

Descriptions		Graphic Summary	Alignments	Taxonomy	Sequences producing significant alignments		Download	New Select columns	Show	100	?
select all		100 sequences selected		GenPept	Graphics	Distance tree of results	Multiple alignment	New	MSA Viewer		
	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession		
<input checked="" type="checkbox"/>	C-SRC (SH2 DOMAIN) COMPLEXED WITH ACE-MALONYL TYR-GLU-(N,N-DIPENTYL AMINE) [Homo sapiens]	Homo sapiens	226	226	100%	2e-78	100.00%	107	1A07_A		
<input checked="" type="checkbox"/>	C-SRC (SH2 DOMAIN WITH C188A MUTATION) COMPLEXED WITH ACE-FORMYL PHOSPHOTYR-GLU-(N,N-DIPENTYL AMINE) [Homo sapiens]	Homo sapiens	223	223	100%	3e-77	99.07%	107	1A1A_A		
<input checked="" type="checkbox"/>	Crystal Structure Of Sh2 In Complex With Ru78300 [Homo sapiens]	Homo sapiens	222	222	98%	1e-76	100.00%	108	1O41_A		
<input checked="" type="checkbox"/>	Triple mutant Src SH2 domain [Homo sapiens]	Homo sapiens	220	220	100%	9e-76	97.20%	112	4F59_A		
<input checked="" type="checkbox"/>	Crystal Structure Of Sh2 In Complex With Phosphate [Homo sapiens]	Homo sapiens	218	218	98%	3e-75	99.05%	108	1O4C_A		
<input checked="" type="checkbox"/>	The structure of 1NA in complex with Src T338G [Homo sapiens]	Homo sapiens	225	225	99%	2e-73	100.00%	448	4K11_A		
<input checked="" type="checkbox"/>	Crystal Structure Of Human Tyrosine-Protein Kinase C-Src [Homo sapiens]	Homo sapiens	225	225	99%	2e-73	100.00%	452	1FMK_A		
<input checked="" type="checkbox"/>	Structure of unphosphorylated c-Src in complex with an inhibitor [Homo sapiens]	Homo sapiens	225	225	99%	2e-73	100.00%	452	1Y57_A		
<input checked="" type="checkbox"/>	Structure of Human c-Src Tyrosine Kinase (Thr338Gly Mutant) in Complex with N6-benzyl ADP [Homo sapiens]	Homo sapiens	225	225	99%	2e-73	100.00%	452	1KSW_A		
<input checked="" type="checkbox"/>	Autoinhibited Src kinase bound to ADP [Mus musculus]	Mus musculus	225	225	99%	2e-73	100.00%	455	6F3F_A		
<input checked="" type="checkbox"/>	Src kinase in complex with a quinazoline inhibitor [Homo sapiens]	Homo sapiens	226	226	99%	5e-73	100.00%	535	2H8H_A		
<input checked="" type="checkbox"/>	Crystal Structure of a Complex of the Src SH2 Domain with Conformationally Constrained Peptide Inhibitor [Rous sarcoma virus]	Rous sarcoma virus	212	212	96%	7e-73	97.09%	106	1IS0_A		
<input checked="" type="checkbox"/>	Crystal Structure Of The Phosphotyrosine Recognition Domain Sh2 Of V-Src Complexed With Tyrosine-Phosphorylated Peptide [Rous sarcoma virus]	Rous sarcoma virus	212	212	96%	9e-73	97.09%	104	1SHA_A		
<input checked="" type="checkbox"/>	COCRYSTAL STRUCTURE OF UREA-SUBSTITUTED PHOSHOPEPTIDE COMPLEX [Rous sarcoma virus]	Rous sarcoma virus	210	210	95%	6e-72	97.06%	113	1SKJ_A		
<input checked="" type="checkbox"/>	Crystal Structure of Src SH2 domain bound to doubly phosphorylated peptide PQpYEpYIPI [Rous sarcoma virus (strain 17B3)]	Rous sarcoma virus (strain 17B3)	209	209	95%	7e-72	97.06%	103	1NZL_A		
<input checked="" type="checkbox"/>	Crystal Structure of the Src SH2 Domain Complexed with Peptide (SDpYANFK) [Gallus gallus]	Gallus gallus	209	209	95%	8e-72	97.06%	102	1P13_A		
<input checked="" type="checkbox"/>	CHICKEN SRC TYROSINE KINASE [Gallus gallus]	Gallus gallus	219	219	99%	4e-71	97.17%	453	2PTK_A		

図4.7 BLAST検索の結果