Confusion-Matrix (Normalized)

	tumor_any	nonTIL_stromal	sTIL	other_nucleus
tumor_any	0.85645	0.06203	0.07709	0.00443
nonTIL_stromal	0.10224	0.64617	0.25	0.0016
sTIL	0.0212	0.09037	0.88174	0.00669
other_nucleus	0.36437	0.40081	0.18623	0.04858

Actual Classes vs Predicted Classes