

Electronic Supplementary Information (ESI)

Chemistry in a graph: Modern Insights into Commercial Organic Synthesis Planning

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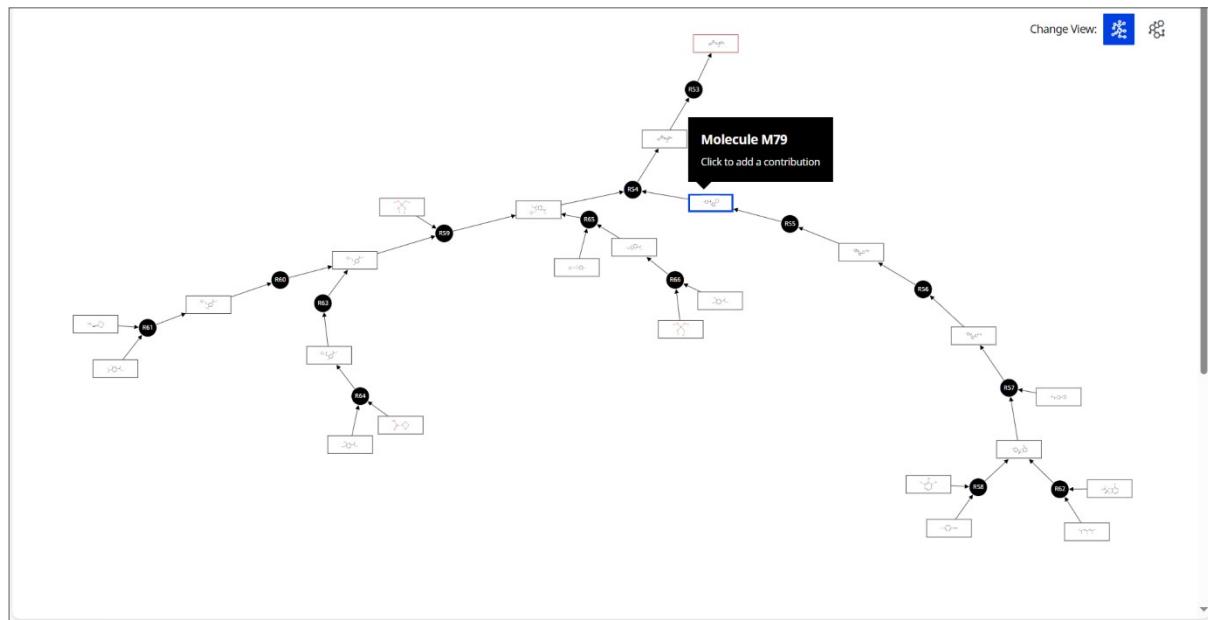
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ESI-1 – Digitisation of human generated ideas (digitalised brainstorm)

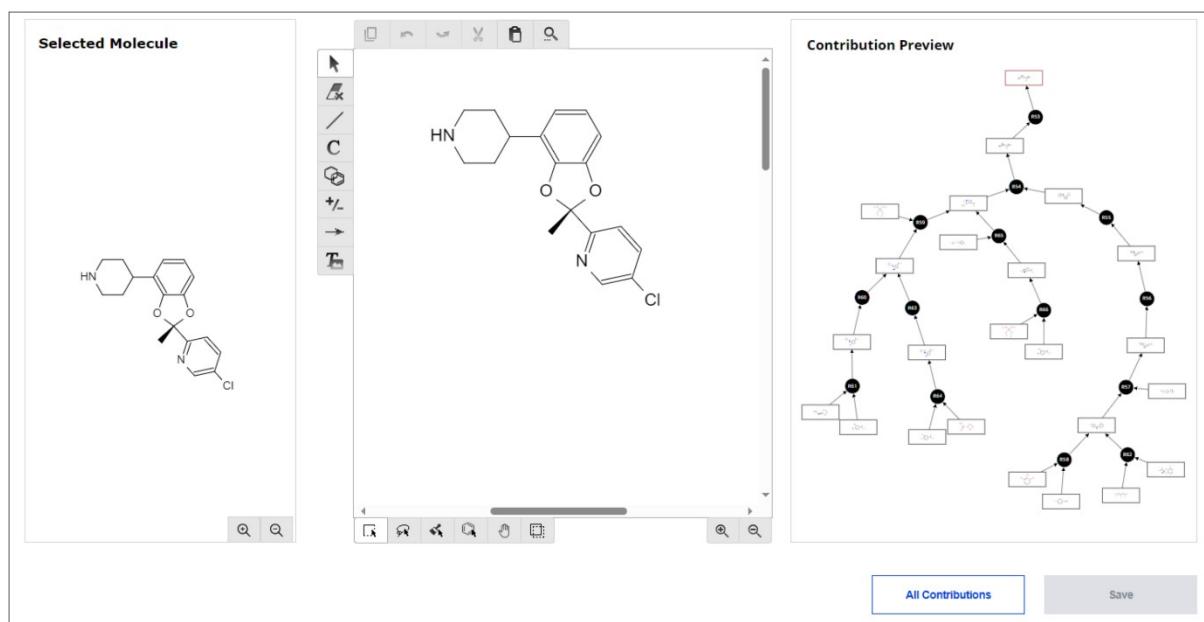
1.1 Target molecule properties

Synonyms	PF-07081532, Lotiglipron
SMILES	<chem>C[C@@]1(OC2=CC=CC(=C2O1)C3CCN(CC3)CC4=NC5=C(N4C[C@@H]6CCO6)C=C(C=C5)C(=O)O)C7=NC=C(C=C7)Cl</chem>
INCHI	InChI=1S/C31H31ClN4O5/c1-31(27-8-6-21(32)16-33-27)40-26-4-2-3-23(29(26)41-31)19-9-12-35(13-10-19)18-28-34-24-7-5-20(30(37)38)15-25(24)36(28)17-22-11-14-39-22/h2-8,15-16,19,22H,9-14,17-18H2,1H3,(H,37,38)/t22-,31-/m0/s1
INCHI Key	SVPYZAJTWFQTSU-UGDMGKLASA-N

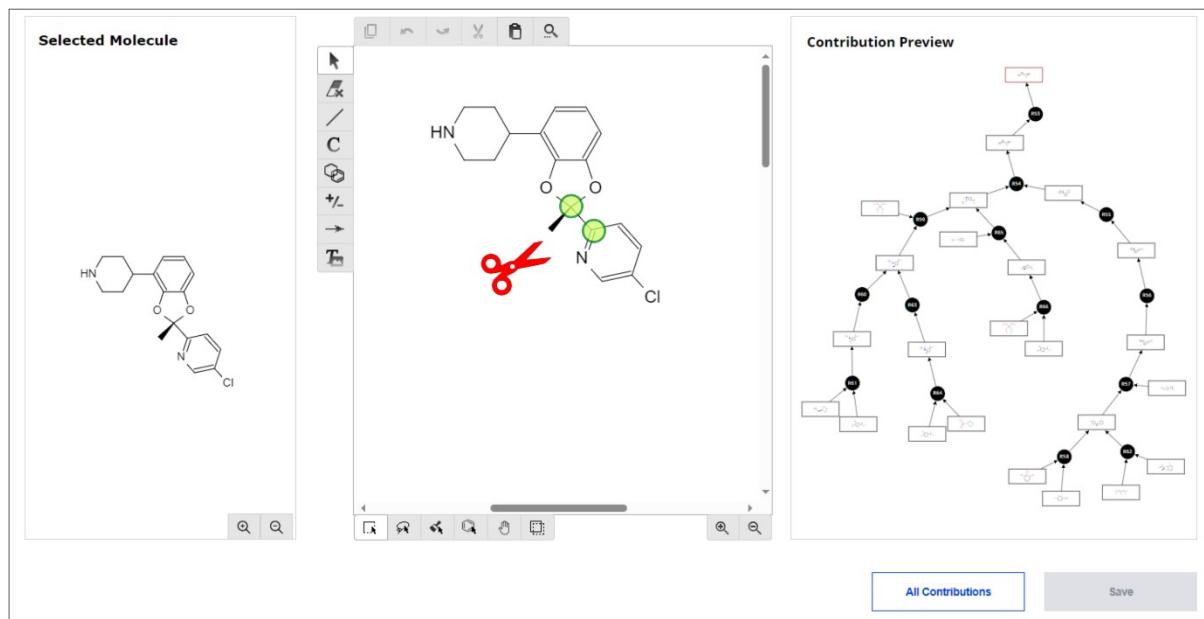
1.2 Initial network where contribution (disconnection) is made, and selection of the specific molecule node to be expanded. Specific molecule details for the entire network are provided in section 2.4.



1.3 Initial rendering of the molecule, with a digital user interface for a chemist to draw the transformation (centre of the picture)



1.4 Resulting chemical transformation to be introduced, generating two individual fragments from the initial selected molecule. This could include leaving groups or only main fragments.



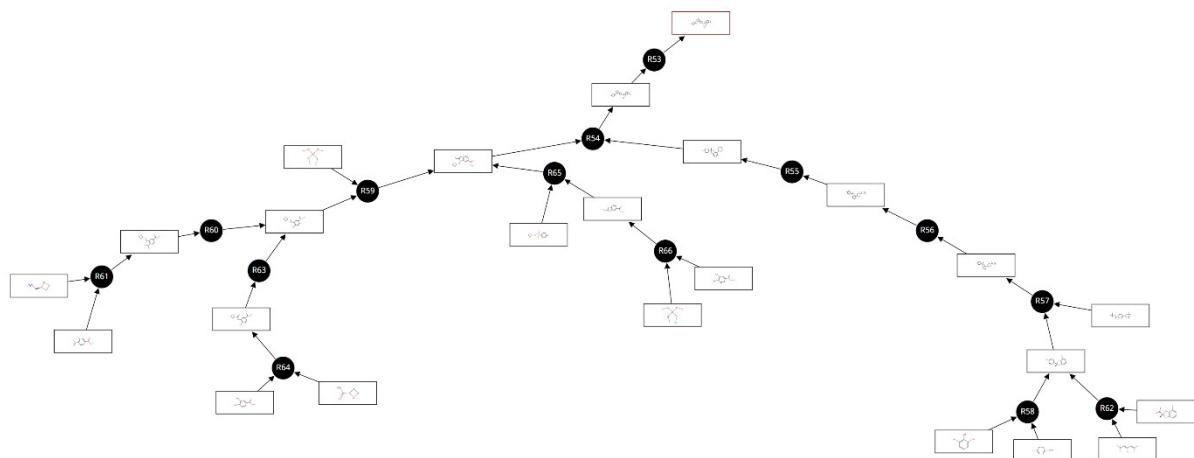
1.5 Summary of the contribution before submitting an entry. Metadata can be added considering chemist or contributor experiences. This can include background information searches to autocomplete information, such as literature references, safety data from modelling or published, etc. When this step is ready, data is written in the graph database supporting the defined data model (Figure 2). Form is displayed alongside to capture additional information, with fields aligned to SELECT criteria (or any other standard used).

Contribution Preview

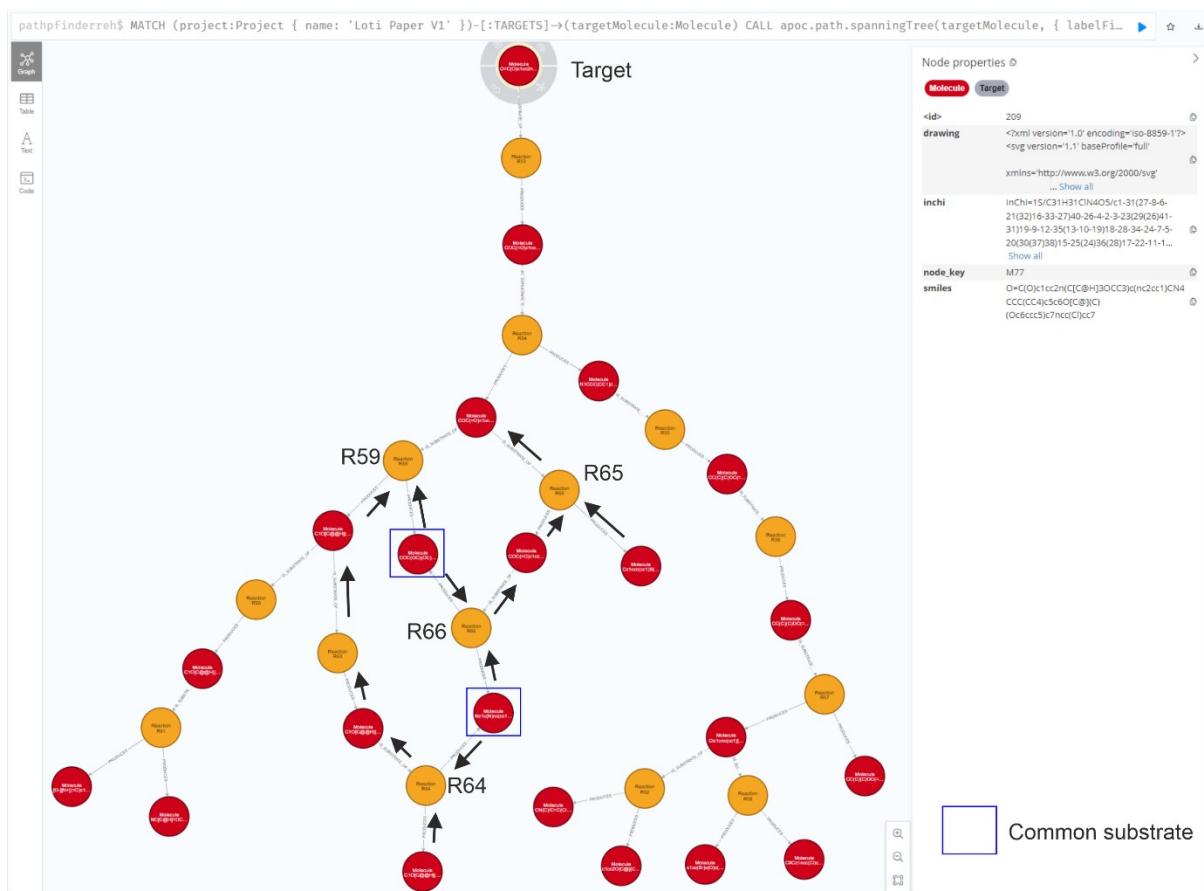
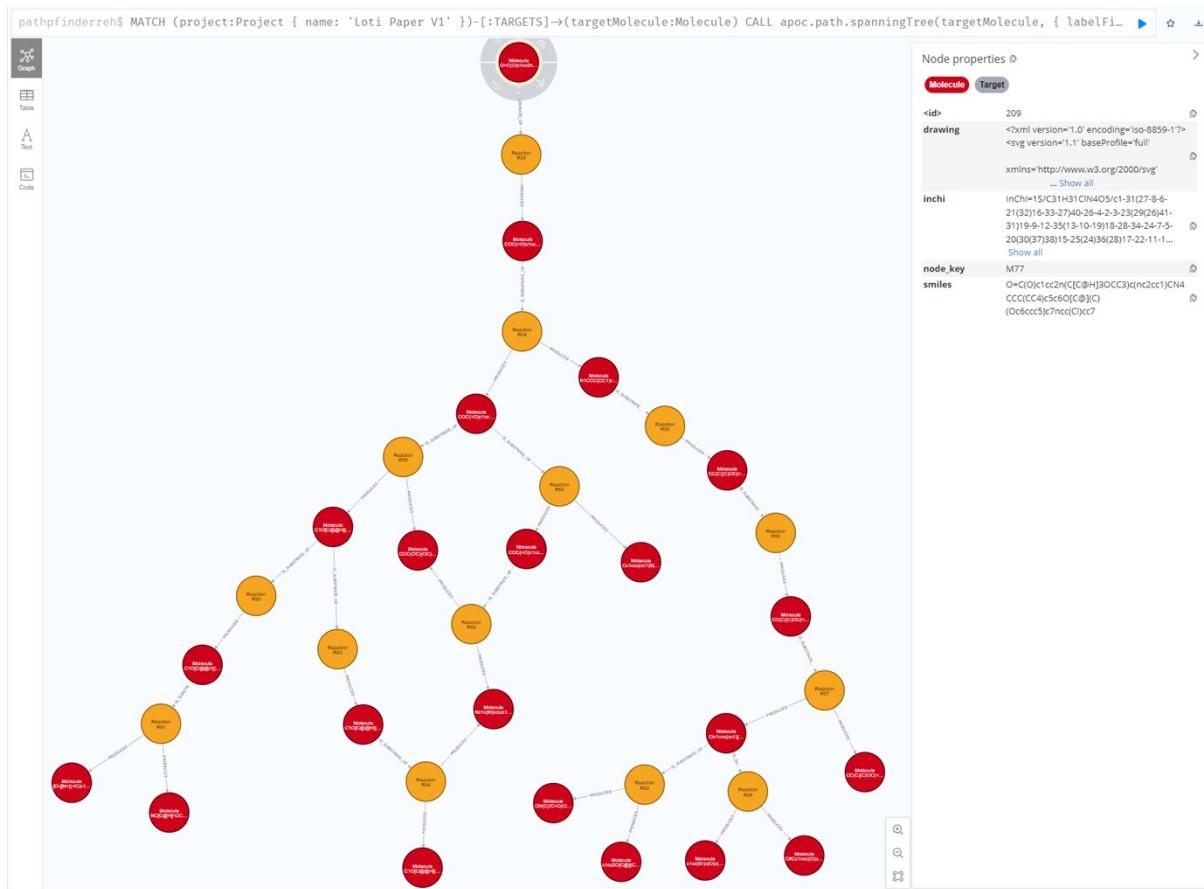
Add Details

Reaction Name	Reaction Type
<input type="text"/>	<input type="button" value="▼"/>
Notes	
<input type="text"/>	
Reference for Transformation	Conditions
<input type="text"/>	<input type="text"/>
Ratings 1 - Low 5 - High	
Similarity Of Ref To Compound	<input type="button" value="▼"/>
Scalability	<input type="button" value="▼"/>
Confidence Of Success	<input type="button" value="▼"/>

1.6 Network view with ideas already included into the graph database and accessible to anyone.

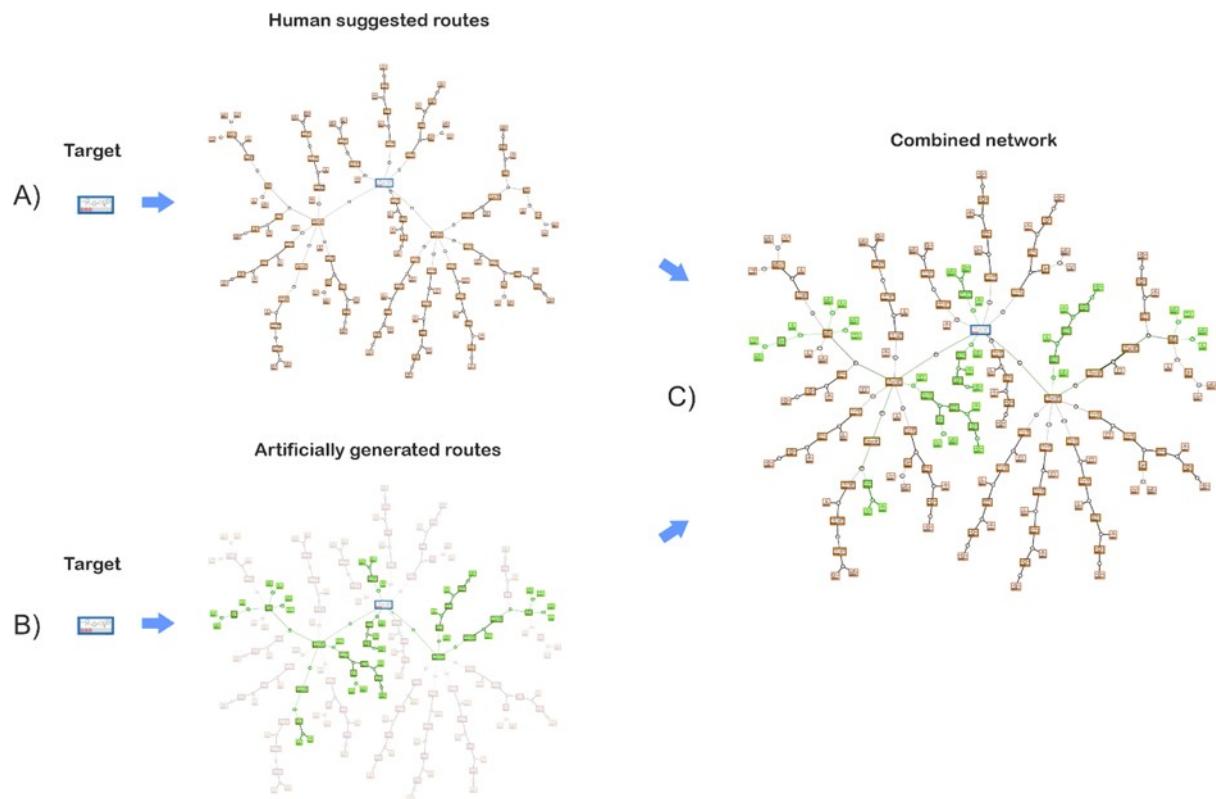


1.7 Same network shown in 1.6, visualised from the graph database directly (Neo4J backend). Original image above, image with indications below. Note the rendering differences. As molecules are unique on the graph database, the substrate feeding R59 and R66 appear only once in the database (same as substrate feeding R64 and R66). This creates a recurrence within the graph (see second image below with indications).



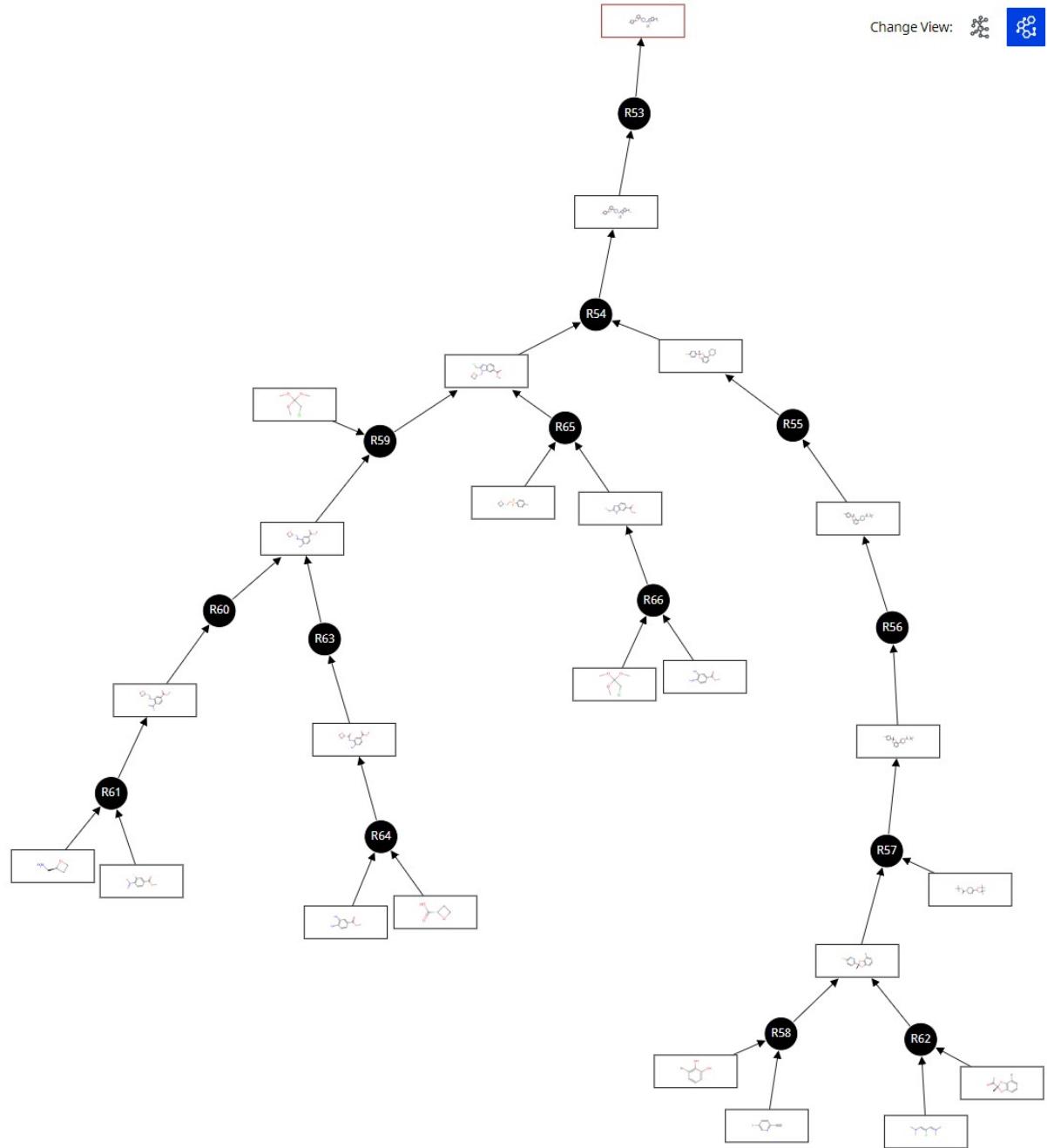
ESI-2 – Synthesis of ***Lotiglipron PF-07081532*** – full network produced.

2.1 Combining human with artificially generated routes – a theoretical network example is shown below. The same procedure was implemented for a real molecule, and it is described in the next sections.

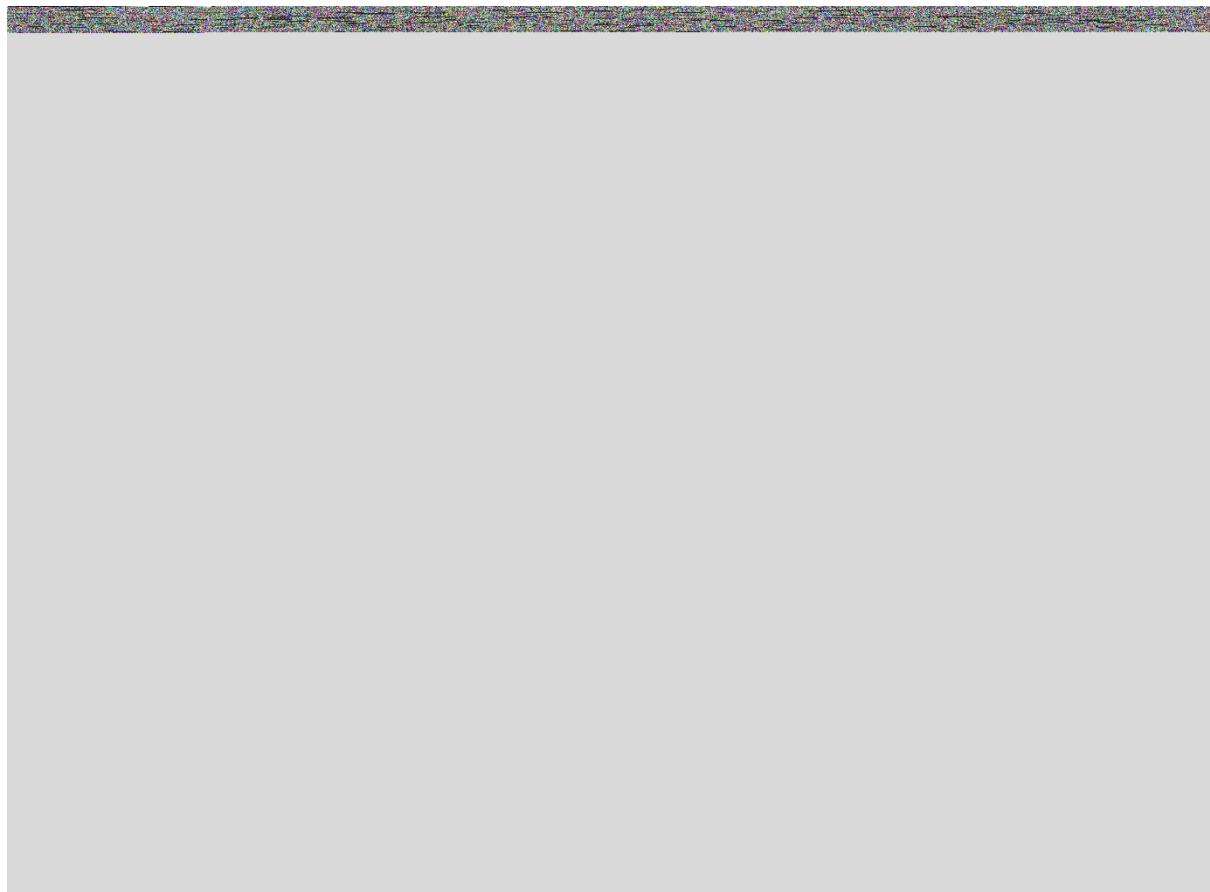


2.2 For the target molecule (*Lotiglipron*), some of the routes generated by human-led retrosynthesis analysis (following the procedure described in ESI-1) are shown in the image

below:



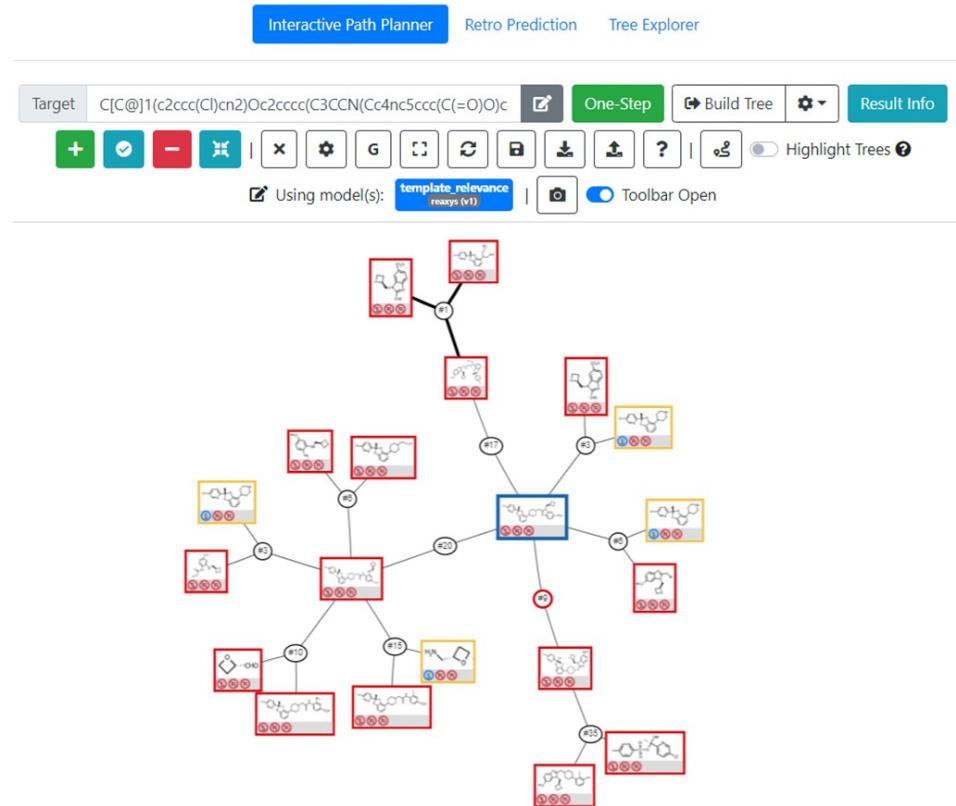
The equivalent graph stored in the graph database is also shown below:



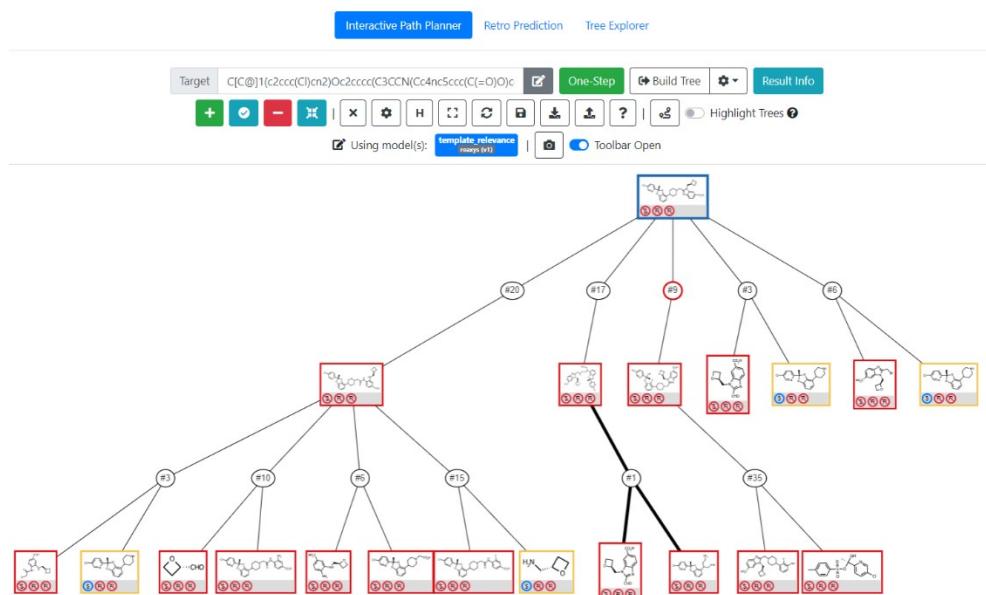
2.3 Some retrosynthesis routes generated by assisting predictive software (ASCKOS)

Details of all the routes digitally suggested are presented in section 3 (ESI-3). A summary figure with the routes filtered from ASCKOS is presented in the graph below.

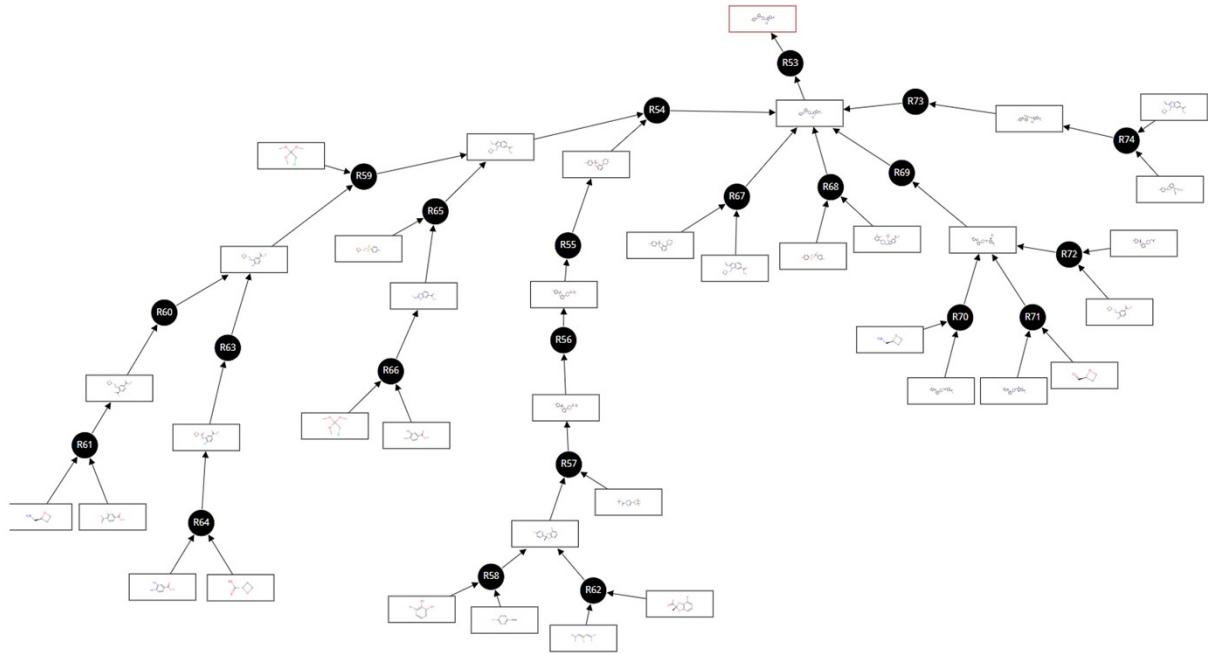
Network view:



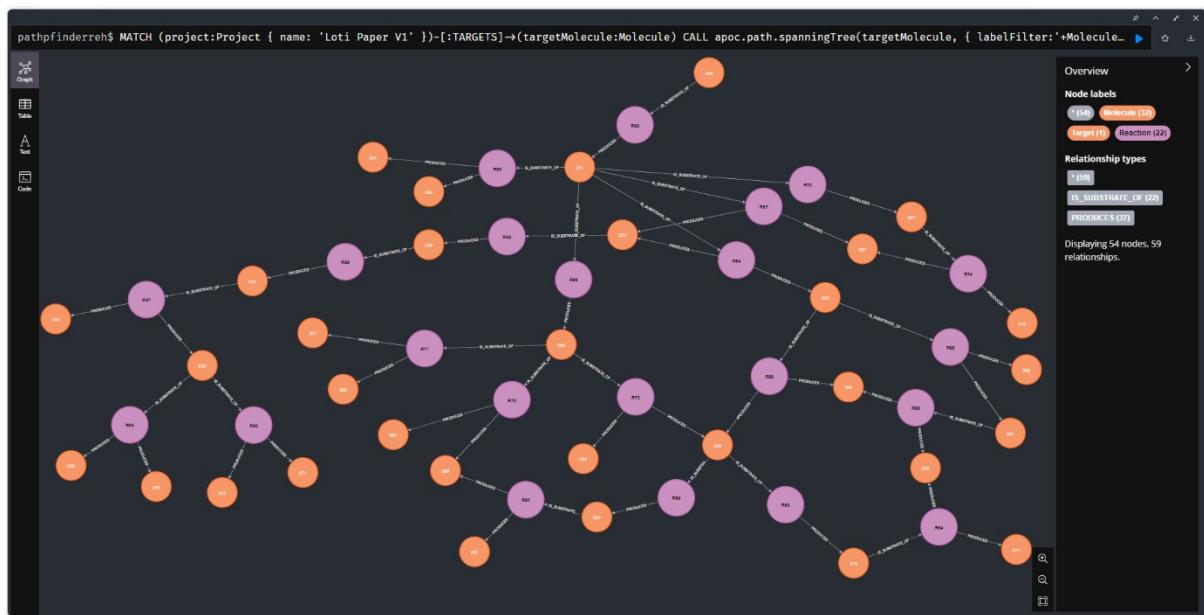
Alternative tree view:



2.4 Combined network – merging human with synthetically generated ideas for Lotiglipron

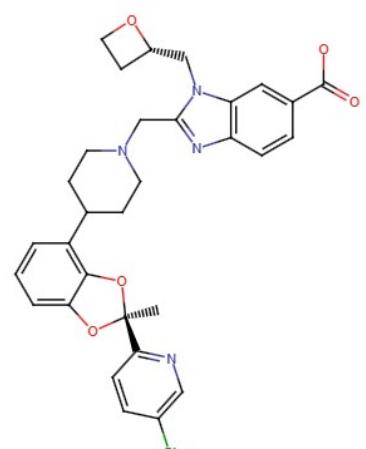
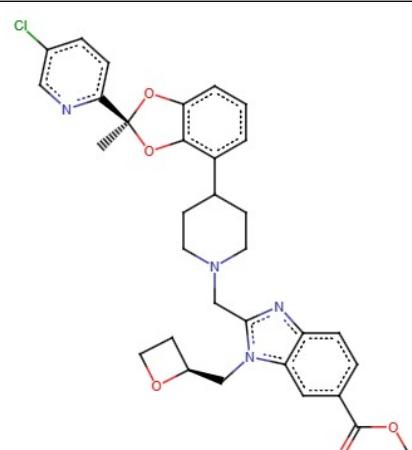
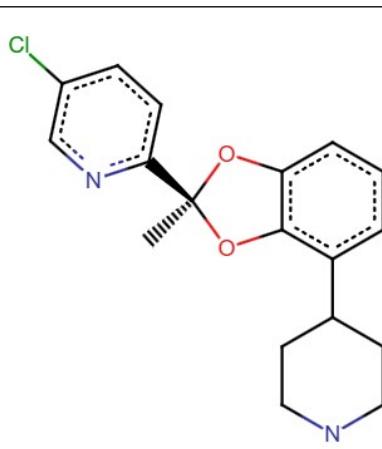
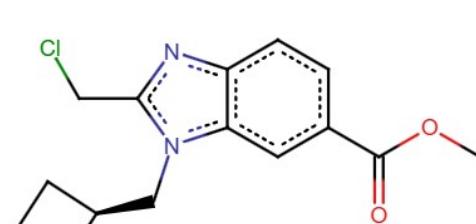


The corresponding graph database representation (native Neo4J) is also shown below:

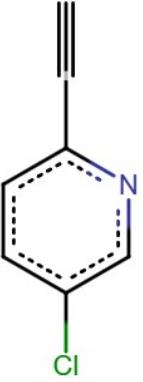
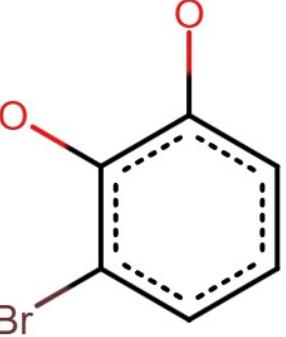
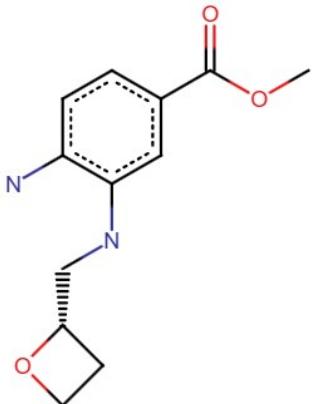
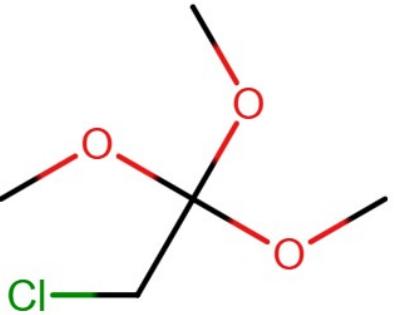


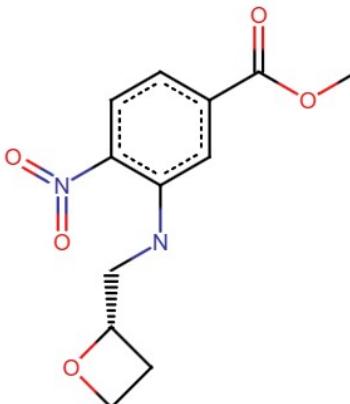
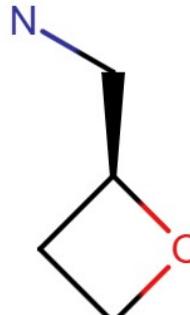
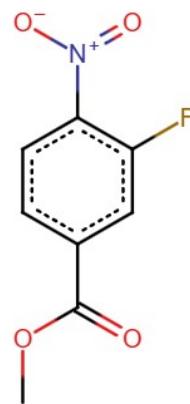
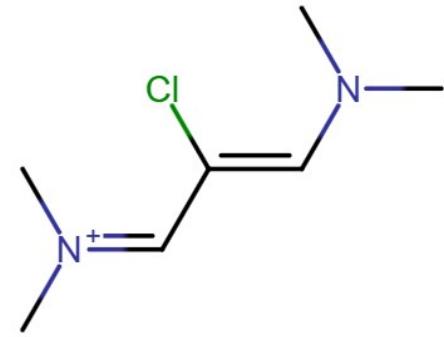
Specific molecule details are provided in the table below:

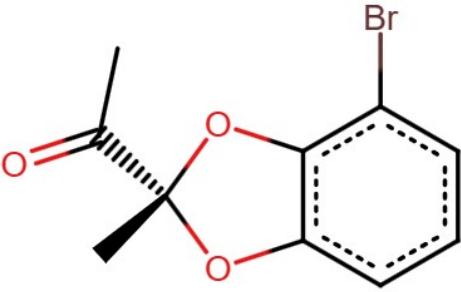
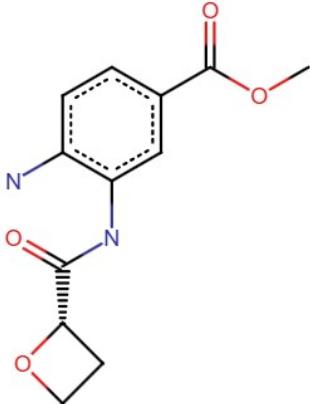
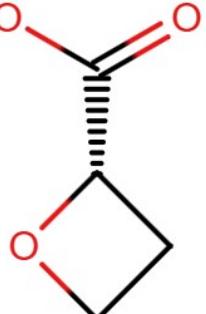
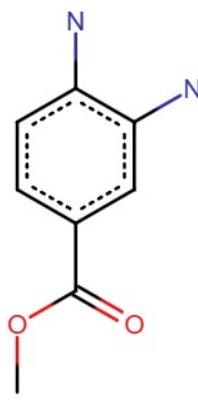
Molecule ID (picture display)	Node_Key Neo4j	Smiles	
-------------------------------	-------------------	--------	--

209	M77	<chem>C[C@@]1(OC2=CC=CC(=C2O1)C3CCN(CC3)CC4=NC5=C(N4C[C@H]6CCO6)C=C(C=C5)C(=O)O)C7=NC=C(C=C7)Cl</chem>	
211	M78	<chem>COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7</chem>	
214	M79	<chem>N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4</chem>	
215	M80	<chem>COC(=O)c1cc2n(C[C@H]3OCC3)c(Cl)nc2cc1</chem>	

218	M81	<chem>CC(C)(C)OC(=O)N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4</chem>	
221	M82	<chem>CC(C)(C)OC(=O)N1CCC(=CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4</chem>	
224	M84	<chem>CC(C)(C)OC(=O)N1CCC(=CC1)B2OC(C)(C)C(C)(O2)C</chem>	
225	M83	<chem>Clc1ncc(cc1)[C@]2(C)Oc3c(O2)cccc3Br</chem>	

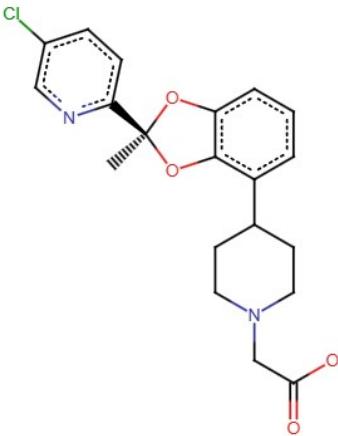
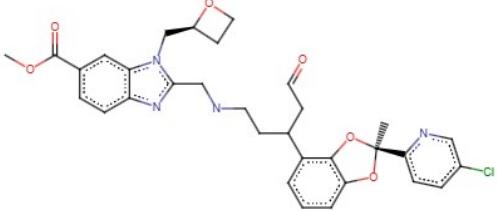
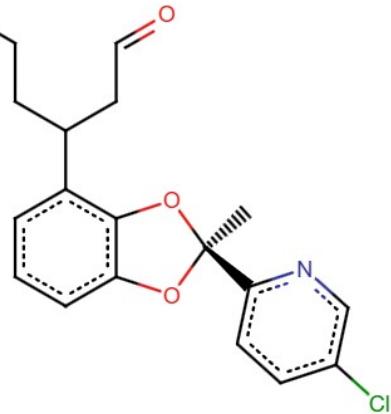
255	M85	C#Cc1ncc(Cl)cc1	
256	M86	c1cc(Br)c(O)c(c1)O	
259	M87	C1O[C@@H](C1)CNc2c(N)ccc(c2)C(=O)OC	
260	M88	COC(OC)(OC)CCl	

263	M89	<chem>C1O[C@@H](C1)CNc2c(N(=O)=O)ccc(c2)C(=O)OC</chem>	
266	M90	<chem>NC[C@H]1OCC1</chem>	
267	M91	<chem>[O-][N+](=O)c1c(F)cc(cc1)C(=O)OC</chem>	
270	M92	<chem>CN(C)/C=C(Cl)/C=[N+](C)C</chem>	

271	M93	<chem>c1cc2O[C@](C)(C(=O)C)Oc2c(c1)Br</chem>	
274	M94	<chem>C1O[C@@H](C1)C(=O)Nc2c(N)cc(c2)C(=O)OC</chem>	
277	M95	<chem>C1O[C@@H](C1)C(O)=O</chem>	
278	M96	<chem>Nc1c(N)cc(cc1)C(=O)OC</chem>	

281	M97	<chem>CC(=O)c1cc2c(cc1)nc([nH]2)C</chem>	
282	M98	<chem>Cc1ccc(cc1)S(=O)(=O)OC[C@H]2OCC2</chem>	
287	M99	<chem>CC(=O)c1cc2n(C[C@H]3OCC3)c(C=O)nc2cc1</chem>	
290	M100	<chem>Fc1c(O)c(ccc1)C2CCN(CC2)Cc3n(C[C@H]4OCC4)c5c(n3)ccc(c5)C(=O)OC</chem>	
291	M101	<chem>Cc1ccc(cc1)S(=O)(=O)O[C@@@](C)(O)c2ncc(Cl)cc2</chem>	

294	M102	<chem>C1O[C@@H](C1)CNc2c(cc(c2)C(OC)=O)NC(=O)CN3CCCC(CC3)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)cc6</chem>	
297	M103	<chem>COc1cc(F)c(cc1)NC(=O)CN2CCC(CC2)c3c4O[C@](C)(Oc4ccc3)c5ncc(Cl)cc5</chem>	
300	M104	<chem>O=C[C@H]1OCC1</chem>	
301	M105	<chem>COc1cc(N)cc(cc1)NC(=O)CN2CCC(CC2)c3c4O[C@](C)(Oc4ccc3)c5ncc(Cl)cc5</chem>	

304	M106	O=C(O)CN1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4	
307	M107	COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CNCCCC(CC=O)c4c5O[C@](C)(C)(Oc5ccc4)c6ncc(Cl)cc6	
310	M108	NCCCC(CC=O)c1c2O[C@](C)(Oc2ccc1)c3ncc(Cl)cc3	

2.5 Individual routes identified.

An algorithm transversed the network automatically identifying 12 routes. Details are shown below and as seen directly from the user interface developed. A form is displayed alongside each route to capture route data and aligned to the SELECT criteria.

Route 001

```

graph LR
    R53[ ] --> R54[ ]
    R54 --> R55[ ]
    R54 --> R56[ ]
    R55 --> R56
    R55 --> R57[ ]
    R56 --> R57
    R56 --> R58[ ]
    R57 --> R58
    R57 --> R59[ ]
    R58 --> R59
    R58 --> R60[ ]
    R59 --> R60
    R59 --> R61[ ]
    R60 --> R61
    R61 --> R62[ ]
    R61 --> R63[ ]
    R62 --> R63
  
```

Critique Comments

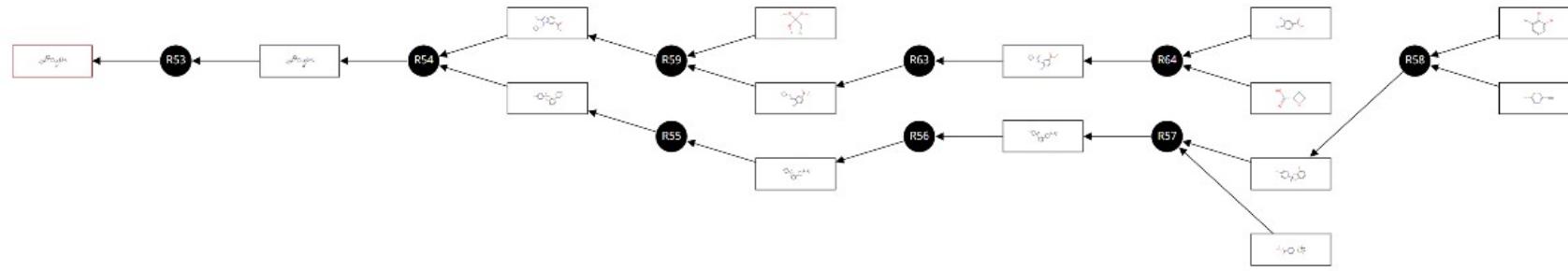
Notes	
Benefits	
Drawbacks	
Killers	

Route Assessment

Confidence Of Success		Value
Priority		Status
Impact	Steps	Convergency
Evaluation Team		Cybernetic

Route 001 Assessment Item

Back Save



Critique Comments

Notes

Benefits

Demands

Kill Sup

Focus Points

Route Assessment

Confidence Of Success

Value

Priority

Status

HQPS

Stops

Emergency

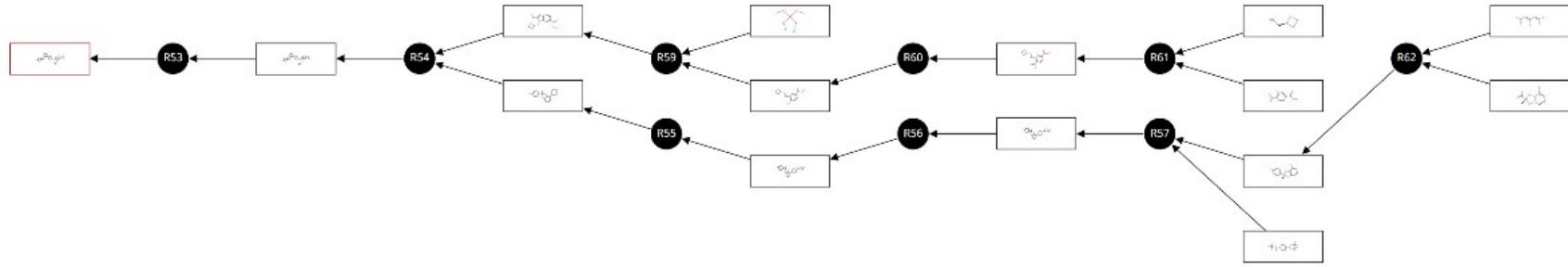
Cryogenic

Evaluation Team

Back

Save

Route 002 Assessment Item



Critique Comments

Notes

Benefits

Downsides

Kill Spur

Focus Points

Route Assessment

Confidence Of Success

Value

Priority

Status

HIPPOs

Stress

Emergency

Cybersecurity

Evaluation Team

Route 003 Assessment Item

Back

Save

Critique Comments

Notes

Benefits

Damages

Kill Stop

Route Assessment

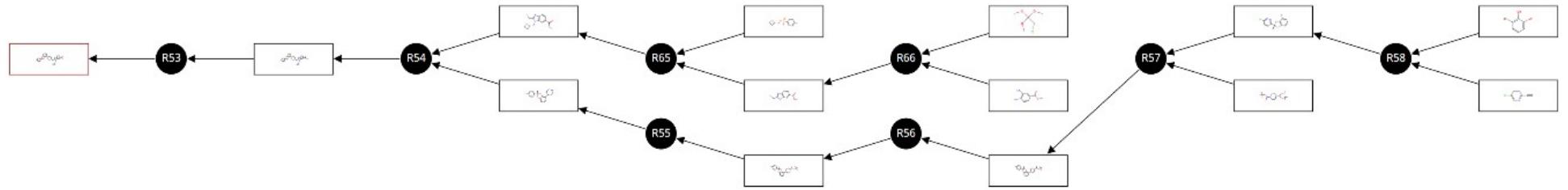
Confidence of success	Value
+	+

Priority	Score
+	+

Hazard	Status	Emergency	Cryogenic
+	+	+	+

Evaluation Team
+

Route 004 Assessment Form
Back
Save



Critique Comments

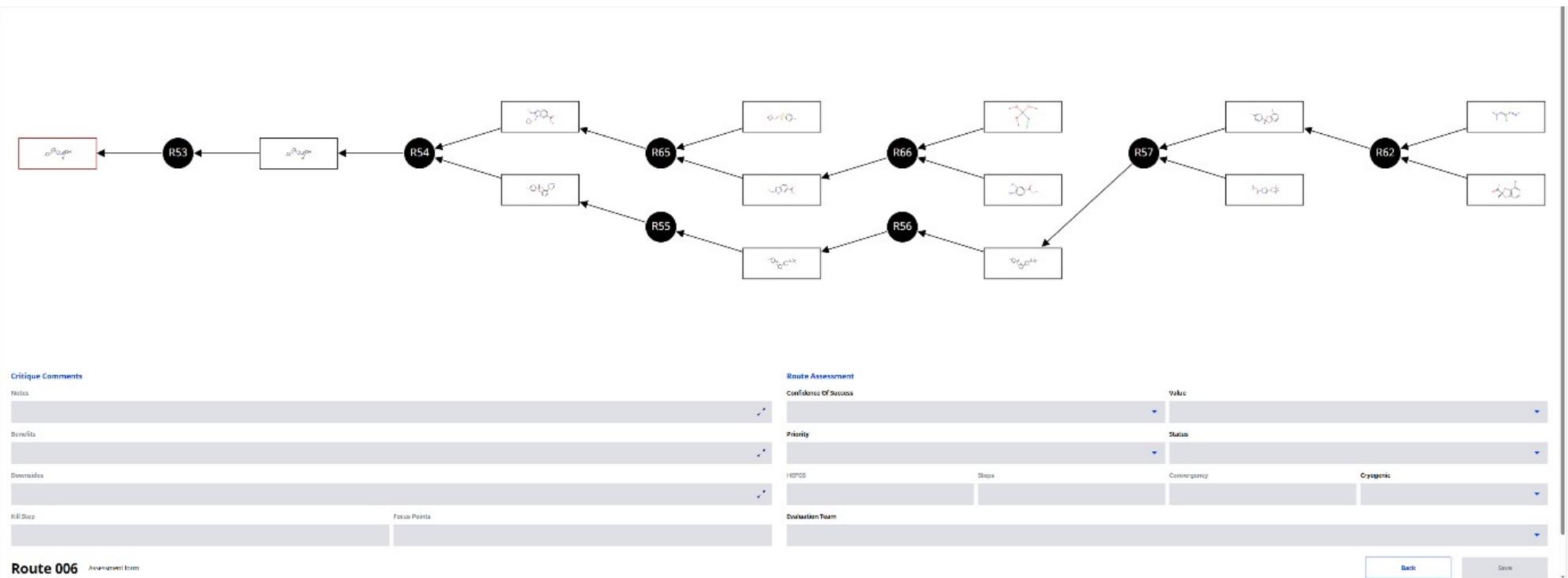
Notes	
General	
Comments	
Kill Step	

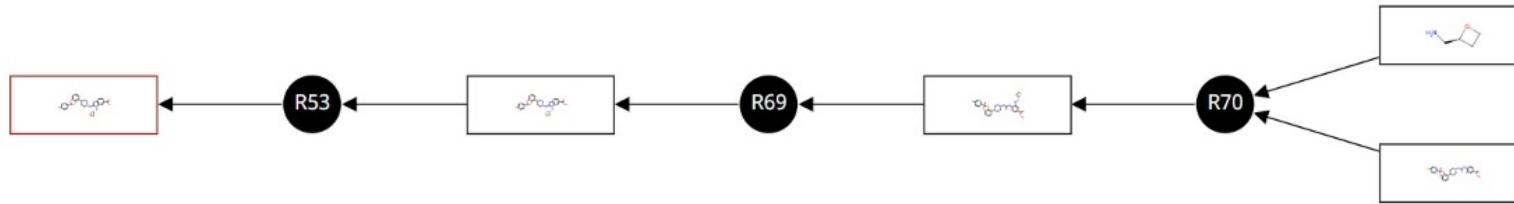
Route Assessment

Confidence Of Success	Value
Priority	Status
Hazard	Steps
Concavity	Cyberpic
Rotation Team	

Route 005 Assessment Form

[Back](#) [Save](#)





Critique Comments

Notes

Benefits

Downsides

Kill Step

Focus Points

Route Assessment

Confidence Of Success

Value

Priority

Status

HEFGS

Steps

Convergency

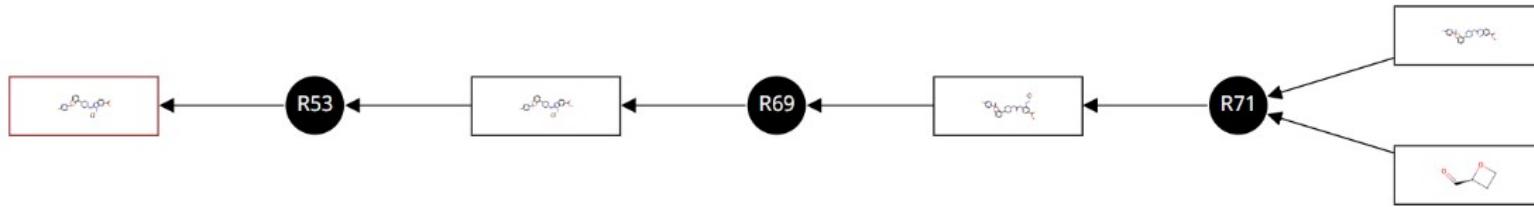
Cryogenic

Evaluation Team

Route 007 Assessment form

Back

Save



Critique Comments

Notes

Benefits

Downsides

Kill Step

Focus Points

Route Assessment

Confidence Of Success

Value

Priority

Status

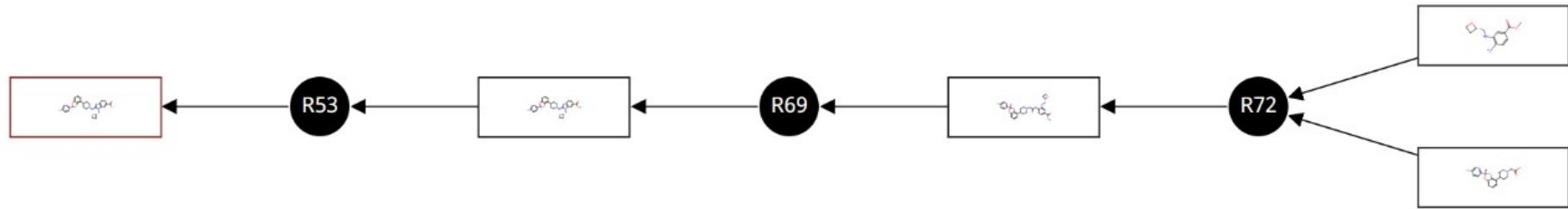
HEFGS

Steps

Convergency

Cryogenic

Evaluation Team



Critique Comments

Notes

Benefits

Downsides

Route Assessment

Confidence Of Success

Value

Priority

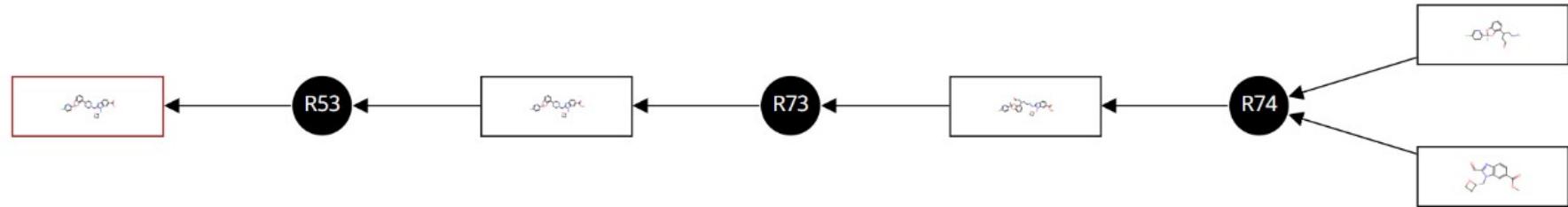
Status

HEFGS

Steps

Convergency

Cryogenic



Critique Comments

Notes

Benefits

Downsides

Kill Step

Focus Points

Route Assessment

Confidence Of Success

Value

Priority

Status

HEFGS

Steps

Convergency

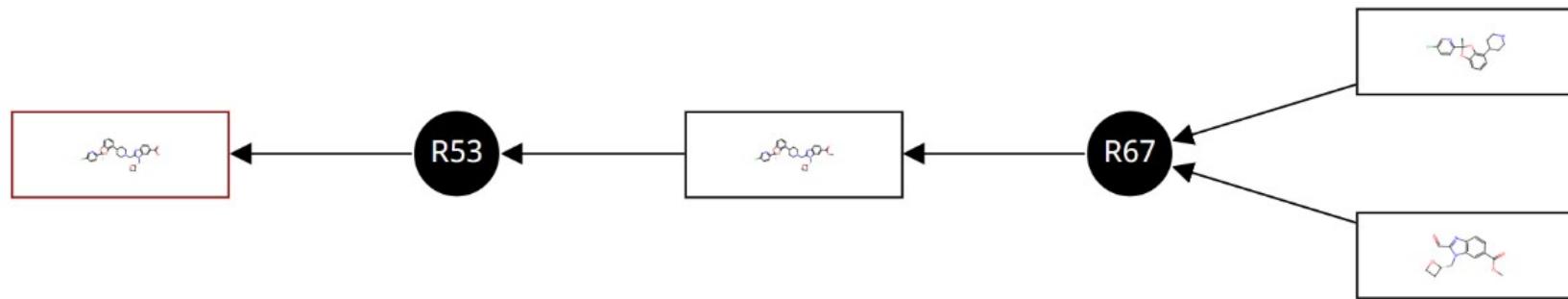
Cryogenic

Evaluation Team

Route 010 Assessment form

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[Save](#)



Critique Comments

Notes

Benefits

Downsides

Kill Step

Focus Points

Route Assessment

Confidence Of Success

Value

Priority

Status

HEFGS

Steps

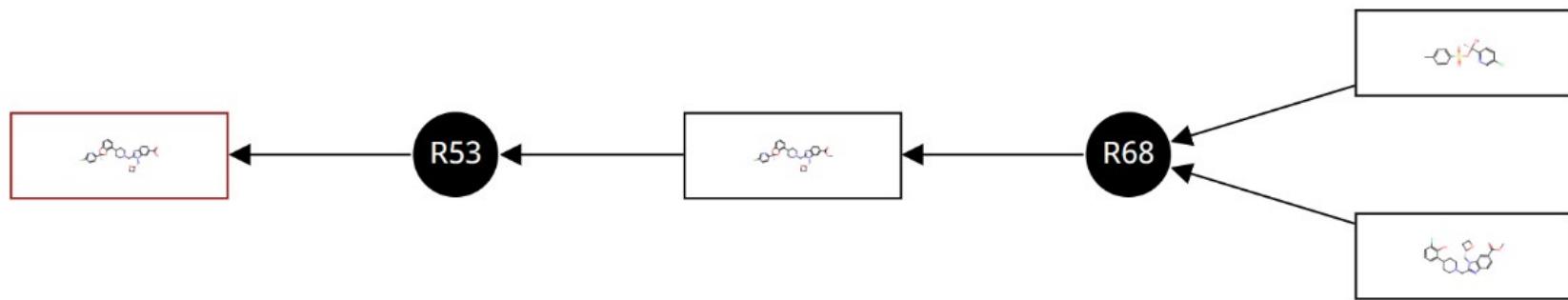
Convergency

Cryogenic

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Critique Comments

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Kill Step

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Route Assessment

Confidence Of Success

Value

Priority

Status

HEFGS

Steps

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Evaluation Team

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ESI-3 – Idea suggestions obtained from predictive software (ASKCOS)

3.1 The target molecule (Lotiglipron) was introduced into ASKCOS software to obtain ideas synthetically generated. Details of the idea search are omitted but they can be found on the MIT website (open to the public)

3.2 ASKCOS model settings used.

ASKCOS stand for Automated System for Knowledge-Based Continuous Organic Synthesis. It is a software output created by the Machine Learning for Pharmaceutical Discovery and Synthesis Consortium (MLPDS), a collaboration between the pharmaceutical and biotechnology industries (including Pfizer) and the departments of Chemical Engineering, Chemistry, and Computer Science at the Massachusetts Institute of Technology (MIT, United States). Additional information available at <https://mlpds.mit.edu/>

- Predictions were generated using Pfizer-ASKCOS (private consortia member version)
- Public version generates similar outputs, and it is available at <https://askcos.mit.edu/>

The model settings used for a basic retrosynthesis search are displayed below:

Settings

Global Settings

Highlight changed atoms

Align node images to target

Align precursors to product

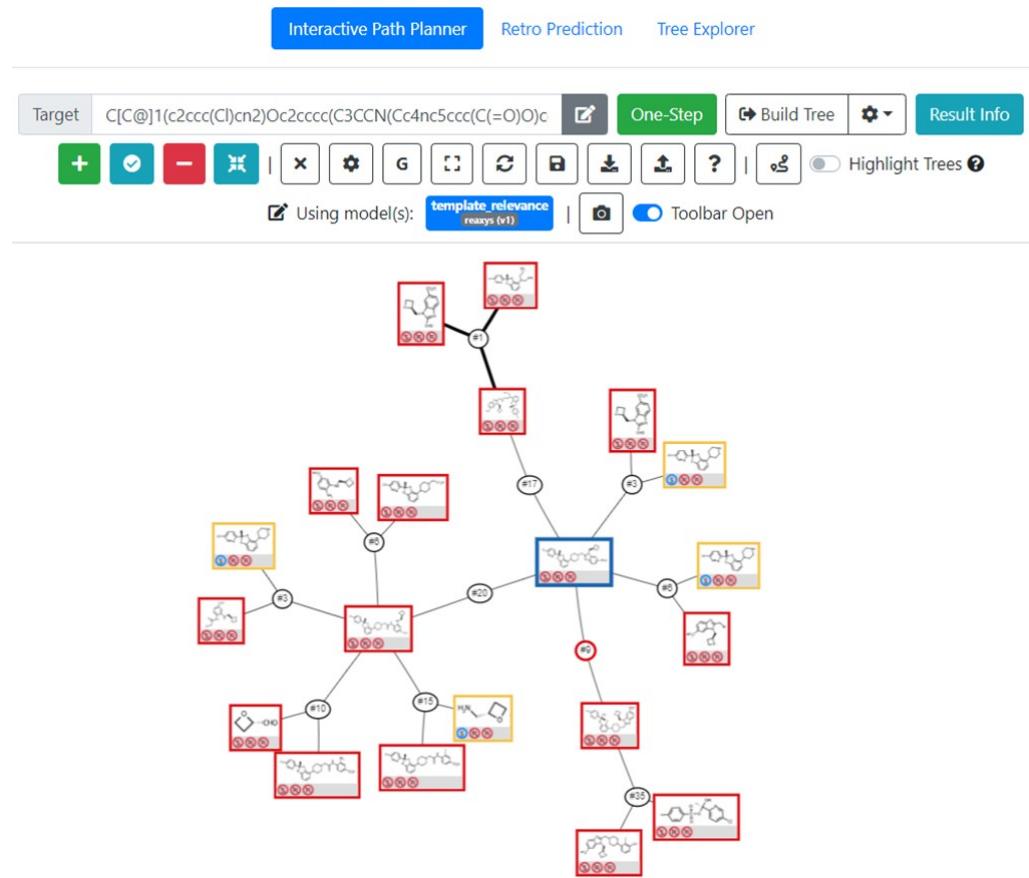
Top-N result to add to graph

Strategy Plan

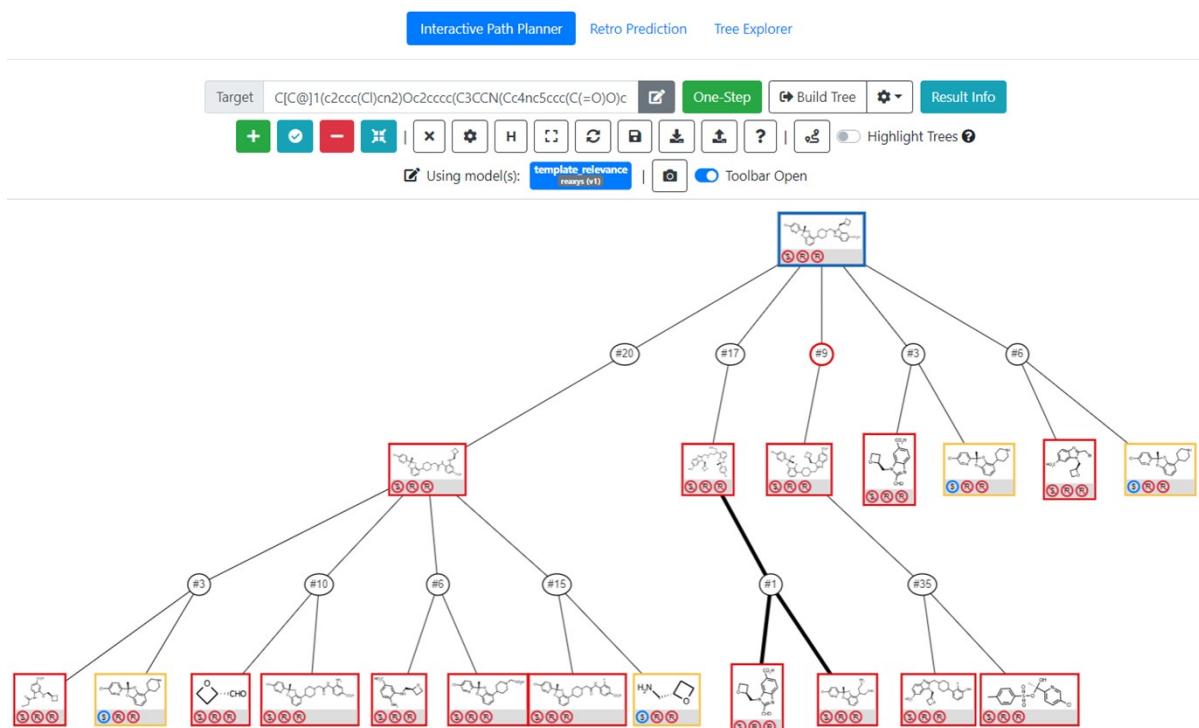
Strategy 1	
<input checked="" type="checkbox"/> Model	<input type="button" value="▼"/>
<input checked="" type="checkbox"/> Template prioritizers	<input type="button" value="Add +"/>
Template Set <small>i</small>	Version <small>i</small>
1 <input type="button" value="▼"/> reaxys	1 <input type="button" value="▼"/> <input type="button" value="X"/>
<i>Note: Template attribute filters are not supported by the tree builder.</i>	
<input checked="" type="checkbox"/> Max. num. templates	<input type="text" value="1000"/>
<input checked="" type="checkbox"/> Max. cum. prob.	<input type="text" value="0.999"/>
<input checked="" type="checkbox"/> Precursor scoring	<input type="button" value="Relevance+Heuristic"/>
<input checked="" type="checkbox"/> Min. plausibility	<input type="text" value="0.1"/>
<input checked="" type="checkbox"/> Regio-selectivity model	<input type="button" value="QM-WLN"/>
<input checked="" type="checkbox"/> Apply regio-selectivity checking	<input checked="" type="checkbox"/>

3.3 Filtered routes obtained by human aided prediction.

The output from a basic search is shown as follow. Network view:



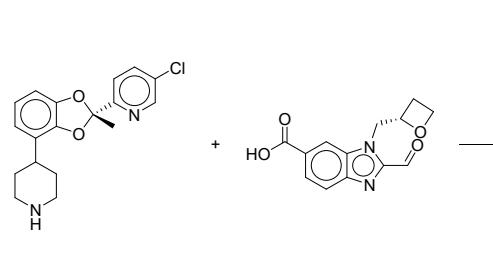
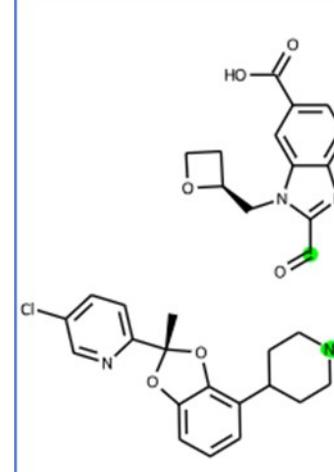
Alternative tree view:



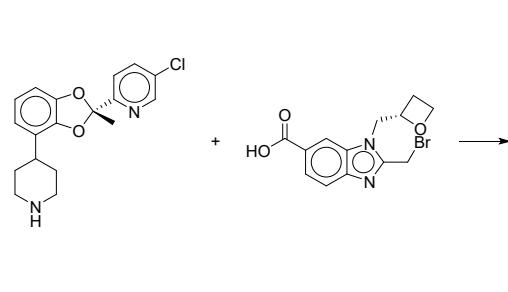
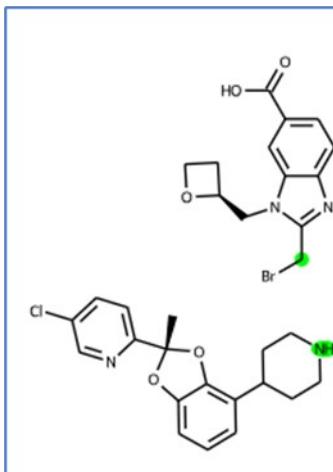
3.4 Description of the individual steps.

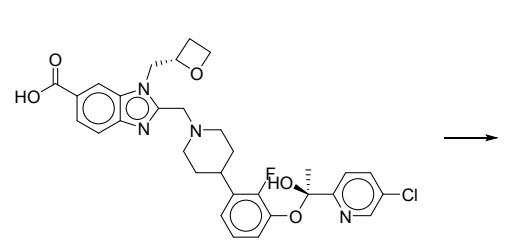
The disconnections shown in the trees above can be reproduced by using the images or reaction smiles provided in the tables below for each of the individual transformations. These details were used later to merge human and digitally generated ideas (Supplementary information 1 and 2).

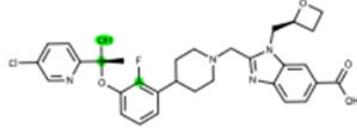
1st level disconnections from target

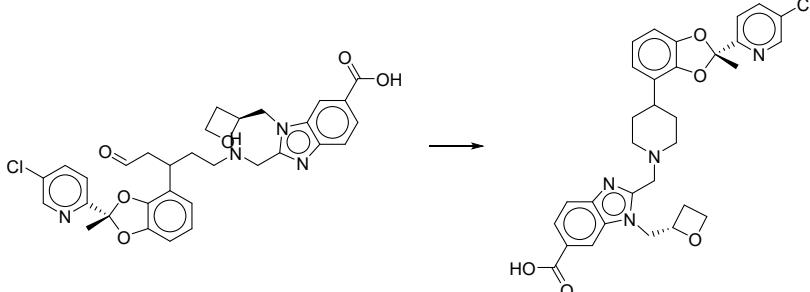
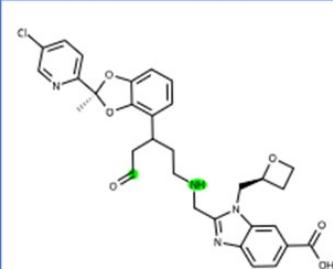
Reaction Node Number	3																
Reaction																	
Reaction SMILES	<chem>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCNCC3)c2O1.O=Cc1nc2ccc(C(=O)O)cc2n1C[C@@H]1CCO1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(Cc4nc5ccc(C(=O)O)cc5n4C[C@@H]4CCO4)CC3)c2O1</chem>																
ASKCOS output	<div style="display: flex; align-items: center;"> <div style="flex: 1; text-align: center;">  </div> <div style="flex: 1; border: 1px solid blue; padding: 5px; margin-left: 10px;"> <table border="1"> <tbody> <tr> <td>Rank</td> <td>#3</td> </tr> <tr> <td>Score</td> <td>-7000</td> </tr> <tr> <td>Synthetic complexity</td> <td>4.1</td> </tr> <tr> <td># Examples</td> <td>7680</td> </tr> <tr> <td>Template rank</td> <td>4</td> </tr> <tr> <td>Template score</td> <td>0.025</td> </tr> <tr> <td>Plausibility</td> <td>1.0</td> </tr> <tr> <td>Reaction cluster</td> <td>Reaction Cluster #3</td> </tr> </tbody> </table> </div> </div>	Rank	#3	Score	-7000	Synthetic complexity	4.1	# Examples	7680	Template rank	4	Template score	0.025	Plausibility	1.0	Reaction cluster	Reaction Cluster #3
Rank	#3																
Score	-7000																
Synthetic complexity	4.1																
# Examples	7680																
Template rank	4																
Template score	0.025																
Plausibility	1.0																
Reaction cluster	Reaction Cluster #3																

Reaction Node Number	6
----------------------	---

Reaction																	
Reaction SMILES	<chem>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCNCC3)c2O1.O=C(O)c1ccc2n c(CBr)n(C[C@@H]3CCO3)c2c1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(Cc4nc5ccc(C(=O)O)cc5n4C[C@@H]4CCO4)CC3)c2O1</chem>																
ASKCOS output	<div style="display: flex; align-items: center;"> <div style="flex: 1; text-align: center;">  </div> <table border="1" style="margin-left: 20px; border-collapse: collapse; text-align: center;"> <tr> <td>Rank</td> <td>#6</td> </tr> <tr> <td>Score</td> <td>-65000</td> </tr> <tr> <td>Synthetic complexity</td> <td>4.1</td> </tr> <tr> <td># Examples</td> <td>5790</td> </tr> <tr> <td>Template rank</td> <td>8</td> </tr> <tr> <td>Template score</td> <td>0.0027</td> </tr> <tr> <td>Plausibility</td> <td>1.0</td> </tr> <tr> <td>Reaction cluster</td> <td>Reaction Cluster #4</td> </tr> </table> </div>	Rank	#6	Score	-65000	Synthetic complexity	4.1	# Examples	5790	Template rank	8	Template score	0.0027	Plausibility	1.0	Reaction cluster	Reaction Cluster #4
Rank	#6																
Score	-65000																
Synthetic complexity	4.1																
# Examples	5790																
Template rank	8																
Template score	0.0027																
Plausibility	1.0																
Reaction cluster	Reaction Cluster #4																

Reaction Node Number	9
Reaction	
Reaction SMILES	<chem>C[C@@](O)(Oc1cccc(C2CCN(Cc3nc4ccc(C(=O)O)cc4n3C[C@@H]3CCO3)CC2)c1F)c1ccc(Cl)cn1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(Cc4nc5ccc(C(=O)O)cc5n4C[C@@H]4CCO4)CC3)c2O1</chem>

ASKCOS output	 <table border="1"> <tr><td>Rank</td><td>#9</td></tr> <tr><td>Score</td><td>-740000</td></tr> <tr><td>Synthetic complexity</td><td>4.8</td></tr> <tr><td># Examples</td><td>3603</td></tr> <tr><td>Template rank</td><td>16</td></tr> <tr><td>Template score</td><td>7.9e-4</td></tr> <tr><td>Plausibility</td><td>0.26</td></tr> <tr><td>Reaction cluster</td><td>Reaction Cluster #6</td></tr> </table>	Rank	#9	Score	-740000	Synthetic complexity	4.8	# Examples	3603	Template rank	16	Template score	7.9e-4	Plausibility	0.26	Reaction cluster	Reaction Cluster #6
Rank	#9																
Score	-740000																
Synthetic complexity	4.8																
# Examples	3603																
Template rank	16																
Template score	7.9e-4																
Plausibility	0.26																
Reaction cluster	Reaction Cluster #6																

Reaction Node Number	17																
Reaction																	
Reaction SMILES	<chem>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C(CC=O)CCNCc3nc4ccc(C(=O)O)cc4n3C[C@@H]3CCO3)c2O1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(Cc4nc5ccc(C(=O)O)cc5n4C[C@@H]4CCO4)CC3)c2O1</chem>																
ASKCOS output	 <table border="1"> <tr><td>Rank</td><td>#17</td></tr> <tr><td>Score</td><td>-2100000</td></tr> <tr><td>Synthetic complexity</td><td>4.7</td></tr> <tr><td># Examples</td><td>3781</td></tr> <tr><td>Template rank</td><td>32</td></tr> <tr><td>Template score</td><td>2.8e-4</td></tr> <tr><td>Plausibility</td><td>1.0</td></tr> <tr><td>Reaction cluster</td><td>Reaction Cluster #7</td></tr> </table>	Rank	#17	Score	-2100000	Synthetic complexity	4.7	# Examples	3781	Template rank	32	Template score	2.8e-4	Plausibility	1.0	Reaction cluster	Reaction Cluster #7
Rank	#17																
Score	-2100000																
Synthetic complexity	4.7																
# Examples	3781																
Template rank	32																
Template score	2.8e-4																
Plausibility	1.0																
Reaction cluster	Reaction Cluster #7																

Reaction Node Number	20																
Reaction																	
Reaction SMILES	<chem>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(CC(=O)Nc4ccc(C(=O)O)c4NC[C@@H]4CCO4)CC3)c2O1>>C[C@]1(c2ccc(Cl)cn2)Oc2ccc(C3CCN(Cc4nc5ccc(C(=O)O)cc5n4C[C@@H]4CCO4)CC3)c2O1</chem>																
ASKCOS output	<table border="1"> <tbody> <tr> <td>Rank</td> <td>#20</td> </tr> <tr> <td>Score</td> <td>-6700000</td> </tr> <tr> <td>Synthetic complexity</td> <td>5.0</td> </tr> <tr> <td># Examples</td> <td>301</td> </tr> <tr> <td>Template rank</td> <td>62</td> </tr> <tr> <td>Template score</td> <td>9.1e-5</td> </tr> <tr> <td>Plausibility</td> <td>1.00</td> </tr> <tr> <td>Reaction cluster</td> <td>Reaction Cluster #8</td> </tr> </tbody> </table>	Rank	#20	Score	-6700000	Synthetic complexity	5.0	# Examples	301	Template rank	62	Template score	9.1e-5	Plausibility	1.00	Reaction cluster	Reaction Cluster #8
Rank	#20																
Score	-6700000																
Synthetic complexity	5.0																
# Examples	301																
Template rank	62																
Template score	9.1e-5																
Plausibility	1.00																
Reaction cluster	Reaction Cluster #8																

1st branch disconnections (from reaction node 9)

Reaction Node Number	35
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Reaction																	
Reaction SMILES	Cc1ccc(S(=O)(=O)O[C@@](C)(O)c2ccc(Cl)cn2)cc1.O=C(O)c1ccc2nc(CN3CCC(c4cccc(O)c4F)CC3)n(C[C@@H]3CCO3)c2c1>>C[C@@](O)(Oc1cccc(C2CCN(Cc3nc4ccc(C(=O)O)cc4n3C[C@@H]3CCO3)CC2)c1F)c1ccc(Cl)cn1																
ASKCOS output	<div style="display: flex; align-items: center;"> <div style="flex: 1; text-align: center;"> </div> <table border="1" style="margin-left: 20px; border-collapse: collapse; width: fit-content;"> <tbody> <tr> <td>Rank</td> <td>#35</td> </tr> <tr> <td>Score</td> <td>-40000000</td> </tr> <tr> <td>Synthetic complexity</td> <td>4.5</td> </tr> <tr> <td># Examples</td> <td>1520</td> </tr> <tr> <td>Template rank</td> <td>155</td> </tr> <tr> <td>Template score</td> <td>1.5e-5</td> </tr> <tr> <td>Plausibility</td> <td>1.00</td> </tr> <tr> <td>Reaction cluster</td> <td>Reaction Cluster #19</td> </tr> </tbody> </table> </div>	Rank	#35	Score	-40000000	Synthetic complexity	4.5	# Examples	1520	Template rank	155	Template score	1.5e-5	Plausibility	1.00	Reaction cluster	Reaction Cluster #19
Rank	#35																
Score	-40000000																
Synthetic complexity	4.5																
# Examples	1520																
Template rank	155																
Template score	1.5e-5																
Plausibility	1.00																
Reaction cluster	Reaction Cluster #19																

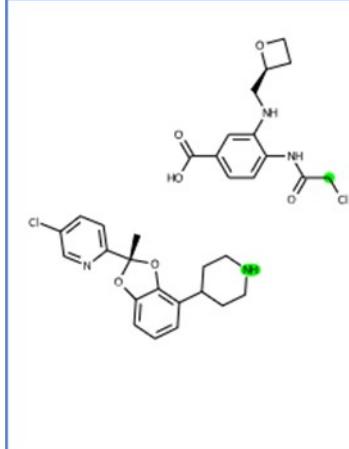
2nd branch disconnections (from reaction node 17)

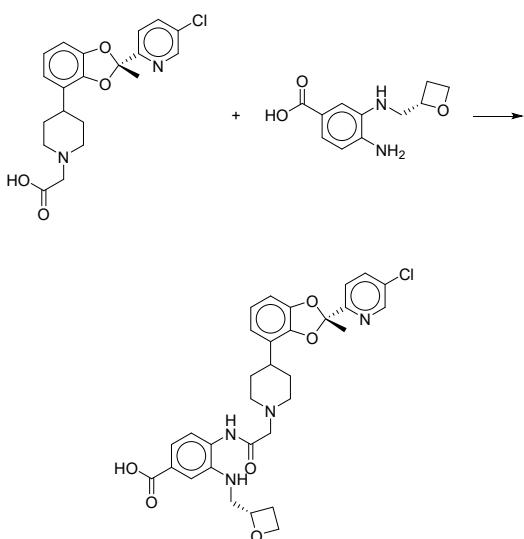
Reaction Node Number	1
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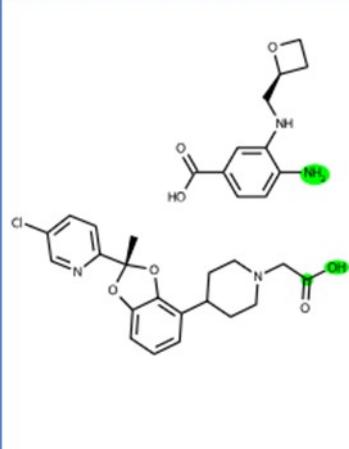
Reaction																	
Reaction SMILES	<chem>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C(CC=O)CCN)c2O1.O=Cc1nc2ccc(C(=O)O)cc2n1C[C@@H]1CCO1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C(CC=O)CCNCC3nc4ccc(C(=O)O)cc4n3C[C@@H]3CCO3)c2O1</chem>																
ASKCOS output	<div style="display: flex; align-items: center;"> <div style="flex: 1; text-align: center;"> </div> <table border="1" style="margin-left: 20px; border-collapse: collapse; text-align: center;"> <thead> <tr> <th>Rank</th> <th>#1</th> </tr> </thead> <tbody> <tr> <td>Score</td> <td>-1500</td> </tr> <tr> <td>Synthetic complexity</td> <td>3.8</td> </tr> <tr> <td># Examples</td> <td>8560</td> </tr> <tr> <td>Template rank</td> <td>2</td> </tr> <tr> <td>Template score</td> <td>0.29</td> </tr> <tr> <td>Plausibility</td> <td>1.00</td> </tr> <tr> <td>Reaction cluster</td> <td>Reaction Cluster #1</td> </tr> </tbody> </table> </div>	Rank	#1	Score	-1500	Synthetic complexity	3.8	# Examples	8560	Template rank	2	Template score	0.29	Plausibility	1.00	Reaction cluster	Reaction Cluster #1
Rank	#1																
Score	-1500																
Synthetic complexity	3.8																
# Examples	8560																
Template rank	2																
Template score	0.29																
Plausibility	1.00																
Reaction cluster	Reaction Cluster #1																

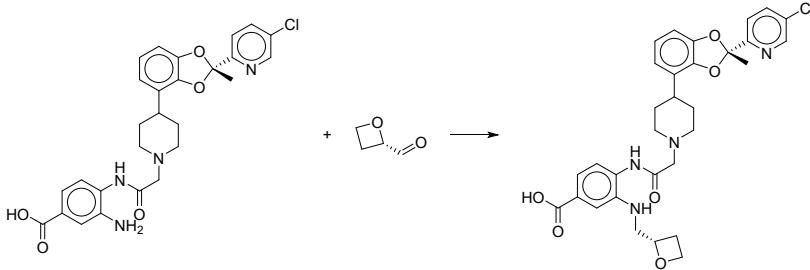
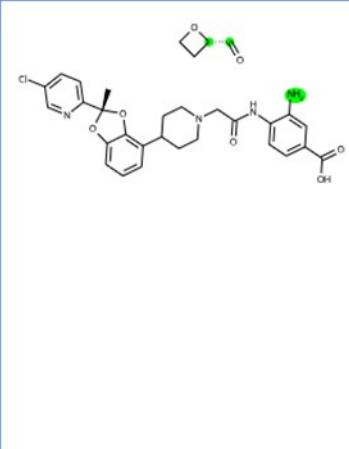
3rd branch disconnections (from reaction node 20)

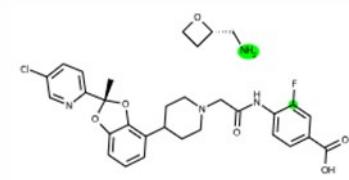
Reaction Node Number	3
Reaction	
Reaction SMILES	<chem>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCNCC3)c2O1.O=C(CCl)Nc1cc(C(=O)O)cc1NC[C@@H]1CCO1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(CC(=O)Nc4ccc(C(=O)O)cc4NC[C@@H]4CCO4)CC3)c2O1</chem>

	2O1																
ASKCOS output	 <table border="1" data-bbox="960 280 1341 729"> <tbody> <tr> <td>Rank</td> <td>#3</td> </tr> <tr> <td>Score</td> <td>-5800</td> </tr> <tr> <td>Synthetic complexity</td> <td>4.1</td> </tr> <tr> <td># Examples</td> <td>17781</td> </tr> <tr> <td>Template rank</td> <td>6</td> </tr> <tr> <td>Template score</td> <td>0.032</td> </tr> <tr> <td>Plausibility</td> <td>1.0</td> </tr> <tr> <td>Reaction cluster</td> <td>Reaction Cluster #2</td> </tr> </tbody> </table>	Rank	#3	Score	-5800	Synthetic complexity	4.1	# Examples	17781	Template rank	6	Template score	0.032	Plausibility	1.0	Reaction cluster	Reaction Cluster #2
Rank	#3																
Score	-5800																
Synthetic complexity	4.1																
# Examples	17781																
Template rank	6																
Template score	0.032																
Plausibility	1.0																
Reaction cluster	Reaction Cluster #2																

Reaction Node Number	6
Reaction	
Reaction SMILES	<chem>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(CC(=O)O)CC3)c2O1.Nc1ccc(C(=O)O)cc1NC[C@@H]1CCO1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(CC(=O)O)Cc4ccc(C(=O)O)cc4NC[C@@H]4CCO4)CC3)c2O1</chem>

ASKCOS output	 <table border="1"> <tbody> <tr><td>Rank</td><td>#6</td></tr> <tr><td>Score</td><td>-120000</td></tr> <tr><td>Synthetic complexity</td><td>4.7</td></tr> <tr><td># Examples</td><td>20006</td></tr> <tr><td>Template rank</td><td>14</td></tr> <tr><td>Template score</td><td>0.0038</td></tr> <tr><td>Plausibility</td><td>1.0</td></tr> <tr><td>Reaction cluster</td><td>Reaction Cluster #4</td></tr> </tbody> </table>	Rank	#6	Score	-120000	Synthetic complexity	4.7	# Examples	20006	Template rank	14	Template score	0.0038	Plausibility	1.0	Reaction cluster	Reaction Cluster #4
Rank	#6																
Score	-120000																
Synthetic complexity	4.7																
# Examples	20006																
Template rank	14																
Template score	0.0038																
Plausibility	1.0																
Reaction cluster	Reaction Cluster #4																

Reaction Node Number	10																
Reaction	 <p>Reaction SMILES: C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(CC(=O)Nc4ccc(C(=O)O)c4N)CC3)c2O1.O=C[C@@H]1CCO1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(CC(=O)Nc4ccc(C(=O)O)cc4NC[C@@H]4CCO4)CC3)c2O1</p>																
ASKCOS output	 <table border="1"> <tbody> <tr><td>Rank</td><td>#10</td></tr> <tr><td>Score</td><td>-580000</td></tr> <tr><td>Synthetic complexity</td><td>4.8</td></tr> <tr><td># Examples</td><td>1770</td></tr> <tr><td>Template rank</td><td>31</td></tr> <tr><td>Template score</td><td>9.0e-4</td></tr> <tr><td>Plausibility</td><td>1.0</td></tr> <tr><td>Reaction cluster</td><td>Reaction Cluster #5</td></tr> </tbody> </table>	Rank	#10	Score	-580000	Synthetic complexity	4.8	# Examples	1770	Template rank	31	Template score	9.0e-4	Plausibility	1.0	Reaction cluster	Reaction Cluster #5
Rank	#10																
Score	-580000																
Synthetic complexity	4.8																
# Examples	1770																
Template rank	31																
Template score	9.0e-4																
Plausibility	1.0																
Reaction cluster	Reaction Cluster #5																

Reaction Node Number	15																
Reaction																	
Reaction SMILES	<chem>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(CC(=O)Nc4ccc(C(=O)O)c4F)CC3)c2O1.NC[C@@H]1CCO1>>C[C@]1(c2ccc(Cl)cn2)Oc2cccc(C3CCN(CC(=O)Nc4ccc(C(=O)O)cc4NC[C@@H]4CCO4)CC3)c2O1</chem>																
ASKCOS output	<div style="display: flex; align-items: center;">  <table border="1" style="margin-left: 20px;"> <tbody> <tr> <td>Rank</td> <td>#15</td> </tr> <tr> <td>Score</td> <td>-1200000</td> </tr> <tr> <td>Synthetic complexity</td> <td>4.9</td> </tr> <tr> <td># Examples</td> <td>3664</td> </tr> <tr> <td>Template rank</td> <td>46</td> </tr> <tr> <td>Template score</td> <td>4.0e-4</td> </tr> <tr> <td>Plausibility</td> <td>1.0</td> </tr> <tr> <td>Reaction cluster</td> <td>Reaction Cluster #7</td> </tr> </tbody> </table> </div>	Rank	#15	Score	-1200000	Synthetic complexity	4.9	# Examples	3664	Template rank	46	Template score	4.0e-4	Plausibility	1.0	Reaction cluster	Reaction Cluster #7
Rank	#15																
Score	-1200000																
Synthetic complexity	4.9																
# Examples	3664																
Template rank	46																
Template score	4.0e-4																
Plausibility	1.0																
Reaction cluster	Reaction Cluster #7																

ESI-4 – Reproducing the graph database (using Neo4J) for Lotiglipron.

A set of queries is provided to reproduce the entire database presented in this paper.

For this, a public or community version of neo4j can be used as the graph database (accessible from <https://neo4j.com/>). From this link, you can create a new account and an initial database. After this, the Lotiglipron network can be created following the instructions below:

The queries in Cypher (Neo4J native language) are provided each inside a box. Follow them step by step, copying and pasting the queries as one chunk, directly on the Neo4J browser. Then, to run the query press the blue arrow at the top right of the browser.

1. Query 1 to create all molecule nodes (copy and paste it into the neo4j browser):

```
CREATE (n1:Molecule { node_key:'M77', INCHI: 'InChI=1S/C31H31ClN4O5/c1-31(27-8-6-21(32)16-33-27)40-26-4-2-3-23(29(26)41-31)19-9-12-35(13-10-19)18-28-34-24-7-5-20(30(37)38)15-25(24)36(28)17-22-11-14-39-22/h2-8,15-16,19,22H,9-14,17-18H2,1H3,(H,37,38)/t22-,31-/m0/s1',  
SMILES:'O=C(O)c1cc2n(C[C@H]3OCC3)c(ne2cc1)CN4CCCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''},  
(n2:Molecule { node_key:'M78', INCHI: 'InChI=1S/C32H33ClN4O5/c1-32(28-9-7-22(33)17-34-28)41-27-5-3-4-24(30(27)42-32)20-10-13-36(14-11-20)19-29-35-25-8-6-21(31(38)39-2)16-26(25)37(29)18-23-12-15-40-23/h3-9,16-17,20,23H,10-15,18-19H2,1-2H3/t23-,32-/m0/s1',  
SMILES:'COC(=O)c1cc2n(C[C@H]3OCC3)c(ne2cc1)CN4CCCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''},  
(n3:Molecule { node_key:'M79', INCHI: 'InChI=1S/C18H19ClN2O2/c1-18(16-6-5-13(19)11-21-16)22-15-4-2-3-14(17(15)23-18)12-7-9-20-10-8-12/h2-6,11-12,20H,7-10H2,1H3/t18-/m0/s1',  
SMILES:'N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''},  
(n4:Molecule { node_key:'M80', INCHI: 'InChI=1S/C14H15ClN2O3/c1-19-14(18)9-2-3-11-12(6-9)17(13(7-15)16-11)8-10-4-5-20-10/h2-3,6,10H,4-5,7-8H2,1H3/t10-/m0/s1',  
SMILES:'COC(=O)c1cc2n(C[C@H]3OCC3)c(Cl)nc2cc1', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''}, (n5:Molecule { node_key:'M81', INCHI: 'InChI=1S/C23H27ClN2O4/c1-22(2,3)30-21(27)26-12-10-15(11-13-26)17-6-5-7-18-20(17)29-23(4,28-18)19-9-8-16(24)14-25-19/h5-9,14-15H,10-13H2,1-4H3/t23-/m0/s1',  
SMILES:'CC(C)(C)OC(=O)N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''},  
(n6:Molecule { node_key:'M82', INCHI: 'InChI=1S/C23H25ClN2O4/c1-22(2,3)30-21(27)26-12-10-15(11-13-26)17-6-5-7-18-20(17)29-23(4,28-18)19-9-8-16(24)14-25-19/h5-10,14H,11-13H2,1-4H3/t23-/m0/s1',  
SMILES:'CC(C)(C)OC(=O)N1CCC(=CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''},  
(n7:Molecule { node_key:'M83', INCHI: 'InChI=1S/C13H9BrClNO2/c1-13(11-6-5-8(15)7-16-11)17-10-4-2-3-9(14)12(10)18-13/h2-7H,1H3/t13-/m0/s1', SMILES:'Clc1ncc(cc1)[C@]2(C)Oc3c(O2)cccc3Br', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''},  
(n8:Molecule { node_key:'M84', INCHI: 'InChI=1S/C16H28BNO4/c1-14(2,3)20-13(19)18-10-8-12(9-11-18)17-21-15(4,5)16(6,7)22-17/h8H,9-11H2,1-7H3',  
SMILES:'CC(C)(C)OC(=O)N1CCC(=CC1)B2OC(C)(C)C(C)(O2)C', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''}, (n9:Molecule { node_key:'M85', INCHI: 'InChI=1S/C7H4ClN/c1-2-7-4-3-6(8)5-9-7/h1,3-5H', SMILES:'C#Cc1ncc(Cl)cc1', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''}), (n10:Molecule { node_key:'M86', INCHI: 'InChI=1S/C6H5BrO2/c7-4-2-1-3-5(8)6(4)9/h1-3,8-9H', SMILES:'c1cc(Br)c(O)c(c1)O', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''}), (n11:Molecule { node_key:'M87', INCHI: 'InChI=1S/C12H16N2O3/c1-12(15)8-2-3-10(13)11(6-8)14-7-9-4-5-17-9/h2-3,6,9,14H,4-5,7,13H2,1H3/t9-/m0/s1',  
SMILES:'C1O[C@@H](C1)CNc2c(N)ccc(c2)C(=O)OC', Safety: '', Environmental: '', Legal: '', Economic: '', Control: '', Throughput: ''}), (n12:Molecule { node_key:'M88', INCHI: 'InChI=1S/C5H11ClO3/c1-7-5(4-6,8-
```

2)9-3/h4H2,1-3H3', SMILES:'COC(OC)(OC)CCl', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n13:Molecule { node_key:'M89', INCHI:'InChI=1S/C12H14N2O5/c1-18-12(15)8-2-3-11(14(16)17)10(6-8)13-7-9-4-5-19-9/h2-3,6,9,13H,4-5,7H2,1H3/t9-/m0/s1', SMILES:'C1O[C@@@H](C1)CNc2c(N(=O)=O)ccc(c2)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n14:Molecule { node_key:'M90', INCHI:'InChI=1S/C4H9NO/c5-3-4-1-2-6-4/h4H,1-3,5H2/t4-/m0/s1', SMILES:'NC[C@H]1OCCI', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n15:Molecule { node_key:'M91', INCHI:'InChI=1S/C8H6FNO4/c1-14-8(11)5-2-3-7(10(12)13)6(9)4-5/h2-4H,1H3', SMILES:'[O-][N+](=O)c1c(F)cc(cc1)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n16:Molecule { node_key:'M92', INCHI:'InChI=1S/C7H14CIN2/c1-9(2)5-7(8)6-10(3)4/h5-6H,1-4H3/q+1', SMILES:'CN(C)/C=C(Cl)/C=[N+](C)C', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n17:Molecule { node_key:'M93', INCHI:'InChI=1S/C10H9BrO3/c1-6(12)10(2)13-8-5-3-4-7(11)9(8)14-10/h3-5H,1-2H3/t10-/m0/s1', SMILES:'c1cc2O[C@](C)(C(=O)C)Oc2c(c1)Br', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n18:Molecule { node_key:'M94', INCHI:'InChI=1S/C12H14N2O4/c1-17-12(16)7-2-3-8(13)9(6-7)14-11(15)10-4-5-18-10/h2-3,6,10H,4-5,13H2,1H3,(H,14,15)/t10-/m0/s1', SMILES:'C1O[C@@@H](C1)C(=O)Nc2c(N)ccc(c2)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n19:Molecule { node_key:'M95', INCHI:'InChI=1S/C4H6O3/c5-4(6)3-1-2-7-3/h3H,1-2H2,(H,5,6)/t3-/m0/s1', SMILES:'C1O[C@@@H](C1)C(O)=O', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n20:Molecule { node_key:'M96', INCHI:'InChI=1S/C8H10N2O2/c1-12-8(11)5-2-3-6(9)7(10)4-5/h2-4H,9-10H2,1H3', SMILES:'Nc1c(N)cc(cc1)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n21:Molecule { node_key:'M97', INCHI:'InChI=1S/C10H9ClN2O2/c1-15-10(14)6-2-3-7-8(4-6)13-9(5-11)12-7/h2-4H,5H2,1H3,(H,12,13)', SMILES:'COC(=O)c1cc2c(cc1)nc([nH]2)CCl', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n22:Molecule { node_key:'M98', INCHI:'InChI=1S/C11H14O4S/c1-9-2-4-11(5-3-9)16(12,13)15-8-10-6-7-14-10/h2-5,10H,6-8H2,1H3/t10-/m0/s1', SMILES:'Cc1ccc(cc1)S(=O)(=O)OC[C@H]2OCC2', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n23:Molecule { node_key:'M99', INCHI:'InChI=1S/C14H14N2O4/c1-19-14(18)9-2-3-11-12(6-9)16(13(8-17)15-11)7-10-4-5-20-10/h2-3,6,8,10H,4-5,7H2,1H3/t10-/m0/s1', SMILES:'COC(=O)c1cc2n(C[C@H]3OCC3)c(C=O)nc2cc1', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n24:Molecule { node_key:'M100', INCHI:'InChI=1S/C25H28FN3O4/c1-32-25(31)17-5-6-21-22(13-17)29(14-18-9-12-33-18)23(27-21)15-28-10-7-16(8-11-28)19-3-2-4-20(26)24(19)30/h2-6,13,16,18,30H,7-12,14-15H2,1H3/t18-/m0/s1', SMILES:'Fc1c(O)c(ccc1)C2CCN(CC2)Cc3n(C[C@H]4OCC4)c5c(n3)ccc(c5)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n25:Molecule { node_key:'M101', INCHI:'InChI=1S/C14H14ClNO4S/c1-10-3-6-12(7-4-10)21(18,19)20-14(2,17)13-8-5-11(15)9-16-13/h3-9,17H,1-2H3/t14-/m1/s1', SMILES:'Cc1ccc(cc1)S(=O)(=O)O[C@@](C)(O)c2ncc(Cl)cc2', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n26:Molecule { node_key:'M102', INCHI:'InChI=1S/C32H35ClN4O6/c1-32(28-9-7-22(33)17-35-28)42-27-5-3-4-24(30(27)43-32)20-10-13-37(14-11-20)19-29(38)36-25-8-6-21(31(39)40-2)16-26(25)34-18-23-12-15-41-23/h3-9,16-17,20,23,34H,10-15,18-19H2,1-2H3,(H,36,38)/t23-,32-/m0/s1', SMILES:'C1O[C@@@H](C1)CNc2c(ccc(c2)C(OC)=O)NC(=O)CN3CCC(CC3)c4c5O[C@](C)(Oc5ccc4)c6n cc(Cl)cc6', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n27:Molecule { node_key:'M103', INCHI:'InChI=1S/C28H27ClFN3O5/c1-28(24-9-7-19(29)15-31-24)37-23-5-3-4-20(26(23)38-28)17-10-12-33(13-11-17)16-25(34)32-22-8-6-18(14-21(22)30)27(35)36-2/h3-9,14-15,17H,10-13,16H2,1-2H3,(H,32,34)/t28-/m0/s1', SMILES:'COC(=O)c1cc(F)c(cc1)NC(=O)CN2CCC(CC2)c3c4O[C@](C)(Oc4ccc3)c5ncc(Cl)cc5', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n28:Molecule { node_key:'M104', INCHI:'InChI=1S/C4H6O2/c5-3-4-1-2-6-4/h3-4H,1-2H2/t4-/m0/s1', SMILES:'O=C[C@H]1OCC1', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n29:Molecule { node_key:'M105', INCHI:'InChI=1S/C28H29ClN4O5/c1-28(24-9-7-19(29)15-31-24)37-23-5-3-4-20(26(23)38-28)17-10-12-33(13-11-17)16-25(34)32-22-8-6-18(14-21(22)30)27(35)36-2/h3-9,14-15,17H,10-13,16,30H2,1-2H3,(H,32,34)/t28-/m0/s1', SMILES:'COC(=O)c1cc(N)c(cc1)NC(=O)CN2CCC(CC2)c3c4O[C@](C)(Oc4ccc3)c5ncc(Cl)cc5', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''}, (n30:Molecule { node_key:'M106', INCHI:'InChI=1S/C20H21ClN2O4/c1-20(17-6-5-14(21)11-22-17)26-16-'

```

4-2-3-15(19(16)27-20)13-7-9-23(10-8-13)12-18(24)25/h2-6,11,13H,7-10,12H2,1H3,(H,24,25)/t20-/m0/s1',
SMILES:'O=C(O)CN1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4', Safety:'', Environmental:'',
Legal:'', Economic:'', Control:'', Throughput:''},
(n31:Molecule { node_key:'M107', INCHI:'InChI=1S/C32H33ClN4O6/c1-32(28-9-7-22(33)17-35-28)42-27-
5-3-4-24(30(27)43-32)20(11-14-38)10-13-34-18-29-36-25-8-6-21(31(39)40-2)16-26(25)37(29)19-23-12-15-
41-23/h3-9,14,16-17,20,23,34H,10-13,15,18-19H2,1-2H3/t20?,23-,32-/m0/s1',
SMILES:'COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CNCCC(CC=O)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)c
c6', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:''},
(n32:Molecule { node_key:'M108', INCHI:'InChI=1S/C18H19ClN2O3/c1-18(16-6-5-13(19)11-21-16)23-15-
4-2-3-14(17(15)24-18)12(7-9-20)8-10-22/h2-6,10-12H,7-9,20H2,1H3/t12?,18-/m0/s1',
SMILES:'NCCCC(CC=O)c1c2O[C@](C)(Oc2ccc1)c3ncc(Cl)cc3', Safety:'', Environmental:'', Legal:'',
Economic:'', Control:'', Throughput:''}

```

Copying this query in neo4j browser will look like this:

The screenshot shows the Neo4j Browser interface. On the left, the 'Database Information' sidebar is visible, showing the selected database 'wf2test02'. The main area displays the results of a query run in the terminal window above. The query creates nodes for molecules M77 and M78, setting their InChI and SMILES properties. The results show the generated InChI and SMILES strings for each molecule. Below the browser window, there is a 'neo4j' logo and two cards: 'Getting started with Neo4j Browser' and 'Try Neo4j with live data'.

Pressing the blue arrow at the top will run the query with the following answer:

The screenshot shows the Neo4j Browser interface after the query has been executed. The terminal window displays the command and the response: 'Added 32 labels, created 32 nodes, set 96 properties, completed after 2 ms.' This indicates that the nodes for molecules M77 and M78 have been successfully created with their respective InChI and SMILES properties.

If at any point, it is required to start all over again, the following query can be used to delete all what has been saved:

```
>> MATCH (n) DETACH DELETE n
```

2. Query 2 to create all the Reaction nodes (implemented in the same way as query 1):

```
CREATE (n1:Reaction { node_key:'R53', reaction_smiles:'COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7>O=C(O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n2:Reaction { node_key:'R54', reaction_smiles:'N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4.COC(=O)c1cc2n(C[C@H]3OCC3)c(Cl)nc2cc1>COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n3:Reaction { node_key:'R55', reaction_smiles:'CC(C)(C)OC(=O)N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4>N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n4:Reaction { node_key:'R56', reaction_smiles:'CC(C)(C)OC(=O)N1CCC(=CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4>CC(C)(C)OC(=O)N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n5:Reaction { node_key:'R57', reaction_smiles:'CC(C)(C)OC(=O)N1CCC(=CC1)B2OC(C)(C)C(C)(O2)C.C1c1nc(cc1)[C@]2(C)Oc3c(O2)cccc3Br>CC(C)(C)OC(=O)N1CCC(=CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n6:Reaction { node_key:'R58', reaction_smiles:'C#Cc1ncc(Cl)cc1.c1cc(Br)c(O)c(c1)O>Cc1cnc(cc1)[C@]2(C)Oc3c(O2)cccc3Br', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n7:Reaction { node_key:'R59', reaction_smiles:'C1O[C@@H](C1)CNc2c(N)ccc(c2)C(=O)OC.COC(OC)(O)CCl>COC(=O)c1cc2n(C[C@H]3OCC3)c(Cl)nc2cc1', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n8:Reaction { node_key:'R60', reaction_smiles:'C1O[C@@H](C1)CNc2c(N(=O)=O)ccc(c2)C(=O)OC>C1O[C@@H](C1)CNc2c(N)ccc(c2)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n9:Reaction { node_key:'R61', reaction_smiles:'NC[C@H]1OCC1.[O-][N+](=O)c1c(F)cc(cc1)C(=O)OC>C1O[C@@H](C1)CNc2c(N(=O)=O)ccc(c2)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n10:Reaction { node_key:'R62', reaction_smiles:'CN(C)/C=C(Cl)/C=[N+](C)C.c1cc2O[C@](C)(C(=O)C)Oc2c(c1)Br>Cc1cnc(cc1)[C@]2(C)Oc3c(O2)cccc3Br', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n11:Reaction { node_key:'R63', reaction_smiles:'C1O[C@@H](C1)C(=O)Nc2c(N)ccc(c2)C(=O)OC>C1O[C@@H](C1)CNc2c(N)ccc(c2)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n12:Reaction { node_key:'R64', reaction_smiles:'C1O[C@@H](C1)C(O)=O.Nc1c(N)cc(cc1)C(=O)OC>C1O[C@@H](C1)C(=O)Nc2c(N)ccc(c2)C(=O)OC', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n13:Reaction { node_key:'R65', reaction_smiles:'COC(=O)c1cc2c(cc1)nc([nH]2)CCl.Cc1ccc(cc1)S(=O)(=O)OC[C@H]2OCC2>COC(=O)c1cc2n(C[C@H]3OCC3)c(Cl)nc2cc1', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n14:Reaction { node_key:'R66', reaction_smiles:'Nc1c(N)cc(cc1)C(=O)OC.COC(OC)(OC)CCl>COC(=O)c1cc2c(cc1)nc([nH]2)CCl', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n15:Reaction { node_key:'R67', reaction_smiles:'COC(=O)c1cc2n(C[C@H]3OCC3)c(C=O)nc2cc1.N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4>COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},  
(n16:Reaction { node_key:'R68', reaction_smiles:'Fc1c(O)c(ccc1)C2CCN(CC2)Cc3n(C[C@H]4OCC4)c5c(n3)ccc(c5)C(=O)OC.Cc1ccc(cc1)S(=O)(=O)O[C@@](C)(O)c2ncc(Cl)cc2>COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7', Safety:'', Environmental:'', Legal:'', Economic:'', Control:'', Throughput:'{}},
```

```

(n17:Reaction { node_key:'R69', reaction_smiles:'C1O[C@@H](C1)CNc2c(cc(c2)C(OC)=O)NC(=O)CN3CC(CC(C3)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)cc6>COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}),
(n18:Reaction { node_key:'R70', reaction_smiles:'NC[C@H]1OCC1.COC(=O)c1cc(F)cc1NC(=O)CN2CC(CC2)c3c4O[C@](C)(Oc4ccc3)c5ncc(Cl)cc5>C1O[C@H](C1)CNc2c(cc(c2)C(OC)=O)NC(=O)CN3CC(CC3)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)cc6', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}),
(n19:Reaction { node_key:'R71', reaction_smiles:'O=C[C@H]1OCC1.COC(=O)c1cc(N)cc1NC(=O)CN2CC(CC2)c3c4O[C@](C)(Oc4ccc3)c5ncc(Cl)cc5>C1O[C@H](C1)CNc2c(cc(c2)C(OC)=O)NC(=O)CN3CC(CC3)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)cc6', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}),
(n20:Reaction { node_key:'R72', reaction_smiles:'C1O[C@H](C1)CNc2c(N)ccc(c2)C(=O)OC.O=C(O)CN1CCCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4>C1O[C@H](C1)CNc2c(cc(c2)C(OC)=O)NC(=O)CN3CCCC(CC3)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)cc6', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}),
(n21:Reaction { node_key:'R73', reaction_smiles:'COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CNCCCC(CC=O)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)cc6>COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}),
(n22:Reaction { node_key:'R74', reaction_smiles:'COC(=O)c1cc2n(C[C@H]3OCC3)c(C=O)nc2cc1.NCCC(CC=O)c1c2O[C@](C)(Oc2ccc1)c3ncc(Cl)cc3>COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CNCCCC(CC=O)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)cc6', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '})

```

Copying this query in neo4j browser will look like this:

The screenshot shows the Neo4j browser interface. On the left, the "Database Information" sidebar is open, showing the database "wf2test02" selected under "Use database". It also lists "Node Labels" (Molecule, *32), "Relationship Types" (none), and "Property Keys" (Control, Economic, Environmental, INCHI, Legal, SMILES, Safety, Throughput, node_key, reaction_smiles). On the right, the main window displays a command-line interface (CLI) with two queries. The top query is a complex CREATE statement for reactions R53, R54, and R55. The bottom query is a CREATE statement for a molecule M77. Both queries are highlighted in yellow. Below the CLI, a message states: "Added 32 labels, created 32 nodes, set 96 properties, completed after 2 ms." A blue arrow icon is located at the top right of the CLI area.

```

wf2test02$ CREATE (n1:Reaction { node_key:'R53', reaction_smiles:'COC(=O)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7>O=C(0)c1cc2n(C[C@H]3OCC3)c(nc2cc1)CN4CCC(CC4)c5c6O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7'}, (n2:Reaction { node_key:'R54', reaction_smiles:'N1CCC(CC1)c2c3O[C@](C)(Oc3ccc2)c4ncc(Cl)cc4>COC(=O)c1cc2n(C[C@H]3OCC3)c(cc1)NC(=O)CN3CCCC(CC3)c4c5O[C@](C)(Oc5ccc4)c6ncc(Cl)cc6'}, (n3:Reaction { node_key:'R55', reaction_smiles:'CCCC(O)C(=O)N1CCC(CC1)c2c3O[C@](C)(Oc6ccc5)c7ncc(Cl)cc7'}), (n3:Reaction { node_key:'M77', INCHI: 'InChI=1S/C31...' })

```

Pressing the blue arrow at the top will run the query with the following answer:

A quick way to double check the creation of all the molecule and reaction nodes is running this short query:

```
MATCH (n:Molecule),(r:Reaction) RETURN n,r
```

Pressing the blue arrow at the top will run the query with the following answer:

3. Query 3 to create all the relationships (substrate and product).

This query will execute one element at the time, wait 10s until all of them are completed.

Copying this query in neo4j browser will look like this:

The screenshot shows the Neo4j browser interface. On the left, the 'Database Information' sidebar shows the database is 'wf2test02'. Under 'Node Labels', 'Molecule' is selected. Under 'Relationship Types', 'Product' is selected. The main area displays a list of six CREATE queries:

- 1 MATCH (n:Molecule),(e:Reaction) WHERE n.node_key='M77' AND e.node_key='R53' CREATE (e)-[:RELTYPE:Product]->(n);
- 2 MATCH (n:Molecule),(e:Reaction) WHERE n.node_key='M78' AND e.node_key='R53' CREATE (n)-[:RELTYPE:Substrate]->(e);
- 3 MATCH (n:Molecule),(e:Reaction) WHERE n.node_key='M78' AND e.node_key='R54' CREATE (e)-[:RELTYPE:Product]->(n);
- 4 MATCH (n:Molecule),(e:Reaction) WHERE n.node_key='M78' AND e.node_key='R67' CREATE (e)-[:RELTYPE:Product]->(n);
- 5 MATCH (n:Molecule),(e:Reaction) WHERE n.node_key='M78' AND e.node_key='R68' CREATE (e)-[:RELTYPE:Product]->(n);
- 6 MATCH (n:Molecule),(e:Reaction) WHERE n.node_key='M78' AND e.node_key='R69' CREATE (e)-[:RELTYPE:Product]->(n); MATCH

Below the queries, a graph visualization shows 54 nodes. The right sidebar shows 'Overview' and 'Node labels' with 'Molecule (32)' and 'Reaction (22)'.

Pressing the blue arrow at the top will run the query with the following answer:

The screenshot shows the Neo4j browser interface again. The 'Connected as' section is expanded, showing the user has roles: admin, PUBLIC, and Admin: server user list. The right panel shows the results of running the previously copied query, displaying a list of nodes and relationships.

Now we can see the whole network already created:

```
MATCH (n:Molecule),(r:Reaction) RETURN n,r
```

Pressing the blue arrow at the top will run the query with the following answer:

In this window you can zoom and reorganise the nodes for a clearer visualisation. Nodes in Neo4j follow a gravity field, so if you pick a hub node and drag it around the canvas, the whole network will reorganise automatically.

Now the nodes contain all the relationships. To visualise the **data model** on the graph database (equivalent to a traditional Database Schema) run the following query:

```
CALL db.schema.visualization()
```

Pressing the blue arrow at the top will run the query with the following answer:

Route nodes

The route nodes illustrated on the paper are created automatically using an algorithm that identify all the routes from the entire network. In this case, as we are not providing access to this specific algorithm, the number of routes nodes already identified (12) will be used to create them on the network. To achieve this, two queries are needed:

To create the 12 Route nodes run this query:

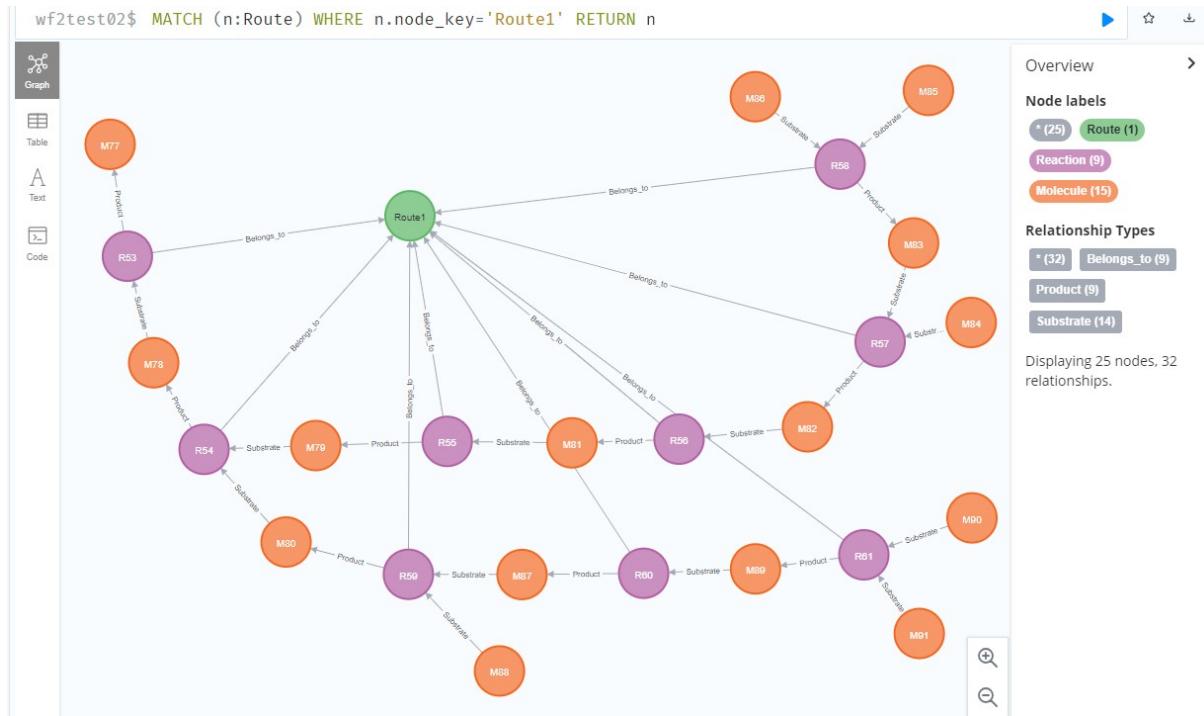
```
CREATE (n1:Route { node_key:'Route1', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n2:Route { node_key:'Route2', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n3:Route { node_key:'Route3', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n4:Route { node_key:'Route4', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n5:Route { node_key:'Route5', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n6:Route { node_key:'Route6', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n7:Route { node_key:'Route7', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n8:Route { node_key:'Route8', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n9:Route { node_key:'Route9', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n10:Route { node_key:'Route10', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n11:Route { node_key:'Route11', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '}), (n12:Route { node_key:'Route12', Safety:' ', Environmental:' ', Legal:' ', Economic:' ', Control:' ', Throughput:' '})
```

To create the relationships run these queries:

Route 1

```
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction),(e4:Reaction),(e5:Reaction),(e6:Reaction),(e7:Reaction),(e8:Reaction),(e9:Reaction) WHERE n.node_key='Route1' AND e1.node_key='R53' AND e2.node_key='R54' AND e3.node_key='R59' AND e4.node_key='R55' AND e5.node_key='R60' AND e6.node_key='R56' AND e7.node_key='R61' AND e8.node_key='R57' AND e9.node_key='R58' CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-[REL2:Belongs_to]->(n), (e3)-[REL3:Belongs_to]->(n), (e4)-[REL4:Belongs_to]->(n), (e5)-[REL5:Belongs_to]->(n), (e6)-[REL6:Belongs_to]->(n), (e7)-[REL7:Belongs_to]->(n), (e8)-[REL8:Belongs_to]->(n), (e9)-[REL9:Belongs_to]->(n)
```

The resulting route can be visualised by unhiding all the adjacent nodes resulting on this:



In the same way, to get all the other routes, run the query below.

Route 2 to 10 – copy and paste whole block

```

MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction),(e4:Reaction),(e5:Reaction),(e6:Reaction),(e7:Reaction),(e8:Reaction),(e9:Reaction)
WHERE n.node_key='Route2' AND e1.node_key='R53' AND e2.node_key='R54'
AND e3.node_key='R59' AND e4.node_key='R55' AND e5.node_key='R63' AND e6.node_key='R56'
AND e7.node_key='R64' AND e8.node_key='R57' AND e9.node_key='R58'
CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-[REL2:Belongs_to]->(n), (e3)-[REL3:Belongs_to]->(n), (e4)-[REL4:Belongs_to]->(n), (e5)-[REL5:Belongs_to]->(n), (e6)-[REL6:Belongs_to]->(n), (e7)-[REL7:Belongs_to]->(n), (e8)-[REL8:Belongs_to]->(n), (e9)-[REL9:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction),(e4:Reaction),(e5:Reaction),(e6:Reaction),(e7:Reaction),(e8:Reaction),(e9:Reaction)
WHERE n.node_key='Route3' AND e1.node_key='R53' AND e2.node_key='R54'
AND e3.node_key='R59' AND e4.node_key='R55' AND e5.node_key='R60' AND e6.node_key='R56'
AND e7.node_key='R61' AND e8.node_key='R57' AND e9.node_key='R62'
CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-[REL2:Belongs_to]->(n), (e3)-[REL3:Belongs_to]->(n), (e4)-[REL4:Belongs_to]->(n), (e5)-[REL5:Belongs_to]->(n), (e6)-[REL6:Belongs_to]->(n), (e7)-[REL7:Belongs_to]->(n), (e8)-[REL8:Belongs_to]->(n), (e9)-[REL9:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction),(e4:Reaction),(e5:Reaction),(e6:Reaction),(e7:Reaction),(e8:Reaction),(e9:Reaction)
WHERE n.node_key='Route4' AND e1.node_key='R53' AND e2.node_key='R54'
AND e3.node_key='R59' AND e4.node_key='R55' AND e5.node_key='R63' AND e6.node_key='R56'
AND e7.node_key='R64' AND e8.node_key='R57' AND e9.node_key='R62'
CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-[REL2:Belongs_to]->(n), (e3)-[REL3:Belongs_to]->(n), (e4)-[REL4:Belongs_to]->(n), (e5)-[REL5:Belongs_to]->(n), (e6)-[REL6:Belongs_to]->(n), (e7)-[REL7:Belongs_to]->(n), (e8)-[REL8:Belongs_to]->(n), (e9)-[REL9:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction),(e4:Reaction),(e5:Reaction),(e6:Reaction),(e7:Reaction),(e8:Reaction)
WHERE n.node_key='Route5' AND e1.node_key='R53' AND e2.node_key='R54'
AND e3.node_key='R65' AND e4.node_key='R55' AND e5.node_key='R66' AND e6.node_key='R56'
AND e7.node_key='R57' AND e8.node_key='R58'
CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-[REL2:Belongs_to]->(n), (e3)-[REL3:Belongs_to]->(n), (e4)-[REL4:Belongs_to]->(n), (e5)-[REL5:Belongs_to]->(n), (e6)-[REL6:Belongs_to]->(n), (e7)-[REL7:Belongs_to]->(n), (e8)-[REL8:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction),(e4:Reaction),(e5:Reaction),(e6:Reaction),(e7:Reaction)

```

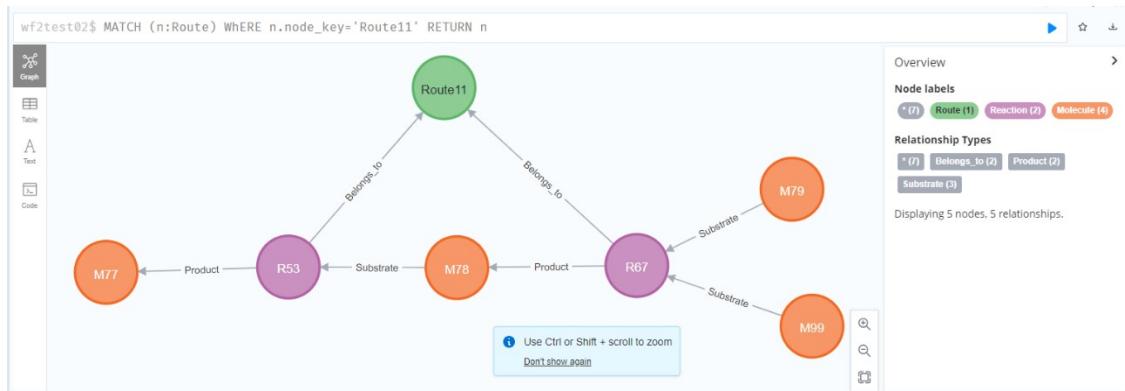
```

Reaction),(e8:Reaction) WHERE n.node_key='Route6' AND e1.node_key='R53' AND e2.node_key='R54'
AND e3.node_key='R65' AND e4.node_key='R55' AND e5.node_key='R66' AND e6.node_key='R56'
AND e7.node_key='R57' AND e8.node_key='R62' CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-
[REL2:Belongs_to]->(n), (e3)-[REL3:Belongs_to]->(n), (e4)-[REL4:Belongs_to]->(n), (e5)-
[REL5:Belongs_to]->(n), (e6)-[REL6:Belongs_to]->(n), (e7)-[REL7:Belongs_to]->(n), (e8)-
[REL8:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction) WHERE n.node_key='Route7' AND e1.node_
key='R53' AND e2.node_key='R69' AND e3.node_key='R71' CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-
[REL2:Belongs_to]->(n) , (e3)-[REL3:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction) WHERE n.node_key='Route8' AND e1.node_
key='R53' AND e2.node_key='R69' AND e3.node_key='R70' CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-
[REL2:Belongs_to]->(n) , (e3)-[REL3:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction) WHERE n.node_key='Route9' AND e1.node_
key='R53' AND e2.node_key='R69' AND e3.node_key='R72' CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-
[REL2:Belongs_to]->(n) , (e3)-[REL3:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction),(e3:Reaction) WHERE n.node_key='Route10' AND e1.node_
key='R53' AND e2.node_key='R73' AND e3.node_key='R74' CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-
[REL2:Belongs_to]->(n) , (e3)-[REL3:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction) WHERE n.node_key='Route11' AND e1.node_key='R53' A
ND e2.node_key='R67' CREATE (e1)-[REL1:Belongs_to]->(n), (e2)-[REL2:Belongs_to]->(n);
MATCH (n:Route),(e1:Reaction),(e2:Reaction) WHERE n.node_key='Route12' AND e1.node_key='R53' A
ND e2.node key='R68' CREATE (e1)-[REL1:Belongs to]->(n), (e2)-[REL2:Belongs to]->(n)

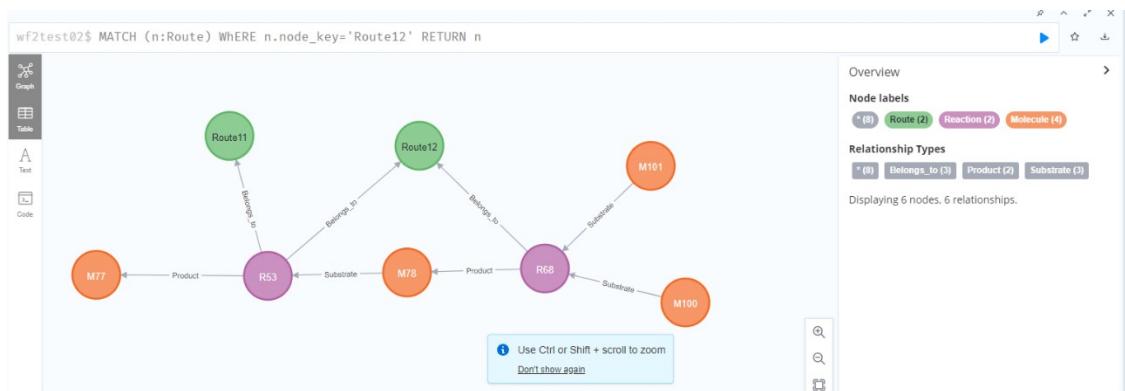
```

And two extra examples of the resulting routes:

Route 11



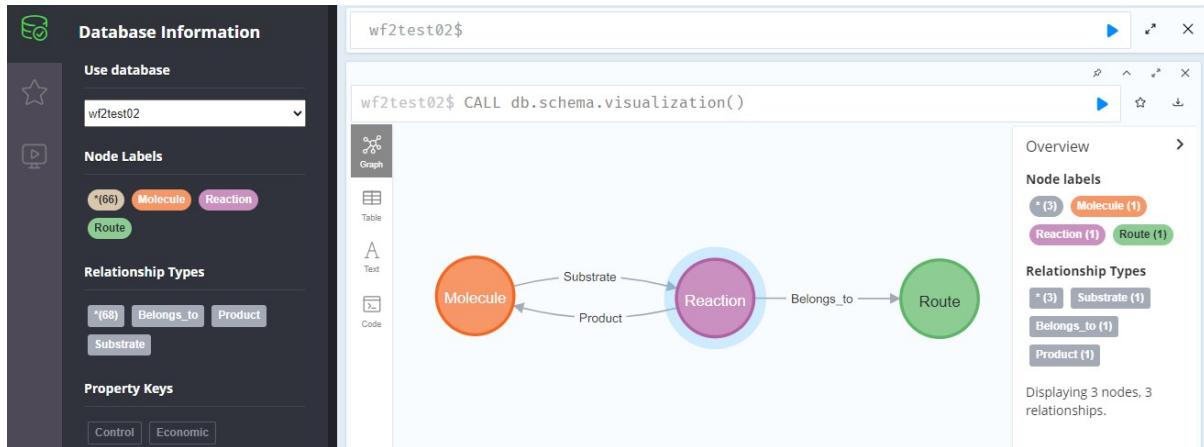
Route 12



Finally, to visualise the **data model** again on the graph database (after introducing the route node) run the following query:

```
CALL db.schema.visualization()
```

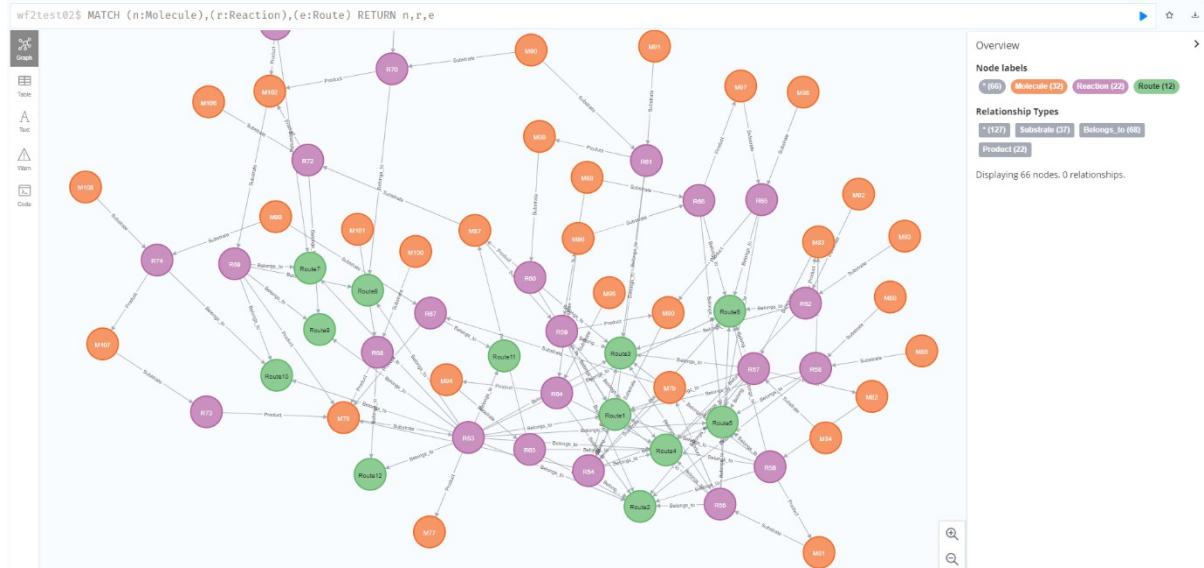
Pressing the blue arrow at the top will run the query with the following answer:



Now the database contains an intricated array of connections and browsing manually become difficult. For this reason, queries to uncover specific parts of the graph are needed.

For instance, this query will reveal the whole graph (this might take time to be resolved):

```
MATCH (n:Molecule),(r:Reaction),(e:Route) RETURN n,r,e
```



For instance, reducing the query will return the reactions and molecule nodes only:

```
MATCH (n:Molecule),(r:Reaction) RETURN n,r
```

