		99N	998	00N	00S	01N	015	02N	02S	03N	035	04N	045	05N	05S	06N	06S	07N	07S	08N	085	09N	098	10N	10S	11N	11 S	12N	12 S	13N	135	14N	145	15N	15S
HA	G5V	0	0	0.828	0.284	0.264	0.152	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	K92T	0	0	0.828	0.284	0.264	0.152	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	D271N	0	0	0.824	0.284	0.264	0.152	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Q33H	0	0	0.808	0.284	0.092	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	R50G	0	0	0	0.6	0.58	0.236	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	S199P	0	0	0	0.6	0.58	0.236	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	P273S	0	0	0.084	0.6	0.58	0.248	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	S247C	0	0	0	0.58	0.58	0.236	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	A106V	0	0	0	0	0.092	0.312				0.02	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	N144D	0	0	0	0	0		0.928			0.02	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	S186G	0	0.072	0	0	0.092	0.312			0.232	0.02	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	R50G	0	0	0.008	0.096	0.064				0.768		0.956	1	0.976	0.968	0.5	0.16	0.28	0.056	0.004	0	0	0	0	0.008	0	0	0	0	0	0	0	0	0	0
	E83K	0	0	0	0	0	0.048			0.768	0.98	1	1	1	0.992	0.96	1	1	0.972	0.82	0.944	1	1	1	1	1	0.992		0.984	0.996	0.984	0.988	0.976	0.98	0.996
	A131T	0	0	0	0	0	0			0.768	0.98	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	0.976		1	1		0.996	1	0.992
	S186G	0	0	0	0	0	0			0.768	0.98	1	1	1	1	1	1	0.988	1	0.972	1	0.984	1	1	1	1	1	0.976	_	0.992		0.968		1	1
	V202I	0	0	0	0	0	0.048			0.768	0.98	1	1	1	1	1	1	1	1	1	1	0.992	1	1	1	1	1	0.976	1	1		0.996		0.992	1
	W222R G225D	0	0	0	0 006	0	0.048			0.768	0.98	1	1	0 0e T	1 0.426	1 0.094	7 T	0.013	7 T	0	7 T	1	1	1	U 000	1	1	0.976 0	1	1	1 0		0.996	0	1
	L251	0	0	0.008	0.096	0.064	0.288 0	0.052 0		0.768	0.98 0.98	1 1	0.936	0.96	0.436 0.952	0.084	0	0.012	0 1	0.984	1	1	0 0.992	0 744	0.008 0.992	0 0.984	1	0.972	0	0	1	0 0.984	0	1	0.98
	H75Q	0	0	0	0	0	0	0		0.744 0.744	0.98	0.988	0.930	0.992	0.952	0.990	1	1	0.992	0.964	1	1		0.744	0.992	0.964	1	0.972	0.992	1	0.992		0.996	1	0.96
	H155T	0	0	0	0	0	0	0		0.744	0.98	0.566	1	1	1	1	1	1	1	1	1	1	0.552	0.504	1	1	1	0.976	1	1	1		0.996	1	1
	Q156H	0	0	0	0	0	0	0	0.048	0.612	0.98	1	1	1	1	0.968	1	0.988	1	0.992	1	1	1	1	1	1	1	0.976	_	1	1	0.996		1	1
	N126D	0	0	0	0	0	0	0	0	0	0.756	0.624	0.008	0.004	0	0.500	0	0.500	0	0.332	0	0	0	0	0	0	0	0	0.552	0	0	0.550	0.550	0	0
	Y105H	0	0	0	0	0	0	0	0	0	0.68	0.008	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Y159F	0	0	0	0	0	0	0	0	0	0.052		0.976	0.984	1	1	1	1	1	0.988	1	1	0.992	1	0.992	1	1	0.976	0.992	1	1	0.744	0.26	0.132	0.012
	S189N	0	0	0	0	0	0	0	0	0	0.104		0.976		1	1	1	1	1	1	1	0.96	0.188	0.024		0	0	0	0	0	0	0	0	0	0
	K145N	0	0	0	0	0	0	0	0	0	0	0.004	0.876	0.968	0.62	0.992	1	0.984	1	1	1	0.972	1	1	1	0.976	0.368	0.244	0.36	0.104	0.004	0	0	0	0
	S227P	0	0	0	0	0	0	0	0	0	0.132	0.2	0.764	0.94	1	1	1	1	1	1	1	1	1	1	1	1	1	0.976	1	1	1	0.996	0.996	1	1
	V226I	0	0	0	0	0	0	0	0	0	0	0	0.664	0.904	1	1	1	1	1	1	1	1	1	1	1	1	1	0.976	1	1	1	0.996	0.996	1	1
	S193F	0	0	0	0	0	0	0	0	0	0	0	0	0.04	0.564	0.916	1	0.988	1	1	1	1	1	1	0.992	1	1	0.976	1	1	1	0.988	0.988	1	1
	D225N	0	0	0	0	0	0	0	0	0	0	0	0	0.04	0.564	0.856	0.84	0.96	1	0.964	0.756	0.86	0.992	1	0.96	0.996	0.992	0.944	0.96	0.988	0.992	0.684	0.208	0.132	0.012
	G50E	0	0	0	0	0	0	0	0	0	0	0	0	0.024	0.032	0.492	0.84	0.012	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	G50E	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.008	0	0.62	0.944	0.996	1	1	1	1	0.98	0.996	0.996	0.976	1	1	0.992	0.996	0.996	1	1

K140I	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.008	0	0.62	0.944	0.988	1	0.948	0.988	1	0.98	0.976	0.992	0.932	0.948	1	0.984	0.996	0.996	1	1
Q261R	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.008	0	0.58	0.912	0.984	0.964	0.984	0.94	0.848	0.96	0.988	0.972	0.976	1	1	1	0.964	0.98	0.904	0.884
K173Q	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.044	0.524	0.816	0.948	1	1	0.992	1	1	0.956	1	1	1	0.996	0.996	1	1
E62K	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.052	0.7	0.576	0.052	0.044	0.004	0	0	0	0	0	0	0	0
K158N	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.052	0.7	0.576	0.052	0.044	0.004	0	0	0	0	0	0	0	0
N189K	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.04	0.7	0.576	0.052	0.044	0.004	0	0	0	0	0	0	0	0
N144K	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.04	0.696	0.576	0.052	0.044	0.004	0	0	0	0	0	0	0	0
V213A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.008	0.564	0.128	0.016	0.032	0	0	0	0	0	0	0	0	0
P194L	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.008	0.56	0.128	0.016	0.032	0	0	0	0	0	0	0	0	0
K158N	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.112	0.4	0.94	0.956	0.996	0.976	1	1	1	0.944	0.996	1	1
N189K	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.112	0.4	0.94	0.956	0.996	0.976	1	1	1	0.996	0.996	1	1
T212A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.112	0.356	0.86	0.596	0.936	0.832	0.784	0.964	1	0.996	0.996	1	1
N45S	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.112	0.4	0.852	0.8	0.876	0.732	0.504	0.104	0.016	0	0.008	0	0
D53N	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.132				0.236				0	0	0	0
1230V	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.092	0.352	0.668	0.356	0.236	0.328	0.096	0.004	0	0	0	0
Y94H	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.044				0.236				0	0	0	0
E280A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.144	0.344			0.236				0	0	0	0
N312S	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.012	0.204	0.48	0.152	0.64	0.672		0.884	0.98		0.988	1	1
V223I	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			0.732						1	1
A198S	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.02		0.672							0.992
N145S	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			0.556					0.008	0	0
S45N	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.008	0.02	0.164					0.98	1	1
T48I	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.02					0.996		1	1
N278K	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.008	0.028					0.936		1	1
S157L	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.02								
N145S	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.004		0.836			0.972		0.992
Q33R	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.02					0.888			0.013
N53D T128A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		0.096					0.176	
	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.004		0.344	0.7	0.604		0.184	
R142G N225D	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.004	0.02	0.344	0.7			0.184 0.816	
	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			0.816	
Q311H L3I	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			0.816	
N144S	0	0	0	0	0	0	0	0	0	0	0	0	n	0	0	0	0	0	0	n	0	0	0	0	0	0	0	0	0	0			0.816	
F159Y	0	0	0	0	0	0	0	0	0	0	0	0	0	n	0	0	0	0	n	n	0	n	n	0	n	n	n	n	n	n			0.816	
K160T	0	0	0	0	0	0	0	0	0	0	0	0	0	n	0	0	0	0	n	0	0	n	0	0	n	0	n	0	n	n			0.772	
KTOOL	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	0.104	0.42	0.772	0.520