Supporting Information

Supporting Table S1.

Sequence identity statistics for RsgI-CBM3 modules.

The identity and high similarity percentage values between the pairs of RsgI-CBM3's are

presented. PDB codes are indicated.

Protein	**0059**CBM3b, 4B9C	**0267**CBM3b, 4B9P
0267CBM3b, 4B9P	Identical: 28.4% Highly similar: 21.3%	
0404CBM3b, 4B97	Identical: 35.1% Highly similar: 23.8%	Identical: 40.5% Highly similar: 20.8%

Supporting Table S2.

R.m.s.d. values of Cα atoms for superposed structures of several CBM3's.

Calculated with PyMOL. The number of Cα atoms used for alignment is shown in parentheses.

Values are in Å.

Protein	CipA CBM3a 1NBC	**0059**CBM3b 4B9C	**0267**CBM3b 4B9P
0059CBM3b 4B9C	0.988 (112)		
0267CBM3b 4B9P	0.661 (113)	0.783 (86)	
0404CBM3b 4B97	0.711 (118)	0.738 (106)	0.737 (115)