

human SRC	MGSNKS KPKDASQRRRSLEPAENVHGAGGGAFFPASQTPSKPASADGHRGPSAAAFAPAAAE	60
murine SRC	MGSNKS KPKDASQRRRSLEPSENVHGAG-GAFFPASQTPSKPASADGHRGPSAAAFVPPAAE	59
2JYQ_1 Chain	-----	0
human SRC	PKLFGGFNSSDTVTSPQRAGPLAGGVTTFFVALYDYESRTETDLSFKKGERLQIVNNTEGD	120
murine SRC	PKLFGGFNSSDTVTSPQRAGPLAGGVTTFFVALYDYESRTETDLSFKKGERLQIVNNTEGD	119
2JYQ_1 Chain	-----	0
human SRC	WWLAHSLSTGQTGYIPSNYVAPSDSIQAE EWYFGKITRRESERLLLLNAENPRGTFVLVRES	180
Murine SRC	WWLAHSLSTGQTGYIPSNYVAPSDSIQAE EWYFGKITRRESERLLLLNAENPRGTFVLVRES	179
2JYQ_1 Chain	-----QAE EWYFGKITRRESERLLLLNPEENPRGTFVLVRES	34
	*****	
human SRC	ETTKGAYCLSVSDFDNAKGLNVKHYKIRKLDSGGFYITSRTQFNSLQQLVAYYSKHADGL	240
Murine SRC	ETTKGAYCLSVSDFDNAKGLNVKHYKIRKLDSGGFYITSRTQFNSLQQLVAYYSKHADGL	239
2JYQ_1 Chain	ETTKGAYCLSVSDFDNAKGLNVKHYKIRKLDSGGFYITSRTQFSSLQQLVAYYSKHADGL	94
	*****	
Human SRC	CHRLTTVCPTSKPQTQGLAKDAWEIPRESLRLEVKLGGQCFGEVWMGTWNGTTRVAIKTL	300
murine SRC	CHRLTTVCPTSKPQTQGLAKDAWEIPRESLRLEVKLGGQCFGEVWMGTWNGTTRVAIKTL	299
2JYQ_1 Chain	CHRLTNVCPTSK-----	106
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Human SRC	KPGTMSPEAFLQEAQVMKKLRHEKLVQLYAVVSEEPYIYVTEYMSKGSLLDFLKGETGKY	360
murine SRC	KPGTMSPEAFLQEAQVMKKLRHEKLVQLYAVVSEEPYIYVTEYMNKGSLLDFLKGETGKY	359
2JYQ_1 Chain	-----	106
human SRC	LRLPQLVDMAAQIASGMAYVERMNYVHRDLRAANILVGENLVCKVADFGLARLIEDNEYT	420
murine SRC	LRLPQLVDMSAQIASGMAYVERMNYVHRDLRAANILVGENLVCKVADFGLARLIEDNEYT	419
2JYQ_1 Chain	-----	106
human SRC	ARQGAKFFIKWTAPEAAALYGRFTIKSDVWSFGILLTELTTKGRVPYPGMVNREVLDQVER	480
murine SRC	ARQGAKFFIKWTAPEAAALYGRFTIKSDVWSFGILLTELTTKGRVPYPGMVNREVLDQVER	479
2JYQ_1 Chain	-----	106
human SRC	GYRMPCPPECPESLHDLMCQCWRKEPEERPTFEYLAFLDYFTSTEPQYQPGENL	536
murine SRC	GYRMPCPPECPESLHDLMCQCWRKEPEERPTFEYLAFLDYFTSTEPQYQPGENL	535
2JYQ_1 Chain	-----	106