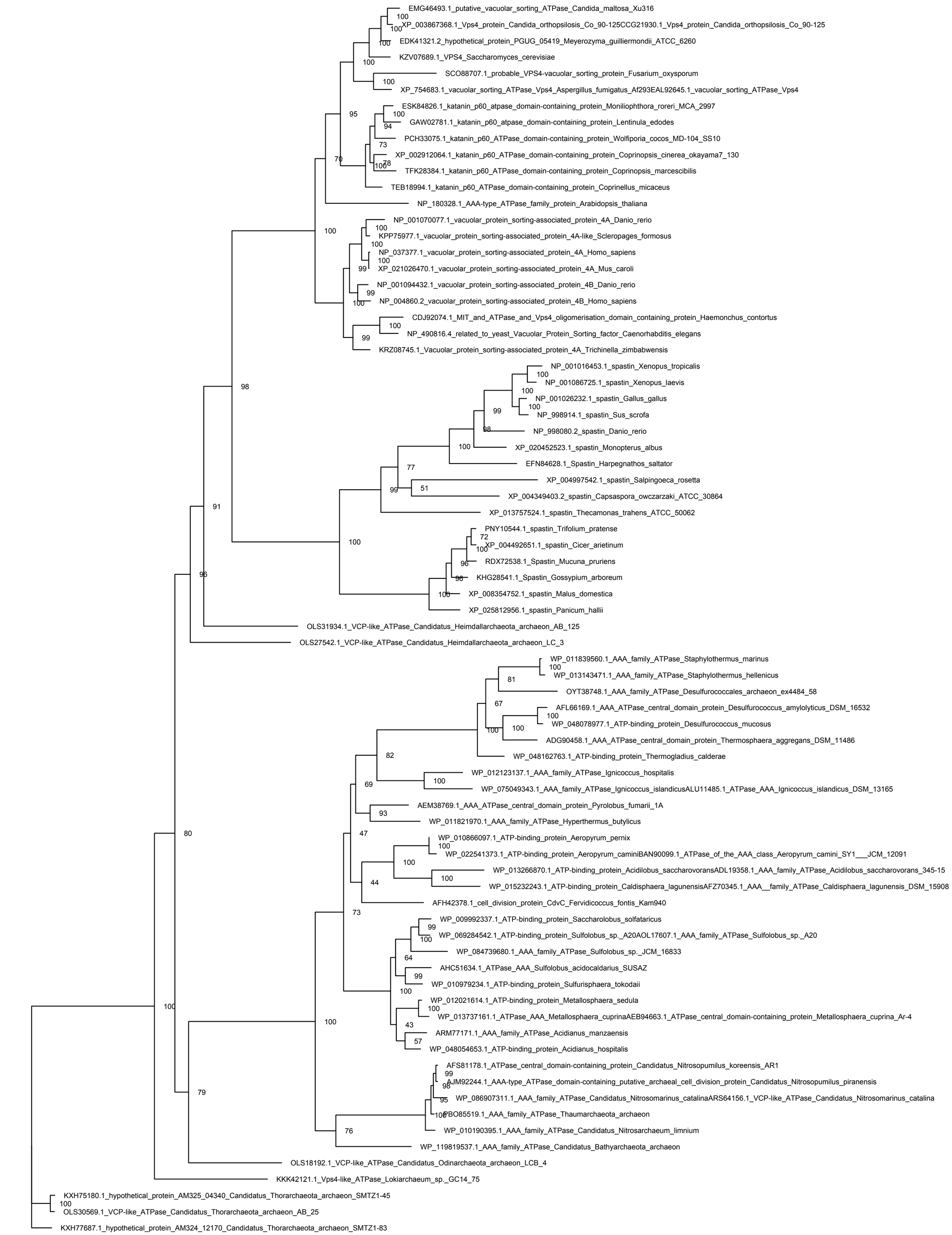
EMG46493.1_putative_vacuolar_sorting_ATPase_Candida_maltosa_Xu316
XP_003867368.1_Vps4_protein_Candida_orthopsilosis_Co_90-125CCG21930.1_Vps4_protein_Candida_orthopsilosis_Co_90-125
EDK41321.2_hypothetical_protein_PGUG_05419_Meyerozyma_guilliermondii_ATCC_6260
KZV07689.1_VPS4_Saccharomyces_cerevisiae
SCO88707.1_probable_VPS4-vacuolar_sorting_protein_Fusarium_oxysporum
XP_754683.1_vacuolar_sorting_ATPase_Vps4_Aspergillus_fumigatus_Af293EAL92645.1_vacuolar_sorting_ATPase_Vps4
ESK84826.1_katanin_p60_atpase_domain-containing_protein_Moniliophthora_roreri_MCA_2997
GAW02781.1_katanin_p60_atpase_domain-containing_protein_Lentinula_edodes
PCH33075.1_katanin_p60_ATPase_domain-containing_protein_Wolfiporia_cocos_MD-104_SS10
XP_002912064.1_katanin_p60_ATPase_domain-containing_protein_Coprinopsis_cinerea_okayama7_130
TFK28384.1_katanin_p60_ATPase_domain-containing_protein_Coprinopsis_marcescibilis
TEB18994.1_katanin_p60_ATPase_domain-containing_protein_Coprinellus_micaceus
NP_180328.1_AAA-type_ATPase_family_protein_Arabidopsis_thaliana
NP_001070077.1_vacuolar_protein_sorting-associated_protein_4A_Danio_rerio
KPP75977.1_vacuolar_protein_sorting-associated_protein_4A-like_Scleropages_formosus
NP_037377.1_vacuolar_protein_sorting-associated_protein_4A_Homo_sapiens
XP_021026470.1_vacuolar_protein_sorting-associated_protein_4A_Mus_caroli
NP_001094432.1_vacuolar_protein_sorting-associated_protein_4B_Danio_rerio
NP_004860.2_vacuolar_protein_sorting-associated_protein_4B_Homo_sapiens
CDJ92074.1_MIT_and_ATPase_and_Vps4_oligomerisation_domain_containing_protein_Haemonchus_contortus
NP_490816.4_related_to_yeast_Vacuolar_Protein_Sorting_factor_Caenorhabditis_elegans
KRZ08745.1_Vacuolar_protein_sorting-associated_protein_4A_Trichinella_zimbabwensis
NP_001016453.1_spastin_Xenopus_tropicalis
NP_001086725.1_spastin_Xenopus_laevis
NP_001026232.1_spastin_Gallus_gallus
NP_998914.1_spastin_Sus_scrofa
NP_998080.2_spastin_Danio_rerio
XP_020452523.1_spastin_Monopterus_albus
EFN84628.1_Spastin_Harpegnathos_saltator
XP_004997542.1_spastin_Salpingoeca_rosetta
XP_004349403.2_spastin_Capsaspora_owczarzaki_ATCC_30864
XP_013757524.1_spastin_Thecamonas_trahens_ATCC_50062
PNY10544.1_spastin_Trifolium_pratense
XP_004492651.1_spastin_Cicer_arietinum
RDX72538.1_Spastin_Mucuna_pruriens
KHG28541.1_Spastin_Gossypium_arboreum
XP_008354752.1_spastin_Malus_domestica
XP_025812956.1_spastin_Panicum_hallii
OLS31934.1_VCP-like_ATPase_Candidatus_Heimdallarchaeota_archaeon_AB_125
OLS27542.1_VCP-like_ATPase_Candidatus_Heimdallarchaeota_archaeon_LC_3
WP_011839560.1_AAA_family_ATPase_Staphylothermus_marinus
WP_013143471.1_AAA_family_ATPase_Staphylothermus_hellenicus
OYT38748.1_AAA_family_ATPase_Desulfurococcales_archaeon_ex4484_58
AFL66169.1_AAA_ATPase_central_domain_protein_Desulfurococcus_amylolyticus_DSM_16532
WP_048078977.1_ATP-binding_protein_Desulfurococcus_mucosus
ADG90458.1_AAA_ATPase_central_domain_protein_Thermosphaera_aggregans_DSM_11486
WP_048162763.1_ATP-binding_protein_Thermogladius_calderae
WP_012123137.1_AAA_family_ATPase_Ignicoccus_hospitalis
WP_075049343.1_AAA_family_ATPase_Ignicoccus_islandicusALU11485.1_ATPase_AAA_Ignicoccus_islandicus_DSM_13165
AEM38769.1_AAA_ATPase_central_domain_protein_Pyrolobus_fumarii_1A
WP_011821970.1_AAA_family_ATPase_Hyperthermus_butylicus
WP_010866097.1_ATP-binding_protein_Aeropyrum_pernix
WP_022541373.1_ATP-binding_protein_Aeropyrum_caminiBAN90099.1_ATPase_of_the_AAA_class_Aeropyrum_camini_SY1___JCM_12091
WP_013266870.1_ATP-binding_protein_Acidilobus_saccharovoransADL19358.1_AAA_family_ATPase_Acidilobus_saccharovorans_345-15
WP_015232243.1_ATP-binding_protein_Caldisphaera_lagunensisAFZ70345.1_AAA__family_ATPase_Caldisphaera_lagunensis_DSM_15908
AFH42378.1_cell_division_protein_CdvC_Fervidicoccus_fontis_Kam940
WP_009992337.1_ATP-binding_protein_Saccharolobus_solfataricus
WP_069284542.1_ATP-binding_protein_Sulfolobus_sp._A20AOL17607.1_AAA_family_ATPase_Sulfolobus_sp._A20
WP_084739680.1_AAA_family_ATPase_Sulfolobus_sp._JCM_16833
AHC51634.1_ATPase_AAA_Sulfolobus_acidocaldarius_SUSAZ
WP_010979234.1_ATP-binding_protein_Sulfurisphaera_tokodaii
WP_012021614.1_ATP-binding_protein_Metallosphaera_sedula
WP_013737161.1_ATPase_AAA_Metallosphaera_cuprinaAEB94663.1_ATPase_central_domain-containing_protein_Metallosphaera_cuprina_Ar-4
ARM77171.1_AAA_family_ATPase_Acidianus_manzaensis
WP_048054653.1_ATP-binding_protein_Acidianus_hospitalis
AFS81178.1_ATPase_central_domain-containing_protein_Candidatus_Nitrosopumilus_koreensis_AR1
AJM92244.1_AAA-type_ATPase_domain-containing_putative_archaeal_cell_division_protein_Candidatus_Nitrosopumilus_piranensis
WP_086907311.1_AAA_family_ATPase_Candidatus_Nitrosomarinus_catalinaARS64156.1_VCP-like_ATPase_Candidatus_Nitrosomarinus_catalina
PBO85519.1_AAA_family_ATPase_Thaumarchaeota_archaeon
WP_010190395.1_AAA_family_ATPase_Candidatus_Nitrosarchaeum_limnium
WP_119819537.1_AAA_family_ATPase_Candidatus_Bathyarchaeota_archaeon
OLS18192.1_VCP-like_ATPase_Candidatus_Odinarchaeota_archaeon_LCB_4
KKK42121.1_Vps4-like_ATPase_Lokiarchaeum_sp._GC14_75
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