SUPPLEMENTARY MATERIAL

Splice2Deep: An Ensemble of Deep Convolutional Neural Networks for Improved Splice Site Prediction in Genomic DNA

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Supplementary Table S6. Accuracy (Acc), Sensitivity (Sn), and Specificity (Sp) from the cross-organism validation obtained by using the *Caenorhabditis elegans* model and testing on the other organisms. N/A indicates that the tool is not available for that specific organism.

Splice site	Model	*Homo sapiens*			*Arabidopsis thaliana*			*Oryza sativa japonica*			*Drosophila melanogaster*			*Caenorhabditis elegans*		
		Sp.	*Sn.*	*Acc.*	*Sp.*	*Sn.*	*Acc.*	*Sp.*	*Sn.*	*Acc.*	*Sp.*	*Sn.*	*Acc.*	*Sp.*	*Sn.*	*Acc.*
AcSS	Genesplicer	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	SplicePredictor	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	DeepSS	0.8904	0.8914	0.8909	0.8904	0.8954	0.8929	0.8704	0.8804	0.8919	0.9404	0.9309	0.9207	0.9049	0.9086	0.9332
	Splicerover	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	Splice2Deep	0.9404	0.9414	0.9409	0.9304	0.9354	0.9329	0.9104	0.9134	0.9119	0.9604	0.9509	0.9407	0.9778	0.9838	0.9808
DoSS	Genesplicer	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	SplicePredictor	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	DeepSS	0.9104	0.8934	0.9019	0.9204	0.9054	0.9129	0.8804	0.8904	0.9019	0.9504	0.9309	0.9405	0.9194	0.9168	0.940
	Splicerover	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A	N/A
	Splice2Deep	0.9604	0.9434	0.9519	0.9604	0.9454	0.9529	0.9204	0.9234	0.9219	0.9704	0.9509	0.9497	0.9774	0.9763	0.9768