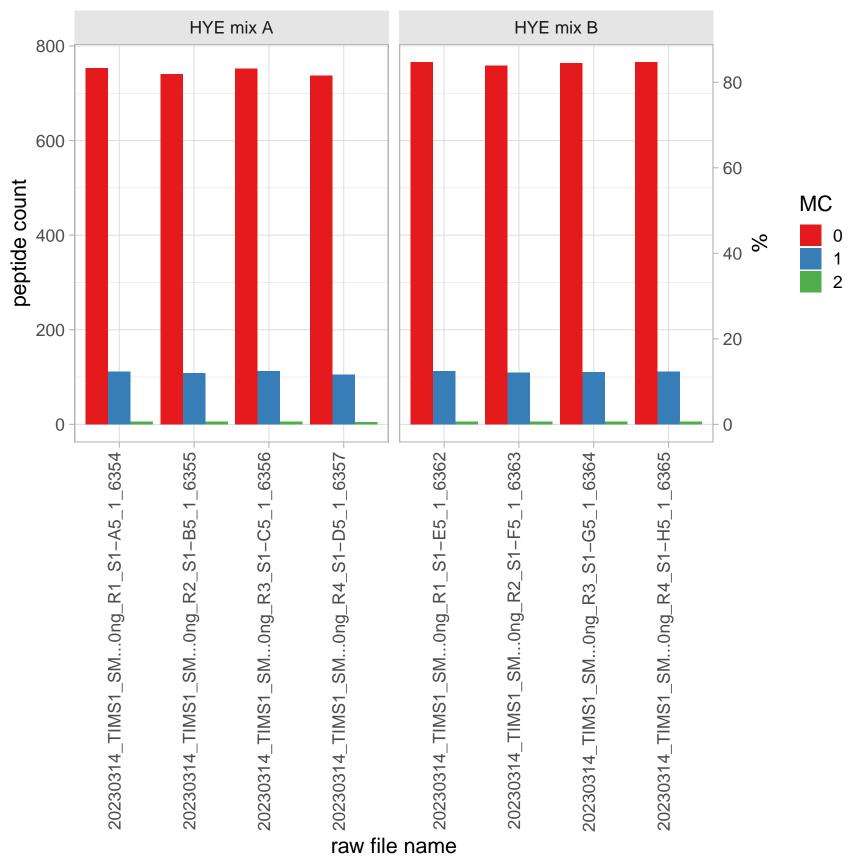count of missed cleavages per sample
Q-value < 0.01