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Preface

This is a Quarto book.

To learn more about Quarto books visit <https://quarto.org/docs/books>.

[1][1] [2]

1 Introduction

This is a book created from markdown and executable code.

See^[3] for additional discussion of literate programming.

2 Summary

In summary, this book has no content whatsoever.

References

1. Powers, R. K., Goodspeed, A., Pielke-Lombardo, H., Tan, A.-C., & Costello, J. C. (2018). GSEA-InContext: identifying novel and common patterns in expression experiments. *Bioinformatics*, 34(13), i555–i564. <https://doi.org/10.1093/bioinformatics/bty271>
2. Subramanian, A., Tamayo, P., Mootha, V. K., Mukherjee, S., Ebert, B. L., Gillette, M. A., Paulovich, A., Pomeroy, S. L., Golub, T. R., Lander, E. S., & Mesirov, J. P. (2005). Gene set enrichment analysis: A knowledge-based approach for interpreting genome-wide expression profiles. *Proceedings of the National Academy of Sciences*, 102(43), 15545–15550. <https://doi.org/10.1073/pnas.0506580102>
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