

Selection temp

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Selection analysis

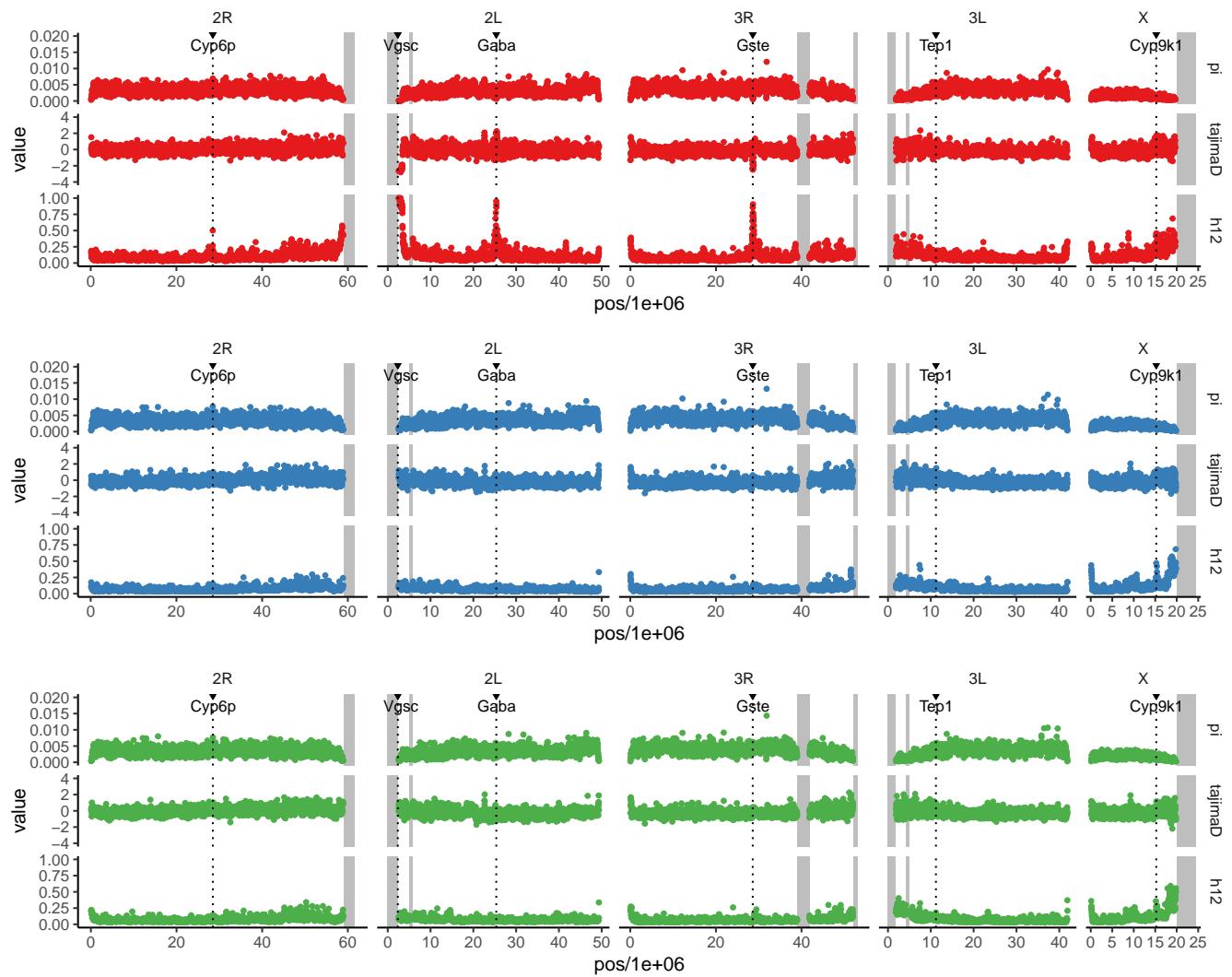
```
popStatsDescWindows.py --snp wilding.unifiedGenotyper.cov14x.passQC.phased.zarr/ -b bed -o LBVwil --keep
popStatsDescWindows.py --snp wilding.unifiedGenotyper.cov14x.passQC.phased.zarr/ -b bed -o LPdom --keep
popStatsDescWindows.py --snp wilding.unifiedGenotyper.cov14x.passQC.phased.zarr/ -b bed -o LPfor --keep

fst_scan.py --zarr wilding.unifiedGenotyper.cov14x.passQC.phased.zarr/ --pop1 LPdom.ind --pop2 LPfor.ind
fst_scan.py --zarr wilding.unifiedGenotyper.cov14x.passQC.phased.zarr/ --pop1 LPdom.ind --pop2 LBVwil.ind
fst_scan.py --zarr wilding.unifiedGenotyper.cov14x.passQC.phased.zarr/ --pop1 LPfor.ind --pop2 LBVwil.ind

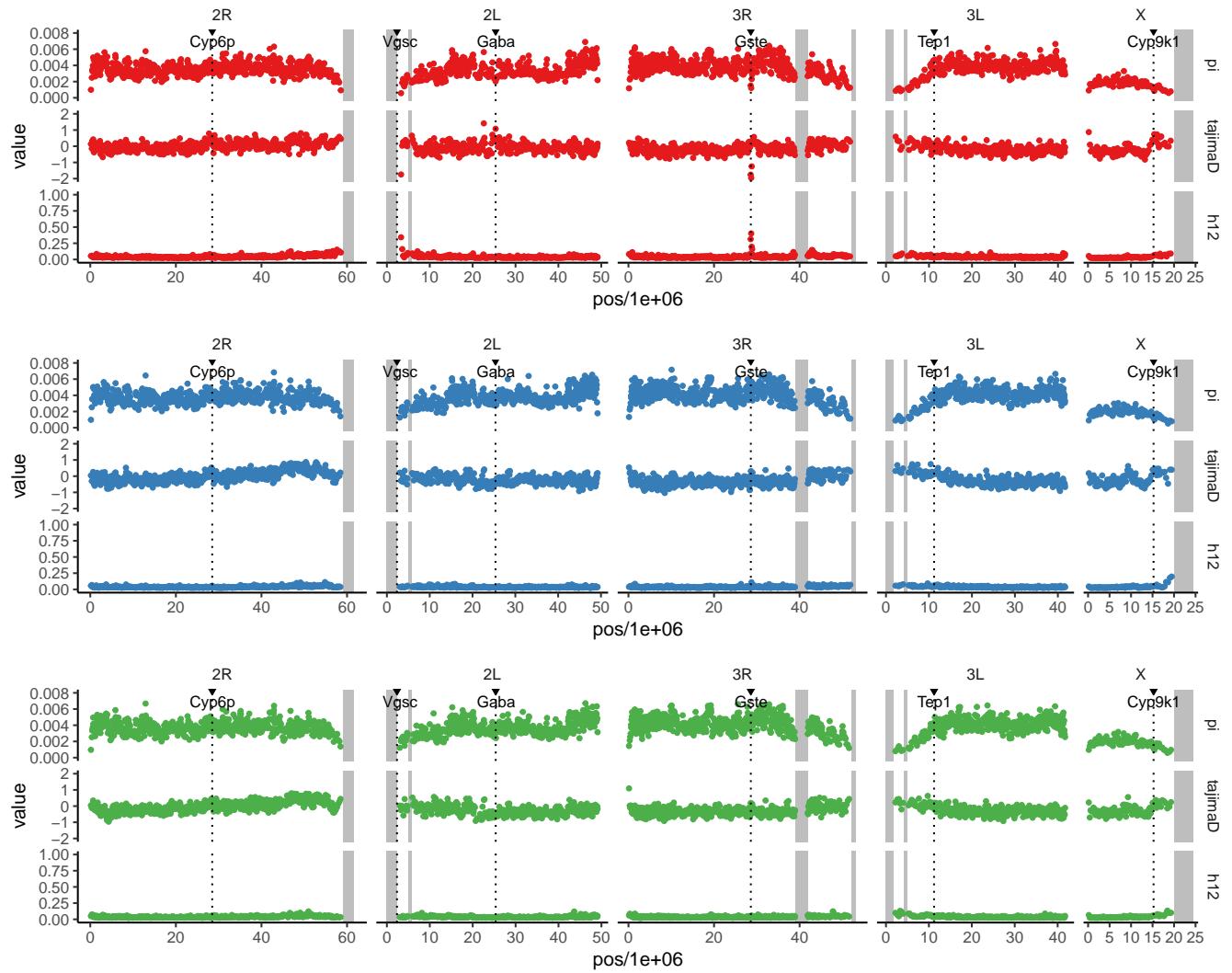
getPBS_scan.py --pop1 LPdom.ind --pop2 LPfor.ind --pop3 LBVwil.ind --zarr wilding.unifiedGenotyper.cov14x.ind
getPBS_scan.py --pop1 LBVwil.ind --pop2 LPdom.ind --pop3 LPfor.ind --zarr wilding.unifiedGenotyper.cov14x.ind
getPBS_scan.py --pop1 LBVwil.ind --pop2 LPfor.ind --pop3 LPdom.ind --zarr wilding.unifiedGenotyper.cov14x.ind
```

1pop stat

windows size = 20kb



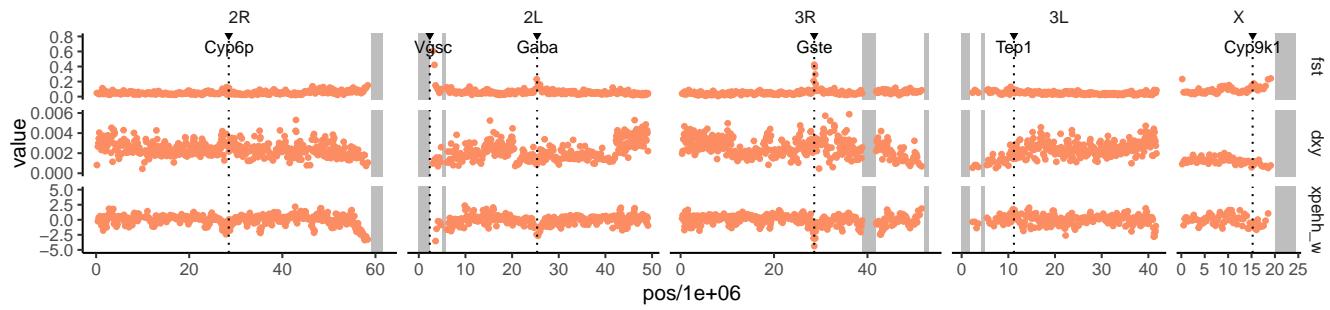
windows size = 1000snp



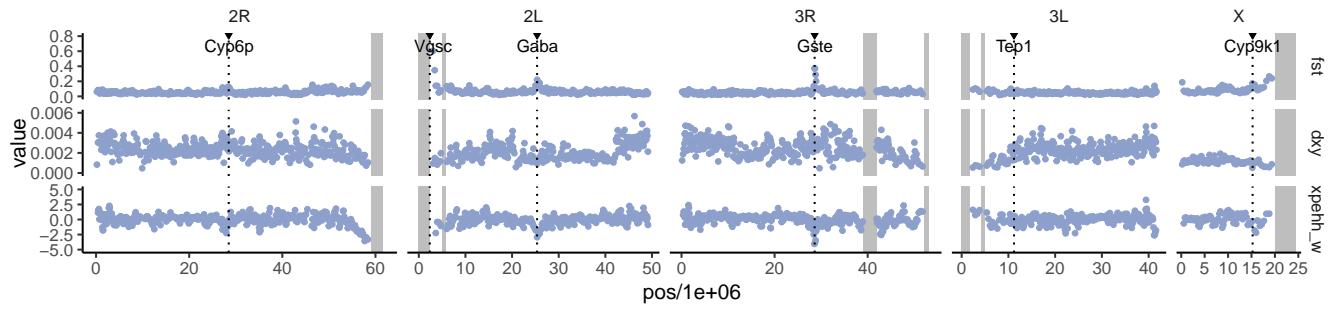
2pop stat

windows size = 1000snp

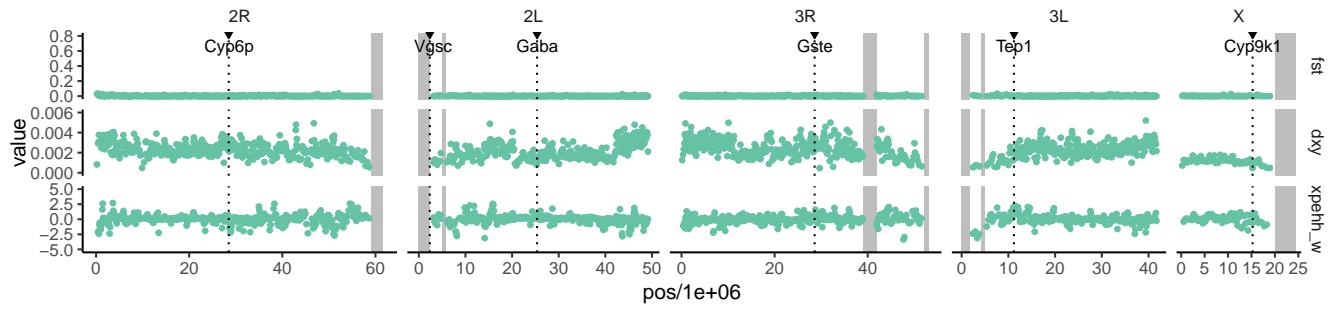
LPdom_LBVwil



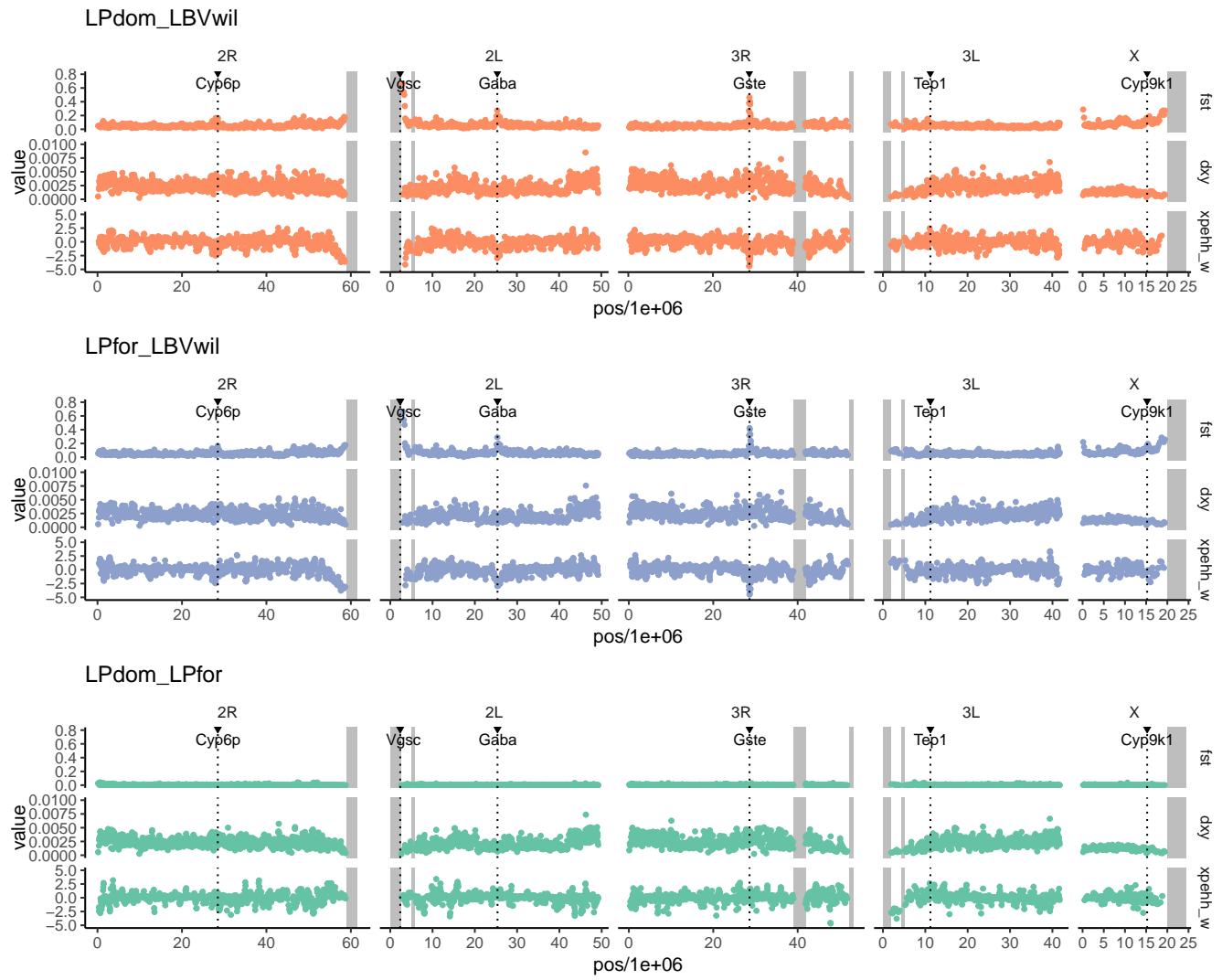
LPfor_LBVwil



LPdom_LPfor

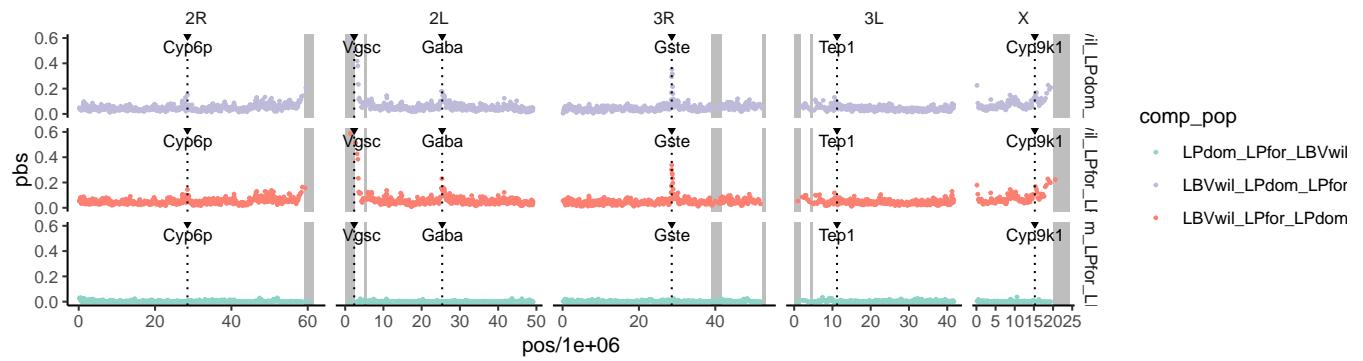


windows size = 500snp

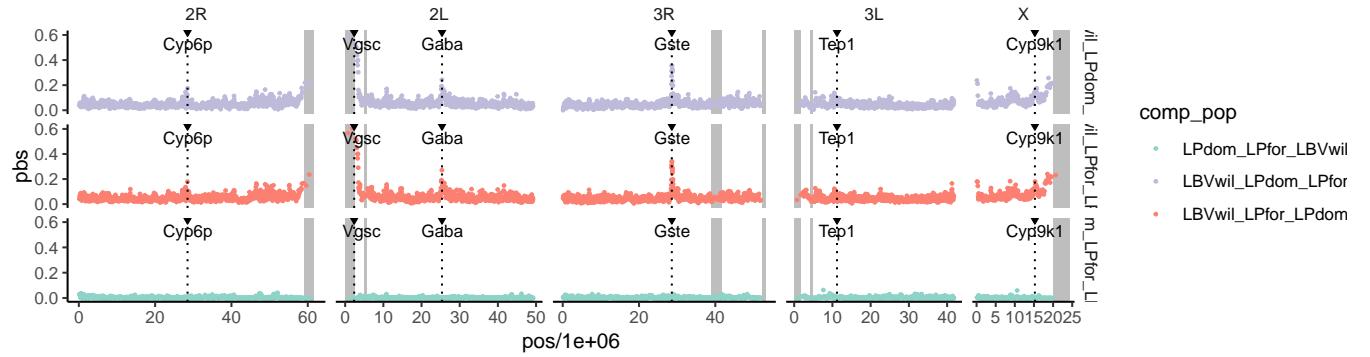


PBS

windows size = 1000snp



windows size = 500snp



windows size = 250snp

