

Supplementary Table S2. *P1* transcripts with wild type and mutant base pairs in the absence and presence of CCC.

Base pair	-CCC	+CCC	Base pair	-CCC	+CCC	Base pair	-CCC	+CCC
+1A	1.00	13.85	-9C	1.08	13.29	-18G	0.89	11.24
+1G	1.09	16.22	-9T	1.01	12.29	-18C	0.62	12.76
+1C	0.70	3.86	-9A	5.28	21.52	-18A	1.27	13.63
+1T	0.74	7.60	-10G	0.35	11.70	-19T	3.11	14.90
-1G	0.60	4.94	-10C	0.21	11.07	-19A	2.13	14.86
-1A	2.29	16.64	-10A	0.67	12.83	-19C	1.46	15.20
-1T	1.36	13.86	-11G	0.09	0.30	-20G	1.23	12.94
-2G	1.17	9.56	-11C	0.03	0.17	-20C	1.51	13.45
-2C	1.15	12.49	-11T	0.08	0.32	-20A	1.79	12.35
-2A	1.30	12.50	-12G	0.13	5.18	-21G	0.98	29.83
-3G	1.48	10.39	-12C	0.13	4.15	-21C	1.04	36.02
-3C	1.17	10.41	-12A	0.10	4.34	-21A	1.15	34.14
-3A	1.08	6.87	-13G	1.49	15.39	-22G	1.25	18.24
-4G	1.17	17.35	-13T	1.32	17.20	-22C	1.15	28.46
-4C	1.24	12.93	-13A	1.29	18.24	-22A	1.01	21.53
-4A	1.45	20.02	-14A	0.38	6.40	-23G	1.03	26.41
-5G	0.99	12.75	-14C	0.32	3.68	-23A	0.91	19.53
-5C	0.78	10.86	-14T	0.32	8.20	-23T	0.91	24.12
-5T	0.80	13.41	-15G	0.13	4.96	-24G	0.83	13.70
-6G	0.80	18.11	-15C	0.33	9.44	-24A	0.70	25.77
-6C	1.01	19.08	-15A	0.37	9.96	-24C	0.77	21.57
-6A	1.18	12.98	-16G	0.89	17.08	-25G	1.52	26.58
-7G	0.20	4.03	-16C	0.47	13.28	-25C	1.65	14.81
-7C	0.25	3.32	-16T	0.28	9.57	-25T	1.53	28.23
-7A	0.34	6.56	-17G	0.25	10.56			
-8C	3.74	11.95	-17C	0.45	10.15			
-8T	1.29	30.36	-17A	0.22	7.17			
-8A	6.65	21.92						