

## Supplementary Material

### Are machine learning based methods suited to address complex biological problems? Lessons from CAGI-5 challenges.

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## Supplementary Tables

**Table S1.** INPS-3D predictions for the *FXN* SAV dataset

SAV	Experimental $\Delta\Delta G$	Predicted $\Delta\Delta G$ (INPS-3D)
p.Asp104Gly	0.255	-0.776491
p.Ala107Val	0.22	-0.1439655
p.Ser202Phe	-0.685	0.4780699
p.Ser181Phe	-2.035	-0.115336
p.Phe109Leu	-2.645	-1.40127
p.Ser161Ile	-3.44	-0.697307
p.Tyr123Ser	-4.48	-2.577015
p.Trp173Cys	-9.5	-2.05509

**Table S3.** Predictions of different methods for the *CHEK2* SAV dataset

SAV	$p_{\text{case}}$	SNPs&GO	$P_d$	$P_p$	INPS
p.Ala584Val	1.00	0.760	0.560	0.750	0.840
p.Ala94Thr	1.00	0.590	0.540	0.594	0.500
p.Asp154Gly	1.00	0.880	0.820	0.654	0.793
p.Asp540Asn	1.00	0.530	0.650	0.570	0.500
p.Ser67Tyr	1.00	0.650	0.640	0.750	0.840
p.Glu547Gln	0.00	0.530	0.560	0.500	0.500
p.Glu64Lys	0.00	0.680	0.760	0.500	0.500
p.Leu555Val	0.50	0.560	0.540	0.834	0.500
p.Pro85Leu	0.00	0.500	0.590	0.714	0.775
p.Arg562Leu	0.00	0.650	0.960	0.500	0.500
p.Arg95Gln	0.00	0.650	0.750	0.500	0.500
p.Ser548Pro	0.00	0.680	0.690	0.500	0.500
p.Val66Met	0.00	0.650	0.630	0.630	0.500
p.Ile158Met	1.00	0.820	0.590	0.750	1.000
p.Ile200Thr	1.00	0.740	0.770	1.000	1.000
p.Ile203Arg	1.00	0.850	0.810	0.834	1.000
p.Arg160Gly	1.00	0.970	0.850	0.882	1.000
p.Arg180Gln	1.00	0.590	0.750	0.500	0.500
p.Arg191Gly	1.00	0.880	0.850	0.882	1.000
p.Arg223Cys	1.00	0.820	0.950	0.750	0.620
p.Asp481Tyr	1.00	0.760	0.910	0.630	0.500
p.Glu282Lys	1.00	0.760	0.760	0.500	0.500
p.Gly385Ser	1.00	0.850	0.820	0.500	0.673
p.Ile264Val	1.00	0.820	0.500	0.500	0.519
p.Leu279Pro	0.82	0.910	0.960	0.702	1.000
p.Asn448Lys	1.00	0.850	0.750	0.630	0.500

p.Pro527Leu	1.00	0.790	0.590	0.714	1.000
p.Arg449His	1.00	0.710	0.800	0.702	0.500
p.Thr519Met	1.00	0.760	0.600	0.500	0.804
p.Glu420Gly	0.00	0.760	0.660	0.870	1.000
p.Ile491Ser	0.33	0.680	0.900	1.000	1.000
p.Ile491Val	0.00	0.740	0.500	0.500	0.708
p.Arg389His	0.00	1.000	0.800	0.702	1.000
p.Thr421Ile	0.00	0.710	0.630	0.558	0.520

**Table S4.** Predictions of different methods for the *PCM1* SAV dataset

SAV	Functional effect	P <sub>d</sub> (numeric)	P <sub>d</sub> (class)	P <sub>p</sub> (numeric)	P <sub>p</sub> (class)	INPS (numeric)	INPS (class)
p.Gly6Asp	P	0.82	P	0.67	P	-0.767	H
p.Glu23Asp	B	0.35	B	0.43	H	-0.618	H
p.Thr77Ala	B	0.27	B	0.5	H	-0.299	B
p.Met146Val	H	0.46	H	0.57	H	-1.078	P
p.Ala156Val	H	0.45	H	0.6	P	0.494	B
p.Met200Thr	H	0.41	H	0.25	B	-1.666	P
p.Met200Ile	H	0.45	H	0.43	H	-0.138	B
p.Asp214Gly	P	0.66	P	0.52	H	-0.802	H
p.Glu248Gln	H	0.45	H	0.25	B	-0.556	H
p.Glu311Gln	H	0.45	H	0.25	B	-0.640	H
p.Glu369Gly	H	0.53	H	0.7	P	-1.134	P
p.Pro390Ser	H	0.48	H	0.5	H	-0.345	B
p.Leu472Val	H	0.43	H	0.67	P	-0.381	B
p.Gly482Val	P	0.82	P	0.7	P	-0.508	H
p.Glu543Lys	P	0.61	P	0.33	B	-0.300	B
p.Asp574Gly	P	0.66	P	0.52	H	-0.803	H
p.Arg604Leu	P	0.77	P	0.4	B	0.056	B
p.Glu624Lys	P	0.61	P	0.33	B	-0.225	B
p.Ile659Met	H	0.47	H	0.6	P	-1.423	P
p.Ser804Arg	P	0.62	P	0.43	H	-0.020	B
p.Arg833Thr	P	0.71	P	0.67	P	-0.976	H
p.Cys876Arg	P	0.76	P	0.72	P	-1.281	P
p.Gly892Trp	P	0.81	P	0.5	H	-0.780	H
p.Glu917Gly	H	0.53	H	0.7	P	-1.099	P
p.Lys954Asn	H	0.5	H	0.57	H	-0.755	H
p.Asn1125Ser	H	0.49	H	0.63	P	-0.296	B
p.Lys1275Glu	H	0.48	H	0.33	B	-0.594	H
p.His1352Tyr	H	0.58	H	0.42	H	-0.038	B
p.Cys1361Tyr	P	0.83	P	0.67	P	0.486	B
p.Ala1490Gly	H	0.47	H	0.58	H	-1.192	P
p.Glu1535Lys	P	0.61	P	0.33	B	-0.467	B
p.Ala1553Gly	H	0.47	H	0.58	H	-0.031	B
p.Gly1556Asp	P	0.82	P	0.67	P	-0.518	H
p.Lys1861Asn	H	0.5	H	0.57	H	-0.648	H
p.Asn1875Lys	P	0.6	P	0.5	H	-0.271	B
p.Arg1907His	P	0.64	P	0.56	H	0.052	B
p.Pro1913Leu	H	0.47	H	0.57	H	0.618	H
p.Ala1979Ser	B	0.33	B	0.27	B	-0.760	H