

c1 ([S@E]) (NC (NCCC) =O) (=O) =O) ccc (C) cc1 TOLBUTAMIDE
c1 ([S] (Nc2sc ([CH] (C) C) nn2) (=O) =O) ccc (N) cc1 GLYPROTHIAZOL
c1 ([S] (NC (NCCC) =O) (=O) =O) ccc (Cl) cc1 CHLORPROPAMIDE
c1 (CCNC (NC (N) =N) =N) cccccc1 PHENFORMIN
c1 ([S@E]) (NC (NCCCC) =O) (=O) =O) ccc (N) cc1 CARBUTAMIDE
c1 ([S@E]) (Nc2ncc (OCCOC) cn2) (=O) =O) cccccc1 GLYMIDINE
c12c (ccc (c1) [S] (NC (N[CH] 1CCCCC1) =O) (=O) =O) CCC2 GLYHEXAMIDE
c1 (sc ([C] (C) (C) C) nn1) N[S] (c1ccc (N) cc1) (=O) =O GLYBUTHIAZOL
c1 ([S] (NC (N[CH] 2CCCCC2) =O) (=O) =O) cc (c (C) cc1) N METAHEXAMIDE
c1 ([S] (NC (N[N] 2CCCC2) =O) (=O) =O) ccc (Cl) cc1 GLYCLOPYRAMIDE
C (NC (N) =N) ([N] (C) C) =N METFORMIN
c1 ([S] (NC (N[CH] 2CCCCC2) =O) (=O) =O) ccc (C) cc1 GLYCYCLAMIDE
N (C (NC (N) =N) =N) CCCC BUFORMIN
c1 ([S] (NC (N[CH] 2CCCCC2) =O) (=O) =O) ccc (C (C) =O) cc1 ACETOHEXAMIDE
c1 ([S] (NC (N[CH] 2CCCC2) =O) (=O) =O) ccc (C) cc1 TOLPENTAMIDE
c1 ([S] (NC (N[CH] 2CCCCC2) =O) (=O) =O) ccc (C) cc1 HEPTOLAMIDE
c1 ([S] (NC (N[CH] 2CCCCCCC2) =O) (=O) =O) ccc (C) cc1 GLYOCTAMIDE
c1 ([S] (NC (N[N] 2CCCCC2) =O) (=O) =O) ccc (C) cc1 TOLAZAMIDE
c1 ([S] (NC (N[N] 2CCCCC2) =O) (=O) =O) ccc (Cl) cc1 GLYPINAMIDE
c1 (sc ([C] (C) (C) C) nn1) N[S] (c1ccccc1) (=O) =O GLYBUZOLE
c1 (c (ccc (c1) OC) OC) [CH] ([CH] (N[C] (C) (C) C) O) BUTOXAMINE
c12c (ccc (c1) [S] (NC (N[N] 1CCCCC1) =O) (=O) =O) CCC2 GLIDAZAMIDE
c1 ([S] (Nc2sc (C[CH] (C) C) nn2) (=O) =O) ccc (OC) cc1 GLYSOBUZOLE
c1 ([S] (NC (N[CH] 2CCCCC2) =O) (=O) =O) ccc (SC) cc1 THIOHEXAMIDE
C (CCCNC (N) =N) (N) =O TIFORMIN
c1 (ccc ([N] (C) C) cc1) NC (N[S] (c1ccc (Cl) cc1) (=O) =O) =O GLYPARAMIDE
c1 ([S@E]) (NC ([N] 2CCCC2) =O) (=O) =O) ccc (C) cc1 TOLPYRRAMIDE
c1 ([S] (Nc2sc (CCCC) nn2) (=O) =O) ccc (Cl) cc1 BUTADIAZAMIDE
c1 (c (ccc (c1) Cl) OC) C (NCCc1ccc ([S] (NC (N[CH] 2CCCCC2) =O) (=O) =O) cc1) =O
GLYBURIDE
c1 ([S] (=O) (=O) [O-]) c (ccc (c1) O) O. c1 ([S] (=O) (=O) [O-]) c (ccc (c1) O) O. [Ca+2]
CALCIUM DOBESILATE
C1 [CH] 2 [CH] (C [N] 1NC (N[S] (c1ccc (C) cc1) (=O) =O) =O) CCC2 GLICLAZIDE
c1 (c (ccc (c1) Cl) OC) NC (Cc1ccc ([S] (Nc2ncc (C[CH] (C) C) cn2) (=O) =O) cc1) =O
GLICETANILE
c1 (ccc (CCNC (c2noc (c2) C) =O) cc1) [S] (NC (N[CH] 1CCCCC1) =O) (=O) =O
GLISOLAMIDE
c1 (ccc (CCNC (c2noc (c2) C) =O) cc1) [S] (NC (N[N] 1CCCCC1) =O) (=O) =O
GLISOXEPIDE
C1 ([N] ([CH] 2CCC=CC2) CC [N] 1 [S] (c1ccc (CCNC (CCC) =O) cc1) (=O) =O) =N
GLIBUTIMINE
C1 [C] 2 ([C] ([C@H] ([C@@H] ([C@@H] 2O) NC (N[S@E]) (c2ccc (C) cc2) (=O) =O) =O) (C1)) (C)
C) C GLIBORNURIDE
c1 (ccc (CCNC (c2cnc (C) cn2) =O) cc1) [S] (NC (N[CH] 1CCCCC1) =O) (=O) =O GLIPIZIDE
c1 (c (ccccc1) OC) C (NCCc1ccc ([S@E]) (NC (N[CH] 2CCCC2) =O) (=O) =O) cc1) =O GLIPENTIDE
c12c (C ([N] (CCC3ccc ([S] (NC (N[CH] 4CCCCC4) =O) (=O) =O) cc3) C ([C] 1 (C) C) =O) =O) cc
(OC) cc2 GLIQUIDONE
c1 (c (ccc (c1) F) OC) [C@H] (NC (Cc1ccc ([S] (Nc2ncc (C[CH] (C) C) cn2) (=O) =O) cc1) =O) (C)
GLIFLUMIDE
c12c ([n] (CC) nc2C) ncc (c1OCC [CH] (C) C) C (NCCc1ccc ([S] (NC (N[CH] 2CCCCC2) =O) (=O)
=O) cc1) =O GLICARAMIDE
C (/NCCCC) (NC (N) =N) =N/CC ETOFORMIN

C1[CH]2[CH](C[CH]1C=C2)CNC(N[S]([N]1CC[CH](CCNC(c2c(nccc2)OC)=O)CC1)(=O)=O)=O GLIAMILIDE
c1([S](NC(N[CH]2CC[CH](C)CC2)=O)(=O)=O)ccc(CCNC([N](c2ccccc2)C)=O)cc1
GLISAMURIDE
c1(c(ccc(c1)C1)OC)C(NCCc1ccc([S](NC(NC)=O)(=O)=O)cc1)=O GLICONDAMIDE
c1(cc(c(N)cc1)C#N)[CH](CN[CH](C)C)O CIMATEROL
c1(c(ccc(c1)C1)OC)C(NCCc1ccc(C(O)=O)cc1)=O MEGLITINIDE
C([N]1CCCC1)(/N=C1\ [N](CCC1)C)=N\c1cccc1 PIROGLIRIDE
c1([CH](c2ccccc2)CC=2NCCN2)cccc1 MIDAGLIZOLE
C1[C](O1)(CCCCCCCCCCCC)C(O)=O PALMOXIRIC ACID
C1[C](O1)(CCCCCCCCCCCC)C(OC)=O METHYL PALMOXIRATE
c12c(C[N](C1=O)C(NCCc1ccc([S@@](NC(N[CH]3CCCC3)=O)(=O)=O)cc1)=O)cccc2
GLISINDAMIDE
C1[N]([C@@H]([C@H]([C@@H](O)[C@H]1O)O)CO)CCO MIGLITOL
C([CH]1SC(=O)NC1=O)c1ccc(OC[C]2(CCCC2)C)cc1 CIGLITAZONE
C([N]1CCOCC1)(/N=C1\ [N](CCC1)C)=N\c1cccc1 LINOGLIRIDE
c1(c2c(cccc2)[nH]c1)C[C@H]1NC([C@H](Cc2ccc(O)cc2)NC([C@@H]([N@](C([C@H](C
c2ccccc2)NC([C@H](NC([C@H](NC1=O)CCCN)=O)[CH](C)C)=O)C)C)=O)=O
SEGLITIDE
C1(/C([N](CC(O)=O)C(S1)=S)=O)=C\C(=C/c1cccc1)C EPALRESTAT
C1[C](O1)(CCCCc1ccc(C1)cc1)C(O)=O CLOMOL
c12[C]3(c4c(ccc(c4)F)c1ccc(c2)F)C(NC(N3)=O)=O IMIRESTAT
C1([N](CC(C)=C1CC)C(NCCc1ccc([S](NC(N[CH]2CC[CH](C)CC2)=O)(=O)=O)cc1)=O)=O
GLIMEPIRIDE
C([CH]1C(NC(S1)=O)=O)c1ccc(OCCc2ccc(CC)cn2)cc1 PIOGLITAZONE
c12c(C[N](C1)NC=1NCCN1)cccc2F ISAGLIDOLE
c12c(c([n](Cc3nc4cc([C@@](F)(F)F)ccc4s3)nc1CC(O)=O)=O)cccc2
ZOPOLRESTAT
C(NN)(N)=N PIMAGEDINE
C1[C@@H]([C@@H]([C@@H](O)[C@@H]([C@]1(CO)O)O)O)N[CH](CO)CO VOGLIBOSE
c12c(CC[C](O1)(COc1ccc(C[CH]3C(NC(S3)=O)=O)cc1)C)c(c(O)c(c2C)C)C
TROGLITAZONE
c12c(O[CH](Cc3ccccc3)CC2)ccc(c1)C[CH]1C([N]([Na])C(S1)=O)=O
ENGLITAZONE
C=1([C]2([N]3c4c(cccc4C2)CC3)CCC)NCCN1 DERIGLIDOLE
C([N]1[C@@H]([C@H]([C@H](O)[C@H](C1)O)O)CO)[C@H]1O[C@@H]([C@H](O)[C@H]([C
@@H]1O)O)OC CAMIGLIBOSE
c1(c2ccccc2)nc(CCC(c2ccc(C[CH]3C([N]([Na])C(S3)=O)=O)cc2)=O)c(o1)C
DARGLITAZONE
c1(CNC(/N=C(/N[P](=O)([O-])[O-])N)=N)cccc1.[Na+].[Na+] BENFOSFORMIN
c12c(c[nH]c1cccc2)C[C@@H](C(N[C@H](C(N[C@H](C(N[C@@H](Cc1ccccc1)C(N)=O)=O
)CC(O)=O)=O)CCSC)=O)N TETRAGASTRIN
C([C@@H](C(NCCCCCCCC)=O)N)S[V](SC[C@@H](C(NCCCCCCCC)=O)N)=O NAGLIVAN
C([N]1CCCC1)[CH](CO\N=C(\c1cccn1)C1)O BIMOCLOMOL
c1([CH](NCc2cc(ccc2)[C](F)(F)F)C(O)=O)cccc1 PHENYLGLYCINE-BASED INSULIN
SENSITIZER
c12c(C[C@H](NC([C@@H](NC([C@@H](NC([C@H](Cc3ccccc3)NC([C@@H](NC([C@@H](NC
([C@@H](NC([C@@H](NC([C@@H](NC([C@@H](NC([C@@H](NC([C@H](Cc3ccc
(O)cc3)NC([C@@H](NC([C@@H](NC([C@H](Cc3ccc(O)cc3)NC([C@@H](NC([C@@H](NC([
C@@H](NC([C@H](Cc3ccccc3)NC([C@@H](NC(CNC([C@@H](NC([C@@H](NC([C@H](Cc3c
[nH]cn3)N)=O)CO)=O)CCC(N)=O)=O)=O)[C@@H](C)O)=O)=O)[C@@H](C)O)=O)CO)=O)CC(
O)=O)=O)=O)CO)=O)CCCCN)=O)=O)C[CH](C)C)=O)CC(O)=O)=O)CO)=O)CCCNC(N)=N)=O)
CCCNC(N)=N)=O)C)=O)CCC(N)=O)=O)CC(O)=O)=O)=O)[CH](C)C)=O)CCC(N)=O)=O)C(N

[illegible]

C ([C@H] 1 [N] (C ([C@@H] (NC ([C@@H] (NC ([C@H] 2 [N] (CCC2) C (CNC ([C@H] (Cc2ccccc2) NC
([C@@H] (NC ([C@@H] (NC ([C@@H] (NC ([C@H] (Cc2c[nH]cn2) NC ([C@@H] (NC ([
C@@H] (NC ([C@H] (Cc2ccccc2) NC ([C@@H] (NC ([C@@H] (NC ([C@@H] (NC ([C@@H] (NC ([C@@H]
(NC ([C@@H] (NC ([C@@H] (NC ([C@H] 2NC ([C@@H] (NC ([C@@H] (NC ([C@@H] (NC ([C@@H] (NC
([C@H] (CSSC2) NC ([C@H] (CCCCN) N)=O)=O CC(N)=O)=O [C@H] (C) O)=O) C)=O [C@H] (
C) O)=O)=O) C)=O [C@@H] (C) O)=O CCC(N)=O)=O CCCNC(N)=N)=O C[CH] (C) C)=O) C)=O
CC(N)=O)=O)=O) C[CH] (C) C)=O [C@@H] (C) C)=O)=O CO)=O CO)=O CC(N)=O)=O CC(N)=
O)=O)=O)=O) [C@H] (CC) C)=O C[CH] (C) C)=O CCC1) ([N] 1 [C@@H] (CCC1) C(N[C@H] (C
(N[C@H] (C(N[C@H] (C(NCC(N[C@H] (C(N[C@H] (C(N[C@H] (C(N[C@@H] (Cc1ccc(O)cc1) C(
N)=O)=O [C@@H] (C) O)=O CC(N)=O)=O CO)=O)=O [C@@H] (C) C)=O CC(N)=O)=O [C@H] (
C) O)=O)=O MCMC00010668

c12c(CC[C@](O1)(COc1ccc(C[C@H]3C(NC(S3)=O)=O)cc1)C)c(c(O)c(c2C)C)C
) S,S)-TROGLITAZONE

c12c(CC[C@](O1)(COc1ccc(C[C@@H]3C(NC(S3)=O)=O)cc1)C)c(c(O)c(c2C)C)C
) S,R)-TROGLITAZONE

c12c(CC[C](O1)(COc1ccc(C[C@H]3C(NC(S3)=O)=O)cc1)C)c(c(O)c(c2C)C)C
) R,S)-TROGLITAZONE

c12c(CC[C](O1)(COc1ccc(C[C@@H]3C(NC(S3)=O)=O)cc1)C)c(c(O)c(c2C)C)C
) R,R)-TROGLITAZONE

S1C(NC([CH]1Cc1ccc(cc1)OCc1nc2cccc2c([n]1C)=O)=O)MCMC00010890

c1(ccccc1)C[C@@H](Oc1c(cc(cc1C)c1c2c(sc(c2C)C)c(c2c1cccc2)Br)C)C(=O)O
MCMC00010916

SI_table 2: Full list of all descriptors used in modeling process and their redundancy within the selected best filters.

Descriptor name	Redundancy
GCUT_SLOGP_0	24
a_ICM	16
PEOE_VSA+4	12
SMR_VSA1	10
logS	9
nmol	9
lip_druglike	9
chi1_C	8
GCUT_PEOE_0	8
opr_leadlike	7
Q_VSA_FPOS	7
SMR_VSA3	7
a_don	6
a_hyd	6
BCUT_SMR_3	4
chi1v_C	4
chiral	4
GCUT_PEOE_3	4
GCUT_SLOGP_3	3
PEOE_VSA_NEG	3
SMR_VSA6	2
a_nH	2
a_nN	2
a_nO	2
b_single	2
chi0v_C	2
chiral_u	2
PEOE_PC+	2
PEOE_VSA+5	2
PEOE_VSA_FNEG	1
a_IC	1
BCUT_SLOGP_3	1
bpol	1
b_count	1
b_double	1
chi0_C	1
density	1
GCUT_PEOE_1	1
GCUT_SMR_3	1
lip_violation	0
logP(o/w)	0
Q_VSA_FNEG	0

SI_table 2: Full list of all descriptors used in modeling process and their redundancy within the selected best filters.

SlogP_VSA8	0
SMR_VSA0	0
vsa_hyd	0
apol	0
a_acc	0
a_acid	0
a_aro	0
a_base	0
a_count	0
a_heavy	0
a_nB	0
a_nBr	0
a_nC	0
a_nCl	0
a_nF	0
a_nI	0
a_nP	0
a_nS	0
balabanJ	0
BCUT_PEOE_0	0
BCUT_PEOE_1	0
BCUT_PEOE_2	0
BCUT_PEOE_3	0
BCUT_SLOGP_0	0
BCUT_SLOGP_1	0
BCUT_SLOGP_2	0
BCUT_SMR_0	0
BCUT_SMR_1	0
BCUT_SMR_2	0
b_1rotN	0
b_1rotR	0
b_ar	0
b_heavy	0
b_rotN	0
b_rotR	0
b_triple	0
chi0	0
chi0v	0
chi1	0
chi1v	0

SI_table 2: Full list of all descriptors used in modeling process and their redundancy within the selected best filters.

diameter	0
FCharge	0
GCUT_PEOE_2	0
GCUT_SLOGP_1	0
GCUT_SLOGP_2	0
GCUT_SMR_0	0
GCUT_SMR_1	0
GCUT_SMR_2	0
Kier1	0
Kier2	0
Kier3	0
KierA1	0
KierA2	0
KierA3	0
KierFlex	0
lip_acc	0
lip_don	0
mr	0
mutagenic	0
opr_brigid	0
opr_nring	0
opr_nrot	0
opr_violation	0
PC+	0
PC-	0
PEOE_PC-	0
PEOE_RPC+	0
PEOE_RPC-	0
PEOE_VSA+0	0
PEOE_VSA+1	0
PEOE_VSA+2	0
PEOE_VSA+3	0
PEOE_VSA+6	0
PEOE_VSA-0	0
PEOE_VSA-1	0
PEOE_VSA-2	0
PEOE_VSA-3	0
PEOE_VSA-4	0
PEOE_VSA-5	0
PEOE_VSA-6	0

SI_table 2: Full list of all descriptors used in modeling process and their redundancy within the selected best filters.

PEOE_VSA_FHYD	0
PEOE_VSA_FPNEG	0
PEOE_VSA_FPOL	0
PEOE_VSA_FPOS	0
PEOE_VSA_FPPOS	0
PEOE_VSA_HYD	0
PEOE_VSA_PNEG	0
PEOE_VSA_POL	0
PEOE_VSA_POS	0
PEOE_VSA_PPOS	0
petitjean	0
petitjeanSC	0
Q_PC+	0
Q_PC-	0
Q_RPC+	0
Q_RPC-	0
Q_VSA_FHYD	0
Q_VSA_FPNEG	0
Q_VSA_FPOL	0
Q_VSA_FPPOS	0
Q_VSA_HYD	0
Q_VSA_NEG	0
Q_VSA_PNEG	0
Q_VSA_POL	0
Q_VSA_POS	0
Q_VSA_PPOS	0
radius	0
reactive	0
rings	0
RPC+	0
RPC-	0
rsynth	0
SlogP	0
SlogP_VSA0	0
SlogP_VSA1	0
SlogP_VSA2	0
SlogP_VSA3	0
SlogP_VSA4	0
SlogP_VSA5	0
SlogP_VSA6	0

SI_table 2: Full list of all descriptors used in modeling process and their redundancy within the selected best filters.

SlogP_VSA7	0
SlogP_VSA9	0
SMR	0
SMR_VSA2	0
SMR_VSA4	0
SMR_VSA5	0
SMR_VSA7	0
TPSA	0
VAdjEq	0
VAdjMa	0
VDistEq	0
VDistMa	0
vdw_area	0
vdw_vol	0
vsa_acc	0
vsa_acid	0
vsa_base	0
vsa_don	0
vsa_other	0
vsa_pol	0
Weight	0
weinerPath	0
weinerPol	0
zagreb	0