

Table S7. Accuracy, sensitivity, specificity and predictive value of hybridizations with the genomic sequences of Sheba and 162.0

Genome	Hybridization	zfp (No. of CDSs)			Accuracy
		>91	<91	Predictive Value	
Sheba	+	1263 (TP)	51 (FP)	Pos = 95.6%	93.3%
	-	58 (FN)	257 (TN)	Neg = 83.4%	
		Sensitivity = 95.6%	Specificity = 83.4%		
162.0	+	1417 (TP)	36 (FP)	Pos = 97.8%	95.8%
	-	32 (FN)	146 (TN)	Neg = 80.2%	
		Sensitivity = 97.8%	Specificity = 80.2%		

NOTE: TP, true positive; FP, false positive; TN, true negative; FN, false negative. Accuracy = (TN+TP)/(TN+TP+FN+FP).

Pos = Positive Predictive Value [TP/(TP+FP)]; Neg = Negative Predictive Value [TN/(FN+TN)].

Sensitivity [TP/(TP+FN)]; Specificity [TN/(FP+TN)]. Hybridization = observed hybridization with microarray.