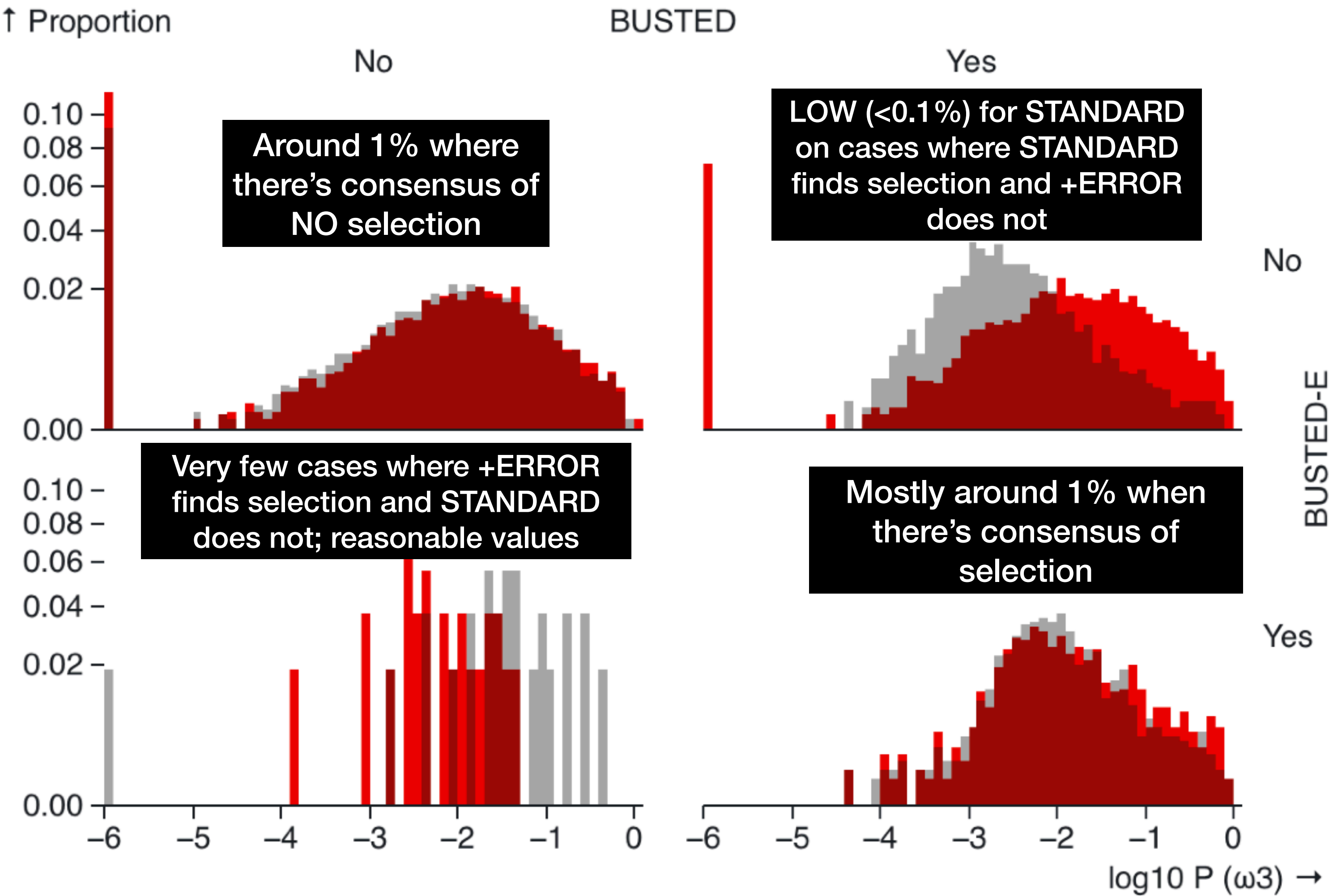


The **weight** of  $\omega$  in the positive selection component

$\omega_3$  **weight** by selection detection type

■ BUSTED ■ BUSTED-E

↑ Proportion



- BUSTED-E can annotate each input alignment with what it identifies as potentially erroneous codon positions (specific sequence, specific column)
- These can be further filtered (e.g. replaced with ‘- - -’) for downstream analyses with other tools
- The stringency of filtering is tunable