

**Fig. S1.** Pairwise identities between the Src homologs presented in Figure 1. Identities were calculated by WU-BLAST (<http://www.proweb.org/proweb/Tools/WU-blast.html>).

**Fig. S2.** Immobilized MbSrc1 SH3 and SH2 domains bind to Cas in SYF cell lysates. The blot presented in Figure 2A was stripped and reprobed with anti-Cas antibody.

**Fig. S3.** Binding of the MbSrc1 SH3 domain to a Pro-rich peptide derived from the sequence of Sam68 (RGAAPPPPPVPRGRG). A representative titration is shown using 20  $\mu$ M SH3 domain and 5  $\mu$ l injections of 2 mM peptide. The dissociation constant derived from this titration was 2.1  $\mu$ M.

**Fig. S4.** Binding of MbSrc1 to the pYEEI-substrate peptide. The pYEEI-substrate peptide used in Figure 5 was coupled to Affi-Gel 15 resin as described previously (ref. 24). The immobilized peptide (or Affi-Gel resin as control) was incubated with lysate from SYF cells expressing FLAG-MbSrc1 (25  $\mu$ g total protein). After washing, the bound protein was detected by anti-FLAG Western blotting.

**Fig. S5.** pYEEI-substrates with shorter linker lengths are not phosphorylated efficiently by MbSrc1. The peptides tested were: the FEEI-substrate control shown in Figure 5; Y(7)pY, a shorter variant of the pYEEI-substrate with a 7-amino acid linker (ref. 42); and Y(3)pY, with a linker length of 3 amino acids. Phosphorylation of the peptides was measured using the phosphocellulose paper assay, as described in Materials and Methods.

**Fig. S6.** Identities between MbCsk and the Csk homologs presented in Figure 6. Identities were calculated by WU-BLAST (<http://www.proweb.org/proweb/Tools/WU-blast.html>).

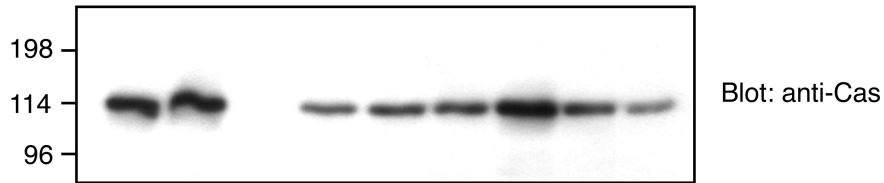
**Fig. S7.** MALDI-TOF analysis of the MbCsk reaction on MbSrc1. The reactions are described in the legend to Figure 8A. This figure shows a larger range of m/z. The arrow in the bottom panel points to the peak arising from the phosphorylated C-terminal tail of MbSrc1. The large peak at ca. 2350 likely represents residual unmodified C-tail peptide.

**Fig. S8.** The indicated FLAG-tagged MbSrc1 constructs were expressed in SYF cells. Lysates were analyzed by SDS-PAGE and Western blotting with anti-phosphotyrosine and anti-FLAG antibodies.

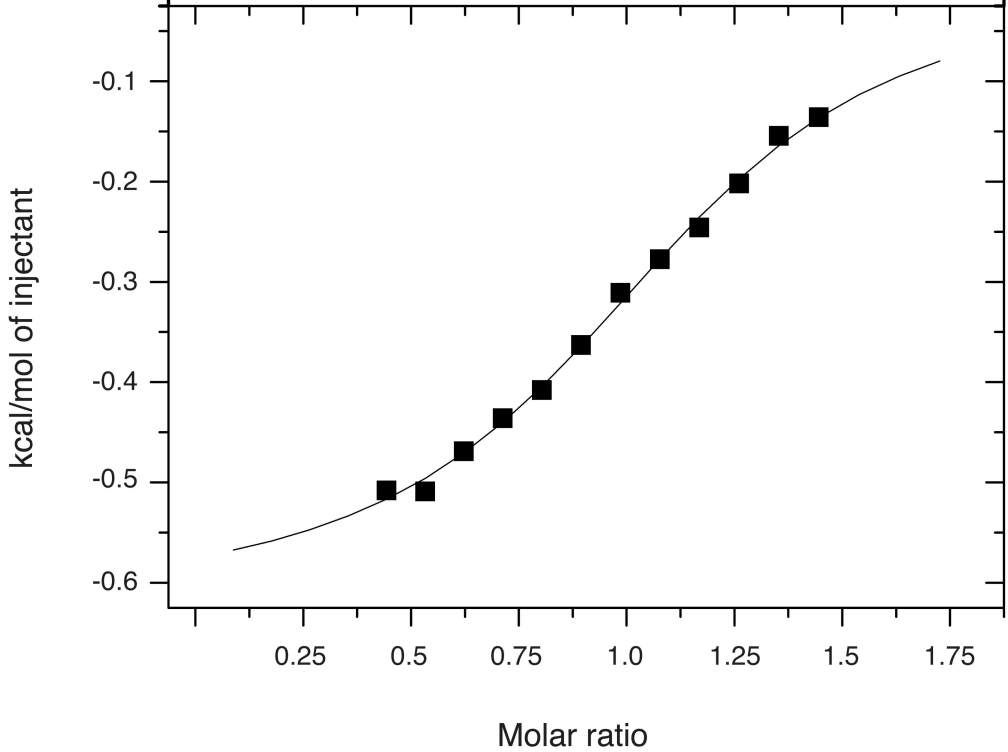
Database	Query					
	Hsap	Dmel	Hvul	Eflu	Mbre	Mova
Hsap		54% Id, 73% Pos	65% Id, 77% Pos	57% Id, 72% Pos	61 % Id, 77% Pos	60% Id, 75 % Pos
Dmel	50% Id, 67% Pos		50% Id, 68% Pos	49% Id, 67%Pos	49% Id, 68% Pos	49% Id, 68% Pos
Hvul	65% Id, 77% Pos	51% Id, 69% Pos		61%Id, 75% Pos	63% Id, 78% Pos	60% Id, 78% Pos
Eflu	56% Id, 71% Pos	49% Id, 67% Pos	57% Id, 72% Pos		58% Id, 75% Pos	58 % Id, 75 % Pos
Mbre	57% Id, 72% Pos	49% Id, 68% Pos	60% Id, 74% Pos	58% Id, 75% Pos		73% Id, 84% Pos
Mova	59% Id, 74% Pos	50% Id, 68% Pos	57% Id, 74% Pos	57% Id, 75% Pos	73%Id, 84%Pos	

Supplemental Figure S1

		GST		GST-SH2		GST-SH3		pulldown	
lysate		-	-	-	-	-	10	100	polyPro peptide ( $\mu\text{M}$ )
+Src	-Src	-	-	10	100	-	-	-	pYEEI peptide ( $\mu\text{M}$ )



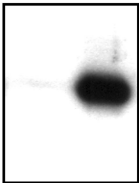
Supplemental Figure S2



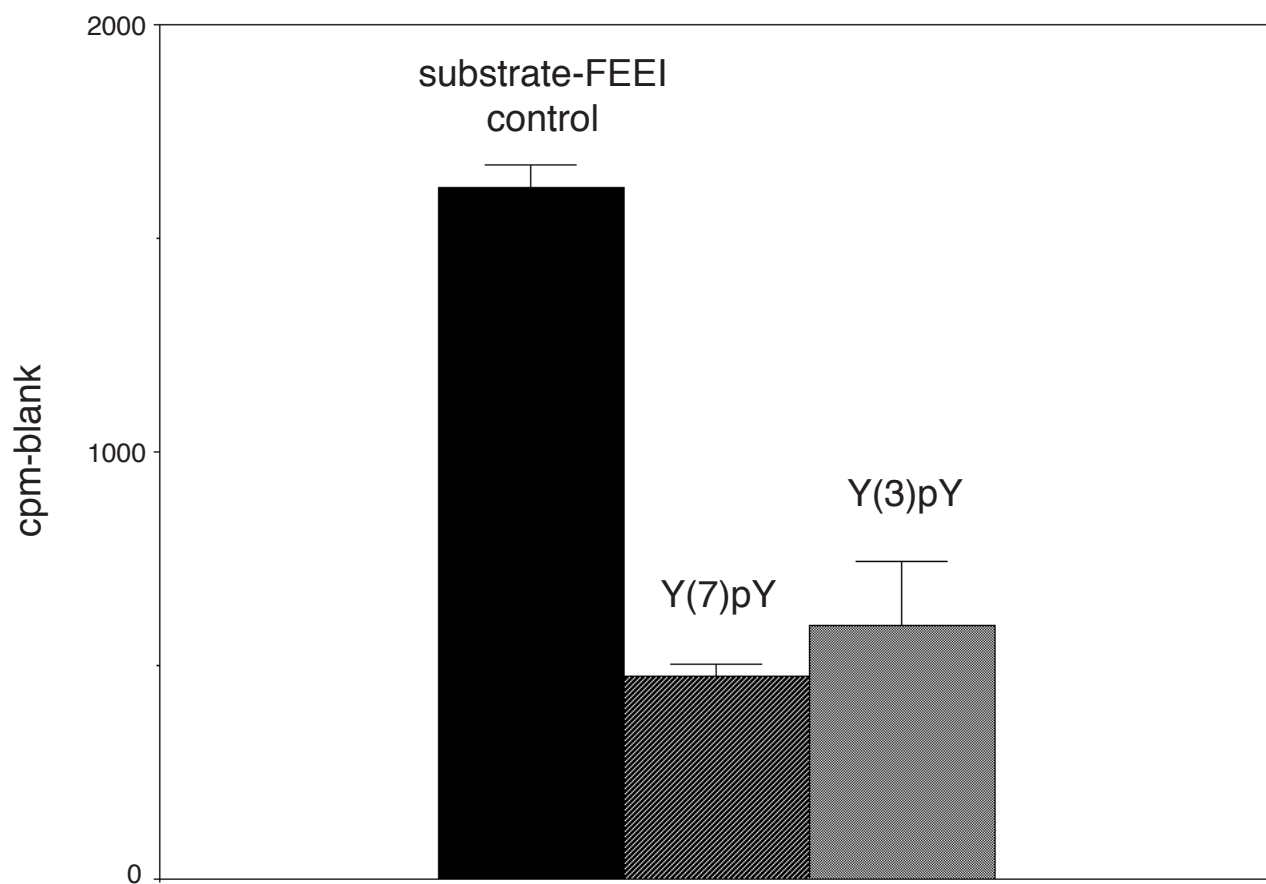
Supplemental Figure S3

control

substrate-pYEEI



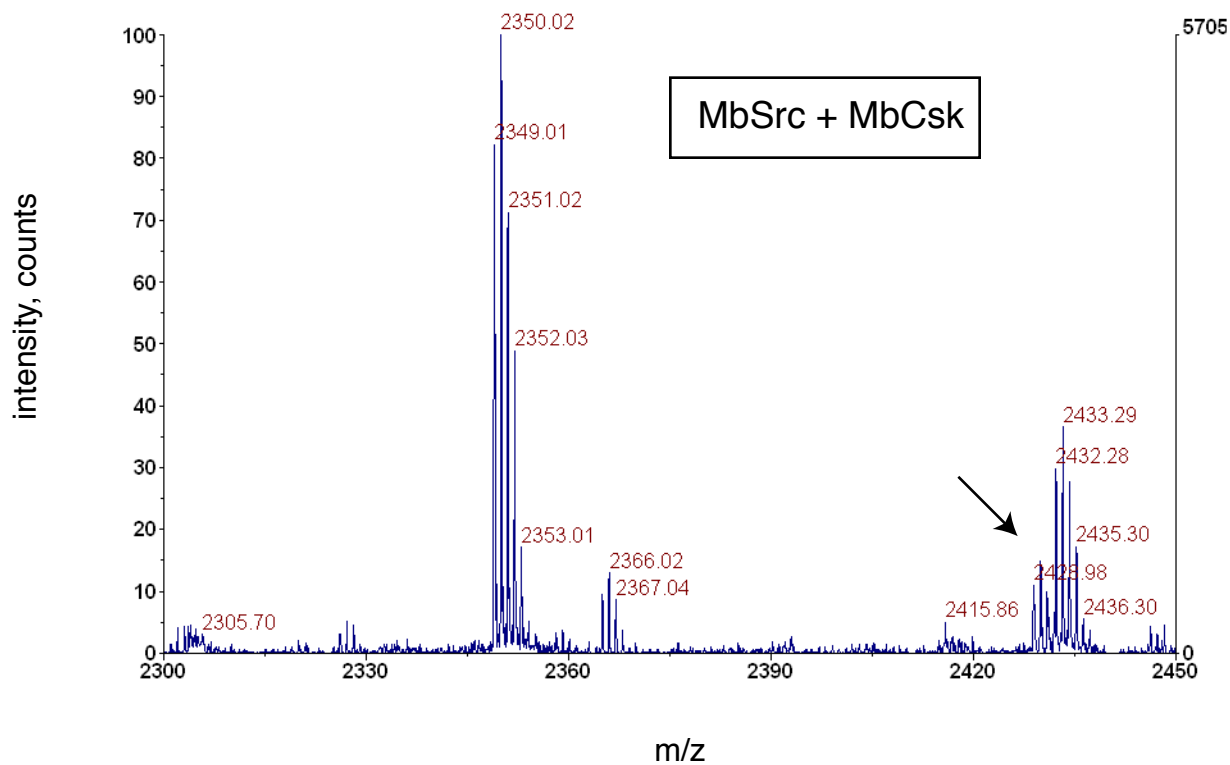
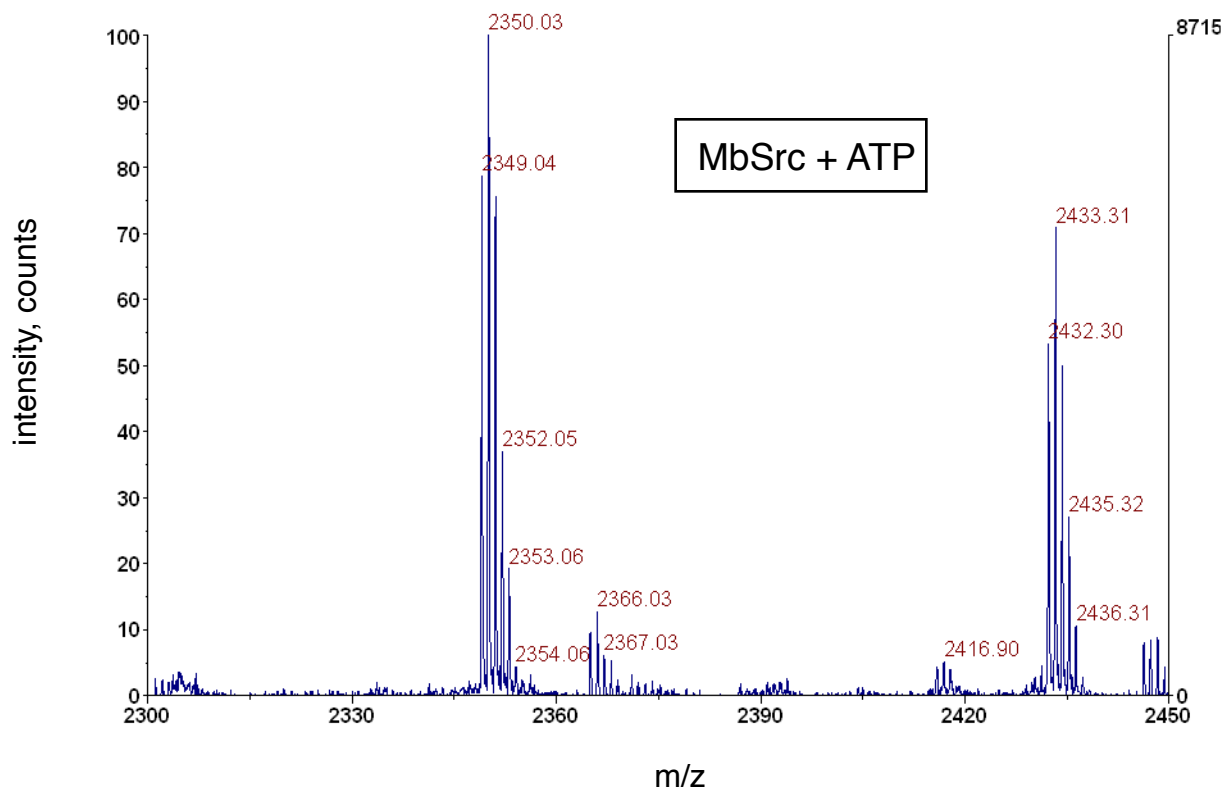
Supplemental Figure S4



Supplemental Figure S5

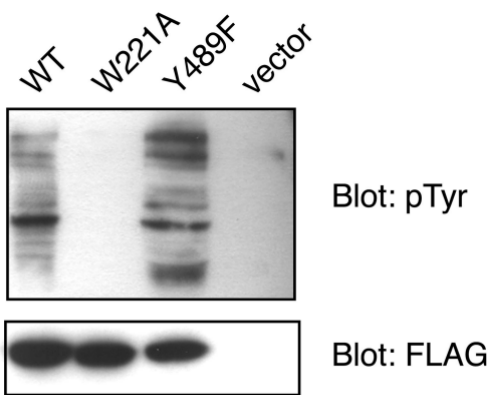
<b>Organism</b>	<b>% Identities</b>	<b>% Positives</b>
<i>H. sapiens</i>	51	66
<i>D. melanogaster</i>	52	68
<i>H. vulgaris</i>	45	66
<i>E. fluviatilis</i>	47	68
<i>M. ovata</i>	56	71

**Supplemental Figure S6. Identities with MbCsk.**



Supplemental Figure S7





Supplemental Figure S8