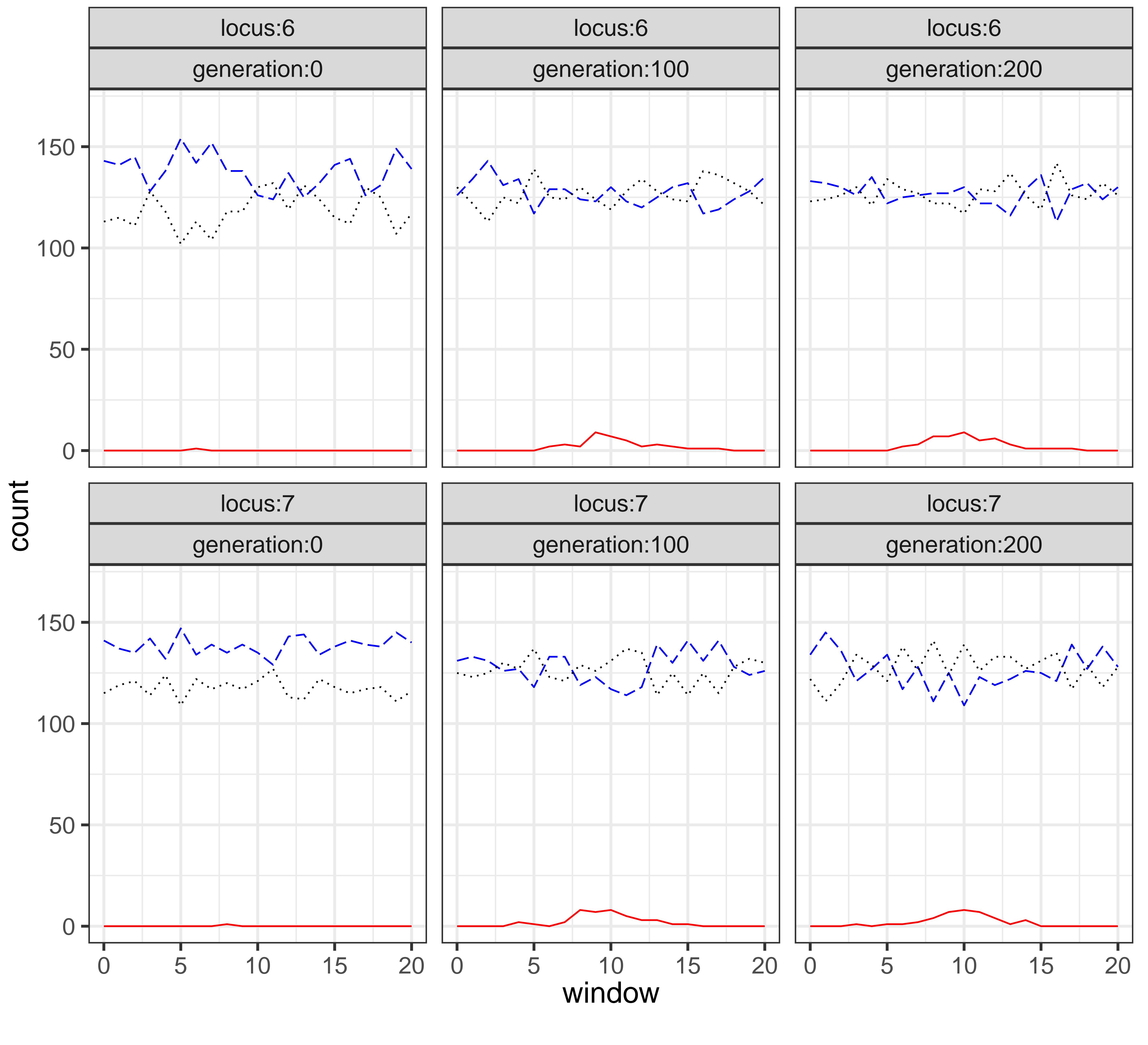
medium mutation rate-high alpha



classification — hard+linkedHard -- neutral soft+linkedSoft