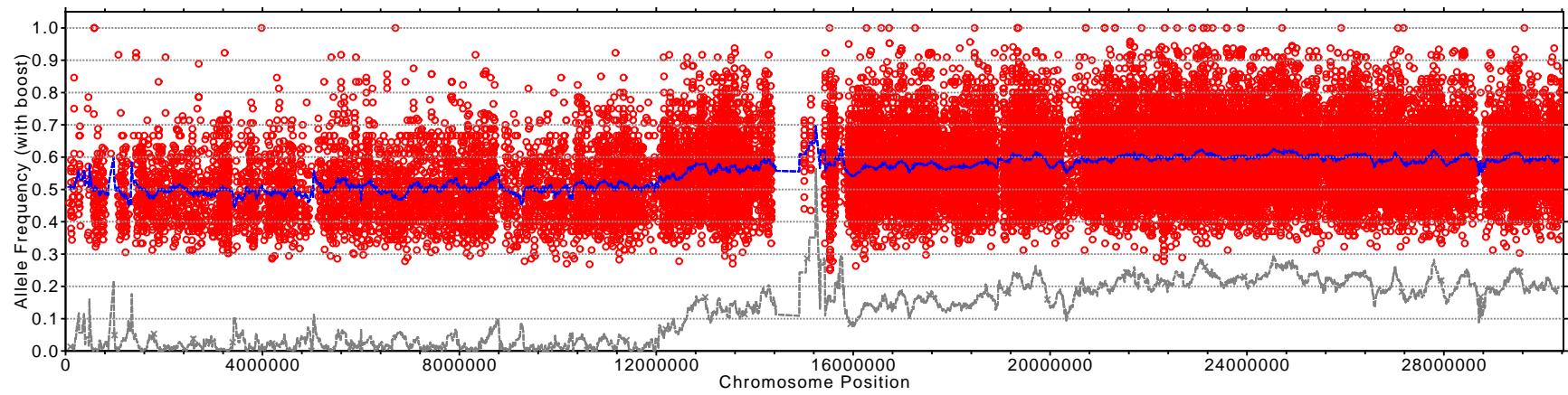


Chromosome 1: 30427671bp



\*Parameter settings:

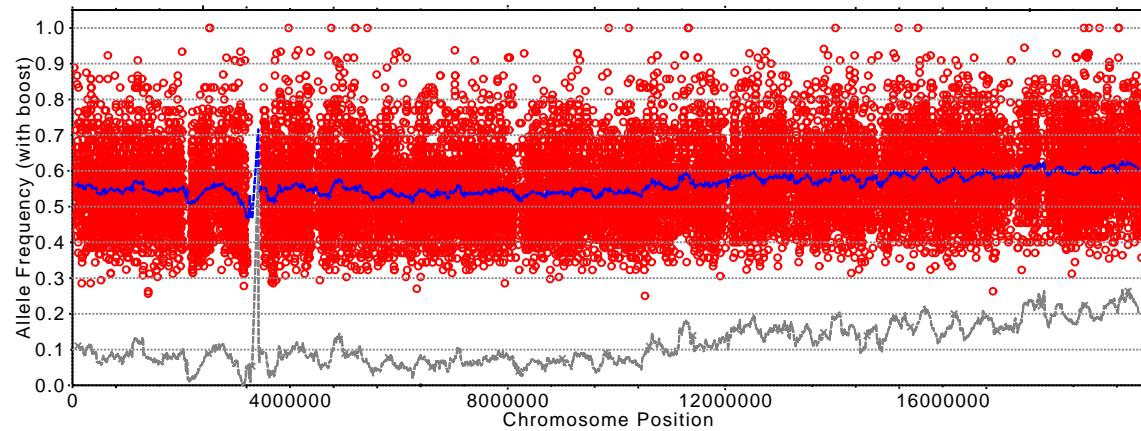
interval-mean-min~max 0.9900~1.0000 with Cv-max 0.0100; window-size 200000 with step 5000; marker-min 5; coverage-min~max 10~80

\*Points in colors: AF (=alt divided by (alt+ref)) at markers. alt or ref: coverage of non-reference or reference allele

\*Dashed line in blue: window-based AF (= summation-of-single-marker-AF divided by number-of-markers with min quality score involved).

\*Dashed line in gray: window-based boost value

Chromosome 2: 19698289bp



\*Parameter settings:

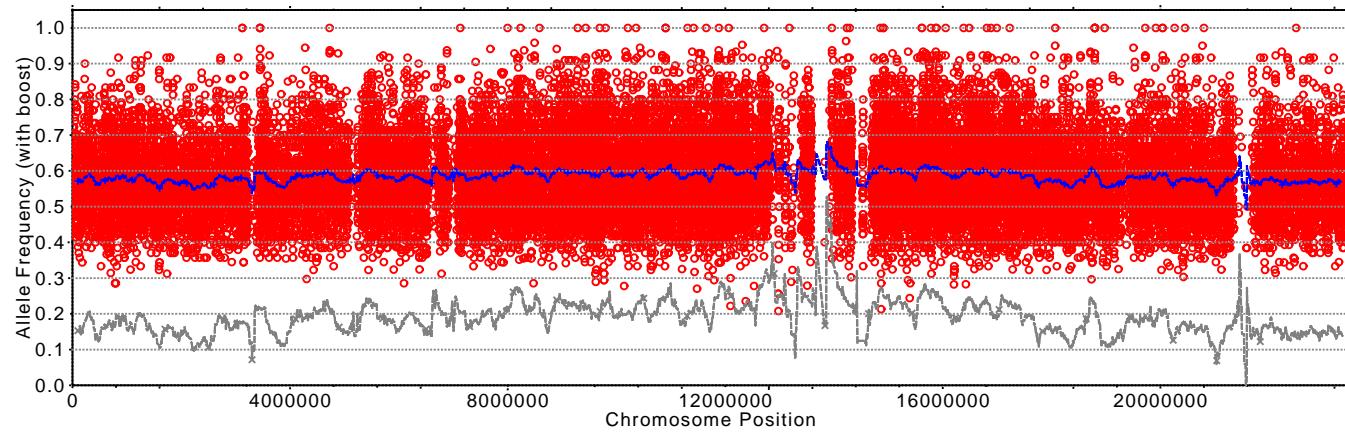
interval-mean-min~max 0.9900~1.0000 with Cv-max 0.0100; window-size 200000 with step 5000; marker-min 5; coverage-min~max 10~80

\*Points in colors: AF (=alt divided by (alt+ref)) at markers. alt or ref: coverage of non-reference or reference allele

\*Dashed line in blue: window-based AF (= summation-of-single-marker-AF divided by number-of-markers with min quality score involved).

\*Dashed line in gray: window-based boost value

Chromosome 3: 23459830bp



\*Parameter settings:

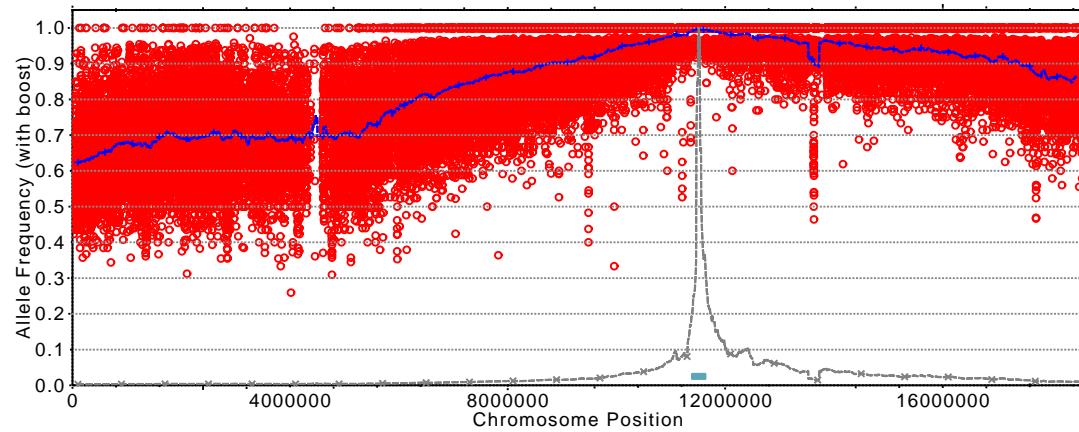
interval-mean-min~max 0.9900~1.0000 with Cv-max 0.0100; window-size 200000 with step 5000; marker-min 5; coverage-min~max 10~80

\*Points in colors: AF (=alt divided by (alt+ref)) at markers. alt or ref: coverage of non-reference or reference allele

\*Dashed line in blue: window-based AF (= summation-of-single-marker-AF divided by number-of-markers with min quality score involved).

\*Dashed line in gray: window-based boost value

Chromosome 4: 18585056bp



\*Parameter settings:

interval-mean-min~max 0.9900~1.0000 with Cv-max 0.0100; window-size 200000 with step 5000; marker-min 5; coverage-min~max 10~80

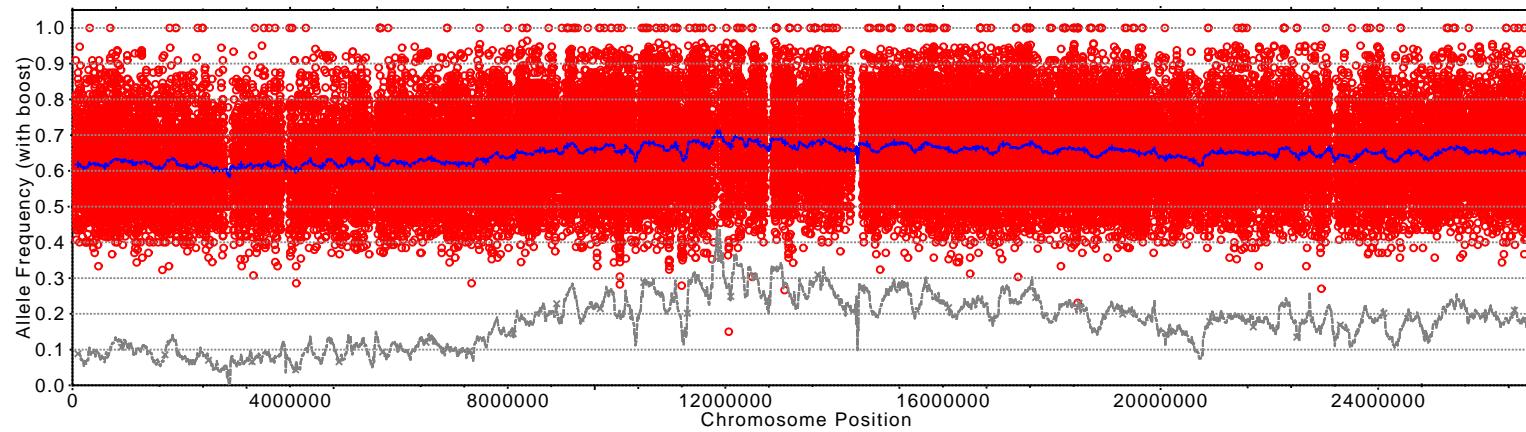
\*Points in colors: AF (=alt divided by (alt+ref)) at markers. alt or ref: coverage of non-reference or reference allele

\*Dashed line in blue: window-based AF (= summation-of-single-marker-AF divided by number-of-markers with min quality score involved).

\*Dashed line in gray: window-based boost value

\*Predicted mapping interval of size 230.00 Kbp (normal): 11400000 ~ 11629999

Chromosome 5: 26975502bp



\*Parameter settings:

interval-mean-min~max 0.9900~1.0000 with Cv-max 0.0100; window-size 200000 with step 5000; marker-min 5; coverage-min~max 10~80

\*Points in colors: AF (=alt divided by (alt+ref)) at markers. alt or ref: coverage of non-reference or reference allele

\*Dashed line in blue: window-based AF (= summation-of-single-marker-AF divided by number-of-markers with min quality score involved).

\*Dashed line in gray: window-based boost value