- What is a key feature of genes that may be falsely flagged as positively selected?

  - (Very) low proportions of (very) high omega values!

The magnitude of ω in the positive selection component

ω_3 **magnitude** by selection detection type

Low when STANDARD and +ERROR agree there's NO selection

~100 (or higher) for STANDARD on cases where STANDARD finds selection and +ERROR does not

Very few cases where +ERROR finds selection and STANDARD does not; reasonable values

Mostly below 10 when STANDARD and +ERROR agree there IS selection