question #335 opened 11 days ago by drc007
- Metabolism** Series 4 #334 opened 12 days ago by drc007
- Trends in logD vs metabolic stability?** Administration question Search/Data Needed Series 4 #333 opened 16 days ago by mattodd

Organization of data

- Historically project data published on web pages
- All in public domain
- But.....
- Challenging to collate all data associated with project
- Multiple compound identifiers
- Multiple data sources
- Not always clear which assay data refers to

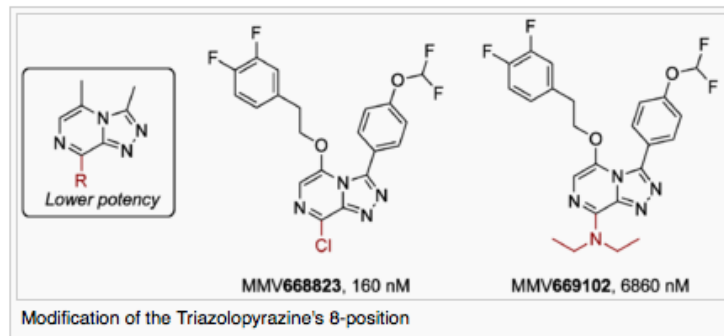
Examples of data

spending triazolopyrazine compounds.

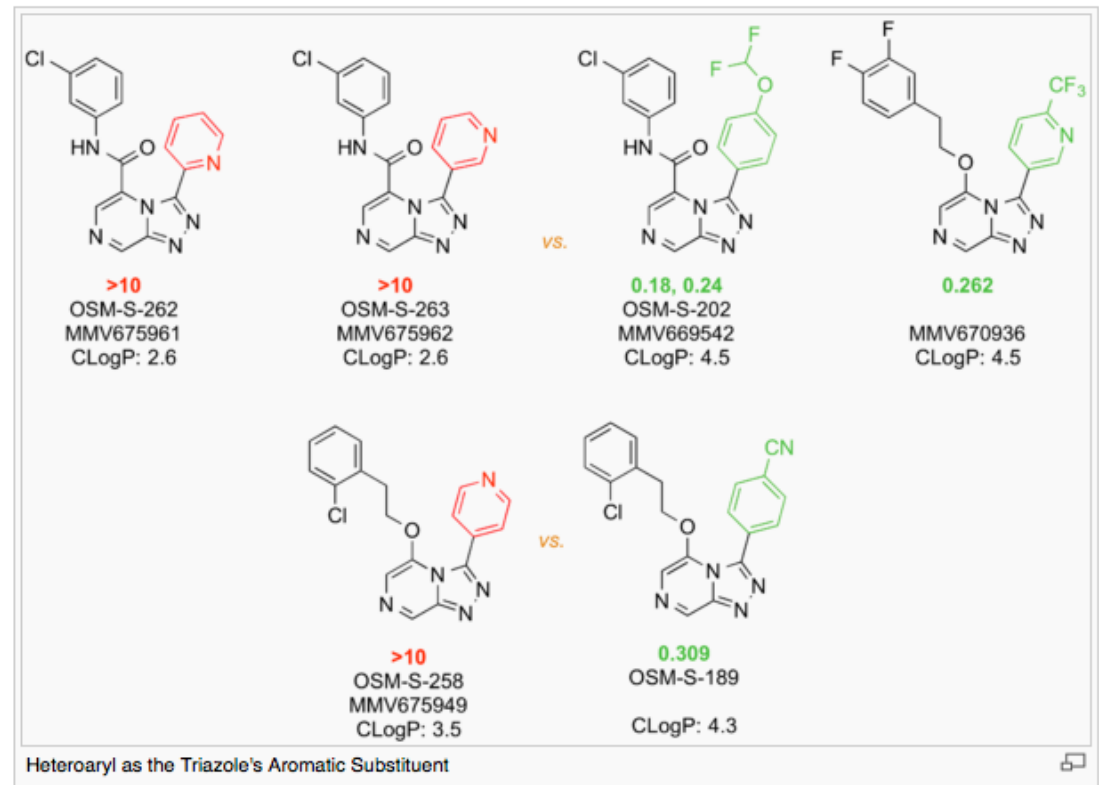


Modification of Pyrazine Substitution Pattern

It is thought possible that the pyrazine moiety of the triazolopyrazine could undergo aldehyde substitution. If modifications were made to the substitution of the "southernmost" (8-) ring C-H, all lowering potency.



...the triazolo-pyrazine moiety was investigated. The side chain on the pyrazine ring was shifted to the



Stage 1:- Implement single data source

- Use Google spreadsheet
- Users comfortable with data entry
- Negligible set up or support required

Google doc spreadsheet

- Simple spreadsheet has allowed multiple users to enter data (BIG THANKS TO ALL!).
- https://docs.google.com/spreadsheets/d/1Rvy6OiM291d1GN_cyT6eSw_C3ISuJ1jaR7AJa8hgGsc/edit#gid=510297618
- Currently contains 265 rows (compounds), and 46 columns (descriptors and expt data).
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 - http://docs.google.com/spreadsheets/d/1Rvy6OiM291d1GN_cyT6eSw_C3ISuJ1jaR7AJa8hgGsc/export?format=tsv
- How can we leverage this data source.

Google doc spreadsheet

Malaria Molecules, Click on "ID" to see visualisation ☆

File Edit View Insert Format Data Tools Add-ons Help All changes saved in Drive

Comments Share

fx Pfal EC50 uMol (Mean)

	A	B	C	D	E	F	G	H	I	J	K	L
1	ID	SMILES	OSM	InChI	IChI Key	Series	Pfal EC50 (Inh) qualifier	Pfal EC50 (Inh)	Pfal IC50 (GSK)	Pfal IC50 (Syngene)	Pfal IC50 (Dundee)	Pfal EC50 uM
2	MMV670944	<chem>c1ncc2n(c1C(=O)Nc1ccnc(c1)C(F)F</chem>	(OSM-S-175)	InChI=1S/C18H11F3		4		0.1404				
3	MMV668958	<chem>c1ncc2n(c1C(NCc1cc(ccc1)Cl)=O)c(r</chem>	(OSM-S-176)	InChI=1S/C20H14Cl		4		0.25				
4	MMV669000	<chem>c1ncc2n(c1C(N1Cc3c(C1)cccc3)=O)x</chem>	(OSM-S-177)	InChI=1S/C21H15F2		4	>	10				
5	MMV669001	<chem>c1ncc2n(c1C(NCc1ccc(cc1)Cl)=O)c(r</chem>	(OSM-S-178)	InChI=1S/C20H14Cl	InChI=1S/C20H1	4		1.34				
6	MMV669003	<chem>c1ncc2n(c1C(NCC1CCOCC1)=O)c(n</chem>	(OSM-S-179)	InChI=1S/C19H19F2		4		10				
7	MMV669010	<chem>c1ncc2n(c1C(N1CCN(CC1)c1ncccc1</chem>	(OSM-S-180)	InChI=1S/C23H22F2	UYWFTHNGLI	4		10				
8	MMV669011	<chem>c1ncc2n(c1C(NCCN1CCOCC1)=O)c</chem>	(OSM-S-181)	InChI=1S/C19H20F2	LYBWZYLEK BK	4		10				
9	MMV669022	<chem>c1ncc2n(c1C(N1CCCC1)=O)c(nn2)c</chem>	(OSM-S-182)	InChI=1S/C17H15F2	HDIHMYGJUUD	4		10				
10	MMV669104	<chem>c1ncc2n(c1C(N1C(CCC1)c1cccc1)=</chem>	(OSM-S-183)	InChI=1S/C23H19F2		4		3.56				
11	MMV669023	<chem>c1ncc2n(c1C(N1CCN(CC1)S(=O)(C)</chem>	(OSM-S-184)	InChI=1S/C18H18F2		4		10				
12	MMV669020	<chem>c1ncc2n(c1C(N1CCOCC1)=O)c(nn2)</chem>	(OSM-S-185)	InChI=1S/C17H15F2	LFRCBCKOIXZI	4		10				
13	MMV669024	<chem>c1ncc2n(c1C(=O)N[C@@H](c1cccc</chem>	(OSM-S-186)	InChI=1S/C21H17F2		4		1.36				
14	MMV639725	<chem>N#CC(C=C1)=CC=C1C2=NN=C3C=</chem>	(OSM-S-189)	InChI=1S/C20H14Cl	DQNFHRXL TAC	4			0.309			
15	MMV669542	<chem>c1ncc2n(c1C(Nc1cccc(c1)Cl)=O)c(nn</chem>	(OSM-S-202)	InChI=1S/C19H12Cl		4		0.18	0.242			
16	MMV669844	<chem>[C@H](COc1cncc2n1c(nn2)c1ccc(cc</chem>	(OSM-S-218)	InChI=1S/C21H15F2		4		0.04				
17	MMV639565	<chem>FC1=C(F)C=CC(COC2=CN=CC3=I</chem>	(OSM-S-272)	InChI=1S/C19H13Cl	PMHUSEXABGI	4		0.038				
18	MMV669846	<chem>c1ncc2n(c1OCCc1cc(c(cc1)F)F)c(on</chem>	(OSM-S-273)	InChI=1S/C20H14Cl	MQHQNFQVEX	4		0.11				
19	MMV670250	<chem>c1ncc2cnc(n2c1OCCc1cc(c(cc1)F)F</chem>	(OSM-S-274)	InChI=1S/C20H14Cl	JHORUMGXLQ	4		0.83				

Column Description

Malaria column description ☆

File Edit View Insert Format Data Tools Add-ons Help All changes saved in Drive

Comments Share

fx

	A	B	C	D	E	F	G
1	columnName	Description	Target	Accession Number	Organism	Strain	Format
2							
3	ID	Unique identifier of the molecule					MMV639965
4	SMILES	Chemical structure as a SMILES					c1ccccc1CN
5	OSM	Identifier					OSM-S-272
6	InChI	Chemical structure as InChI useful for uniqueness check and some search engines					InChI=1S/C19H13
7	InChi Key	Chemical structure as a InChi Key useful for some search engines					DQNFHRXLTAF
8	Series	Number assigned to structural class					
9	Pfal EC50 (Inh)	In vitro EC50, micromolar, inherited data, NF54, 3H-Hypoxanthine, 72 h timepoint			Plasmodium falciparum		
10	Pfal IC50 (GSK)	In vitro IC50, micromolar, GSK			Plasmodium falciparum		
11	Pfal IC50 (Syngene)	In vitro IC50, micromolar, Syngene			Plasmodium falciparum		
12	Pfal IC50 (Dundee)	In vitro IC50, micromolar, Dundee			Plasmodium falciparum		
13	Single Shot Inhibition %	Single shot Pf, 2 micromolar, % inhibition, GSK Tres Cantos			Plasmodium falciparum		

Google doc spreadsheet

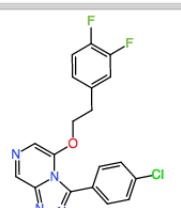
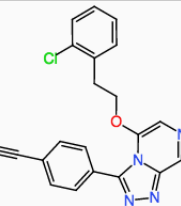
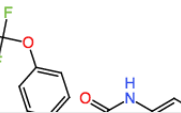
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- How can we leverage this data source?

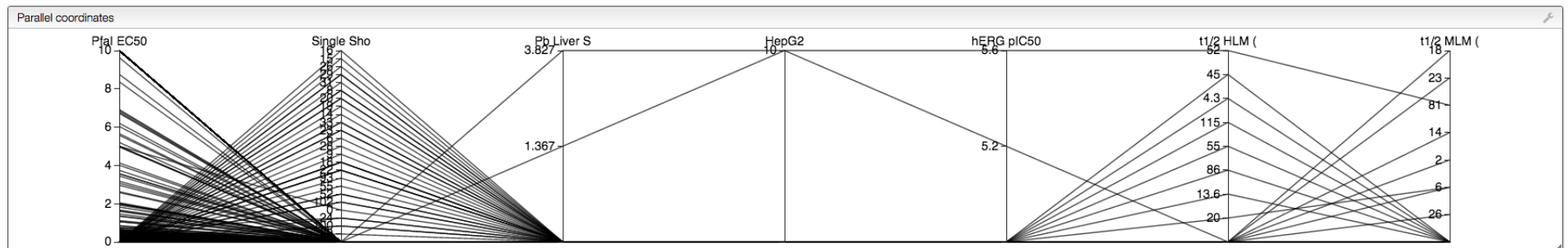
Web View

- One click access to data provided by chemifo.org (huge thanks to Luc Patiny)
- http://www.cheminfo.org/fluor/malaria/Display_data.html
- Imports data from live Google spreadsheet
- Calculates a variety of physiochemical properties on the fly.
- Can display data as table and a variety of different plots.

Web View

DISPLAY DATA

Mofie	ID	smiles	OSM	InChI	InChI Key	Series	Pfal EC50 (l...	Pfal EC50 (lnh)	Pfal IC50 (G...	Pfal IC50 (S...	Pfal IC50 (D...	Pfal EC50 u...	Single Shot L...	Ion Regulati...	Pb Liver Stage	HepG2	hERG pIC50	t1/2 HLM (M...	t1/2 HLM (lnh)	t1/2 N	
	MMV639565	FC1=C(F)C=...	(OSM-S-272)	InChI=1S/C1...	PMHUSEXA...	4		0.038				0.038									
	MMV639725	N#CC(C=C)...	(OSM-S-189)	InChI=1S/C2...	DQNFHRXLT...	4			0.309				99								
	MMV657963	O=C(NC1=C...	OSM-S-271	InChI=1S/C1...	KPPYAXXED...	4		0.234				0.234									



Vortex

- Vortex is a high performance data visualisation and analysis platform
- Written script to automatically import from Google spreadsheet
- http://macinchem.org/reviews/vortex/tut26/scripting_vortex26.php

Vortex

```
# Python imports
```

```
import urllib2
import urllib
import csv
import sys
from com.xhaus.jyson import JysonCodec as jsn
```

```
# Vortex imports
```

```
import com.dotmatics.vortex.util.Util as Util
import com.dotmatics.vortex.mol2img.jni.genImage as genImage
import com.dotmatics.vortex.mol2img.Mol2Img as mol2Img
import com.dotmatics.vortex.table.VortexTableModel as vtm
import jarray
import binascii
import string
import os
```

```
mystr = "http://docs.google.com/spreadsheets/d/
1Rvy6OiM291d1GN_cyT6eSw_C3ISuJ1jaR7AJa8hgGsc/export?format=tsv"
```

```
myreturn = urllib2.urlopen(mystr).read()
list1 = myreturn.split("\n")
```

```
TableName = "OSMData"
```

```
# Get column names
```

```
column_names = list1[0].split("\t")
```

```
rows = []
```

```
for i in list1[1:]:
    row = i.split("\t")
    rows.append(row)
```

```
arrayToWorkspace(rows, column_names, TableName)a
```

ID	SMILES	OSM	InChI	IChI Key	Series	Pfal ECSO qualifier	Pfal ECSO uMol (Mean)	Pfal ECSO uMol (results)
MMV669848			InChI= 1S/C21H17F...		4		0.11	0.11
MMV670659			InChI= 1S/C22H14F...		4		0.1147	0.1147
MMV672727			InChI= 1S/C21H15F...		4		0.1228	0.1228
MMV672687			InChI= 1S/C20H14F...		4		0.124	0.124
MMV670944		OSM-S-17...	InChI= 1S/C18H11F...		4		0.1404	0.1404
MMV668823			InChI= 1S/C20H13C...		4		0.16	0.16
MMV672941			InChI= 1S/C22H12F...		4		0.1734	0.1734
MMV669542		OSM-S-20...	InChI= 1S/C19H12C...		4		0.18	0.18
MMV670767			InChI= 1S/C19H11C...		4		0.1911	0.1911
MMV657963		OSM-S-271	InChI= 1S/C19H11C... KPPYAXXEDYTONY...		4		0.234	0.234
MMV668958		OSM-S-17...	InChI= 1S/C20H14C...		4		0.25	0.25

iPython Notebook

- iPython Notebooks are becoming increasingly popular formats for combining code execution, rich text, mathematics, plots, chemical structures etc
- <http://www.macinchem.org/reviews/osm/osmipython.php>

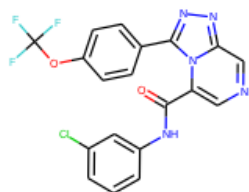
iPython Notebook

```
!wget http://docs.google.com/spreadsheets/d/1Rvy6OiM291d1GN_cyT6eSw_C3ISuJ1jaR7AJa8hgGsc/export?format=tsv
```

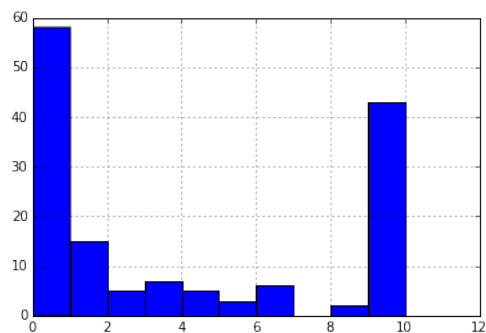
```
from rdkit.Chem import AllChem as Chem
from rdkit.Chem.Draw import IPythonConsole
from rdkit.Chem import PandasTools
from rdkit.Chem import Draw
import pybel
```

```
import pandas as pd
datafile = pd.read_table('./export?format=tsv')
smiles = datafile['SMILES'].loc[2]
```

```
mol = Chem.MolFromSmiles(smiles)
mol
```



```
datafile['Pfal EC50 uMol (Mean)'].hist()
```



Everybody has access

- Everybody has live up to the minute access to the data
- All the data
- Why not give it a try and then contribute your findings and suggestions to the [Open Source Malaria project.](#)