Multiple Sequence Alignment of IDH1 with WebPRANK

>Arabidopsis thaliana MSRRSLTLLKNLARNANGSGIQTRSVTYMPRPGDGAPRAVTLIPGDGIGPLVTNAVEQVM EAMHAPIFFEKYDV-HGEMS----RV-PPEVMESIRKNKVCLKGGLKTPVG-GGV--------SSLNVOLRKELD-LFASLVNCFNLPGLPTRHENVDIVVIRENTEGEYAGLEHEV VPGV-----VESLK-VITKF-----CSERIAKYA---FEYAYLNNR KKVTAVHKANIMKLADGLFLESCREV-AKKYPS-----ITYNEIIVDNCCMOLVAKPEO FDVMVTPNLYGNLVANTAAGIAGGTGVM-----PGGNVGADHAVFEQGASAGNVGKDKI VLENKANPVALLLSSAMMLRHL------QFPSFADRLETAVKKVIAEGKCRTKDLG----GTSTTQ------EVVDAVIAKL-----D->Cyberlindnera jadinii -----MLRTQVKSQSKRSLATVSEALLPKK-YGGRYTVTLLPGDGIGKEITDSVVEIF KAERIPVDWETVNVTNSEKS-----KEGVKEAVESLKRNKVGLKGIWKTDVSQSGH--------GSLNVAFRKELD—IYASLTLIKNIPGVKTRMDGIDLVLIRENTEGEYSGLEHES VPGV-----KSERIAKFA---FDFAKKNNR QTVTAVHKANIMKLADGLFRKTVKDVGAEQYPG-----IEVKDIIVDNASMQAVSNPQQ FDVLVTPNLYGAILSNIGTALVGGPGLV-----PGANYGREYAVFEPGSR--HVGLE-I AGKGVANPTGMILSSVLMLRHL-----GLDSSADRISNAVYKVIADGKKTTRDIG----GPASTK-----EFTQAVIDEL-----ASA

>Homo_sapiens

IFPYVELDLHSYDL-GIENRDATNDQV-TKDAAEAIKKHNVGVKCATITPDE-KRVEEFK
LKQMWKSPNGTIRNILGGTVFREAIICKNIPRLVSGW--VKPIIIGRHAYGDQYRATDFV
VPGPGKVEMTYTPSDGTQKVTYLVHNFEEGGGVAMGMYNQDKSIEDFAHSSFQMA-LSKG
WPLYLSTKNTILKKYDGRFKDIFQEIYDKQYKSQFEAQKIWYEHRLIDDMVAQAMKSEGG
F-IWACKNYDGDVQSDSVAQGYGSLGMMTSVLVCPDGKTVEAEAAHGTVTR--HYRMYQK
GQETSTNPIASIFAWTRGLAHRAKLDNNKELAFFANALEEVSIETIEAG-FMTKDLAACI
KGLPNVQRSDYLNTFEFMDKLGENLKIKL-AQAKL

>Cricetulus_griseus

-----MSKKIHGG-SVVEMQGDEMTRIIWELIKEKL ILPYV------TKDAAEAIKKYNVGVKCATITPDE-KRVEEFK LKQMWKSPNGTIRNILGGTVFREAIICKNIPRLVTGW--VKPIIIGRHAYGDQYRATDFV VPGPGKVEITYTPKDGTQKVTYVVHNFEEGGGVAMGMYNEDKSIEDFAHSSFQMA-LSKG WPLYLSTKNTILKKYDGRFKDIFQEIYDKQYKSQFEAQKIWYEHRLIDDMVAQAMKSEGG F-IWACKNYDGDVQSDSVAQGYGSLGMMTSVLICPDGKTVEAEAAHGTVTR--HYRMYQK GQETSTNPIASIFAWSRGLAHRAKLDNNTELSFFAKALEDVCIETIEAG-FMTKDLAACI KGLPNVQRSDYLNTFEFMDKLGENLKAKL-AQAKL

>Zebrafish

------MDSSRNRMRFGGFKMSQKVKAG—SVVEMQGDEMTRVIWELIKEKL IFPYLELDLHSYDL-GMENRDATDDKV-TVEAAEAVRRYNVGIKCATITPDE-KRVEEFK LKQMWRSPNGTIRNILGGTVFREAIICKNIPRLVPGW--IKPIIIGRHAHGDQYKATDFV VPGPGTVEMTYKPKNGGETLKFVVHDFEGTGGVALGMYNTDKSIRDFAHSSFQMG-LNKG WPMYLSTKNTILKKYDGRFKDIFQEIYDKEYKAKYEAMGIWYEHRLIDDMVAQAMKSEGG F-IWACKNYDGDVQSDSVAQGYGSLGMMTSVLVCPDGRTVEAEAAHGTVTR--HYRMHQQ GKETSTNPIASIFAWTRGLLHRAELDKNAELRVFAEALEVVCVETIEAG-FMTKDLAICI KGMSGVTRSDYLNTFEFLDKLAENLKIKLSSQPKL

