Programs

Sequence	Filename	Description
1	weswgs.sh	WES/WGS preprocessing
2.1	rva.sh	Rare-variant analysis
2.2	spa.sh	Single-point analysis

idmap.do, ngs.wrap, weswgs.R, prune.wrap, rva.sb, spa.sb are subprograms; and remarks on variant lists submitted centrally are described in INTERVAL.md.

The natural order is therefore
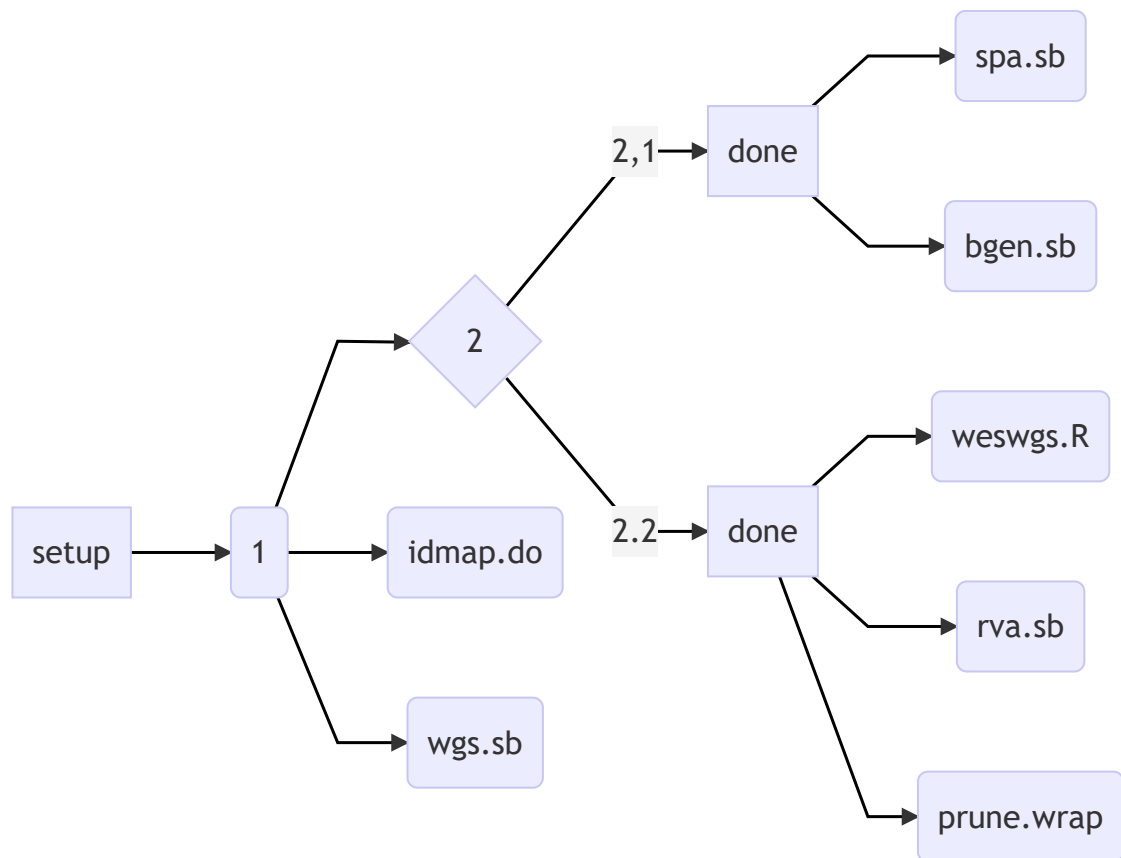
noting in particular that sbatch implicates the --wait option as the succeeding steps would require its full results.

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Slack channel

- https://scallop-seq.slack.com

URLs

- https://github.com/hmgu-itg/burden_testing
- https://sites.google.com/site/jpopgen/dbNSFP
- https://sites.google.com/site/jpopgen/wgsa
- http://web.corral.tacc.utexas.edu/WGSAdownload/resources/precomputed_hg38/
- http://www.columbia.edu/~ii2135/Eigen_functions_112015.R