

Drcd-1r/Dsec	MSEEP I PVMSPQRQAEREKLYQLI IELAYPATREVALLELSKKNPDLAPMLWNSFGTSY	60
Drcd-1r/Dsim	MSEEP I PVMTPQRQAEREKLYQLMIELAYPASREVALLELSKKNPDLAPMLWNSFGTSY	60
Drcd-1r/Dmel	MSAEPSPVMSPQQQAEREKVYQLI IELAYPATRETALLELSKNTYADLAPMLWKSVMGTTTC	60
	** ** ***:***:*****:***:*****:***.*****:.. *****:*.**:	
Drcd-1r/Dsec	TLLQEVVNVYPFISNPILKANQSNRVCYALTLLQCVAHPGTRPAFLRGQIPVYVYPFIS	120
Drcd-1r/Dsim	TLLQEVVNVYPFISNPILKANQSNRVCYALTLLQCVAHPGTRPAFLRGQIPVYVYPFIS	120
Drcd-1r/Dmel	TLLQEIVNIYPIITTPVLKANQSNRVCYALTLLQCVASHPETRPAFLRDQIPMYLYPFLS	120
	*****:***:***:*.*:..*:*****:*****:*** *****.***:*.***:*	
Drcd-1r/Dsec	TTFKSKPFQRLRLNSLRVIGALVETDDTEAISFLLSTEVVPLCLSNMLKGSMLTKFAAIK	180
Drcd-1r/Dsim	TTFKSKPFQRLRLNSLGVIGALVETDDTEAISFLLSSEVVPLCLSNMLKGSMLTKSAAIK	180
Drcd-1r/Dmel	TTFKSRPFQRLRLTTLGVINALAETGDTEVLIFLIWSEVVPCLTNMVRGSKLTKIAATS	180
	*****:*****.:* **.*.*.*.*.*.*. : ** : ***** **.*.*.:** *** ** .	
Drcd-1r/Dsec	IIEKILLNETGLSYICETHDRFAGVAITLGKMTIRMMKSPCFRVLKHVVR CYLLLTENEI	240
Drcd-1r/Dsim	IIEKILLNKTGLSYICETHDRFARVATTLGKMTIRMMKYPCFRVLKRVVRCYLLITENEL	240
Drcd-1r/Dmel	IIEKILLDEMGLTYICENHDRFSQVAITLGKMTIHMVKFPCFRVLKHVVR CYLLLTENAR	240
	*:*****.: **:*****.*****: ** *****.*:*:* **:******:*****:***	
Drcd-1r/Dsec	ARSALGVCLPDQLRDGTFSSAIQHDTCTKQWLKTLKLNLEPRAPAAVQQVVE SPQDS	297
Drcd-1r/Dsim	GRSALRVCLPDPLRDGTFSSAMQHDPCAQWLKTLKLNLESRAPAVMLQVVEST EGS	297
Drcd-1r/Dmel	ARSALRVCLPDLRLRDGTFSTLVQHDTCTKQWLQMLLKNLQTNVAVNPMGSS-----	290
	.**** ***** *****:* :***.*:*****: *****:..* : .	