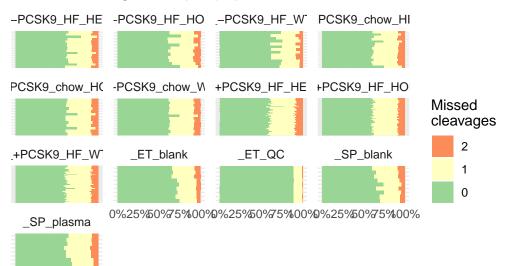
## Missed cleavage sites per peptide



0%25%50%75%100%