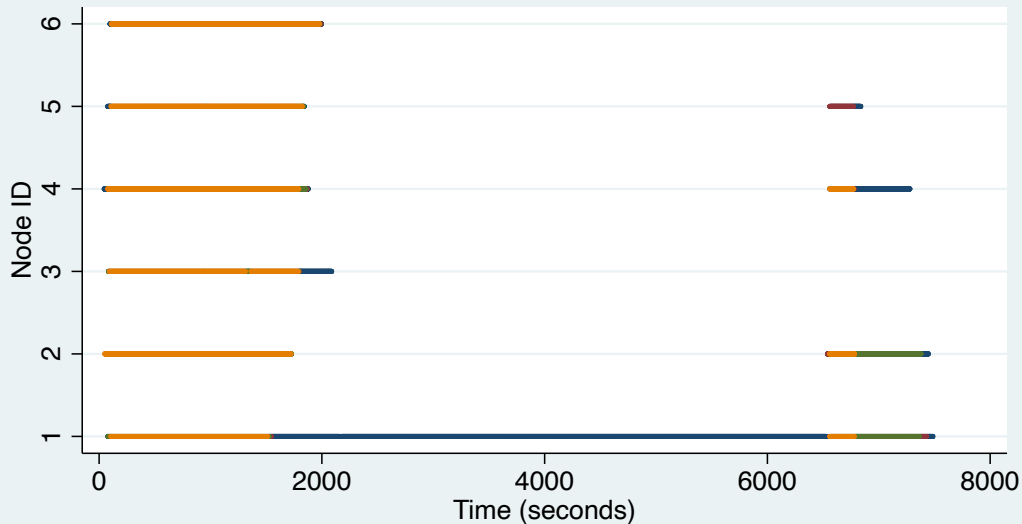


# Genome Workflow

20k data and 6 nodes



• 1 Active Thread

• 2 Active Thread

• 3 Active Thread

• 4 Active Thread