

Table S1

Construct	Length (bp)	%SG accumulation	Predicted number of protein binding sites
FL NORAD	5287	70.9	51
NORAD-PREmut	5287	50	33 (assuming SAM68 sites not affected)
1/2 NORAD	2644	61.8	24
2/2 NORAD	2643	58.9	29
1/4 NORAD	1322	42.6	9
2/4 NORAD	1322	56	17
3/4 NORAD	1322	50	22
4/4 NORAD	1321	33.2	9
1/8 NORAD	661	20.9	2
2/8 NORAD	661	38.2	9
3/8 NORAD	661	35.8	7
4/8 NORAD	661	57.5	12
5/8 NORAD	661	41.3	15
6/8 NORAD	661	31.8	8
7/8 NORAD	661	34.8	6
8/8 NORAD	660	45.6	5
2/2 asNORAD	2643	25	8
1/4 asNORAD	1322	20.7	4
2/4 asNORAD	1322	26.2	5
3/4 asNORAD	1322	45.4	3
4/4 asNORAD	1321	56.25	7
1/8 asNORAD	661	20	2
2/8 asNORAD	661	27.45	4
3/8 asNORAD	661	33.3	4
4/8 asNORAD	661	21.2	3
5/8 asNORAD	661	29.8	3
6/8 asNORAD	661	30.4	2
7/8 asNORAD	661	21.1	4
8/8 asNORAD	660	32	5
660bp Luc	660	20.2	3
1300 bp Luc	1300	24.2	4
FL Luc	1635	21.1	4
2xLuc	3340	23.25	6
3xLuc	5045	30.5	8

*Assuming Luciferase has 2 protein binding sites