

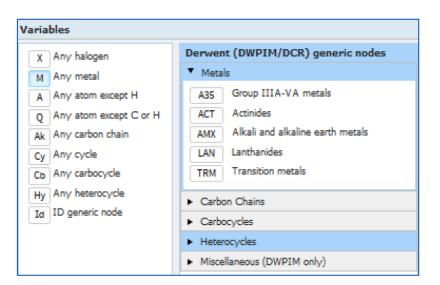
This document summarizes all known issues that can affect the results of structure searches in the Derwent Markush Resource (DWPIM) database.

If you need support concerning your structure query or regarding DWPIM in general, please contact EMEAhelp@cas.org.

1.Use M (any metal) generic node for Metallacycles

Derwent metal generic nodes A35, ACT, AMX, LAN and TRM do not work correctly within ring systems and lead to 0 hits.

Possible Workaround: Use generic node M (any metal) within rings.







2. Within rings only one repeating group (rpg) per ring is allowed

In case of two or more repeating groups within the same ring incorrect results are obtained.

There is no workaround other than drawing out all possible combinations for every additional repeating group.

Example:



3.DWPIM Hitstructures in WPIX/WPINDEX after Cross-Over

DWPIM hitstructures are only preserved in WPIX/WPINDEX answer sets within the same session (including log off hold). Otherwise hitstructures are lost and the full Markush record is diplayed. This limitation may also affect results from SDI's and batch searches.

Recommendation: hitstructure information within DWPIM is preserved. Therefore all answer sets should always be also saved in DWPIM.

Records from DWPIM and WPIX/WPINDEX can be mutually allocated by Markush number (AN in DWPIM corresponds to MCN in WPIX/WPINDEX).



4.STNext: Adjust bond value for Carboxylic acid derivates to exact/normalized (e/n)

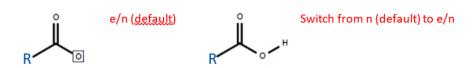
Carboxylic acid derivates as well as corresponding phosphoric, sulfonic and selenic acid derivates are defined with normalized bonds in the STNext structure editor (as shortcuts or drawn, see table below). In order to achieve correct matching in DWPIM those bonds need to be set to exact/normalized. This can be done either by drawing a respective group with locked heteroatom (bonds are automatically set to e/n in the editor) or by changing the normalized bonds of the Hydrogen containing group manually.

The bond value of the shortcuts cannot be changed and therefore those shortcuts should not be used at all!

The following groups are affected:

-соон	-OPO3H2	-SO3H	-SeO3H
-CO2H	-OSO3H	-SO2H	-SeO2H
-COSH	-PO3H2		
-CSSH	-PO2H		
-CS2H			

How to draw those groups correctly:





5.Usage of generic ring and chain nodes via variable attachment points (VPA) can lead to incorrect results

The problem only applies to directly VPA connected generic ring or chain nodes (example A). If there is at least one atom in the VPA between the generic ring or chain node and the main molecule, the problem does not exist (example B).

The following nodes are affected:

- Cy, Cb, Hy, CYC, ARY, HEA, HET, HEF
- Ak, CHK, CHE, CHK

The only practicable workaround for this issue is to use the respective individual structures encoded in the VPA query (example C).

Examples:



6.R-groups within repeating groups may lead to incomplete results

The variants within R groups within repeating groups do not provide mixed results but only results of one of the occurring variants.

Examples:

Query structure A with R1=C,O does only yield results with either C's (example B) or O's but not mixed results (example C).

The only practicable workaround is to search for structures with mixed variants separately as individual structures.



7. Rings with unspecified bonds do not match on aromatic generic ring nodes (ARY, HEA)

Examples:

Query structure A does not match on hit records containing HEA (e.g. record A003-X0H04).

The only practicable workaround is to use query structures with alternating double and single bonds (example B, Match level Class) or to use the respective generic aromatic ring node HEA (example C, Match level Class).

The same applies for carbocycles. In these cases hit records with ARY are missed. For the workaround the respective benzene ring or the ARY node have to be used.



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