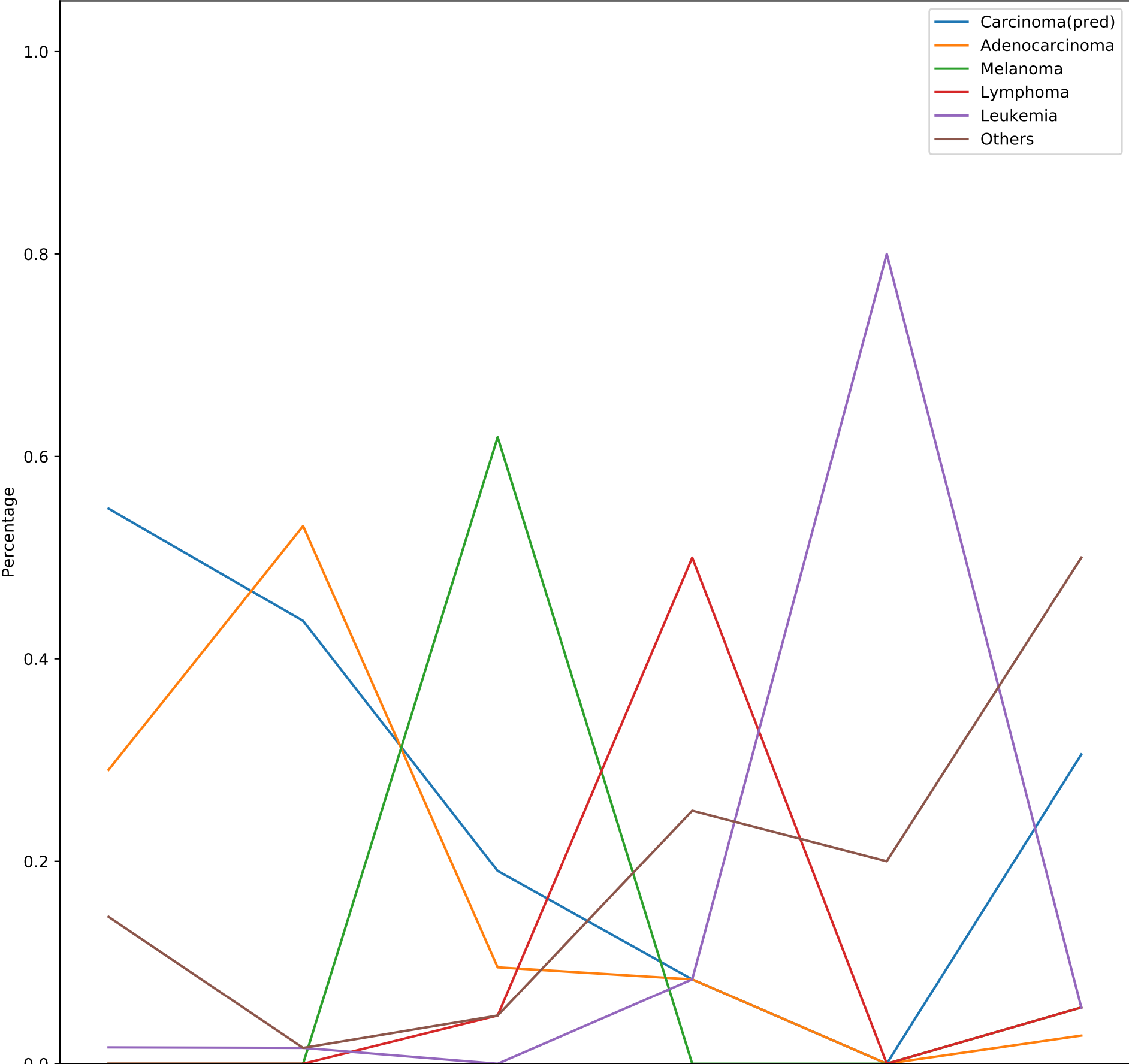


Confusion matrix of combined RNA-seq and copy number variation data



	Carcinoma(pred)	Adenocarcinoma	Melanoma	Lymphoma	Leukemia	Others
Carcinoma(true)	0.55	0.29	0.0	0.0	0.02	0.15
Adenocarcinoma	0.44	0.53	0.0	0.0	0.02	0.02
Melanoma	0.19	0.1	0.62	0.05	0.0	0.05
Lymphoma	0.08	0.08	0.0	0.5	0.08	0.25
Leukemia	0.0	0.0	0.0	0.0	0.8	0.2
Others	0.31	0.03	0.06	0.06	0.06	0.5