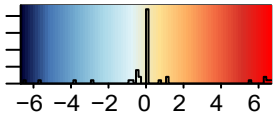

COBALSYN−PWY: adenosylcobalamin salvage from cobinamide I
PWY−6969: TCA cycle V (2−oxoglutarate:ferredoxin oxidoreductase)
PWY0−1298: superpathway of pyrimidine deoxyribonucleosides degradation
PWY0−862: (5Z)−dodecenoate biosynthesis I
PWY−6606: guanosine nucleotides degradation II
P164−PWY: purine nucleobases degradation I (anaerobic)
PWY−6608: guanosine nucleotides degradation III
PWY−241: C4 photosynthetic carbon assimilation cycle, NADP−ME type
PWY−7117: C4 photosynthetic carbon assimilation cycle, PEPCK type
PWY0−1586: peptidoglycan maturation (meso−diaminopimelate containing)
PWY−6527: stachyose degradation
PWY−6545: pyrimidine deoxyribonucleotides de novo biosynthesis III
THRESYN−PWY: superpathway of L−threonine biosynthesis
PWY−7383: anaerobic energy metabolism (invertebrates, cytosol)
PWY0−162: superpathway of pyrimidine ribonucleotides de novo biosynthes
FUCCAT−PWY: fucose degradation
PYRIDOXSYN−PWY: pyridoxal 5'−phosphate biosynthesis I
HEME−BIOSYNTHESIS−II: heme b biosynthesis I (aerobic)
PWY−7282: 4−amino−2−methyl−5−diphosphomethylpyrimidine biosynthesis
PWY0−1241: ADP−L−glycero−β−D−manno−heptose biosynthesis
PWY0−845: superpathway of pyridoxal 5'−phosphate biosynthesis and salva