SUPPLEMENTARY MATERIAL

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

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Supplementary Table S3. Accuracy (Acc), Sensitivity (Sn), and Specificity (Sp) from the cross-organism validation obtained by using the *Arabidopsis thaliana* 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	0.9734	0.4175	0.6955	0.9767	0.7785	0.8776	0.9859	0.6837	0.8348	0.9791	0.4178	0.6984	0.9910	0.3950	0.6930
	SplicePredictor	0.9566	0.5229	0.7398	0.9683	0.8744	0.9213	0.9792	0.7902	0.8847	0.9626	0.6496	0.8061	0.9863	0.6287	0.8075
	DeepSS	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
	Splicerover	0.8847	0.8837	0.8857	0.9312	0.9402	0.9435	0.9030	0.8745	0.8730	0.8787	0.8502	0.8487	0.9049	0.8764	0.8749
	Splice2Deep	0.8993	0.8983	0.9003	0.9486	0.9553	0.9521	0.9131	0.8846	0.8851	0.8872	0.8587	0.8572	0.9201	0.8916	0.8901
DoSS	Genesplicer	0.9685	0.4222	0.6954	0.9758	0.8412	0.9085	0.9866	0.7439	0.8652	0.9817	0.5466	0.7641	0.9928	0.5252	0.7590
	SplicePredictor	0.9546	0.4266	0.6906	0.9604	0.8894	0.9249	0.9776	0.7551	0.8663	0.9651	0.6444	0.8048	0.9850	0.6846	0.8348
	DeepSS	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
	Splicerover	0.8837	0.8827	0.8847	0.9412	0.9502	0.9425	0.9040	0.8755	0.8740	0.8807	0.8522	0.8507	0.9069	0.8784	0.8769
	Splice2Deep	0.8983	0.8973	0.8993	0.9567	0.9550	0.9559	0.9141	0.8856	0.8841	0.8902	0.8617	0.8602	0.9231	0.8946	0.8931