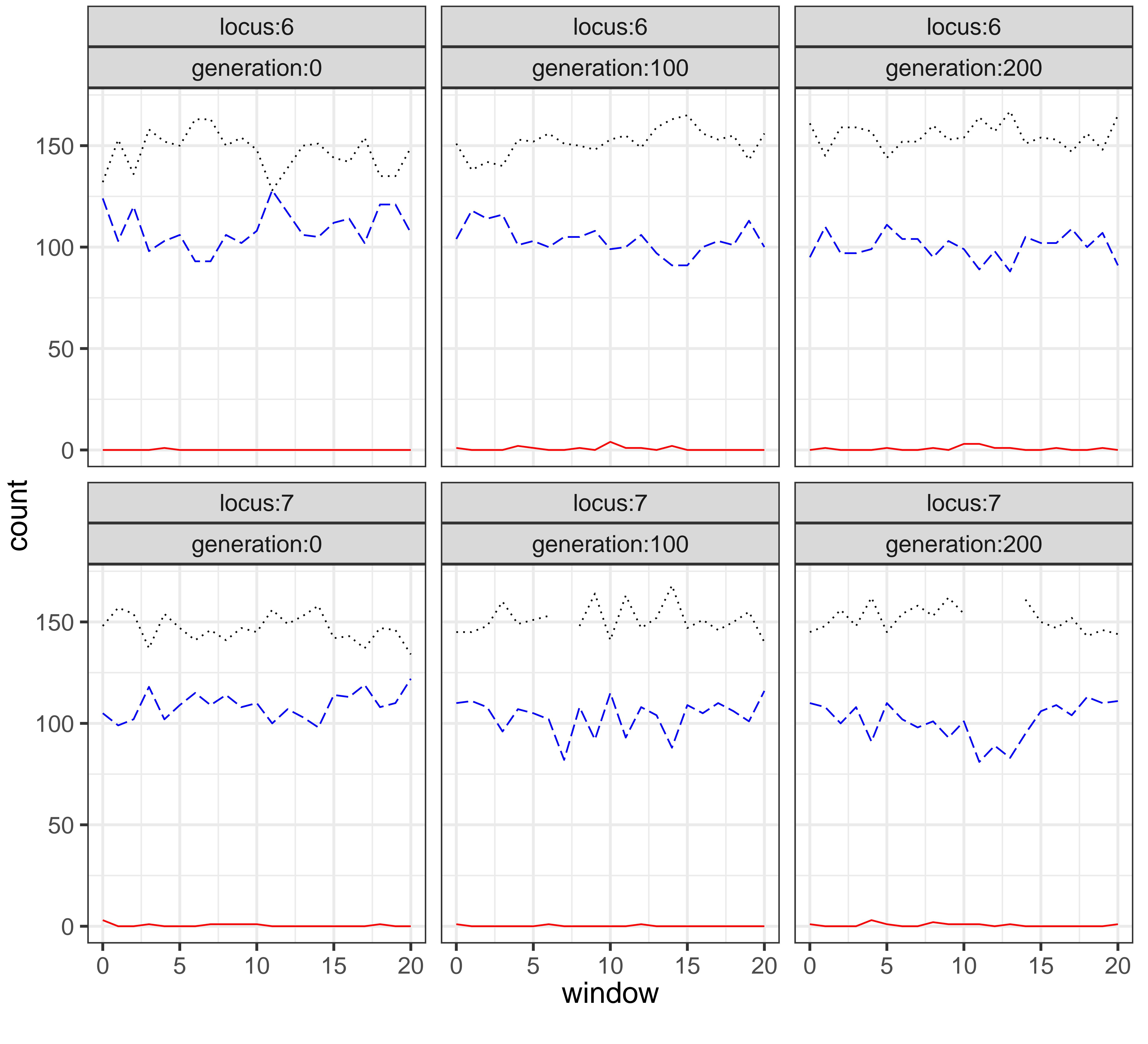
high mutation rate-medium alpha



classification — hard+linkedHard -- neutral soft+linkedSoft