

# Supplementary Table 1 – Task design for crowdsourced glioma cell annotation in microscopy images

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## ABSTRACT

### Methods – Machine learning

Supplementary Table 1 shows an estimation about computational complexity of the single processing steps (for machine learning processes, we used the required time for training instead of complexity).

**Table 1.** Computational complexity of each image analysis step and time to process machine learning tools. The process steps are described in Methods – Machine learning section of main document. Training of the models was performed for 1000 epochs.

Process step	Complexity	Time
Background removal	O(n)	–
Flood-fill algorithm	O(n)	–
Preliminary models (single cell type)	–	~1.5 hours
Combining both results	O(n)	–
Final model	–	~4 hours