

Table S2. Protein sequences used to construct the phylogenetic tree.

>KAJ0960429.1

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>KAJ0961056.1

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>KAJ0961057.1

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>KAJ0961255.1(DzCYP72A23)

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>KAJ0961256.1(DzCYP72A24)

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>KAJ0961259.1(DzCYP72A25)

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>KAJ0962634.1

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>KAJ0962805.1

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>KAJ0964198.1(DzCYP94N8)

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>KAJ0965356.1

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>KAJ0966589.1

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>KAJ0966999.1

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>KAJ0967766.1

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>KAJ0977987.1

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>KAJ0978714.1

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>KAJ0978746.1

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>KAJ0978747.1

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>KAJ0978942.1

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RLLNIGSERELRESIATVHGATQIIRSKKEKLRSDDLLS RFISNEDNSDEF LR DIVIS FILAGRDTTSSALT WFFWLL
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SYNA YSMGRM ESIWG KDC VEFK PERW LDGE GA FRPEN PYR PVFHAG PRMCLG KEMAYIQ MKSIA CVL ER FMVD VLHKD
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>KAJ0980322.1

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CMKELFTLCNSCDQNP NHVAKV DMDKQWFGH LNYNIVVQM VAGK RYFGSGSASEE AWFR DAA TQFFHLLSFVPS DMF P
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DTTSTS LT KVL ANLL NHREVLKKVQEELDEQV GRDRV NVNESDT KKL VYLQAVIKESFRMTPPDPLIR RATQ KDCI LAGY
HVPKG TQVL VN AWKV HNDPNV WPEP SEFR PERFL SSQKG IDVKG QHYELI PFGT GRRV CPGV TMA LQVMN LTAR LIQGF
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>KAJ0981901.1

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>KAJ0981902.1(42)

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>KAJ0981907.1

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>KAJ0982068.1

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>KAJ0985657.1

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SSTIISVQDIMQECKTFFFAGKQTSSLT WATVLLAMHSDWQELARQEVLSVC GSSDIPT RDHLSKLKTLGMVLYETLR
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>KAJ0985668.1

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FFKKI

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>KAJ0985925.1(DzCYP72A2/9)

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FQT IYIPGYRFLPTPMNKRRSQVYNEMKRILKGMIKREKAIRMGDSSKNDLLGLLDSNMKEGEEHGKSQNKG M STEDV
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>KAJ0985926.1(DzCYP72A3)

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>KAJ0985933.1(DzCYP72A5)

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>KAJ0985934.1(DzCYP72A6)

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TMILLSMYPNWQANAREEVLQVFGKSTPDIEGLSHLKIVTMILYEVRLYPPGVFLDSKTSKAMELGGITYPSGVILSLP
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>KAJ0985935.1(DzCYP72A7)

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>KAJ0985941.1(DzCYP72A8)

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>KAJ0985942.1(DzCYP72A2/9)

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>KAJ0985943.1(DzCYP72A10)

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