

Table 1. Design of relaxed consensi .

	Phage display Consensus	Relaxed consensus	N of peptides screened
Rvs167	RxFPxxPP PP□ PPR	(R/K) xxPxxP PxxPx (R/K)	735 729
Yfr024c	PxLPxRP	(R/K) xxPxxP PxxPx (R/K)	735 729
Ysc84	PxLPxR	(R/K) xxPxxP PxxPx (R/K)	735 729
Boi1	RxxPxxP pPRxPrR□	(R/K) xxPxxP PxxPx (R/K)	735 729
Boi2	ppRnPxR□	(R/K) xxPxxP PxxPx (R/K)	735 729
Abp1	+P□ xxPx+P PxxPxRP	xxPx (K/R) P Pxxx (K/R) P	687 714
Myo5	Px@pPPxxP	(R/K) xxPxxP (F/P/L/W/A/E) xx (W/Y/L/M/F/H) x xPxxP	735 470
Amph1	PxRPxR	(P/F/L/I) (K/R) RP (P/F/L/I) xRPx (R/K) (P/L/R/F/S/I/V/K/G) Px (R/K) PP	1684 1139 951
End1	P+RPPxP	(P/F/L/I) (K/R) RP (P/F/L/I) xRPx (R/K) (P/L/R/F/S/I/V/K/G) Px (R/K) PP	1684 1139 951

@: aromatic residues; □ : hydrophobic residues; + : arginine or lysine; x: any residue.

The phage display *consensus* sequences were obtained by Tong et al (2002) and by Cestra et al (2001).