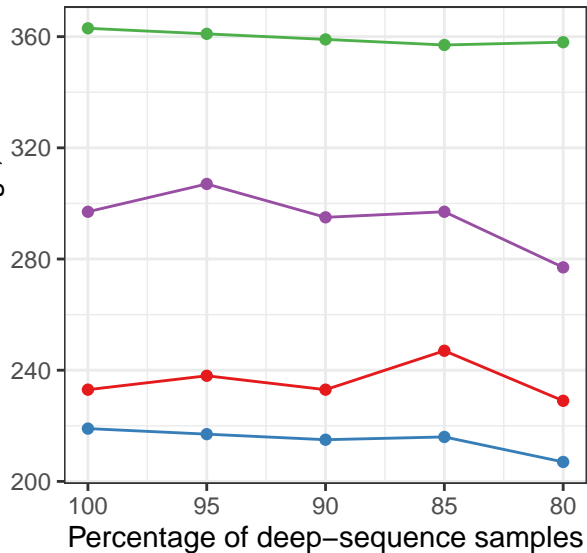
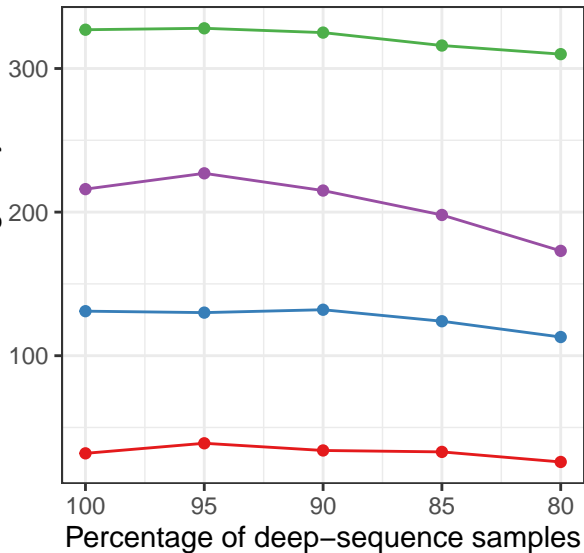


Number of findings, raw P



Number of findings, adjusted P



Gene stratum

- EE_index
- mean_index
- mean_var_index
- var_index