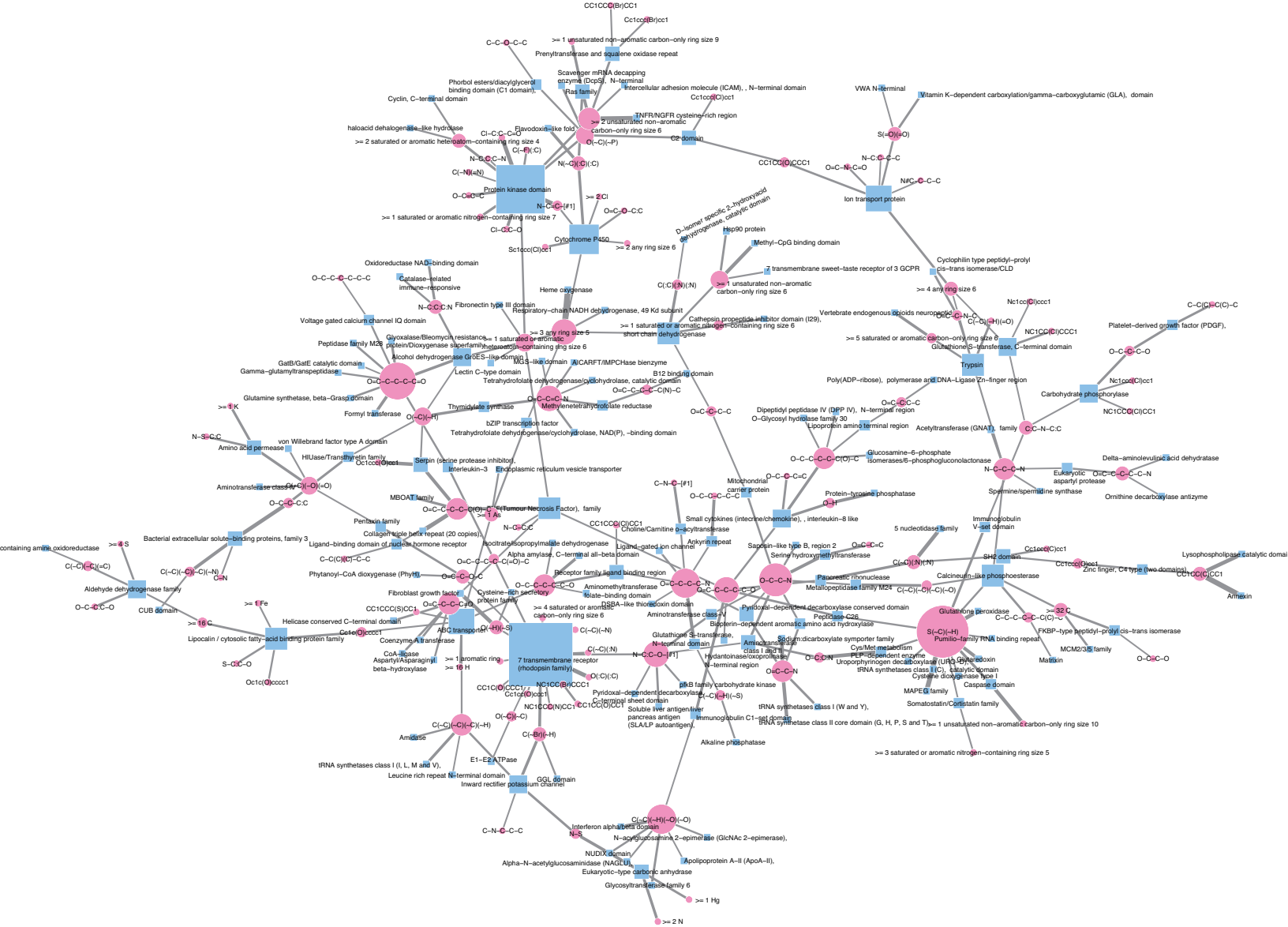
CC1CCC(Br)CC1
Cc1ccc(Br)ccc1
>= 1 unsaturated non-aromatic carbon-only ring size 9
C-C-O-C-C
Prenyltransferase and squalene oxidase repeat
Scavenger mRNA decapping enzyme4 (DcpS), N-terminal
Phorbol esters/diacylglycerol binding domain (C1 domain),
Ras family
Intercellular adhesion molecule (ICAM), , N-terminal domain
Cyclin, C-terminal domain
Cc1ccc(Cl)ccc1
TNFR/NGFR cysteine-rich region
>= 2 unsaturated non-aromatic carbon-only ring size 6
Flavodoxin-like fold
haloacid dehalogenase-like hydrolase
>= 2 saturated or aromatic heteroatom-containing ring size 4
Cl-C-C-C-C
O-C(-C)(-P)
C2 domain
C(-F)(:C)
N-C(:C)-C-N
N(-C)(:C)(:C)
C(-N)(=N)
Protein kinase domain
O-C-C-C
>= 2 Cl
N-C-C-[#1]
>= 1 saturated or aromatic nitrogen-containing ring size 7
O=C-O-C:C
Cl-C:C-O
Cytochrome P450
Sc1ccc(Cl)cc1
>= 2 any ring size 6
CC1CC(O)CCC1
VWA N-terminal
Vitamin K-dependent carboxylation/gamma-carboxyglutamic (GLA), domain
S(=O)(=O)
N-C-C-C-C
O=C-N-C=O
N#C-C-C-C
Ion transport protein
O-isomer specific 2-hydroxyacid dehydrogenase, catalytic domain
Hsp90 protein
Methyl-CpG binding domain
7 transmembrane sweet-taste receptor of 3 GCPR
C(:C):N):N)
>= 1 unsaturated non-aromatic carbon-only ring size 6
Cyclophilin type peptidyl-prolyl cis-trans isomerase/CLD
>= 4 any ring size 6
Oxidoreductase NAD-binding domain
O-C-C-O-C-C-C-C
Catalase-related immune-responsive
N-C-C:C:N
Fibronectin type III domain
Heme oxygenase
Respiratory-chain NADH dehydrogenase, 49 Kd subunit
Cathepsin propeptide inhibitor domain (I29), short chain dehydrogenase
>= 1 saturated or aromatic nitrogen-containing ring size 6
Vertebrate endogenous opioids neuropeptide
C-C-C-N-C
C(-O)(-H)=O)
Nc1ccc(Cl)ccc1
Voltage gated calcium channel IQ domain
>= 3 any ring size 5
>= 1 saturated or aromatic heteroatom-containing ring size 6
Glyoxalase/Bleomycin resistance protein/Dioxygenase superfamily
Peptidase family M28
Alcohol dehydrogenase GroES-like domain
>= 5 saturated or aromatic carbon-only ring size 6
NC1CCC(Cl)CCC1
Glutathione S-transferase, C-terminal domain
Platelet-derived growth factor (PDGF),
C-C(C)=C(C)-C
O-C-C-C-O
GatB/GatE catalytic domain
Gamma-glutamyltranspeptidase
MGS-like domain
AICARFT/IMPCHase bienzyme
Trypsin
Lectin C-type domain
Tetrahydrofolate dehydrogenase/cyclohydrolase, catalytic domain
B12 binding domain
Poly(ADP-ribose), polymerase and DNA-Ligase Zn-finger region
O=C-C-C-C-C-C-O
Glutamine synthetase, beta-Grasp domain
>= 1 K
Thymidylate synthase
Methylenetetrahydrofolate reductase
O=C-C-C-N
O=C-C-C-C-C(-N)-C
Carbohydrate phosphorylase
Nc1ccc(Cl)cc1
NC1CCC(Cl)CC1
Formyl transferase
O(-C)(-H)
bZIP transcription factor
Tetrahydrofolate dehydrogenase/cyclohydrolase, NAD(P), -binding domain
Dipeptidyl peptidase IV (DPP IV), N-terminal region
O-Glycosyl hydrolase family 30
Lipoprotein amino terminal region
Acetyltransferase, (GNAT) family
C-C-N-C-C
N-S-C:C
Amino acid permease
von Willebrand factor type A domain
HIUase/Transthyretin family
Oc1ccc(O)cc1
Serpin (serine protease inhibitor),
Interleukin-3
Endoplasmic reticulum vesicle transporter
Glucosamine-6-phosphate isomerases/6-phosphogluconolactonase
O-C-C-C-C-C(O)-C
Delta-aminolevulinic acid dehydratase
Eukaryotic aspartyl protease
Aminotransferase class-I
C(-C)(-N)(=O)
O-C-C-C:C
MBOAT family
Tumour Necrosis Factor), family
>= 1 As
Mitochondrial carrier protein
C-N-C-[#1]
O-C-C-C-C
Protein-tyrosine phosphatase
O-H
Spermine/spermidine synthase
Ornithine decarboxylase antizyme
Pentaxin family
Collagen triple helix repeat (20 copies)
Ligand-binding domain of nuclear hormone receptor
Isocitrate/isopropylmalate dehydrogenase
Alpha amylase, C-terminal all-beta domain
CC1CCC(Cl)CC1
Choline/Carnitine o-acyltransferase
Small cytokines (intecrine/chemokine), , interleukin-8 like
Ankyrin repeat
Ligand-gated ion channel
Immunoglobulin V-set domain
5 nucleotidase family
Bacterial extracellular solute-binding proteins, family 3
>= 4 S
containing amine oxidoreductase
C(-O)(-C)(=O)
Aldehyde dehydrogenase family
O-C-C:C-O
CUB domain
C(-C)(-C)(-C)-N
C=N
C-C(C)(C)-C-C
Phytanoyl-CoA dioxygenase (PhyH),
Receptor family ligand binding region
Aminomethyltransferase folate-binding domain
DSBA-like thioredoxin domain
Aminotransferase class-V
Sarcosin-like type B, region 2
Serine hydroxymethyltransferase
Pancreatic ribonuclease
Metallopeptidase family M24
SH2 domain
Calcineurin-like phosphoesterase
Lysophospholipase catalytic domai
Zinc finger, C4 type (two domains)
CC1CC(C)CC1
Annexin
Fibroblast growth factor
Cysteine-rich secretory protein
Pyridoxal-dependent decarboxylase conserved domain
Peptidase C26
Bioplerin-dependent aromatic amino acid hydroxylase
Glutathione peroxidase
>= 32 C
C-C-C-C-C-C(O)-C
FKBP-type peptidyl-prolyl cis-trans isomerase
MCM2/3/5 family
Matrixin
O-C-C-O
Helicase conserved C-terminal domain
>= 1 Fe
>= 16 C
Lipocalin / cytosolic fatty-acid binding protein family
S-C:C-O
Oc1c(O)cccc1
ABC transporter
>= 4 saturated or aromatic carbon-only ring size 6
Coenzyme A transferase
CoA-ligase
Aspartyl/Asparaginyl beta-hydroxylase
>= 1 aromatic ring
>= 16 H
7 transmembrane receptor (rhodopsin family),
Glutathione S-transferase, N-terminal domain
Aminotransferase class I and II
Hydantoinase/oxoprolinase N-terminal region
Sodium:dicarboxylate symporter family
Cys/Met metabolism PLP-dependent enzyme
Uroporphyrinogen decarboxylase (URO-D)
tRNA synthetases class II (C), catalytic domain
Cysteine dioxygenase type I
Caspase domain
MAPEG family
Somatostatin/Corstatin family
Glutathione S-transferase, C-terminal domain
Pumilio-family RNA binding repeat
tRNA synthetases class I (W and Y),
tRNA synthetase class II core domain (G, H, P, S and T)
>= 1 unsaturated non-aromatic carbon-only ring size 10
>= 3 saturated or aromatic nitrogen-containing ring size 5
pkB family carbohydrate kinase
Pyridoxal-dependent decarboxylase, C-terminal sheet domain
Soluble iver antigen/liver pancreas antigen (SLA/LP autoantigen),
Immunoglobulin C1-set domain
Alkaline phosphatase
Amidase
E1-E2 ATPase
tRNA synthetases class I (I, L, M and V),
Leucine rich repeat N-terminal domain
Inward rectifier potassium channel
GGL domain
Interferon alpha/beta domain
N-S
C-N-C-C-C
C(-C)(-H)(-O)(-O)
N-acylglucosamine 2-epimerase (GlcNAc 2-epimerase),
NUDIX domain
Alpha-N-acetylglucosaminidase (NAGLU)
Apolipoprotein A-II (ApoA-II),
Eukaryotic-type carbonic anhydrase
Glycosyltransferase family 6
>= 1 Hg
>= 2 N