# **Supporting Information**

New substructure filters for removal of pan assay interference compounds [PAINS] from screening libraries and for their exclusion in bioassays

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TABLE S1. Functional Group Filters(#) Used For the WEHI 93K HTS Library to Remove Inappropriate Compounds. Subsequent Groups Used for the CTX 136K HTS Library are double hashed (##). These are currently blocked out but it is recommended that they be activated (unblocked) prior to the purchase of new libraries as a more expansive set of filters

```
#SYBYL/3DB HITLIST
#@CLASS REGLIST
#@DATABASE NONE
#@SOURCE in-house
#One hash means in original filter for WEHI 93K HTS Library
##Two hashes mean added for CTx 136K HTS Library (but blocked out here)
##The added ones may be new or simply broadened
##Many definitions are dependent on others (e.g. if allow sulfonates, others defs need modification)
#acid halide
C(=O)Hal
##...and related
##C(=S)Hal
#any het halide (includes sulfonyl halides)
HetHal
##acrylamide and related, including vinylpyridines etc:
##C(=O)C=CH2
##S(=O)C=CH2
##N#CC=CH2
##C(=CH2)C[5]:N:Hev:Hev:Hev:Hev:@5
##....and related
##C(=S)C=CH2
##acyl carbamate and related hot linear aryl carbonates/carbamates
##OK to go
\#C(=O)NC(=O)Any[IS=O,S]
HetC(=O)O-[!R]C[8]: Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): Hev: Hev(Any[IS=Hal,C\#N,C(F)(F)F,S(=O)=O,C(=O)\&NOT=C(=O)OH]): Hev: Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): @8 \\ HetC(=O)O-[!R]C[8]: Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): @8 \\ HetC(=O)O-[!R]C[8]: Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): @8 \\ HetC(=O)O-[!R]C[8]: Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): @8 \\ HetC(=O)O-[!R]C[8]: Hev(Any[NOT=O,S[TAC=4],N[TAC=3]): Hev: Hev(Any[NOT=O,S[TAC=4],N[TAC=3])): @8 \\ HetC(=O)O-[!R]C[8]: Hev(Any[NOT=O,S[TAC=4],N[TAC=3]): Hev: Hev(Any[NOT=O,S[TAC=4],N[TAC=3])): @8 \\ HetC(=O)O-[!R]C[8]: Hev(Any[NOT=O,S[TAC=4],N[TAC=4],N[TAC=3]): Hev: Hev(Any[NOT=O,S[TAC=4],N[TAC=4],N[TAC=3])): @8 \\ HetC(=O)O-[!R]C[8]: Hev(Any[NOT=O,S[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[TAC=4],N[T
HetC(=O)O-[!R]C[8]: Hev(Any[IS=Hal,C\#N,C(F)(F)F,S(=O)=O,C(=O)\&NOT=C(=O)OH]): Hev: Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): Hev: Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]): \#(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]): \#(Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any[NOT=O,S[TAC=3],Any
##acyl hydrazone (and oxime) not in ring:
##leave in (as well as carbazides, semicarbazides,
##acylsemicarbazides, hydrazones, acyl hydrazides)
##despite some reservation of Metal chelating for some of these
##O=CHetN=[!R]C(C)Any[IS=C,H]
##acyl hydrazide not in ring -leave in: see above
##O=CNH-[!R]NHC(=O)Het
##acyl imide - taken care of by imide filter
##O=CN(C(=O))C=O
##N-alkyl acyl sulfonamide
##O=C-[!R]N(C)S(=O)(=O)C
#acyl and ##sulfo cyanide
C(=O)C#N
##S(=O)C#N
##C(=S)C#N taken care of by thiocarbonyl filter
##acylimines
\#Any[NOT=O,N]C(Any[NOT=O,N])=[!R]NC=O
#aldehyde
C(=O)H
##C(=S)H taken care of by thiocarbonyl filter
##aliphatic long chain
C[!r]H2CH2CH2C[!r]H2CH2CH3
#alkyl halide
BrC[TAC=4]H
ClC[TAC=4]H
##BrC[TAC=4]
##ClC[TAC=4]H2
##amidotetrazole
##N[1](C=NN=N@1)C=O
###aminals etc not in ring unless basic amine - harshest criteria - could revisit
##HetC[TAC=4]-[!R]Het
```

##HetC[TAC=4]N(Any[IS=H,C[TAC=4]])Any[IS=H,C[TAC=4]]

#anhydride/thioanhydrides C(=Any[IS=O,S])Any[IS=O,S]C=Any[IS=O,S]##anilines - not removed; medchem issue: specifics dealt with in frequent hitter filter ##azepanes - not removed #azido/diazonium/diazo - specifics also in frequent hitters:  $N\sim N[f]$ ##azo (and carbazones etc) ##N[!r]=N #aziridines, epoxides etc N[1]CC@1 ##Hev[1]HetC@1 ##benzidine - not removed; medchem issue ##betalactams/lactones, not thietanes, oxetanes, azetidines etc ##Hev[1]C(=O)HetC~@1 ##beta-keto - in diketos ##beta-amino ketones and related ##CC[!r](=O)CH2CH2Het ##CS[!r](=O)CH2CH2Het ##biotin - not now ##S[1]CH2C[5]HCH(CH@1CH2CH2CH2CH2C=O)NHC(NH@5)=O #boron В #carbamic acids NC(=O)OH ##carbazide - leave in ##NHC(=O)-[!R]NHNH ##carbocation/anion and other wrongly charged atoms ##checked to make sure sulfoxide retained ##C[+1] ##C[-1] S[+1] O[+1] O[TAC=3] Hal[+1] S(Any)(Any)(Any)(Any)Any ##catechol ##C[1]:C(:C:C:C(:C:@1)OH)OH ##chromones - reactive; in Michael acceptors ##crown ethers - main ones - not now ##HetC~CHetC~CHetC~CHetC~CHetC~C ##coumarins - reactive; in Michael Acceptors; other specific ones in frequent hitters #cyanamides CH2NC#N

#cyanate/thiocyanate OC#N

SC#N

#cyanohydrin

N#CCOH ##cycloalkanes - not removed

#cyclohexadienes

C[1]HC=CC=CCH@1

C[1]HC=CCH2C=C@1

##dialkynes

##C#CC#C

##diaminobenzenes - specific ones in frequent hitters

##diketo (includes o-quinones) and beta-diketo and related hot keto

##CC(=Het[!r])C(=Het[!r])C

#CC(=O)C(=O)Any[IS=O,N]

#CC(=Het[!r])S(=O)(=O)

##CC(=O)C(=S)C - covered by thiocarbonyl ##CC(=O)C[TAC=4]C(=O)C

##CC(=O)C[TAC=4]S(=O)C

##CC(=O)C[TAC=4]C#N

##CC(=O)C[TAC=4]C(Hal)(Hal)Hal

##disulfide and related

S[TAC=2&!r]S[TAC=2&!r]

##enamines - in frequent hitters

##ester of hobt and su (and carbamates/carbonates) A C(=O)O[!r]N:Hev ##C(=O)O[!r]NC(=O) ##esters - thio-type; some accounted for by thiocarbonyl removal ##CC(=S)O[!r]C CC(=O)S[!r]C##C(=S)S ##esters - aryl B CC(=O)O[!r]C[TAC=3] ##ester - hot benzyl, non aryl C  $\#\#C[TAC=4]C(=0)O[!r]CH2C[8]: \\ Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): \\ Hev:Hev(Any[S=Hal,C\#N,C(F)(F)F,S(=O)=O,C(=O)Hev[NOT=OH]]): \\ Hev:Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): \\ \#\#C[TAC=4]C(=0)O[!r]CH2C[8]: \\ Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): \\ \#\#C[TAC=4]C(=0)O[!r]CH2C[8]: \\ Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]): \\ \#\#C[TAC=4]C(=0)O[!r]CH2C[8]: \\ \#\#C[TAC=4]C[8]: \\$ ##esters - other labile D ##CC(=O)OCHC(Hal)(Hal)Hal ##C[TAC=4]C(=O)O[!r]CHC(=O)Any[NOT=OH] ##C[TAC=4]C(=O)O[!r]CHC#N ##esters - other labile D ##COC[!r](=O)C(=O) ##COC(=O)S(=O)(=O) ##COC[!r](=O)C[TAC=4]S(=O)=O##COC[!r](=O)C[TAC=4]C#N ##COC[!r](=O)C[TAC=4]C(Hal)(Hal)Hal ##COC[!r](=O)C[TAC=4]N(C(=O))C(=O) ##over halogenated rings ##1,2 ##C[1]:C(:C(:Hev:Hev:Hev:@1)Br)Any[IS=Cl,Br] ##1,3 ##C[1]:C(:Hev:C(:Hev:Hev:@1)Br)Any[IS=Cl,Br] ##1.4 ##C[1]:C(:Hev:Hev:C(:Hev:@1)Br)Any[IS=Cl,Br] ##1,2,4 ##Hev[1]:C(:C(:Hev:C(:Hev:@1)Hal)Hal)Hal ##1,2,3 ##C[1](:C(:C(:Hev:Hev:Hev:@1)Hal)Hal)Hal ##C[1](:C:C(:C:C(:C:@1)Cl)Cl)Cl ##hydrazine and other nucleophilic NH2 Any[IS=O,N]NH2 ##hydroxylamine ##HON[TAC=3]C[TAC=4] ##hydroquinone - more related ones in frequent hitters ##C[1]:C:C:C(:C(:C:@1)OH)OH ##hydroxamic acid ##C(=O)NOH ##imide ##CC(=O)N-[!R]C(=O)C ###imidoyl chlorides N=CHal #imines -reactive ones ##Any[NOT=N,O]N=C[!r](Any[NOT=N])Any[NOT=N] ###imines - not completely clear whether ok or not; may need to reexamine in second pass ##Any[NOT=O,N]N=[!R]C(Any[IS=C,H])Any[IS=C,H] ##Any[NOT=O,N]N=[!R]C(C[TAC=4])C ##iodine ##I #isocyanate, isothiocyanate - dealt with in ketenes #ketene - includes allenes, carbodimides, isocyanates etc Any=C=Any ##maleimides and surrogates ##C[1]C(NC(CH=@1)=O)=O ##C[1](CH2C(NC@1=O)=O)HetC[TAC=3] ##michael acceptors - more in frequent hitter file CCH=C[!r]HC(=O)C CCH=C[!r](C#N)C#N CCH=C[!r](C(=O)Hev~Hev)C#N ##first, alpha-beta unsubstituted, unsaturated nitriles and ketones ##CCH=[!R]C(Any[IS=H,C])Any[IS=C#N,C(=O)&NOT=C(=O)Any[IS=N,O]]##second, linear diactivated A:

# # Any[IS=C,O]CH=[!R]C(Any[IS=C(=O),S(=O),C#N,Hal,C(Hal)(Hal)(Hal)&NOT=C(=O)OH])Any[IS=C(=O),S(=O),C#N&NOT=C(=O)OH]## linear diactivated B: ##Hev[NOT=OH]C(=O)CH=[!R]CHC(=O)Hev[NOT=OH] ## linear diactivated C: # Any[IS=C#N,C(=O)Hev[NOT=OH],S(=O)]CH=[!R]C(Any[IS=H,C])C[8]:Hev(Any[NOT=O,S]TAC=2],C[TAC=4],N[TAC=3]]):@ 8# Any[IS=C#N,C(=O)Hev[NOT=OH],S(=O)] CH=[!R]C(Any[IS=H,C])C[8] : Hev(Any[IS=Hal,C#N,C(F)(F)F,S(=O)=O,C(=O)Hev[NOT=OH]]) : Hev: Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]) : @ 8###thirdly, cyclic: activated coumarins; could revisit some of the less activated amides # C[1](C(=O)OC[5]:C:C:C:C:@5CH=@1)Any[IS=C(=O),C(=S),S(=O),C#N,Hal,C(Hal)(Hal)Hal&NOT=C(=O)OH]###and activated chromones;- could revisit some amides ##C[1](=CHOC[5]:C:C:C:C:@5C@1=O)Any[IS=C(=O),C(=S),S(=O),C#N,Hal,C(Hal)(Hal)Hal&NOT=C(=O)OH]##mercaptothiazoles and related ##S[TAC=2]C[2]:S:C:Any:N:@2 ##N-acyl and quaternary alkyl pyridines etc ##CN[+1]:Hev -redundant by definition below ##Broaden to include N-nitros, N oxides etc: can review ##N-acyl pyrroles, imidazoles etc not in ring, but not ureas, carbamates ##C[!r](=O)N(:Hev):Hev ##naphthylamines -alpha and beta; leave in; medchem issue ##C[1]:C:C[3]:C(:C:C:@1):C:C:C:C:@3-[!R]N ##C[1]:C:C[3]:C(:C:C:@1):C:C:C(:C:@3)[!R]N ##C[1]H:CH:C[5]:C(:CH:CH:@1):CH:CH:CH:C:@5NH ##C[1]H:CH:C[5]:C(:CH:CH:@1):CH:CH:CH:C:@5N(Any[IS=H,C(=O)C])Any[IS=H,C(=O)C] ##C[1]H:CH:C[5]:C(:CH:CH:@1):CH:CH:C(:CH:@5)NH ##nitro, including N-NO2 etc: ##HevN[!r]( $\sim$ O[f]) $\sim$ O[f]  $CN[!r](\sim O[f])\sim O[f]$ #nitroso N[TAC=2]=O ##N oxide and N-hydroxypyridine ##Hev:N[+1]~[!R]O ##N-oxide: broader ##NO[f] ##oxetanes, thietanes (see beta lactams) ###oximes: not removed ##O-N single bond: not removed #peroxide OO ##phenyl carbonate and carbamate - hot ones removed above ##phosphorous ##phthalimides: not removed (except hot ones - above in triaryls) ##polycyclic aromatic: not removed ##polyene: not removed ##quaternary nitrogen ##CN[+1](C)(C)C ##saponins - not removed #silicon Si ##stilbenes - not removed ##sulfur oxygen single bond (includes sulfinic/sulfonic acids/esters) ##SO[!f] ##sulfur-nitrogen single bond, exluding sulfonamides ##NS[TAC=2] ##sulfur-nitrogen single bond surrogate ##C[1](C(=O)N):C(:C:C:C:C:@1)SCCC(=O)Any[NOT=OH] ##thiocarbonyl ##C=S #thiols SH##trichloromethyl ##tri (and di-) haloketones and amides (+esters) and related #C(=O)C[!r](Hal)Hal#trihaloimidate OC(=NH)C(Hal)(Hal)Hal

##isonitrile

##N[TAC=2]~C[TAC=1]

##sulfone-activated rings and related below ##2-halo and 2-sulfonylpyrimidines, but deactivated triazines OK ##Any[IS=Hal,S(=O)(=O)]C[1]:N:Any(Any[NOT=N]):Any:Any(Any[NOT=N]):N:@1 ##4-halo and 4-sulfonylpyrimidines not too deactivated ##C[1](Any[NOT=N]):N:Any(Any[NOT=N]):Any:Any(Any[IS=Hal,S(=O)(=O)]):N:@1 ##4-activated 2-halo and 2-sulfonyl pyridines ##C[1]:N:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,C(C)=NO,C#N,C(=O),C(F)(F)F,S(=O)=O&NOT=C(=O)OH]):Any:@1(Any[IS=Hal,S(=O)(=O)C)):Any:@1(Any[IS=Hal,S(=O)(=O)C)):Any:@1(Any[IS=Hal,S(=O)(=O)C)):Any:@1(Any[IS=Hal,S(=O)(=O)C)):Any:Any(Any[IS=Hal,S(=O)(=O)C)):Any##2-activated 4-halo and 4-sulfonyl pyridines ##C[1]:N:Any(Any[IS=Hal,C(C)=NO,C#N,C(=O),C(F)(F)F,S(=O)=O&NOT=C(=O)OH]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:@1(Any[IS=Hal,C(C)=NO,C#N,C(=O),C(F)(F)F,S(=O)=O&NOT=C(=O)OH]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:@1(Any[IS=Hal,C(C)=NO,C#N,C(=O),C(F)(F)F,S(=O)=O&NOT=C(=O)OH]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:@1(Any[IS=Hal,S(=O)(=O)C)):Any:@1(Any[IS=Hal,S(=O)(=O)C)):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:@1(Any[IS=Hal,S(=O)(=O)C)):Any:@1(Any[IS=Hal,S(=O)(=O)C)):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,S(=O)(=O)C]):Any(Any[IS=Hal,S(=O)(=O)C##3-activated 2-halo and 2-sulfonyl pyridines ##5-activated 2-halo and 2-sulfonyl pyridines ##C[1]:N:Any(Any[IS=Hal,S(=O)(=O)C]):Any:Any(Any[IS=Hal,C(C)=NO,C#N,C(=O),C(F)(F)F,S(=O)=O&NOT=C(=O)OH]):@1##6-activated 2-halo and 2-sulfonyl pyridines and not too deactivated ##C[1](Any[IS=Hal,C(C)=NO,C#N,C(=O),C(F)(F)F,S(=O)=O&NOT=C(=O)OH]): N:Any(Any[IS=Hal,S(=O)(=O)C]): Any(Any[NOT=N]): Any(Any##3-activated 4-halo and 4-sulfonyl pyridines ##1,2,4-activated aryls (including activated phthalimides ##C[1](Any[NOT=N,O]):C(Any[N##1,2,3-activated aryls ##C[1](Any[IS=Hal,C#N,C(=O),C(F)(F)F,S(=O)=O&NOT=C(=O)OH]):C(Any[IS=Hal,S(=O)(=O)C,C[r](=O)NC]):C(Any[IS=Hal,C#N,C(=O),C(F)(F)F,S(=O)=O&NOT=C(=O)OH]):C(Any[NOT=N,O]):C(Any[##activated pyrimidine phenol ethers: ##C[1](Any[NOT=O[TAC=2],N[TAC=3])):N:C(:Hev:Hev:N:@1)-[!R]Anv[IS=O,S]C[8]:Hev(Anv[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3])):@8##C[1](Any[NOT=O[TAC=2],N[TAC=3])):N:C(:Hev:Hev:N:@1)-[!R]Any[IS=O,S]C[8]:Hev(Any[IS=Hal,C#N,C(F)(F)F,S(=O)=O]):Hev:Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3])):Hev:Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=4],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=4],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=4],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=4],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=4],N[TAC=3],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=4],N[TAC=3],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=4],N[TAC=3]##C[1](Any[NOT=O[TAC=2],N[TAC=3])):N:C(:N:Hev:Hev[NOT=N]:@1)-[!R]Any[IS=O,S]C[8]:Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]):Hev:Hev(Any[IS=Hal,C#N,C(F)(F)F,S(=O)=O]):Hev:Hev(Any[NOT=O,S[TAC=2],C[TAC=4],N[TAC=3]]):@8##C[1](Any[NOT=O[TAC=2],N[TAC=3])):N:C(:N:Hev:Hev[NOT=N]:@1)-[!R]Any[IS=O,S]C[8]:Hev(Any[IS=Hal,C#N,C(F)(F)F,S(=O)=O]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3])):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=2],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=3],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=3],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=3],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=3],N[TAC=3],N[TAC=3]):Hev:Hev(Any[NOT=O,S[TAC=3],N[TAC=##activated 4-fluorosulphonamides ##C[1](S(=O)(=O)NC:N):CH:CH:C(F):CH:CH:@1 ##thiazolinium halides and related ##Het[1]:C(Hal):N:C:C:@1 ##reactive haloenamines ##CC(C)=C(Hal)N(Any[IS=H,C[TAC=4]])Any[IS=H,C[TAC=4]] ##terminal acetylenes ##C#CH ##Limiting number of certain slns# ##furans and thiophenes## ##S[1]:C:CH:CH:CH:@1<max=1> ##O[1]:C:CH:CH:CH:@1<max=1> ##nitriles ##C#N<max=2> #acids C(=O)OH<max=2>##esters ##C(=O)OC<max=2> #thioethers C[TAC=4]S[TAC=2]C[TAC=4] < max=2 >#halogens Br<max=1> Cl<max=3> F<max=5>

**Supplementary Table S2**. The six high throughput screening campaigns using the WEHI 93K HTS library selected for analysis<sup>c</sup>

HTS	Target Type	<b>Detection method</b>	Nature of	Number of hits <sup>b</sup>
Campaign <sup>a</sup>		/ anchor	interaction	
A	Protein-protein	AlphaScreen® / anti-GST Mab- conjugated acceptor beads, streptavidin-coated donor beads	Binding of GST- protein target to an amphipathic helical ligand	3,006
В	Protein-protein	AlphaScreen® / protein A- conjugated acceptor beads, streptavidin-coated donor beads.	Binding of receptor construct IL13Rα1-Fc to biotinylated cytokine IL13	4,086
С	Protein-protein	AlphaScreen® / anti-GST Mab- conjugated acceptor beads, streptavidin-coated donor beads	Binding of GST- protein target to an amphipathic helical ligand	3,145
<b>D</b> <sup>c</sup>	Protein-protein	AlphaScreen® / anti-GST Mab- conjugated acceptor beads, streptavidin-coated donor beads	Binding of GST- protein target to an amphipathic helical ligand	746
Е	Protein-protein	AlphaScreen® / nickel-coated acceptor beads, streptavidin-coated donor beads	SH2-binding domain: hexaHis-SOCS2 binding to biotinylated gp130	9,309
F	Protein-protein	AlphaScreen® / nickel-coated acceptor beads, streptavidin-coated donor beads	SH2-binding domain: hexaHis-SOCS3 binding to biotinylated gp130	14,745

a. We are not at liberty to reveal the targets for HTS Campaigns A, C and D.

b. Primary hits, classified as any compound that gave an inhibitory signal of 50% or more at the screening concentration tested (50 $\mu$ M except for HTS Campaign D {25 $\mu$ M}, single point determination).

c. A typical assay protocol is described in the standard operating procedure below. This particular protocol is used for follow-up from HTS Campaign D, where we delineate between the host protein as protein D and the competitive ligand as ligand D or peptide D. However, the protocol is essentially the same as that used as the primary screen against the WEHI 93K HTS library in HTS Campaign D, and is closely related to that used for HTS Campaigns A and C, whereas HTS Campaigns E and F used a different linking methodology. This involved the use of Nickel-HexaHis

and we have found that this gives rise to the highest hit rates most plausibly through the additional detection of mildly chelating compounds that interfere with the Nickel anchor. We have therefore abandoned this approach in our screening centre. HTS Campaign B also used a different linking technology as indicated.

Standard Operating Procedure: Routine Alphascreen<sup>TM</sup> Assay using Proxiplates

#### 1. Introduction

The aim of the routine Protein D AlphaScreen<sup>TM</sup> assay is to identify active small molecules against protein D. The compounds tested are generally synthesized in-house, however some outsourced compounds are also investigated. The screens are used to determine the potency of each compound. A GST-counter screen assay is also done to determine the verity of the results ie. protein D inhibition. To determine an accurate estimation of the  $IC_{50}$ , the compounds are routinely tested at starting concentrations,  $100\mu M$  and/or  $1\mu M$  and serially titrated 3 fold over 11 dilutions.

The assay was developed using Alphascreen™ technology which relies on hydrogel coated acceptor and donor beads which have functional groups for conjugation to a protein (GST-protein D or GST-Biotin) and a peptide (Biotin-peptide D) respectively. The beads come in close proximity when the protein and the peptides interact. Donor beads contain a photosensitiser that converts oxygen to an excited form of O₂ at an excitation of 680 nm. Energy is transformed from the singlet oxygen and reacts with chemiluminescers on the acceptor bead, resulting in light emission at 520 - 620 nm. Active library compounds when added to the reaction, reduce the intensity of the luminescence, dependent on the inhibition of proximity of the acceptor and donor beads. With this information, the IC₅0 of each compound can be calculated

#### 2. Assay Summary

# 2.1 Reagents and Materials

GST-protein D and biotinylated GST proteins were prepared and provided by Peter Czabotar from Parkville which were stored as stock solutions at -80°C. The biotinylated peptide D was purchased from Auspep and stored as 500 $\mu$ M stock solutions in 100% DMSO at -80°C. The Alphascreen<sup>TM</sup> GST (Glutathione-S-Transferase) Detection Kit was obtained from Perkin Elmer Lifesciences (Cat #6760603R). The Proxiplates, white 384 well flat-Bottom plates were purchased from Interpath Services, Melbourne (Cat #784075). The seals to cover the plates were purchased from Proscience, Melbourne (Cat#784075). DMSO was purchased from AnalaR. The 384 deep well plates and the Polypropylene 50  $\mu$ L, V bottom polypropylene compound plates were purchased from Matrical.

#### 2.2 Preparation of compounds

The chemists provided the compounds in the form of dry powder which were stored at -20°C. 10mM stocks were made with 100% DMSO the day before the assay was scheduled to be run. 12 $\mu$ L of 100% DMSO and 6 $\mu$ L of 10mM compound (ie. 3.333mM, final 100 $\mu$ M) was added to columns 1 and 12 in the Polypropylene 50  $\mu$ L, V bottom compound plates. To achieve a final compound concentration of 1 $\mu$ M, in a separate matrical plate, 28 $\mu$ L of 100% DMSO and 2 $\mu$ L of 10mM compound was added to a well, mixed well, 2 $\mu$ L of this solution was taken and added to 38 $\mu$ L of 100% DMSO. 20 $\mu$ L of this solution was added to the test metrical plate. For the control wells 15 $\mu$ L 100% DMSO only was added to Lanes 23 and 24 of each plate.

## 2.3 Control Compound Addition

Several control compounds were included in the test plates. The control compounds used routinely are listed as follows.

[Positive Control]	[Stock]	[Start Conc]	[IC <sub>50</sub> , Protein D]
Control 1	0.5mM	15µM	$0.06 - 0.2 \mu M$
Control 2	10mM	100μΜ	3-18µM
Control 3	5mM	150μΜ	$0.7-1\mu M$

# 2.4 Titration of Compounds\_Automation

The compound plates were serially diluted 2 fold using the MiniTrak *Titration\_11pt\_P30.wpt* protocol located in the *C:Packard/Plate Track/Bin* folder.

- a. The plates were centrifuged prior to dilution.
- b. The Mini Trak program was opened and executed and the robot was set up with the following parameters:

Enter number of plates to be titrated:

#### **Parameters for Volumes**

 $\begin{array}{ll} \mbox{Diluent to aspirate:} & 16 \mu L \\ \mbox{Sample to aspirate:} & 8 \mu L \\ \mbox{Volume to Dispense:} & 24 \mu L \end{array}$ 

Volume to mix: 20µL

#### Parameters for Stacker Setup

Compound Plate/s: Stacker 1 front

#### **Parameters for Deck Setup**

Tip Carrier: MPD 4 (hooks facing left)

P30 Tips (only in column 1 and 12): MPD 4 DMSO reservoir: 100% DMSO MPD 6

c. Once titrations are complete, the compound plate was immediately covered with a foil seal to prevent evaporation.

#### 2.5 Buffer Preparation

The assay and bead buffers were prepared fresh on the day. Each titrated compound plate was assayed in duplicate. The following volumes were sufficient to run 12 Proxiplates (4 assay plates run in duplicate in each of protein D and counter assays)

#### Assay Buffer

[Stock]	[Final]	[Volume for 100 mL]
1M Hepes pH 7.4	50mM	5mL
1M DTT	10mM	1mL
4M NaCl	100mM	2.5mL
10% Tween-20	0.05%	0.5mL
10mg/mL Casein	0.1  mg/mL	1mL
Milli-Q H <sub>2</sub> O		90mL

#### **Bead Buffer**

[Stock]	[Final]	[Volume for 100 mL]
1M Tris-HCL pH 7.5	50mM	5mL
10% Tween-20	0.01%	0.1mL
10mg/mL Casein	0.1  mg/mL	1mL
Milli-Q H <sub>2</sub> O		93.9mL

#### 2.6 Protein and Peptide Preparation

- 1. The assay and bead buffers were used to prepare the acceptor and donor solutions. Alphascreen  $^{TM}$  beads are light sensitive and therefore prepared in a darkened room. 2.5  $\mu$ L of beads were added per 1 mL of buffer.
- 2. The volume of protein or peptide added was calculated using the following formula:

$$\frac{C1}{C2} \times V1 \times 2 = V2$$

 $C_I$  = Final Concentration of protein/peptide

 $C_2$  = Stock Concentration of protein/peptide

 $V_I$  = Total Volume of Acceptor/Donor Solution

 $V_2$  = Volume of stock protein/Peptide to add to Acceptor/Donor solution

<sup>\*50%</sup> DMSO is required to wash tips

3. The assay components were prepared as separate Acceptor and Donor Solutions. The Acceptor Solution contains Acceptor beads and target protein, whilst the Donor Solution contains Donor beads and biotinylated peptide.

#### **Protein D**

[Acceptor Solution]	[mL]	[Donor Solution]	[mL]
Assay buffer	10mL	Assay buffer	10mL
Bead buffer	10mL	Bead buffer	10mL
Acceptor Beads	50μL	Donor Beads	50μL
114μM GST-protein D	0.35μL	500μM Bt-Peptide D	$0.32 \mu L$

(or 3.51µL of a 1:10 dilution)

Final Protein [1.0nM] Final Peptide [4nM]

#### Counter-GST

[Acceptor Solution]	[mL]	[Donor Solution]	[mL]
Assay buffer	10mL	Assay buffer	8mL
Bead buffer	10mL	Bead buffer	8mL
Acceptor Beads	50μL	Donor Beads	50μL
77μM B-GST	$1.04 \mu L$		

Final Protein [2nM]

- 4. When the solutions were prepared, they were left to incubate for 30 minutes at room temperature to allow the beads to bind to the protein and the peptide.
- 5. 5 μL of protein D solution or biotinylated-GST was added to columns 1-23 of the appropriate assay plates using the Multi-drop Combi (cassette #5). 5μL Assay/Bead buffer was added to column 24 (no protein).
- 6. Then selected 0.3 uL Tip transfer proxiplat\_stationary.wpt program to Transfer 0.3μL of sample from the compound plate into each assay plate.

# **Parameters for Stacker Setup**

Assay Plates: Stacker 3 front Compound Plates: Stacker 2 rear

# **Parameters for Deck Setup**

P10 Tips: MPD 4 (hooks facing left)

\*50% DMSO was needed to wash tips

- 7. Plates were then incubated for 30 mins at RT
- 8.  $5\mu L$  of peptide D solution or donor solution (GST-counter) were added into assay plates (columns 1-24) using the Multi-drop Combi.
- 9. After final addition, tapped plates gently and sealed individually with adhesive film.
- 10. Plate were then incubated at RT for ~4hrs

## 2.7 Assay Measurement

The plates were then loaded onto the right stacker of the Envision 2103 plate reader. Protocol \*Alphascreen 384well proxiplate Auto was used to read the plates.

## 2.8 Data Analysis

Data was displayed in an Excel format spreadsheet and imported into ActivityBase database. IC<sub>50</sub> for each compound calculated by using the *PROTEIN\_D\_IC50* and *COUNTER\_GST\_IC50* templates. The percent inhibition was calculated using the following equation:

%Inhibition = 
$$100*(1-\left[\frac{(x-\mu^{-})}{(\mu^{+}-\mu^{-})}\right])$$

x = RFU obtained after compound treatment

 $\mu^-$  = RFU obtained for the negative controls (no protein controls)

 $\mu^+$  = RFU obtained for the positive controls (DMSO vehicle controls)

IC<sub>50</sub> values were then obtained by non-linear least squares fitting of the above data to XLfit3 equation 205:  $y=A+((B-A)/(1+((C/x)^D)))$ .

The quality of the assay results were monitored by determination of the Z Prime factor for each assay plate, where Z Prime => 0.5 for the results was considered as reliable (Zhang *et al*, J Biomol Screening, 4:67-73, 1999).

**Supplementary Table S3**. Data for one particular validated hit, showing that the profile of an optimizable hit may not always be clean

% inhibition at test concentration (50 $\mu M$ )					Count <sup>a</sup>	
Screen A (50 µM)	Screen B (50 μM)         Screen C (50 μM)         Screen D (25 μM)         Screen E (50 μM)         Screen F (50 μM)					
74	58	58 <50 81 67 <50				

a. In other words, this compound registered as a hit in four out of the six assays studied.

**Supplementary Table S4** - Structural classes with examples of the 362 compounds that qualified as hits all six HTS campaigns selected for study. Many of these compounds contain more than one identified assay interference moiety.

Class	Number 51	Examples
Quinone-like	51	O-nBu
		$\begin{array}{c ccccccccccccccccccccccccccccccccccc$
		HON HON Br
Rhodanine- like	50	CI N N N N N N N N N N N N N N N N N N N
		MeO NH O SH NH COOH
2-alkenyl furans	15	CO <sub>2</sub> Me S CO <sub>2</sub> Me O N N NH
		O H NH

N-alkylated aromatic (charged) nitrogen {often dyes}	17	N+ N+ OH N+ OMe N+ OMe N+ OMe N+ N+ OMe N+ N+ OMe N+
Conjugated and aryl tertiary amines	65	N N N N N N N N N N N N N N N N N N N
		H <sub>2</sub> N N N N N N N N N N N N N N N N N N N
Anisidines	17	Me OMe OMe O-nBu
2,6-dioxo pyridine	3	OH NO NO NO NO NO NO NO NO NO NO
Fused amino- pyridines	5	MeO NH NH CF <sub>3</sub>

DI I	2.1	
Phenylene-diamine,	21	HO HO
hydroquinone,		NIII Y. Y
catechol-like		
		OH N NH N
		HN NH <sub>2</sub>
		$\dot{\mathbb{N}}$
		$NH_2$
		CO <sub>2</sub> Et
		NH HN NH HN NH H2N NH
		H <sub>2</sub> N NH
		NH <sub>2</sub> N <sub>2</sub> N <sub>N</sub> N
Certain aryl	16	N-
sulfonamides		HO O O
with donating		
group (either end)		H Br H N
Cita)		H T
		N-O
		NC N N O O
		OMe H
Alpha keto	13	
and isatin-like		NH O O
hydrazones and imines		N O HN HN HN HN
and mines		CI NH NH NH NH
		S MeO
		· Web
4 4 1 1 1 1	12	" * '8
1,1-dinitrile / diketone /	12	NC NC
meldrum's		1 \( \sigma \) -(1
acids		NC ON NH
		CI NC N-NH
		CI (
		ОН
		но—
		N <sup>-3</sup> \ OA \\ \\ \>OH \\
		EtO <sub>2</sub> C =O N CF <sub>2</sub> O N O
		CF <sub>3</sub>
		a V

Biarylazo and	25	
conjugated	25	N-N OH
azo-imines		$N=N$ HO $H_2N$
		N=N N=N
		O O N HN OH  O S N H N N CN  O CI  O
		F
		N NH <sub>2</sub> N NH <sub>2</sub>
		S 1 H N=N
		N=N $N=N$ $N+N$ $N+1$ $N+1$
		N=N
Thiohydrazide	14	
and electron		NH \
rich phenyl- thioureas		$N = \langle \ \rangle_{NH} \rangle_{0}$
		S NH
		HN HN S HN S
		HN O HN O
		NEt <sub>2</sub> NEt <sub>2</sub>
		F HN N H
		HN S HN S OME HN S OME
Unsaturated	12	
cyclic hydrazides		N-N H
and esters		
		l N N
		HN AcN N
		F F
		S N-N H <sub>2</sub> N N-N
		HN HN HN
		OH N S
		`

Certain 2- amino- thiophenes	8	S NH <sub>2</sub> NC S NH HN S NH <sub>2</sub> SOOH
Five- membered unsaturated hydrazides and lactone- like with exocyclic alkene	18	H N O O O MeO O O
Benzothiazine and miscellaneous sulfur- containing compounds	12	$\begin{array}{c c} CO_2BzI & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & \\ & & & \\ & &$
		OMe HO N S NH S NH S N N N N N N N N N N N N N
		NH S CN CF <sub>3</sub>
		MeO O S MeO N N N OMe MeS N N

Allegaryl	Q	Cl
Alkenyl barbiturates	8	
Electron defficient Benzoxa- diazole-like	5	N-S, N-S, N-S, N-S, N-S, N-S, N-S, N-S,
Cyclic thioureas, thiocarbamate s	10	O NH NH HO N N N N N N N N N N N N N N N
		N S N S N S N S N S N S N S N S N S N S
Electron rich indoles, benzoxathiolones, benzimidazole s	9	MeO OS MeO OMe NH <sub>2</sub> NH <sub>2</sub> OMe OMe OMe NH <sub>2</sub> NH NH

Hydrazones of pyrrole aldehydes and similar	7	NH N
1,2,5- carbopyrroles	7	CF <sub>3</sub> CF <sub>3</sub> CF <sub>3</sub>
Certain dihydropyrans / pyridines	5	NC CN CO <sub>2</sub> Et O F O O O O O O O O O O O O O O O O O
Certain 1,3-di- t-Butyl aromatics	3	OH OH HO N'N'N
Azulenes	2	NC CN
Fused cyclopentene-quinolines	2	OCF <sub>3</sub> COOH  NH COOH  F
Certain 2,3- dialkyl indoles	4	NOH NH HN H

**Supplementary Table S5**. Assay counts for benign versus problematic groups showing the greater number that hit 2-6 assays when expressed as a percentage of those that hit no assays ("Enrichment")

ASSAY COUNT								SLN	Enrichment	
Group	6	5	4	3	2	1	0		2-6/0	
BENIGN STRUCTURES										
amide	30	76	109	185	1142	3384	19625	CNHC(=O)C	7.9%	
aminopyridine	0	3	4	5	18	34	167	C[1]:C(NH2):N:C:C:C:@1	18%	
benzothiazole	5	8	10	14	66	163	746	C[1]:N:C:S:C(:@1):C:C:C:C:@1	14%	
chlorophenyl	44	139	138	220	763	2030	12013	C[1]:C(Cl):C:C:C:@1	11%	
Aryl N	33	64	76	159	1191	1757	9621	C[1]:C:N:C:C:C:@1	16%	
nitrogenpyridine	3	11	13	28	127	278	1811	C[1]:C(N):N:C:C:C:@1	10%	
									Av. 13%	
			1	SUSPE	CTED	PROBL	EMATI	C STRUCTURES		
p_quinones	32	39	36	28	32	35	59	C[1]C(=O)C:=CC(=O)C:=@1	283%	
rhodanine	16	41	21	26	32	39	60	C=C[1]SC(=S)NC(=O)@1	234%	
rhodanines	17	50	30	41	85	109	305	C=C[1]SC(=Het)NC(=O)@1	41%	
Imidazolin-2,5-										
dione	4	7	8	9	13	5	0	C=C[1]C(=O)NNC(=O)@1	infinity	
123carbopyrrole		_			-	_			- <b>-</b> - 0 /	
S	4	5	10	0	6	2	4	C[1]:C(CH2):N(C:C):C(C:C):CH:@1	625%	
aminothiophenes	4	2	6	8	5	18	54	C[1]:C(NH2):S:C:C:@1	43%	

# TABLE S6 FILTER FAMILY A (Freq\_Hit\_5\_morethan150.hits #SYBYL/3DB HITLIST #@CLASS REGLIST #@DATABASE NONE #@SOURCE in-house #the numbers refer to number of members in the 93K wehi library #The first line in each case represents the historical name #sixes\_A7.txt 483 $C[1](C(\sim Het \sim CHetC@1=Het)\sim Het)=C[!r]H< regId="ene_six_het_A(483)">$ #BB\_OHzone\_2.txt 479 C[1]:C:C:C(:C(:C:@1)OH)C=NN<regId="hzone phenol A(479)"> #CC Me2 p anils 2.txt 478 CH2N(CH2)C[1]:C:C(Any[IS=H,CH2,OCH2CH2]):C(N):CH:C:@1<regId="anil\_di\_alk\_A(478)"> #indole\_3yl\_alk\_A.txt 461 #p\_quinones.txt 370 Het=C[1]C=:CC(=Het)C=:C@1<regId="quinone\_A(370)"> #HH\_azo.txt 324 $N[!r]=N< regId="azo_A(324)">$ #iminone\_3.txt 321 $CC(=Het[!r])C(=Het[!r])Any[IS=C,S(=O)=O] < regId="imine_one_A(321)" >$ #New\_Mannichs\_F.txt 296 NC[TAC=4]C[1]:C(OH):C:C:C:C:@1<regId="mannich\_A(296)"> #CC\_Me2\_N\_1.txt 251 C[1]:C(:C:C:C(:C:@1)C=C)N(C[TAC=4])C[TAC=4]<regId="anil\_di\_alk\_B(251)"> #CCpN\_OC\_1.txt 246 C[1]:C(:C:C:C(:C:@1)N(C[TAC=4])Any[IS=H,C[TAC=4]])OC[TAC=4]<regId="anil\_di\_alk\_C(246)"> #KK rhodaninesC 2.txt 235 $N[1]C(=S)SC(=C)C(=O)@1<regId="ene_rhod_A(235)">$ #BB\_OHzone\_1.txt 215 C[1](:C:C:C(:C:C:@1)C=NN)OH<regId="hzone\_phenol\_B(215)"> #fives\_K\_6NO.txt 201 $C[1](=C)C=NHetC(=O)@1<regId="ene_five_het_A(201)">$ #CC Me2 N 5.txt 198 $C[1]: C(:C:C:C:C:@1) C[TAC=4] \\ Any[IS=OH,C=CH,NC[TAC=4]] \\ N(C[TAC=4]) C[TAC=4] \\ < regId="anil_di_alk_D(198)"> regId="anil_di_alk_D(198)">$ #isatins\_1.txt 189 O=C[2]C(=[!R]NN)C[5]:C(:C:C:C:C:@5)N@2<regId="imine\_one\_isatin(189)"> #CC\_Me2\_C\_new.txt 186 CHN(CH2)C[1]:CH:C(Any[IS=H,CH2]):C(CHAny[IS=H,CH]):CH:CH:@1<regId="anil\_di\_alk\_E(186)"> #TABLE S7 FILTER FAMILY B (Freq\_Hit\_5\_lessthan150.hits #SYBYL/3DB HITLIST #@CLASS REGLIST #@DATABASE NONE #@SOURCE in-house #With number of members remaining (in parentheses) in WEHI library after Freq\_Hit\_5\_morethan150.filter has already been applied #For historical reasons, the early name is include before the sln with the revised regid name.

#158\_thiazene.txt 128 C[1](Any[IS=CH2,C:C])=C(Any[IS=H,CH2,C=O])SC(N@1(Any[IS=H,CH,C:C]))=N[!r]<regId="thiaz\_ene\_A(128)"> #pyrrole\_ArAlk.txt 118  $N[1](C[2]:Hev:C:C:C:C:@2)C(C[TAC=4]) = CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = "pyrrole_A(118)" > CCH = C(C[TAC=4]) @1 < regId = (118)" > CCH = C(C[TAC=4]) @1 < regId = (118)" > CCH = C(C[TAC=4]) @1 < regId = (118)" > CCH = C(C[TAC=4]) < regId = (118)" > CCH = C(C[TAC=4]) < regId = (118)" > CCH = C(C[T$ #BBhydroquin\_2.txt 92 C[1]:C:C:C(:C(:C:@1)OH)OH<regId="catechol\_A(92)"> #fives B2ene.txt 90  $C[1](=C)C(N=CS@1)=O< regId="ene_five_het_B(90)">$ #5quin.txt 89 C[1]C(=Het[!r])C(=O)HetHev=@1<regId="imine\_one\_fives(89)"> #furanone.txt 85  $C[1](C(C=CHet@1)=C)=Het<regId="ene_five_het_C(85)">$ #misc9.txt 79 CN[1]CH2CH2N(CH2CH2@1)N=CHC:Hev<regId="hzone\_pipzn(79)"> #diketo\_1.txt 68 C[1]:C:C:C:C:C(:@1)C(=O)C[TAC=4]C(=O)@1<regId="keto\_keto\_beta\_A(68)"> #pyrrole\_zone1.txt 64 N[1](C)CH=CHCH=C(C=NN)@1<regId="hzone\_pyrrol(64)"> #enone A1mod1.txt 57  $C=[!R]C(Hev)-[R]C(=[!R]Het)-[R]C(=[!R]C)Hev<regId="ene_one_ene_A(57)">$ #amin\_cyano\_1.txt 56 C(C#N)(C#N)C(NH2)=CC#N<regId="cyano\_ene\_amine\_A(56)"> #diketo ene 1.txt 55 C[1]:C:C:C:C:C(:@1)C(=O)C(=C)C(=O)@1<regId="ene\_five\_one\_A(55)"> #DD sixNS 4.txt 54 C[1](C#N)C(=S)NHHev=HevHev=@1<regId="cyano\_pyridone\_A(54)"> #EE\_tricycle.txt 51 C[1]:C:C:C[4]:C(:C:@1)NCC[9]C@4C=CC@9<regId="anil\_alk\_ene(51)"> #FF acridines.txt 46 C[1]:C:C:C:C[5]:C:@1:C(:C[8]:C(:N:@5):C:C:C:C:@8)N<regId="amino\_acridine\_A(46)"> #fives\_Nene.txt 46  $C[1](=C)C(=O)NNC(=O)@1<\text{regId}="ene_five_het_D(46)">$ #KK\_EE\_amino\_5hets\_1.txt 45 NH2C[1]:S:C(Hev):C(Hev):C(C=O):@1<regId="thiophene\_amino\_Aa(45)"> #oxind ene 1.txt 44  $NC = [!R]C[2]C(=O)C[5]:C(:C:C:C:C:@5)Het@2 < regId = "ene_five_het_E(44)" > (44) = ($ #DD sulfonOH.txt 43 C[1](:C(:C(:CH:C(:CH:@1)Hal)Hal)OH)S(=O)(=O)N<regId="sulfonamide\_A(43)"> ##S\_thiketone.txt 43 CC(=S)C<regId="thio\_ketone(43)"> #CCpsulfon OH.txt 41  $C[1]:C(:C:C:C:C:@1)NHS(=O)=O)OH < regId="sulfonamide_B(41)">$ #BB\_anils.txt 40 C[1](NH2):CH:CH:C(Any[IS=O,N,CH2]):CH:CH:@1<regId="anil\_no\_alk(40)"> #KK EE amino 5hets 3b.txt 40 Any[IS=H,CH2,C:C]C[1]:S:C(NHC(=O)C):C(C(=O)O):C(Any[IS=C[2]:C:C:C:C:@2,C[2]:S:C:C:C:@2]):@1<regId="thiophene\_amino\_Ab(40)"> #pyridiniums\_A.txt 39 N[1:+1](Any[IS=CH3,O[TAC=1],CH2CH=CH2,CH2CH2OH,CH2C(=O)C,CH2C(=O)NHC:C,CH2CH3]):C:C:Hev:C[2]:C(:@1):CH:C(Any[IS=H,N]):C:C:@2<regId="het\_pyridiniums\_A(39)"> #anthrone.txt 38 C[1]:C(:C(:C:C:C:@1)C(C:C)=O)NHAnv[NOT=C=O]<regId="anthranil one A(38)"> #FF\_cyano\_3.txt 37 NHN=C(C#N)C=Het[!r]<regId="cyano\_imine\_A(37)"> #EE\_diazox\_2.txt 36 #DiMeAnil\_zone.txt 35 #KK\_rhodaninesC\_6.txt 33  $N[1]C(=S)SC[TAC=4]C(=O)@1<regId="rhod_sat_A(33)">$ #EE\_enamine\_2.txt 30  $NHN=CC(Any[IS=H,C])=C(C)-[!R]Any[IS=N,OH] < regId="hzone_enamin(30)"> regId="hzone_enamin(30)"$ #pyrrole\_ArAr.txt 29  $N[1](C[2]:Hev:C:C:C:C:@2)C(C[TAC=4]) = CCH = C(C:C)@1 < regId = "pyrrole_B(29)" > CCH = C(C:C)@1 < regId = "p$ #FF\_OH\_thiophene.txt 28  $S[1]C = CC(=C@1)OH < regId = "thiophene_hydroxy(28)" >$ #FF\_cyano\_8.txt 27  $C[1](=C(C(=O)NC(=N@1)Het)C\#N)C< regId="cyano_pyridone_B(27)">$ #barb\_imine\_1.txt 27  $C[1](C(=O)NC(=O)NC(=O)@1)=N<\text{regId}=\text{"imine_one_sixes}(27)">$ 

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#dyes5A.txt 27
CH2N(:-C:C)~C-:C=:CC~C-:N<regId="dyes5A(27)">
#FF Naphth 4.txt 25
C[1]:C:C:C[4]:C[5]:C(:C:C:C:C:@1:@5)N=CN@4<regId="naphth_amino_A(25)">
#FF_Naphth_1.txt 25
C[1]:C:C:C[4]:C[5]:C(:C:C:C:C:@1:@5)NHC[TAC=4]NH@4<regId="naphth_amino_B(25)">
#enone 2.txt 24
CC(=O)CH=C(NHC)C(=O)OC<regId="ene_one_ester(24)">
#misc3.txt 23
S=C[1]C=:CHetC=:C@1<regId="thio_dibenzo(23)">
#cyanos_1.txt 23
C(C#N)(C#N)C(Any[IS=C#N,C=N])C#N<regId="cyano_cyano_A(23)">
#naphthol_A.txt 22
C[1]:C(:CH:CH:CH:CH:@1):CH:C(:C(:CH:@1)OH)C(=O)NHN=C < regId="hzone_acyl_naphthol(22)"> regId="hz
#misc11.txt 21
O=CC[1]=C[2]N=C(CH2)C=C(OH)N(@2)N=C@1<regId="het_65_A(21)">
#AAmisc2.txt 19
N[1]:C(C[2]:C:C:C:C:C:@2):C(C[3]:C:C:C:C:@3):NH:C(C:Hev):@1<regId="imidazole_A(19)">
#ene_cyano_1.txt 19
C(C#N)(C#N)=CC[1]:C:C:C:C:C:@1<regId="ene_cyano_A(19)">
#GG_acidzone.txt 19
C[1](C(=O)OH):C(NHN=C):C:C:C:C:@1<regId="anthranil_acid_A(19)">
#dyes3A.txt 19
N[+1](:-C:C):=CCH=CN(C[TAC=4])C<regId="dyes3A(19)">
#dhp_bis_amino_CN.txt 19
NH2C[1]=C(C#N)CH(C:C)C(C#N)=C(NH2)S@1<regId="dhp_bis_amino_CN(19)">
#BBparanils_2_mod.txt 18
N~C[1]:Hev[IS=N,C(N)]:Hev[IS=N,CH,C(NH)]:C(Hev[IS=NH,OCH2]):N:N:@1<regId="het_6_tetrazine(18)">
#EE Xenone.txt 17
CC=C(Hal)C(=O)C<regId="ene_one_hal(17)">
#imino_cyano_1.txt 17
C(C#N)(C#N)=NNHC[1]:C:C:C:C:@1<regId="cyano_imine_B(17)">
#158_thiazene_A5b.txt 17
C[1](Any[IS=NC(=0)C:C,NH2])=C(-[!R]C(=0)NCH2)SC(N(@1)Any[IS=CH2CH=CH2,C:C])=S<regId="thiaz_ene_B(17)">
#P rhodanines 4E.txt 16
S[1]C(=O)NC(=O)C(=CHAny[IS=CBr,C:CH:C(Hal):C:CHal,C:CH:CSCH2,C:C:C:C:C:C:C:C:C:C:CCCH2,C[2]:C(CH2):N(CH2):C(CH2):C:@2])@1<regId="ene_rhod_B(16)">
#DD_65_OOS_1_1.txt 15
O[1]C(SC[4]:C:C(:C:C:@1:@4)Any[IS=N,O])=Any[IS=O,S]<regId="thio_carbonate_A(15)">
#KK_EE_amino_5hets_10a.txt 15
N(CH2)(CH2)C[1]:O:C(C=NNHC(=Het)):CH:CH:@1<regId="anil_di_alk_furan_A(15)">
#oxinde ene 4.txt 15
C[1](:C:C:C:C:@1)CH=[!R]C[2]C(=O)C[5]:C(:C:C:C:@5)S@2<regId="ene_five_het_F(15)">
#TABLE S8
FILTER FAMILY C (Freq_Hit_5_lessthan15.hits
#SYBYL/3DB HITLIST
#@CLASS REGLIST
#@DATABASE NONE
#@SOURCE in-house
#Numbers in parenthases refer to hits on database Lessthan15
#For historical reasons, the early name is include before the sln with the revised regid name.
#Me2 N 5.txt 14
C[1]:C:C:C:C(N(Any[IS=H,C[TAC=4]))(Any[IS=H,C[TAC=4])):C:@1)C[TAC=4]:C:C:C:C:(C:C:C:@23)N(Any[IS=H,C[TAC=4]))(Any[IS=H,C[TAC=4]))\\
#CC anilzone.txt 14
C[1](:CH:CH:C(NH2):CH:CH:@1)C=NNH<regId="hzone_anil(14)">
#pyraz OH.txt 14
C[1](C[TAC=4])C(Any[IS=H,CH])=C(OH)N(C[2]:CH:CH:CH:CH:CH:@2)N=@1<regId="het_5_pyrazole_OH(14)">
#JJ_DD_phenothz.txt 13
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C[1]H:CH:CH:CH:C|5]N(Any[IS=H,CH3,CH2CH])C[7]:C(~Any[IS=H,NHC[TAC=4],C:C]):C(Any[IS=H,C:C]):C(Any[IS=H,NH2,OC[TAC=4]]):CH:C:@7SC:@1:@5<regId="het\_thio\_666\_A(13)"> #GG tricvcle.txt 13 C[1]CC[3]:C:C:C:C:@3C(C[10]:C@1:C:C:C:C:@10)=CC<regId="styrene\_A(13)"> #P rhodaninesC 4CB.txt 13 S[1]C(=NC:C)N(Any[IS=H,CH2CH2O,C:C])C(=O)C(=CHAny[IS=C:C:CCl,C:Het])@1<regId="ene\_rhod\_C(13)"> #dhp\_amino\_CNene.txt 13 NH2C[1]=C(C#N)CH(C:C)C(CH2)=C(C=C)O@1<regId="dhp\_amino\_CN\_A(13)"> #FF\_cyano\_12.txt 12 O=S(=O)C(C#N)=NNH<regId="cyano\_imine\_C(12)"> #KK\_thiourea\_3.txt 12 C[1]:C(NHC(=S)NHCH2CH2CH2N(CH2)C[2]:C:C:C:C:C:@2):C:C:C:C:@1<regId="thio\_urea\_A(12)"> #KK EE amino 5hets 2.txt 12  $C[2]: C: C: C: C: C: C: (2) NHC[1]: S: C(Anv[IS = C = O, C # N, C(OH) = C]): C(N): C(Anv[IS = C \# N, C(:N): N]): @1 < regId = "thiophene_amino_B(12)" > (2) C(N): C(N)$ #fives\_Ntac4.txt 12  $C[1:TAC=4]C(=O)NNC(=O)@1<regId="keto_keto_beta_B(12)">$ #DD suber.txt 11 C[1]:C:C:C:C[5]:C:@1C(C[8]:C@5:N:Hev:C:C:@8)=O<regId="keto\_phenone\_A(11)"> #sixes A6.txt 11  $C[1](C(CH2)=C(C\#N)C(\sim O)\sim N\sim C(\sim O)@1)=CHC:C< regId="cyano_pyridone_C(11)">$ #158\_thiazene\_A2.txt 11  $C[1](Any[IS=Hal,N[+1](:C):C])=C(-[!R]C=N)SC(N@1)=O<regId="thiaz_ene_C(11)">$ #KK\_DD\_thiophenezone\_1A.txt 11 C[1]:CH:CH:CH:CH:C:(@1):Het:C(Any[IS=H,OH,CH2]):C(:@1)CH=NNHAny[IS=C[2]:N:C:CH:S:@2,C:CH:CH,C:N:C:N;C:N;C:N;C:N;N:N:N] #FF cvano 18.txt 10  $Hev: HevC(Any[IS=H,C\#N]) = C[1]C: = CC(=Any[IS=O,N[!r]])C: = C@1 < regId="ene_quin_methide(10)"> C(1)C: = C(1$ #DD\_phenothz\_3.txt 10 C[1]:C:C:C[4]:C:C:C[4]:C(:C:@1)SC[8]:CH:CH:C(:CH:C:@8C(C@4)N(Any[IS=H,C[TAC=4]])(Any[IS=H,C[TAC=4]])Any[IS=H,C[TAC=4]])Any[IS=H,C[TAC=4]])Any[IS=H,C[TAC=4]])Any[IS=H,C[TAC=4]]Any[IS=H,C[TAC=#KK\_110\_fives\_6A 10 C[1](=O)C(=CHAny[IS=C[2]:C:C:C:C:@2,C[2]:C:C:C:Het:@2])N=C(Hev:Hev:Hev)Any[IS=S,NHev:Hev]@1<regId="ene\_five\_het\_G(10)"> #JJ\_ureimiums.txt 9  $N[+1](:Hev:Hev:Hev)Hev=O< regId="acyl_het_A(9)">$ #KK\_CC\_ME2\_p\_im\_1D.txt 9 C[TAC=4]N(C[TAC=4])C[1]:CH:CH:C(C[2]:=N:-C:C:Hev:-@2):CH:CH:@1<regId="anil\_di\_alk\_G(9)"> #JJ FF dhp 1.txt 9 N[1](Any[IS=C[TAC=4],H])C[2]:=C(C(=0)C:C:C)C(C)C(Any[IS=C=0,C#N])=C(CH3)@1<regId="dhp\_keto\_A(9)"> #KK thiourea 4.txt 9 C[1]:C(NHC(=S)NHCH2CH2N(CH2)C[2]:C:C:C:C:C:@2):C:C:C:C:@1<regId="thio\_urea\_B(9)"> #JJ bim.txt 9 C[1]:CH:CH:C(NHCH2C[2]:C:C:C:C:@2):CH:C(:@1):N:CH:N(C):@1<regId="anil\_alk\_bim(9)"> #GG indolene.txt 9 C[1]:C:C:C[4]:C(:C:@1)C(C=N@4)=N[!r]<regId="imine\_imine\_A(9)"> #thiocarbz 2.txt 9 C[1](:C:C:C:C:C:@1)NHC(=S)NNHC(=O)C[2]:Hev:Het:C:C@2<regId="thio\_urea\_C(9)"> #isatins 2.txt 9 N[!r]=C[2]C(=O)C[5]:C(:C:C:C:C:@5)S@2<regId="imine one fives B(9)">#dhp amino CNpyrz.txt 9 #CC\_ON\_3.txt 8 NH2C[4]:CH:N:C(OC[10]:C:C:C:C:@10):CH:CH:@4<regId="anil\_OC\_no\_alk\_A(8)"> #DD phenothz 2.txt 8 C(=O)C[3]SC[5]:C:C:C:C:C:@5NC=@3<regId="het\_thio\_66\_one(8)"> #DD\_phenothz\_4.txt 8  $C[1]:C:C:C[4]:C(:C:@1)\\ Any[IS=S[TAC=2],C[TAC=4]]\\ C[8]:C:C:C(:C:C:@8C@4=CC)\\ Any[IS=H,Cl,C[TAC=4]]\\ < regId="styrene_B(8)">1000$ #KK\_107\_SIM.txt 8 CH2S[TAC=2]C[1]:N(-[!R]CH2):C(C[2]:C:C:C:C:C:@2):CH:N:@1<regId="het\_thio\_5\_A(8)"> #NN\_CC\_Me2\_N\_2.txt 8 CH2N(CH2)C[1]=CHC[2]:C:C:C:C:C(:@2)S[TAC=2]C[3]:C:C:C:C:C(:@3)@1<regId="anil\_di\_alk\_ene\_A(8)"> #P\_rhodaninesC\_4CA.txt 8  $S[1]C(=[!R]NAny[IS=H,NHC:C])N(Any[IS=H,C:N:C:C:S])C(=O)C(=CHC:C(Any[IS=C1,OCH])) @ 1 < regId="ene_rhod_D(8)" > 1$ #P\_rhodaninesC\_4D.txt 8  $S[1]C(=O)NC(=S)C(=CHC:C)@1<regId="ene_rhod_E(8)">$ #p\_BnNH\_phenol.txt 8 C[1]:C:C:C:C:C(:@1)CH2NHC[2]:CH:CH:C(OH):CH:CH:@2<regId="anil\_OH\_alk\_A(8)"> #pyrrole\_AlkAralk.txt 8  $N[1](C[TAC=4])C(CH2)=CHCH=C(C:C)@1<regId="pyrrole_C(8)">$ #thiocarbz\_1.txt 8 C[1](:C:C:C:C:C:@1)NHC(=S)NNHC[2]:C:C:C:C:@2<regId="thio\_urea\_D(8)"> #AAthiazole.txt 8

N(C[2]:C:C:C:C:@2)C[8]SC=C(N[charge=+1]=@8)C[13]:C:C:C:C:@13<regId="thiaz\_ene\_D(8)"> #pyrrole\_ene2b.txt 8  $N[1]:C:C:C(CH=C[2]C(=O)NC(=Het)N@2):C(CH2):@1<regId="ene_rhod_F(8)">$ #158 thiazene C1.txt 8  $C[1](SC[r])=C(CH(C)C)SC(N@1(Any[IS=H,CH2]))=O< regId="thiaz_ene_E(8)">$ #fives\_O.txt 7 Hev[1]:Hev[3]:Hev(:Hev:Hev:Hev:@1)NHN(C@3=O)C<regId="het 65 B(7)"> #FF misc 3.txt 7  $C[1]:C:C:C[7]:C(:C:@1)C(C(=C@7OH)C)=O < regId="keto_keto_beta_C(7)">$ #FF\_misc11.txt 7 C[1]:C:C:C[7]:C(:C:@1):N:N:C(:N:@7)CH2C=O<regId="het\_66\_A(7)"> #KK thiourea 6.txt 7 C[1]:C(NHC(=S)NHCH2C[2]:N:C:C:C:C:@2):C:C:C:@1<regId="thio\_urea\_E(7)"> #KK\_EE\_amino\_5hets\_6a.txt 7  $CHCH2C[1]:S:C(NHC(=O)CCC=O):C(Any[IS=C(=O)O,C\#N]):C(CH2):@1< regId="thiophene_amino_C(7)">$ #AA misc11.txt 7 #acid ar rhod.txt 7  $HOC(=O)C[1]:C:C:C:C:C:@1)C:Hev:CCH=C[2]C(=Het)NC(=Het)Het@2 < regId="ene_rhod_G(7)" > regId="ene_rho$ #FF\_cyano\_13.txt 7  $C[1](C[2]:C(:C:C(:C:N:@2)NH2)C\#N)C(C=@1)=C)C\#N < regId="ene_cyano_B(7)"> regId= regId$ #dhp\_amino\_CNCN.txt 7 NH2C[1]=C(C#N)CH(C:C)C(C#N)=C(C:C)O@1<regId="dhp amino CN C(7)"> #166\_het\_5\_q.txt 7  $N[1](C[2]:C:C:C:C:C:@2)N = C(C=O)C[TAC=4]C(=O)@1 < regId = "het_5\_A(7)" > C(C=O)C[TAC=4]C(C=O)@1 < regId = "het_5\_A(7)" > C(C=O)C[TAC=4]C(C=O)C$ #fives\_K\_11.txt 6  $N[1]=CC(C(N@1)=S)=C<regId="ene_five_het_H(6)">$ #EE\_5hets10.txt 6 C[1]H=COC(=CH@1)C(=S)N[10]CH2CH2HevCH2CH2@10<regId="thio\_amide\_A(6)"> #FF cyano 2.txt 6  $C=C(C#N)C(=NH)NN< regId="ene_cyano_C(6)">$ #KK\_111\_furzone\_1.txt 6 C[1](:O:C(Any[IS=H,CH2]):CH:CH:@1)C(Any[IS=H,CH2])=NNHC[2]:N:C:C:S:@2<regId="hzone\_furan\_A(6)"> #KK sulfonam 1.txt 6 C[1]H:C(NHS(=O)(=O)C[2]:C:Hev:C:C:C:@2):CH:C(N(CH2)CH2):CH:CH:@1<regId="anil di alk H(6)"> #FF\_Ncycle.txt 6 N[1]C(=C(NC:C)N[6]C=CC=CC=@1@6)Any[IS=C(Hev)=CHC:C,C:O:C]<regId="het\_65\_C(6)"> #thiocarbz 5.txt 6 C[1]NHNHC(=S)NNH@1<regId="thio\_urea\_F(6)"> #oxind ene 2.txt 6 C[1](:C:C:C:O:@1)CH=[!R]C[2]C(=O)C[5]:C(:C:C:C:C:@5)Het@2<regId="ene five het I(6)"> #diketo B.txt 5  $O=C[2]C[TAC=4]CC(=O)C[7]:C@2:C:C:C:@7<regId="keto_keto_gamma(5)">$ #5quin2.txt 5 C[1]:C:C:C[4]:C(:C:@1)C[7]=C[8]C(C@4=O)=CC=CC@8=NO@7<regId="quinone B(5)"> #DD OHPyr 1.txt 5 HOC[2]:N:C(OH):C:C:C:@2<regId="het\_6\_pyridone\_OH(5)"> #naphth\_D.txt 5 C[1]:C(:CH:CH:CH:CH:@1):CH:C(:CH(:CH:@1))C=NNHAny[IS=C:C,C=S]<regId="hzone\_naphth\_A(5)"> #DD thioest 5.txt 5  $C[1]=C(SC(C=C@1)=S)N< regId="thio_ester_A(5)">$ #EE\_ketal.txt 5  $C[1]C(OC(C=@1)(O)C)O < regId = "ene_misc_A(5)" >$ #FF\_cyano\_17.txt 5 O=C[2]C(=CC(=NN@2)C=O)C#N<regId="cyano\_pyridone\_D(5)"> #FF\_misc\_7.txt 5 C[1]C=CC[4]N(C=@1)C(NC)=C(N=@4)C[12]:C:C:C:C:N:@12<regId="het\_65\_Db(5)"> #FF\_Ncycle\_2.txt 5 N[1]C[2]:C(C(=N)C[6]:C:C:C:C:C@1:@6):C:C:C:C:@2<regId="het\_666\_A(5)"> #KK\_108\_diazox\_3a.txt 5 #KK\_113\_dianiline\_2.txt 5  $C[1](NHCH2):CH:CH:C[2]:C(NHC(=O)NH@2):CH:@1<regId="anil_NH_alk_A(5)">$ #KK\_sulfonam\_2.txt 5  $C[1]H:C(NHS(=O)(=O)C[2]:C:C:C(Het):C:C:@2):CH:C[3]:C(OCH2O@3):CH:@1 < regId="sulfonamide_C(5)"> regId="sulfonamide_C(5)"$ #KK\_107\_SIM\_4a.txt 5 CHC[1]:N:N(C[3]:C:C:C:C:C:@3):S[2]:Het:Hev:C:C(:@2):@1<regId="het\_thio\_N\_55(5)"> #DD\_enone\_1.txt 5

O=CC=CHOH<regId="keto\_keto\_beta\_D(5)"> #P rhodaninesC 4B.txt 5 N[1]C(=NC(=O)C(CH2)=N@1)SC(=CHC:C)C(=O)@1<regId="ene\_rhod\_H(5)"> #chalcimine.txt 5 C:CCH=CHCH=NN(C[TAC=4])C[TAC=4]<regId="imine\_ene\_A(5)"> #carbaz\_NSallyl.txt 5 C[1]:C:C:C:C:C(:@1):C[2]:N:N:C(SCH2C=O):N:C(:@2):N(CH2CH=CH2):@1<regId="het\_thio\_656a(5)"> #pyrrole\_thiourea.txt 5  $N[1](C)CH=CHCH=C(CH2NHC(=S)NH)@1<regId="pyrrole_D(5)">$ #pyrrole\_HetAr.txt 5 N[1](C[2]:CH:Hev:Het:Hev:@2)C(C[TAC=4])=CHCH=C(C[TAC=4])@1<regId="pyrrole\_E(5)"> #thiocarbz 4.txt 5 C[1](:C:C:C:C:C:@1)NHC(=S)NNHC(=:N[r])-:N[r]<regId="thio\_urea\_G(5)"> #KK\_113\_dianisole\_2.txt 5 #pyrrole\_ArCN\_1.txt 5 N[1](C[2]:C(C#N):C:C:Het:@2)CH=CHCH=C@1<regId="pyrrole\_F(5)"> #dhp amino CNS.txt 5 NH2C[1]=C(C#N)CH(C:C)C[2]:C(:C:C:S:@2)O@1<regId="dhp\_amino\_CN\_D(5)"> #GG\_thiophene\_3.txt 4 NHC[1]:N:C(C[2]:C:N:C(NH2):S:@2):C:S:@1<regId="thiazole\_amine\_A(4)"> #BBparanils\_2\_1.txt 4 N=C[1]NHC(NH)=C(NH)N=N@1<regId="het 6 imidate A(4)">#CCpCN O2.txt 4  $C[1]:C:C:C[4]:C(:C:@1OC[8]:CH:CH:C(:CH:CH:@8)NH):C:C:C:C:@4 < regId = "anil_OC_no_alk_B(4)" > regId = "anil_OC_no_alk_B(4)"$ #DD\_phenothz\_5.txt 4 C[1]:C:C:C[4]:C(:C:@1)C(C[8]=C(SC@4)SC=C@8)=C<regId="styrene\_C(4)"> #DD\_apine\_1.txt 4 C[1]:C:C:C[4]:C(:C:C:@1):C:C:C:@4<regId="azulene(4)"> #EE 5hets29.txt 4 C[1](:O:C(CH2):C(CH2OC:C):CH:@1)C(=O)OH<regId="furan\_acid\_A(4)"> #FF\_cyano\_19.txt 4 Hev:CC[1]=CHC(C:O)=C(C#N)C(=O)NH@1<regId="cyano\_pyridone\_E(4)"> #FF SS.txt 4 C[1]=C(C(NC[5]:C@1:C:C:C:C:@5)(C)C)SSC@1=Hev<regId="anil\_alk\_thio(4)"> #KK benzanilid 1.txt 4 C[1]H:C(NHC(=0)C[2]:C:C:C:C:@2):CH:C(N(CH2)CH2):CH:CH:@1<regId="anil\_di\_alk\_I(4)"> #NN sixesB8 2.txt 4 CH2S[TAC=2]C[1]:N:C(C[2]:O:CH:CH:CH:@2):C(C[3]:O:CH:CH:CH:@3):N:N:@1<regId="het\_thio\_6\_furan(4)"> #NN CC Me2 N 1.txt 4 CH2N(CH2)C[1]=CC[2]:C:C:C:C:C:(:@2)CH2@1<regId="anil di alk ene B(4)"> #NN DD NNCON 1 1a.txt 4 NH(C[1]:C:C:C:C:C:@1)N=C(C(=0)CH2)NHAny[IS=NH,C:C]<regId="imine\_one\_B(4)"> #P fur anis.txt 4 C[1]:CH:CH:CH:CH:C(:@1):O:C[2]:CH:C(NHCH2):C(OCH2):CH:C(:@2):@1<regId="anil OC alk A(4)"> #KK 116 thipyr new.txt 4 S=C[1]NHC=CC[2]=C(@1)C(=O)OC(@2)=CH<regId="ene\_five\_het\_J(4)"> #pyrrole\_ArH.txt 4 N[1](C[2]:C:CH:C(Any[IS=NH2,C:N]):C:CH:@2)CH=CHCH=CH@1<regId="pyrrole\_G(4)"> #pyrrole ene.txt 4 N[1](C)CH=CHCH=C(CH=C[2]C(=O)HetC=:Hev@2)@1<regId="ene\_five\_het\_K(4)"> #amino\_cyano\_2a.txt 4 C=CC(C#N)(C#N)C(C#N)=CNH2<regId="cyano\_ene\_amine\_B(4)"> ##Ar\_thiono\_S.txt 4 C:CC(=S[f])S[TAC=2]CHAny[IS=CH2,C:C]<regId="thio\_ester\_B(4)"> #isatin\_ene.txt 4 O=C[1]C(=[!R]CHC[3]:C:C:N:C:@3)C[2]:C(:C:C:C:C:@2)N@1<regId="ene\_five\_het\_L(4)"> #KK\_111\_thiophzone\_2a.txt 4 #dhp\_amino\_CNone.txt 4 CH2S[f]C[1]=C(C#N)CH(C:C)C(C#N)C(=O)N@1<regId="dhp\_amino\_CN\_E(4)"> #166\_het\_5\_u.txt 4  $N[1](C[2]:C:C:C:C:@2)N=C(NHC=O)CH2C(=O)@1<regId="het_5_B(4)">$ #JJ\_FF\_misc\_10\_B.txt 3 C:CCH=CHCH=NN=C<regId="imine\_imine\_B(3)"> #fives\_V.txt 3 C[1](:C:C:C(CH2):C:C:@1)C[2]:N:C(NH2):S:C(CH3):@2<regId="thiazole\_amine\_B(3)"> #sixesB3.txt 3

C[1](C=NC[4]:C(N@1):C:C:C:C:@4)=CHC=O<regId="imine\_ene\_one\_A(3)"> #EE diazox 1a.txt 3 O(C[2]:C:C:C:C:C:@2)C[3]:C:C[1]:N:O:N:C(:@1):C:C:@3<regId="diazox\_A(3)"> #chalc1.txt 3 Hev[1]:Hev:Hev:Hev:Hev:@1)CH=CHC(NC[13]:C:C[15]:C(:C:C:@13):N:C(:C:C:@15)N(C)C)=O<regId="ene\_one\_A(3)"> #BB\_o\_anils\_3.txt 3 NH2C[1]:N:C:C:C:C(OCH2C:C):@1<regId="anil\_OC\_no\_alk\_C(3)"> #DD S thiaz2.txt 3 CS[TAC=2]C[1]:N:CH:C:S:@1<regId="thiazol\_SC\_A(3)"> #DD\_phenox.txt 3 C[1]:C:C:C:C[9]:C:@1OC[12]:C(N@9CH2CH2):C:C:C:C:@12<regId="het\_666\_B(3)"> #EE 5hets21.txt 3 C[1](:O:C(CH2):CH:CH:@1)CH(OH)C#CC[TAC=4]<regId="furan\_A(3)"> #FF\_azep.txt 3 C[1](C(=CC=CC=C@1)NH)=NC<regId="colchicine\_A(3)"> #JJ EE enamine 1.txt 3 CH2N(CH2)CH=CC(=O)C[1]:C(S[TAC=2]):S:C(Any[IS=C#N,C=O]):C:@1<regId="thiophene\_C(3)"> #JJ enone Da.txt 3 #KK 107 SIM 3a.txt 3 C[1]:C:C:C:C:C(:@1)CH2CH2N=C(@1)S[TAC=2]CH2C(=O)C[2]:C:C:C:C:C:@2<regId="het\_thio\_66\_A(3)"> #KK 110 rhod 6.txt 3 N[1](C[2]:C(CH2):C:C:C:C:@2)C(=S)N(CH2Hev:Hev:Hev:Hev:Hev)CH2C(=O)@1<regId="rhod sat B(3)"> #KK 110 rhod 9.txt 3 N[1](CH2)C(=S)NHC(=CHC[2]:C:C(Br):C:C:C:@2)C(=O)@1<regId="ene\_rhod\_I(3)"> #KK\_111\_benzthioph\_1.txt 3 C[1](:S:C[2]:C:C:C:C:@2:C(CH2):@1)C(=O)CH2CH2<regId="keto\_thiophene(3)"> #KK\_113\_polyimine\_3.txt 3 N(CH2)(CH2)CH=NC(CH2)=NN(CH2)C:C<regId="imine\_imine\_C(3)"> #KK\_114\_hetpyrdones\_1.txt 3 C[1](CH2CH2):C(CH2):C[2]C(=Het[TAC=1])N(CHAny[IS=C(=O)O,C:C])C(Any[IS=H,SCH2])=NC(:@2):Het[TAC=2]:@1<regId="het\_65\_pyridone\_A(3)"> #KK\_115\_Furthiaz\_1a.txt 3 C[1](NHCH2C[2]:CH:CH:CH:O:@2):N:C(Hev:Hev:Hev(Any[IS=OCH2,CH2]):Hev:Hev):CH:S:@1<regId="thiazole\_amine\_C(3)"> #KK 116 thipyr 1a.txt 3 N[1]:C(SCH):C(C#N):C(C[2]:C:C:C(OCH2):C:C:@2):CH:C(C:C):@1<regId="het\_thio\_pyr\_A(3)"> #KK\_108\_melamime\_3.txt 3 #KK 113 dianiline.txt 3 NH(C[1]:C:C:Hev:C:C:@1)C[2]:C:C:(NHCH):C:C:@2<regId="anil\_NH\_alk\_B(3)"> #KK 110 rhod 3.txt 3 N[1](C[2]:C:C:C:C:@2)C(=NC=O)SCH2C(=O)@1<regId="rhod sat C(3)"> #KK EE amino 5hets 8a.txt 3 C=CC(=O)NC[1]:S:C(C(=O)O):C(CH):C(C#N):@1<regId="thiophene\_amino\_D(3)"> #KK CC o anisidine 5C.txt 3 Any[IS=H,CH2]OC[1]:CH:CH:CH:CH:C(:@1)NHCH2C[2]:N:C:C:N:@2<regId="anil OC alk C(3)"> #NN sixesB8 1.txt 3 CH2S[TAC=2]C[1]=NC[2]C(CH2)=NN(C[3]:C:C:C:C:@3)C(=@2)N=N@1<regId="het\_thio\_65\_A(3)"> #NN\_sixesB8\_3.txt 3 CC(=0)CH2S[TAC=2]C[1]:N:C[2]:NH:C[3]:CH:CH:CH:CH:C(:@3):C(:@2):N:N:@1<regId="het\_thio\_656b(3)"> #NN JJ thiziniums 1.txt 3 S[1]:C(NHC[2]:C(-:Any[IS=CH2,C:C]):C:C:C:C:@2):N[+1](CH2):C(C):CH:@1<regId="thiazole\_amine\_D(3)"> #M1\_DD\_NNNS2a.txt 3 C[1](=S)N(CH2C[2]:O:C:C:C:@2)C(C:C)=NNH@1<regId="thio\_urea\_H(3)"> #KK\_116\_pyr\_2.txt 3  $N[1](C[2]:C:C:C:C:@2)C(=O)C(C\#N) = CC(C\#N) = N@1 < regId = "cyano_pyridone_F(3)" > 1 < regId = "cyan$ #KK\_110\_rhod\_8.txt 3 N[1](C[2]:C:C:C:C:C:@2)C(=O)SCH(CH2C(=O)NHC:C)C(=O)@1<regId="rhod\_sat\_D(3)"> #P\_rhodaninesC\_4A.txt 3  $CH2N[1]C(=Any[IS=S,N])HetC(=C[2]CH=CHC:CN(CH2)@2)C(=O)@1 < regId="ene_rhod_J(3)" > regId="ene_rhod_J$ #iminophenol.txt 3 C=N[!r]C[3]:C(OH):C:C:C:C:@3<regId="imine\_phenol\_A(3)"> #JJ\_EE\_CNene\_1.txt 3 O=C[1]SC[2]:C:C:C:C(OCH2):C(:@2)O@1<regId="thio\_carbonate\_B(3)"> #QQ\_EE\_NS\_1.txt 3  $N=C[1]N=CNS@1<regId="het_thio_N_5A(3)">$ #QQ\_EE\_NS\_2.txt 3 N[1]SC[2]CNC:CC(=@2)C(=S)@1<regId="het\_thio\_N\_65A(3)"> #QQ\_CC\_ME2\_pim\_1E.txt 3

CH2N(CH2)C[1]:CH:CH:C(CH=NN=C(C)C:C):CH:CH:@1<regId="anil\_di\_alk\_J(3)"> #pyrrole\_iminium.txt 3  $N[1]C=CC=C(C=N(C)CC@1)@1<regId="pyrrole_H(3)">$ #ene cyano.txt 3  $C(C#N)(C#N)=C(S)S < regId = "ene_cyano_D(3)" >$ #cyano\_cprop\_1a.txt 3 C[1](C#N)(C#N)CH(C(=O)C)CH@1<regId="cyano\_cyano\_B(3)"> #fives\_I.txt 3  $C[1]Any[IS=O,S]C(C(C=:@1)=O)=CC=O<regId="ene_five_het_M(3)">$ #KK\_116\_conjugcyano.txt 3 C:CC(=O)NHC(=O)C(C#N)=CHNHC:C<regId="cyano\_ene\_amine\_C(3)"> #thiocarbz 6.txt 3 C[1](:C:C:C:C:C:@1)NHC(=S)NHN=CC[2]:C:C:N:C:@2<regId="thio\_urea\_I(3)"> #dhp\_amino\_CNCO.txt 3 NH2C[1]=C(C#N)CH(C[2]:C:C:C:S:@2)C(C(=O)OC)=C(CH2)O@1<regId="dhp\_amino\_CN\_F(3)"> #KK phthalimide 3.txt 3 C[3]:C:C(C(=O)NHC[4]:C:C:C:C(C(=O)OH):@4):C:C[1]:C(:@3)C(=O)N(CH2)C(=O)@1<regId="anthranil\_acid\_B(3)"> #EE diazox 3.txt 3 ClC[2]:C:C[1]:N:O:N:C(:@1):C:C:@2<regId="diazox\_B(3)"> #misc5.txt 3 CC(=S)H<regId="thio\_aldehyd\_A(3)"> #KK\_FF\_misc\_10\_1.txt 2 C[TAC=4]NHC(C:C)=CHC(=S)NHC[1]:C:C:C:C:C:@1<regId="thio\_amide\_B(2)"> #fives U.txt 2 CH2CH2SCH2C[1]N=CNHC=@1<regId="imidazole\_B(2)"> #fives\_W.txt 2 O=CNHC[1]:C(C:C):N:C(CH2C#N):S:@1<regId="thiazole\_amine\_E(2)"> #fivesY.txt 2 CHNHC[1]:N:C(C[2]=CN=C[3]N(C=CS@3)@2):C:S:@1<regId="thiazole\_amine\_F(2)"> #sixesB10.txt 2  $N[1]C(=O)CH=C(C)SC(=S)@1<regId="thio_ester_C(2)">$ #enone\_E.txt 2 C(S)(N)=CHC=CHC=O<regId="ene\_one\_B(2)"> #quinones2.txt 2 O=C[1]C[2]:C:C:C:C:C(:@2)C[3]=C(OH)C(=O)NC[4]:C:C:C:C(@1):C(@3):@4<regId="quinone\_C(2)"> #naphthol\_C.txt 2  $C[1]:C(:C:C:C:@1)C(C)=O):C(:C:C:C:@1)Any[IS=OH,NH2] < regId="keto_naphthol_A(2)"> re$ #AAmisc5.txt 2 CH(C[1]:C:C:C:C:@1)(C[2]:C:C:C:C:@2)C(=S)NH<regId="thio\_amide\_C(2)"> #AA misc13.txt 2 N[1](C(=O)C[2]:CH:C(C(=O)OH):CH:CH:C(:@2)C(=O)@1)(C[3]:C:CH:C(O):C:CH:@3)<regId="phthalimide misc(2)"> #BBparanils 5.txt 2 C[1]:C(:C:C:C(:C:@1)NHS(=O)(=O))NHS(=O)=O<regId="sulfonamide\_D(2)"> #BBorthoanils 2.txt 2 CHNHC[1]:CH:CH:CH:CH:C(NHCH):@1<regId="anil NH alk C(2)"> #BB thiophenelactam.txt 2 S[1]C(=C(C[4]=C@1NHC(C(=CH@4)C(=O)OH)=O)NH2)C(=O)NH<regId="het\_65\_E(2)"> #CC\_naphth\_hydraz.txt 2 C[1]H:CH:CH:CH:C[9]:CH:C(:CH:CH:C:@1:@9)NHNHC=O<regId="hzide\_naphth(2)"> #CC ON 6.txt 2 CH2(C[4]:C(:CH:CH:C(:CH:@4)CH2NHC[TAC=4])OCH2)<regId="anisol\_B(2)"> #DD\_thiocarbam\_5.txt 2  $C[1]C[TAC=4](SC(NC=@1)=S)< regId="thio_carbam_ene(2)">$ #DD\_thioamid\_2.txt 2  $C(N(CH)CH):CNHC(=S)CH< regId="thio_amide_D(2)">$ #DD\_phenaz.txt 2 N[1]N=C(C[4]C(=C@1C)C=CC=@4)C<regId="het\_65\_Da(2)"> #EE\_5hets20.txt 2  $S[1]:CH:CH:C(OCH2):C(C(=O)NHNH):@1<regId="thiophene_D(2)">$ #EE\_sixNS.txt 2  $C[1]:CN=CC(=CNC)S@1<regId="het_thio_6_ene(2)">$ #FF\_cyano\_A1.txt 2 CH2CH(C#N)C(=O)C<regId="cyano\_keto\_A(2)"> #FF\_cyano\_11.txt 2  $C[1](=C(NH2)N(C[7]:C(:C:C:C:@7)C(=O)OH)N=C@1C=O)Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[7]:C(:C:C:C:@7)C(=O)OH)N=C@1C=O)Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[7]:C(:C:C:C:@7)C(=O)OH)N=C@1C=O)Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[7]:C(:C:C:C:@7)C(=O)OH)N=C@1C=O)Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[7]:C(:C:C:C:@7)C(=O)OH)N=C@1C=O)Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[7]:C(C:C:C:C:@7)C(=O)OH)N=C@1C=O)Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[1]:C(1))Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[1]:C(1))Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[1]:C(1))Any[IS=C\#N,C=S] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[1]:C(1))Any[IS=C[1]:C(1)] \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[1]:C(1))Any[IS=C[1]:C(1)" \\ < regId="anthranil_acid_C(2)"> C[1](=C(NH2)N(C[1]:C(1)") \\ < regId="anthranil_$ #FF\_Naphth\_2.txt 2 C[1]:C:C:C[4]:C[5]:C(:C:C:C:C:@1:@5)C=NNH@4<regId="naphth\_amino\_C(2)"> #FF\_Naphth\_3.txt 2

C[1]:C:C:C[4]:C[5]:C(:C:C:C:C:@1:@5)N=NN@4<regId="naphth\_amino\_D(2)"> #FF misc 4.txt 2  $C[1]N=C(SC=@1)NNS(=O)=O< regId="thiazole_amine_G(2)">$ #FF misc8.txt 2 C[1]:C:C:C[4]:C(:C:@1):N:C(:N:C:@4)NHC[13]=NC(C=CNH@13)(CH2)CH2<regId="het\_66\_B(2)"> #FF\_chromen\_2\_1.txt 2 C[1]:C:C(OCH):CH:CH:C(:@1)C[2]:CH:CH:C(OCH2):CH:C(:@2)C(=O)O@1<regId="coumarin A(2)"> #GG\_bicycle\_2.txt 2  $C[1]:N:C:C:C:C(:@1)C(NH2) = C(C(=O) \sim O[f])S@1 < regId = "anthranil_acid_D(2)" > C(I) = C(I$ #GG\_bicycle\_5.txt 2 C[1]:Hev:C:C:C:C(:@1):N:C(CH=C(OH)C):C(CH=C(OH)C):N:@1<regId="het\_66\_C(2)"> #GG thiophene bis.txt 2 C[1]=CSC(=C@1NH2)CH=CHC[13]=CC=CS@13<regId="thiophene\_amino\_E(2)"> #GG\_4cycle.txt 2 C[1]:C:C:C[4]:C(:C:@1):N:C[8]:C(:N:@4):C:C:C[13]:C:@8:C:C:C:C:@13<regId="het\_6666\_A(2)"> #GG NSO2N.txt 2 C:CNHS(=O)(=O)NHC:C<regId="sulfonamide\_E(2)"> #CC\_Ar2\_p\_anils\_1.txt 2 C[1]:C(:C:C:C:(:C:@1)N(C[TAC=3])C[TAC=3])NH2<regId="anil\_di\_alk\_K(2)"> #JJ AAmisc1.txt 2 #JJ enone D3.txt 2 C[2]:C:C:C:C:C(:@2)C(=O)CH=C[1]C(=O)NHC(=O)C(=CHC[3]:C:C:C:C:(@3)NH@1<regId="ene six het B(2)"> #JJ AA steroid.txt 2 O=C[1]CCC[2]C[3]C(=O)CC[4]CCCC(@4)C(@3)CCC(@2)=C@1<regId="steroid\_A(2)"> #JJ\_ON\_indol\_4.txt 2 C[1]:C[2]:C(:N:C:N:@2):C(OH):CH:C(:@1):C(C=O):C:N(C):@1<regId="het\_565\_A(2)"> #JJ\_thiiniums\_3.txt 2 C[TAC=4]N[+1](C[TAC=4]OH)=CSCH<regId="thio\_imine\_ium(2)"> #KK 110 fives 7.txt 2  $C[1](=O)C(=CHNHC[2]:C(C(=O)OH):C:C:C:C:@2)N=C(C[3]:C:C:C:C:@3)O@1<=regId="anthranil_acid_E(2)">C(E(E(B)C(E(B)E(E(B)E(E(B)C(E$ #KK\_111\_furzone\_2.txt 2 C[1](:O:C(Any[IS=H,CH2]):CH:CH:@1)C(Any[IS=H,CH2])=NNHC[2]:C:C:N:C:C:@2<regId="hzone\_furan\_B(2)"> #KK 111 thiophbtz.txt 2 #KK\_114\_maleimide\_1.txt 2 C:CCH2CH(C=O)N[1]C(=O)C[2]HCH2C=CCH2CH(@2)C(=O)@1<regId="ene\_misc\_B(2)"> #KK 107 SIM 2.txt 2  $C[1](C=O)(C:C)S[TAC=2]C=NNH@1<regId="het_thio_5_B(2)">$ #KK EE amino 5hets 4a.txt 2 NH2C[1]:S:C(NHC(=O)C[3]:C:C:C:C:(@3):C(C#N):C(C[2]:Hev:Hev:Hev:Hev:Hev:@2):@1<regId="thiophene amino F(2)"> #KK CC o anisidine 5A.txt 2 CH2OC[1]:CH:CH:C(CH2):CH:C(:@1)NHCH2C[2]:C(Any[IS=CH2,OCH2]):C:C:C:C:@2<regId="anil\_OC\_alk\_D(2)"> #KK DD t Bu 1C.txt 2 CH3C(CH3)(CH3)C[1]:CH:C(C(CH3)(CH3)CH3):C(OCHN):C:CH:@1<regId="tert butyl A(2)"> #KK DD thiophenezone 1C.txt 2 C[1](:C:C:O:C(CH):@1)C=NNHC(=S)NH<regId="thio\_urea\_J(2)"> #FF\_purine\_1.txt 2 NHC[3]=NC(=NC[7]=NN=C(N@3@7)SC)NHC<regId="het\_thio\_65\_B(2)"> #NN FF chromen 7 4.txt 2 C[1]:C(CH2CH=CH2):C:C:C:C(:@1)CH=C(C(=O)NHC:C)C(=O)O@1<regId="coumarin\_B(2)"> #M1\_DD\_NNNS.txt 2 C[1](=S)N[2]C:CN=NC(@2)=NNH@1<regId="thio\_urea\_K(2)"> #KK\_EE\_amino\_5hets\_5a.txt 2  $C:C:C:C:C:C:C[1]:S:C(NHC(=O)C):C(C(=O)OH):C:@1 < regId = "thiophene_amino_G(2)" > 1 < regId = 1$ #QQ\_BB\_orthoanils\_1.txt 2 NH2C[1]:C(NHCH(C)CHCH2):C:CH:CH:CH:@1<regId="anil\_NH\_alk\_D(2)"> #QQ\_DD\_NNSO\_1.txt 2 S=C[1]NHN=C(C[2]:CH:CH:C(OCH2):CH:CH:@2)O@1<regId="het\_thio\_5\_C(2)"> #QQ\_DD\_thi\_azene.txt 2 S=CC[1]:N[2]:C:C:C:C:C:(:@2):C:C:@1<regId="thio\_keto\_het(2)"> #KK\_115\_polyimine.txt 2 C[1]~C(~N~N~C(~CH2)~CH2)~N~S~C~@1<regId="het\_thio\_N\_5B(2)"> #o\_quinones.txt 2  $C[1](C:=CC:=CC@1=Het)=Het<regId="quinone_D(2)">$ #amino\_furan\_ene.txt 2 CH2N(CH2)C[1]:CH:CH:C(:O:@1)CH=CC#N<regId="anil\_di\_alk\_furan\_B(2)"> #sixes\_last2.txt 2

O=C[1]C:CCH2NC(@1)=CH<regId="ene\_six\_het\_C(2)"> #five\_pz\_sulf.txt 2 C:CN[1]:N:C[2]CH2S[TAC=2]CH2C(:@2)C(:@1)NHC(=O)CH=CH<regId="het\_55\_A(2)"> #bim SCH2.txt 2 N[1]:C[2]:C:C:C:C:(:@2):N(CH2):C(:@1)SCH2C(=O)NHN=CHCH=CH<regId="het\_thio\_65\_C(2)"> #hydroquin\_enone.txt 2 C[1](OH):C:C:C(OH):C(:C:@1)C(=[!R]CN)C=O<regId="hydroquin\_A(2)"> #anthran OH.txt 2  $C[1](OH):C:C:C(NHC(=O)C:C):C(:C:@1)C(=O)OH < regId="anthranil_acid_F(2)" > regId="acid_F(2)" > regId="ac$ #pyrrole\_2one.txt 2 N[1](CH2)C[2]=C(C:CC(=O)@2)C=C(CH2)@1<regId="pyrrole\_I(2)"> #KK EE amino 5hets 7a.txt 2 CHNHC[1]:S:C(CH):C(CH):C(C(=O)NHC:C):@1<regId="thiophene\_amino\_H(2)"> #isatins\_3.txt 2 C:CN[!r]=C[2]C(=Het)C[5]:C(:C:C:C:C:@5)N@2<regId="imine\_one\_fives\_C(2)"> #DD\_acylzone\_3.txt 2 C[1]:C(:C:C:C:@1)C(=O)NHN=C[2]C[3]:C(:C:C:C:@3)C[4]:C@2:C:C:C:@4<regId="keto\_phenone\_zone\_A(2)"> #dves7A.txt 2 C[1]:C:C:C:C:C:(:@1)N(CH2)CH=CHC=[!R]CHCH=CC=[R]NC[2]:C:C:C:C:C:@2<regId="dyes7A(2)"> #pyridiniums\_B.txt 2  $C[1]:C:C:C:C:Hev(:@1):C(Any):Hev:N[+1](\sim C:C):Hev:@1< regId="het_pyridiniums_B(2)"> regId="het$ #166\_het\_5\_t.txt 2 N[1](C[2]:C:C:C:C:@2)N=C(CH2)CH(SC)C(=O)@1<regId="het 5 D(2)"> #fives P 1.txt 1  $C[1]: C(:C:C(C(=0)OH): C:C:@1)NHC[12] = NC(=CHS@12)C[18]: C:C(C(H(CH)(CH)): C:C:@18 < regId = "thiazole_amine_H(1)" > regId$ #fives\_P2.txt 1 CH2NHC=NNHC[1]=NC(C:C)=CHS@1<regId="thiazole\_amine\_I(1)"> #fives\_P3.txt 1 C:CNHC(=O)C[1]N=NSC(NHC:C)=@1<regId="het\_thio\_N\_5C(1)"> #fives\_Q.txt 1  $O=S(=O)(C:C)NHC[7]=NC(=C(S@7))C:C< regId="sulfonamide_F(1)">$ #fives\_Q2.txt 1  $O=S(=O)(C:C)NHNHC[7]=NC(=C(S@7))C:C< regId="thiazole_amine_J(1)">$ #sixesB5.txt 1 S[1]C[2]:C(C(CH2)=C@1CH2):C(:N:C:N:@2)NN=CC[19]=CC=CO@19<regId="het 65 F(1)"> #enone H.txt 1 C(=O)CH=C(OH)C(OH)=CHC(=O)C<regId="keto\_keto\_beta\_E(1)"> #enone H2.txt 1 C[1]H:CH:CH:CH:C[2]CH2C(=O)C(=C(CH2)CH2)C(:@2):@1<regId="ene\_five\_one\_B(1)"> #meldrums2.txt 1 C:CNHN=C(CH2)CH2C(CH2)=NNHC:C<regId="keto\_keto\_beta\_zone(1)"> #thioimidate 2.txt 1 C[TAC=4]S[TAC=2]C(=N-Hev:Hev:Hev:Hev)NHN=C<regId="thio\_urea\_L(1)"> #fivesZ.txt 1 C[1](SC(=CHC:C)N(C:C)N=@1)C=O<regId="het thio urea ene(1)"> #multiCN.txt 1  $C[1](:C(:N:C[4]:C(:C:@1C\#N):C(:C(:C:C:@4)C\#N)NH2)NH2)C\#N < regId="cyano_amino_het_A(1)"> regId="cyano_het_A(1)"> regId="cyano_het_A(1)"> regId="cyano_het_A(1)"> regId="cyano_het_A(1)"> regId="cyano_het_A(1)"> regId="cyano_het_A(1)"> regId="cyano_het_A(1)"> regId="cyano_het_A(1)"> regId="cyano_het_A(1)">$ #chalc2.txt 1 Hev[1]:Hev:Hev:Hev:Hev:@1)CH=CHC(NHNHC[16]N(N=NN=@16)C)=O<regId="tetrazole\_hzide(1)">  $C[1]:C(:CH:CH:CH:@1):C(:C(:CH:CH:@1)C(=NC:C)CH2)OH < regId="imine_naphthol_A(1)"> regId="imine_naphth$ #misc1.txt 1 C[1](:C(:CH:C[5]:C(:CH:@1):C(:CH:C(:CH:@5)C[15]:CH:CH:C(:CH:@15)OCH2)OCH2)NHCH3)OCH2)OCH2regId="misc\_anisole\_A(1)"> #misc2.txt 1 C[1]:C:C:C[7]:C(:C:@1)NC[13]=C(S@7)SC(=C@13)CH2<regId="het\_thio\_665(1)"> #misc7.txt 1 #misc8.txt 1  $C[1](:C:C:C:C:@1)C[7] = CC(C[10]C(C(=C@7)OH) = COC=@10) = O)SCH2 < regId = "colchicine_B(1)" > COC=@10) = O(CCC) = O(C$ #misc12.txt 1 C[TAC=4]C[1]:CH:CH:C(C(=0)NHCH(CH2CH2SCH3)C(=O)OH):CH:CH:@1<regId="misc\_aminoacid\_A(1)"> #AAmisc3.txt 1  $N[1]: C(C[2]: C: C: C: C: C: @2): C(C[3]: C: C: C: C: @3): N(N=[!R]C): C(NH2): @1 < regId = "imidazole_amino_A(1)" > (NH2): @1 < regId = "imidazole_Amino_$ #AAmisc4.txt 1  $C(C[1]:C:C:C(OH):C:C:@1)(C[2]:C:C:C(OH):C:C:@2)OS(=O) = O < regId = "phenol_sulfite_A(1)" > O < regI$ #AAmisc7.txt 1 C[1]:C:C:C[7]:C(:C:@1):N:C(:C(:N:@7)CH2C(=O)C:C)CH2C(=O)C:C<regId="het\_66\_D(1)"> #AAmisc8.txt 1

 $C[1](:CH:C(OCH2):C(OCH2):CH:CH:@1)C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2)C:C:@2<regId="misc_anisole_B(1)">C(=O)CH2N(CH2)C[2]:C:C(CH2$ CH2C[1]=NN=NN(C[2]:CH:CH:C(OCH3):CH:CH:@2)@1<regId="tetrazole\_A(1)"> #AAmisc9 A.txt 1  $C[1](C)CH2C(=O)N[2]N=C(NH2)C(NH2)=C(@2)N=@1<regId="het_65_G(1)">$ #AA\_misc14.txt 1  $C(C:C)(C:C)(C:C)SC:CC(=O)OH < regId = "misc_trityl_A(1)" >$ #AA misc15.txt 1  $O=C(C[1]:CH:C(OCH3):N:C(OCH3):CH:@1)NHCH(CH2)CH2 < regId="misc_pyridine_OC(1)">1)CH:C(OCH3):CH:@1)NHCH(CH2)CH2 < regId="misc_pyridine_OC(1)">1)CH:C(OCH3):CH:@1)NHCH(CH2)CH2 < regId="misc_pyridine_OC(1)">1)CH:C(OCH3):CH$ #AA\_misc17.txt 1 N[1]CH(C:C)CH2C(=O)NHC(NH)=@1<regId="het\_6\_hydropyridone(1)"> #AA misc18.txt 1 C[1](C:C)CH2CH(C(=O)C)CH(C(=O)OH)CH2C(C:C)=@1<regId="misc\_stilbene(1)"> #AA\_misc19.txt 1  $CH(C[1]:CH:CH:C(Cl):CH:CH:@1)(C[2]:CH:CH:C(Cl):CH:CH:@2)OCH2CH2CH2C[3]N(CH3)CH=CHN=@3 < regId="misc_imidazole(1)"> regId="misc_$ #BBparanils 6.txt 1 N[1]:CH:C(NH2):CH:CH:C(NHC:C):@1<regId="anil\_NH\_no\_alk\_A(1)"> #BB o anils 5.txt 1 NH(C[1]:CH:CH:CH:CH:C(OH):@1)C[2]N=NC(=N)OC(NH2)=@2<regId="het\_6\_imidate\_B(1)"> #BB anils misc.txt 1 NH(C[1]:CH:CH:C(CH2):CH:CH:@1)CH2CH2C[2]:CH:CH:C(OCH2):CH:CH:@2<regId="anil\_alk\_B(1)"> #BB anils2.txt 1 C[1]:C:C:C[7]:C(:C:@1)C(C[12]:C@7:C:C:C(:C:@12)NH2)=CHC<regId="styrene anil A(1)"> #BB aminal1.txt 1 C[1]:C:C:C[4]:C(:C:@1)COC(N@4C(=0)CH2)(C(=0)OH)CH2<regId="misc\_aminal\_acid(1)"> #BB\_meta\_anil2.txt 1 N[1]:C(NH2):C(CH2):CH:C(CH2):C(NH2):@1<regId="anil\_no\_alk\_D(1)"> #CC\_pCN\_O3.txt 1 NH(C[1]:C:C:C:C:C:@1)C(C)(C)C[2]:CH:CH:C(OCH2):CH:CH:@2<regId="anil\_alk\_C(1)"> #CCpNH OC 1 1.txt 1 #CC\_CH2dioxy3.txt 1 C[1]:C(:C:C[4]:C(:C:@1)CC@4)OCO@1<regId="het\_465\_misc(1)"> #CC ON 1 3 1.txt 1 C[1](:CH:CH:C(OCH2):CH:C(C(=O)OH):@1)NHC:C<regId="anthranil acid G(1)"> #CC N 7.txt 1  $C[1]H:C(:CH:N:C[7]:C:@1:CH:CH:CH:CH:CH:@7)CH2N[20]C[21]:C(CH2CH2@20):CH:CH:CH:CH:CH:CH:@21 < regId="anil_di_alk_M(1)">10 < r$ #CC 7CO2H indol.txt 1 C[1]:C:C:C[4]:C(:C:@1C(=O)OH)NHC(=C@4C:C)C:C<regId="anthranil\_acid\_H(1)"> #DD thiourea.txt 1 C:CN(CH2)CH2CH2CH2NHC(=S)NHC[1]:C(Hal):CH:C(CH2):CH:CH:@1<regId="thio urea M(1)"> #DD thiobenz misc2.txt 1 N[1]:C[2]:C(:C:C[5]:C:@1N=C(S@5)N)SC(=N@2)N<regId="thiazole\_amine\_K(1)"> #DD NOS 4.txt 1 N=C[1]SC(=N)N=C@1<regId="het thio 5 imine A(1)">#DD thioamid.txt 1 C[1]:C:C:C:N:C(:@1)C(=S)NHC[2]:C:C:C:C:C(:@2)OCH2<regId="thio\_amide\_E(1)"> #DD\_phenothz\_6.txt 1 C[1]CH2SC[2]:CH:CH:CH:CH:C(:@2)C(OC:C)C(:@1):CH:CH:CH:CH:@1<regId="het\_thio\_676\_B(1)"> #DD sulfon pyr.txt 1 CH3C[5]:N:C(NHS(C[11]:CH:CH:CH:CH:@11)OCH2CH2CH2()=O)=O):CH:CH:@5<regId="sulfonamide\_G(1)"> #DD \_thionomorph.txt 1 C(=O)(N[3]CCSCC@3)C[9]:C(:CH:CH:CH:CH:@9)SCH2<regId="thio\_thiomorph\_Z(1)"> #DD\_enone\_3.txt 1  $C[1]:C:C[3]:C[4]:C(:C:@1):C:C:C:C:@4C(=CC@3=O)OCH2 < regId="naphth_ene_one_A(1)">$ #DD\_cf3.txt 1 C[1]: C: C: C: C: C: C: 0: C: C: 0: C: C: 0: C#DD\_polycyc.txt 1  $C[1]:C:C:C:C[5]:C:@1:C:C[8]:C(:N:@5):N:C[12]:C(:C:@8N):C:C:C:C:@12 < regId = "amino_acridine_A(1)" > regId = "amino_acridine$ #DD\_suber2.txt 1 C[1]:C:C:C:C[5]:C:@1C(C[8]C@5=NHev=CC@8)=O<regId="keto\_phenone\_B(1)"> #DD\_suber\_4.txt 1 C[1]:CH:CH:C(OCH2):CH:C(:@1)C(=NNHC[2]:CH:CH:C(C(=O)OH):CH:CH:@2)C[3]:CH:C(OCH2):CH:CH:C(:@3)@1<regId="hzone\_acid\_A(1)"> #DD\_sulfonAZ.txt 1 C[1](:CH:CH:C(:CH:CH:@1)NH2)S(=O)(=O)NHC[19]:CH:CH:CH:N:N:@19<regId="sulfonamide\_H(1)"> #EE\_pyrrole\_5.txt 1 C[1]H=CHN(CH2)C[7]:C@1:C[9]CH=CHN(C:@9:C(:C:@7OCH2)OCH2)CH2<regId="het\_565\_indole(1)"> #EE\_pyrrole\_6.txt 1

C[1](=C[2]C(=C(N@1C(O)=O)CH2)SCH2S@2)CH2<regId="pyrrole\_J(1)"> #EE pz thz.txt 1 S[1]C=CN=C@1C[6]CH=NNHC=@6NH2<regId="pyrazole\_amino\_B(1)"> #EE 5hets9.txt 1 C[1](NHC(C[2]:CH:CH:CH:CH:CH:@2)=C(CH2)CH=@1)C(=O)OH<regId="pyrrole\_K(1)"> #EE\_5hets30.txt 1 C[1](:O:C(C):CH:CH:@1)C(=O)NHC[1]:CH:CH:CH:CH:C(=O)OH):@1<regId="anthranil acid I(1)"> #EE thiocarbz.txt 1 Hev:CC(=S)NHNHC:Hev<regId="thio\_amide\_F(1)"> #FF\_cyano\_10.txt 1  $C[1](=O)C(C(C\#N)=CHN)C(N)C=C@1<regId="ene_one_C(1)">$ #FF cyano 14a.txt 1 C[1](=C[2]N(C(C=CN@2)=O)N=C@1C[11]NC=CC=@11)C#N<regId="het\_65\_H(1)"> #FF\_cyano\_15.txt 1  $O=C[1]C(C\#N)=NNCC@1< regId="cyano_imine_D(1)">$ #FF\_cyano\_16.txt 1 C[1](:C[2]:C:C:C:C:@2:N:N:C:@1)C(C:C)C#N<regId="cyano\_misc\_A(1)"> #FF misc8.txt 1 #FF misc12.txt 1 #FF misc13.txt 1 CH2C(OH)=C(C(=O)CH2)CHC#C<regId="keto keto beta F(1)"> #FF zene.txt 1 #FF\_naphth\_imine.txt 1  $C[1](:C:C:C[4]:C[5]:C(:C:C:C:C:@1:@5)C(C=C@4)=N)N < regId="naphth_ene_one_C(1)">$ #FF\_fluorenone.txt 1 C[1](:C:C[3]:C(:C[5]:C:C:C:C:@1:@5)C[11]:C(C@3=O):C:C:C:C:@11)OH<regId="keto\_phenone\_C(1)"> #FF chromen 6.txt 1 C[1](C=NC[4]:C(O@1):C:C:C(Cl):C:@4)=O<regId="coumarin\_C(1)"> #FF\_misc\_8\_1.txt 1 C[1]=CN(C(C[5]:C:C:C:C:C@1:@5)(C#N)C(=S)S)C=O<regId="thio\_est\_cyano\_A(1)"> #FF Bim urea.txt 1 C[1](=NC[3]:CH:CH:CH:CH:C:@3N@1C)NHC(NHC[20]:CH:C:C:C:CH:@20)=O<regId="het 65 imidazole(1)"> #FF\_Ncycle\_3.txt 1  $NH(C:C)C[5]:C(C(=O)OH):C:C:C(:N:@5)C:C < regId = "anthranil_acid_J(1)" >$ #GG bicycle 3.txt 1 C[1]:C:C:C:C:C(:@1)SC(=NN=C[2]C=CC=CC=C@2)N(CH2)@1<regId="colchicine\_het(1)"> #GG bicvcle 4.txt 1 C[1]:CH:C(OCH2):C(OCH2):CH:C(:@1)C(C)=C(C)SCH2@1<regId="ene misc D(1)"> #GG carbazole 2.txt 1 C[1]:CH:CH:CH:CH:C(:@1)C(CH2)=C(C:C)N(-[!R]C:C)@1<regId="indole\_3yl\_alk\_B(1)"> #GG anthril SO3 2.txt 1 NH2C[1]:C(OH):C:C(S(=O)(=O)OH):C:C:@1<regId="anil OH no alk A(1)"> #GG thiophene 2.txt 1 S[1]:C:C:C(C[2]:C:S:C(NH2):N:@2):CH:@1<regId="thiazole\_amine\_L(1)"> #GG\_thiophene\_4.txt 1 C[1]=C(NH2)NN=C(@1)C[2]=C(CH2)OCH=CH@2<regId="pyrazole\_amino\_A(1)"> #GG thiophene 6.txt 1 N[1]=NSC=C(@1)C[2]ON=C(C:C)N=@2<regId="het\_thio\_N\_5D(1)"> #GG\_indoline.txt 1 C[1](:C:C:C[4]:C(C[6]CC[8]:C(C@6N@4):C:C:C:C:@8):C:@1)C[TAC=4]<regId="anil\_alk\_indane(1)"> #GG\_tricycle3.txt 1 C[1]:CH:CH:CH:CH:C(:@1)CH=CHC[2]C(C#N)CH2CHN(@2)@1<regId="anil\_di\_alk\_N(1)"> #GG\_tricycle5.txt 1 C[1]:C:C[2]:C:C:C:C:C:(:@2):C:C(:@1)N(CH2)C(=O)C(C:CNHCH2)=N@1<regId="het\_666\_C(1)"> #enone\_F.txt 1  $C(OH):CC(=O)CH=C(C)C< regId="ene_one_D(1)">$ #JJ\_CC\_ON\_indol\_2.txt 1 C[1]:CH:CH:C(N(CH2)CH2):CH:C(:@1):C(S(=O)=O):CH:NH:@1<regId="anil\_di\_alk\_indol(1)"> #JJ\_CC\_ON\_indol\_3.txt 1  $C[1]: CH: CH: C(NH2): CH: C(:@1): CH: CH: N(CH2): @1 < regId = "anil_no_alk_indol_A(1)" > regId = "anil_no_alk_indol_A($ #JJ\_FF\_dhp\_2.txt 1  $S[1:TAC=2]C=C(C\#N)C(C)(C=O)C(Any[IS=C=O,C\#N])=C(NH2)@1 < regId="dhp_amino_CN_G(1)">1 < regId="$ #JJ\_FF\_dhp\_3.txt 1  $N[1]C = C(C = O)C(C[3] : C:C:C(N(CH2)CH2) : C:C:@3)C[2] = C(\sim N \sim C(\sim S) \sim N \sim C(\sim N) \sim @2)@1 < regId = "anil_di_alk_dhp(1)" > C(\sim N) \sim (N \sim C(\sim N) \sim N))$ #JJ\_FF\_anthrilNN\_1.txt 1

C[1]:C:C:C:C:C(:@1)C(=O)NHC[2]:C:C:C:C(:@2)C(=O)NHNHC[3]:N:C:C:S:@3<regId="anthranil\_amide\_A(1)"> #JJ DD anthran.txt 1 C[1]:C:C:C:C[5]:C:@1:C:C[8]:C(:C:@5C=NNHC[2]:C:C:C:C:C:@2):C:C:C:C:@8<regId="hzone\_anthran\_Z(1)"> #JJ enone D2.txt 1 C[1]:C:C:C:C:C(:@1)CHNC(=O)C(NHCH2)=CHC(=O)C[2]:C:C:C(OCH2):C:C:@2<regId="ene\_one\_amide\_A(1)"> #JJ\_EE\_5hets17.txt 1 S[1]:CH:CH:CH:C(:@1)C[2]NHN=C(C[4]:C:C:N:C:C:@4)C[3]:C(:C:C:C:C:@3)N=@2<regId="het 76 A(1)"> #JJ EE 5hets18.txt 1 #JJ\_FF\_chromen\_5.txt 1  $C[2]: C: C: C: C: C: C: (@2)N(CH)CHCHNHC(=O)C[1]C(CH2) = CHC(=O)OC(CH2) = @1 < regId = "anil_di_alk_coum(1)" > CHC(CH2) = (@1 < regId$ #JJ FF misc 10.txt 1 C[1]:C:C:C[2]:C:C:C:C:C:C(:@2):C(@1)CHC[TAC=4]NC(@1)=CHC(=0)N(CH2)CH2<regId="ene\_one\_amide\_B(1)"> #JJ\_thiiniums\_2.txt 1 #KK 110 fives 3.txt 1 C[1](=O)C(=C(CH2)NHCH2CH2CH2)N=C(C[2]:C:C:C:C:C:C:@2)O@1<regId="het\_5\_ene(1)"> #KK 110 imide 1 C[2]:C:C:C:C:C:(:@2)N[1]C(=O)C(SC[3]:C:C:C:C:C:@3)=CHC(=O)@1<regId="thio\_imide\_A(1)"> #KK\_116\_thidhp\_1.txt 1  $N[1]HN=C(NH)SC(C:C)=C(C:C)@1<regId="dhp_amidine_A(1)">$ #KK thiourea 2.txt 1 C[1]H:C(NHC(=S)NHCHC[3]:O:C(CH):CH:CH:@3):CH:C[2]:C(OCH2O@2):CH:@1<regId="thio urea O(1)"> #KK thiourea 8.txt 1 C[1]H:C(NHC(=S)NHC[2]:C:C:C:C:C:@2):CH:C(N(CH2)CH2):CH:CH:@1<regId="anil\_di\_alk\_O(1)"> #KK\_107\_indole.txt 1 O=C-[!R]N[1]:C:C:C[2]:C(NHC(=S)NH@2):@1<regId="thio\_urea\_P(1)"> #KK\_107\_pyrazol\_1.txt 1 C(F)(F)C(=O)NHC[1]:CH:N(CH2CH2OCH2C:C):N:CH:@1<regId="het\_pyraz\_misc(1)"> #KK 108 diazox 4.txt 1 N[2]=NC[1]:N:Het:N:C(:@1)N=NC:C@2<regId="diazox\_C(1)"> #KK\_108\_diazox\_5.txt C[1]H(OH)C[2]:N:Het:N:C(:@2)CH(OH)C=C@1<regId="diazox\_D(1)"> #KK AA misc10.txt 1 #KK\_109\_enam.txt  $C[1]:CC(=O)C=C(@1)N=CHN(C[TAC=4])C[TAC=4] < regId="imine_ene_one_B(1)">$ #KK 109 coumarin 2.txt 1 C[1]:C[2]C(C[3]:C:C:C:C:@3)=CHC(=O)OC(:@2):C:C(OCH2C:O:C):C:@1<regId="coumarin\_D(1)"> #KK 109 furmorph 1.txt 1 C[3]:C(CH2):C(CH2):O:C(:@3)CH2NCH2CH(OCH2)CH2OC[1]:C:C:C[2]:C(OCH2O@2):C:@1<regId="misc furan A(1)"> #KK 110 rhod 2.txt 1 N[1](C[2]:C:C:C:C:@2)C(=O)SCH(NHC[3]:C[4]:C:C:C:C:(:@4):C:C:C:@3)C(=O)@1<regId="rhod\_sat\_E(1)"> #KK 110 rhod 5.txt 1 N[1](C(=0)C[2]:C:C:C:C:@2)C(=NC[3]:C:C:C:C:@3)SCH2C(=0)@1<regId="rhod sat imine A(1)"> #KK 110 rhod 7.txt 1 N[1](C[2]:C:C:C:C:@2)C(=O)SCH2C(=S)@1<regId="rhod\_sat\_F(1)"> #KK\_110\_rhod\_13.txt 1 N[1](CH2)C(=S)N(C:C)C(=NC:C)C(=NC:C)@1<regId="het\_thio\_5\_imine\_B(1)"> #KK 110 rhod 14.txt 1 S[1]C(=NNH)SC(=NC:C)C(=NC:C)@1<regId="het\_thio\_5\_imine\_C(1)"> #KK\_110\_fives\_5.txt 1 C[1](=O)C(=CHC[2]:C:C(Hal):C:C:C(:@2)OCH2)N=C(SCH2)S@1<regId="ene\_five\_het\_N(1)"> #KK\_112\_thiocarbam.txt 1 CH2SC(=S)NHCH2C:C<regId="thio\_carbam\_A(1)"> #KK\_113\_dianilide\_1.txt 1  $C[1](NHC(=O)CH2CH2C:C):CH:CH:C(CH2):C(NHC(=O)CH2CH2C:C):CH: @ 1 < regId = "misc_anilide_A(1)" > 1 < regId$ #KK\_113\_lutidine\_1.txt 1  $C[1](NHC(=O)NHCH2CH2CH2):C(CH2):CH:C(Br):CH:C(CH2):@1 < regId="misc_anilide_B(1)">1 < regId="m$ #KK\_113\_mannich.txt 1 C[1]:C(OCH2N(C:COCH2)CH2@1):CH:C:C:CH:@1<regId="mannich\_B(1)"> #KK\_113\_mannich\_1.txt 1 C[1]:C(OCH2N(CH2)CH2@1):CH:C(O):C(O):CH:@1<regId="mannich\_catechol\_A(1)"> #KK\_113\_anilpip.txt 1 NH(C[1]:CH:CH:C(CH(CH2)(CH2)):CH:CH:@1)CH2CH2N(CH2)(CH2)<regId="anil\_alk\_D(1)"> #KK\_114\_acid.txt 1 N[1]:C:C:C(C(~O)~O):C:C(:@1):C:C(C:C):C(C(=O)C:C):@1<regId="het\_65\_I(1)"> #KK\_114\_alkylarylurea.txt 1

#KK 114 imipram 1.txt 1 #KK 114 arylstilbene.txt 1 C(=CHCH2N[1]:CH:N:CH:CH:@1)(C:C)C:C<regId="styrene\_imidazole\_A(1)"> #KK\_115\_Arthiaz\_2.txt 1 C[1](NHC:CCH2):N:C(C[2]:C:C:N:C:C:@2):CH:S:@1<regId="thiazole\_amine\_M(1)"> #KK\_115\_Arthiaz\_3.txt 1 #KK\_115\_pyrrolester.txt 1 N[1]H:C(C(=O)OCH2):C(CH2):C(CH2CH2):C(CH2CH2):@1<regId="pyrrole\_L(1)"> #KK 115 thiobim.txt 1 C[1](SCH2C(=O)NHC:C):N:C[2]:CH:C(Hal):C:CH:C(:@2):NH:@1<regId="het\_thio\_65\_D(1)"> #KK\_116\_thianisole\_1.txt 1 C[1](OCH2):C(OCH2):CH:C[2]C=CCHSC(:@2):CH:@1<regId="ene\_misc\_E(1)"> #KK\_116\_thipyr\_2.txt 1 N[1]HC(=S)CH(C#N)CH(C:C)CH=C(C:C)@1<regId="thio\_cyano\_A(1)"> #KK 116 thipvr 3.txt 1  $N[1]: C(S[TAC=2]C[3]: C(NH2): C:C:C:C:@3): C(C\#N): C(C[2]:C:C:C:C:@2): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C:C:C:C:C:C): @2): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C:C:C:C:C): @2): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C:C:C): @2): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C:C): @2): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C:C): @2): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C):C): (C(C\#N): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C):C): (C(C\#N): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C):C): (C(C\#N): C(C\#N): C(NH2): @1 < regId = "cyano_amino_het_B(1)" > (C(C[2]:C):C): (C(C\#N): C(C\#N): C(C\#N):$ #KK\_116\_pyr\_1.txt 1  $N[1](C[2]:C:C:C(OCH2):C:C:@2)C(=O)C(C\#N) = CC(N[3]:C:N:C:C:@3) = N@1 < regId = "cyano_pyridone_G(1)" > 1 < regId = 1 < regId = "cyano_pyridone_G(1)" > 1 < regId = "cyano_pyridone_G(1)" > 1 < regId = "cyano_pyridone_G(1)" > 1 < regId = "cyano_py$ #KK\_116\_benzofuran\_1.txt 1 O[1]:C(C(~O)~O):C:C[2]:CH:C(OCH2):C(OCH2):CH:C(:@2):@1<regId="het 65 J(1)"> #KK 116 divneone.txt 1  $C\#CC(=O)C\#C< regId="ene_one_yne_A(1)">$ #KK\_116\_aminonaphthol.txt 1 C[1](NH2):C(OH):C(C=O):CH:C[2]:CH:CH:CH:CH:C(:@2):@1<regId="anil\_OH\_no\_alk\_B(1)"> #KK\_111\_furzid\_3.txt 1 C[1](:O:C(Any[IS=H,CH2]):CH:CH:@1)C(=O)NHN=C(Any[IS=H,CH2])C[2]:C:C(AnyAnyAnyC[3]:O:C:C::@3):C:C:C:@3:C:C:C:@2<regId="hzone\_acyl\_misc\_A(1)"> #KK EE amino 5hets 9a.txt 1  $S(=O)(=O)NHC[1]:S:C(CH):C(CH):C(C(=O)NH):@1<regId="thiophene_F(1)">$ #KK\_CC\_o\_anisidine\_5B.txt 1 CH2OC[1]:CH:CH:CH:CH:C(:@1)NHCH2CH(OH)CH2<regId="anil\_OC\_alk\_E(1)"> #KK CC o anisidine 5E.txt 1 CH2OC[1]:CH:CH:CH:CH:C(:@1)NHCH(C=O)S<regId="anil OC alk F(1)"> #KK\_DD\_t\_Bu\_1A.txt 1  $N[1]=NNN=C[2]C=CC=C(@2)@1<regId="het_65_K(1)">$ #KK DD thiophenezone 1B.txt 1 C[1]:CH:S:C(C=O):C(:@1)C(=O)NN=C(@1)NH2<regId="het\_65\_L(1)"> #NN FF chromen 7 1.txt 1 C[1]:C(Br):C:C[2]:C(:C:C:O:@2):C(:@1)CH=CC(=O)O@1<regId="coumarin E(1)"> #NN FF chromen 7 2.txt 1 C[1]:C:C:C:C(:@1)CH=C(C(=O)NHC[2]:C(Br):C:O:N:@2)C(=O)O@1<regId="coumarin\_F(1)"> #NN\_FF\_chromen\_7\_3.txt 1 C[1]:C(Hal):C:C(Hal):C:C(:@1)CH=C(C(=O)NH2)C(=NH)O@1<regId="coumarin G(1)"> #NN FF chromen 7 5.txt 1 C[1]:C:C:C:C:(:@1)CH=C(C(=O)NHC[2]:S:C:C(C:S:CH):N:@2)C(=O)O@1<regId="coumarin\_H(1)"> #NN\_sixesB8\_4.txt 1 CH2S[TAC=2]C[1]:N:C[2]OC=NC:CC(:@2):N:N:@1<regId="het\_thio\_67\_A(1)"> #NN CC ON 8 1.txt 1 S(=O)(=O)(C[1]:C:N(CH2):C:N:@1)NHC[2]:C:N:N(CH2C:COCH2):C:@2<regId="sulfonamide\_I(1)"> #NN\_CC\_ON\_8\_2.txt 1 C[1]:C(OCH2O@1):C(CH2N[2]CH2CH2C:C@2):CH:CH:CH:@1<regId="het\_65\_mannich(1)"> #NN\_CC\_ON\_8\_3.txt 1 CH2OC:CCH2NHC[1]:CH:C[2]:N:CH:N(CH):C(:@2):CH:CH:@1<regId="anil\_alk\_A(1)"> #NN\_JJ\_thiziniums\_2.txt 1 #NN\_CC\_Me2\_p\_anils\_5.txt 1  $CH2N(CH2)C[1]:C:C[2]:N:C(SCH2):S:C(:@2):C:C:@1 < regId = "anil_di_alk_P(1)" > regId = "anil_di_alk_P($ #M15\_naphth\_F.txt 1 C[1]:C(:CH:CH:CH:CH:@1):C(:CH:CH:CH:@1)C(CH2)=NNHC(=S)NHC:C:C<regId="thio\_urea\_Q(1)"> #M1\_DD\_sixNS.txt 1 C[1](:N:Hev:N:C(C[2]:O:CH:CH:CH:@2):N:@1)SC[TAC=4]<regId="thio\_pyridine\_A(1)"> #KK\_108\_melamime\_1.txt 1 N[1]:C(N(CH2)CH2):N:C(N(CH2)CH2):N:C(N(CH)C=O):@1<regId="melamine\_B(1)"> #P\_isothidiaz\_1.txt 1 C[1](C[2]:C:C:C:C:C:@2):N:C(N(CH2)CH2CH2NC(=O)C[3]:C(C(=O)OH):C:C:C:C:@3):S:N:@1<regId="misc\_phthal\_thio\_N(1)"> #P\_pyracylzone.txt 1

N[1]:CH:CH:CH:CH:C(:@1)C(=O)NHN=CHC[2]:C(OCH2C(=O)OH):C:C:C:C:@2<regId="hzone\_acyl\_misc\_B(1)"> #KK DD t Bu 1B.txt 1 #OO BB orthoanils.txt 1 NH2C[1]:C(NH2):CH:CH:C[2]:N:O:N:C(:@1):@2<regId="diazox\_E(1)"> #QQ\_BB\_orthoanils\_2.txt 1 NH2C[1]:C(NHS(=O)=O):CH:C(NHCH2):C(Hal):CH:@1<regId="anil\_NH\_no\_alk\_B(1)"> #QQ\_BB\_orthoanils\_3.txt 1 NH2C[1]:C(N=C[2]CH=C~C~C=C@2):CH:CH:CH:CH:@1<regId="anil\_no\_alk\_A(1)"> #QQ\_BB\_orthoanils\_3.txt 1 NH2C[1]:C(N[2]:C:C:C:C:@2):CH:C(CH2):C(CH2):CH:@1<regId="anil\_no\_alk\_B(1)"> #OO DD thi eneaz.txt 1 S=CC(CH2)=C(CH2)N(CH2)CH2<regId="thio\_ene\_amine\_A(1)"> #QQ\_EE\_furane.txt 1  $C[1]:COC[2]CH2C(=O)OC(@2)@1<regId="het_55_B(1)">$ #thioimidate 4.txt 1 OC(=O)CH2S[TAC=2]C(=NC#N)NHC[1]:C:C:C:C:@1<regId="cyanamide\_A(1)"> #thioacetene.txt 1  $O=CC[1]=C(SC(=CHC)S@1)C=O<regId="ene_one_one_A(1)">$ #sixes\_last1.txt 1  $O=C[1]NNC(Hev:Hev)=NC(@1)=CH<regId="ene_six_het_D(1)">$ #acceptor.txt 1 O=CCH=C(C#N)C<regId="ene\_cyano\_E(1)"> #fur acid CN.txt 1 #acid\_ar\_zone.txt 1  $C[1]:C:C:C:C:C:(@1)N(C[2]:C:C:C:C:@2)N=CHC[3]:Hev:C(:CH:CH:@3)C[4]:C:C(C(=O)OH):C:C:C:@4<regId="hzone_furan_C(1)">-1000-furan_C(1)=-1000-fur$ NH2C[1]:CH:CH:C(:CH:CH:@1)C[2]:CH:C(C=O):C(CH2):O:@2<regId="anil\_no\_alk\_C(1)"> #acid zone fur.txt 1 HOC(=0)C[1]:C:C:C(:C:C:@1)NN=CHC[2]:Hev:C(:CH:CH:@2)C[3]:C:C:C:C:@3<regId="hzone\_acid\_D(1)"> #acid\_furzonide.txt 1 HOC(=O)C[1]:C:C(:C:C:@1)C:Hev:CC=NNHC(=O)CH2O<regId="hzone\_furan\_E(1)"> #DD OHPyr 2.txt 1 HOC[2]:N:C(NH2):Hev:C:C(CH2C(=O)O):@2<regId="het\_6\_pyridone\_NH2(1)"> #fives\_B2het.txt 1  $C[1](=Het)C(N=CS@1)=O< regId="imine_one_fives_D(1)">$ #pyrrole Aroxime.txt 1 N[1](C[2]:C:C:C:C:@2)CH=CHCH=C(C=NOH)@1<regId="pyrrole\_M(1)"> #pyrrole AlkAlkAlk.txt 1  $N[1](CHC[2]:CH:CH:CH:CH:@2)C(CH) = CHCH = C(CH)@1 < regId = "pyrrole_N(1)" > regId = "pyrrole_$ #pyrrole\_pentsub.txt 1  $N[1](CH2)C(CH2) = C(C(=O)C)C(C) = C(C:C) @ 1 < regId = "pyrrole_O(1)" > (CH2)C(CH2) = C(C(=O)C)C(C) = C(C:C) @ 1 < regId = "pyrrole_O(1)" > (CH2)C(CH2) = C(C(=O)C)C(C) = C(C:C) @ 1 < regId = "pyrrole_O(1)" > (CH2)C(CH2) = C(C:C) @ 1 < regId = "pyrrole_O(1)" > (CH2)C(CH2) = C(C:C) @ 1 < regId = "pyrrole_O(1)" > (CH2)C(CH2) = C(C:C) @ 1 < regId = "pyrrole_O(1)" > (CH2)C(CH2) = C(CH2)C(CH2) = C(CH2)C(CH2)C(CH2) = C(CH2)C(CH2)C(CH2) = C(CH2)C(CH$ #pyrrole acrCN.txt 1 N[1](C)CH=CHCH=C(CH=C(C#N)C[2]:N:C:C:S:@2)@1<regId="ene cyano G(1)"> #pyrrole\_Arsulfon A.txt 1 N[1](C[2]:C(NHS(=O)(=O)C[3]:S:C:C:C:@3):C:C:C:@2)CH=CHCH=CH@1<regId="sulfonamide\_J(1)"> #pyrrole\_benzamide.txt 1 N[1](C[2]:CH:CH:CH:CH:C(C(=O)NHCH(CH2)CH2OC:C):@2)CH=CHCH=CH@1<regId="misc\_pyrrole\_benz(1)"> #thiocarbz 3.txt 1 C[1](:C:C:C:C:C:@1)NHC(=S)NNHCH=CHC=O<regId="thio\_urea\_R(1)"> #meldrum\_1a.txt 1  $C[1](C(=O)CH2CCH2C(=O)@1)=C(NH)C=O< regId="ene_one_one_B(1)">$ #dhp\_amino\_CNSS.txt 1  $NH2C[1] = C(C\#N)CH(C:C)SC[TAC=4]S@1 < regId = "dhp_amino_CN_H(1)" > TAC(C)C(TAC=4)S(C)C(TAC=4)S(C)C(TAC=4)C($ #anis\_122\_A.txt 1 CH2OC[1]:CH:CH:CH:CH:C(:@1)NHC[2]:C[3]:C(OCH2):C:C(OCH2):C(:@3):N:C:C:@2<regId="het\_66\_anisole(1)"> #anis\_112\_B.txt 1  $CH2OC[1]: CH: CH: C(OCH2): CH: C(:@1)NHC[2]: S: C: C(C[3]: C: C(OCH2): C: C: @3): N: @2 < regId = "thiazole_amine_N(1)" > CH2OC[1]: CH: C(:@1)NHC[2]: S: C: C(C[3]: C: C(OCH2): C: C: @3): N: @2 < regId = "thiazole_amine_N(1)" > CH2OC[1]: CH2OC[1$ #pyridiniums\_C.txt 1  $C[1] \sim N(C:C) \sim C \sim C \sim C(-@1) \sim C[2] \sim N \sim C \sim C \sim N[+1] (-@2) \sim N \sim @1 < regId = "het_pyridiniums_C(1)" > C[1] \sim N(C:C) \sim C \sim C \sim C(-@1) \sim C[2] \sim N \sim C \sim C \sim N[+1] (-@2) \sim N \sim (-C) \sim C(-@1) \sim C[2] \sim N \sim C \sim C \sim N[+1] (-@2) \sim N \sim (-C) \sim C(-C) \sim C(-C)$ #166\_het\_5\_p.txt 1  $N[1](C[2]:C:C:C:C[3]:C:C:C:C:(@3):@2)N=C(CH2)CH2C(=0)@1 < regId="het_5_E(1)" > regId="het_5$ 

**Supplementary Table S9**. Enrichment calculations for all problematic substructures with 4 or more member compounds. These are listed in order of population size. This value can be used to match up with the structure as listed in Figure S1.

Search WEHI DB for structures in Freq Hit 5 norethan150.hits

								Total	Enrichment <sup>a</sup>
Filename	6	5	4	3	2	1	0		
ene_six_het_A	10	20	21	30	69	10 5	228	483	66%
hzone_phenol_A	5	4	7	17	20 8	82	156	479	154%
anil_di_alk_A	31	66	27	35	28	59	232	478	81%
indol_3yl_alk	6	15	23	18	51	80	268	461	42%
quinone_A	40	57	48	41	42	56	86	370	265%
azo_A	29	30	33	43	24	55	110	324	145%
imine_one_A	17	20	18	15	38	49	164	321	60%
mannich_A	2	4	13	15	59	57	146	296	64%
anil_di_alk_B	6	18	7	18	22	43	137	251	52%
anil_di_alk_C	15	23	13	10	26	29	130	246	67%
ene_rhod_A	16	41	21	26	32	39	60	235	227%
hzone_phenol_B	2	2	9	6	38	54	104	215	55%
ene_five_het_A	6	14	24	14	39	40	64	201	152%
anil_di_alk_D	12	10	18	11	24	28	95	198	79%
imine_one_isatin	1	6	12	8	35	46	81	189	67%
anil_di_alk_E	8	12	17	10	15	27	97	186	64%

Filter WEHI DB using

Freq\_Hit\_5\_morethan150.hits

Then Search Remaining Compounds using

Freq Hit 5 lessthan150.hits

thiaz ene A	4	10	9	10	12	27	56	128	80%
pyrrole A	1	16	13	14	11	21	42	118	131%
catechol A	4	7	10	4	10	21	36	92	97%
ene five het B	0	4	4	2	14	22	44	90	55%
imine one fives	7	11	9	5	9	18	30	89	137%
ene_five_het_C	3	9	7	7	7	13	39	85	85%
hzone pipzn	1	7	8	13	8	9	33	79	112%
keto_keto_beta_A	3	2	0	3	10	15	35	68	51%
hzone_pyrrol	4	4	1	2	9	10	34	64	59%
ene_one_ene_A	1	2	3	5	6	16	24	57	71%
cyano_ene_amine_A	0	8	6	7	5	19	11	56	236%
ene_five_one_A	1	2	1	1	11	11	28	55	57%
cyano_pyridone_A	1	3	3	4	4	16	23	54	65%
anil_alk_ene	1	6	6	3	7	11	17	51	135%
amino_acridine_A	3	8	4	4	4	7	16	46	144%
ene_five_het_D	4	7	8	9	13	5	0	46	na
thiophene_amino_Aa	2	2	5	4	3	11	18	45	89%
ene_five_het_E	1	8	3	4	4	11	13	44	154%
sulfonamide_A	0	3	4	5	10	2	19	43	116%

thio_ketone sulfonamide_B anil_no_alk thiophene amino Ab	4 5 2 0	4 1 1 2	6 2 1 2	5 1 3 1	3 3 5 5	4 14 8 7	17 15 20 23	43 41 40 40	135% 80% 60% 43%
het_pyridiniums_A anthranil_one_A	7 0 1	3 2 1	5 2 1	3 2 1	2 4 5	5 9 7	14 19 21	39 38 37	143% 53% 43%
cyano_imine_A diazox_sulfon_A	1	4	2	2	4	6	17	36	76%
hzone_anil_di_alk rhod_sat_A	1 0	4 6	5 6	0 6	5 6	7 7	13 2	35 33	115% 1200%
hzone_enamin pyrrole_B	0 4	0 5	3 9	2 0	9 0	1 2	15 3	30 29	93% 600%
thiophene_hydroxy cyano_pyridone_B	$0 \\ 0$	4 1	2 2	1 2	3 7	7 4	11 11	28 27	91% 109%
imine_one_sixes dyes5A	0 8	1 3	1 4	0	5	7 2	13 9	27 27	54% 178%
naphth_amino_A naphth_amino_B	0	1 9	2 3	2 5	2 5	3	15 1	25 25	47% 2300%
ene_one_ester thio_dibenzo cyano cyano A	0 2 0	1 2 1	0 5 6	1 3 3	4 1 4	11 2 5	7 8 4	24 23 23	86% 163% 350%
hzone_acyl_naphthol het_65_A	1 0	1 0	0 2	3 2	5 2	7 7	5 8	22 21	200% 75%
imidazole_A ene_cyano_A	2 3	2 2	2 1 2	2 0	2 3	2 4	7 6	19 19	143% 150%
anthranil_acid_A dyes3A dhp_bis_amino_CN	1 5 0	2 0 1	2 1 5	4 2 7	2 3 1	5 1 4	3 7 1	19 19 19	375% 157% 1400%
het_6_tetrazine ene_one_hal	2	2 0	3 0	2 2	0 2	4 4	5	18 17	180% 63%
cyano_imine_B thiaz_ene_B	0 1	2 2	3 1	4 0	2 1	2 1	4 11	17 17	275% 36%
ene_rhod_B thio_carbonate_A	0	1 9	5 1	1 3	4 0	4 2	4 0	16 15	275% na
anil_di_alk_furan_A ene_five_het_F	0 1	0 1	2 0	1 1	6 2	4 6	2 4	15 15	450% 125%
Filter WEHI DB using Freq_Hit_5_lessthan150.h Then Search Remaining C	ompo	unds ı	ısing						
Freq_Hit_5_lessthan15.hiv anil_di_alk_F	2	1	3	2	0	1	5	14	160%
hzone_anil het_5_pyrazole_OH het_thio_666_A	0 1 1	1 0 3	1 1 3	0 3 0	1 4 0	1 3 3	10 2 3	14 14 13	30% 450% 233%
styrene_A ene_rhod_C dhp_amino_CN_A	0 0 0	0 3 0	2 0 2	1 0 2	4 2 6	1 2 0	5 6 3	13 13 13	140% 83% 333%
cyano_imine_C thio_urea_A	$\begin{array}{c} 0 \\ 0 \end{array}$	3 4	1 3	2	2 0	2 3	2 1	12 12	400% 800%

thiophene_amino_B	2	2	1	0	1	3	3	12	200%
keto keto beta B	$\overset{2}{0}$	0	2	1	3	0	6	12	100%
keto_keto_beta_b keto phenone A	0	2	0	0	1	1	7	11	43%
cyano_pyridone_C	2	2	0	3	3	1	0	11	
thiaz ene C	1	1	1	0	4	1	3	11	na 233%
<u> </u>	2	3	2	1	0	0	3	11	267%
hzone_thiophene_A ene_quin_methide	$\stackrel{\scriptstyle 2}{0}$	2	2	1	3	2	0	10	
het thio 676 A	0	3	$\overset{2}{0}$	1	0	$\stackrel{\scriptstyle 2}{0}$	6	10	na 67%
	0	0	2	1	1	1	5	10	80%
ene_five_het_G acyl_het_A	0	2	2	1	0	2	2	9	250%
	2	1	$\overset{2}{0}$	0	0	2	4	9	75%
anil_di_alk_G	4					1			
dhp_keto_A		0	2	0	1		1	9	700%
thio_urea_B	0	0	2 2	2	2	0	3	9	200%
anil_alk_bim	1	1			1	0	3	9	200%
imine_imine_A	0	3	0	0	0	1	5	9	60%
thio_urea_C	0	1	1	1	3	2	1	9	600%
imine_one_fives_B	3	0	1	1	0	1	3	9	167%
dhp_amino_CN_B	0	2	2	1	1	1	2	9	300%
anil_OC_no_alk_A	0	1	0	1	2	1	3	8	133%
het_thio_66_one	1	0	1	1	2	2	1	8	500%
styrene_B	1	0	3	3	1	0	0	8	na 1220/
het_thio_5_A	1	0	1	2	0	1	3	8	133%
anil_di_alk_ene_A	0	2	1	0	2	1	2	8	250%
ene_rhod_D	0	1	1	1	0	1	4	8	75%
ene_rhod_E	0	3	1	1	1	0	2	8	300%
anil_OH_alk_A	0	4	3	0	1	0	0	8	na
pyrrole_C	0	1	2	2	1	1	1	8	600%
thio_urea_D	1	1	3	0	1	1	1	8	600%
thiaz_ene_D	1	1	1	1	0	1	3	8	133%
ene_rhod_F	0	2	1	1	1	0	3	8	167%
thiaz_ene_E	0	1	3	1	0	1	2	8	250%
het_65_B	0	1	2	0	0	3	1	7	300%
keto_keto_beta_C	0	1	1	1	1	3	0	7	na
het_66_A	0	2	1	1	0	1	2	7	200%
thio_urea_E	0	0	1	2	1	2	1	7	400%
thiophene_amino_C	0	1	1	0	1	0	4	7	75%
hzone_phenone	0	0	3	2	1	0	1	7	600
ene_rhod_G	0	2	1	2	0	2	0	7	na
ene_cyano_B	0	1	1	2	2	0	1	7	600%
dhp_amino_CN_C	1	0	0	2	1	2	1	7	400%
het_5_A	0	1	0	1	1	1	3	7	100%
ene_five_het_H	0	1	0	0	2	3	0	6	na
thio_amide_A	0	2	2	0	0	1	1	6	400%
ene_cyano_C	1	0	1	1	1	1	1	6	400%
hzone_furan_A	0	2	1	0	0	1	2	6	150%
anil_di_alk_H	1	1	0	2	1	0	1	6	500%
het_65_C	1	1	0	0	0	0	4	6	50%
thio_urea_F	0	1	0	1	4	0	0	6	na
ene_five_het_I	0	1	0	0	1	3	1	6	200%
keto_keto_gamma	1	1	1	0	1	1	0	5	na
quinone_B	0	1	2	2	0	0	0	5	na

het_6_pyridone_OH	1	0	0	0	3	0	1	5	400%
hzone_naphth_A	0	1	1	0	0	2	1	5	200%
thio_ester_A	0	1	3	0	0	0	1	5	400%
ene_misc_A	0	1	0	0	1	0	3	5	67%
cyano_pyridone_D	1	0	0	1	1	0	2	5	150%
het_65_Db	0	1	0	0	3	1	0	5	na
het_666_A	1	0	3	0	0	0	1	5	400%
diazox_sulfon_B	0	2	0	0	1	0	2	5	150%
anil_NH_alk_A	0	1	0	0	0	2	2	5	50%
sulfonamide_C	0	0	2	0	0	2	1	5	200%
het_thio_N_55	0	2	2	1	0	0	0	5	na
keto_keto_beta_D	0	2	2	1	0	0	0	5	na
ene_rhod_H	0	2	0	1	1	1	0	5	na
imine_ene_A	2	0	1	2	0	0	0	5	na
het_thio_656a	0	0	1	0	1	2	1	5	200%
pyrrole_D	1	0	0	0	1	3	0	5	na
pyrrole_E	0	2	0	0	0	0	3	5	67%
thio_urea_G	2	0	0	0	0	0	3	5	67%
anisol_A	0	1	1	0	0	0	3	5	67%
pyrrole_F	0	1	1	0	0	1	2	5	100%
dhp_amino_CN_D	0	1	1	0	0	1	2	5	100%
thiazole_amine_A	0	1	0	1	0	2	0	4	na
het_6_imidate_A	1	0	0	0	0	1	2	4	50%
anil_OC_no_alk_B	0	1	1	0	0	0	2	4	100%
styrene_C	0	1	0	0	0	1	2	4	50%
azulene	2	0	1	0	0	0	1	4	300%
furan_acid_A	0	0	1	0	1	0	2	4	100%
cyano_pyridone_E	0	1	0	0	1	1	1	4	200%
anil_alk_thio	1	3	0	0	0	0	0	4	na
anil_di_alk_I	1	2	0	0	0	0	1	4	300%
het_thio_6_furan	0	1	1	1	1	0	0	4	na
anil_di_alk_ene_B	0	2	0	0	1	0	1	4	300%
imine_one_B	2	1	0	1	0	0	0	4	na
anil_OC_alk_A	1	0	0	0	2	0	1	4	300%
ene_five_het_J	0	0	1	0	1	0	2	4	50%
pyrrole_G	0	1	1	0	0	0	2	4	100%
ene_five_het_K	0	2	0	0	0	1	1	4	200%
cyano_ene_amine_B	0	0	2	1	0	0	1	4	300%
thio_ester_B	0	4	0	0	0	0	0	4	na
ene_five_het_L	0	0	1	1	0	1	1	4	200%
hzone_thiophene_B	0	1	1	1	0	1	0	4	na
dhp_amino_CN_E	0	0	3	1	0	0	0	4	na
het_5_B	0	1	1	0	0	1	1	4	200%

a. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays. na = not applicable (enrichment = infinity because of absence of any compound that hit no assays). Note that a high count of 2 is usually due to interference with HTS assays E and F and may usefully point to chelators

**Supplementary Table S10**. A comparison of functional group prevalence in MDDR 2008.1, the inaugural WEHI 93K HTS Library, and the current new CTX 153K HTS Library (comprising a collaboration library, and compounds from Vendor A and Vendor B).<sup>a</sup>

Entry	Group	Prevalence in (%)											
				CTX 1	53K HTS I	ibrary							
		MDDR <sup>a</sup>	WEHI <sup>b</sup>	16K	87K	49K							
				sub-	sub-	sub-							
				Library <sup>c</sup>	Library <sup>d</sup>	Library <sup>e</sup>							
1	Aromatic N	62	43	65	56	52							
2	Amides	50	32	42	52	47							
3	Alcohols	29	4.5	2.1	1.8	7.8							
4	Carboxylic Acid	26	6.0	2.3	4.3	2.5							
5	Lactams (not beta)	23	9.7	23	17	11							
6	Phenols	12	3.8	1.7	1.8	2.1							
7	Sulfonamides	10	6.8	19	19	8.7							
8	Amines	10	7.1	6.4	10	36							
9	Linear Ureas	6.3	4.0	4.6	4.6	2.8							
10	Linear Carbamates	5.7	1.8	0	1.0	1.2							
11	ketones	4.3	1.0	0	5.7	6.1							
12	sulfones	3.8	1.6	3.8	2.8	1.2							
13	Anilines (1°, 2°, 3° and cyclic or linear)	2.4	7.3	6.4	5.2	4.7							
14	Cyclic acyl ureas	2.4	1.9	3.6	2.0	0.3							
15	Cyclic carbamates	1.9	0.8	1.2	0.72	0.4							
16	Cyclic imides (phthalimides)	1.6 (0.6)	2.7	0.06 (0)	1.5	1.2							
17	Cyclic ureas	1.1	0.8	1.8	1.5	0.4							
18	Linear Imides	0.18	0.6	0.7	0.08	0.13							

a. MDDR 2008.1

It can be seen that there are similar overall trends in functional group prevalence for the WEHI 93K HTS Library compared with the new CTX 153K HTS Library, which we have segregated according to compound source (Collaborative Library, and Vendor A and Vendor B). That is, our filters for problematic compounds do not appear to have markedly altered functional group prevalence. Aromatic nitrogens are extremely common (entry 1) as are linear amides (entry 2) and then lactams (entry 5) and to some extent sulfonamides (entry 7), amines (entry 8) and then

b. Inaugural WEHI 93K HTS Library.

c. Collaborative HTS Library of 15,667 compounds established using separate criteria to those in Table 8, which we term the CTX 16K HTS Sub-Library.

d. 87,059 compounds from Vendor A, which we term the CTX 87K HTS Sub-Library

e. 49,144 compounds from Vendor B, which we term the CTX 49K HTS Sub-Library

anilines (entry 13) and alcohols (entry 3). Thereafter the remaining functional groups become significantly less prevalent. Even though trending the same way, for many functional groups there are significant fold differences depending on compound source. For example, the CTX 16K HTS Sub-Library has twice as many lactams as the CTX 16K HTS Sub-Library, whereas the latter has a very high percentage of amines (36%) and much higher than any other compound source. Similar variations are noted for alcohols, sulfonamides, ketones, cyclic carbamates, cyclic imides and linear imides.

We have also included analysis of MDDR 2008.1. Of note is the lower prevalence of alcohols, carboxylic acids, phenols, and linear carbamates in ordered screening compounds compared with MDDR, but the latter may be biased with highly populated classes (e.g. angiotensin converting enzyme inhibitors, Sweeteners, natural products etc) and this further analysis has not been undertaken.

**Table S11.** Rhodanines as problematic compounds.

Substructure <sup>a</sup>		Num	ber of	Alpha	Screen®	assays	hit	Total	Enrichment <sup>b</sup>
	6	5	4	3	2	1	0	Cpds	
ON S	16	41	21	26	22	20	60	225	2270/
ene_rhod_A	16	41	21	26	32	39	60	235	227%
R S S									
rhod_sat_A	0	6	6	6	6	7	2	33	1200%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

Rhodanines frequently occur as primary hits in our AlphaScreen® assays. There are alkylidene rhodanines with an enrichment of 227% for a total of 235 compounds, and saturated rhodanines, fewer in number (33) but with an astonishingly high enrichment of 1,200%. Focusing on the more common alkylidine rhodanines, these have been reported as inhibitors of anthrax lethal factor,  $^{21,22}$  glycosyltransferase MurG,  $^{23}$  SARS coronavirus,  $^{24}$  PRL-3,  $^{25}$  glycogen synthase kinase-3 $\beta$ ,  $^{26,27}$  HIV-1 integrase,  $^{28,29}$  extracellular signal-regulated kinase 2,  $^{30}$  tau aggregation,  $^{31}$  botulinum neurotoxin type A,  $^{32}$  *Plasmodium falciparum* enoyl-acyl carrier protein reductase,  $^{33}$  leucocyte migration (by stabilizing activated  $\alpha_{\rm M}\beta_{\rm 2}$  integrin),  $^{34}$  hepatitis C RNA polymerases,  $^{35}$  tumour necrosis factor- $\alpha$ ,  $^{38,39}$  UDP-galactopyranose mutase,  $^{40}$  tyrosine kinase p56 Lck SH2 domain,  $^{41}$  VHR phosphatase,  $^{42}$  formylpeptide receptor,  $^{43}$  RNA polymerase assembly,  $^{44}$  cholesterol accumulation,  $^{45}$  peptide deformylase,  $^{46}$  human apurinic/apyrimidinic endonuclease I,  $^{47}$  aggrecanase-2,  $^{48}$  *Trypanosoma brucei* dolicholphosphate mannose synthase  $^{49}$ , *Heliobacter pylori* shikimate kinase,  $^{50}$  protein tyrosine phosphatase-1B,  $^{51}$  *Yersinia* tyrosine phosphatase,  $^{52}$  retinoid x receptor- $\alpha$ ,  $^{53}$  *Yersinia* protein kinase A,  $^{54}$  and bacterial DNA adenine methyltransferases.

Despite extensive downstream investigation reported by many of these studies and attempts at describing SAR, these rhodanines are often bereft of meaningful SAR and poor correlation is often observed with cell-based activity where investigated. Clues as to why can be gleaned from the work of Carlsson et al. 40 who report that rhodanines can undergo extremely facile reaction with nucleophiles and so reactivitybased SAR is often observed. Indeed, workers at Amgen<sup>36,37</sup> have elucidated the crystal structure of a rhodanine-hepatitis C NS5b RNA complex revealing the formation of a covalent bond. Furthermore, work by DuPont<sup>38</sup> and Bristol-Myers Squibb<sup>39</sup> shows that rhodanines can undergo light-induced irreversible binding to tumor necrosis factor receptor-1. Disturbingly, two different reaction sites on the rhodanine scaffold are involved. Moreover, rhodanines are known chelators of transition metals and this has also been shown in the context of a protein's active site.<sup>22</sup> Common substituents include phenolic OH, carboxylic acid, halogen and nitro. It is possible that the former two can contribute more to chelation and the latter to reactivity (including light induced) but this currently remains speculative. Abbott have recently provided supporting evidence that rhodanines could be problematic protein-reactive screening compounds.<sup>6,7</sup>

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

We have additionally found that variations of rhodanines such as those shown below can also be frequent hitters in our AlphaScreen® assays (See Supplementary Tables S6-S8 and Figure S1).

These also often appear as screening hits in the literature alongside the more common rhodanine parent structures in studies we have already cited above. <sup>21,23,25,26,38-42,51,52,55</sup> We expand briefly for the "2-oxo" rhodanines and the "2-imino" rhodanines. The "2-oxo" rhodanines (thiazolidin-2,4-diones) have also been extensively studied for their peroxisome proliferator activated receptor gamma agonist activity as troglitazone-like compounds. and for their aldose reductase inhibition as epalrestat-like compounds. They have also been reported as inhibitors of protein tyrosine phosphatase-1b and low molecular weight protein tyrosine phosphatase are reported to also inhibit CFTR and DYRK1A. They are reported to also inhibit CFTR and DYRK1A. The SAR is generally confusing but especially so when the nitrogen is substituted and hence the ring not given the possibility of being deactivated through deprotonation of the relatively acidic NH proton.

Similar comments apply to "2-imino" rhodanines, which have also been reported as inhibitors of cyclooxygenase and lipooxygenase, HCV NS5B polymerase, PTP1B, mitogen-activated protein kinase phosphatase-1, NS5B polymerase, and cysteine proteases. They can also be highly cytotoxic. We have not undertaken an in-depth analyses of this sub-class, but it is plausible that these compounds could also interfere in assays in many of the ways that the parent rhodanines do, and indeed light-induced protein reactivity has been reported. 38,39

**Table S12.** Phenolic mannich bases as problematic compounds

Substructure <sup>a</sup>		Num	ber of	Alpha	hit	Total	Enrichment <sup>b</sup>		
	6	5	4	3	Cpds				
N OH									
mannich_A	2	4	13	15	59	57	146	296	64%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

Compounds with a 2-hydroxybenzylamine moiety frequently occur as primary hits in our AlphaScreen® assays with an enrichment of 64% for a total of 296 compounds. A brief survey of the literature revealed apparent hits for a number of different targets, including Hsp90,<sup>78</sup> L-type calcium channels,<sup>79</sup> bacterial ribosomal A-site,<sup>80</sup> human murine double mutant 2,<sup>81</sup> Mcl-1<sup>82</sup> and E2F.<sup>83</sup> As for the rhodanines, the molecules are often strikingly similar to each other. Disturbingly, this class of compounds can readily form highly reactive quinone methides<sup>84,85</sup> and they are also known to be metal chelators. Perhaps not surprisingly, these compounds can be potently cytotoxic towards bacteria. These compounds can also be highly cytotoxic to mammalian cells and susceptible to form intensely colored solutions (Theola Louie, personal communication).

While this paper was under review, the ability of this class to covalently modify proteins via the quinone methide was confirmed by Sanofi-Aventis and firms up these as problematic screening compounds. 92

b Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S13**. Hydroxyphenylhydrazones as problematic compounds.

Substructure <sup>a</sup>		Num	ber of	Alpha	Screen®	assays	hit	Total	Enrichment <sup>b</sup>
	6	5	4	3	2	1	0	Cpds	
N OH									
hzone_phenol_A	5	4	7	17	208	82	156	479	154%
N N N hzone_phenol_B	•	2	9	6	38	54	104	215	55%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

Hydrazones per se are close to being problematic in our assays (enrichment 28%) but scrutiny reveals certain sub-types of hydrazones are far more problematic than others and cart blanche removal of all hydrazones is unnecessary and likely to limit chemical diversity. Thus, 2- and 4-hydroxyphenylhydrazones frequently occur as primary hits in our AlphaScreen® assays with a respective enrichment of 154% and 55% for a total of 479 and 215 compounds respectively. These compounds often coexist in the literature with the other screening hits reported above 21,41,74,75,78,91 and have also been reported as inhibitors of type II topoisomerase, <sup>93</sup> anthrax lethal factor, <sup>94</sup> methionyl-tRNA synthetase, <sup>95</sup> 17β-hydroxysteroid dehydrogenase-1, <sup>96</sup> Mycobacterium tuberculosis<sup>97</sup> and Yersinia Type III secretion.<sup>98</sup> Closely related compounds such as the corresponding 2-hydroxybenzamides and the like are also usually co-reported screening hits. <sup>30,75,78-80,91,92,94,99,100</sup> These compounds can not only have unusual spectroscopic properties, 101,102 but they are also well known chelators. 103,104 Furthermore, the 2-hydroxyphenylhydrazones have been postulated to be able to form transient but highly reactive tautomeric quinone methide warheads 105 and unsurprisingly can be potently cytotoxic. 104 Additionally, they are known to interfere in bioassays through the formation of robust macromolecular aggregates.<sup>2</sup> disproportionately Intriguingly, the high count for the hydroxyphenylhydrazones is dominated by hits for HTS Campaigns E and F, which use a Nickel-based anchor and which would strongly support a component of interference through chelation. This is a good example where a compound may appear to be a hit in a primary assay and may give a readout in a cell-based secondary assay due to entirely separate mechanisms, the first in this example being chelation (or potentially aggregation under certain conditions) and the second mechanism being reactivity. Together, however, the two results in any given academic HTS laboratory may be convincing and publishable. This is also a good example why the selection of primary hit sets from six of our early screens that used relatively high screening concentrations, with the inclusion of two that would be susceptible to chelation interference in addition to other possible mechanisms of interference (apart from aggregation), from a library of typically commercially available sources, allows for maximum capture of compound classes likely to be discovered similarly and reported by HTS academics.

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S14.** Alkylidene barbiturates as problematic compounds

Substructure <sup>a</sup>	N	lumber	Total	Enrich-					
	6	5	4	3	2	1	0	Cpds	ment <sup>b</sup>
Het Het O,S NO									
ene_six_het_A	10	20	21	30	69	105	228	483	66%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

The substructure shown in Table S14 encodes for compounds that frequently occur as primary hits in our AlphaScreen® assays with an enrichment of 66% for a total of 483 compounds. These compounds are dominated by alkylidene barbiturates. These compounds are conspicuously prevalent as screening hits co-reported in studies already cited. Page 23,26,29,41,42,51-55,74,100 A brief survey of the literature reveals that they are also reported as inhibitors for Omi/HtrA2 protease, potentiators of  $\Delta F508$ -CFTR chloride channel gating, for inhibitors of ERK, dimethylarginine dimethylaminohydrolase, fatty acid synthase, for PPM1D, for acglucosidase, acglucosidase, and  $\alpha$ 4 $\beta$ 7-MAdCAM and RNA Polymerase. The structures involved are often very similar to each other and the SAR confusing. They are also Michael Acceptors and oxidants and undergo light-induced irreversible covalent binding with proteins. The thioxo counterparts in particular are prevalent in photoactive dyes that give rise to long-lived cytotoxic photoproducts. We note that the exocyclic alkene renders these compounds distinct from marketed barbituric acid-based pharmaceuticals that have dialkyl substitution instead of the alkene and so would not be reactive.

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S15.** Alkylidenes of 5-membered heterocycles as problematic compounds

Substructure <sup>a</sup>						assays		Total	Enrichment <sup>b</sup>
Substructure	6	5	4	3	2	assays	0	Cpds	Emicinient
0	0	3	4	3	2	1	U	Cpus	
<b>■ J</b> ĭ									
Hot									
Het									
ene_five_het_A	6	14	24	14	39	40	64	201	152%
N N									
S N								0.0	<b>7.7</b> 0 /
ene_five_het_B	0	4	4	2	14	22	44	90	55%
Het									
Het									
ene_five_het_C	3	9	7	7	7	13	39	85	85%
Q									
] N									
O N									
ene_five_het_D	4	7	8	9	13	5	0	46	na
0									
N Het									
7			2	1	,	,	_	10	000/
ene_five_het_G	0	0	2	1	1	1	5	10	80%
ene_five_het_H	0	1	0	0	2	3	0	6	na
	U	1	U	U	2	3	U	U	IIα

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

A number of 5-membered heterocycles bearing an exocyclic double bond interfere in our AlphaScreen® assays. In particular, the pyrazolidin-3,5-diones have a very high enrichment with no members that are clean. These sorts of compounds are prevalent in the screening literature already cited, <sup>21,23,28,40-42,45,50,51,66</sup> but are also reported in other screening literature as dual phosphodiesterase 1 and 5 inhibitors, <sup>119</sup> dual c-Src/Abl kinase inhibitors, <sup>120</sup> ubiquitin activating enzyme E1 inhibitors, <sup>121</sup> SARS-3CL pro inhibitors, <sup>122</sup> farnesoid X receptor inhibitors, <sup>123</sup> HIV-1 integrase inhibitors, <sup>124</sup> angiogenin and 5-lipoxygenase. <sup>127</sup> Notably, these hits are dominated by pyrazolidin-3,5-diones. Similar to the other compound classes discussed above, these compounds tend to all look very similar and where reported, there is no conventional SAR apparent.

It is also somewhat disconcerting that the 5-alkenylimidazolin-4-one core in the **ene\_5\_het\_G** family of assay interference compounds corresponds precisely to that in the chromophore of green fluorescent protein.<sup>128</sup>

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays. na = not applicable (enrichment = infinity because of absence of any compound that hit no assays).

**Table S16**. Fused tetrahydroquinolines as problematic compounds

Substructure <sup>a</sup>	ĺ	Num	ber of	Alpha	hit	Total	Enrichment <sup>b</sup>		
	6	5	4	3	Cpds				
N									
anil_alk_ene	1	6	6	3	7	11	17	51	135%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

Alkyl anilines feature strongly as problematic compounds in our assays. Amongst these, a distinctive class of tricyclic tetrahydroquinolines frequently appeared as primary hits in our AlphaScreen® assays and as shown, these totalled 51 in number and give rise to an enrichment value of 135%. These compounds also commonly occur in screening-based publications and are listed in works already cited. They are also conspicuous in publications as inhibitors of anthrax edema factor, low molecular weight protein tyrosine phosphatase, MIF, and protein-tyrosine phosphatases, Cdc25B dual specificity phosphatase and polo-like kinase 1. The structures involved are often very similar to each other and the SAR confusing. For example, weak but consistent inhibition against protein tyrosine phosphatase 1B is observed for the 6-F, 6-COOH and 6-OH analogs as well as the 8-H, 8-Br, 8-COOH and 8-Ac analogs.

**Table S17**. Pyrroles as problematic compounds

Table S17. 1 yrroles as									
Substructure <sup>a</sup>		Num	ber of	Alpha	Screen®	assays	hit	Total	Enrichment <sup>b</sup>
	6	5	4	3	2	1	0	Cpds	
pyrrole_A	1	16	13	14	11	21	42	118	131%
pyrrole_B	4	5	9	0	0	2	3	29	600%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

Certain pyrrole-containing compounds were prevalent as primary hits in our assays. A particularly troublesome group comprised N-aryl-2,5-dialkylpyrroles and N-alkyl-2-aryl-5-alkylpyrroles and as shown, these numbered some 118 and 29 respectively in our screening library with respective enrichment values of 131% and 600%. Very similar compounds are conspicuous in references already cited, 21,47,55,73,114,126,130,132 but have also been reported as inhibitors of HIV-1 fusion, 35 M. tuberculosis protein tyrosine phosphatase A inhibitors, 36 sphingosine 1-phosphate receptor agonists, 57 EphA4/EphA2 receptor inhibitors and inhibitors of metabolic glutamate receptor 1. An aryl carboxylate substituent frequently appears to be important for apparent

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

biological activity, for reasons that are unclear. As for the other PAINS described here, follow-up studies on close scrutiny reveal confusing SAR. 140

**Table S18**. Benzofurazans as problematic compounds.

Tuble 510. Benzolaruzun as problematie compounds.									
Substructure <sup>a</sup>		Num	ber of	Alpha	Total	Enrichment <sup>b</sup>			
	6	5	4	3	Cpds				
R <sub>2</sub> N S O									
diazox_sulfon_A	1	4	2	2	4	6	17	36	78%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1.

Certain activated benzofurazans frequently occurred as hits in our primary screens. As shown, an exemplary class of benzofurazan sulfonamides, while not that common in our library and numbering 36, registered an enrichment of 78%. Of note, similar compounds are reported as hits in screening papers already cited 33,93,98,130 and are also reported to be inhibitors of HIV-1 reverse transcriptase inhibitors, 141 *M. tuberculosis* CYP51,142 Yersinia type III secretion,143 measles virus RNA-dependent RNA polymerase complex activity,144 cholecystokinin-2 receptor antagonists and thymidine monophosphate kinase.146 SAR can be highly confusing.147

These compounds are highly unstable in vivo<sup>145</sup> and can be highly reactive sulfhydryl electrophiles, <sup>148</sup> used as fluoregenic and fluorescent labels, <sup>149</sup> and react rapidly with singlet oxygen. <sup>150</sup> Interestingly, though, this system can complex non-covalently with proteins <sup>142</sup> and has been the subject of a successful pharmacokinetic optimisation campaign, albeit one that required complete replacement of the benzofurazan with a phthalazine. <sup>145</sup> Abbott have recently provided supporting evidence that benzofurazans could be problematic protein-reactive screening compounds. <sup>6,7</sup>

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S19**. 2-Amino-3-carbonylthiophenes as problematic compounds

Substructure <sup>a</sup>		Num	ber of	Alpha	Screen®	hit	Total	Enrichment <sup>b</sup>	
	6	5	4	3	2	1	0	Cpds	
R S S N H thiophene_amino_Aa	2	2	5	4	3	11	18	45	94%
HO S O H N O thiophene_amino_Ab	0	2	2	1	5	7	23	40	43%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

These compounds are less common in our screening library but as shown, register frequently as hits with an enrichment value of up to 94%. This class is conspicuous as hits in references already cited, <sup>27,74,78,96,107,128,134</sup> but has already been reported as inhibitors of tubulin, <sup>151</sup> reverse transcriptase-associated ribonuclease H, <sup>152</sup> HCV RNA polymerase <sup>153</sup> and FLT3 tyrosine kinase. <sup>154</sup> Notably, in two studies, this class appeared to comprise promiscuous, non-specific hits when investigated in more detail. <sup>155,156</sup> We note that these compounds can form intensely blue solutions (Luigi Aurelio, personal communication). Abbott have recently provided supporting evidence that 2-amino-3-carbonylthiophenes could be problematic protein-reactive screening compounds that cause protein thiol oxidation. <sup>157</sup>

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S20**. Catechols as problematic screening compounds.

Substructure <sup>a</sup>		Num	ber of	Alpha	Total	Enrichment <sup>b</sup>			
	6	5	4	Cpds					
HO									
OH									
catechol A	4	7	10	4	10	21	36	92	97%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

Catechols are numerous (92) and problematic (enrichment 97%) in our screening campaigns. Catechols and hydroquinones are noticeable in screening literature already cited. <sup>2,25,75,78,94,125,131</sup> This activity may be due to oxidation to protein-reactive quinones. <sup>158</sup> Catechol-like templates are present in redox-reactive dyes such as pyocyanin. <sup>159</sup> Abbott have recently provided supporting evidence that catechols could be problematic protein-reactive screening compounds. <sup>6,7,157</sup>

**Table S21**. Quinones as problematic screening compounds

Substructure <sup>a</sup>		Num	ber of	Alpha	Total	Enrichment <sup>b</sup>			
	6	5	4	3	2	1	0	Cpds	
Het O									
quinone_A	40	57	48	41	42	56	86	370	265%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

Quinones and quinone-like compounds are amongst the most problematic of the compounds we have in our library, being both numerous - numbering some 370, and giving rise to a very high enrichment factor of 265%. Quinones are conspicuous in screening literature already cited 54,73,107,111,130,133 and are known to be protein-reactive. Abbott have recently provided supporting evidence that quinones could be problematic protein-reactive screening compounds. We also find that many quinones in our library are substituted with electron donating groups, giving rise to strongly colored compounds.

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S22**. Azo compounds as problematic compounds.

Substructure <sup>a</sup>		Num	ber of	Alpha	Total	Enrichment <sup>b</sup>			
	6	6 5 4 3 2 1 0							
R' <b></b> N <b>=</b> N-R"									
azo_A	29	30	33	43	24	55	110	324	145%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

There are significant numbers of azo compounds in our screening library, numbering some 324 and these lead to a very high enrichment value of 145%. Of note, these compounds are conspicuous in references already cited, 125,126,134 but are also reported as screening hits that inhibit Alzheimer's-related fibril formation, 162-164 avian influenza neuraminidase 165 and Bcl-2. 166 These compounds are often strongly colored and in fact are often dyes in reported assay studies 162-164 and can also undergo photoisomerization and reaction with biological nucleophiles. 167 They have also been reported to give rise to false positive results mediated via aggregation phenomena and can interfere with AlphaScreen® technology. 168

**Table S23**. Cyanopyridones as problematic compounds.

Table 525. Cyanopyridones as problematic compounds.									
Substructure <sup>a</sup>		Num	ber of	Alpha	Screen®	assays	hit	Total	Enrichment <sup>b</sup>
	6	5	4	3	0	Cpds			
H									
N ✓S									
CN									
cyano_pyridone_A	1	3	3	4	6	16	23	54	65%
H									
N <b>→</b> O									
CN									
cyano_pyridone_B	0	1	2	2	7	4	11	27	109%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

There are significant numbers of cyanopyridones, the most common numbering 54 and 27 with respective enrichment values of 65% and 109% respectively. Similar compounds are co-reported in references already cited. We note that such compounds can be effective thiol-reactive electrophiles. We note that such

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S24**. Divinylketones as problematic screening compounds

							1		
Substructure <sup>a</sup>		Num	ber of	Alpha	Total	Enrichment <sup>b</sup>			
	6	5	4	3	2	1	0	Cpds	
0									
ene_one_ene_A	1	2	3	5	6	16	24	57	71%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

Divinylketones are problematic screening compounds, numbering 57 in our Library and registering an enrichment value of 71%. We note that these compounds are reported in screening literature already cited. Abbott have recently provided supporting evidence that these could be problematic protein-reactive screening compounds. Of relevance here is work from the Cravatt group that suggests vinyl ketones readily and selectively react with protein cysteines, whereas tosylate esters readily react with a variety of residue types, including Asp, Glu and Tyr (Weerapana et al., Nat. Chem. Biol. 2008, 4, 405-407).

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S25**. Certain indoles as problematic screening compounds.

Substructure <sup>a</sup>	Νι					assay	s hit	Total	Enrichment <sup>b</sup>
	6	5	4	3	2	1	0	Cpds	
H,CH <sub>2</sub> R <sup>1</sup> H Any Any									
R <sup>1</sup> = CH <sub>2</sub> , C=Het,C:Het, CHN,CH(CH <sub>2</sub> )CH <sub>2</sub> NCH <sub>2</sub> indol_3yl_alk	6	15	23	18	51	80	268	461	42%
HN CO <sub>2</sub> H H,Alkyl Hal,H Allkyl/Aryl	0	0	2	3	7	4	14	30	86%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

One of our largest classes of problematic compounds comprises 3-alkylindoles, numbering some 461 compounds although as shown the enrichment value of 42% is not so high. We have not refined this class further, but it seems likely that several subclasses are involved. Some of these may specifically interfere in the AlphaScreen® technology, but it is likely that other subclasses may be PAINS. For example, we have noted that a particular subclass, indole-3-acetamide-2-carboxylic acids, number thirty within this broad class and as shown, have a high assay interference enrichment of 86%. It can be envisaged that the indoline tautomer of this class, with the enecarbonyl projecting from the indoline 3-position, could act as a Michael Acceptor (with or without further oxidative activation) as shown below.

$$NR_2$$
 $Nu^ NR_2$ 
 $Nu^ Nu^ NR_2$ 
 $Nu^ Nu^ NR_2$ 
 $Nu^ Nu^ Nu^-$ 

We have noticed these compounds in the screening literature reported as catalysts of peptide exchange, <sup>170</sup> mitogen-activated protein kinase phosphate-1 dual-specificity protein phosphatase inhibitors, <sup>171</sup> *Trypanosoma cruzi trans*-sialidase inhibitors, <sup>172</sup> and nitrosoglutathione reductase inhibitors. <sup>173</sup> As before, other PAIN structures already discussed are conspicuous in the latter three of these reports. We note that the broader class of indole-3-acetamides allowing any substituent off the 2 position (including H) yields 138 compounds with a relatively high enrichment 26%. This broader class is thus close to meeting our definition of being problematic in its own right, but like hydrazones, it is particular substituents that are likely to result in sub-classes that are particularly problematic. We believe there is the opportunity for many of our substructure filters to be thus refined further.

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

Table S26. Tertiary anilines as putative AlphaScreen®-specific problematic

screening compounds.

Anii_di_alk_B  anii_di_alk_C  15 23 13 10 26 29 130 246 67%  HC=C, R <sub>3</sub> CN,HO anii_di_alk_D  12 10 18 11 24 28 95 198 79%	screening compounds.									
Anil_di_alk_A  31 66 27 35 28 59 232 478 81%  Anil_di_alk_B  6 18 7 18 22 43 137 251 52%  Anil_di_alk_C  15 23 13 10 26 29 130 246 67%  HC=C, R <sub>3</sub> CN,HO anil_di_alk_D  12 10 18 11 24 28 95 198 79%	screening compounds.  Substructure <sup>a</sup>	N	umbe		phaSc	reen®	assays		Total	Enrichment <sup>b</sup>
Anil_di_alk_B  Anil_di_alk_B  Anil_di_alk_C  Anil_di_alk_C  Anil_di_alk_D  Anil_d		6	5	4	3	2	1	0	Cpds	
anil_di_alk_B 6 18 7 18 22 43 137 251 52%  NN*H,CR3  anil_di_alk_C 15 23 13 10 26 29 130 246 67%  HC=C, R3CN,HO anil_di_alk_D 12 10 18 11 24 28 95 198 79%	H	31	66	27	35	28	59	232	478	81%
## HC=C, R3CN,HO anil_di_alk_D 12 10 18 11 24 28 95 198 79%	>\								., 5	0170
## HC=C, R3CN,HO anil_di_alk_D 12 10 18 11 24 28 95 198 79%	anil di alle R	6	10	7	10	22	42	127	251	520/
anil_di_alk_C       15       23       13       10       26       29       130       246       67%         HC=C, R <sub>3</sub> CN,HO       12       10       18       11       24       28       95       198       79%         H,CH <sub>2</sub> H       H	anii_di_aik_B	6	18	/	18	22	43	13/	251	52%
HC=C, R <sub>3</sub> CN,HO anil_di_alk_D 12 10 18 11 24 28 95 198 79%  H,CH <sub>2</sub> H H,CH <sub>2</sub> H		15	23	13	10	26	29	130	246	67%
anil_di_alk_D         12         10         18         11         24         28         95         198         79%           H,CH2         H	\	10		13	10	20		150	2.0	0770
H,CH <sub>2</sub> H		12	10	10	11	24	20	05	109	709/
H,CH <sub>2</sub> H	anii_di_aik_D	12	10	18	11	24	28	95	198	79%
anil_di_alk_E   8   12   17   10   15   27   97   186   64%	H,CH <sub>2</sub> H									
	anil_di_alk_E	8	12	17	10	15	27	97	186	64%

a. Name corresponds to name in Supplementary Table S6-S8 and Figure S1

These compounds are less conspicuous in the screening literature that we have thus far cited. We propose that many of these compounds could interfere specifically with AlphaScreen®-technology, possibly through efficient quenching of singlet oxygen. We note the involvement of such compounds in singlet oxygen reaction in the AlphaScreen® beads themselves (Figure 3). We felt it important to investigate this further and so deliberately sourced and tested individual compounds related to the AlphaScreen® bead chemistry. The results are shown in Table S27.

b. Total number of compounds that hit from 2-6 assays expressed as a percentage of those compounds that have hit none of the six assays.

**Table S27**. Activity of certain anilines and olefins in the AlphaScreen® assay

Entry	Structure	Color of Solution	IC 50 (n=6)
1	Me <sub>2</sub> N NMe <sub>2</sub>	Colorless	$2.9 \pm 0.2 \mu\text{M}$
2	H <sub>2</sub> N NMe <sub>2</sub>	Rose	$30 \pm 3 \mu M$
3	Me NMe <sub>2</sub>	Colorless	>>100 μM
4	S S	Colorless	>>100 μM

Here is it apparent that tertiary anilines do not interfere in AlphaScreen® assay technology *per se* as the 4-methyl analogue is inactive (entry 3). However, when suitably functionalised, they can do so and bis-(*N*,*N*-dimethylaminophenyl)methane, entry 1, consistently gives rise to an IC<sub>50</sub> of around 3μM. It is plausible that this is due to quenching singlet oxygen, which would make it around 30 times as effective in this regard as DABCO (Table 9) under the same conditions. However, in none of the classes of tertiary anilines in Table S26 is it exclusive that all six assays are hit, which would be expected if singlet oxygen quenching was the only mechanism involved. Thus it is plausible that other unknown mechanisms are also involved.

## Drugs containing PAIN substructures - extended discussion

Epalrestat, which is marketed in Japan as an aldose reductase inhibitor for treating diabetic neuropathy (and more recently launched in India), contains a rhodaninebased PAIN structure but does not appear to have poor pharmacokinetics. compound contains an extended arylene group with a y-methyl substituent. It may be that our filters have the potential to be refined further to identify in more detail specific components that give rise to reactivity and discriminate from those that do not within a given class. Similarly, troglitazone, rosiglitazone and pioglitazone are rhodanine-like thiazolidinedione-based orally active antidiabetic agents for the treatment of type 2 diabetes. However, in these cases, there is serious concern associated with metabolically-activated toxic by-products, some of which may derive from the thiazolidinedione core. 190 Similarly, we have found several marketed drugs to contain PAIN substructures (in the following discussion the compound numbers refer to those listed in Supplementary Figure S2). This is because they contain catechols or hydroquinones (levodopa (25), adrenaline (30), apomorphine (31), dopamine (32), isoprenaline (33), noradrenaline (34), dobutamine (36), carbidopa (37), methyldopa (38), rifampicin (6), benserazide (18)) or guinones (menadione (1), phytomenadione (2), epirubicin (5), mitozantrone (8) (mitoxantrone), atavaguone (11), mitomycin (22), daunorubicin (23), idarubicin, (24)) or a masked quinoid (amsacrine (35)). These moieties are strongly associated with in vivo toxicity for reasons that also likely relate to their assay interference properties discussed herein. <sup>190</sup> In a significant study in 2002, <sup>191</sup> Hofmann La-Roche undertook a related exercise employing automated techniques as opposed to ours that took the more traditional route, relying more on the ability of an experience medicinal chemist to recognize and classify molecular structures. Despite the different approach, several catechol and quinone-based drugs were also identified as frequent hitters by the Roche group. These included dopamine, methyldopamine (methyldopa), benserazide, apomorphine and idebenone and so there is considerable overlap with those drugs similarly identified by us. The quinone-like clofazimine is also identified in both studies as a potentially problematic screening hit. Abbott has also identified the quinone-based cytotoxic doxorubicin as an ALARM-NMR-reactive drug. In a number of cases, therefore, there appears to be a direct link between poorly tolerated marketed drugs and a PAIN-containing structure, where protein-reactivity, either directly or indirectly after metabolic activation, may be a common underlying mechanism. The work by Abbott supports this notion.<sup>7</sup>

There are a number of other drugs in Supplementary Figure S2 that are also recognised by our assay interference filters but which we have not specifically categorized herein as PAINS because there is not the literature evidence to do so. Even so, the substructure involved in many of these also appears to be that which in the drug is also associated with its major metabolic liabilities. For example, the ethylthiomethyleneimidazole in cimetidine (41) is recognized as an assay interference moiety by our filters and it this group in cimetidine which is a major metabolic liability (oxidation of the sulfur atom and engagement and inhibitions of CYPs by the imidazole ring). Similarly, olsalazine (13), balsalazide (14) and sulfasalazine (7) are recognized by our filters because they contain an azo group. This group is unstable in vivo and effectively acts as a prodrug by being metabolised by intestinal

bacteria. <sup>194</sup> An azo group is recognised in the triazene of dacarbazine (20) and the azide in zidovudine (12, AZT) and in both cases is associated with instability or reactivity, though in the case of dacarbazine this reactivity also represents its principle mechanism of antitumor action (DNA methylation). Tricyclics such as promethazine (43) and trimeprazine (44) are recognized by our assay interference filters because they contain a phenothiazine group, and it is the sulfur atom of this which is also one of the major metabolic liabilities. <sup>197</sup> The phenothiazine core has also been linked to phototoxicity. <sup>198</sup> Phenindione (26) is an old and dirty drug <sup>199</sup> and is recognized as a potential assay interference compound primarily by virtue of its 1,3-diketone group. Topotecan (9) is an antitumour drug that is recognized by our filters by virtue of the phenolic mannich base, which has been associated with aqueous instability with the formation of a reactive quinone methide. <sup>200</sup> Similarly, pyrroles related to those that we find are assay interference compounds can be DNA-reactive<sup>201</sup> and that in Atorvastatin has been linked to phototoxicity. <sup>202</sup>

However, there are some drugs whose substructures recognised by our filters do not appear to impart any metabolic liability. These include the antifungals itraconazole (3), posaconazole (4), ketoconazole (17) and the antibiotic linezolid (16), all of which are recognized by our filters because of the presence of a tertiary aniline but this does not appear to introduce a metabolic liability in these particular cases, though we note that the electron-rich aromatic rings tend to be disfavoured by medicinal chemists as starting points because they are prone to metabolic oxidation. A similar argument applies for the tricyclics amitriptyline (40), nortriptyline (42), flupenthixol (45), zuclopenthixol (46) and thiothixene (47) which are also filtered out and so also appear in Supplementary Figure S2 because they contain a particular type of styrene functionality but which appears to be metabolically stable; though we note here too that the styrene core has been linked to the possibility of metabolically-induced toxicity.<sup>203</sup> Thiothixene, amitryptiline and nortriptyline also comprise three of the four CNS-active drugs that were filtered out in Table 7. The fourth CNS-active drug filtered out was frovatriptan, due to its indole-3-alkyl-containing substructure, but the pharmacokinetics of this compound appear to be acceptable and the indole nucleus not a metabolic liability.<sup>204</sup>

grouped according to ty	pe		
Hev Hev Hev O			O NH NH
1:acyl_het_A(9)	2:amino_acridine_A(1)	3:amino_acridine_A(46)	4:anil_NH_alk_A(5)
NH NH Hev	NH NH	NH NH2	N.H2 N
5:anil_NH_alk_B(3)	6:anil_NH_alk_C(2)	7:anil_NH_alk_D(2)	8:anil_NH_no_alk_A(1)
9:anil_NH_no_alk_B(1)	10:anil_OC_alk_A(4)	11:anil_OC_alk_B(3)	Any O 12:anil_OC_alk_C(3)
NH NH NA	ONH OH	S NH S O S N F (4)	NH2
13:anil_OC_alk_D(2)	14:anil_OC_alk_E(1)	15:anil_OC_alk_F(1)	16:anil_OC_no_alk_A(8)
NH O	NH2 O	NH OOH	OH NH2
17:anil_OC_no_alk_B(4)	18:anil_OC_no_alk_C(3)	19:anil_OH_alk_A(8)	20:anil_OH_no_alk_A(1)

Hits: 1 - 20

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O OH NH2 21:anil_OH_no_alk_B(1)	NH- NH- NH- O- 22:anil_alk_A(1)	23:anil_alk_B(1)	O NH 24:anil_alk_C(1)
		= = (/	= = \( \tau \)
NH	N NH O		
25:anil_alk_D(1)	26:anil_alk_bim(9)	27:anil_alk_ene(51)	28:anil_alk_indane(1)
Hev N N 29:anil_alk_thio(4)	Any N N N N N N N N N N N N N N N N N N N	31:anil_di_alk_B(251)	O N Any 32:anil_di_alk_C(246)
33:anil_di_alk_D(198)	Any N Any N 34:anil_di_alk_E(186)	Any Any Any Any 35:anil_di_alk_F(14)	*N Hev* 36:anil_di_alk_G(9)
Hev O NH  37:anil_di_alk_H(6)	38:anil_di_alk_I(4)	39:anil_di_alk_J(3)	NH2 NH2 40:anil_di_alk_K(2)

Hits: 21 - 40

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41:anil_di_alk_L(1)	42:anil_di_alk_M(1)	43:anil_di_alk_N(1)	S NH NH NH NH A4:anil_di_alk_O(1)
ş	Ĵ		,
	NH O		
45:anil_di_alk_P(1)	46:anil_di_alk_coum(1)	47:anil_di_alk_dhp(1)	48:anil_di_alk_ene_A(8)
	Het NH		O O O O O O O O O O O O O O O O O O O
49:anil_di_alk_ene_B(4)	50:anil_di_alk_furan_A(1~	51:anil_di_alk_furan_B(2)	52:anil_di_alk_indol(1)
Any NH2	NH2	NH2	NH2
53:anil_no_alk(40)	54:anil_no_alk_A(1)	55:anil_no_alk_B(1)	56:anil_no_alk_C(1)
NH2 NH2	N H2	Any	NH
57:anil_no_alk_D(1)	58:anil_no_alk_indol_A(1)	<b>59:anisol_A(5)</b>	60:anisol_B(2)

Hits: 41 - 60

**S63** Page: 3

NH OH  61:anthranil_acid_A(19)	62:anthranil_acid_B(3)	OH NH2 OH NH2 OH Any 63:anthranil_acid_C(2)	O NH2 S S S S S S S S S S S S S S S S S S S
	()	(-)	
O NHOON	о он он	OH OH	OH OH
65:anthranil_acid_E(2)	66:anthranil_acid_F(2)	67:anthranil_acid_G(1)	68:anthranil_acid_H(1)
O OH OH	OH O NH	O NH NH NH	O NH NH
69:anthranil_acid_I(1)	70:anthranil_acid_J(1)	71:anthranil_amide_A(1)	72:anthranil_one_A(38)
!RNN 73:azo_A(324)	74:azulene(4)	OH OH 75:catechol_A(92)	76:colchicine_A(3)
_ (===)	-(-)	()	·\-/
OH OH			O O O O O O O O O O O O O O O O O O O
77:colchicine_B(1)	78:colchicine_het(1)	79:coumarin_A(2)	80:coumarin_B(2)

Hits: 61 - 80

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Page: 4

81:coumarin_C(1)	82:coumarin_D(1)	83:coumarin_E(1)	84:coumarin_F(1)
			_
Hal ONH NH2	S NH NH	N $N$ $N$ $N$ $N$ $N$ $N$ $N$ $N$ $N$	NH2 NH2 NH2
85:coumarin_G(1)	86:coumarin_H(1)	87:cyanamide_A(1)	88:cyano_amino_het_A(1)
NH2 NH2 NH2  S NH2 NH2 NH2 NH2 NH2 NH2	Any——N 90:cyano_cyano_A(23)	91:cyano_cyano_B(3)	NH2 NH2 92:cyano_ene_amine_A(56)
93:cyano_ene_amine_B(4)	O NH ONH NH NH NH Square (3)	NH 95:cyano_imine_A(37)	96:cyano_imine_B(17)
NH N O		$ \longrightarrow \hspace{-1.5cm} \bigcirc \hspace{-1.5cm} \bigcirc \hspace{-1.5cm} \backslash \hspace$	N
97:cyano_imine_C(12)	98:cyano_imine_D(1)	99:cyano_keto_A(2)	100:cyano_misc_A(1)

Hits: 81 - 100

**S65** Page: 5

NH——N Hev—Hev  101:cyano_pyridone_A(54)	Het N = N N N N N N N N N N N N N N N N N	ONN  103:cyano_pyridone_C(11)	N N O 104:cyano_pyridone_D(5)
Hev NH O			NH NH S
105:cyano_pyridone_E(4)	106:cyano_pyridone_F(3)	107:cyano_pyridone_G(1)	108:dhp_amidine_A(1)
ONH2	Any N N N N N N N N N N N N N N N N N N N	NH2	NH2 NH2
109:dhp_amino_CN_A(13)	110:dhp_amino_CN_B(9)	111:dhp_amino_CN_C(7)	112:dhp_amino_CN_D(5)
	NH2	Any O NH2-N	S——N NH2
113:dhp_amino_CN_E(4)	114:dhp_amino_CN_F(3)	115:dhp_amino_CN_G(1)	116:dhp_amino_CN_H(1)
N-H2—N N-H2	Any		NO N
117:dhp_bis_amino_CN(19)	118:dhp_keto_A(9)	119:diazox_A(3)	120:diazox_B(3)
	1 = - 1	= \ /	= \ /

Hits: 101 - 120

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Het  N  N  N  N  N  N  N  121:diazox_C(1)	Неt N ОН—ОН 122:diazox_D(1)	NH2—NH2  123:diazox_E(1)	Any O O O O O O O O O O O O O O O O O O O
	= ` `	_ ` `	= = , /
	sirir '* 'N+* · N	**************************************	
125:diazox_sulfon_B(5)	126:dyes3A(19)	127:dyes5A(27)	128:dyes7A(2)
	NH2	NH=\(\big \)N	S $S$ $S$
129:ene_cyano_A(19)	130:ene_cyano_B(7)	131:ene_cyano_C(6)	132:ene_cyano_D(3)
0 N	OH OH OH ON NON		Het
133:ene_cyano_E(1)	134:ene_cyano_F(1)	135:ene_cyano_G(1)	136:ene_five_het_A(201)
S NO	Het	o N O	Het—(NR)
137:ene_five_het_B(90)	138:ene_five_het_C(85)	139:ene_five_het_D(46)	140:ene_five_het_E(44)
<del></del>			

Hits: 121 - 140

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141:ene_five_het_F(15)	Hev Hev Any Any  142:ene_five_het_G(10)	S N—N 143:ene_five_het_H(6)	Het (NR) O 144:ene_five_het_I(6)
	1121010_1110_1101_0(10)	Tiorene_iive_net_ii(0)	TTMENC_ITYC_ICC_I(0)
O S NH	Het *		O Any
145:ene_five_het_J(4)	146:ene_five_het_K(4)	147:ene_five_het_L(4)	148:ene_five_het_M(3)
Hal  S  S  O  149:ene_five_het_N(1)	150:ene_five_one_A(55)	151:ene_five_one_B(1)	0 0 0 152:ene_misc_A(5)
	NH OO		
153:ene_misc_B(2)	154:ene_misc_C(1)	155:ene_misc_D(1)	156:ene_misc_E(1)
Hey-Hev Hev-Hev ON	s No	N S(4)	он О
157:ene_one_A(3)	158:ene_one_B(2)	159:ene_one_C(1)	160:ene_one_D(1)

Hits: 141 - 160

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NH O O O 161:ene_one_amide_A(1)	162:ene_one_amide_B(1)	Hev Hev	0 NH 0 0 164:ene_one_ester(24)
Tor.enc_onc_amuc_A(1)	102:ene_one_amide_b(1)	103.cnc_onc_enc_A(37)	104.ene_one_ester(24)
Hal	S O	O NH	
165:ene_one_hal(17)	166:ene_one_one_A(1)	167:ene_one_B(1)	168:ene_one_yne_A(1)
Hev	SN	O N O O N O O O O O O O O O O O O O O O	Any S
169:ene_quin_methide(10)	170:ene_rhod_A(235)	171:ene_rhod_B(16)	172:ene_rhod_C(13)
Any O NRN Any Any	s s	O N N Het	Het Het Hev Hev OH
173:ene_rhod_D(8)	174:ene_rhod_E(8)	175:ene_rhod_F(8)	176:ene_rhod_G(7)
	Br. NH S	Het	Het Het Het
177:ene_rhod_H(5)	178:ene_rhod_I(3)	179:ene_rhod_J(3)	180:ene_six_het_A(483)
	<u>-</u> <del></del> (-)	<u>-</u> (-)	

Hits: 161 - 180

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181:ene_six_het_B(2)	182:ene_six_het_C(2)	Hev. Hev. No. 183:ene_six_het_D(1)	OH ====================================
		s—	
о он		NH O	
185:furan_acid_A(4)	186:het_465_misc(1)	187:het_55_A(2)	188:het_55_B(1)
189:het_565_A(2)	190:het_565_indole(1)	0 N N 191:het_5_A(7)	0 NH NH 192:het_5_B(4)
107.Hct_303_A(2)	170.nct_505_mdote(1)	171.nct_3_A(1)	172.nct_5_b(4)
F F F Any			NH O O
193:het_5_C(2)	194:het_5_D(2)	195:het_5_E(1)	196:het_5_ene(1)
	OH N————Any	OH N	NH Hev Hev Hev Hev
197:het_5_inium(1)	198:het_5_pyrazole_OH(14)	199:het_65_A(21)	200:het_65_B(7)

Hits: 181 - 200

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201:het_65_C(6)	202:het_65_Da(2)	203:het_65_Db(5)	OH NH S NH ONH S ON S
205:het_65_F(1)	NH2 NH2 N N N N N N N N N N N N N N N N	207:het_65_H(1)	208:het_65_I(1)
209:het_65_J(1)	N—N 210:het_65_K(1)	207:Het_05_H(1)  0 NH2  N_N_N  211:het_65_L(1)	208:het_05_l(1)  NH NH NH 212:het_65_imidazole(1)
213:het_65_mannich(1)	Any Het  N Het  214:het_65_pyridone_A(3)	215:het_6666_A(2)	216:het_666_A(5)
217:het_666_B(3)	218:het_666_C(1)	219:het_66_A(7)	220:het_66_B(2)

Hits: 201 - 220

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OH OH 221:het_66_C(2)	222:het_66_D(1)	он 223:het_66_E(1)	224:het_66_anisole(1)
O NH NH	NH NH N	N H2 NH2 OH	N.H2 OH Hev
225:het_6_hydropyridone(~  OH OH OH 229:het_6_pyridone_OH(5)	Hev N	227:het_6_imidate_B(1)  NH  NH  231:het_76_A(1)	228:het_6_pyridone_NH2(1)  NN NN NH OFF F  232:het_pyraz_misc(1)
Hev	Hev Hev Any  234:het_pyridiniums_B(2)	N	N N N S
NH S 237:het_thio_5_B(2)	238:het_thio_5_C(2)	N N S N 239:het_thio_5_imine_A(1)	240:het_thio_5_imine_B(1)

Hits: 221 - 240

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241:het_thio_5_imine_C(1)	242:het_thio_656a(5)	243:het_thio_656b(3)	244:het_thio_656c(1)
	NH NH	NON S ON NH N	Hal O NH- S NH- O
245:het_thio_65_A(3)	246:het_thio_65_B(2)	247:het_thio_65_C(2)	248:het_thio_65_D(1)
249:het_thio_665(1)	Any Any 250:het_thio_666_A(13)	251:het_thio_66_A(3)	252:het_thio_66_one(8)
247.nct_tmo_003(1)	230:hct_tmo_000_A(13)	251:hct_tmo_00_A(5)	232.net_tmo_oo_one(o)
Any		N S S	
253:het_thio_676_A(10)	254:het_thio_676_B(1)	255:het_thio_67_A(1)	256:het_thio_6_ene(2)
	Het—S N N N S5(5)	N N N N N S A (2)	S. N. N. S.
257:het_thio_6_furan(4)	258:het_thio_N_55(5)	259:het_thio_N_5A(3)	260:het_thio_N_5B(2)

Hits: 241 - 260

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262:het_thio_N_5D(1)	263:het_thio_N_65A(3)	O— N S 264:het_thio_pyr_A(3)
OH (NR) N		0 HV
Any O	OH OO	268:hzone_acid_A(1)  NH O OH 272:hzone_acyl_naphthol(~
274:hzone_anil_di_alk(35)	275:hzone_anthran_Z(1)	Any (NR) Any 276:hzone_enamin(30)
Any NH N Any 278:hzone_furan_B(2)	OH O OH O OH O O O O O O O O O O O O O	ONT Hev OH  280:hzone_furan_E(1)
	266:hydroquin_A(2)  Any Any Any Any Any Any 270:hzone_acyl_misc_A(1)  Any Any Any Any Any Any Any Any Any An	266:hydroquin_A(2)  267:hzide_naphth(2)  Any Any Any OH

Hits: 261 - 280

S74 Page: 14

Any NH	N OH  282:hzone_phenol_A(479)	OH  283:hzone_phenol_B(215)	Any Any 284:hzone_phenone(7)
Hev		Any Het NH Any	Any
285:hzone_pipzn(79)	286:hzone_pyrrol(64)	287:hzone_thiophene_A(11)	288:hzone_thiophene_B(4)
Hev	NH	Br Br	(NR) N NH2
289:imidazole_A(19)	290:imidazole_B(2)	291:imidazole_C(1)	292:imidazole_amino_A(1)
N N			IR N
293:imine_ene_A(5)	294:imine_ene_one_A(3)	295:imine_ene_one_B(1)	296:imine_imine_A(9)
		OH OH	Het 'R'Het Any
297:imine_imine_B(3)	298:imine_imine_C(3)	299: $imine_naphthol_A(1)$	300:imine_one_A(321)

Hits: 281 - 300

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301:imine_one_B(4)	Het Hev :R:Het  302:imine_one_fives(89)	S O S N N N N N N N N N N N N N N N N N	Het N Het N 304:imine_one_fives_C(2)
S Het	(NR)N N	O N O	OH ;R <sup>N</sup>
305:imine_one_fives_D(1)	306:imine_one_isatin(189)	307:imine_one_sixes(27)	308:imine_phenol_A(3)
Any	N. (NR)		N O
309:indol_3yl_alk(461)	310:indole_3yl_alk_B(1)	311:keto_keto_beta_A(68)	312:keto_keto_beta_B(12)
ОН	ОМОН	о ОН ОН	ОН
313:keto_keto_beta_C(7)	314:keto_keto_beta_D(5)	315:keto_keto_beta_E(1)	316:keto_keto_beta_F(1)
NH NH		Any	Hev
317:keto_keto_beta_zone(~		319:keto_naphthol_A(2)	320:keto_phenone_A(11)

Hits: 301 - 320

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321:keto_phenone_B(1)	322:keto_phenone_C(1)	323:keto_phenone_zone_A(~	324:keto_thiophene(3)
0.17	$\rangle$	0	Ġ?
OH			NH NON NH Any Any
325:mannich_A(296)	<b>326:</b> mannich_B(1)	327:mannich_catechol_A(1)	328:melamine_A(3)
	OH O	ONH	NHO NHO A 222 min a suitide A(1)
329:melamine_B(1)	330:misc_aminal_acid(1)	331:misc_aminoacid_A(1)	332:misc_anilide_A(1)
O NH NH	NH NH		O NH NH
333:misc_anilide_B(1)	334:misc_anisole_A(1)	335:misc_anisole_B(1)	336:misc_anisole_C(1)
NH O		CI CI	N.H2 NH2 NH
337:misc_cyclopropane(1)	338:misc_furan_A(1)	339:misc_imidazole(1)	340:misc_naphthimidazole~

Hits: 321 - 340

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OH SON	O NH	O NH	
341:misc_phthal_thio_N(1)	342:misc_pyridine_OC(1)	343:misc_pyrrole_benz(1)	344:misc_pyrrole_thiaz(1)
OH OH	S OH	O+NHN	
345:misc_stilbene(1)	346:misc_trityl_A(1)	347:misc_urea_A(1)	348:naphth_amino_A(25)
NH NH	NH N		
349:naphth_amino_B(25)	350:naphth_amino_C(2)	351:naphth_amino_D(2)	352:naphth_ene_one_A(1)
F F	N N N N N N N N N N N N N N N N N N N	OH OH	OH OO
353:naphth_ene_one_B(1)	354:naphth_ene_one_C(1)	355:phenol_sulfite_A(1)	356:phthalimide_misc(2)
N N H2	N NH NH2	Hev	Hev
357:pyrazole_amino_A(1)	358:pyrazole_amino_B(1)	359:pyrrole_A(118)	360:pyrrole_B(29)

Hits: 341 - 360

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361:pyrrole_C(8)	NH NH S 362:pyrrole_D(5)	N Hev Hev 363:pyrrole_E(5)	N N 364:pyrrole_F(5)
Any	N N	O N	s s
365:pyrrole_G(4)	366:pyrrole_H(3)	367:pyrrole_I(2)	368:pyrrole_J(1)
NH O 369:pyrrole_K(1)	370:pyrrole_L(1)	OH  371:pyrrole_M(1)	372:pyrrole_N(1)
373:pyrrole_O(1)	Het  * Het  374:quinone_A(370)	375:quinone_B(5)	OH ON 376:quinone_C(2)
Het	SNS	Hev Hev E Hev Hev Hev Hev S O N S	
377:quinone_D(2)	378:rhod_sat_A(33)	379:rhod_sat_B(3)	380:rhod_sat_C(3)

Hits: 361 - 380

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381:rhod_sat_D(3)	382:rhod_sat_E(1)	383:rhod_sat_F(1)	384:rhod_sat_imine_A(1)
		Any	
385:steroid_A(2)	386:styrene_A(13)	387:styrene_B(8)	388:styrene_C(4)
NH2		Hal OH ON N	OH NH S 0
389:styrene_anil_A(1)	390:styrene_imidazole_A(~	391:sulfonamide_A(43)	392:sulfonamide_B(41)
O O O Het	O NH ONH ONH ONH ONH ONH ONH ONH ONH ONH O	O	S O NH S O
393:sulfonamide_C(5)	394:sulfonamide_D(2)	395:sulfonamide_E(2)	396:sulfonamide_F(1)
O NH N	NH2	ON NH	NHO SO SO
397:sulfonamide_G(1)	398:sulfonamide_H(1)	399:sulfonamide_I(1)	400:sulfonamide_J(1)

Hits: 381 - 400

S80 Page: 20

401:tert_butyl_A(2)	OH OH OH OH 402:tert_butyl_B(1)	403:tetrazole_A(1)	HevHev Hev
S Any Any Any	S (NR) O	O S (NR) N	S N±- N-O
405:thiaz_ene_A(128)	406:thiaz_ene_B(17)	407:thiaz_ene_C(11)	408:thiaz_ene_D(8)
Any S RC  409:thiaz_ene_E(8)	S S S 410:thiazol_SC_A(3)	NH2 NH S NH 411:thiazole_amine_A(4)	412:thiazole_amine_B(3)
Hev Hev Any Hev	**Any NH  414:thiazole_amine_D(3)	NH N	SNN NH 416:thiazole_amine_F(2)
	NHOOH OH	N <sub>NH</sub> -NH	NH NH O
417:thiazole_amine_G(2)	418:thiazole_amine_H(1)	419:thiazole_amine_I(1)	420:thiazole_amine_J(1)

Hits: 401 - 420

S81 Page: 21

421:thiazole_amine_K(1)	NH2 S NH2 S 422:thiazole_amine_L(1)	NH-NH-NH-NH-NH-NH-NH-NH-NH-NH-NH-NH-NH-N	ON NHO ON
ZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZZ			.2
s	Hev	NH NH	N.H. S
425:thio_aldehyd_A(3)	426:thio_amide_A(6)	427:thio_amide_B(2)	428:thio_amide_C(2)
429:thio_amide_D(2)	430:thio_amide_E(1)	Hev NH Hev S  431:thio_amide_F(1)	SNH S 432:thio_carbam_A(1)
		/	= = :/
S	Any	S	NH—N
433:thio_carbam_ene(2)	434:thio_carbonate_A(15)	435:thio_carbonate_B(3)	436:thio_cyano_A(1)
Het *	8 A (1)	s s s	S
437:thio_dibenzo(23)	438:thio_ene_amine_A(1)	439:thio_est_cyano_A(1)	440:thio_ester_A(5)

Hits: 421 - 440

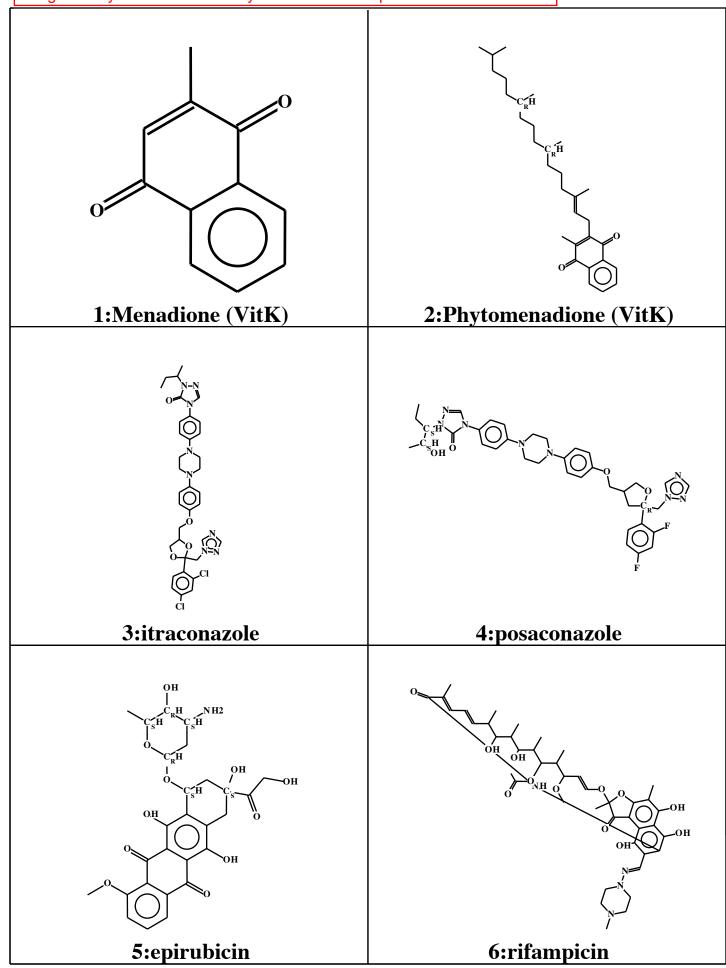
S82 Page: 22

S Any S Any 441:thio_ester_B(4)	O N S S 442:thio_ester_C(2)	443:thio_imide_A(1)	OH N <sup>y</sup> S  444:thio_imine_ium(2)
< /	/	/	/
	s	N Hev	S S
445:thio_keto_het(2)	446:thio_ketone(43)	447:thio_pyridine_A(1)	448:thio_thiomorph_Z(1)
NH SNH O 449:thio_urea_A(12)	450:thio_urea_B(9)	Het Hev NH S O  451:thio_urea_C(9)	SNH NH 452:thio_urea_D(8)
NH SNH	NH NH NH	NH R'N	NH O
453:thio_urea_E(7)	454:thio_urea_F(6)	455:thio_urea_G(5)	456:thio_urea_H(3)
NH S	NH NH	NH-\S	Hev Hev NNHNNH
457:thio_urea_I(3)	458:thio_urea_J(2)	459:thio_urea_K(2)	460:thio_urea_L(1)

Hits: 441 - 460

S83 Page: 23

Hal NH NH S  461:thio_urea_M(1)	SNH NH NH A62:thio_urea_N(1)	Volume NH NH S	NH NN (NR) O  464:thio_urea_P(1)
401.0110_01@a_W(1)	402.0110_01 ea_1\(1)	405.mio_urea_O(1)	404.tmo_urea_F(1)
S NH NH	S NH	Any S S	ONH—NH
465:thio_urea_Q(1)	466:thio_urea_R(1)	467:thiophene_C(3)	468:thiophene_D(2)
Any O S O NH O S O NH O S O S O S O S O S O S O S O S O S O	S NH—S 0 470:thiophene_F(1)	Hev NH2 471:thiophene_amino_Aa(4	O NH S S Any Any 472:thiophene_amino_Ab(4~
Any NH S Any NH 473:thiophene_amino_B(12)	ONH Any S 474:thiophene_amino_C(7)	475:thiophene_amino_D(3)	476:thiophene_amino_E(2)
Hev'lev Hev'lev Hev'lev Hev'lev Hev NH2 NH2 NH0	O NH S OH S	ONH NH	OH
477:thiophene_amino_F(2)	478:thiophene_amino_G(2)	479:thiophene_amino_H(2)	480:thiophene_hydroxy(28)



Hits: 1 - 6

S85 Page: 1

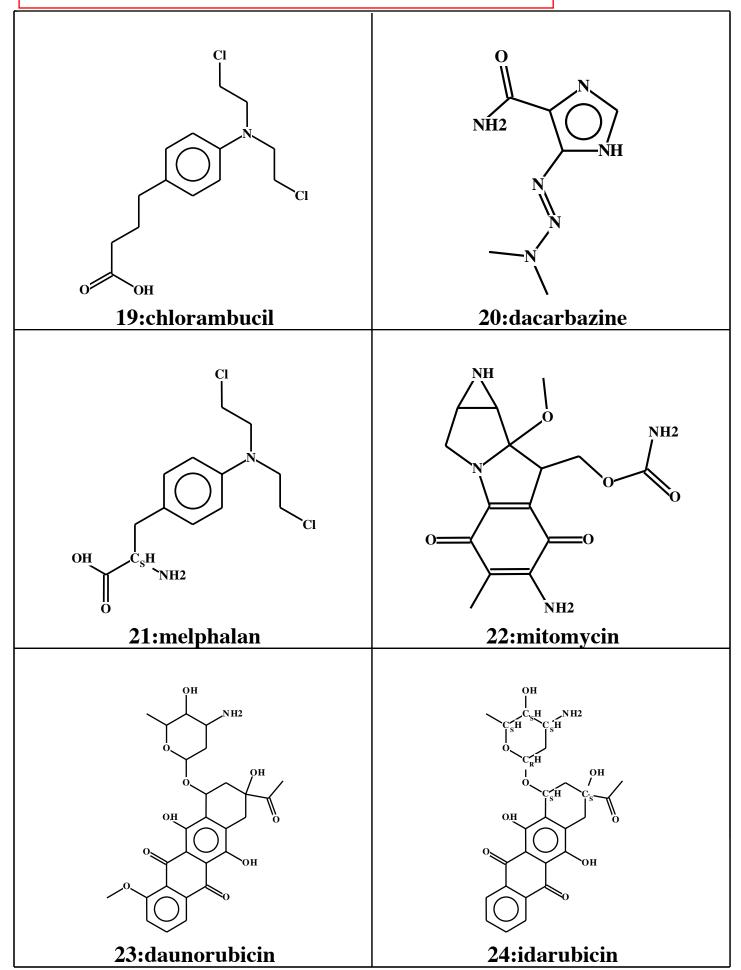
	7 AVIII_Inttel5.Ints
OH O	он он он NH NH он 8:mitozantrone
OH OH	O C'H O C'SH
9:topotecan	10:rifabutin
$OH \qquad C_1H \qquad C_1H$	$\begin{array}{c} N = N^{+} = N \\ OH \\ C_{S}H \\ C_{R}H \\ NH \\ O \end{array}$
11.stavaanana	12.5idomidino
11:atovaquone	12:zidovudine

Hits: 7 - 12

S86 Page: 2

Hits: 13 - 18

S87 Page: 3

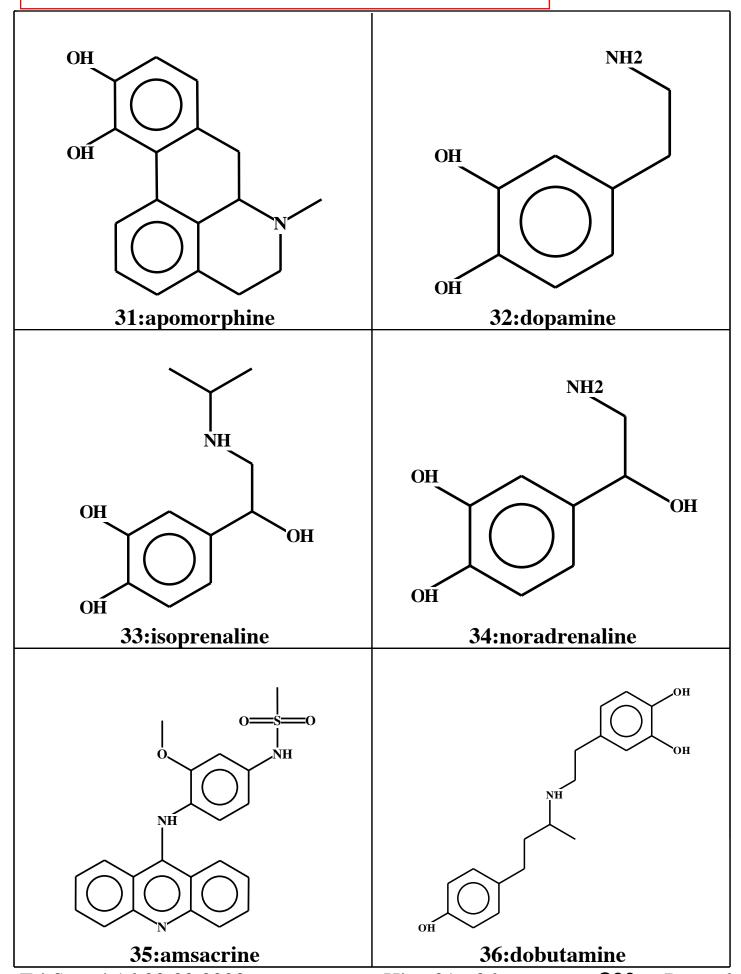


Hits: 19 - 24

S88 Page: 4

Hits: 25 - 30

S89 Page: 5



Hits: 31 - 36

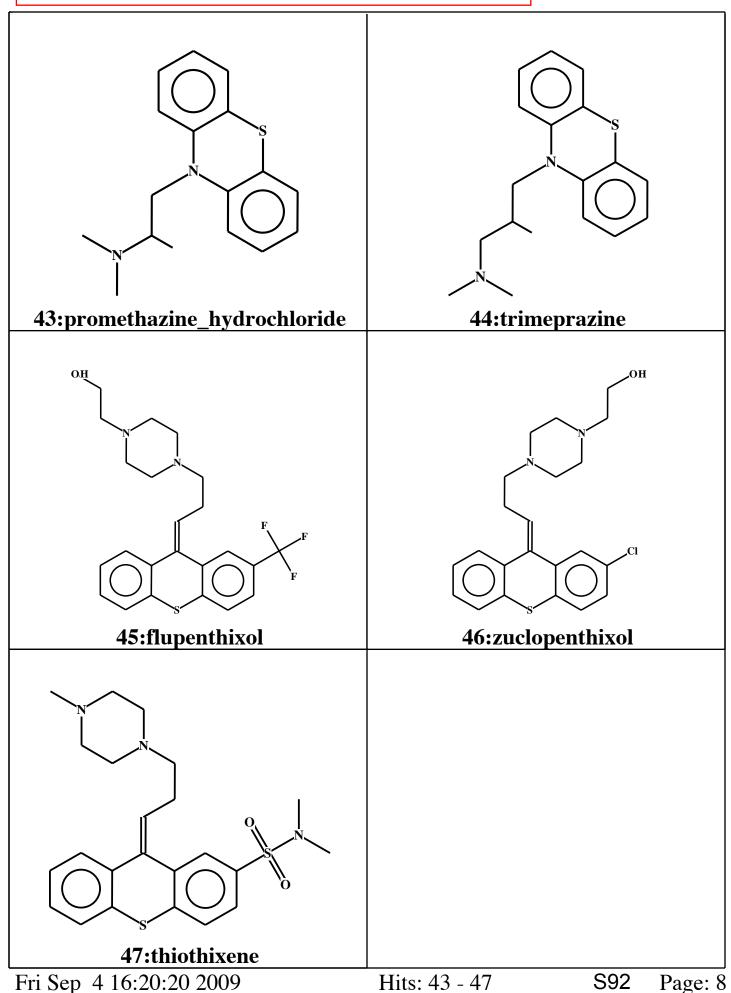
S90 Page: 6

## NH2 -OH NH OH OH -OH NH2 37:carbidopa 38:methyldopa 39:cortisone acetate 40:amitriptyline 41:cimetidine 42:nortriptyline

Fri Sep 4 16:20:20 2009

Hits: 37 - 42

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