hypothetical protein AMSG_12403 [Thecamonas trahens ATCC 50062]

hypothetical protein CEUSTIGMA_g12406.t1 [Guillardia theta CCMP2712]

hypothetical protein GUITHDRAFT_149851 [Guillardia theta CCMP2712]

hypothetical protein CAOG_05132 [Capsaspora owczarzaki ATCC 30864]

serine incorporator 3, putative [Acanthamoeba castellanii str. Neff]

hypothetical protein PROFUN_14547 [Planoprotostelium fungivorum]

membrane protein tms1d [Thecamonas trahens ATCC 50062]

membrane protein TMS1 [Thecamonas trahens ATCC 50062]

eukaryotes | 5 leaves

cellular slime molds | 7 leaves

hypothetical protein PTSG_04461 [Salpingoeca rosetta]

hypothetical protein [Monosiga brevicollis MX1]

hypothetical protein SARC_05479 [Sphaeroforma arctica JP610]

Multiple organisms | 114 leaves

hypothetical protein CAOG_09056 [Capsaspora owczarzaki ATCC 30864]

hypothetical protein PTSG_01886 [Salpingoeca rosetta]

club-mosses | 3 leaves

club-mosses | 5 leaves

club-mosses | 3 leaves

mosses | 5 leaves

eudicots | 5 leaves

eudicots | 10 leaves

eudicots | 4 leaves

eudicots | 11 leaves

eudicots | 4 leaves

monocots | 6 leaves

monocots | 21 leaves

eudicots | 28 leaves

mosses | 3 leaves

hypothetical protein CEUSTIGMA_g4603.t1 [Chlamydomonas eustigma]

predicted protein CHLRE_09g412803v5 [Chlamydomonas reinhardtii]

probable serine incorporator [Chlamydomonas reinhardtii]

probable serine incorporator [Selaginella moellendorffii]

probable serine incorporator [Physcomitrella patens]

club-mosses | 3 leaves

monocots | 6 leaves

eudicots | 3 leaves

eudicots | 7 leaves

monocots | 6 leaves

eudicots | 3 leaves

eudicots and unknown | 14 leaves

Serine incorporator 3 [Hondaea fermentalgiana]

hypothetical protein SteCoe_7204 [Stentor coeruleus]

hypothetical protein SteCoe_8598 [Stentor coeruleus]

hypothetical protein SteCoe_27403 [Stentor coeruleus]

TMS membrane protein/tumor differentially protein [Tetrahymena thermophila SB210]

membrane protein tms1, putative [Ichthyophthirius multifiliis]

TMS membrane protein tms1, putative [Ichthyophthirius multifiliis]

hypothetical protein PPERSA_09332 [Pseudocohnilembus persalinus]

hypothetical protein [Paramecium tetraurelia strain d4-2]

hypothetical protein [Paramecium tetraurelia strain d4-2]

hypothetical protein PPERSA_05317 [Pseudocohnilembus persalinus]

hypothetical protein PPERSA_12077 [Pseudocohnilembus persalinus]

TMS membrane protein/tumor differentially protein [Tetrahymena thermophila SB210]

hypothetical protein SteCoe_32950 [Stentor coeruleus]

hypothetical protein SteCoe_4975 [Stentor coeruleus]

membrane protein [Stylonychia lemnae]

hypothetical protein FGO68_gene6558 [Halteria grandinella]

hypothetical protein FGO68_gene8497 [Halteria grandinella]

hypothetical protein FGO68_gene17333 [Halteria grandinella]

UNKNOWN [Stylonychia lemnae]

putative serine incorporator [Strobomastix strix]

hypothetical protein FDP4L_006279 [Naegleria fowleri]

predicted protein [Chlamydomonas reinhardtii]

0.5