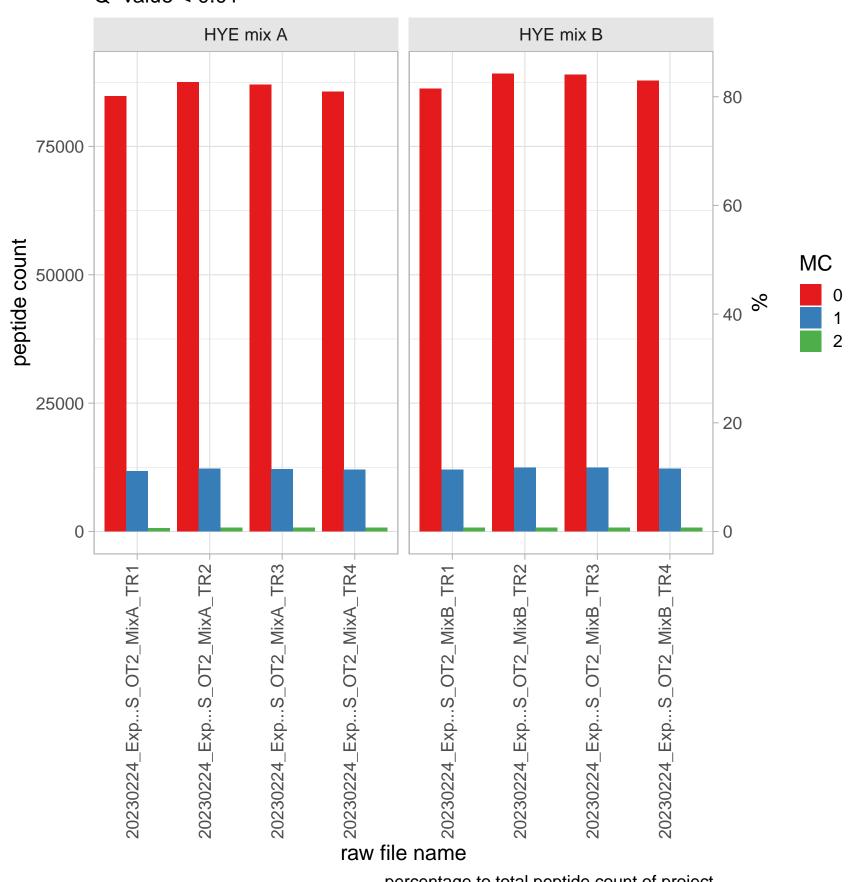
count of missed cleavages per sample Q-value < 0.01



percentage to total peptide count of project