



	720	730	740	750	760	77 ọ
U5WHY8_9CHIR						I <mark>girdr</mark> rktd q ars e
E2DHI4_9CHIR						F <mark>GIRDR</mark> R <mark>K</mark> TD <mark>Q</mark> ARS E
E2DHI7_9CHIR						F <mark>GIRDR</mark> R <mark>KTDQ</mark> ARS E
E2DHI9_9CHIR	DDNSLEFI	.GIQPTLGPPY	QPPVTIWL:	IVFGVVMAVV	7VGIVVLII:	F <mark>GIRDR</mark> R <mark>K</mark> AD Q ARS E
E2DHI3_RHIMR						F <mark>GIRDR</mark> R <mark>K</mark> TD <mark>Q</mark> ARS E
B6ZGN7_RHIFE						I <mark>girdr</mark> r <mark>kk</mark> d <mark>q</mark> ars <mark>e</mark>
E2DHI2_RHIFE	DDNSLEFI	GIQPTLGPPY	QPPVTIWL	IVFGVVMAVV	VGIVV <mark>LI</mark> I:	F <mark>GIRDR</mark> R <mark>K</mark> KD Q ARS E
A4PIG8_ROULE	DDDTLEFI	GIEPTLGTPY	QPPVTIWL	I VFGVVM GLV <mark>V</mark>	VGIVL <mark>LI</mark> FV	/GIRDRRKKNQERSE
D8WU01_ROULE	DDDTLEFI	GIEPTLGTPY	QPPVTIWL	IVFGVVMGLV	VGIVL <mark>LI</mark> FV	/GIRDRRKKNOERSE
G1PXH7_MYOLU	DDNTLEFI	.GIQPTLGPPN	QPPVTIWL	I VFGVVM GVA	/IGIAV <mark>LI</mark> F:	I <mark>GIRDR</mark> K <mark>K</mark> KK <mark>Q</mark> AGN <mark>E</mark>

	780	790	8 O Ö
U5WHY8_9CHIR	ENPYSS	VDLSKGENNP	GFQNGDDVQTSF
E2DHI4_9CHIR	ENPYSS	VDLSKGENNP	GFQNGDDVQTSF
E2DHI7_9CHIR	ENPYSS	V D L S K G E N N P	GFQNGDDVQTSF
E2DHI9_9CHIR	ENPYPS	V D L S K G E N N P	GFQNGDDVQTSF
E2DHI3_RHIMR	ENPYPS	V D L S K G E N N P	GFQNGDDVQTSF
B6ZGN7_RHIFE	ENPYSS	V D L S K G E N N P	GFQNGDDVQTSF
E2DHI2_RHIFE	ENPYSS	V D L S K G E N N P	GFQNGNDVQTSF
A4PIG8_ROULE			GFQNNDDVQTSF
D8WU01_ROULE			GFQNNDDVQTSF
G1PXH7_MYOLU	ENPYSSY	VNLSKGENNP	GFQNGDDVQTSF