diaminopimelate decarboxylase [EC:4.1.1.20] diaminopimelate epimerase [EC:5.1.1.7] LL-diaminopimelate aminotransferase [EC:2.6.1.83] 4-hydroxy-tetrahydrodipicolinate reductase [EC:1.17.1.8] para-aminobenzoate synthetase [EC:2.6.1.85] avdrolysing) (EC:6) 35 1 rtuin 4 [EC:7.3-dibyeroneopterin aldolase/epimerase oxygenase [EC:4.1.2.25 5.1.99.8 1.13.11.81] urate oxida<mark>se [</mark>EC:1.7.3.3] proteinyseigsterieneyaldystecraerkoo<mark>xyet</mark>hyttrydessterrasse [EC:2.1.1.100] alkaline phosphatase D [EC:2.7/1/23]

xanthine dehydrogenase/pxides/clipsing phosphatase D [EC:2.7/1/23]

nicotinamide mononucleotide die hybritalisterase [EC:2.7/1/23]

\$2231.1.2.3 allantoate deiminase [EC:3.5.3.9] psrefix)4 perrode prepe<mark>iptas</mark>e se (ES:48.2442824-]] dibydrgerntesenEG (EC?237.1.39) nicotinate phosphoribosylfolyglytamate synthase [EC:6.3.2.17]

nicotinate phosphoribosylfolyglytamate synthase ditrans,polycis-polyprenyl diphosphate synthase (EC:2.5.1.87) anthranilate phosphoribosyltransferase [EC:2.4.2.18] (S)-ureidoglycine aminohydrolase [EC:3.5.3.26] acylpyruvate hydrolase [EC:3.7.1.5] protein farnesyltransferase/geranylderanyltransferase type-1 subunit alpha [EC:2.5.1.58 2.5.1.59] isopentenyl phosphate kinase [EC:2.7.4.26]

isopentenyl phosphate argininosuccinate lyase [EC:4.3.2.1] formyltetrahydrofolate deft fumarylacetoacetase [EC:3.7.1.2] ureidoglycolate amidohydrolase [EC:3.5.1.116] phosphomevalonate kinase (EC:1.3.1.83 1.3.1.111) Panestyranorers (a) 13 1,14 cm as (FC) 2.1.3 5 Administration (a) 1.3 Administration (a) 1. urease [EC:3.5.1.5] armacic desulfoglucosinolar sufficience (ECS.1.1.10) and the suffi coproporphyrinogen III oxidase [EC:1.3.3.3] alpha-1,3-glucosyltransferase [EC:2.4.1.265]
glucosamine-phosphate N-ace
zeaxanthin epoxida phosphatidylinositol phospholipase C, delta/[EC:3.1.4.11] beta-fructofurandshtase [EC:3.1.3.7 3.4.3.57]

beta-fructofurandshtase project state of the second state o 3'(2'), 5'-bisphosphate nucleotidase / inositol polyphosphate 1-phosphatase [EC:3.1.37] phosphatidylinositol-3,4,5-trisphosphate 3-phosphatase and dual-specificity protein phosphatase PTEN [EC:3.1.3.16 3.1.3.48 3.1.3.67] chyose synthetase [EC:2/2] GDPmannose 4,6-dehydratase [EC:4.2.1.47] lactosylceramide 4-ahrihaagada ECOS Lirans Perase [EC:2.4.1.228] 1-phosphatidylinositol-4-phosphate 5-kinase [EC:2.7.1.68] GDP-L-fucose synthase [EC:1.1.1.271] digalactosyldiacylglyce<mark>rol s</mark>ynthase [EC:2.4.1.241]
phosphatidylinositol 4-kinase A [EC:2.7.1.6/2-diacylglycerol 3-beta-galactosyltrans@pase_[F6q2iflov6645] allene oxide gyclase (fc.5.5.3996) ase [decurpnokinase [EC:2.74][43]glucose 1-phosphate uridylyltransferase [EC:2.7.7.9] phosphatidylglygerophosphatase GEP4 [EC:3.1.3.27] neutral ceramidase/[atc1345ble2a-kylosidase [EC:3.2.1.37] L-arabinokinase [EC:2.7.1.46]

Constitution of the control of the serine palmitoyltransferase [#C:2.3.1.50] sphinganine-1-phosphate aldolase [EC:4.1.2.27] dihydroceram<mark>idas</mark>e [EC:3.5.1.-] alpha-L-arabinofuranosidase [EC:3.2.1.55] ceramide kina<mark>se [</mark>EC:2.7.1.138] 1p/myo-inositol 3-kinase [EC:2.7.1.64] alpha-dioxygenase LEC:1.14.99 3-phosphate acyltransferase [EC:2.3.1.15 2.3.1.198] AF acetyitransferase [EC:2.3.1.23 2.3.1De/IJa24-sterol reductase [EC:1.3.1.72 1.3.1.-] *Stero 22-desaturase [EC:1.14.19.41] glycerol kinase [EC:2.7.1.30] Delta7-sterol 5-desaturase [EC:1.14.19.20] sterol 24-C-methyltransferase [EC:2.1.1.41] ethanolamine-phosphate The second store of Q-4algha-methyl-monooxygenase [EC:1.14.18.10] alcohol dehydrogen<mark>ase class-P [EC:1.1.1.1]</mark> (Z)-3-hexen-1-ol acety<mark>ltrans</mark>ferase [EC:2.3.1.195] vnthase 2 [EC:5.4.9676] Post Short The self of Styles wild -4 alpha-carbox ylate 3-dehydrogenase [EC:1.1.1.418]

124 methylegie (1.2.1.1.1.418)

10 peol synthyse (1.2.1.1.1.1.14.14.17)

10 squalene monooxygenase [EC:1.14.14.14.17] farnesyl-diphosphate farnesyltransferase [EC:2.5.1.21] cycloeucalenqlzgy៤<mark>៦តែមានទៀត ទៀប ទី ទ</mark>្វាក់ជាងse [EC:4.2.3.40] (+)-alpha-barbatene/beta-chamigrene/thujo<mark>psene synthase [EC:4.2.3.69 4.2.3.78 4.2.3.79]</mark>

alpha-humulene/beta-caryophyllene synthase [EC:4.2.3.104 4.2.3.57]