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Supporting information for article:

Structure of a lectin from the sea mussel *Crenomytilus grayanus* (CGL)

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Table S1 Models used in the MR searches, identified by *SALAMI* server

PDB code	Resolution [Å]	Size of the alignment	Identity	Coverage	Rmsd [Å]
4efr	2.50	140	14	1.00	2.98
3vsf	2.76	138	49	0.99	0.72
1t9f	2.00	138	17	0.99	2.92
2x2s	1.60	134	24	0.96	2.08
3mal	1.95	136	15	0.97	2.97
3nbc	1.01	132	18	0.94	2.20
4g9n	2.20	132	25	0.94	1.91
2y9g	1.67	132	16	0.94	2.95
3phz	1.70	131	20	0.94	2.50
4i4q	1.51	130	15	0.93	2.97
4i4u	1.57	130	13	0.93	2.96
2f2f	2.40	129	17	0.92	2.37
3n0k	2.80	129	10	0.92	2.99
1ups	1.82	128	22	0.91	2.94
3a22	1.90	127	28	0.91	1.98
4jp0	1.80	127	20	0.91	2.71
4a7k	2.00	135	13	0.96	2.95
2fdb	2.28	126	12	0.90	2.91
1qqk	3.10	126	8	0.90	2.99
1nun	2.90	125	11	0.89	2.97
4jpz	3.02	125	11	0.89	2.96
1g82	2.60	125	11	0.89	2.96
4oeg	1.60	124	10	0.89	2.98
3k1x	1.98	124	10	0.89	2.91
3q7y	1.45	123	18	0.88	2.43
3f1r	2.50	125	11	0.89	2.95
4jq0	3.84	125	10	0.89	2.97
3o49	1.45	122	16	0.87	2.45
3kmv	1.80	121	15	0.86	2.94
3o3q	1.60	121	14	0.86	2.39
2p39	1.50	123	13	0.88	3.00
2p23	1.80	124	8	0.89	2.99
1dqg	1.70	117	15	0.84	2.97
4gai	1.49	128	8	0.91	2.99
1mc9	1.70	116	40	0.83	2.86
3a07	1.19	112	29	0.80	2.65
3p6j	1.35	110	17	0.79	2.43
2wry	1.58	128	9	0.91	2.99
8i1b	2.40	127	9	0.91	2.98