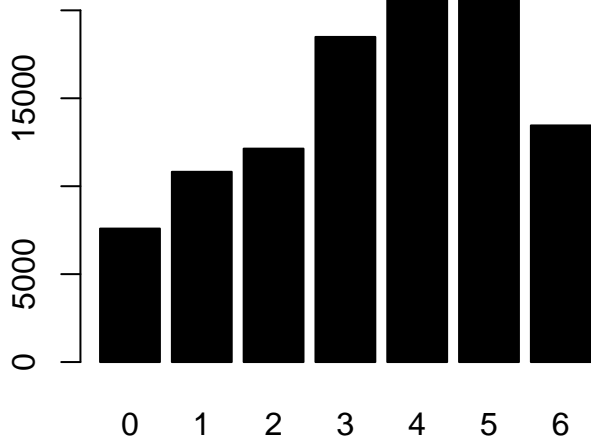


Count of Possible Sequencing Sites



Number of Individuals with at Least One Read