## Mouse / human DYSF protein sequence alignment

Human	DYSF sequence accession number: NM_021469 DYSF sequence accession number: NM_003494 uman sequence is colored by its domain structure.	
Ferlin Region		
This	alignment was done using NCBI BLAST.	
Posit	ities = 1918/2062 (93%) ives = 1984/2062 (96%) = 16/2062 (0%)	
Mouse 21	SDPVASLIFRGVKKRTKVIKNSVNPVWNEGFEWDLKGIPLDQSSELLVVVKDHETMGRNR	80
20 Human	SD S +F GVKKRTKVIKNSVNPVWNEGFEWDLKGIPLDQ SEL VVVKDHETMGRNR SDAYCSAVFAGVKKRTKVIKNSVNPVWNEGFEWDLKGIPLDQGSELHVVVKDHETMGRNR	79
Mouse 81	FLGEAKIPLQEVLATPSLSASFNAPLLDAKQQPTGASLVLQVSYTPPPGAVPLFPPPASL FLGEAK+PL+EVLATPSLSASFNAPLLD K+QPTGASLVLQVSYTP PGAVPLFPPP L FLGEAKVPLREVLATPSLSASFNAPLLDTKKQPTGASLVLQVSYTPLPGAVPLFPPPTPL	140 139
Human		
Mouse 141	APSPTLPDMDLVPDTGGEEDTEDQGLTGDEAEPFLDQSAAVGPGGPTTPRKPPSHPPPHY PSPTLPD+D+V DTGGEEDTEDQGLTGDEAEPFLDQS PG PTTPRK PS PPPHY	200
140 Human	EPSPTLPDLDVVADTGGEEDTEDQGLTGDEAEPFLDQSGGPGAPTTPRKLPSRPPPHY	197
Mouse 201	PGAKRKRSSAPPRKLLSDKPQDFQIRVQVIEGRQLPGVNIKPVVKVTAAGQTKRTRIQKG PG KRKRS+ RKLLSDKPODFOIRVOVIEGROLPGVNIKPVVKVTAAGOTKRTRI KG	260
198 Human	PG KRARS+ RELESDAPQDFQIRVQVIEGRQLPGVNIKPVVKVIAAGQIKRIRI KG PGIKRKRSAPTSRKLLSDKPQDFQIRVQVIEGRQLPGVNIKPVVKVTAAGQTKRTRIHKG	257
Mouse 261	NSPLFNETLFFNVFDSPLELFDEPIFITVVDSRSLRTDALLGEFRMDVGTVYREPRHAYL	320
258 Human	NSPLFNETLFFN+FDSP ELFDEPIFITVVDSRSLRTDALLGEFRMDVGT+YREPRHAYL NSPLFNETLFFNLFDSPGELFDEPIFITVVDSRSLRTDALLGEFRMDVGTIYREPRHAYL	317
Mouse		200
321	RKWLLLSDPDDFSAGARGYLKASLCVLGPGDEAPLDKKDPSEDKEDIEGNLLRPTGVALR RKWLLLSDPDDFSAGARGYLK SLCVLGPGDEAPL++KDPSEDKEDIE NLLRPTGVALR	380
318	RKWLLLSDPDDFSAGARGYLKTSLCVLGPGDEAPLERKDPSEDKEDIESNLLRPTGVALR	377

Human

Mouse 381	GAHFCLKLFRAEDLPQMDDAVMDNVKQIFGFDSNKKNLVDPFVEVSFAGKMLCSKILEKT GAHFCLK+FRAEDLPQMDDAVMDNVKQIFGF+SNKKNLVDPFVEVSFAGKMLCSKILEKT GAHFCLKVFRAEDLPQMDDAVMDNVKQIFGFESNKKNLVDPFVEVSFAGKMLCSKILEKT	440
378 Human		437
Mouse 441	ANPQWNQNITLPAMFPSMCEKMRIRVMDWDRLTHNDIVATTYLGMSKISATGGEIEANPQWNQNITLPAMFPSMCEKMRIR++DWDRLTHNDIVATTYL MSKISA GGEIE ANPQWNQNITLPAMFPSMCEKMRIRIIDWDRLTHNDIVATTYLSMSKISAPGGEIEEEPA	496
438 Human		497
Mouse 497	VDDNLGFLPTFGPCYVNLYGSPREFTGFPDPYAELNTGKGEGVAYRGRVL +DD LGFLPTFGPCY+NLYGSPREFTGFPDPY ELNTGKGEGVAYRGR+L GAVKPSKASDLDDYLGFLPTFGPCYINLYGSPREFTGFPDPYTELNTGKGEGVAYRGRLL	546
498 Human		557
Mouse 547	LSLETKLVEHSEQKVEDLPADDILRVEKYLRRRKYSLFAAFYSATMLQDVDDAIQFEVSI LSLETKLVEHSEQKVEDLPADDILRVEKYLRRRKYSLFAAFYSATMLQDVDDAIQFEVSI LSLETKLVEHSEQKVEDLPADDILRVEKYLRRRKYSLFAAFYSATMLQDVDDAIQFEVSI	606
558 Human		617
Mouse 607	GNYGNKFDTTCLPLASTTQYSRAVFDGCHYYYLPWGNVKPVVVLSSYWEDISHRIETQNQ GNYGNKFD TCLPLASTTQYSRAVFDGCHYYYLPWGNVKPVVVLSSYWEDISHRIETQNQ GNYGNKFDMTCLPLASTTQYSRAVFDGCHYYYLPWGNVKPVVVLSSYWEDISHRIETQNQ	666
618 Human		677
Mouse 667	LLGIADRLEAGLEQVHLALKAQCSTEDVDSLVAQLTDELIADCSQPLCDIHEIPSATHLD LLGIADRLEAGLEQVHLALKAQCSTEDVDSLVAQLTDELIA CSQPL DIHE PSATHLD	726
678 Human	LLGIADRLEAGLEQVHLALKAQCSTEDVDSLVAQLTDELIAGCSQPLGDIHETPSATHLD	737
Mouse 727	QYLLRLRTRHLSQIKEAALALKLGHSELSTALEQAEDWLLHLRALAEEPQNSLPDIIIWM OYL +LRT HLSOI EAALALKLGHSEL ALEOAEDWLL LRALAEEPONSLPDI+IWM	786
738 Human	QYLYQLRTHHLSQITEAALALKLGHSELPAALEQAEDWLLRLRALAEEPQNSLPDIVIWM	797
Mouse 787	LQGDKRVAYQRVPAHEVLFSRRGPSYCGRNCGKLQTIFLKYPMEGMPGARMPVQIRIKLW LQGDKRVAYQRVPAH+VLFSRRG +YCG+NCGKLQTIFLKYPME +PGARMPVQIR+KLW	846
798 Human	LQGDKRVAYQRVPAHQVLFSRRGANYCGKNCGKLQTIFLKYPMEKVPGARMPVQIRVKLW	857
Mouse 847	FGLSVDEKEFNQFAEGKFSVFAETYENQTKLALVGNWGTTGFSYPKFSDVTGKIKLPKDS FGLSVDEKEFNQFAEGK SVFAETYEN+TKLALVGNWGTTG +YPKFSDVTGKIKLPKDS	906
858 Human	FGLSVDEKEFNQFAEGKLSVFAETYENETKLALVGNWGTTGLTYPKFSDVTGKIKLPKDS	917
Mouse 907	FRPTAGWAWAGDWFVCPEKTLLHDAAAGHLSFVEEVFENQTRLPGGQWIYMSDNYTDVNG FRP+AGW WAGDWFVCPEKTLLHD AGHLSFVEEVFENQTRLPGGQWIYMSDNYTDVNG FRPSAGWTWAGDWFVCPEKTLLHDMDAGHLSFVEEVFENQTRLPGGQWIYMSDNYTDVNG	966
918 Human		977

Mouse 967	EKVLPKDDIECPLGWKWEDEEWSTDLNRAVDEQGWEYSITIPPDRKPKHWVPVEKMYYTH EKVLPKDDIECPLGWKWEDEEWSTDLNRAVDEQGWEYSITIPP+RKPKHWVP EKMYYTH EKVLPKDDIECPLGWKWEDEEWSTDLNRAVDEQGWEYSITIPPERKPKHWVPAEKMYYTH	1026
978 Human		1037
Mouse 1027	RRRRWVRLRRRDLSQMEALKRHRQAEAEGEGWEYASLFGWKFHLEYRKTDAFRRRWRRR RRRRWVRLRRRDLSQMEALKRHRQAEAEGEGWEYASLFGWKFHLEYRKTDAFRRRWRRR	1086
1038 Human	RRRRWVRLRRRDLSQMEALKRHRQAEAEGEGWEYASLFGWKFHLEYRKTDAFRRRWRRR	1097
Mouse 1087	MEPLEKTGPAAVFALEGALGGMVDDKSEDSMSVSTLSFGVNRPTISCIFDYGNRYHLRCY MEPLEKTGPAAVFALEGALGG++DDKSEDSMSVSTLSFGVNRPTISCIFDYGNRYHLRCY MEPLEKTGPAAVFALEGALGGVMDDKSEDSMSVSTLSFGVNRPTISCIFDYGNRYHLRCY	1146
1098 Human		1157
Mouse 1147	LYQARDLAAMDKDSFSDPYAIVSFLHQSQKTVVEKNTLNPTWDQTLIFYEIEIFGEPASI +YQARDLAAMDKDSFSDPYAIVSFLHQSQKTVV KNTLNPTWDQTLIFYEIEIFGEPA++	1206
1158 Human	MYQARDLAAMDKDSFSDPYAIVSFLHQSQKTVVVKNTLNPTWDQTLIFYEIEIFGEPATV	1217
Mouse 1207	AEHPPCIVVELYDHDTYGADEFMGRCICQPSLERMPRLAWFPLTRGSQPAGELLAAFELI AE PP IVVELYDHDTYGADEFMGRCICQPSLERMPRLAWFPLTRGSQP+GELLA+FELI	1266
1218 Human	AEQPPSIVVELYDHDTYGADEFMGRCICQPSLERMPRLAWFPLTRGSQPSGELLASFELI	1277
Mouse 1267	QREKPAIHHIPGFEMHETSRILDETEDTDLPYPPPQREANIYMVPQNIKPALQRTAIEIL QREKPAIHHIPGFE+ ETSRILDE+EDTDLPYPPPQREANIYMVPQNIKPALQRTAIEIL	1326
1278 Human	QREKPAIHHIPGFEVQETSRILDESEDTDLPYPPPQREANIYMVPQNIKPALQRTAIEIL	1337
Mouse 1327	AWGLRNMKSYQMASISSPSLVVECGGQTVQSCVIRNLRKNPNFDVCTLFMEVMLPREDLY AWGLRNMKSYO+A+ISSPSLVVECGGOTVOSCVIRNLRKNPNFD+CTLFMEVMLPRE+LY	1386
1338 Human	AWGLRNMKSYQLANISSPSLVVECGGQTVQSCVIRNLRKNPNFDICTLFMEVMLPREELY	1397
Mouse 1387	CPPIVVKVIDNRQFGRRPVVGQCTIRSLENFLCDPYSAESPSPQGGPDDVSLLSPGEDVL CPPI VKVIDNRQFGRRPVVGQCTIRSLE+FLCDPYSAESPSPQGGPDDVSLLSPGEDVL CPPITVKVIDNRQFGRRPVVGQCTIRSLESFLCDPYSAESPSPQGGPDDVSLLSPGEDVL	1446
1398 Human		1457
Mouse 1447	IDIDDKEPLIPVQEEEFIDWWSKFFASVGEREKCGSYLEKDFDTLKVYDTQLENVEAFGG IDIDDKEPLIP+QEEEFIDWWSKFFAS+GEREKCGSYLEKDFDTLKVYDTQLENVEAF G	1506
1458 Human	IDIDDKEPLIPIQEEEFIDWWSKFFASIGEREKCGSYLEKDFDTLKVYDTQLENVEAFEG	1517
Mouse 1507	LSDFCNTFKLYRGRTQEETDDPSVIGEFKGLFKIYPLPEDPAIPMPPRQFHQLAAQGPQE LSDFCNTFKLYRG+TQEET+DPSVIGEFKGLFKIYPLPEDPAIPMPPRQFHQLAAQGPQE LSDFCNTFKLYRGKTQEETEDPSVIGEFKGLFKIYPLPEDPAIPMPPRQFHQLAAQGPQE	1566
1518 Human		1577

Mouse 1567	CLVRIYIVRAFGLQPKDPNGKCDPYIKISIGKKSVSDQDNYIPCTLEPVFGKMFELTCTL CLVRIYIVRAFGLQPKDPNGKCDPYIKISIGKKSVSDQDNYIPCTLEPVFGKMFELTCTL	1626
1578 Human	CLVRIYIVRAFGLQPKDPNGKCDPYIKISIGKKSVSDQDNYIPCTLEPVFGKMFELTCTL	1637
Mouse 1627	PLEKDLKITLYDYDLLSKDEKIGETVIDLENRLLSKFGARCGLPQTCCVSGPNKWRDKLR PLEKDLKITLYDYDLLSKDEKIGETV+DLENRLLSKFGARCGLPQT CVSGPN+WRD+LR	1686
1638 Human	PLEKDLKITLYDYDLLSKDEKIGETVVDLENRLLSKFGARCGLPQTYCVSGPNQWRDQLR	1697
Mouse 1687	PSQLLHLFCQQHRIKAPVYRTDRVTFQDKDYTIEEIEAGRLPNPHLGPVEERLALHVLQQ PSQLLHLFCQQHR+KAPVYRTDRV FQDK+Y+IEEIEAGR+PNPHLGPVEERLALHVLQQ	1746
1698 Human	PSQLLHLFCQQHRVKAPVYRTDRVMFQDKEYSIEEIEAGRIPNPHLGPVEERLALHVLQQ	1757
Mouse 1747	QGLVPEHVESRPLYSPLQPDIEQGKLQMWIDIFPKVLGQPGPPFNITPRKARRFFLRCII QGLVPEHVESRPLYSPLQPDIEQGKLQMW+D+FPK LG+PGPPFNITPR+ARRFFLRCII	1806
1758 Human	QGLVPEHVESRPLYSPLQPDIEQGKLQMWVDLFPKALGRPGPPFNITPRRARRFFLRCII	1817
Mouse 1807	WNTKDVILDDLSLTGEKMSDIYVKGWMVGFEEHKQKTDVHYRSLGGEGNFNWRFVFPFDY WNT+DVILDDLSLTGEKMSDIYVKGWM+GFEEHKQKTDVHYRSLGGEGNFNWRF+FPFDY	1866
1818 Human	WNT-DVILDDLSLTGEKMSDIYVKGWMIGFEEHKQKTDVHYRSLGGEGNFNWRFIFPFDY	1877
Mouse 1867	LPAEQVCAVAKKDAFWRLDKTESKIPARVVFQIWDNDKFSFDDFLGSLQLDLNRMPKPAK	1926
1878 Human	LPAEQVC +AKKDAFWRLDKTESKIPARVVFQIWDNDKFSFDDFLGSLQLDLNRMPKPAK LPAEQVCTIAKKDAFWRLDKTESKIPARVVFQIWDNDKFSFDDFLGSLQLDLNRMPKPAK	1937
Mouse 1927	TAEKCSLDQLDDTFHPEWFVSLFEQKTVKGWWPCVTEEGEKKMLAGKLEMTLEIVAESEH TA+KCSLDQLDD FHPEWFVSLFEQKTVKGWWPCV EEGEKK+LAGKLEMTLEIVAESEH	1986
1938 Human	TAKKCSLDQLDDAFHPEWFVSLFEQKTVKGWWPCVAEEGEKKILAGKLEMTLEIVAESEH	1997
Mouse 1987	EERPAGQGRDEPNMNPKLEDPRRPDTSFLWFTSPYKTMKFILWRRFRCAIILFIILFILL	2046
1998 Human	EERPAGQGRDEPNMNPKLEDPRRPDTSFLWFTSPYKTMKFILWRRFR AIILFIILFILL EERPAGQGRDEPNMNPKLEDPRRPDTSFLWFTSPYKTMKFILWRRFRWAIILFIILFILL	2057
Mouse 2047	LFLGVFVYAFPNYAAMKLVKPF 2068	
2058 Human	LFL +F+YAFPNYAAMKLVKPF LFLAIFIYAFPNYAAMKLVKPF 2079	