Running head: SUBTITLE

1

Sally E. Claridge¹, Daniel M. Charytonowicz¹, & Adam A. Margolin^{1, 2}

¹ Department of Genetics and Genomic Sciences, Icahn Institute for Data Science and

Manuscript

1

- Genomic Technology, Icahn School of Medicine at Mount Sinai, New York, NY
- ² Cancer Target Discovery and Development Network (Oregon Health and Science
- 6 University, Portland, OR), National Cancer Institute, National Institutes of Health

Author Note

- Add complete departmental affiliations for each author here. Each new line herein
- 9 must be indented, like this line.
- Enter author note here.
- 11 Correspondence concerning this article should be addressed to Adam A. Margolin,
- 12 Icahn School of Medicine at Mount Sinai, Department of Genetics and Genomic Sciences,
- One Gustave L. Levy Place, Box 1498, New York, NY 10029. E-mail:
- adam.margolin@mssm.edu

15 Abstract

One or two sentences providing a **basic introduction** to the field, comprehensible to a scientist in any discipline.

- Two to three sentences of **more detailed background**, comprehensible to scientists in related disciplines.
- One sentence clearly stating the **general problem** being addressed by this particular study.
- One sentence summarizing the main result (with the words "here we show" or their equivalent).
- Two or three sentences explaining what the **main result** reveals in direct comparison to what was thought to be the case previously, or how the main result adds to previous knowledge.
- One or two sentences to put the results into a more **general context**.
- Two or three sentences to provide a **broader perspective**, readily comprehensible to a scientist in any discipline.
- 30 Keywords: keywords
- Word count: X

32	Manuscript
33	(Aksoy et al., 2017)
34	Introduction
35	Results
36	Discussion

37 Methods

VennDiagram (R-VennDiagram) for all our analyses.

```
We used R (R-base) and the R-packages bsselectR (R-bsselectR), cowplot

(R-cowplot), data.table (R-data.table), dplyr (R-dplyr), forcats (R-forcats),

futile.logger (R-futile.logger), ggplot2 (R-ggplot2), ggpubr (R-ggpubr), ggrepel

(R-ggrepel), ggsignif (R-ggsignif), gplots (R-gplots), gridExtra (R-gridExtra),

htmlwidgets (R-htmlwidgets), kableExtra (R-kableExtra), magrittr (R-magrittr),

papaja (Version 0.1.0.9842; Aust & Barth, 2018), plotly (R-plotly), purrr (R-purrr),

readr (R-readr), reshape2 (R-reshape2), scales (R-scales), shiny (R-shiny), stringr

(R-stringr), tibble (R-tibble), tidyr (R-tidyr), tidyverse (R-tidyverse), and
```

47 References

 $_{\tt 48}$ $\,$ Finally, the APA guidelines require a note at the start of the reference section that explains

what an asterisk means. This note can be added at the end of the document as

follows.

F1	References
51	References

- Aksoy, B. A., Dančík, V., Smith, K., Mazerik, J. N., Ji, Z., Gross, B., . . . Clemons, P. A.
- 53 (2017). CTD2 Dashboard: A searchable web interface to connect validated results
- from the Cancer Target Discovery and Development Network. *Database (Oxford)*,
- ⁵⁵ 2017. doi:10.1093/database/bax054
- ⁵⁶ Aust, F., & Barth, M. (2018). papaja: Create APA manuscripts with R Markdown. R
- package version 0.1.0.9842. Retrieved from https://github.com/crsh/papaja