SUPPLEMENTARY MATERIAL

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

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Supplementary Table S1. Sensitivity (Sn) and Specificity (Sp) comparison of Splice2Deep and the state-of-the-art tools for the prediction of SS on the five selected 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.	Sp.	Sn.	Sp.	Sn.	Sp.	Sn.	Sp.	Sn.
AcSS	Genesplicer	97.3	69.28	97.67	77.85	97.5	70.92	93.4	83.93	N/A	N/A
	SplicePredictor	91.86	84.16	96.83	87.44	98.2	80.65	94.92	82.47	N/A	N/A
	DeepSS	92.93	90.18	N/A	N/A	N/A	N/A	N/A	N/A	90.49	90.86
	Splicerover	97.39	90.77	93.12	94.02	N/A	N/A	N/A	N/A	N/A	N/A
	Splice2Deep	97.8	95.61	94.86	95.53	93.62	94.16	95.04	94.09	97.78	98.38
DoSS	Genesplicer	98.6	60.36	97.58	84.12	97.71	74.62	93.7	86.66	N/A	N/A
	SplicePredictor	97.24	79.15	96.04	88.94	97.86	77.14	96.12	81.46	N/A	N/A
	DeepSS	93.63	91.18	N/A	N/A	N/A	N/A	N/A	N/A	91.94	91.68
	Splicerover	96.44	90.01	94.12	95.02	N/A	N/A	N/A	N/A	N/A	N/A
	Splice2Deep	98.83	95.93	95.67	95.5	94.41	94.25	93.71	90.46	97.74	97.63