

<b>Supplementary Table 21 additional</b>		
<b>All binding sites at &gt; 66% knockdown</b>		
<b>Length</b>	<b>Not significantly downregulated</b>	<b>Significantly downregulated</b>
<b>Short</b>	28	18
<b>Long</b>	74	80
$\chi^2(1, N=200) = 1.84, p = .17$		
<b>Single binding site at &gt; 66% knockdown</b>		
<b>Length</b>	<b>Not significantly downregulated</b>	<b>Significantly downregulated</b>
<b>Short</b>	27	13
<b>Long</b>	51	50
$\chi^2(1, N=141) = 2.7, p = .10$		
<b>Multiple binding sites at &gt; 66% knockdown</b>		
<b>Length</b>	<b>Not significantly downregulated</b>	<b>Significantly downregulated</b>
<b>Short</b>	1	5
<b>Long</b>	23	30
$\chi^2(1, N=59) = 0.68, p = .41$		
Fisher's Exact Test for Count Data, $p = .39$		

Supplementary Table 12 investigation of genes showing a degree of knockdown of more than 66%. The number of genes that have one or more (all binding sites), one (single binding site), two or more (multiple binding sites) in a short or long gene and are either not significantly downregulated or significantly downregulated is shown. The numbers are too small to draw any meaningful conclusion.