

Reproduced Metrics for ZM data in Sumukh et al. 2019.

	NM P0	ONNX P0	NM P1	ONNX P1
Precision	0.95	0.95	0.9	0.9
Recall	0.90	0.90	0.95	0.95
F1-Score	0.93	0.93	0.93	0.93
Support	519	519	481	481

Reference:

Deshpande S, Shuttleworth J, Yang J, Taramonli S, England M. PLIT: An alignment-free computational tool for identification of long non-coding RNAs in plant transcriptomic datasets. Comput Biol Med. 2019 Feb;105:169-181. doi: 10.1016/j.combiomed.2018.12.014. Epub 2019 Jan 4. PMID: 30665012.