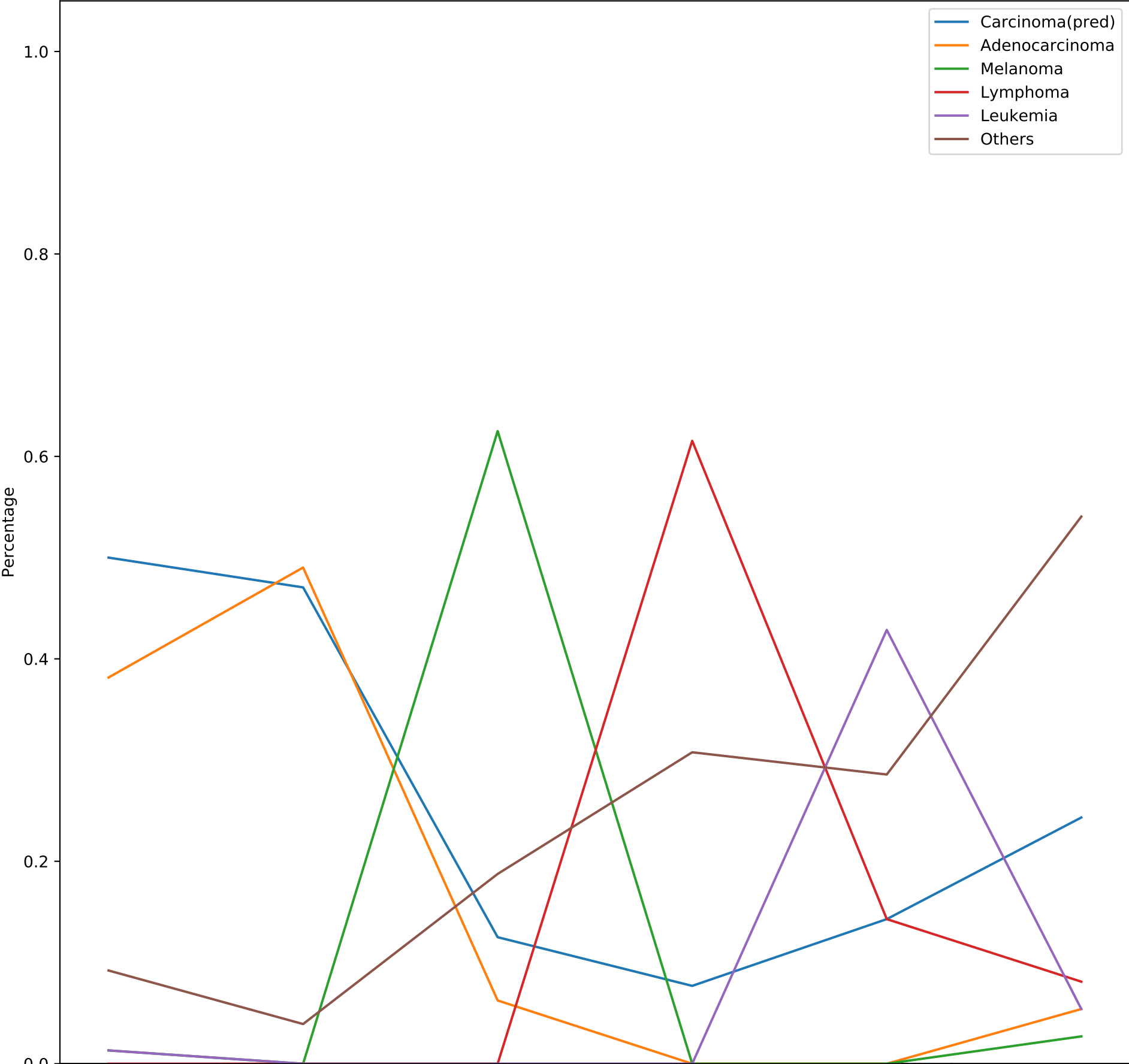


Confusion matrix of combined RNA-seq and long non-coding RNA-seq data



	Carcinoma(pred)	Adenocarcinoma	Melanoma	Lymphoma	Leukemia	Others
Carcinoma(true)	0.5	0.38	0.01	0.0	0.01	0.09
Adenocarcinoma	0.47	0.49	0.0	0.0	0.0	0.04
Melanoma	0.12	0.06	0.62	0.0	0.0	0.19
Lymphoma	0.08	0.0	0.0	0.62	0.0	0.31
Leukemia	0.14	0.0	0.0	0.14	0.43	0.29
Others	0.24	0.05	0.03	0.08	0.05	0.54