









































































































A

SV	Location (Mb)	SVRE	GASVPro	SVDetect	Pindel	breseq	DELLY
phage inversion	0.91	 		 	 	 	 
<i>hyxS</i> inversion	0.35						
phage del/dup	1.2				 		
phage dup	1.6	 		 	 	 	 
phage inversion	2.1	 			 		
phage inversion	2.9	 		 		 	 
<i>fimS</i> inversion	4.9	 	 	 	 	 	 
phage del/dup	5.0						 

 *fimB*  *fimX*

B

SV	Location (Mb)	SVRE	GASVPro	SVDetect	Pindel	breseq	DELLY
<i>hyxS</i> inversion	0.41						
phage del/dup	0.9	   			    		   
<i>ipuS</i> inversion	2.75	    		  		 	   
<i>fimS</i> inversion	5.1	    		     	   	    	   

 *fimB*  *ipuA*  *fimX*
 *fimE*  *ipuB*

Figure S3. Comparison of SVRE calls to that of other SV prediction programs. SV predictions for (A) UTI89 and (B) CFT073 are listed in the first columns of each table. Whether that SV was detected in a given sample by a program is indicated by a filled box following the color code indicated in the legend.