



Apr 15, 2019

aCGH results extraction from TIFF file

 [PLOS One](#)

DOI

<https://dx.doi.org/10.17504/protocols.io.zj7f4rn>

Hendrik F van Essen¹

¹Amsterdam UMC



Hendrik F HFvE Van Essen

Amsterdam UMC

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.zj7f4rn>

External link: <https://doi.org/10.1371/journal.pone.0223827>

Protocol Citation: Hendrik F van Essen 2019. aCGH results extraction from TIFF file. **protocols.io**

<https://dx.doi.org/10.17504/protocols.io.zj7f4rn>

Manuscript citation:

Vincenten JPL, Essen HFv, Lissenberg-Witte BI, Bulkman NWJ, Krijgsman O, Sie D, Eijk PP, Smit EF, Ylstra B, Thunnissen E (2019) Clonality analysis of pulmonary tumors by genome-wide copy number profiling. PLoS ONE 14(10): e0223827. doi: [10.1371/journal.pone.0223827](https://doi.org/10.1371/journal.pone.0223827)



License: This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working

We use this protocol and it's working

Created: March 27, 2019

Last Modified: April 15, 2019

Protocol Integer ID: 21855



- 1 Copy TIFF file to "output folder"
- 2 Open "Feature Extraction" (Fe) program
- 3 File → 'Add extraction set'
- 4 Select TIFF files in "output folder"
- 5 Select under 'project properties' tab the correct grid name
 - * 4×44K
 - * 2×105K
 - * 4×180k
- 6 Under Project select 'Start extracting'