

Table S1. Results of ANOVA showing no significant differences in the percentage cover of *Chthamalus stellatus* and *Rissoella verruculosa* in, respectively, five patches (50 × 50 cm) and plots (8 × 8 cm) allocated to each combination of levels of treatments *Chthamalus* (unmanipulated vs. removed vs. killed) and *Rissoella* (unmanipulated vs. removed), sampled before the application of any experimental manipulation.

Source of variation	<i>C. stellatus</i>				<i>R. verruculosa</i>		
	df	MS	<i>F</i>	p	MS	<i>F</i>	p
<i>Chthamalus</i>	2	110.53	0.77	>0.4	483.70	1.63	>0.2
<i>Rissoella</i>	1	80.03	0.55	>0.4	1.63	0.01	>0.9
<i>Chthamalus</i> × <i>Rissoella</i>	2	212.13	1.47	>0.2	161.03	0.59	>0.5
Residual	24	144.33			296.13		
Cochran's test		<i>C</i> = 0.40			<i>C</i> = 0.31		
Transformation		None			None		

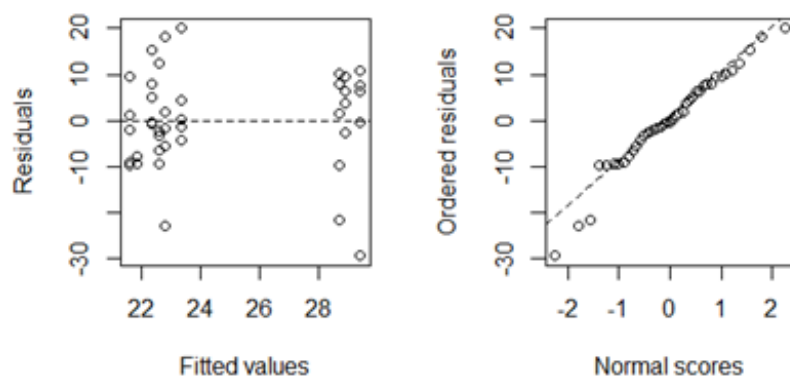


Figure S1. Fitted values vs. residuals and normal scores vs. ordered residuals for the analysis on frond size of *Rissoella verruculosa*.

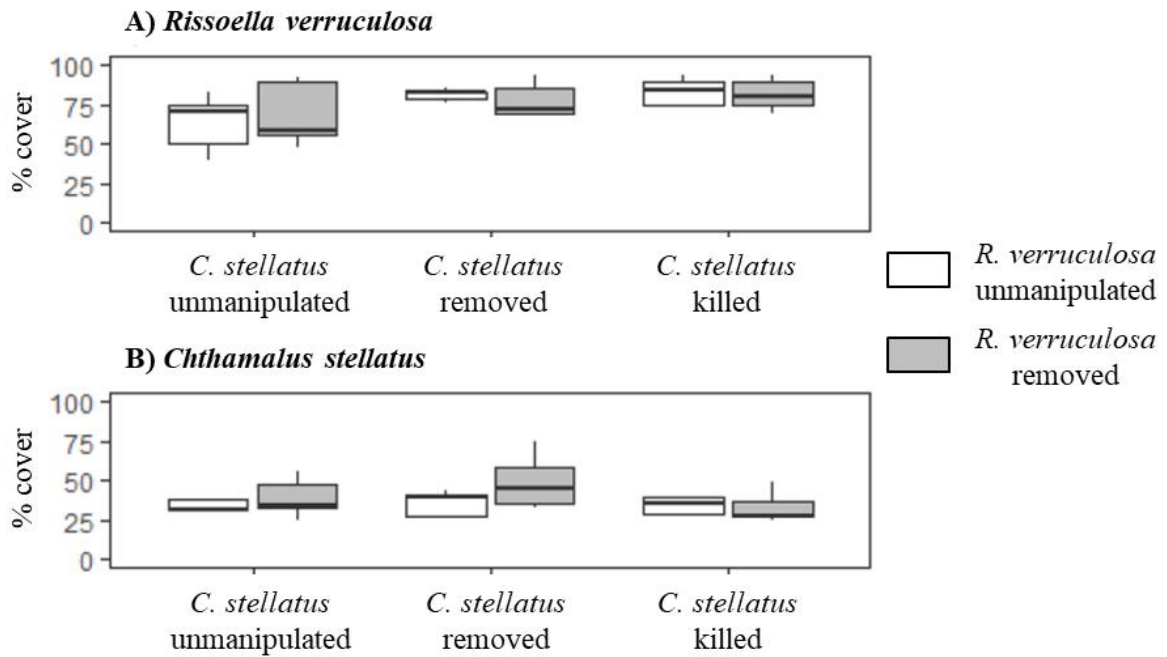


Figure S2. Mean percentage cover ($n = 5$) of (A) *R. verruculosa* and (B) *C. stellatus* in each combination of levels of *Chthamalus* and *Rissoella* treatments, sampled before the application of any experimental manipulation. Boxes span the interquartile range, with the horizontal black line within each box indicating the median value.