

Accession	Protein description
W0TFR3	Nonsense-mediated decay protein 4 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=NMD4 PE=4 SV=1
W0TDX5	3'(2'),5'-bisphosphate nucleotidase OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=MET22 PE=3 SV=1
W0TH38	Eukaryotic translation initiation factor 2 subunit alpha OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SUI2 PE=3 SV=1
Q24FB1	Squalene-tetrahymanol cyclase THC1 OS=Tetrahymena thermophila (strain SB210) OX=312017 GN=THC1 PE=1 SV=2
W0T6E8	Sm_like super family OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=LSM7 PE=3 SV=1
W0TFD2	ATPase inhibitor, mitochondrial OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=KLMA_60492 PE=3 SV=1
W0TI22	Vacuolar protein sorting-associated protein OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=VPS13 PE=3 SV=1
W0T8I7	DUF3535 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=MOT1 PE=4 SV=1
W0T8D8	V-type proton ATPase proteolipid subunit OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=VMA3 PE=3 SV=1
W0TEM1	DNA damage-binding protein CMR1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=KLMA_70244 PE=3 SV=1
W0T858	GrpE protein homolog OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=mge1 PE=3 SV=1
W0TGZ2	ATP-dependent RNA helicase DHH1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=DHH1 PE=3 SV=1
W0T4R4	Chromatin structure-remodeling complex subunit RSC7 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=NPL6 PE=4 SV=1
W0TA87	Uncharacterized protein OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SAE2 PE=4 SV=1
W0TF89	Chromatin structure-remodeling complex subunit RSC9 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RSC9 PE=4 SV=1
W0T4Z8	Protein RMD9 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RMD9 PE=4 SV=1
W0TBR8	U6 snRNA-associated Sm-like protein Lsm3 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=LSM3 PE=4 SV=1
W0TEY9	C-1-tetrahydrofolate synthase OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=MIS1 PE=3 SV=1

W0TGR6	Conserved hypothetical membrane protein OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=KLMA_70420 PE=4 SV=1
W0TDK4	Target of rapamycin complex 2 subunit AVO1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=AVO1 PE=3 SV=1
W0T7V3	ISWI chromatin-remodeling complex ATPase ISW1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=ISW1 PE=3 SV=1
W0TK27	Protein MKT1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=MKT1 PE=3 SV=1
W0TAP1	Uncharacterized protein YPR085C OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=ASA1 PE=4 SV=1
W0T5B2	RSC chromatin remodeling complex subunit RSC8 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SWI3 PE=4 SV=1
W0T460	37S ribosomal protein NAM9 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=KLMA_10573 PE=3 SV=1
W0TGJ8	Ubiquinone biosynthesis protein COQ4, mitochondrial OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=COQ4 PE=3 SV=1
W0THZ6	U6 snRNA-associated Sm-like protein LSm4 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=LSM4 PE=3 SV=1
W0TA74	Putative aldehyde dehydrogenase-like protein YHR039C OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=ALDH22A1 PE=3 SV=1
W0TEU0	Sorting nexin-41 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SNX41 PE=3 SV=1
W0T7P3	Rtt102p super family OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=KLMA_10502 PE=4 SV=1
W0TA33	Antiviral helicase SKI2 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SKI2 PE=3 SV=1
W0T4T9	Replication protein A subunit OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RFA1 PE=3 SV=1
W0TH80	Chromatin-remodeling ATPase INO80 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=INO80 PE=3 SV=1
W0TBC9	Checkpoint serine/threonine-protein kinase BUB1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=BUB1 PE=4 SV=1
W0T7F9	Chromatin structure-remodeling complex protein RSC8 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RSC8 PE=4 SV=1

W0T9X2	Replication factor A protein 2 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RFA2 PE=3 SV=2
W0T916	Uncharacterized bolA-like protein YAL044W-A OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=KLMA_30590 PE=3 SV=1
W0TDR8	H2A super family protein OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=BUR6 PE=4 SV=1
W0TJ61	Heterogeneous nuclear rnp K-like protein 2 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=HEK2 PE=3 SV=1
W0T941	Chromatin structure-remodeling complex subunit RSC4 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RSC4 PE=4 SV=1
W0TER7	Nuclear protein STH1/NPS1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=STH1 PE=4 SV=1
W0TFQ1	Mitotic spindle-associated protein SHE1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SHE1 PE=4 SV=1
W0TA47	DNA-directed RNA polymerase III subunit RPC3 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RPC82 PE=3 SV=1
W0T4U9	Reduced growth phenotype protein 1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RGP1 PE=4 SV=1
W0TG55	Protein phosphatase 2C homolog 4 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=PTC4 PE=3 SV=1
W0TKL4	DNA-directed RNA polymerase subunit OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RPO31 PE=3 SV=1
W0TFW3	Actin patch protein 1 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=APP1 PE=4 SV=1
W0TCE3	Coatomer subunit epsilon OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SEC28 PE=3 SV=1
W0TCZ4	Antiviral protein SKI8 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SKI8 PE=4 SV=1
W0TEN0	Zf-C2H2 super family protein OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=AZF1 PE=4 SV=1
W0T601	U6 snRNA-associated Sm-like protein LSM5 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=LSM5 PE=3 SV=1
W0T9K4	Chromatin structure-remodeling complex protein RSC58 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=RSC58 PE=4 SV=1
W0T789	NAD-dependent histone deacetylase SIR2 OS=Kluyveromyces marxianus (strain DMKU3-1042 / BCC 29191 / NBRC 104275) OX=1003335 GN=SIR2 PE=3 SV=1