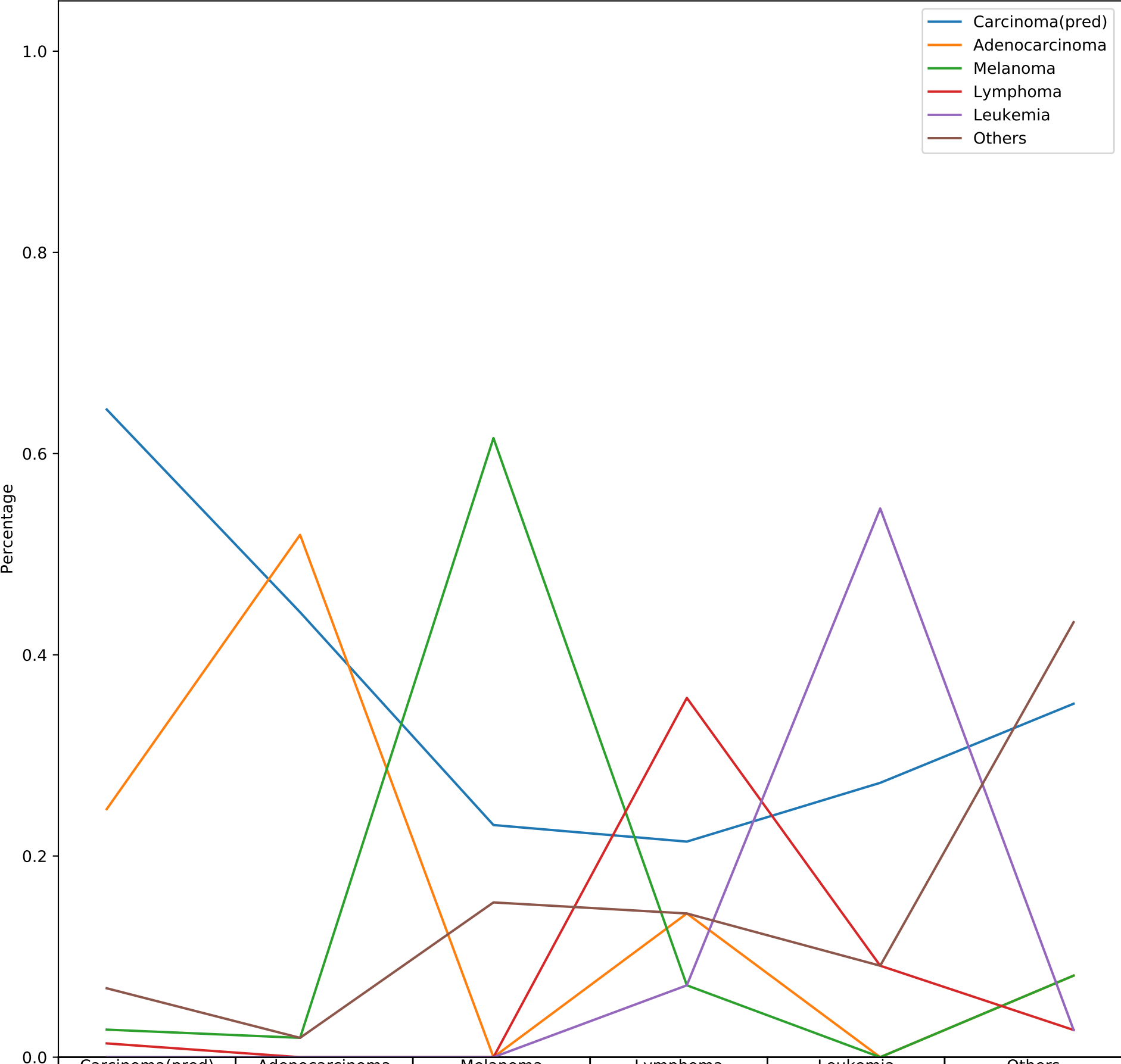


Confusion matrix of combined copy number variation and long non-coding RNA-seq data



| | Carcinoma(pred) | Adenocarcinoma | Melanoma | Lymphoma | Leukemia | Others |
|-----------------|-----------------|----------------|----------|----------|----------|--------|
| Carcinoma(true) | 0.64 | 0.25 | 0.03 | 0.01 | 0.0 | 0.07 |
| Adenocarcinoma | 0.44 | 0.52 | 0.02 | 0.0 | 0.0 | 0.02 |
| Melanoma | 0.23 | 0.0 | 0.62 | 0.0 | 0.0 | 0.15 |
| Lymphoma | 0.21 | 0.14 | 0.07 | 0.36 | 0.07 | 0.14 |
| Leukemia | 0.27 | 0.0 | 0.0 | 0.09 | 0.55 | 0.09 |
| Others | 0.35 | 0.08 | 0.08 | 0.03 | 0.03 | 0.43 |