

Attachment 1A: The Cancer Genome Atlas (TCGA) Data User Platform Survey

Cover Email: In order to help data repository managers at CGHub and The Cancer Genome Atlas (TCGA) develop the best client tool, please answer the following one question survey.

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Public reporting burden for this collection of information is estimated to average 1minute per response, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing the collection of information. **An agency may not conduct or sponsor, and a person is not required to respond to, a collection of information unless it displays a currently valid OMB control number.** Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden, to: NIH, Project Clearance Branch, 6705 Rockledge Drive, MSC 7974, Bethesda, MD 20892-7974, ATTN: PRA (0925-0642). Do not return the completed form to this address.

Question 1: When downloading the raw sequencing files from the Cancer Genome

Atlas, which operating system do you utilize? Check all that apply

- a) Mac
- b) PC
- c) Unix or Linux
- d) Other. Specify_____.

Thank you for your participation.