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| | COBALSYN-PWY: adenosylcobalamin salvage from cobinamide |
| | PWY-6969: TCA cycle V (2-oxoglutarate:ferredoxin oxidoreducta |
| | PWY0-1298: superpathway of pyrimidine deoxyribonucleosides o |
| | 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-M |
| | PWY-7117: C4 photosynthetic carbon assimilation cycle, PEPCK |
| | PWY0-1586: peptidoglycan maturation (meso-diaminopimelate o |
| | PWY-6527: stachyose degradation |
| | PWY-6545: pyrimidine deoxyribonucleotides de novo biosynthes |
| | THRESYN-PWY: superpathway of L-threonine biosynthesis |
| | PWY-7383: anaerobic energy metabolism (invertebrates, cytosol |
| | PWY0-162: superpathway of pyrimidine ribonucleotides de novo |
| | 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 b |
| | PWY0-1241: ADP-L-glycero-β-D-manno-heptose biosyn |
| | PWY0-845: superpathway of pyridoxal 5'-phosphate biosynthesi |