Confusion-Matrix (Normalized)

	tumor_any	nonTIL_stromal	sTIL	other_nucleus
tumor_any	0.88923	0.03633	0.07444	0.0
nonTIL_stromal	0.21885	0.54313	0.23802	0.0
sTIL	0.06062	0.08479	0.85459	0.0
other_nucleus	0.47773	0.2996	0.22267	0.0

Actual Classes / Predicted Classes