

DeepVF: a deep learning-based hybrid framework for identifying virulence factors using the stacking strategy

Supplementary Material

Table of Contents

1. Supplementary Tables

Table S1. Detailed information of existing methods for VF prediction.

Table S2. Performance comparison of multiple models for predicting VFs with different CD-HIT thresholds on the 10-fold cross-validation test.

Table S3. Explanation of the seven parameters of XGBoost.

Table S4. Performance comparison of ML baseline models with traditional ML features (without combining with *seqsim*) and *seqsim* for predicting VFs on the 10-fold cross-validation test.

Table S5. Performance comparison of DL baseline models with their optimal features using different dropout rates on the 10-fold cross-validation test.

Table S6. Performance comparison of baseline models with different cut-off thresholds on the 10-fold cross-validation test.

Table S7. Performance comparison of different algorithms to construct the final meta model on the 10-fold cross-validation test.

Table S8. Performance comparison between models the meta model using the prediction label and the prediction score of baseline models based on the 10-fold cross validation test.

Table S9. Detailed contributions of different baseline models to the performance of the ultimate meta XGBoost classifier.

Table S10. Performance comparison of baseline models for VF prediction on the independent test.

Table S11. Performance comparison of DeepVF trained based on VirulentPred's training dataset and VirulentPred on different independent datasets.

Table S12. Summary of the top 100 sequences predicted by DeepVF and the BLAST-based baseline predictor based on three proteomes.

Table S13. Detailed information of DeepVF and the existing toolkits for VF prediction from the user's viewpoint.

2. Supplementary Figures

Fig. S1. Distribution of sequence lengths of proteins in the curated dataset.

Fig. S2. The pipeline of constructing a baseline model based on the training dataset, and the prediction process on the independent dataset.

Fig. S3. Performance comparison of different DL models trained using different sizes of protein sequence segments based on the sequence-to-vector feature encoding method.

Fig. S4. ROC curves of DeepVF, the BLAST-based baseline predictor and three existing state-of-the-art methods for VF prediction on the independent test.

3. Supplementary References

1. Supplementary Tables

Table S1. Detailed information of existing methods for VF prediction.

Method	Data size (Positive/Negative samples)	Negative sample selection	Features	Models	Ensemble learning strategy used	Web server or software accessibility	Origin
SPAAN (predicting the adhesins)	469/703	Bacterial enzymes and other non-virulent proteins	amino acid frequencies, multiplet frequencies, dipeptide frequencies, charge composition, hydrophobic composition	Neural network	NO	-	[1]
VirulentPred	1025/1030	Bacterial enzymes and other non-virulent proteins	AAC, DPC, higher order dipeptide composition, PSSM, sequence similarity	SVM	YES	http://bioinfo.icgeb.res.in/virulent/	[2]
E_SVM_VF ^a	1025/1030	Bacterial enzymes and other non-virulent proteins	Amino acid (AA) composition, Weighted AA composition, Five-level grouping composition	SVM	YES	-	[3]
Virulent-GO	1025/1030	Bacterial enzymes and other non-virulent proteins	GO terms	SVM, Naïve Bayes, KNN, C4.5 Decision tree	NO	-	[4]
VF_PAAC_EI ^a	1025/1030	Bacterial enzymes and other non-virulent proteins	2-Grams, RC, PA, GP, PSSM, AA	SVM	YES	-	[5]
ACCM_VFs ^a	803/803*5	Random sampling from the VF-contributing proteomes	AAC, PAAC	STRING network, BLAST, NNA classifier	NO	-	[6]
VF_KEGG ^a	514/514*5	Random sampling from the VF-contributing proteomes	KEGG pathway	RF	NO	-	[7]
MP3	1708/5815	Proteins deemed ‘essential’ for survival	AAC, DPC	SVM, HMM	NO	http://metagenomics.iisrb.ac.in/mp3/index.php	[8]
PBV ^a	3891/10025	A novel strategy for selecting negatives that relies on protein function annotation data.	DPC, sequence similarity	SVM, RF, BLAST	NO	-	[9]

^aAs the names of these methods were not explicitly provided in the original work, here we named them according to the titles of their published papers.

Table S2. Performance comparison of multiple models for predicting VFs with different CD-HIT thresholds on the 10-fold cross-validation test.

Model ^a	Threshold	SN	SP	ACC	F-value	MCC
RF-AAC	0.3	0.641±0.029	0.699±0.011	0.670±0.017	0.660±0.019	0.341±0.033
	0.4	0.655±0.025	0.710±0.021	0.682±0.014	0.673±0.016	0.366±0.027
	0.5	0.653±0.016	0.726±0.026	0.689±0.009	0.678±0.011	0.380±0.019
	0.6	0.670±0.020	0.738±0.015	0.704±0.013	0.693±0.016	0.408±0.025
	0.7	0.663±0.031	0.751±0.013	0.707±0.017	0.693±0.019	0.416±0.032
	0.8	0.667±0.013	0.755±0.025	0.711±0.015	0.698±0.016	0.424±0.030
	0.9	0.671±0.026	0.761±0.018	0.716±0.016	0.702±0.016	0.433±0.031
SVM-AAC	0.3	0.663±0.037	0.700±0.016	0.681±0.022	0.675±0.026	0.363±0.044
	0.4	0.685±0.034	0.710±0.027	0.697±0.016	0.693±0.018	0.396±0.032
	0.5	0.687±0.016	0.717±0.024	0.702±0.013	0.697±0.017	0.403±0.025
	0.6	0.706±0.018	0.729±0.024	0.718±0.016	0.714±0.019	0.435±0.031
	0.7	0.716±0.019	0.741±0.012	0.728±0.012	0.725±0.010	0.456±0.023
	0.8	0.714±0.011	0.740±0.016	0.727±0.010	0.723±0.015	0.453±0.020
	0.9	0.723±0.025	0.752±0.017	0.737±0.011	0.733±0.014	0.475±0.022
XGBoost-AAC	0.3	0.651±0.022	0.672±0.016	0.661±0.009	0.657±0.012	0.323±0.019
	0.4	0.663±0.027	0.704±0.026	0.683±0.013	0.676±0.018	0.368±0.026
	0.5	0.666±0.021	0.702±0.025	0.684±0.014	0.678±0.019	0.368±0.029
	0.6	0.667±0.019	0.721±0.017	0.694±0.014	0.685±0.020	0.388±0.028
	0.7	0.673±0.024	0.737±0.018	0.705±0.019	0.695±0.018	0.411±0.036
	0.8	0.663±0.011	0.737±0.029	0.700±0.016	0.688±0.013	0.401±0.032
	0.9	0.676±0.029	0.747±0.018	0.711±0.016	0.700±0.018	0.424±0.031
MLP-AAC	0.3	0.571±0.082	0.790±0.066	0.681±0.015	0.639±0.043	0.374±0.025
	0.4	0.667±0.064	0.711±0.069	0.690±0.010	0.682±0.027	0.381±0.021
	0.5	0.625±0.066	0.765±0.069	0.694±0.009	0.670±0.023	0.397±0.020
	0.6	0.637±0.058	0.770±0.046	0.705±0.014	0.682±0.033	0.412±0.025
	0.7	0.624±0.035	0.795±0.046	0.711±0.013	0.683±0.018	0.427±0.029
	0.8	0.652±0.051	0.783±0.054	0.718±0.009	0.697±0.021	0.441±0.020
	0.9	0.681±0.072	0.747±0.091	0.716±0.015	0.704±0.026	0.434±0.033
CNN-L500	0.3	0.647±0.082	0.694±0.076	0.670±0.012	0.660±0.032	0.346±0.022
	0.4	0.616±0.057	0.739±0.059	0.677±0.012	0.655±0.020	0.360±0.025
	0.5	0.638±0.055	0.750±0.050	0.695±0.008	0.675±0.028	0.392±0.017

	0.6	0.661±0.056	0.747±0.048	0.704±0.009	0.690±0.027	0.411±0.019
	0.7	0.687±0.066	0.745±0.060	0.717±0.011	0.706±0.026	0.436±0.022
	0.8	0.688±0.044	0.756±0.037	0.722±0.006	0.712±0.022	0.446±0.013
	0.9	0.688±0.055	0.774±0.050	0.731±0.010	0.718±0.023	0.465±0.020
LSTM-L500	0.3	0.567±0.033	0.754±0.016	0.660±0.014	0.625±0.021	0.327±0.025
	0.4	0.568±0.057	0.767±0.045	0.668±0.012	0.630±0.030	0.343±0.021
	0.5	0.608±0.040	0.745±0.044	0.677±0.008	0.652±0.020	0.358±0.016
	0.6	0.612±0.050	0.767±0.036	0.690±0.014	0.662±0.032	0.384±0.027
	0.7	0.647±0.053	0.742±0.053	0.694±0.013	0.678±0.020	0.392±0.029
	0.8	0.641±0.048	0.748±0.050	0.696±0.010	0.677±0.023	0.393±0.022
	0.9	0.636±0.079	0.772±0.065	0.705±0.012	0.680±0.037	0.415±0.021
	0.3	0.611±0.038	0.748±0.041	0.680±0.012	0.656±0.019	0.364±0.025
DNN-L500	0.4	0.632±0.048	0.736±0.043	0.685±0.016	0.667±0.028	0.372±0.033
	0.5	0.616±0.029	0.765±0.029	0.691±0.011	0.665±0.022	0.385±0.025
	0.6	0.607±0.039	0.792±0.037	0.699±0.010	0.668±0.024	0.406±0.024
	0.7	0.623±0.036	0.774±0.041	0.699±0.013	0.674±0.020	0.403±0.026
	0.8	0.615±0.048	0.784±0.047	0.701±0.012	0.672±0.027	0.407±0.024
	0.9	0.622±0.029	0.785±0.036	0.704±0.013	0.677±0.017	0.413±0.027

Note: For ML, we used the simplest feature AAC (without combining with *seqsim*) to train multiple ML models with different CD-HIT thresholds. For DL, we used the L-500 to train multiple DL models with different thresholds of CD-HIT as the length of majority of sequences in the curated dataset were less than 500 (**Fig. S1**). ^aModel are denoted as model_feature. For example, RF_AAC means that the model was trained based on RF with AAC features. Values are expressed as mean±standard deviation. The best performance value for each metric across different CD-HIT thresholds within each model is highlighted in bold.

Table S3. Explanation of the seven parameters of XGBoost.

Parameters	Description ^a	Parameter tuning range
<i>max_depth</i>	max depth of the tree	[4,8], step: 1
<i>eta</i>	step size shrinkage	[0.01, 0.03], step: random
<i>gamma</i>	minimum loss reduction	[0, 0.2], step: random
<i>subsample</i>	subsample ratio of the training instances	[0.6, 0.9], step: random
<i>colsample_bytree</i>	subsampling ratio of columns	[0.5, 0.8], step: random
<i>min_child_weight</i>	minimum sum of instance weight	[1, 40], step: 1
<i>max_delta_step</i>	maximum delta step we allow each leaf output to be	[1,10], step: 1

Note: ^aThe description of the above parameters comes from the official XGBoost document (<https://xgboost.readthedocs.io/en/latest/>).

Table S4. Performance comparison of ML baseline models with traditional ML features (without combining with *seqsim*) and *seqsim* for predicting VFs on the 10-fold cross-validation test.

Algorithm	Encoding	SN	SP	ACC	F-value	MCC
RF	AAC	0.643±0.019	0.694±0.028	0.668±0.017	0.659±0.015	0.337±0.033
	DPC	0.599±0.029	0.739±0.025	0.669±0.020	0.644±0.022	0.341±0.040
	DDE	0.610±0.036	0.720±0.024	0.665±0.022	0.645±0.027	0.331±0.044
	PAAC	0.641±0.021	0.698±0.025	0.669±0.017	0.660±0.017	0.339±0.034
	QSO	0.641±0.022	0.717±0.024	0.679±0.014	0.666±0.015	0.359±0.029
SVM	PSSM-composition	0.660±0.022	0.830±0.024	0.745±0.015	0.721±0.017	0.498±0.031
	S-FPSSM	0.661±0.022	0.816±0.021	0.739±0.017	0.717±0.018	0.484±0.033
	RPM-PSSM	0.652±0.020	0.833±0.024	0.743±0.015	0.717±0.016	0.493±0.030
	seqsim	0.824±0.017	0.751±0.020	0.787±0.011	0.795±0.012	0.576±0.022
	AAC	0.651±0.022	0.710±0.025	0.680±0.016	0.671±0.018	0.362±0.033
XGBoost	DPC	0.626±0.021	0.695±0.027	0.660±0.017	0.648±0.017	0.321±0.035
	DDE	0.647±0.026	0.683±0.029	0.665±0.020	0.658±0.020	0.330±0.041
	PAAC	0.657±0.017	0.704±0.026	0.680±0.016	0.672±0.018	0.361±0.033
	QSO	0.653±0.022	0.708±0.029	0.681±0.016	0.672±0.014	0.362±0.032
	PSSM-composition	0.695±0.034	0.846±0.022	0.770±0.021	0.751±0.025	0.547±0.040
MLP	S-FPSSM	0.718±0.026	0.783±0.023	0.750±0.019	0.742±0.021	0.502±0.038
	RPM-PSSM	0.692±0.033	0.819±0.026	0.756±0.019	0.739±0.023	0.516±0.038
	seqsim	0.859±0.022	0.737±0.025	0.798±0.016	0.810±0.017	0.601±0.030
	AAC	0.647±0.020	0.678±0.029	0.662±0.017	0.657±0.015	0.325±0.033
	DPC	0.631±0.024	0.701±0.026	0.666±0.017	0.654±0.019	0.333±0.034
	DDE	0.631±0.025	0.699±0.026	0.665±0.017	0.653±0.020	0.331±0.033
	PAAC	0.646±0.018	0.680±0.027	0.663±0.015	0.657±0.014	0.327±0.030
	QSO	0.651±0.021	0.702±0.027	0.676±0.018	0.668±0.017	0.353±0.035
	PSSM-composition	0.715±0.025	0.810±0.024	0.762±0.016	0.751±0.017	0.528±0.032
	S-FPSSM	0.700±0.025	0.801±0.025	0.750±0.018	0.737±0.019	0.504±0.036
	RPM-PSSM	0.711±0.025	0.793±0.024	0.752±0.016	0.741±0.017	0.506±0.031
	seqsim	0.825±0.019	0.761±0.026	0.793±0.017	0.799±0.019	0.587±0.033
	AAC	0.634±0.082	0.711±0.092	0.673±0.017	0.658±0.032	0.351±0.034
	DPC	0.615±0.079	0.734±0.082	0.675±0.015	0.652±0.034	0.356±0.031
	DDE	0.614±0.074	0.700±0.068	0.657±0.016	0.639±0.036	0.318±0.033
	PAAC	0.625±0.093	0.712±0.092	0.669±0.017	0.651±0.039	0.344±0.033
	QSO	0.607±0.079	0.740±0.071	0.674±0.016	0.648±0.039	0.353±0.031

PSSM-composition	0.746±0.041	0.809±0.040	0.778±0.015	0.770±0.018	0.557±0.030
S-FPSSM	0.700±0.062	0.821±0.048	0.761±0.015	0.744±0.027	0.529±0.028
RPM-PSSM	0.718±0.047	0.824±0.037	0.771±0.017	0.758±0.023	0.547±0.032
seqsim	0.856±0.031	0.738±0.034	0.797±0.012	0.808±0.014	0.600±0.025

Note: Values are expressed as mean±standard deviation. The best performance value for each metric across different encoding methods within each algorithm is highlighted in bold.

Table S5. Performance comparison of DL baseline models with their optimal features using different dropout rates on the 10-fold cross-validation test.

Model ^a	Dropout rate	SN	SP	ACC	F-value	MCC
CNN-L900	0.01	0.598±0.087	0.724±0.084	0.661±0.017	0.635±0.042	0.329±0.032
	0.05	0.609±0.057	0.711±0.053	0.660±0.015	0.641±0.027	0.324±0.029
	0.10	0.585±0.122	0.728±0.111	0.658±0.015	0.624±0.064	0.325±0.029
	0.15	0.608±0.057	0.709±0.045	0.658±0.016	0.639±0.029	0.320±0.031
	0.20	0.620±0.097	0.696±0.101	0.658±0.012	0.642±0.039	0.322±0.024
	0.25	0.578±0.078	0.738±0.068	0.658±0.014	0.625±0.042	0.323±0.026
	0.30	0.585±0.035	0.731±0.037	0.659±0.013	0.631±0.022	0.320±0.027
LSTM-L1000	0.01	0.560±0.102	0.748±0.093	0.654±0.028	0.613±0.062	0.318±0.051
	0.05	0.544±0.115	0.746±0.103	0.646±0.017	0.599±0.064	0.303±0.029
	0.10	0.487±0.091	0.802±0.056	0.645±0.023	0.573±0.064	0.307±0.037
	0.15	0.516±0.136	0.773±0.094	0.645±0.027	0.583±0.082	0.305±0.041
	0.20	0.545±0.107	0.747±0.071	0.646±0.025	0.601±0.064	0.302±0.047
	0.25	0.571±0.100	0.739±0.062	0.654±0.030	0.618±0.070	0.316±0.059
	0.30	0.540±0.073	0.759±0.077	0.649±0.014	0.603±0.032	0.309±0.028
DNN-L500	0.01	0.620±0.054	0.730±0.050	0.675±0.019	0.655±0.027	0.353±0.036
	0.05	0.612±0.069	0.729±0.057	0.670±0.014	0.648±0.033	0.345±0.024
	0.10	0.603±0.045	0.741±0.036	0.672±0.013	0.647±0.021	0.348±0.021
	0.15	0.596±0.077	0.751±0.070	0.673±0.009	0.644±0.034	0.354±0.013
	0.20	0.610±0.027	0.732±0.026	0.671±0.011	0.649±0.013	0.344±0.020
	0.25	0.584±0.064	0.756±0.050	0.670±0.013	0.637±0.032	0.347±0.022
	0.30	0.613±0.056	0.736±0.053	0.674±0.011	0.652±0.023	0.354±0.021

Note: Values are expressed as mean±standard deviation. The best performance value for each metric across different dropout rates within each algorithm is highlighted in bold.

Table S6. Performance comparison of baseline models with different cut-off thresholds on the 10-fold cross-validation test.

Threshold	Model	Encoding	SN	SP	ACC	F-value	MCC
0.3	RF	AAC	0.924±0.013	0.631±0.031	0.777±0.017	0.806±0.015	0.581±0.029
		DPC	0.986±0.011	0.154±0.099	0.570±0.047	0.697±0.024	0.244±0.086
		DDE	0.974±0.018	0.274±0.130	0.624±0.059	0.723±0.031	0.345±0.093
		PAAC	0.925±0.014	0.628±0.038	0.777±0.020	0.806±0.017	0.580±0.034
		QSO	0.938±0.014	0.524±0.059	0.731±0.028	0.777±0.020	0.507±0.043
		PSSM-composition	0.947±0.011	0.459±0.073	0.703±0.035	0.762±0.023	0.465±0.055
		S-FPSSM	0.956±0.010	0.434±0.035	0.695±0.019	0.758±0.015	0.458±0.031
		RPM-PSSM	0.964±0.013	0.323±0.088	0.644±0.041	0.731±0.023	0.373±0.065
0.5	SVM	AAC	0.916±0.014	0.567±0.032	0.742±0.018	0.780±0.016	0.516±0.032
		DPC	0.923±0.016	0.417±0.038	0.670±0.021	0.737±0.016	0.394±0.039
		DDE	0.915±0.017	0.435±0.031	0.675±0.018	0.738±0.015	0.399±0.032
		PAAC	0.918±0.015	0.552±0.035	0.735±0.020	0.776±0.017	0.505±0.035
		QSO	0.923±0.013	0.495±0.032	0.709±0.018	0.760±0.015	0.463±0.032
		PSSM-composition	0.892±0.023	0.652±0.025	0.772±0.017	0.796±0.016	0.560±0.035
		S-FPSSM	0.905±0.021	0.542±0.054	0.723±0.027	0.766±0.020	0.480±0.047
		RPM-PSSM	0.890±0.017	0.582±0.030	0.736±0.018	0.771±0.016	0.496±0.034
0.7	XGBoost	AAC	0.921±0.014	0.658±0.026	0.789±0.014	0.814±0.013	0.600±0.024
		DPC	0.908±0.015	0.677±0.023	0.793±0.014	0.814±0.015	0.601±0.027
		DDE	0.906±0.017	0.693±0.024	0.800±0.015	0.819±0.015	0.614±0.028
		PAAC	0.913±0.013	0.682±0.026	0.797±0.016	0.818±0.015	0.611±0.029
		QSO	0.919±0.014	0.675±0.037	0.797±0.018	0.819±0.015	0.614±0.030
		PSSM-composition	0.890±0.017	0.800±0.024	0.845±0.012	0.852±0.012	0.694±0.023
		S-FPSSM	0.903±0.018	0.764±0.029	0.833±0.015	0.844±0.014	0.673±0.028
		RPM-PSSM	0.900±0.014	0.763±0.020	0.832±0.012	0.842±0.012	0.670±0.023
0.9	MLP	AAC	0.915±0.028	0.602±0.116	0.759±0.048	0.793±0.027	0.545±0.084
		DPC	0.909±0.025	0.637±0.099	0.773±0.040	0.801±0.022	0.568±0.069
		DDE	0.865±0.042	0.697±0.089	0.781±0.030	0.798±0.020	0.574±0.050
		PAAC	0.916±0.029	0.601±0.121	0.758±0.051	0.793±0.028	0.545±0.085
		QSO	0.901±0.026	0.657±0.088	0.779±0.037	0.803±0.023	0.576±0.057
		PSSM-composition	0.875±0.030	0.772±0.073	0.824±0.025	0.833±0.017	0.654±0.042
		S-FPSSM	0.859±0.039	0.762±0.073	0.811±0.025	0.820±0.019	0.627±0.045
		RPM-PSSM	0.884±0.029	0.746±0.075	0.815±0.028	0.828±0.018	0.639±0.045

CNN	L100	0.856±0.118	0.268±0.176	0.562±0.035	0.660±0.024
	L200	0.843±0.105	0.326±0.163	0.585±0.033	0.669±0.020
	L300	0.787±0.115	0.426±0.180	0.607±0.036	0.665±0.022
	L400	0.806±0.104	0.408±0.171	0.607±0.037	0.671±0.021
	L500	0.828±0.093	0.381±0.170	0.604±0.044	0.676±0.017
	L600	0.809±0.101	0.403±0.169	0.607±0.039	0.672±0.021
	L700	0.818±0.101	0.383±0.173	0.600±0.043	0.671±0.017
	L800	0.811±0.099	0.404±0.166	0.607±0.039	0.673±0.019
	L900	0.837±0.100	0.363±0.167	0.600±0.038	0.676±0.020
	L1000	0.819±0.106	0.386±0.170	0.603±0.038	0.672±0.022
LSTM	L100	0.930±0.069	0.147±0.117	0.538±0.030	0.668±0.015
	L200	0.916±0.075	0.187±0.129	0.552±0.029	0.671±0.015
	L300	0.929±0.074	0.149±0.147	0.539±0.039	0.668±0.014
	L400	0.939±0.067	0.118±0.125	0.529±0.033	0.666±0.014
	L500	0.874±0.113	0.226±0.198	0.550±0.046	0.659±0.019
	L600	0.935±0.082	0.111±0.154	0.523±0.042	0.662±0.017
	L700	0.933±0.103	0.105±0.189	0.518±0.046	0.659±0.015
	L800	0.967±0.063	0.054±0.117	0.511±0.031	0.664±0.014
	L900	0.965±0.074	0.066±0.147	0.516±0.040	0.666±0.012
	L1000	0.956±0.093	0.086±0.169	0.521±0.041	0.666±0.014
DNN	L100	0.959±0.027	0.110±0.054	0.534±0.019	0.673±0.011
	L200	0.937±0.032	0.167±0.074	0.553±0.025	0.677±0.015
	L300	0.944±0.027	0.173±0.066	0.559±0.025	0.681±0.014
	L400	0.937±0.034	0.182±0.081	0.560±0.026	0.680±0.013
	L500	0.941±0.029	0.172±0.066	0.556±0.026	0.680±0.014
	L600	0.944±0.029	0.169±0.064	0.557±0.022	0.681±0.012
	L700	0.943±0.033	0.167±0.069	0.555±0.022	0.680±0.012
	L800	0.947±0.031	0.154±0.066	0.551±0.024	0.678±0.014
	L900	0.951±0.026	0.150±0.055	0.551±0.020	0.679±0.013
	L1000	0.948±0.039	0.151±0.074	0.550±0.023	0.678±0.014
0.4	RF	AAC	0.888±0.016	0.735±0.025	0.812±0.016
		DPC	0.923±0.018	0.554±0.084	0.739±0.037
		DDE	0.913±0.018	0.595±0.047	0.754±0.025
		PAAC	0.890±0.015	0.735±0.026	0.813±0.016

SVM	QSO	0.892±0.016	0.716±0.025	0.804±0.015	0.820±0.014
	PSSM-composition	0.865±0.019	0.765±0.038	0.815±0.022	0.824±0.020
	S-FPSSM	0.897±0.017	0.685±0.025	0.791±0.016	0.811±0.014
	RPM-PSSM	0.883±0.015	0.710±0.042	0.797±0.023	0.813±0.019
	AAC	0.875±0.015	0.673±0.029	0.774±0.018	0.795±0.017
	DPC	0.846±0.022	0.589±0.030	0.717±0.019	0.749±0.017
	DDE	0.843±0.017	0.590±0.036	0.716±0.019	0.748±0.016
	PAAC	0.880±0.016	0.666±0.030	0.773±0.017	0.795±0.016
XGBoost	QSO	0.880±0.016	0.630±0.031	0.755±0.018	0.782±0.016
	PSSM-composition	0.844±0.027	0.762±0.023	0.803±0.018	0.811±0.018
	S-FPSSM	0.857±0.023	0.669±0.047	0.763±0.024	0.784±0.020
	RPM-PSSM	0.828±0.023	0.737±0.028	0.782±0.017	0.792±0.016
	AAC	0.886±0.017	0.742±0.025	0.814±0.016	0.826±0.016
	DPC	0.884±0.016	0.744±0.023	0.814±0.014	0.826±0.014
	DDE	0.878±0.018	0.760±0.021	0.819±0.015	0.829±0.016
	PAAC	0.878±0.017	0.757±0.027	0.817±0.018	0.828±0.018
MLP	QSO	0.886±0.015	0.755±0.025	0.820±0.015	0.831±0.015
	PSSM-composition	0.858±0.018	0.851±0.019	0.854±0.012	0.855±0.013
	S-FPSSM	0.873±0.017	0.825±0.025	0.849±0.015	0.852±0.015
	RPM-PSSM	0.869±0.015	0.829±0.019	0.849±0.013	0.852±0.014
	AAC	0.890±0.024	0.683±0.039	0.787±0.016	0.807±0.015
	DPC	0.882±0.025	0.705±0.038	0.794±0.015	0.810±0.014
	DDE	0.838±0.037	0.758±0.058	0.798±0.018	0.806±0.014
	PAAC	0.890±0.027	0.677±0.088	0.784±0.038	0.805±0.023
CNN	QSO	0.878±0.027	0.712±0.050	0.795±0.020	0.810±0.016
	PSSM-composition	0.844±0.027	0.843±0.039	0.843±0.013	0.843±0.012
	S-FPSSM	0.829±0.039	0.825±0.047	0.827±0.016	0.827±0.016
	RPM-PSSM	0.852±0.029	0.817±0.038	0.835±0.011	0.838±0.011
	L100	0.744±0.109	0.445±0.142	0.594±0.025	0.645±0.030
	L200	0.742±0.096	0.502±0.117	0.622±0.020	0.661±0.030

		Performance Metrics							
		Model A			Model B			Model C	
		Parameter	Value	Value	Value	Value	Value	Value	Value
LSTM	L700		0.729±0.082	0.544±0.123	0.636±0.031	0.666±0.021	0.283±0.049		
	L800		0.720±0.081	0.560±0.107	0.640±0.022	0.665±0.024	0.288±0.038		
	L900		0.728±0.092	0.543±0.135	0.636±0.030	0.665±0.027	0.281±0.056		
	L1000		0.729±0.097	0.539±0.133	0.635±0.027	0.664±0.027	0.280±0.042		
	L100		0.773±0.100	0.400±0.138	0.587±0.025	0.650±0.026	0.191±0.038		
	L200		0.747±0.093	0.479±0.129	0.614±0.027	0.657±0.028	0.239±0.047		
	L300		0.768±0.140	0.426±0.218	0.598±0.046	0.653±0.031	0.216±0.076		
	L400		0.769±0.114	0.441±0.183	0.605±0.043	0.659±0.025	0.228±0.070		
	L500		0.671±0.113	0.585±0.155	0.627±0.031	0.640±0.034	0.263±0.052		
	L600		0.724±0.155	0.494±0.268	0.609±0.062	0.646±0.030	0.226±0.117		
DNN	L700		0.729±0.167	0.468±0.293	0.598±0.067	0.642±0.026	0.203±0.132		
	L800		0.720±0.140	0.505±0.236	0.612±0.055	0.647±0.026	0.232±0.104		
	L900		0.721±0.181	0.490±0.302	0.605±0.065	0.641±0.035	0.220±0.126		
	L1000		0.736±0.145	0.500±0.259	0.618±0.061	0.656±0.024	0.244±0.118		
	L100		0.793±0.055	0.393±0.080	0.593±0.018	0.660±0.013	0.204±0.032		
	L200		0.777±0.058	0.465±0.067	0.622±0.015	0.672±0.022	0.258±0.029		
	L300		0.773±0.061	0.468±0.076	0.621±0.021	0.670±0.023	0.255±0.040		
	L400		0.785±0.050	0.482±0.086	0.634±0.024	0.682±0.016	0.282±0.046		
	L500		0.776±0.057	0.491±0.083	0.633±0.022	0.679±0.016	0.281±0.040		
	L600		0.765±0.053	0.509±0.079	0.637±0.022	0.678±0.017	0.285±0.041		
0.6	L700		0.768±0.050	0.498±0.078	0.633±0.021	0.676±0.016	0.277±0.041		
	L800		0.778±0.058	0.485±0.074	0.631±0.017	0.678±0.017	0.277±0.034		
	L900		0.775±0.053	0.485±0.078	0.630±0.018	0.677±0.015	0.274±0.034		
	L1000		0.777±0.081	0.471±0.125	0.624±0.029	0.673±0.019	0.264±0.045		
	RF	AAC	0.743±0.023	0.884±0.018	0.813±0.013	0.799±0.015	0.633±0.025		
	DPC		0.476±0.082	0.953±0.018	0.714±0.037	0.621±0.068	0.489±0.057		
	DDE		0.573±0.094	0.926±0.020	0.749±0.042	0.692±0.069	0.535±0.068		
	PAAC		0.739±0.023	0.886±0.017	0.812±0.013	0.797±0.015	0.632±0.025		
	QSO		0.682±0.031	0.906±0.019	0.794±0.015	0.767±0.019	0.603±0.027		
	PSSM-composition		0.635±0.035	0.957±0.012	0.796±0.017	0.757±0.024	0.626±0.028		
	S-FPSSM		0.635±0.025	0.954±0.011	0.795±0.013	0.755±0.018	0.622±0.023		
	RPM-PSSM		0.600±0.040	0.963±0.011	0.781±0.019	0.732±0.030	0.604±0.030		
SVM	AAC		0.734±0.023	0.848±0.023	0.791±0.016	0.778±0.018	0.586±0.032		
	DPC		0.605±0.022	0.851±0.019	0.728±0.016	0.690±0.018	0.471±0.030		

	DDE	0.614±0.019	0.840±0.022	0.727±0.014	0.692±0.016
	PAAC	0.731±0.021	0.847±0.022	0.788±0.016	0.775±0.018
	QSO	0.697±0.024	0.860±0.021	0.778±0.016	0.758±0.017
	PSSM-composition	0.728±0.026	0.889±0.019	0.809±0.015	0.792±0.017
	S-FPSSM	0.687±0.080	0.873±0.032	0.780±0.031	0.755±0.050
	RPM-PSSM	0.685±0.040	0.891±0.020	0.788±0.022	0.764±0.028
XGBoost	AAC	0.761±0.022	0.870±0.019	0.816±0.012	0.805±0.014
	DPC	0.780±0.024	0.860±0.021	0.820±0.019	0.812±0.021
	DDE	0.791±0.021	0.862±0.017	0.826±0.013	0.820±0.015
	PAAC	0.768±0.016	0.867±0.019	0.818±0.011	0.808±0.013
	QSO	0.763±0.025	0.877±0.016	0.820±0.013	0.809±0.016
	PSSM-composition	0.804±0.018	0.912±0.017	0.858±0.011	0.850±0.012
	S-FPSSM	0.806±0.019	0.908±0.017	0.857±0.013	0.849±0.014
	RPM-PSSM	0.798±0.020	0.909±0.017	0.854±0.013	0.845±0.015
MLP	AAC	0.752±0.110	0.803±0.051	0.778±0.035	0.767±0.071
	DPC	0.768±0.069	0.806±0.047	0.787±0.020	0.781±0.032
	DDE	0.755±0.052	0.832±0.043	0.794±0.014	0.785±0.022
	PAAC	0.730±0.126	0.808±0.056	0.770±0.041	0.753±0.086
	QSO	0.758±0.090	0.813±0.055	0.786±0.023	0.777±0.043
	PSSM-composition	0.783±0.031	0.907±0.024	0.845±0.012	0.834±0.015
	S-FPSSM	0.758±0.042	0.905±0.027	0.832±0.015	0.818±0.021
	RPM-PSSM	0.787±0.042	0.889±0.034	0.838±0.011	0.829±0.016
CNN	L100	0.362±0.152	0.829±0.107	0.596±0.027	0.452±0.132
	L200	0.446±0.122	0.808±0.090	0.627±0.027	0.535±0.090
	L300	0.493±0.125	0.776±0.098	0.635±0.024	0.565±0.084
	L400	0.468±0.118	0.807±0.088	0.638±0.025	0.554±0.084
	L500	0.473±0.127	0.811±0.092	0.642±0.026	0.559±0.086
	L600	0.481±0.116	0.804±0.090	0.642±0.023	0.565±0.076
	L700	0.486±0.105	0.797±0.094	0.642±0.021	0.569±0.069
	L800	0.480±0.102	0.807±0.086	0.644±0.022	0.567±0.068
	L900	0.442±0.112	0.839±0.081	0.641±0.025	0.543±0.079
	L1000	0.475±0.118	0.814±0.092	0.645±0.021	0.563±0.079
LSTM	L100	0.364±0.125	0.836±0.098	0.601±0.020	0.464±0.098
	L200	0.412±0.083	0.842±0.059	0.627±0.022	0.520±0.066

		Performance Metrics				
		ROC-AUC				
Model Type	Feature Type	Performance Metrics				
		Prec.	Recall	F1	AUC	Acc.
DNN	L300	0.422±0.136	0.830±0.097	0.626±0.031	0.518±0.099	0.284±0.049
	L400	0.438±0.104	0.808±0.082	0.623±0.028	0.530±0.075	0.269±0.050
	L500	0.439±0.103	0.815±0.076	0.627±0.026	0.533±0.076	0.279±0.040
	L600	0.444±0.104	0.827±0.060	0.636±0.030	0.542±0.082	0.296±0.050
	L700	0.415±0.127	0.827±0.086	0.621±0.038	0.511±0.105	0.271±0.067
	L800	0.437±0.122	0.825±0.089	0.631±0.027	0.532±0.087	0.290±0.046
	L900	0.436±0.108	0.833±0.073	0.635±0.030	0.536±0.087	0.298±0.054
	L1000	0.450±0.107	0.825±0.072	0.638±0.028	0.546±0.081	0.301±0.047
	DNN	0.361±0.059	0.852±0.045	0.607±0.016	0.476±0.047	0.246±0.031
	L100	0.411±0.060	0.851±0.037	0.631±0.016	0.525±0.048	0.293±0.027
0.7	L200	0.406±0.060	0.854±0.041	0.630±0.017	0.521±0.046	0.293±0.029
	L300	0.438±0.064	0.839±0.045	0.639±0.018	0.545±0.048	0.305±0.032
	L400	0.447±0.057	0.849±0.041	0.648±0.017	0.558±0.042	0.325±0.031
	L500	0.447±0.055	0.845±0.047	0.646±0.017	0.556±0.038	0.320±0.032
	L600	0.431±0.062	0.858±0.043	0.645±0.019	0.546±0.046	0.322±0.032
	L700	0.440±0.062	0.848±0.048	0.644±0.020	0.551±0.047	0.318±0.035
	L800	0.435±0.059	0.856±0.042	0.646±0.020	0.549±0.045	0.323±0.036
	L900	0.422±0.059	0.863±0.040	0.643±0.017	0.539±0.045	0.320±0.030
	RF	0.585±0.036	0.942±0.014	0.764±0.017	0.712±0.027	0.564±0.028
	DPC	0.204±0.054	0.992±0.006	0.598±0.027	0.334±0.073	0.317±0.049
0.7	DDE	0.242±0.075	0.988±0.007	0.615±0.036	0.380±0.094	0.343±0.063
	PAAC	0.583±0.041	0.942±0.013	0.763±0.018	0.710±0.030	0.564±0.029
	QSO	0.480±0.046	0.964±0.013	0.722±0.021	0.632±0.039	0.507±0.033
	PSSM-composition	0.502±0.041	0.980±0.009	0.741±0.021	0.659±0.035	0.549±0.031
	S-FPSSM	0.447±0.027	0.988±0.005	0.718±0.015	0.613±0.025	0.518±0.022
	RPM-PSSM	0.418±0.046	0.987±0.006	0.703±0.024	0.583±0.045	0.493±0.035
	SVM	0.599±0.024	0.918±0.019	0.758±0.013	0.712±0.019	0.546±0.027
	DPC	0.445±0.026	0.931±0.013	0.688±0.016	0.587±0.024	0.430±0.028
0.7	DDE	0.453±0.027	0.921±0.017	0.687±0.016	0.591±0.026	0.423±0.031
	PAAC	0.593±0.028	0.921±0.019	0.757±0.014	0.709±0.021	0.544±0.027
	QSO	0.537±0.029	0.937±0.016	0.736±0.017	0.670±0.023	0.516±0.030
	PSSM-composition	0.650±0.026	0.932±0.013	0.791±0.014	0.757±0.019	0.607±0.026
	S-FPSSM	0.533±0.063	0.943±0.017	0.738±0.028	0.668±0.051	0.522±0.044
	RPM-PSSM	0.595±0.037	0.937±0.016	0.766±0.020	0.718±0.029	0.567±0.035

Performance Metrics for Various Models and Feature Sets						
		AUC	ROC AUC	Precision	Recall	F1 Score
XGBoost	AAC	0.638±0.032	0.925±0.016	0.782±0.015	0.745±0.021	0.589±0.025
	DPC	0.677±0.026	0.916±0.014	0.796±0.016	0.768±0.020	0.610±0.030
	DDE	0.703±0.022	0.908±0.015	0.805±0.013	0.783±0.015	0.624±0.024
	PAAC	0.674±0.022	0.915±0.016	0.794±0.013	0.766±0.015	0.607±0.024
	QSO	0.657±0.047	0.925±0.017	0.791±0.020	0.758±0.030	0.605±0.032
	PSSM-composition	0.765±0.022	0.935±0.016	0.850±0.011	0.836±0.012	0.710±0.020
	S-FPSSM	0.756±0.024	0.936±0.015	0.846±0.012	0.831±0.015	0.704±0.023
	RPM-PSSM	0.745±0.018	0.940±0.015	0.843±0.011	0.826±0.013	0.699±0.021
MLP	AAC	0.527±0.158	0.897±0.049	0.712±0.061	0.630±0.151	0.456±0.114
	DPC	0.590±0.122	0.886±0.050	0.738±0.040	0.685±0.080	0.504±0.061
	DDE	0.701±0.078	0.859±0.047	0.780±0.022	0.759±0.041	0.571±0.036
	PAAC	0.498±0.182	0.901±0.059	0.700±0.066	0.601±0.164	0.439±0.107
	QSO	0.583±0.141	0.890±0.056	0.737±0.047	0.678±0.099	0.503±0.071
	PSSM-composition	0.736±0.046	0.926±0.025	0.831±0.017	0.813±0.023	0.676±0.027
	S-FPSSM	0.714±0.052	0.927±0.023	0.820±0.020	0.798±0.030	0.657±0.033
	RPM-PSSM	0.739±0.059	0.913±0.035	0.826±0.018	0.808±0.028	0.664±0.029
CNN	L100	0.232±0.172	0.903±0.111	0.567±0.034	0.315±0.172	0.194±0.051
	L200	0.301±0.150	0.885±0.096	0.593±0.034	0.403±0.142	0.240±0.044
	L300	0.381±0.159	0.839±0.109	0.610±0.033	0.472±0.136	0.260±0.041
	L400	0.348±0.151	0.870±0.099	0.610±0.033	0.451±0.133	0.268±0.044
	L500	0.352±0.161	0.875±0.095	0.614±0.037	0.454±0.143	0.277±0.050
	L600	0.362±0.156	0.866±0.098	0.614±0.035	0.463±0.131	0.274±0.050
	L700	0.366±0.143	0.868±0.101	0.618±0.031	0.471±0.123	0.282±0.048
	L800	0.358±0.140	0.877±0.092	0.618±0.031	0.467±0.121	0.285±0.042
	L900	0.305±0.138	0.906±0.085	0.606±0.036	0.417±0.131	0.273±0.051
	L1000	0.348±0.153	0.880±0.096	0.614±0.034	0.453±0.134	0.280±0.044
LSTM	L100	0.218±0.132	0.919±0.084	0.570±0.027	0.317±0.134	0.201±0.042
	L200	0.281±0.083	0.917±0.047	0.599±0.023	0.405±0.084	0.260±0.037
	L300	0.293±0.148	0.901±0.088	0.597±0.039	0.400±0.141	0.251±0.062
	L400	0.302±0.118	0.883±0.071	0.593±0.033	0.413±0.110	0.232±0.054
	L500	0.325±0.111	0.874±0.063	0.599±0.033	0.437±0.103	0.241±0.049
	L600	0.315±0.119	0.891±0.054	0.603±0.039	0.429±0.120	0.253±0.066
	L700	0.295±0.120	0.898±0.060	0.596±0.039	0.408±0.122	0.242±0.070
	L800	0.304±0.140	0.900±0.083	0.602±0.035	0.416±0.122	0.260±0.056

	L900	0.326 ± 0.110	0.897 ± 0.047	0.612 ± 0.035	0.445 ± 0.117	0.272 ± 0.062
	L1000	0.337 ± 0.130	0.889 ± 0.069	0.613 ± 0.036	0.450 ± 0.128	0.275 ± 0.056
DNN	L100	0.206 ± 0.046	0.937 ± 0.024	0.571 ± 0.017	0.321 ± 0.055	0.211 ± 0.028
	L200	0.267 ± 0.054	0.930 ± 0.022	0.599 ± 0.019	0.397 ± 0.060	0.264 ± 0.034
	L300	0.259 ± 0.048	0.938 ± 0.023	0.599 ± 0.018	0.390 ± 0.055	0.269 ± 0.032
	L400	0.271 ± 0.050	0.937 ± 0.025	0.604 ± 0.015	0.404 ± 0.055	0.280 ± 0.029
	L500	0.297 ± 0.057	0.931 ± 0.022	0.614 ± 0.020	0.432 ± 0.060	0.295 ± 0.034
	L600	0.307 ± 0.052	0.925 ± 0.029	0.616 ± 0.016	0.442 ± 0.052	0.297 ± 0.027
	L700	0.296 ± 0.059	0.929 ± 0.026	0.613 ± 0.020	0.430 ± 0.062	0.292 ± 0.033
	L800	0.302 ± 0.057	0.924 ± 0.028	0.613 ± 0.020	0.435 ± 0.060	0.290 ± 0.035
	L900	0.293 ± 0.059	0.930 ± 0.024	0.612 ± 0.020	0.427 ± 0.063	0.290 ± 0.032
	L1000	0.286 ± 0.055	0.931 ± 0.024	0.609 ± 0.017	0.419 ± 0.059	0.285 ± 0.032

Note: Values are expressed as mean \pm standard deviation.

Table S7. Performance comparison of different algorithms to construct the final meta model on the 10-fold cross-validation test.

Algorithms	Model	SN	SP	ACC	F-value	MCC
XGBoost	ML model	0.836±0.021	0.916±0.017	0.876±0.013	0.871±0.015	0.755±0.027
	DL model	0.623±0.036	0.739±0.030	0.681±0.020	0.661±0.024	0.365±0.038
	Hybrid model	0.837±0.023	0.918±0.015	0.878±0.014	0.872±0.016	0.758±0.027
MLP	ML model	0.767±0.107	0.946±0.019	0.856±0.053	0.837±0.094	0.726±0.088
	DL model	0.540±0.081	0.792±0.059	0.666±0.022	0.615±0.053	0.346±0.040
	Hybrid model	0.772±0.067	0.947±0.021	0.860±0.026	0.844±0.043	0.732±0.044
DNN	ML model	0.804±0.021	0.908±0.016	0.856±0.013	0.848±0.015	0.716±0.025
	DL model	0.633±0.037	0.730±0.035	0.682±0.020	0.665±0.023	0.366±0.038
	Hybrid model	0.762±0.033	0.864±0.025	0.813±0.015	0.802±0.018	0.630±0.028

Note: Values are expressed as mean±standard deviation. The best performance value is highlighted in bold.

Table S8. Performance comparison between models the meta model using the prediction label and the prediction score of baseline models based on the 10-fold cross validation test.

Feature	SN	SP	ACC	F-value	MCC
Prediction label	0.837±0.023	0.918±0.015	0.878±0.014	0.872±0.016	0.758±0.027
Prediction score	0.837±0.022	0.906±0.019	0.872±0.015	0.867±0.016	0.745±0.030

Table S9. Detailed contributions of different baseline models to the performance of the ultimate meta XGBoost classifier.

Feature ^a	Gain	Cover	Frequency
MLP-S-FPSSM	0.334	0.048	0.016
MLP-RPM-PSSM	0.113	0.031	0.014
XGBOOST-S-FPSSM	0.101	0.035	0.016
CNN-L100	0.096	0.017	0.007
XGBOOST-RPM-PSSM	0.067	0.063	0.029
RF-DDE	0.025	0.047	0.043
SVM-S-FPSSM	0.025	0.039	0.029
SVM-DDE	0.022	0.049	0.053
RF-AAC	0.016	0.080	0.127
SVM-PAAC	0.014	0.024	0.039
MLP-DDE	0.011	0.023	0.015
SVM-PSSM-composition	0.010	0.014	0.022
XGBOOST-AAC	0.009	0.021	0.019
RF-S-FPSSM	0.007	0.015	0.016
MLP-DPC	0.005	0.020	0.013
LSTM-L100	0.005	0.011	0.015
CNN-L600	0.005	0.019	0.019
SVM-AAC	0.005	0.012	0.014
SVM-QSOrder	0.005	0.012	0.015
XGBOOST-PAAC	0.005	0.020	0.016
CNN-L1000	0.005	0.009	0.012
SVM-RPM-PSSM	0.005	0.014	0.018
CNN-L300	0.004	0.017	0.024
CNN-L400	0.004	0.017	0.023
LSTM-L700	0.004	0.010	0.012
LSTM-L1000	0.004	0.017	0.020
CNN-L800	0.004	0.016	0.017
LSTM-L800	0.004	0.011	0.010
RF-PAAC	0.004	0.012	0.012
CNN-L200	0.004	0.014	0.021
CNN-L900	0.004	0.006	0.008
XGBOOST-QSOrder	0.004	0.016	0.017
SVM-DPC	0.004	0.008	0.012
MLP-QSOrder	0.003	0.012	0.008

LSTM-L500	0.003	0.006	0.007
LSTM-L600	0.003	0.013	0.013
DNN-L700	0.003	0.004	0.005
LSTM-L200	0.003	0.010	0.011
CNN-L700	0.003	0.009	0.012
RF-RPM-PSSM	0.003	0.011	0.010
LSTM-L300	0.003	0.010	0.012
LSTM-L400	0.003	0.012	0.013
MLP-PAAC	0.003	0.007	0.007
RF-PSSM-composition	0.003	0.009	0.010
XGBOOST-DPC	0.003	0.011	0.013
CNN-L500	0.003	0.012	0.015
DNN-L900	0.003	0.009	0.011
DNN-L600	0.003	0.007	0.009
LSTM-L900	0.002	0.010	0.010
DNN-L300	0.002	0.009	0.011
DNN-L500	0.002	0.010	0.011
DNN-L200	0.002	0.009	0.011
MLP-AAC	0.002	0.007	0.008
DNN-L100	0.002	0.004	0.005
DNN-L400	0.002	0.005	0.006
XGBOOST-DDE	0.002	0.008	0.009
DNN-L800	0.002	0.006	0.008
DNN-L1000	0.001	0.003	0.004
RF-DPC	0.001	0.005	0.004
MLP-PSSM-composition	0.001	0.006	0.005
RF-QSOrder	0.001	0.004	0.005
XGBOOST-PSSM-composition	0.001	0.004	0.005

Note: Feature^a are denoted as model_feature. For example, MLP_PSSM-composition means that the feature is composed of the predictive labels of the base model constructed by MLP with the PSSM-composition features.

Table S10. Performance comparison of baseline models for VF prediction on the independent test.

Algorithm	Encoding	SN	SP	ACC	F-value	MCC
RF	AAC	0.804	0.809	0.806	0.806	0.613
	DPC	0.774	0.814	0.794	0.790	0.589
	DDE	0.788	0.809	0.799	0.796	0.597
	PAAC	0.804	0.814	0.809	0.808	0.618
	QSO	0.806	0.832	0.819	0.816	0.637
	PSSM-composition	0.726	0.872	0.799	0.783	0.604
	S-FPSSM	0.766	0.861	0.813	0.804	0.63
	RPM-PSSM	0.726	0.847	0.786	0.773	0.577
SVM	AAC	0.868	0.682	0.775	0.794	0.56
	DPC	0.776	0.717	0.747	0.754	0.494
	DDE	0.804	0.658	0.731	0.749	0.467
	PAAC	0.877	0.677	0.777	0.797	0.565
	QSO	0.826	0.743	0.785	0.793	0.571
	PSSM-composition	0.694	0.891	0.793	0.770	0.597
	S-FPSSM	0.819	0.769	0.794	0.799	0.589
	RPM-PSSM	0.785	0.809	0.797	0.794	0.594
XGBoost	AAC	0.802	0.804	0.803	0.803	0.606
	DPC	0.814	0.802	0.808	0.809	0.616
	DDE	0.812	0.800	0.806	0.808	0.613
	PAAC	0.799	0.806	0.802	0.801	0.604
	QSO	0.806	0.818	0.812	0.810	0.623
	PSSM-composition	0.799	0.832	0.815	0.812	0.631
	S-FPSSM	0.795	0.83	0.812	0.809	0.625
	RPM-PSSM	0.797	0.828	0.812	0.810	0.625
MLP	AAC	0.814	0.778	0.796	0.800	0.592
	DPC	0.816	0.785	0.800	0.803	0.601
	DDE	0.806	0.814	0.81	0.809	0.620
	PAAC	0.821	0.778	0.799	0.804	0.600
	QSO	0.809	0.792	0.800	0.802	0.601
	PSSM-composition	0.767	0.851	0.809	0.801	0.62
	S-FPSSM	0.776	0.868	0.822	0.813	0.647
	RPM-PSSM	0.797	0.837	0.817	0.813	0.634
CNN	L100	0.571	0.740	0.655	0.624	0.315
	L200	0.616	0.764	0.69	0.665	0.384
	L300	0.635	0.764	0.700	0.679	0.403

	L400	0.644	0.780	0.712	0.691	0.428
	L500	0.655	0.788	0.721	0.701	0.447
	L600	0.649	0.814	0.732	0.708	0.470
	L700	0.648	0.793	0.72	0.699	0.446
	L800	0.648	0.800	0.724	0.701	0.453
	L900	0.634	0.806	0.72	0.693	0.446
	L1000	0.639	0.799	0.719	0.694	0.443
LSTM	L100	0.59	0.738	0.664	0.637	0.332
	L200	0.587	0.766	0.676	0.644	0.358
	L300	0.592	0.781	0.687	0.654	0.38
	L400	0.639	0.762	0.701	0.681	0.404
	L500	0.594	0.811	0.702	0.666	0.414
	L600	0.609	0.821	0.715	0.682	0.441
	L700	0.606	0.835	0.720	0.684	0.453
	L800	0.62	0.823	0.721	0.690	0.452
	L900	0.594	0.849	0.721	0.681	0.458
	L1000	0.623	0.819	0.721	0.691	0.451
DNN	L100	0.575	0.717	0.646	0.619	0.295
	L200	0.592	0.712	0.652	0.630	0.306
	L300	0.618	0.766	0.692	0.667	0.388
	L400	0.637	0.747	0.692	0.674	0.386
	L500	0.635	0.776	0.706	0.683	0.416
	L600	0.641	0.783	0.712	0.690	0.428
	L700	0.628	0.795	0.712	0.686	0.430
	L800	0.635	0.781	0.708	0.685	0.421
	L900	0.632	0.786	0.709	0.685	0.423
	L1000	0.630	0.793	0.712	0.686	0.429

Note: The best performance value for each metric across different encoding methods within each algorithm is highlighted in bold.

Table S11. Performance comparison of DeepVF trained based on VirulentPred's training dataset and VirulentPred on different independent datasets.

Testing dataset	Algorithm	SN	SP	ACC	F-value	MCC
VirulentPred Ind-I ^a	VirulentPred	0.868	-	-	-	-
	DeepVF	0.868	-	-	-	-
VirulentPred Ind-II ^b	VirulentPred	0.787	-	-	-	-
	DeepVF	0.801	-	-	-	-
The DeepVF independent dataset	VirulentPred	0.641	0.573	0.607	0.620	0.214
	DeepVF	0.668	0.623	0.646	0.654	0.292

Note: ^aVirulentPred Ind-I refers to the independent dataset I of VirulentPred, which contained 38 positive samples (Note that we removed the 22nd whose sequence length was less than 10 and the 25th sample whose sequence had illegal character X) and was downloaded from the VirulentPred webserver (no negative samples available).

^bVirulentPred Ind-II refers to the independent dataset II of VirulentPred, which contained 141 positive samples and was downloaded from the VirulentPred webserver (no negative samples available). The best performance value within each test case is highlighted in bold.

Table S12. Summary of the top 100 sequences predicted by DeepVF and the BLAST-based baseline predictor based on three proteomes.

Proteomes	Number of Proteins ^a	Overlap ^b	DeepVF ^c	BLAST ^d
<i>Escherichia coli</i> O157:H7	4893 (19/3)	12	5	0
<i>Streptococcus pneumoniae</i> (strain ATCC BAA-255 / R6)	1987 (1/2)	14	1	1
<i>Mycobacterium tuberculosis</i> (strain ATCC 25618 / H37Rv)	3616 (58/8)	11	3	0

Note: ^aNumber of proteins within each proteome was counted after removing those sequences that have appeared in the training dataset. The values in the parenthesis represent positive/negative samples contained in the independent dataset/contained negative samples on independent dataset). ^bOverlap represents the number of predicted sequences that appeared in both top 100 ranking lists of DeepVF and the BLAST-based baseline predictor. ^cDeepVF represents the number of positive samples of the independent dataset that appeared in the top 100 ranking list of DeepVF. ^dBLAST represents the number of positive samples of the independent dataset that appeared in the top 100 ranking list of the BLAST-based baseline predictor.

Table S13. Detailed information of DeepVF and the existing toolkits for VF prediction from the user's viewpoint.

Toolkit	Web server reliability ^a	Standalone toolkit	User learning document	Input data format	Number of sequences per submission	Computational cost ^b	Output presentation	Output analysis	Output download
VirulentPred	Sustained	NO	NO	FASTA	<500	~ 11 minutes	Table	None	NO
MP3	Intermittent ^a	YES	YES	FASTA	Unlimited	< 5 seconds	None	None	YES
DeepVF	Sustained	NO	YES	FASTA/ Raw sequence	<5000	~ 0.6 minutes	Table	Result search and rank	YES

^aThe MP3 server has relatively low reliability and could be intermittently accessed.

^bThe computational cost of a web server was evaluated by its executing time per protein sequence, which was obtained by averaging the computational time of 500 test sequences.

2. Supplementary Figures

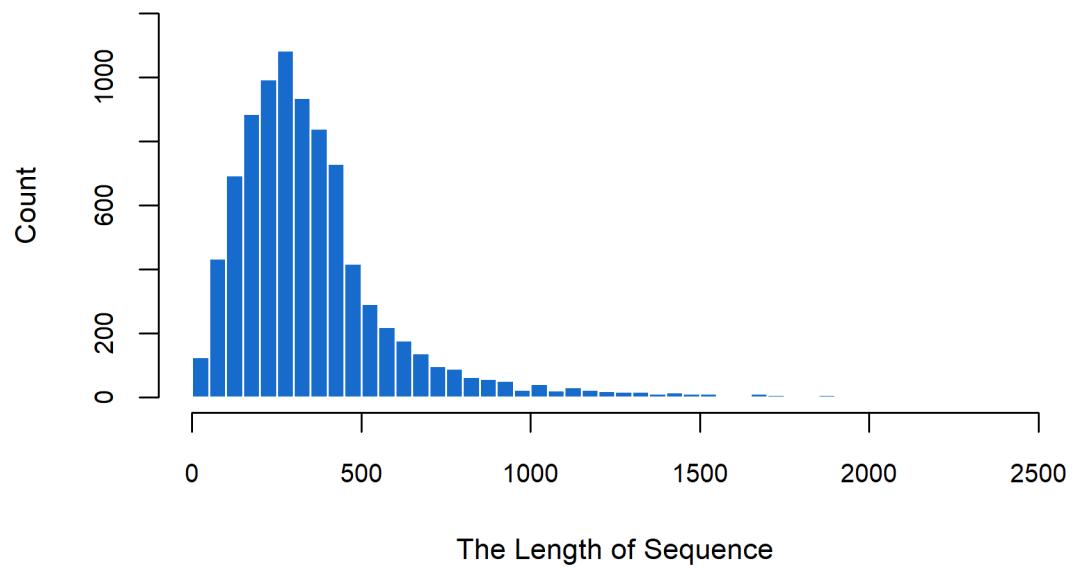


Fig. S1. Distribution of sequence lengths of proteins in the curated dataset.

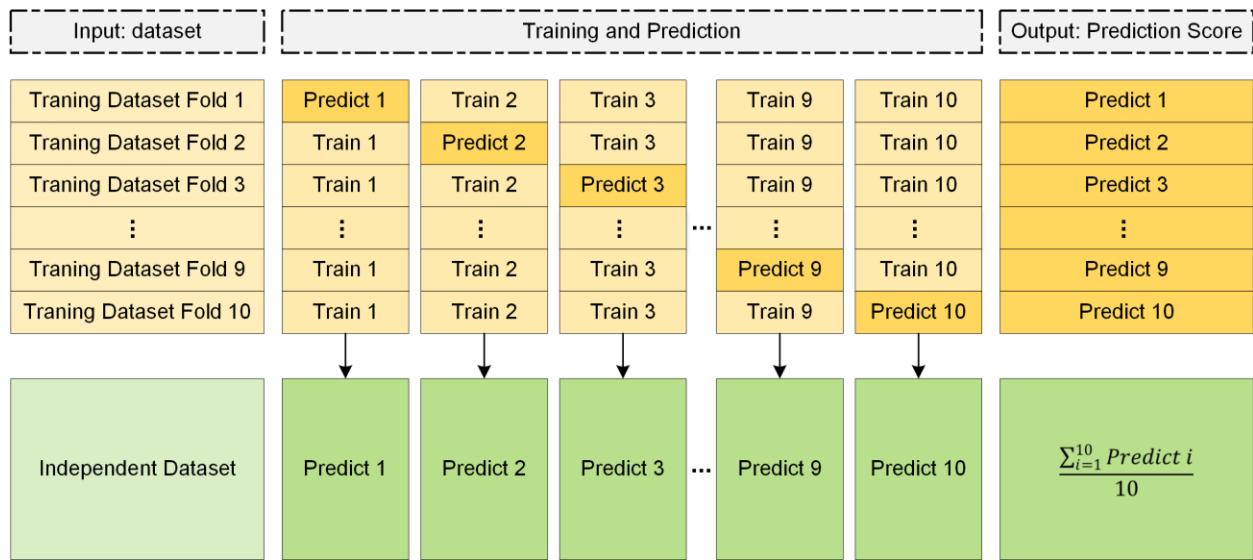


Fig. S2. The pipeline of constructing a baseline model based on the training dataset, and the prediction process on the independent dataset.

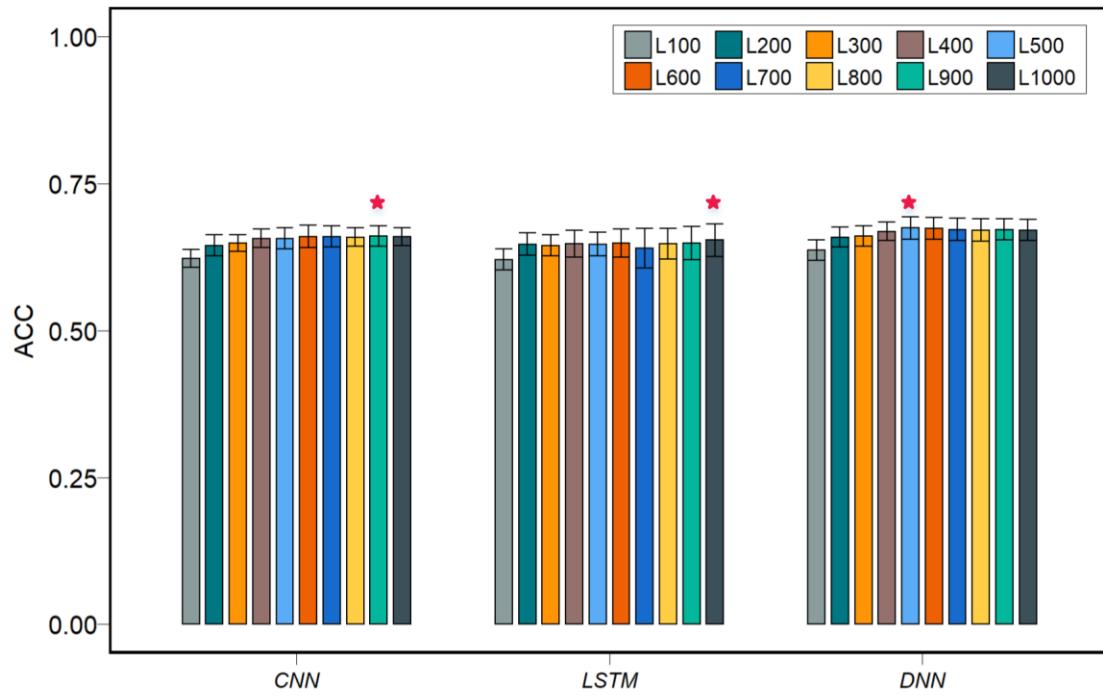


Fig. S3. Performance comparison of different DL models trained using different sizes of protein sequence segments based on the sequence-to-vector feature encoding method.

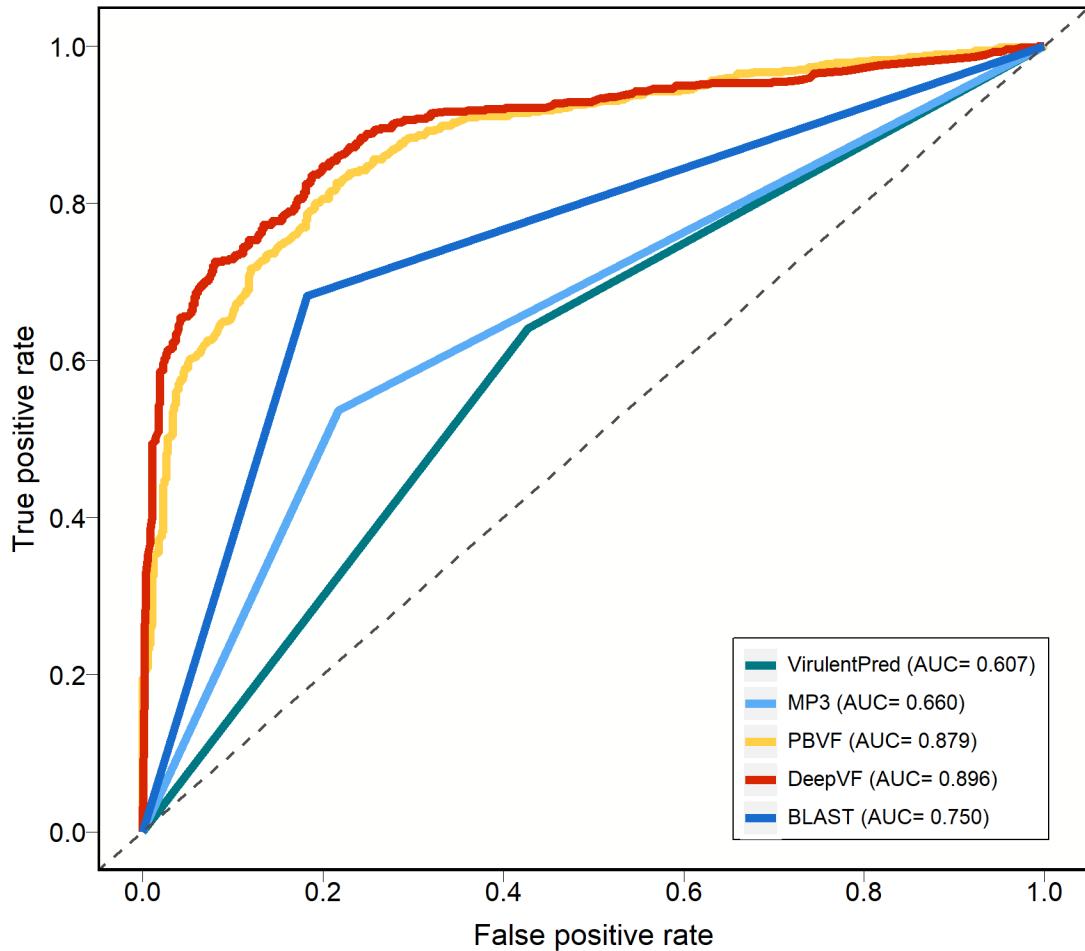


Fig. S4. ROC curves of DeepVF, the BLAST-based baseline predictor and three existing state-of-the-art methods for VF prediction on the independent test. Note that the range of predicting scores of VirulentPred and MP3 are different from that of DeepVF and PBVF, while the BLAST-based baseline predictor only provided a predictive label. Therefore, the ROC curves of those methods were generated by only using their predictive labels, which resulted in three broken lines in this figure.

3. Supplementary References

1. Sachdeva G, Kumar K, Jain P et al. SPAAN: a software program for prediction of adhesins and adhesin-like proteins using neural networks, *Bioinformatics* 2005;21:483-491.
2. Garg A, Gupta D. VirulentPred: a SVM based prediction method for virulent proteins in bacterial pathogens, *BMC bioinformatics* 2008;9:62.
3. Nanni L, Lumini A. An ensemble of support vector machines for predicting virulent proteins, *Expert Systems with Applications* 2009;36:7458-7462.
4. Tsai C-T, Huang W-L, Ho S-J et al. Virulent-GO: prediction of virulent proteins in bacterial pathogens utilizing gene ontology terms, *development* 2009;1:3.
5. Nanni L, Lumini A, Gupta D et al. Identifying bacterial virulent proteins by fusing a set of classifiers based on variants of Chou's pseudo amino acid composition and on evolutionary information, *IEEE/ACM Trans Comput Biol Bioinform* 2012;9:467-475.
6. Zheng LL, Li YX, Ding J et al. A comparison of computational methods for identifying virulence factors, *PLoS One* 2012;7:e42517.
7. Cui W, Chen L, Huang T et al. Computationally identifying virulence factors based on KEGG pathways, *Mol Biosyst* 2013;9:1447-1452.
8. Gupta A, Kapil R, Dhakan DB et al. MP3: a software tool for the prediction of pathogenic proteins in genomic and metagenomic data, *PLoS One* 2014;9:e93907.
9. Rentzsch R, Deneke C, Nitsche A et al. Predicting bacterial virulence factors – evaluation of machine learning and negative data strategies, *Brief Bioinform* 2019.