

Codon no.	WT	Mutant 1	Mutant 2	Mutant 3	Mutant 4	Mutant 5	Mutant 6	Mutant 7	No. of models
	Off site deviation	153(21.223)	153(16.534)	153(21.223)	WT	148(16.121)	148(16.121)	153(16.534)	
25	UGG	CGG	CAG	CGA	CAA	UAG	UGA	UAA	
	W	R	Q	R	Q	STOP	STOP	STOP	3
	Deviationat mutant site	1.065	0.093	1.065	0.093	-	-	-	
	Off site deviation	153(18.31)	136(0.183)	153(18.31)	136(0.183)	21(0.514)	21(0.514)	21(0.514)	
26	AAC	GCG	GAC	AGC	AAU	GAU	AGU	GGU	
	N	G	D	S	N	D	S	G	3
	Deviationat mutant site	0.041	0.056	0.037	WT	0.056	0.037	0.041	
	Off site deviation	27(0.056)	<0.056	128(0.159)	WT	<0.056	128(0.159)	27(0.056)	
27	AGG	GAA	GAG	GGG	AAA	AGA	AAQ	GGA	
	R	E	E	G	K	R	K	G	3
	Deviationat mutant site	0.077	0.077	0.265	0.042	WT	0.042	0.265	
	Off site deviation	700(128)	700(128)	<0.265	141(0.179)	WT	141(0.179)	<0.265	
28	AAG	GAG	AGG	AAA	GGG	AGA	GAA	GGG	
	K	E	R	K	G	R	E	G	3
	Deviationat mutant site	0.036	0.045	WT	0.127	0.045	0.036	0.127	
	Off site deviation	141(0.213)	141(0.207)	WT	128(0.166)	141(0.207)	141(0.213)	128(0.166)	
29	AGA	AAG	AAA	GAA	GGG	GGA	AGG	GAG	
	R	K	K	E	G	G	R	E	3
	Deviationat mutant site	0.051	0.051	0.028	0.055	0.055	WT	0.028	
	Off site deviation	<0.051	<0.051	310(151)	128(0.128)	128(0.128)	WT	310(151)	
30	AUC	GUC	GCC	GUU	ACC	AUU	ACU	GCU	
	I	V	A	V	T	I	T	A	3
	Deviationat mutant site	0.047	0.058	0.047	0.059	WT	0.059	0.058	
	Off site deviation	<0.047	29(0.059)	<0.047	141(0.211)	WT	141(0.211)	29(0.059)	
31	AGC	GAC	AAC	GGC	AGU	GAU	AAU	GGU	
	S	D	N	G	S	D	N	G	3
	Deviationat mutant site	0.157	0.051	0.042	WT	0.157	0.051	0.042	
	Off site deviation	<0.157	<0.051	310(151)	128(0.128)	128(0.128)	WT	310(151)	
32	AAC	GGC	GAC	AGC	AAU	GAU	AGU	GGU	
	N	G	D	S	N	D	S	G	3
	Deviationat mutant site	0.033	0.05	0.037	WT	0.05	0.037	0.033	
	Off site deviation	128(0.109)	<0.05	141(0.204)	WT	<0.05	141(0.204)	128(0.109)	
33	UGU	UAU	CGU	CGC	UGC	CAU	CAC	UAC	
	C	Y	R	R	C	H	H	Y	3
	Deviationat mutant site	0.478	0.17	0.17	WT	0.188	0.188	0.478	
	Off site deviation	<0.478	6(0.254)	6(0.254)	WT	<0.188	<0.188	<0.478	
34	GUU	AUU	ACC	AUC	ACU	GCC	GUC	GCU	
	V	I	T	I	T	A	V	A	3
	Deviationat mutant site	0.032	0.03	0.032	0.03	0.047	WT	0.047	
	Off site deviation	4(0.088)	4(0.039)	4(0.088)	4(0.039)	141(0.185)	WT	141(0.185)	
35	CCU	AUC	ACC	AUU	GCC	GUC	GUU	ACU	
	A	I	T	I	A	V	V	T	3
	Deviationat mutant site	0.187	0.18	0.187	WT	0.256	0.256	0.18	
	Off site deviation	196(0.467)	<0.18	196(0.467)	WT	<0.256	<0.256	<0.18	
36	GAU	AAU	GGU	AGU	GAC	AAC	GGC	AGC	
	D	N	G	S	D	N	G	S	3
	Deviationat mutant site	0.041	0.129	0.075	WT	0.041	0.129	0.075	
	Off site deviation	141(0.209)	<0.129	198(0.158)	WT	141(0.209)	<0.129	198(0.158)	
37	UUU	UAU	CAU	UAC	CAC	CGU	UGC	CGC	
	Y	C	H	Y	H	R	C	R	3
	Deviationat mutant site	0.063	0.048	WT	0.048	0.072	0.063	0.072	
	Off site deviation	141(0.206)	141(0.16)	WT	141(0.16)	35(0.127)	141(0.206)	35(0.127)	
38	UCU	CCU	CUU	CCC	UCC	UUC	UUU	CUC	
	S	P	L	P	S	F	F	L	3
	Deviationat mutant site	0.251	0.094	0.251	WT	0.139	0.139	0.094	
	Off site deviation	36(0.314)	<0.094	36(0.314)	WT	36(0.195)	36(0.195)	<0.094	
39	GUC	AUC	ACC	AUU	ACU	GCC	GCU	GUU	
	V	I	T	I	T	A	A	V	3
	Deviationat mutant site	0.043	0.013	0.043	0.013	0.031	0.031	WT	
	Off site deviation	<0.043	141(0.209)	<0.043	141(0.209)	141(0.206)	141(0.206)	WT	
40	CUA	CUG	UUA	UCG	UCA	CCG	CCA	UUG	
	L	L	S	S	P	P	P	L	2
	Deviationat mutant site	WT	WT	0.02	0.02	0.118	0.118	WT	
	Off site deviation	WT	WT	38(0.067)	38(0.067)	<0.118	<0.118	WT	
41	UAU	UGU	CAU	UAC	CAC	CGU	UGC	CGC	
	Y	C	H	Y	H	R	C	R	3
	Deviationat mutant site	0.039	0.055	WT	0.055	0.019	0.039	0.019	
	Off site deviation	94(0.044)	<0.055	WT	<0.055	57(0.125)	94(0.044)	57(0.125)	
42	AAU	AGU	GAU	GGU	AGC	GAC	GGC	AAC	
	N	S	D	G	S	D	G	N	3
	Deviationat mutant site	0.088	0.05	0.058	0.088	0.05	0.058	WT	
	Off site deviation	128(0.175)	38(0.085)	<0.058	128(0.175)	38(0.085)	<0.058	WT	
43	UCC	CUU	CCU	CCC	UUC	UUU	CUU	UCU	
	S	L	P	P	F	L	S	3	
	Deviationat mutant site	0.042	0.455	0.455	0.028	0.028	0.042	WT	
	Off site deviation	45(0.214)	44(0.455)	44(0.453)	141(0.161)	141(0.161)	45(0.214)	WT	
44	GCA	GAC	ACA	GCG	GCA	GUU	AUA	GGU	
	A	D	I	A	A	V	I	V	4
	Deviationat mutant site	0.081	0.063	WT	WT	0.055	0.026	0.055	
	Off site deviation	<0.081	141(0.204)	WT	WT	<0.055	141(0.039)	<0.055	
45	UCA	UCG	CCA	CUA	UUA	CGG	CGU	UUG	
	S	P	L	P	F	L	S	3	
	Deviationat mutant site	WT	0.314	0.054	0.054	0.314	0.054	0.054	
	Off site deviation	WT	<0.314	<0.054	<0.054	<0.314	<0.054	<0.054	
46	UUU	CUU	UUU	UCU	CUU	UUC	UUC	CUU	
	F	L	S	P	P	S	L	P	3
	Deviationat mutant site	0.093	0.099	WT	0.249	0.099	0.093	0.249	
	Off site deviation	128(0.136)	<0.099	WT	<0.249	<0.099	128(0.136)	<0.249	
47	UCC	CUU	CCU	CCC	UUC	UUU	CUU	UCU	
	S	L	P	P	F	L	S	3	
	Deviationat mutant site	0.059	0.567	0.567	0.077	0.077	0.059	WT	
	Off site deviation	128(0.14)	<0.567	<0.567	141(0.14)	141(0.14)	128(0.14)	WT	
48	ACU	GCU	GCC	GUU	GUU	ACC	AUJ	AUC	
	T	A	A	V	V	T	I	I	3

Codon no.	WT	Mutant 1	Mutant 2	Mutant 3	Mutant 4	Mutant 5	Mutant 6	Mutant 7	No. of models
	F	L	S	F	P	S	L	P	3
73	GUA	AUA	GUU	AUG	GCA	ACA	CGC	ACG	
74	AUU	V	I	V	M	A	T	A	4
75	AGA	AAG	AAA	GAA	GGG	GGA	AGG	GAG	
76	GGU	GCG	GAC	AGC	AAC	GAU	AGU	AAU	
77	GAU	AAU	GGU	AGU	GAC	AAC	GGC	AGC	
78	GAA	GAG	AAA	GGG	AGG	AAG	AGA	AGG	
79	GUC	AUC	ACC	AUJ	ACU	GCC	GCU	GUU	
80	AGA	AAG	AAA	GAA	GGG	GGA	AGG	GAG	
81	CAA	CAG	CGG	UGA	UAG	UGG	UAA		
82	Q	R	Q	R	Stop	Stop	W	Stop	3
83	GGG	AAA	GAG	GAA	AGG	AAG	GGA	AGA	
84	CCA	CCG	UUA	UCA	UUG	UCG	CUG		
85	P	P	L	L	S	L	S	L	2
86	CAA	CGA	CAG	CGG	UGA	UAG	UGG	UAA	
87	Q	R	Q	R	Stop	Stop	W	Stop	3
88	GGA	GAA	GGG	AAA	GGA	AAG	AGG	GAG	
89	K	E	R	E	R	K	G	R	3
90	AUU	I	V	I	T	A	A	V	3
91	AUU	A	A	V	V	T	I	I	3
92	GAU	AAC	GGU	AGU	GAC	AAC	GGC	AGC	
93	UAU	Y	C	H	A	U	U	C	3
94	AAG	AAU	UAU	GAA	AGU	GAC	GGC	AGC	
95	UAU	Y	C	H	A	U	U	C	3
96	UAA	UUA	UAG	UAA	UAC	UAC	UAU	CGC	
97	UAG	UAC	UUC	UAA	UUG	UUC	UGC	CGC	
98	UUG	UAC	UUA	UAA	UUC	UUA	UGC	CGC	
99	UUA	UUC	UUC	UAA	UUA	UUA	UGC	CGC	

Codon no.	WT	Mutant 1	Mutant 2	Mutant 3	Mutant 4	Mutant 5	Mutant 6	Mutant 7	No. of models
120	Off site deviation	WT	154 (18.052)	154 (18.025)	155 (14.677)	154 (18.052)	154 (18.025)	155 (14.677)	
	AAU	AGU	GAU	GGU	AGC	GAC	GGC	AAC	
	N	S	D	G	S	D	G	N	3
	Deviation at mutant site	1.341	1.354	1.302	1.341	1.354	1.302	WT	
121	Off site deviation	WT	154 (18.095)	153 (18.304)	154 (18.11)	154 (18.095)	153 (18.304)	WT	
	UAU	UGU	CAU	UAC	CAC	CGU	UGC	CGC	
	Y	C	H	Y	H	R	C	R	3
	Deviation at mutant site	0.656	0.639	WT	0.639	0.594	0.656	0.594	
122	Off site deviation	WT	151 (9.284)	156 (17.552)	WT	156 (17.552)	156 (17.557)	151 (9.284)	156 (17.557)
	AAU	AGU	GAU	GGU	AGC	GAC	GGC	AAC	
	N	S	D	G	S	D	G	N	3
	Deviation at mutant site	2.004	0.955	2.431	2.004	0.955	2.431	WT	
123	Off site deviation	WT	146 (13.423)	154 (17.931)	148 (21.328)	146 (13.423)	154 (17.931)	148 (21.328)	WT
	UAC	UGC	CGC	CGU	UGU	CAC	CAU	UAU	
	Y	C	R	R	C	H	H	Y	3
	Deviation at mutant site	1.364	0.39	0.39	1.364	1.068	1.068	WT	
124	Off site deviation	WT	154 (20.422)	153 (17.029)	153 (17.029)	154 (20.422)	154 (18.114)	154 (18.114)	WT
	CUG	CUA	UCG	UCA	UUA	CCG	CCA	UUG	
	L	L	S	S	L	P	P	L	2
	Deviation at mutant site	WT	0.494	0.494	WT	0.1779	0.1779	WT	
125	Off site deviation	WT	155 (16.581)	155 (16.581)	WT	154 (17.589)	154 (17.589)	WT	
	UAU	UGU	CAU	UAC	CAC	CGU	UGC	CGC	
	Y	C	H	Y	H	R	C	R	3
	Deviation at mutant site	1.106	2.378	WT	2.378	0.373	1.106	0.373	
126	Off site deviation	WT	156 (18.996)	153 (21.877)	WT	153 (21.877)	155 (16.501)	156 (18.996)	155 (16.501)
	AGA	AAG	AAA	GAA	GGG	GGA	AGG	GAG	
	R	K	K	E	G	G	R	E	3
	Deviation at mutant site	0.972	0.972	0.714	1.116	1.116	WT	0.714	
127	Off site deviation	WT	151 (9.576)	151 (9.576)	147 (16.486)	154 (16.938)	154 (16.938)	WT	147 (16.486)
	UUG	UUA	CUU	UCG	CCG	CUA	UCA	CCA	
	L	L	S	P	L	S	P	2	
	Deviation at mutant site	WT	WT	1.637	2.732	WT	1.637	2.732	
128	Off site deviation	WT	WT	153 (19.069)	153 (16.857)	WT	153 (19.069)	153 (16.857)	
	UUU	CUU	UCU	UUC	CCU	UCC	UCU	CCC	
	F	L	S	F	P	S	L	P	3
	Deviation at mutant site	1.893	1.668	WT	1.848	1.668	1.893	1.848	
129	Off site deviation	WT	154 (20.067)	154 (19.026)	WT	155 (16.196)	154 (19.026)	154 (20.067)	155 (16.196)
	AGG	GAA	GAG	GGG	AAA	AGA	AAG	GGG	
	R	E	E	G	K	R	K	G	3
	Deviation at mutant site	1.66	1.66	1.463	1.944	WT	1.944	1.463	
130	Off site deviation	WT	146 (17.839)	146 (17.839)	153 (18.203)	153 (15.732)	WT	153 (15.732)	153 (18.203)
	AAG	GAG	AGG	AAA	GGG	AGA	GAA	GGG	
	K	E	R	K	G	R	E	G	3
	Deviation at mutant site	1.759	2.598	WT	1.321	2.598	1.759	1.321	
131	Off site deviation	WT	153(17.601)	154(16.458)	WT	153(16.264)	154(16.458)	153(17.601)	153(16.264)
	UCU	CCU	CUU	CCC	UCC	UUC	UUU	CUC	
	S	P	L	P	S	F	F	L	3
	Deviation at mutant site	3.49	2.482	3.49	WT	1.009	1.003	2.482	
132	Off site deviation	WT	153(19.302)	154(22.587)	153(19.302)	WT	153(14.798)	153(14.798)	154(22.587)
	AAU	AGU	GAU	GGU	AGC	GAC	GGC	AAC	
	N	S	D	G	S	D	G	N	3
	Deviation at mutant site	0.685	1.495	2.162	0.685	1.495	2.162	WT	
133	Off site deviation	WT	156(17.704)	156(13.419)	148(13.782)	156(17.704)	156(13.419)	148(13.782)	WT
	CUC	UUC	CCC	CCU	CUU	UCU	UUU	UCC	
	L	F	P	P	L	S	F	S	3
	Deviation at mutant site	1.765	2.126	1.216	WT	1.554	1.765	1.554	
134	Off site deviation	WT	145(13.622)	148(15.246)	148(15.246)	WT	156(15.787)	145(13.622)	156(15.787)
	AAA	GAA	AGA	AAG	GGG	AGG	GAG	GGG	
	K	E	R	K	G	R	E	G	3
	Deviation at mutant site	1.828	0.087	WT	0.374	0.087	1.828	0.374	
135	Off site deviation	WT	159(20.81)	132(0.139)	WT	151(11.308)	132(0.139)	155(20.81)	151(11.308)
	CCU	UCU	UUU	UUC	CUU	CCC	CUC	UCC	
	P	S	F	F	L	P	L	S	3
	Deviation at mutant site	0.076	1.534	1.534	1.524	WT	1.524	0.076	
136	Off site deviation	WT	141(0.205)	153(16.156)	153(16.156)	WT	153(16.147)	WT	153(16.147)
	UUU	CUU	UCU	UUC	CCU	UCC	CUC	CCC	
	F	L	S	F	P	S	L	P	3
	Deviation at mutant site	0.028	1.158	WT	0.192	1.158	0.028	0.192	
137	Off site deviation	WT	98(0.333)	153(18.309)	WT	<0.192	153(18.309)	98(0.333)	<0.192
	GAG	AGA	AAG	GAA	GGG	AGG	GGA	AAA	
	E	R	K	E	G	R	G	K	3
	Deviation at mutant site	0.046	0.129	WT	1.623	0.046	1.623	0.129	
138	Off site deviation	WT	92(0.642)	92(0.614)	WT	153(18.934)	92(0.642)	153(18.934)	92(0.614)
	AGA	AAG	AAA	GAA	GGG	AGG	GAG	GGG	
	R	K	K	E	G	R	E	G	3
	Deviation at mutant site	2.037	2.037	1.515	1.341	1.341	WT	1.515	
139	Off site deviation	WT	153(21.476)	153(21.476)	153(20.243)	153(20.162)	153(20.162)	WT	153(20.243)
	GAU	AAU	GGU	AGU	GAC	AAC	GGC	AGC	
	D	N	G	S	D	N	G	S	3
	Deviation at mutant site	1.608	0.631	1.153	WT	1.608	0.631	1.153	
140	Off site deviation	WT	155(15.023)	148(16.056)	147(15.585)	WT	155(15.023)	148(16.056)	147(15.585)
	AUU	GUU	ACC	AUC	ACU	GCU	GCC	GUC	
	V	T	I	T	A	V	I	V	3
	Deviation at mutant site	2.457	1.78	WT	1.76	0.658	0.658	2.457	
141	Off site deviation	WT	153(17.811)	154(20.4)	WT	154(20.4)	156(17.197)	156(17.197)	153(17.811)
	UUA	UUA	CUA	CUA	UUA	CGU	CUU	UUG	
	S	P	L	L	P	L	2		
	Deviation at mutant site	3.984	4.598	4.598	3.984	4.598	4.598		
142	Off site deviation	WT	153(20.687)	153(17.333)	153(17.333)	WT	153(20.887)	153(17.333)	153(17.333)
	ACU	GUU	GCC	GUU	GUC	ACC	AUC	AUC	
	T	A	V	V	T	I	A	I	3
	Deviation at mutant site	4.428	4.428	4.982	4.982	WT	1.694	1.694	
143	Off site deviation	WT	156(13.201)	156(13.201)	156(2.374)	156(2.374)	WT	153(18.596)	153(9.596)
	GAA	GAG	AAA	GGA	GGG	AAG	AGA	AGG	
	E	E	K	G	K	R	R	3	

Codon no.	WT	Mutant 1	Mutant 2	Mutant 3	Mutant 4	Mutant 5	Mutant 6	Mutant 7	No. of models
	Y	C	H	Y	H	R	C	R	3
Deviation at mutant site	1.604	1.419	WT	WT	1.419	0.053	1.604	0.053	
Off site deviation	(155) 15.746	(153) 18.403	WT	(153) 18.403	128(0.156)	(155) 15.746	128(0.156)		
168	GGU	GAC	AGC	AAC	GAU	AGU	AAU		
	G	G	D	S	N	D	S	N	3
Deviation at mutant site	WT	0.197	0.238	0.177	<0.197	0.197	0.238	0.177	
Off site deviation	WT	<0.197	<0.238	<0.177	<0.197	<0.238	<0.177		
169	UUC	UCU	CUU	UUU	CCC	CUC	UCC		
	F	P	S	L	F	P	L	S	3
Deviation at mutant site	0.285	0.304	0.308	WT	0.285	0.308	0.304		
Off site deviation	<0.285	118 (4.594)	118 (4.555)	WT	<0.285	118 (4.555)	118 (4.594)		
170	CAA	CGA	CAG	CGG	UGA	UAG	UGG	UAA	
	Q	R	Q	R	Stop	Stop	W	Stop	3
Deviation at mutant site	0.022	WT	0.022	-	-	0.009	-		
Off site deviation	173 (0.057)	WT	173 (0.057)	117 (3.286)	117 (3.286)	174 (0.213)	117 (3.286)		
171	CCC	UUU	UCC	UUC	CCU	CUC	UCU		
	P	F	S	F	P	L	L	S	3
Deviation at mutant site	0.101	0.065	0.101	WT	0.187	0.187	0.065		
Off site deviation	173(0.676)	170(0.178)	173(0.676)	WT	173(0.651)	173(0.651)	170(0.178)		
172	ACU	GCU	GCC	GUU	GUC	ACC	AUU	AUC	
	T	A	A	V	V	T	I	I	3
Deviation at mutant site	2.593	2.593	0.183	0.183	WT	0.341	0.341		
Off site deviation	<2.593	<2.593	174(0.568)	174(0.568)	WT	174(0.823)	174(0.823)		
173	AAU	AGU	GAU	GGU	AGC	GAC	GGC	AAC	
	N	S	D	G	S	D	G	N	3
Deviation at mutant site	0.092	0.056	0.655	0.092	0.056	0.655	0.655	WT	
Off site deviation	128(0.149)	168(0.153)	175(0.469)	128(0.149)	168(0.153)	175(1.469)	WT		
174	GGU	GGC	GAC	AGC	AAC	GAU	AGU	AAU	
	G	G	D	S	N	D	S	N	3
Deviation at mutant site	WT	1.692	1.779	0.71	1.692	1.779	0.71		
Off site deviation	WT	<1.692	<1.779	<0.71	<1.692	<1.779	<0.71		
175	GUU	AUU	ACC	AUC	ACU	GCC	GUU	GUU	
	V	I	T	I	T	A	V	A	3
Deviation at mutant site	0.157	0.198	0.157	0.198	0.163	WT	0.163		
Off site deviation	173(0.554)	174(0.633)	173(0.554)	174(0.633)	174(0.425)	WT	174(0.425)		
176	GGU	GGC	GAC	AGC	AAC	GAU	AGU	AAU	
	G	G	D	S	N	D	S	N	3
Deviation at mutant site	WT	0.344	0.306	0.344	0.344	0.306	0.344		
Off site deviation	WT	174(0.439)	174(0.412)	174(0.423)	174(0.439)	174(0.412)	174(0.423)		
177	UAC	UGC	CGC	CGU	UGU	CAC	CAU	UAU	
	Y	C	R	C	H	H	Y	3	
Deviation at mutant site	0.045	0.079	0.045	0.045	0.032	0.032	0.032	WT	
Off site deviation	141(0.201)	170(0.153)	170(0.153)	141(0.201)	141(0.205)	141(0.205)	WT		
178	CAA	CGA	CAG	CGG	UGA	UAG	UGG	UAA	
	Q	R	Q	R	Stop	Stop	W	Stop	3
Deviation at mutant site	0.046	WT	0.046	-	-	0.129	-		
Off site deviation	128(0.121)	WT	128(0.121)	117(3.198)	117(3.198)	128(0.158)	117(3.198)		
179	CCA	CCG	UUU	CUA	UCU	UUG	UCG	CUG	
	P	P	L	L	S	L	S	L	2
Deviation at mutant site	WT	0.334	0.334	0.1	0.334	0.1	0.334		
Off site deviation	WT	73 (0.42)	73 (0.42)	<0.1	73 (0.42)	<0.1	73 (0.42)		
180	UAC	UGC	CGC	CGU	UGU	CAC	CAU	UAU	
	Y	C	R	R	C	H	H	Y	3
Deviation at mutant site	0.034	0.054	0.054	0.034	0.044	0.044	0.044	WT	
Off site deviation	76 (0.068)	109 (0.227)	109 (0.227)	76 (0.068)	141 (0.213)	141 (0.213)	WT		
181	AAG	AAA	GAA	GGG	GGG	AGG	AGG	GAG	
	R	K	K	E	G	G	R	E	3
Deviation at mutant site	0.065	0.065	0.068	0.096	0.096	0.096	WT	0.088	
Off site deviation	141 (0.212)	141 (0.212)	114 (0.202)	114 (0.16)	114 (0.16)	WT	114 (0.202)		
182	GUA	AUA	GUG	AUG	GCA	ACA	GCG	ACG	
	V	I	V	M	A	T	A	T	4
Deviation at mutant site	0.096	WT	0.164	0.037	0.07	0.037	0.07	0.07	
Off site deviation	82 (0.098)	WT	<0.164	141 (0.181)	199 (0.144)	141 (0.181)	199 (0.144)		
183	GUA	AUA	GUG	AUG	GCA	ACA	GCG	ACG	
	V	I	V	M	A	T	A	T	4
Deviation at mutant site	0.095	WT	0.214	0.014	0.077	0.014	0.077	0.077	
Off site deviation	71, 128 (0.14)	WT	13 (0.225)	141 (0.206)	<0.077	141 (0.206)	<0.077		
184	GUA	AUA	GUG	AUG	GCA	ACA	GCG	ACG	
	V	I	V	M	A	T	A	T	4
Deviation at mutant site	0.084	WT	0.187	0.039	0.045	0.039	0.045		
Off site deviation	141 (0.203)	WT	184(0.187)	141 (0.185)	184 (0.045)	141 (0.185)	184 (0.045)		
185	CUC	CUC	CCC	UCC	UUU	UCU	UCU		
	L	L	P	S	F	S	F	3	
Deviation at mutant site	WT	0.183	0.045	0.045	0.193	0.045	0.193		
Off site deviation	WT	104 (0.195)	104 (0.195)	141 (0.173)	69 (0.21)	141 (0.173)	69 (0.21)		
186	UCU	CUU	CCC	UCC	UUC	UCU	UCU	CUC	
	S	P	L	S	F	F	L	3	
Deviation at mutant site	0.209	0.083	0.209	WT	0.052	0.052	0.083		
Off site deviation	<0.209	<0.083	<0.209	WT	<0.052	<0.052	<0.083		
187	UUU	CUU	UUC	CCU	UUU	UCU	UCU	CCC	
	F	L	S	F	S	L	F	3	
Deviation at mutant site	0.13	0.114	WT	0.374	0.314	0.13	0.374		
Off site deviation	103 (0.173)	114 (0.218)	WT	<0.374	114 (0.218)	103 (0.173)	<0.374		
188	GAA	GAG	AAA	GGG	GGG	AAG	AGA	AGG	
	E	K	S	G	K	R	R	3	
Deviation at mutant site	WT	0.087	0.276	0.276	0.067	0.067	0.087		
Off site deviation	WT	186 (0.075)	<0.276	<0.276	186 (0.075)	128 (0.126)	128 (0.126)		
189	CUU	CUC	CCC	CCU	CCU	UUC	UCU	UCU	
	L	P	S	F	S	F	S	3	
Deviation at mutant site	WT	0.149	0.149	0.107	0.106	0.107	0.106		
Off site deviation	WT	<0.149	<0.149	<0.107	<0.106	<0.107	<0.106		
190	CUA	CUG	UUA	UCG	UCA	CGG	CGA	UUG	
	L	L	S	P	P	P	L	2	
Deviation at mutant site	WT	WT	0.087	0.067	0.067	0.301	0.301	WT	
Off site deviation	WT	WT	<0.087	<0.067	<0.067	<0.301	<0.301	WT	

Codon no.	WT	Mutant 1	Mutant 2	Mutant 3	Mutant 4	Mutant 5	Mutant 6	Mutant 7	No. of models													
191	CAU	CGU	UAC	UAU	CAC	UGC	CGC	UGU														
	H	R	Y	Y	H	C	R	C	3													
Deviation at mutant site		0.057	0.052	0.052	WT	0.039	0.057	0.039														
Off site deviation		<0.057	<0.052	<0.052	WT	141 (0.179)	<0.057	141 (0.179)														
192	GCA	ACG	ACA	GCG	AUG	GUA	UAU	GUG														
	A	T	T	A	M	V	I	V	4													
Deviation at mutant site		0.224	0.224	WT	0.057	0.213	0.273	0.213														
Off site deviation		<0.224	<0.224	WT	141 (0.217)	<0.213	<0.273	<0.213														
193	CCA	CCG	UUA	CUA	UCA	UUG	UCG	CUG														
	P	P	L	L	S	L	S	L	2													
Deviation at mutant site		WT	0.052	0.052	0.056	0.052	0.058	0.052														
Off site deviation		WT	141 (0.215)	141 (0.215)	192 (0.073)	141 (0.215)	192 (0.073)	141 (0.215)														
194	GCA	ACG	ACA	GCG	AUG	GUA	UAU	GUG														
	A	T	T	A	M	V	I	V	4													
Deviation at mutant site		0.066	0.066	WT	0.016	0.044	0.197	0.044														
Off site deviation		128 (0.147)	128 (0.147)	WT	128 (0.2)	128 (0.133)	114 (0.316)	128 (0.133)														
195	ACU	CGU	GCC	GUU	GUU	ACC	AUU	AUC														
	T	A	A	V	V	T	I	I	3													
Deviation at mutant site		0.045	0.045	0.109	0.109	WT	0.249	0.249														
Off site deviation		128(0.131)	128(0.131)	128 (0.14)	128 (0.14)	WT	31 (0.266)	31 (0.266)														
196	GUU	AUU	ACC	AUC	ACU	GCC	GUC	GCU														
	V	I	T	I	T	A	V	A	3													
Deviation at mutant site		0.098	0.064	0.098	0.064	0.08	WT	0.08														
Off site deviation		66 (0.252)	141 (0.211)	66 (0.252)	141 (0.211)	141 (0.185)	WT	141 (0.185)														
197	UGU	UAU	CGU	CGC	UGC	CAU	CAC	UAC														
	C	Y	R	R	C	H	H	Y	3													
Deviation at mutant site		0.156	0.201	0.201	WT	0.101	0.101	0.156														
Off site deviation		141 (0.254)	141 (0.208)	141 (0.208)	WT	141 (0.237)	141 (0.237)	141 (0.254)														
198	GGA	GAA	GGG	AAA	AGA	AAG	AGG	GAG														
	G	E	G	K	R	K	R	E	3													
Deviation at mutant site		0.205	WT	0.222	0.471	0.222	0.471	0.205														
Off site deviation		60 (0.587)	WT	60 (0.773)	60 (0.927)	60 (0.773)	60 (0.927)	60 (0.587)														
199	CCU	UCU	UUU	UUC	CUU	CCC	CUC	UCC														
	P	S	F	F	L	P	L	S	3													
Deviation at mutant site		0.163	0.148	0.148	0.158	WT	0.158	0.163														
Off site deviation		<0.163	<0.148	<0.148	<0.158	WT	<0.158	<0.163														
200	AAA	GAA	AGA	AAG	GGG	AGG	GAG	GGG														
	K	E	R	K	G	R	E	G	3													
Deviation at mutant site		0.131	0.103	WT	-	0.103	0.131	-														
Off site deviation		<0.131	141 (0.22)	WT	141 (0.207)	141 (0.22)	<0.131	141 (0.207)														