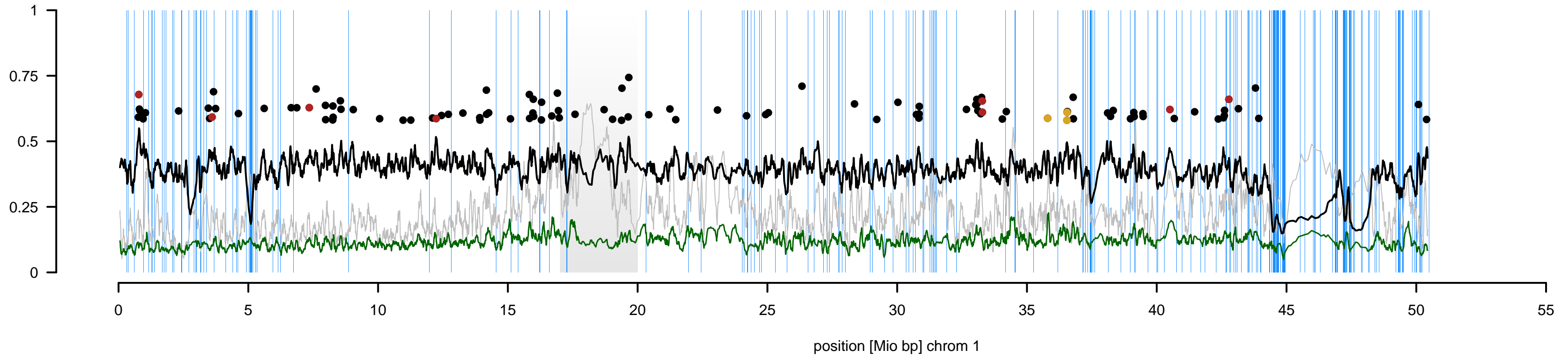
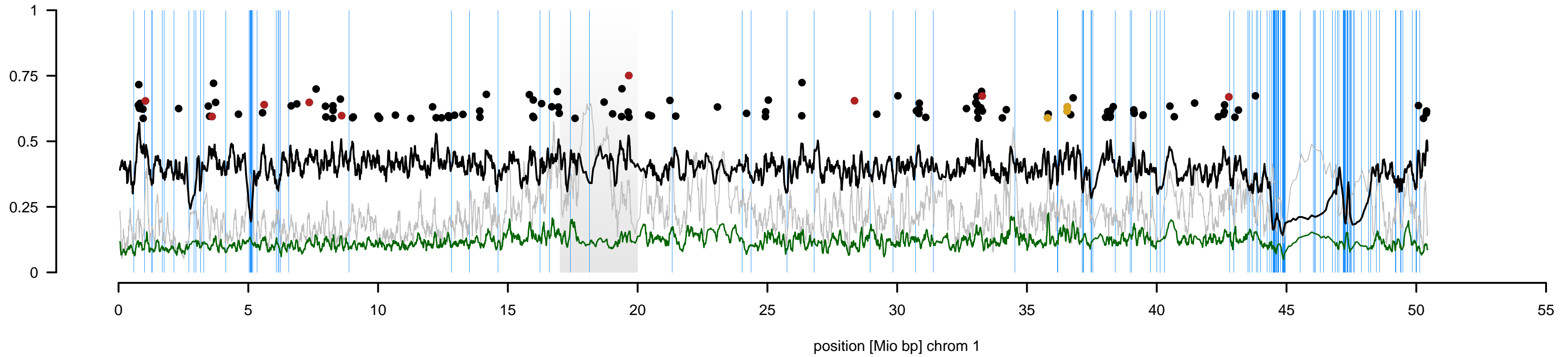


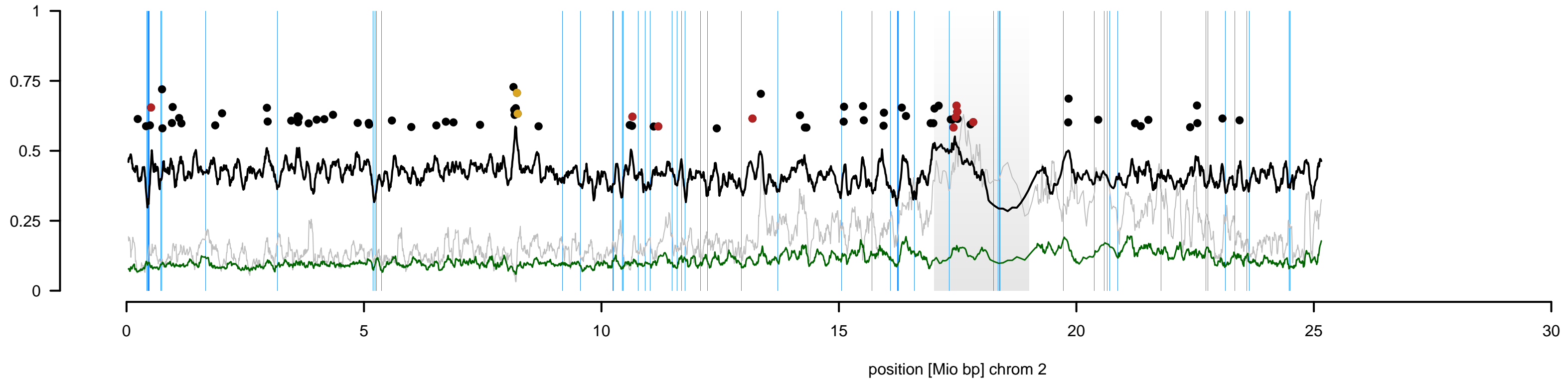
Italy interspecific comparisons – 8k windows



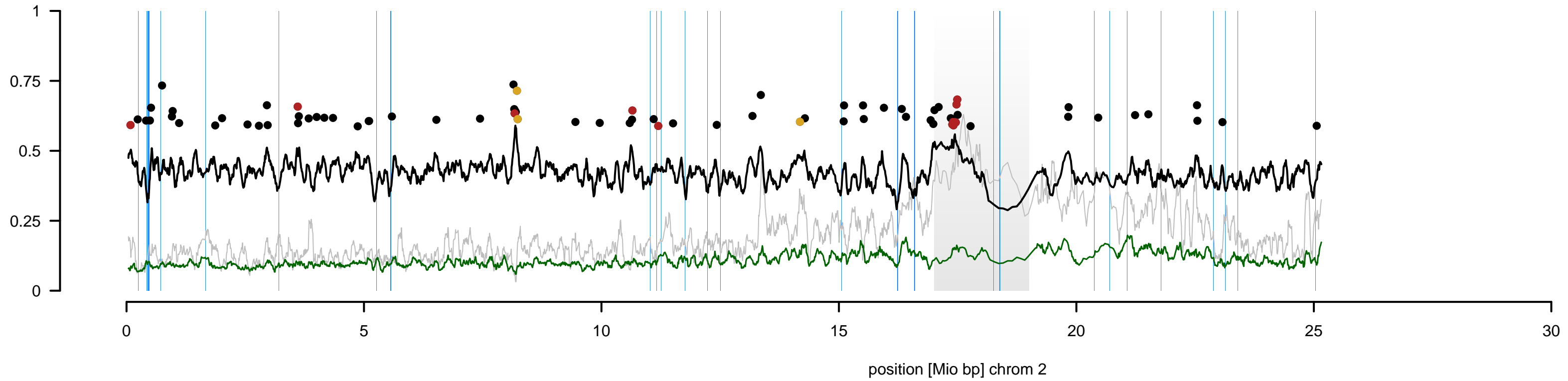
Hungary interspecific comparisons – 8k windows



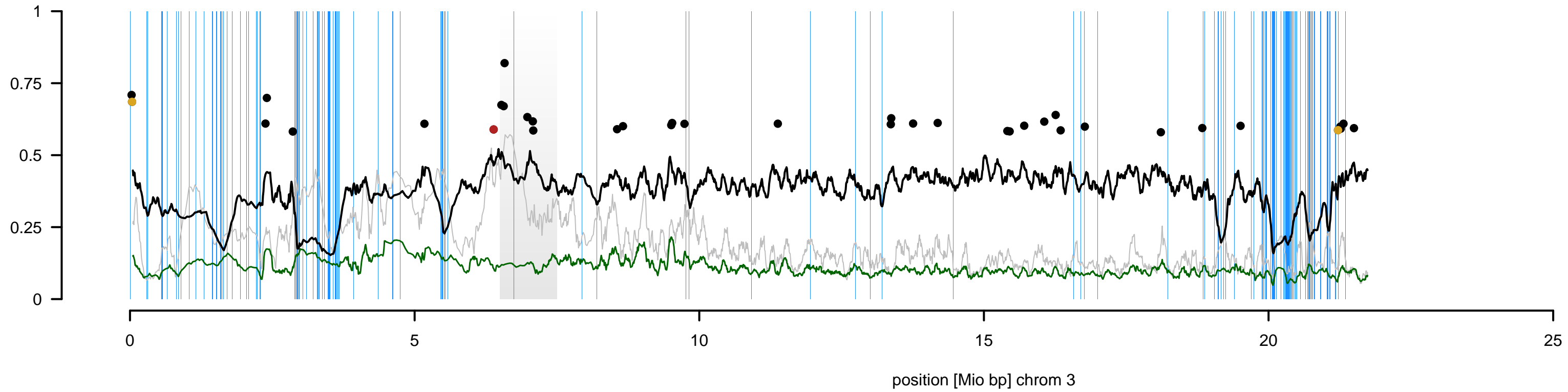
Italy interspecific comparisons – 8k windows



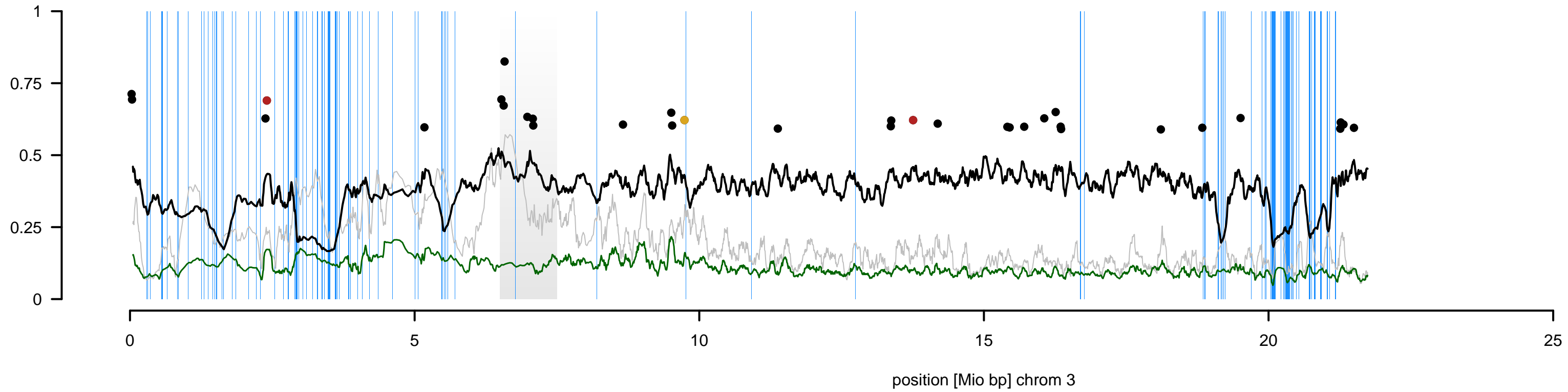
Hungary interspecific comparisons – 8k windows



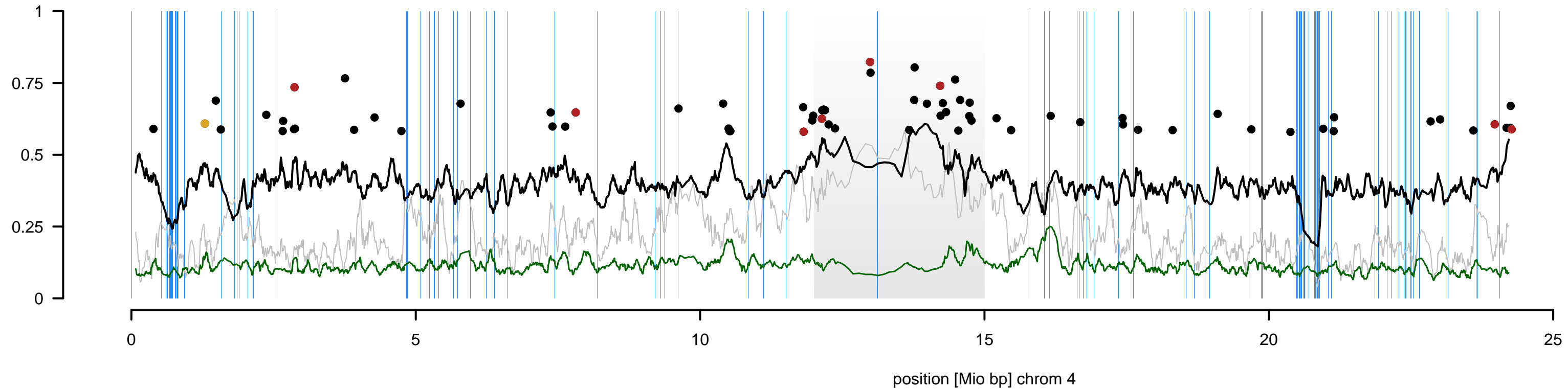
Italy interspecific comparisons – 8k windows



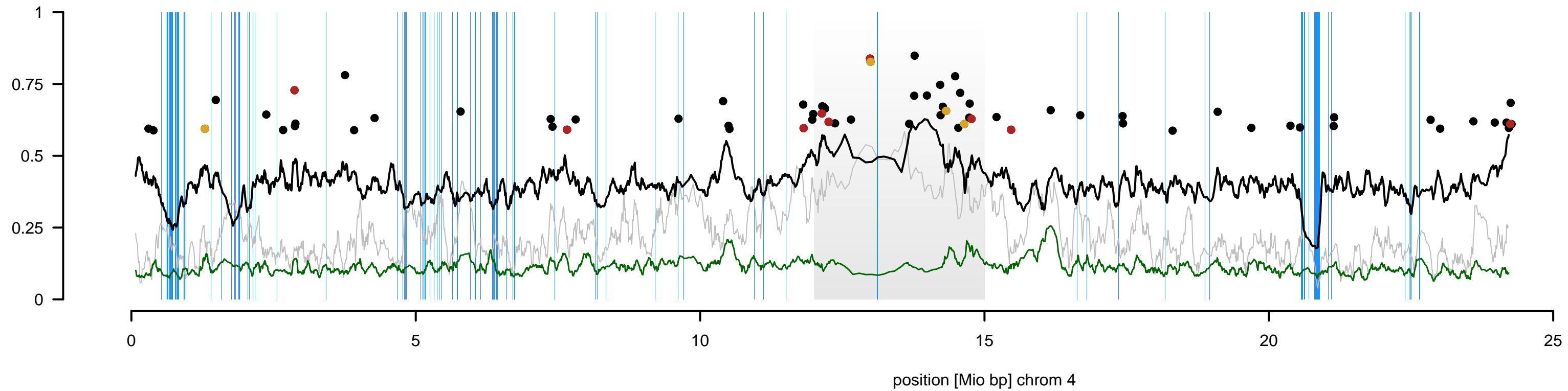
Hungary interspecific comparisons – 8k windows



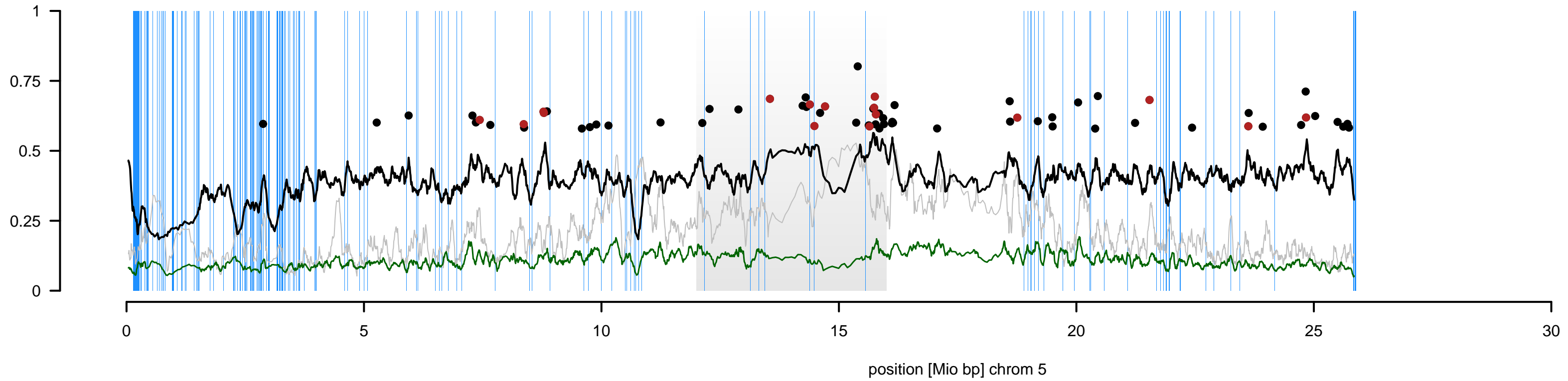
Italy interspecific comparisons – 8k windows



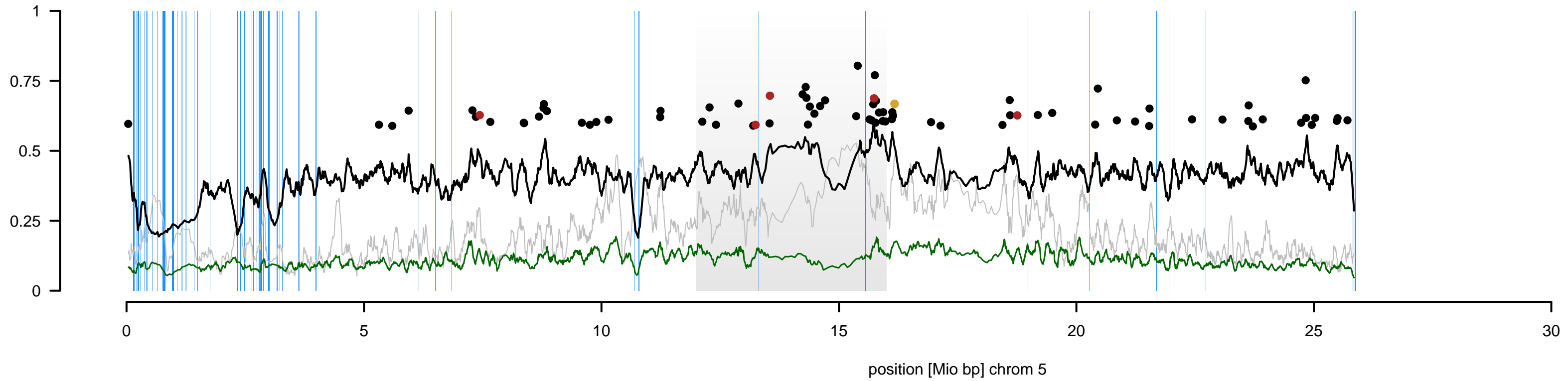
Hungary interspecific comparisons – 8k windows



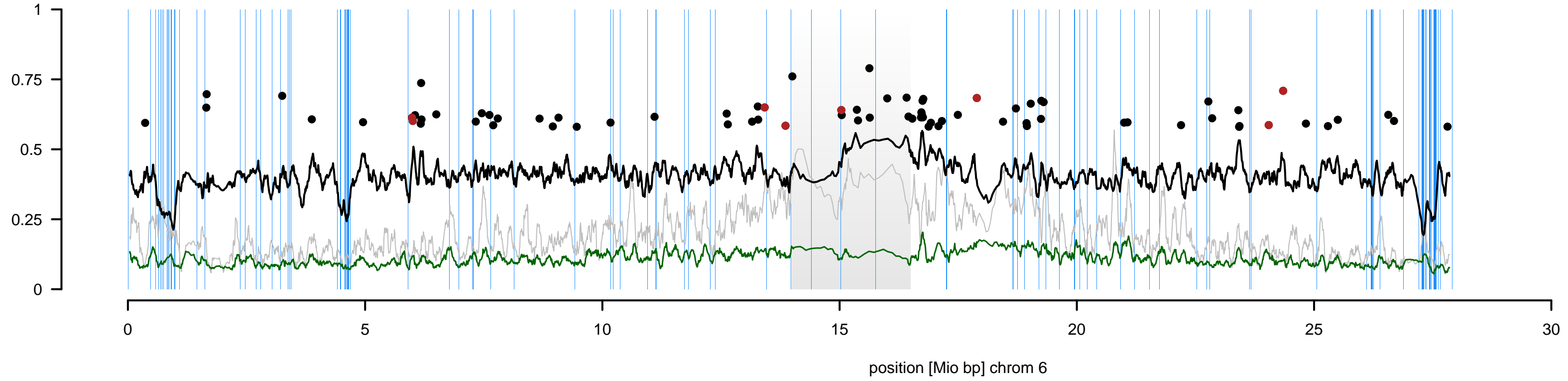
Italy interspecific comparisons – 8k windows



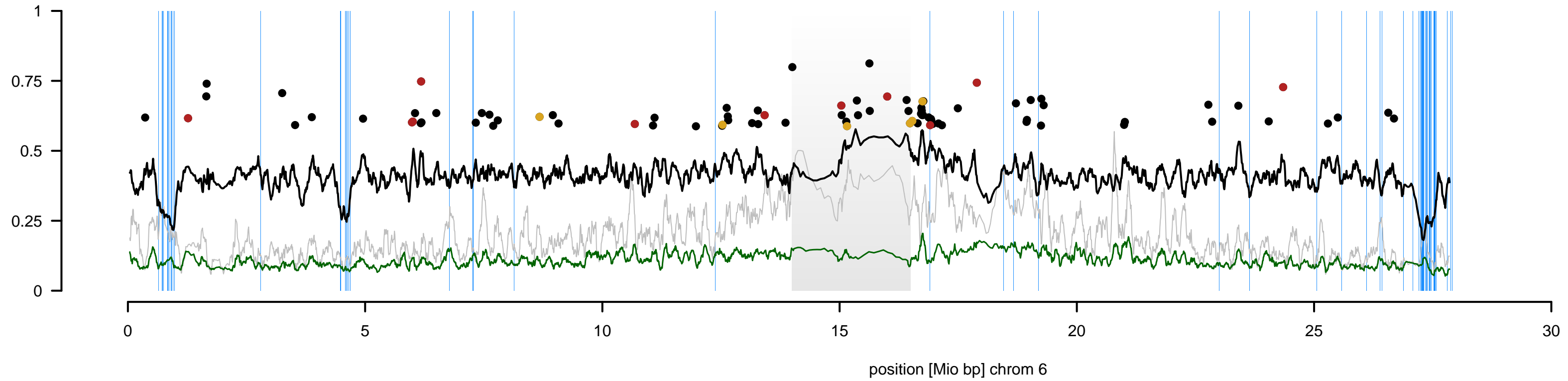
Hungary interspecific comparisons – 8k windows



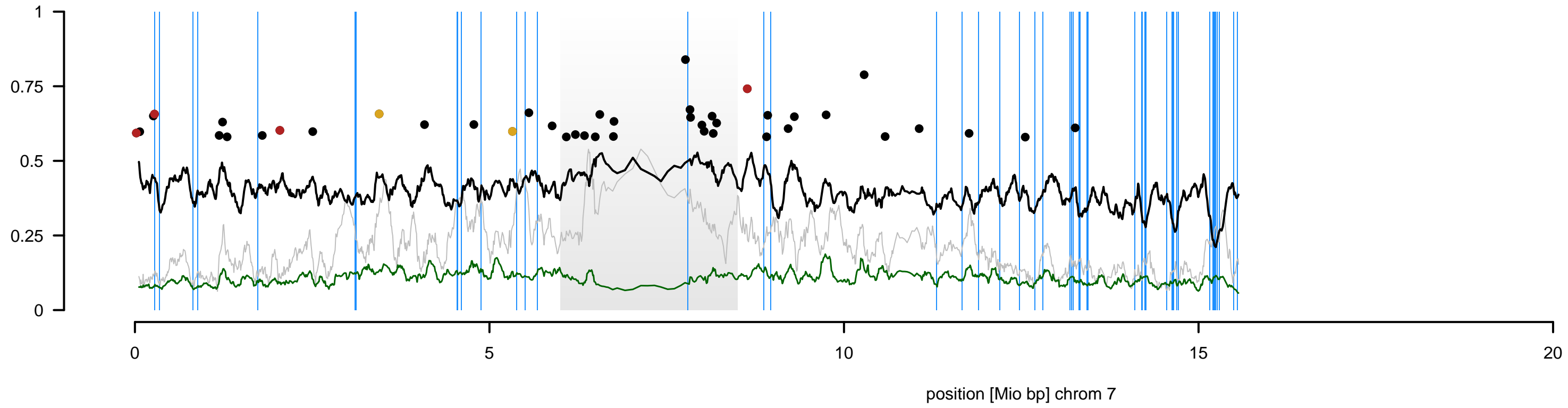
Italy interspecific comparisons – 8k windows



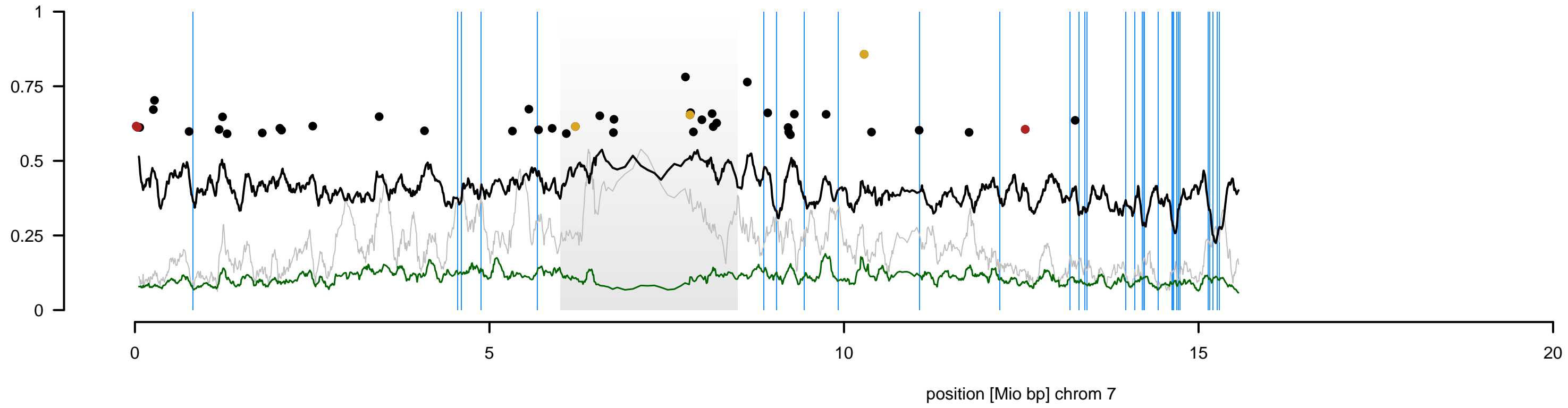
Hungary interspecific comparisons – 8k windows



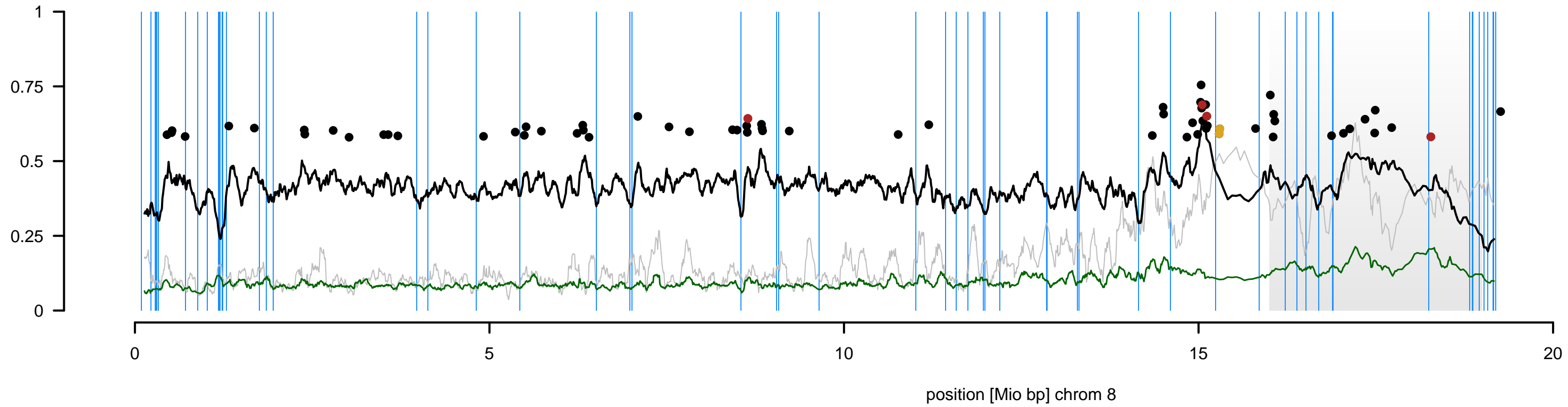
Italy interspecific comparisons – 8k windows



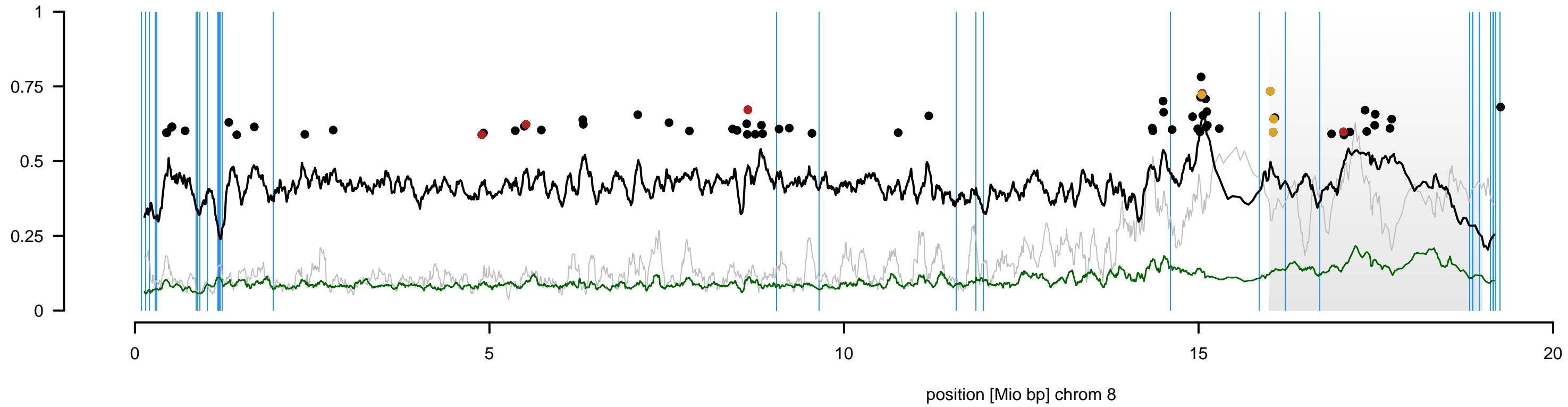
Hungary interspecific comparisons – 8k windows



Italy interspecific comparisons – 8k windows

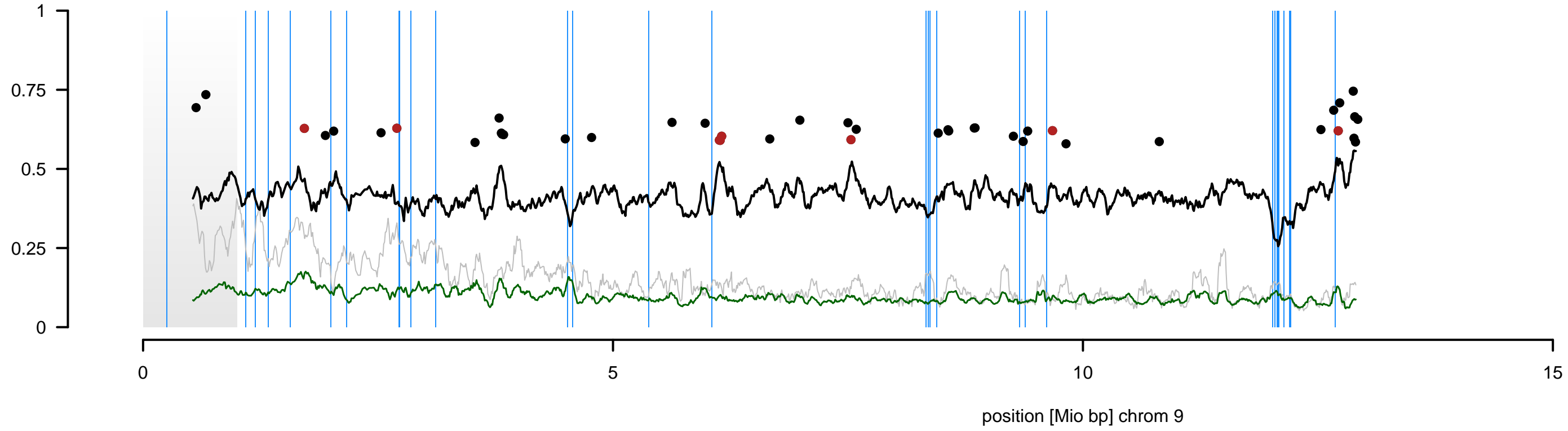


Hungary interspecific comparisons – 8k windows

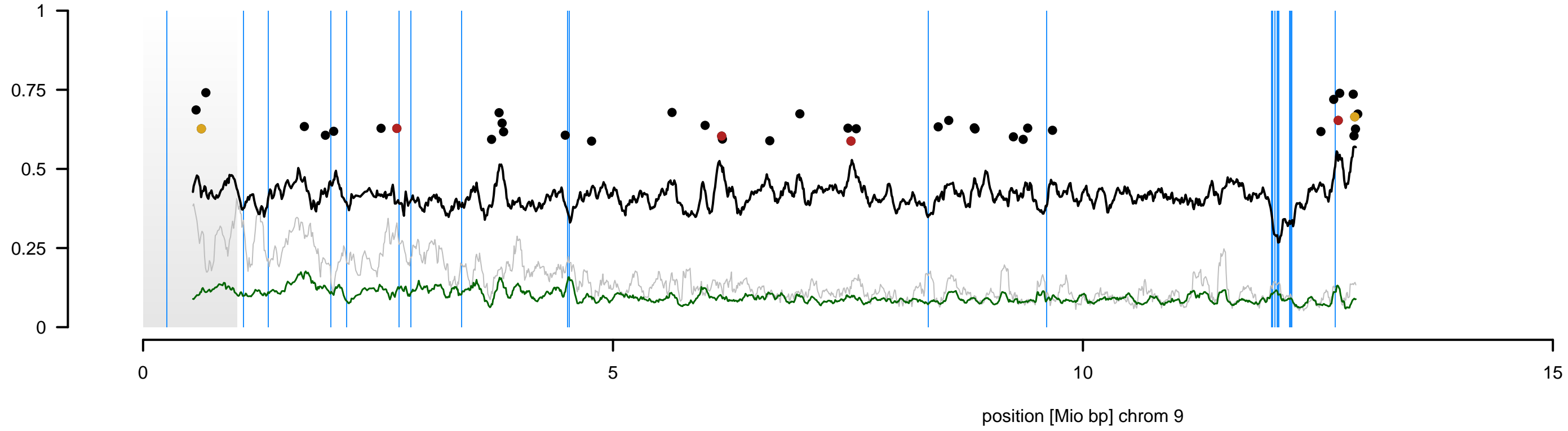




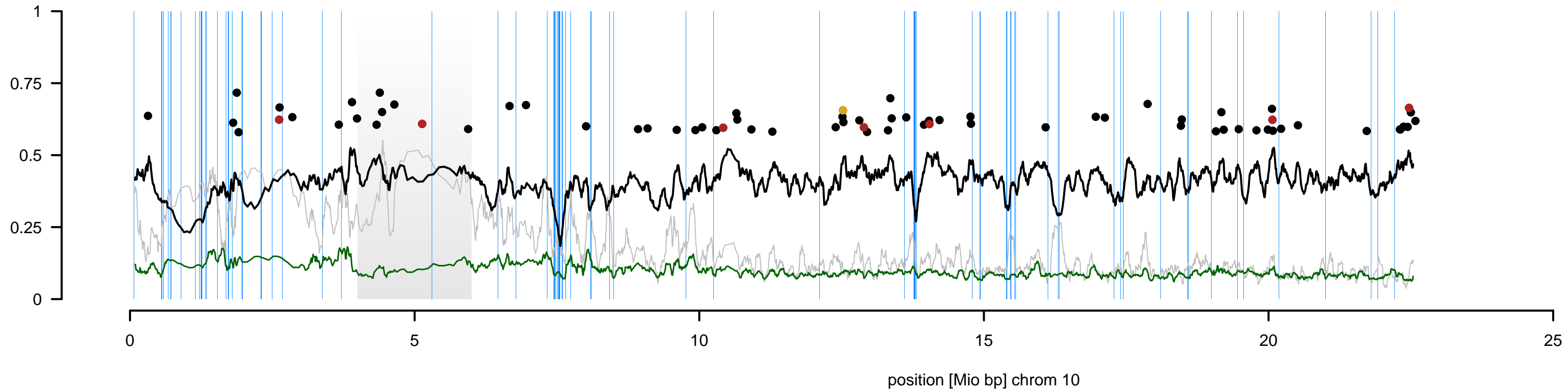
Italy interspecific comparisons – 8k windows



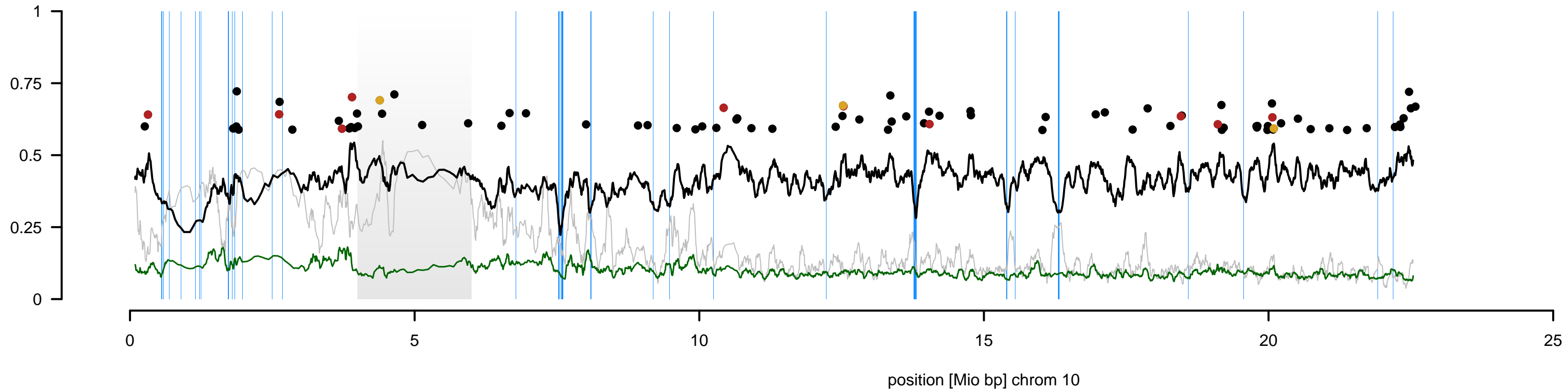
Hungary interspecific comparisons – 8k windows



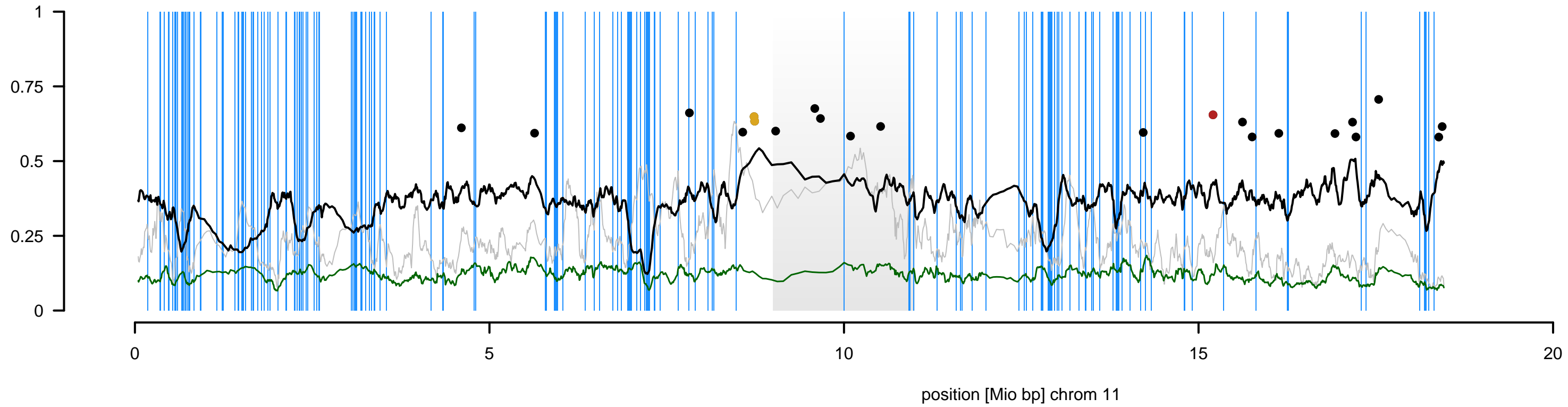
Italy interspecific comparisons – 8k windows



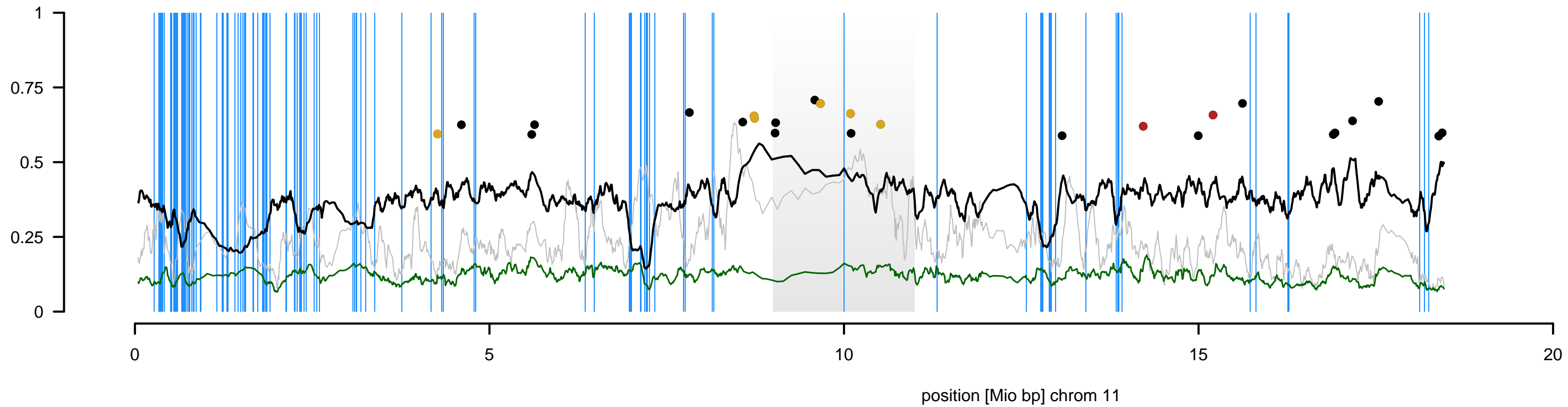
Hungary interspecific comparisons – 8k windows



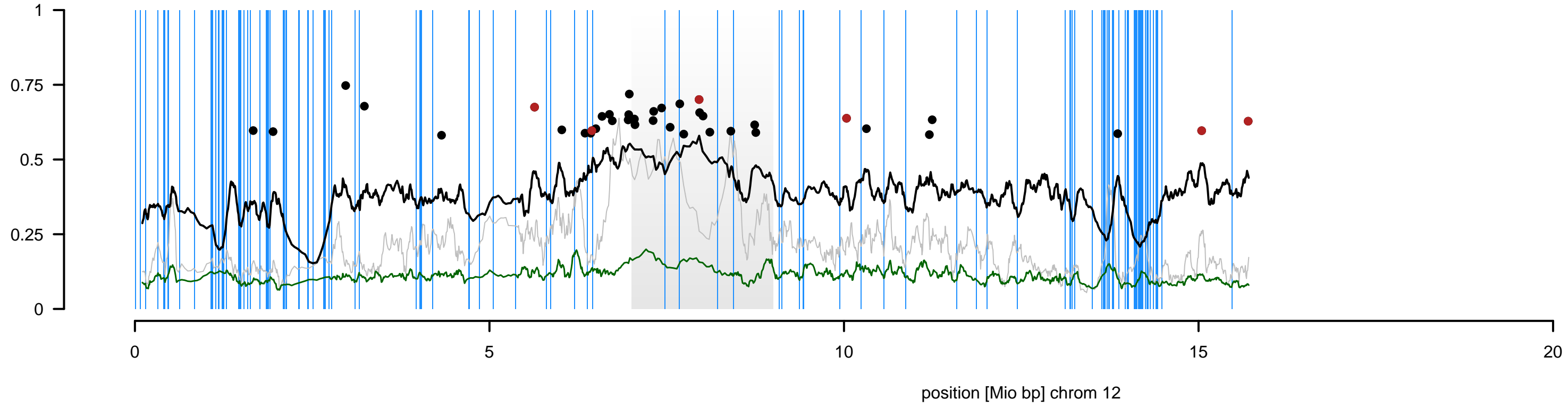
Italy interspecific comparisons – 8k windows



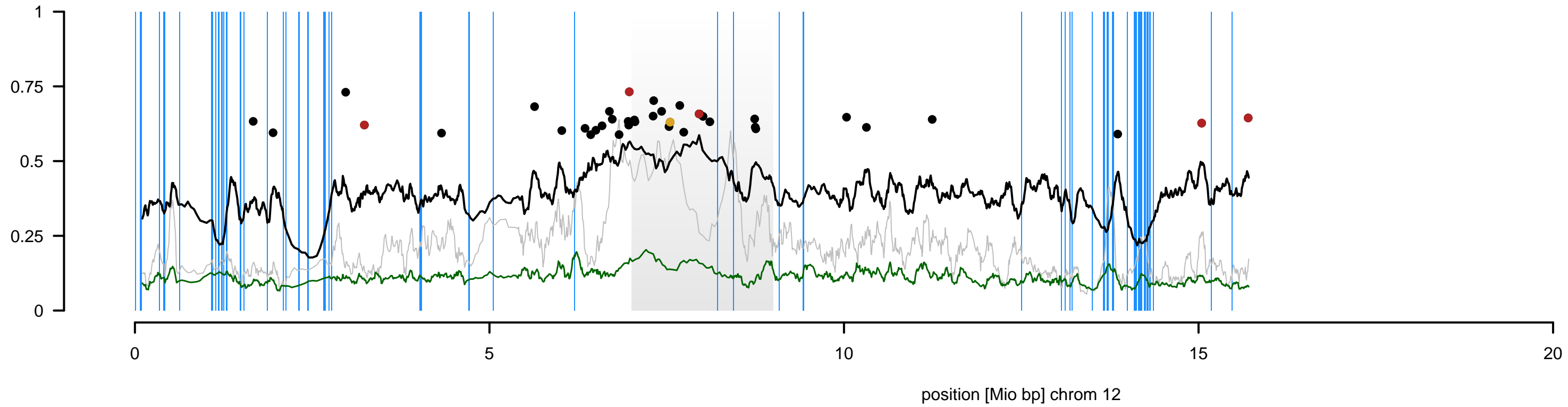
Hungary interspecific comparisons – 8k windows



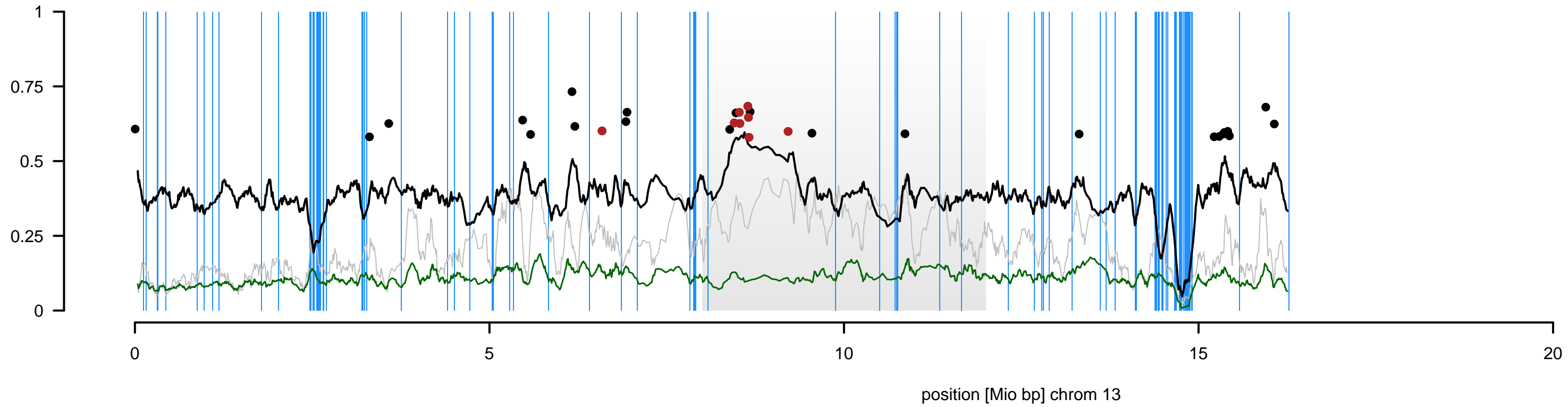
Italy interspecific comparisons – 8k windows



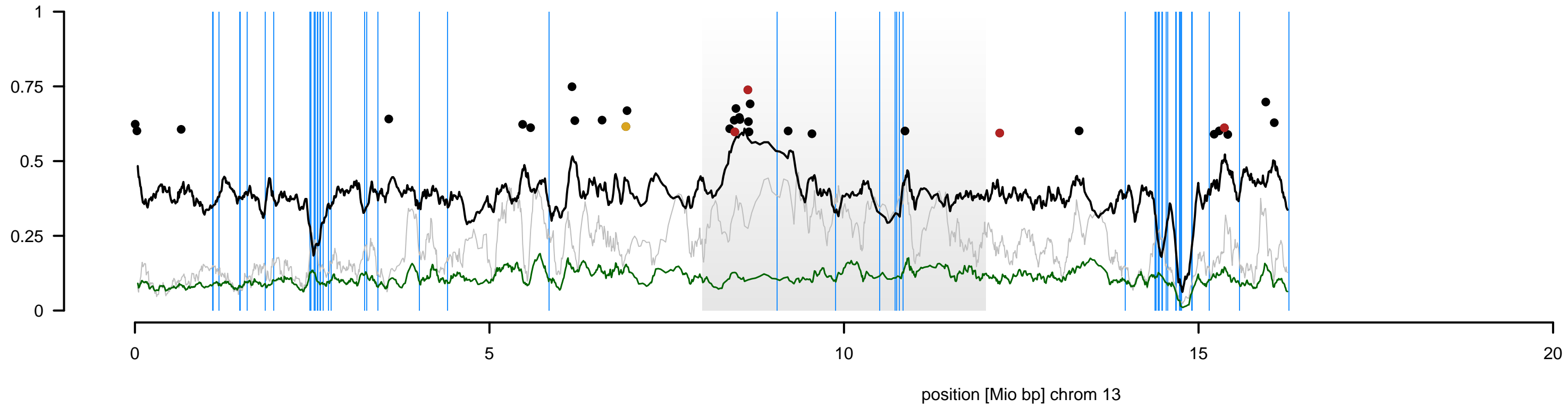
Hungary interspecific comparisons – 8k windows



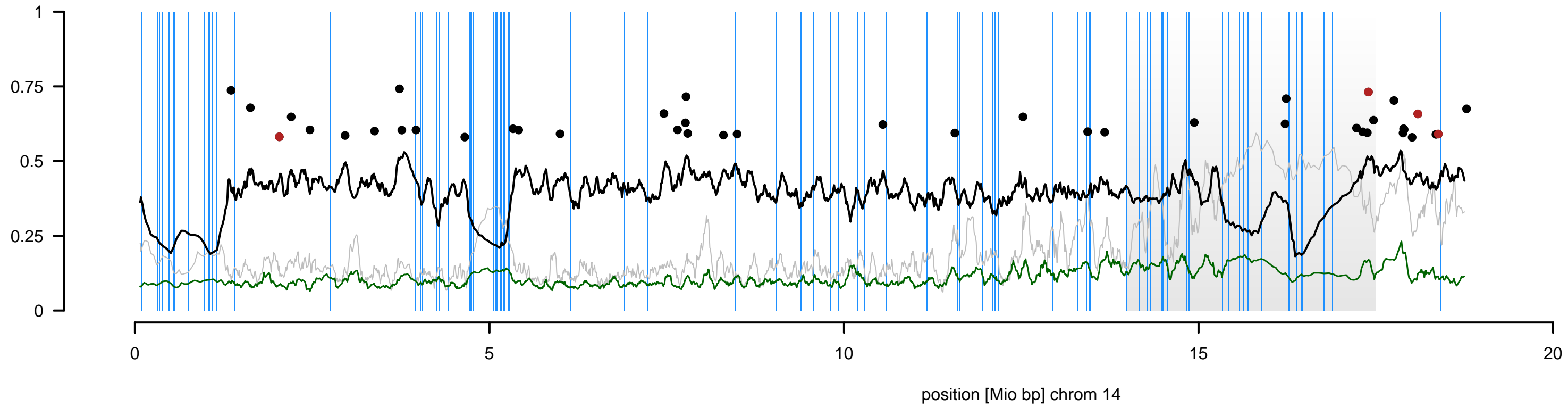
Italy interspecific comparisons – 8k windows



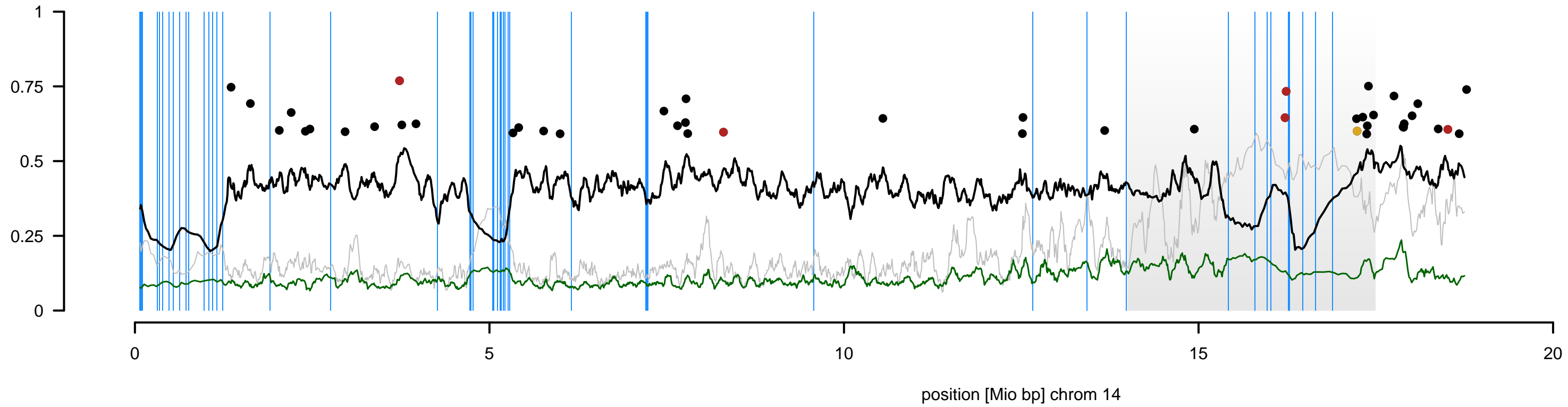
Hungary interspecific comparisons – 8k windows



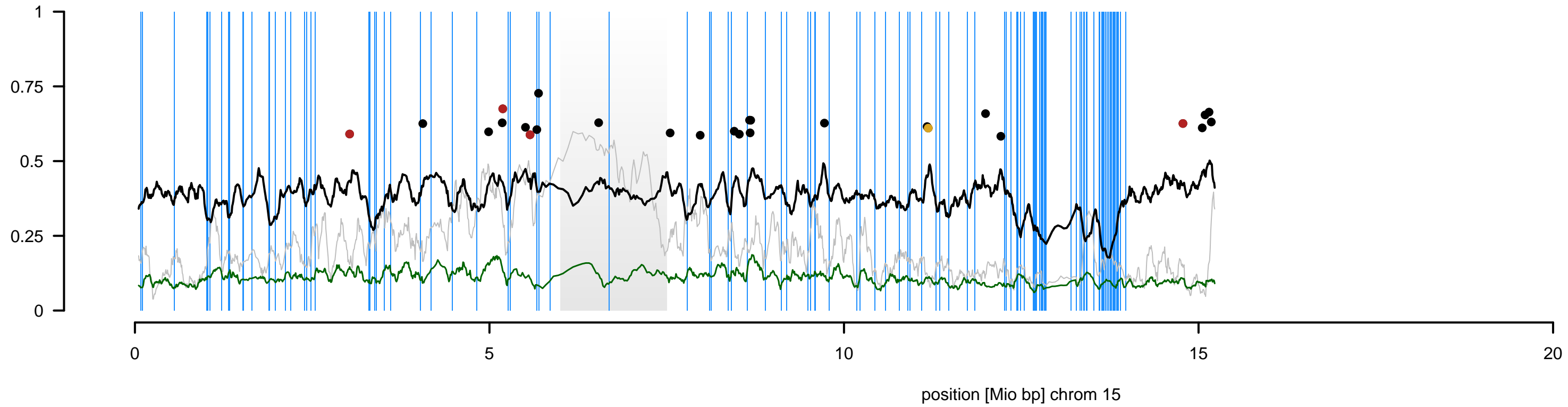
Italy interspecific comparisons – 8k windows



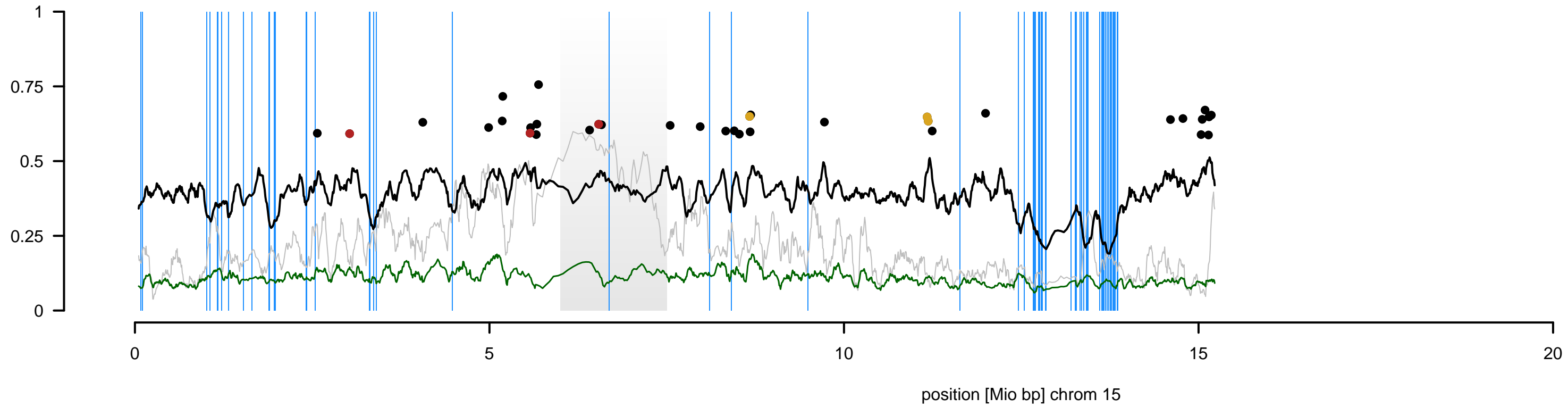
Hungary interspecific comparisons – 8k windows



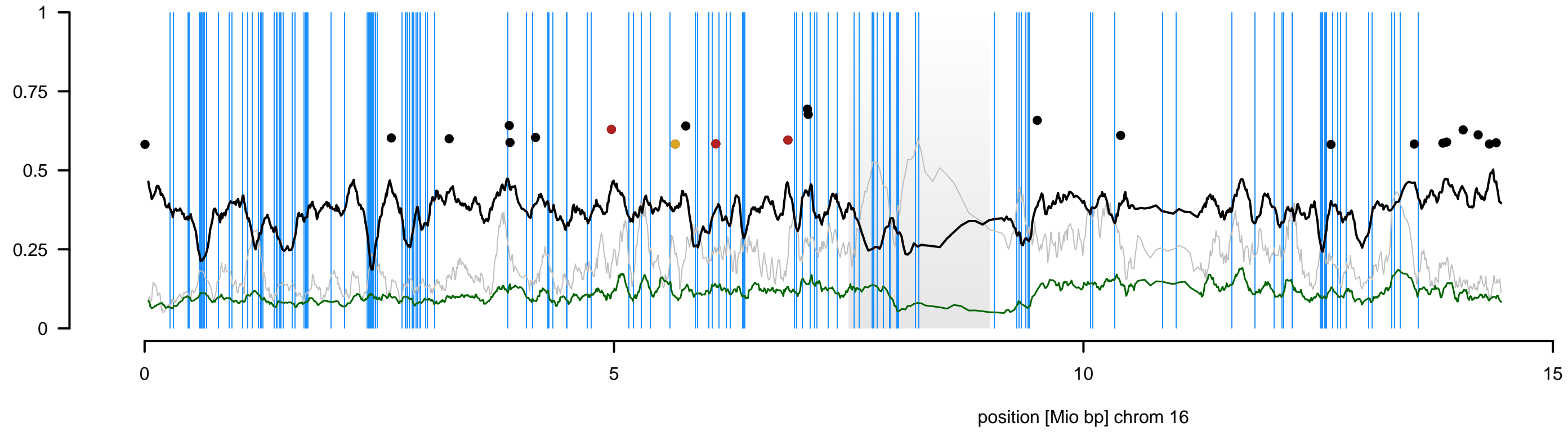
Italy interspecific comparisons – 8k windows



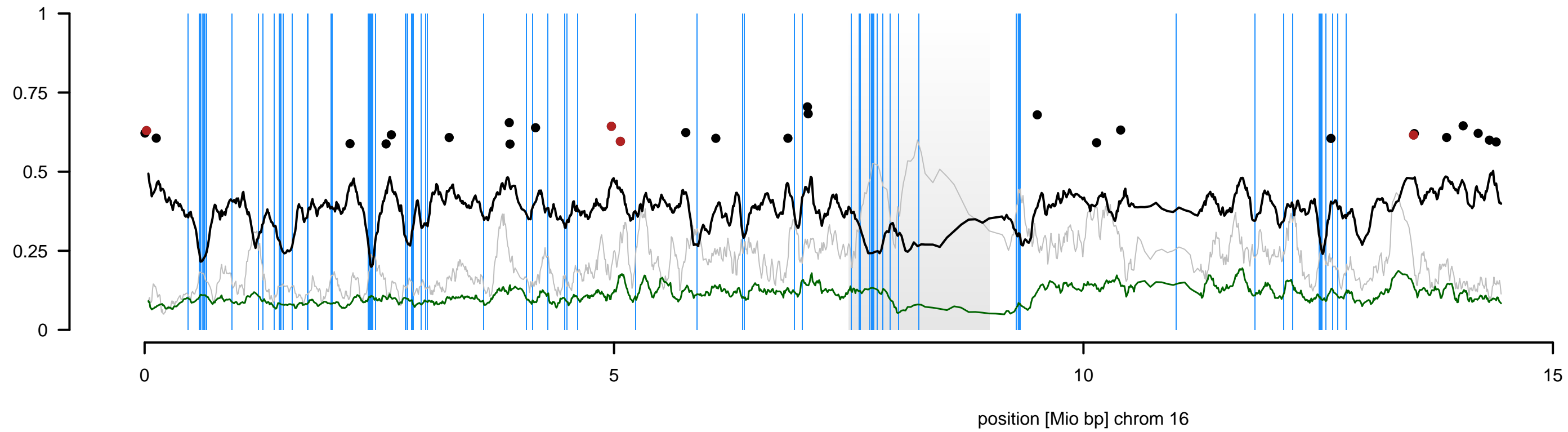
Hungary interspecific comparisons – 8k windows



Italy interspecific comparisons – 8k windows

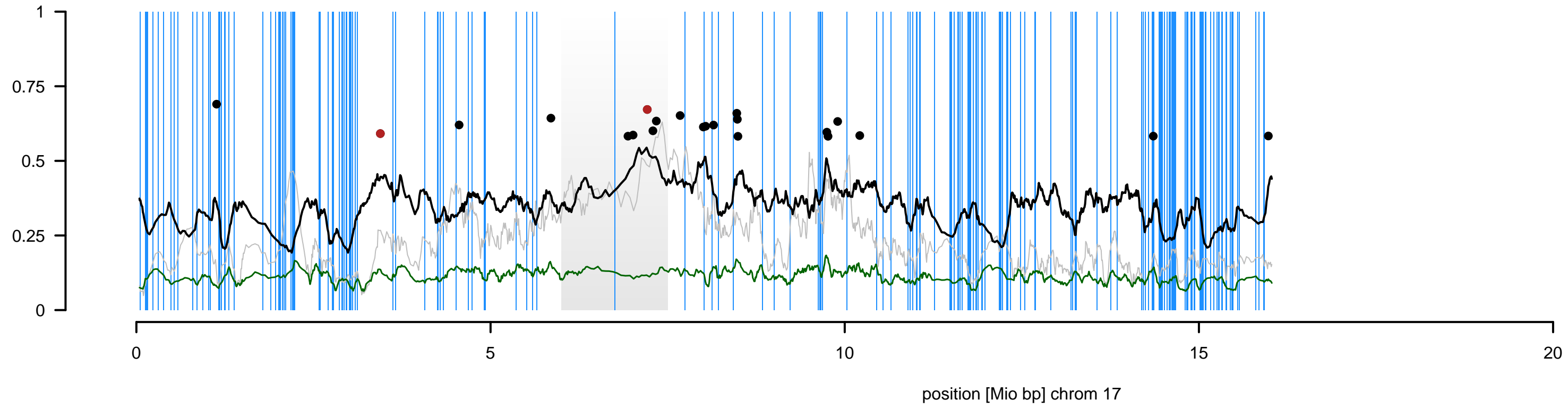


Hungary interspecific comparisons – 8k windows

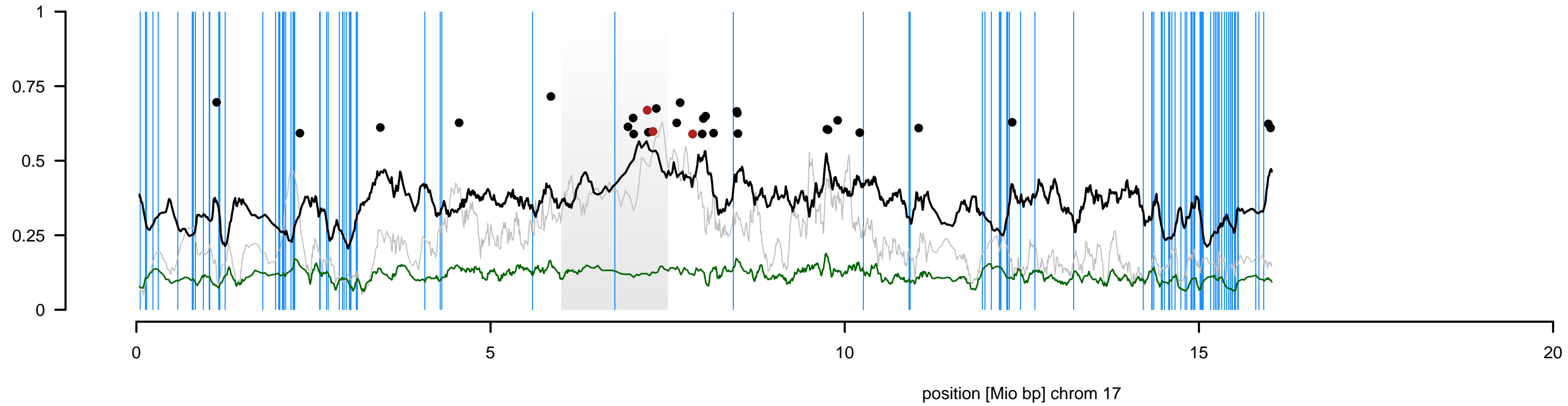




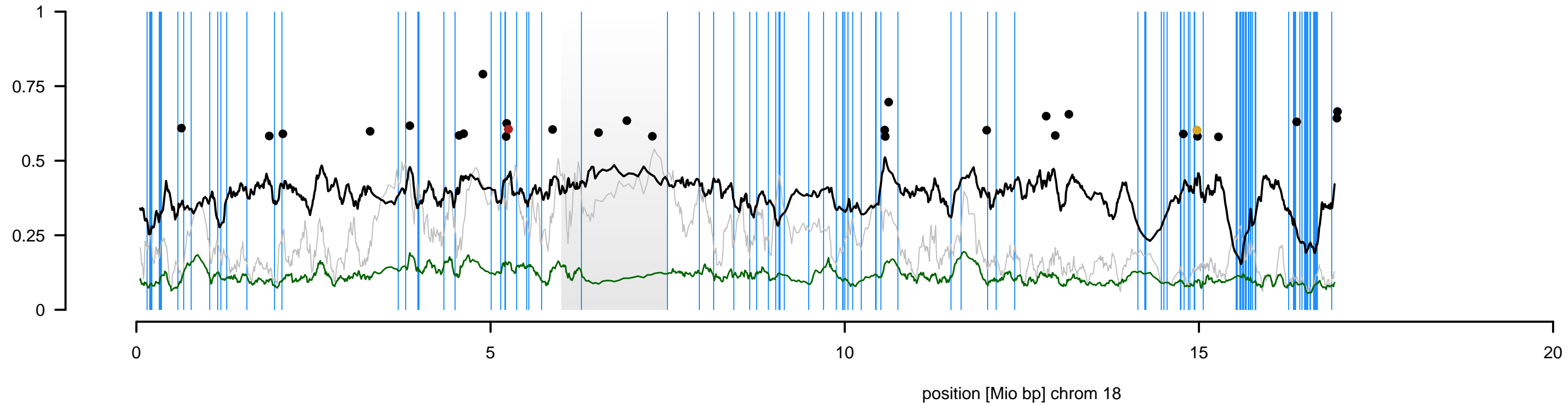
Italy interspecific comparisons – 8k windows



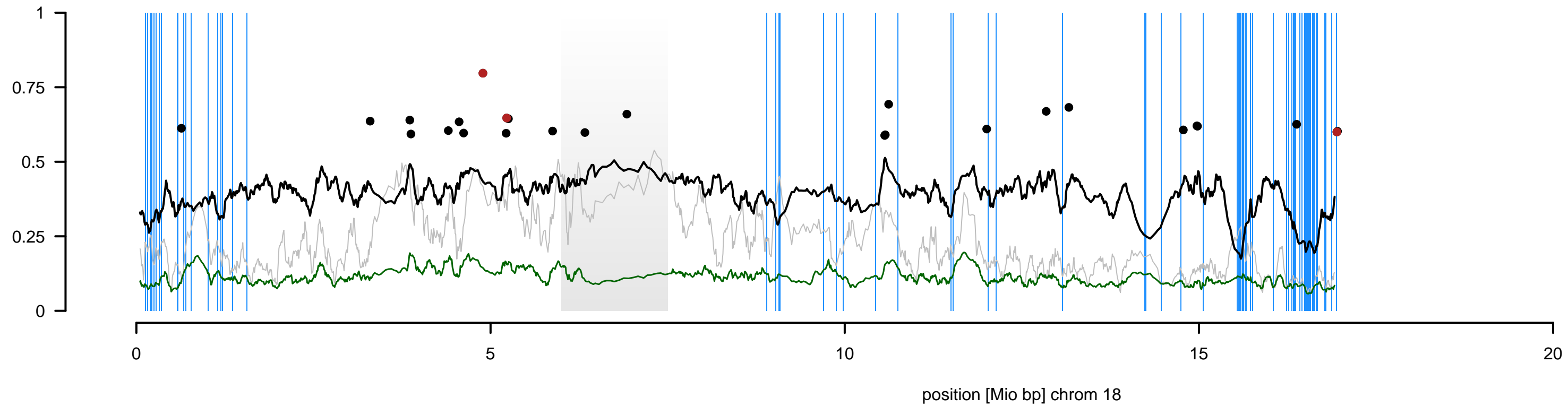
Hungary interspecific comparisons – 8k windows



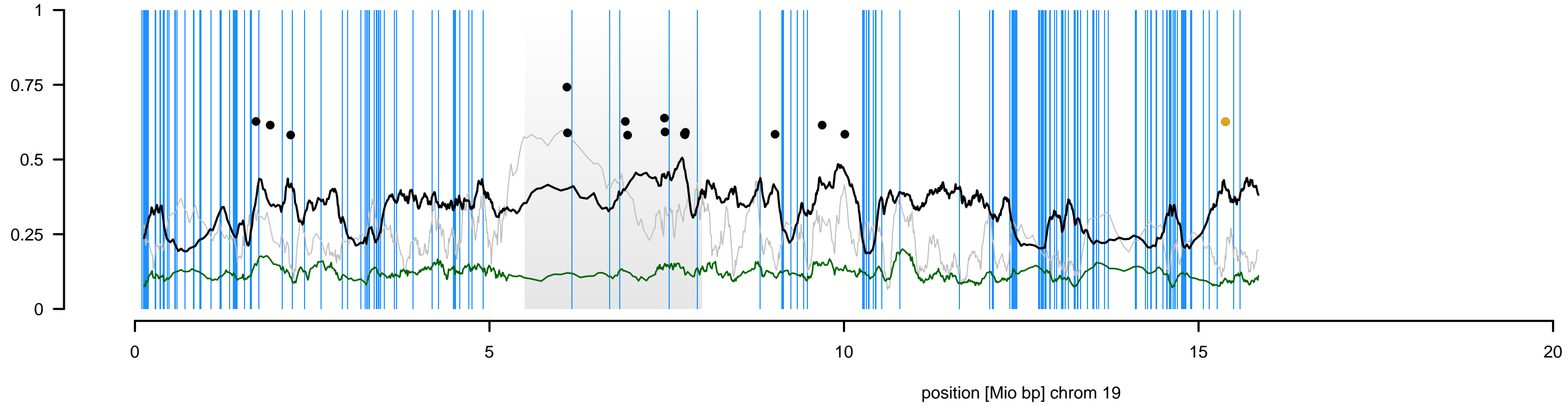
Italy interspecific comparisons – 8k windows



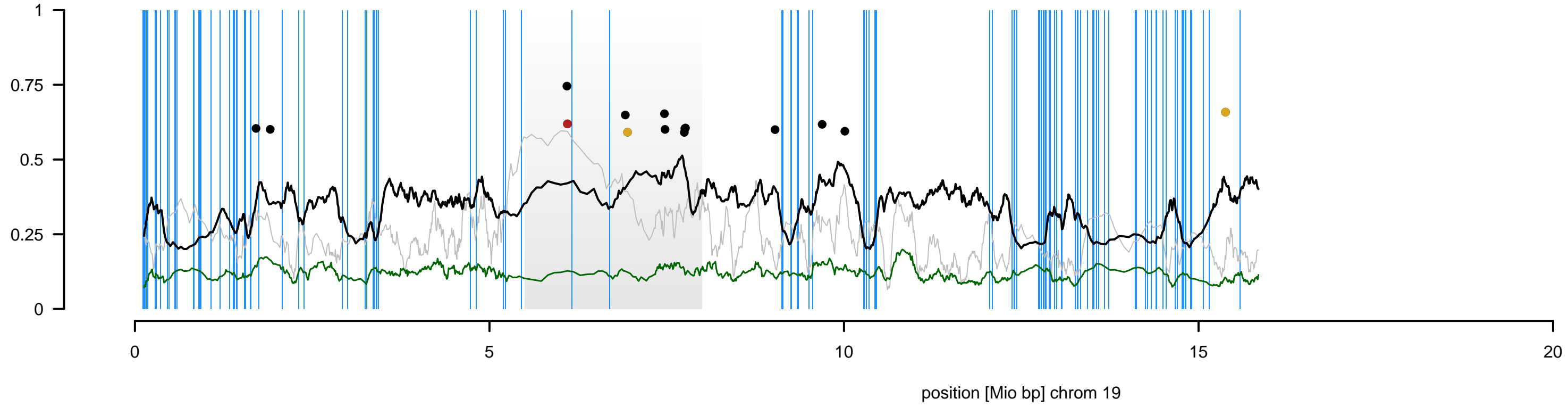
Hungary interspecific comparisons – 8k windows



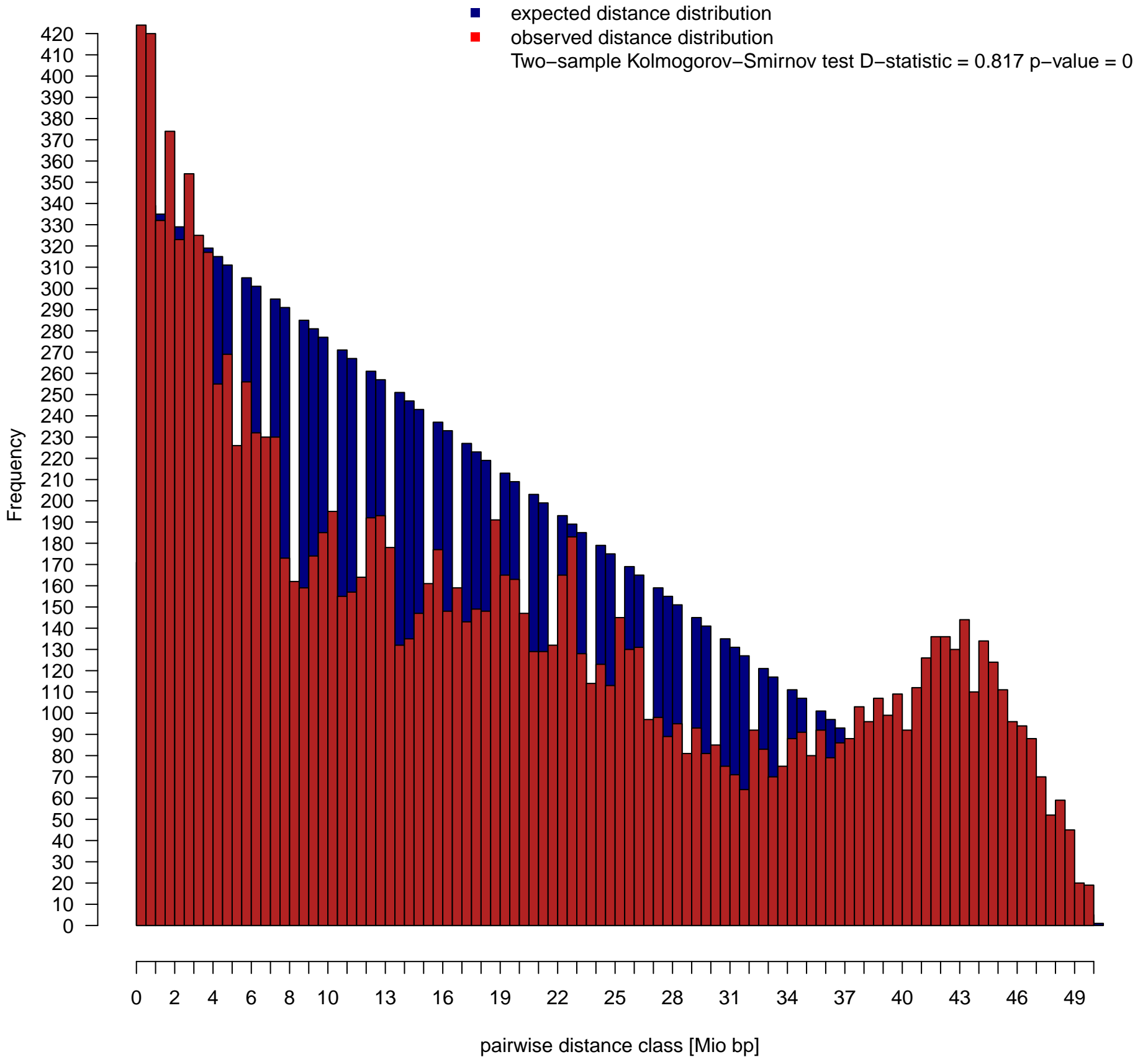
Italy interspecific comparisons – 8k windows



Hungary interspecific comparisons – 8k windows

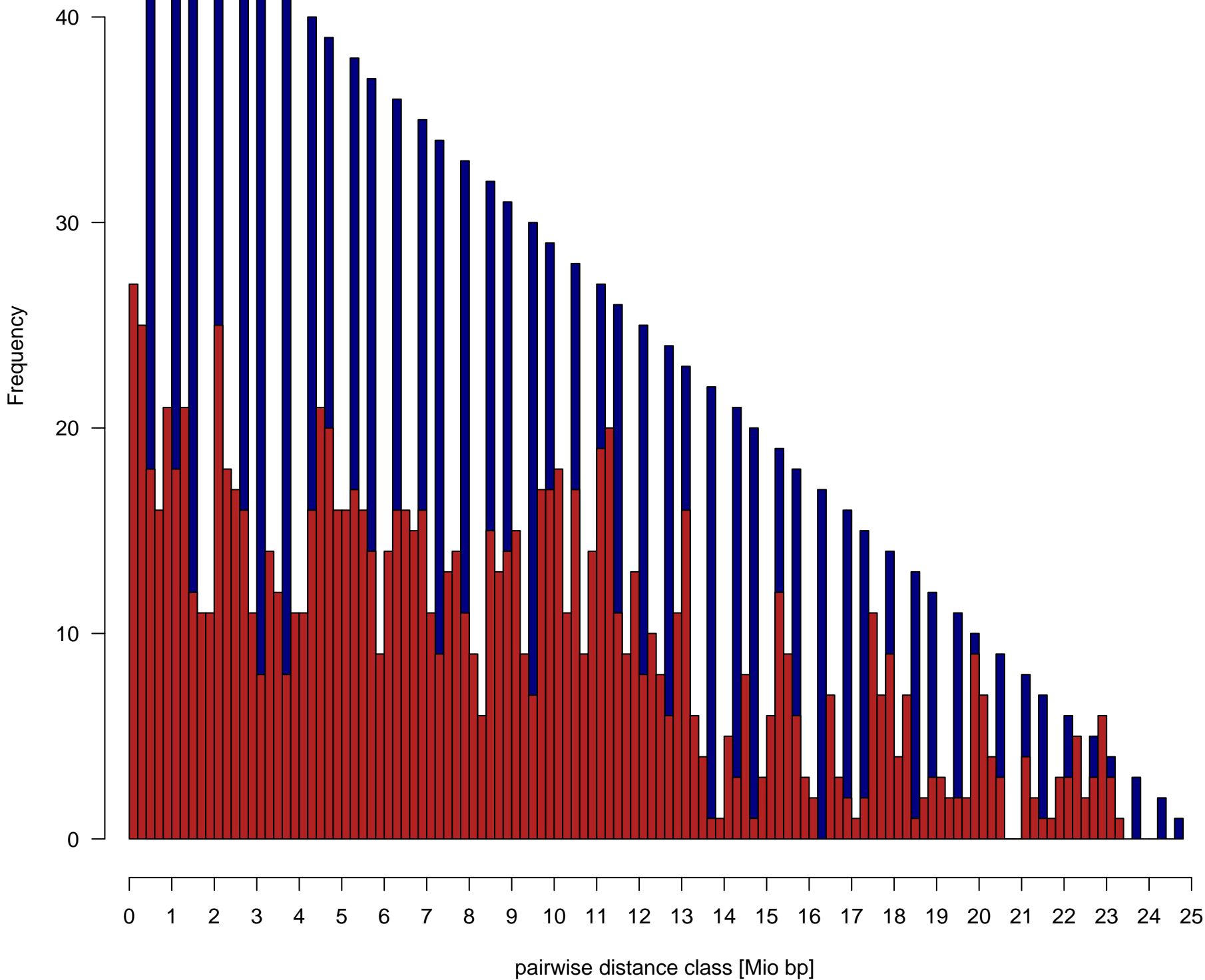


# Observed and expected distances between no-fixation regions on chromosome 1 in Italy

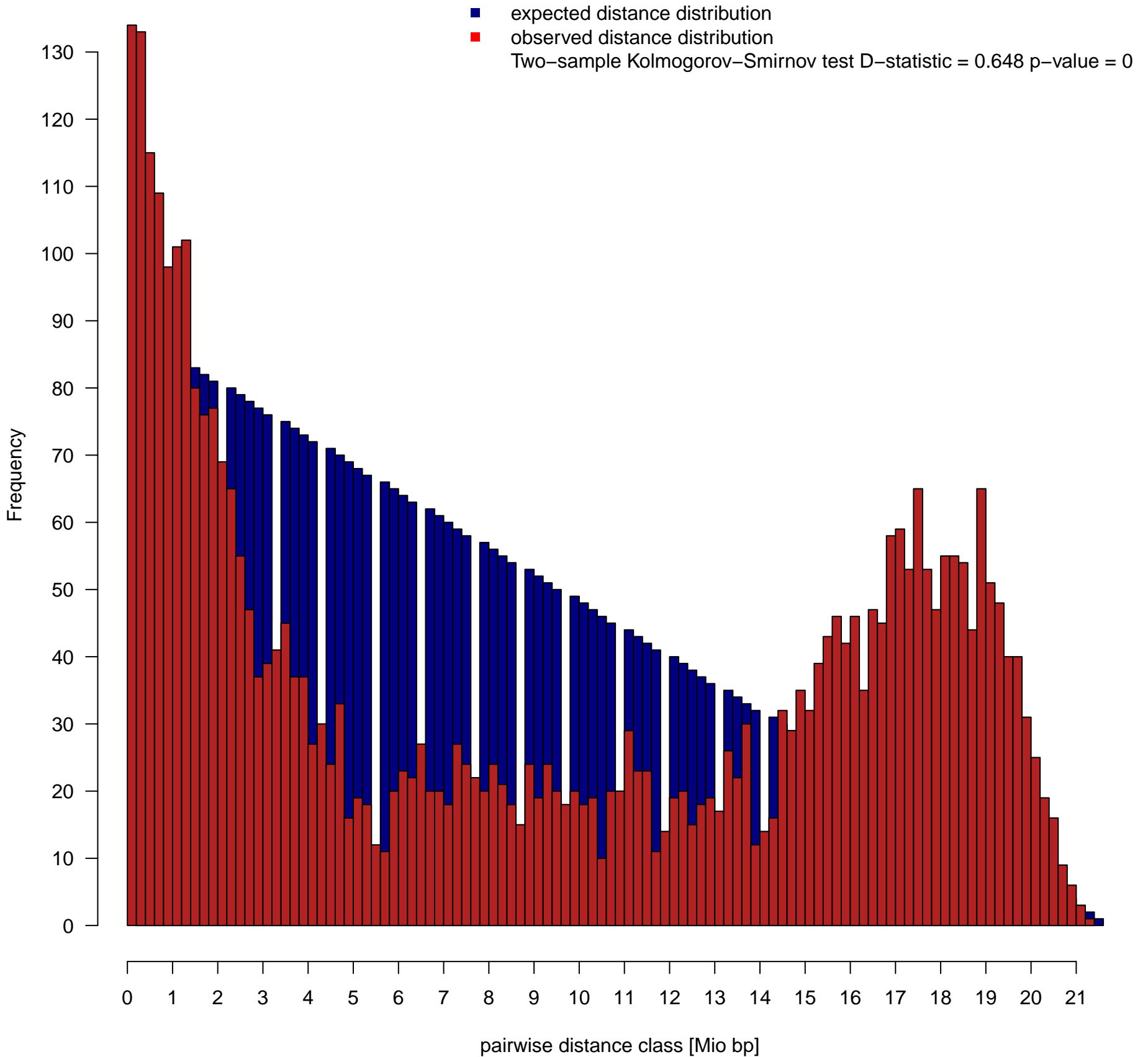


# Observed and expected distances between no-fixation regions on chromosome 2 in Italy

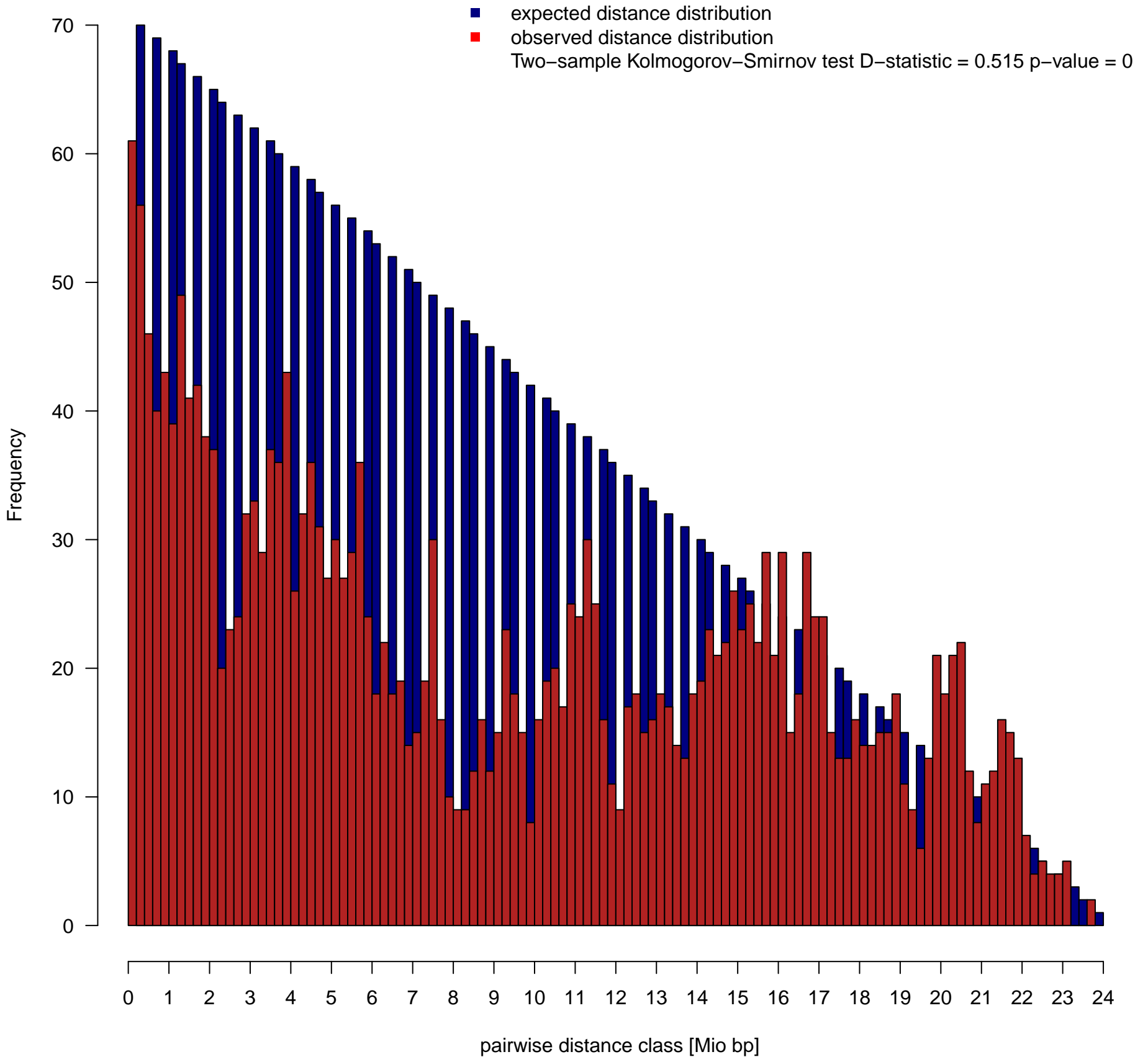
■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.271 p-value = 0



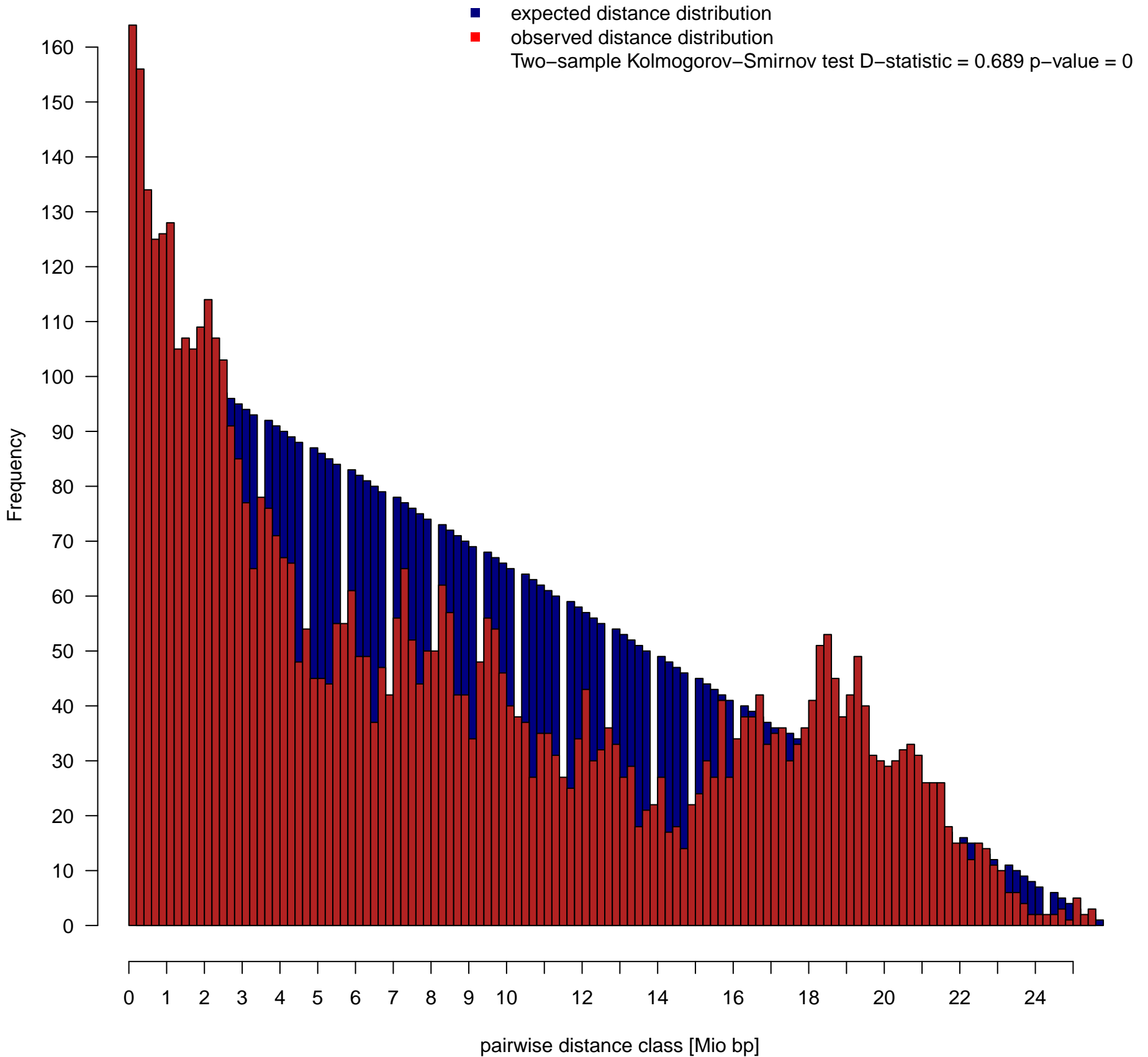
# Observed and expected distances between no-fixation regions on chromosome 3 in Italy



# Observed and expected distances between no-fixation regions on chromosome 4 in Italy

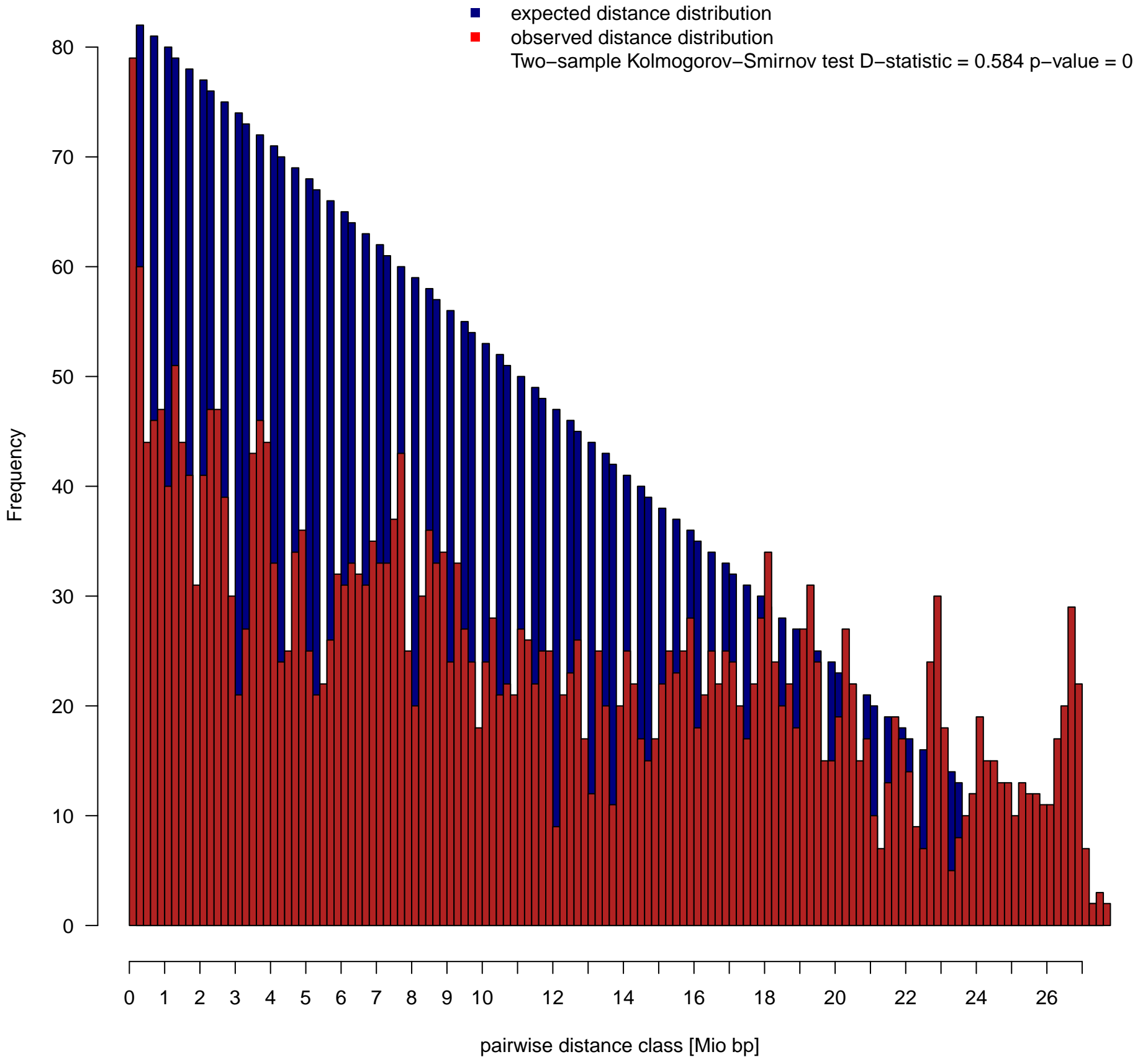


# Observed and expected distances between no-fixation regions on chromosome 5 in Italy



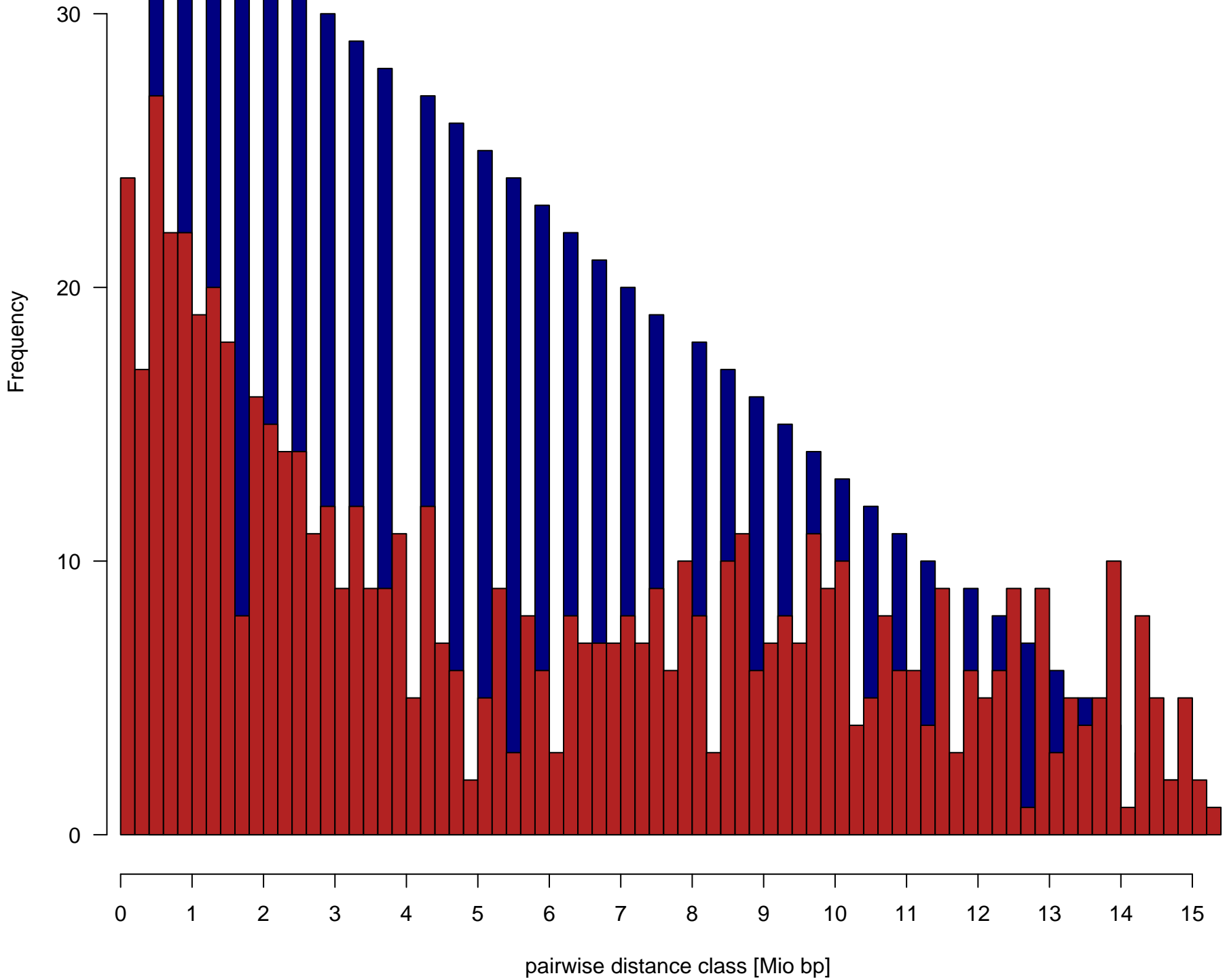


# Observed and expected distances between no-fixation regions on chromosome 6 in Italy



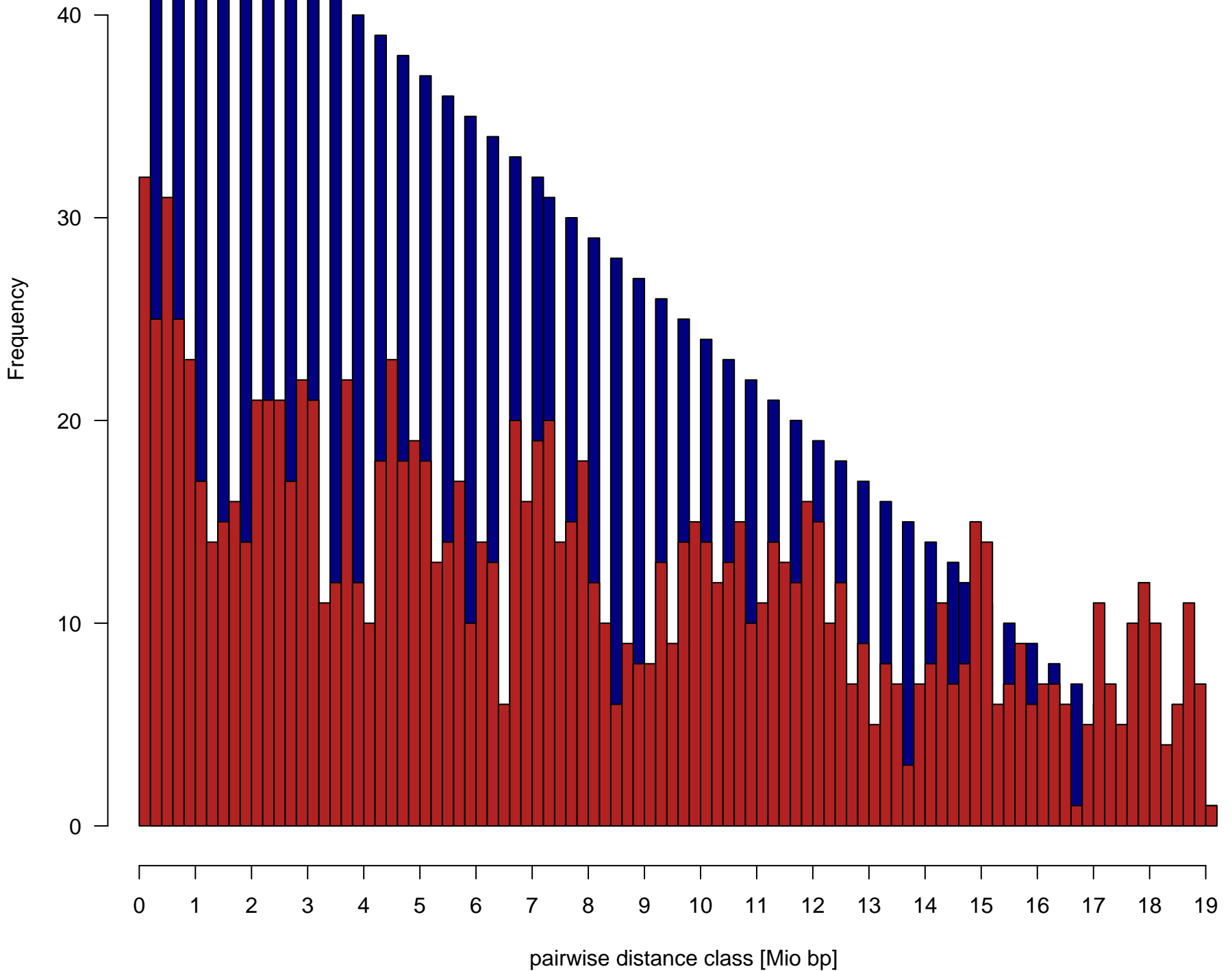
# Observed and expected distances between no-fixation regions on chromosome 7 in Italy

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.264 p-value = 0



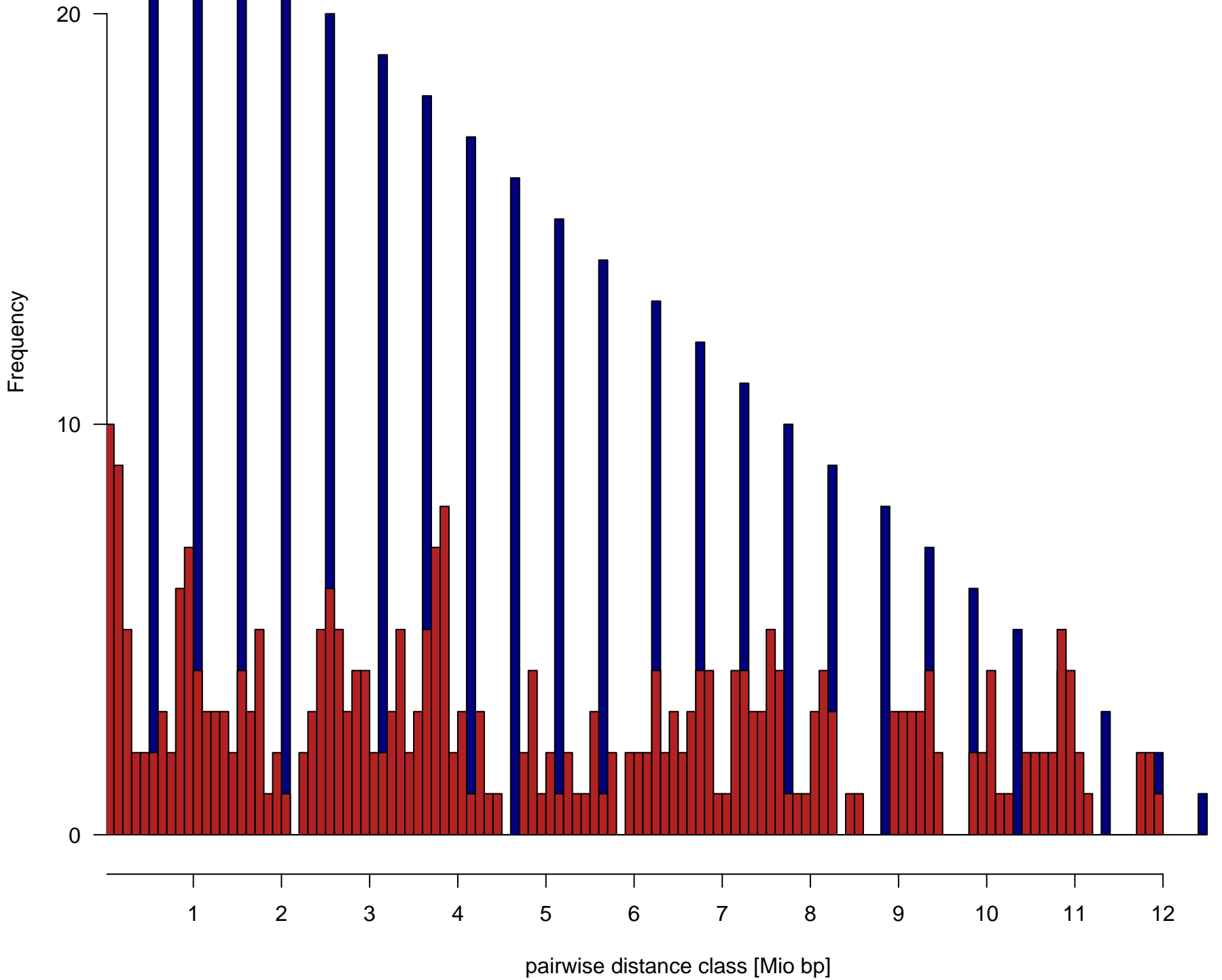
# Observed and expected distances between no-fixation regions on chromosome 8 in Italy

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.367 p-value = 0



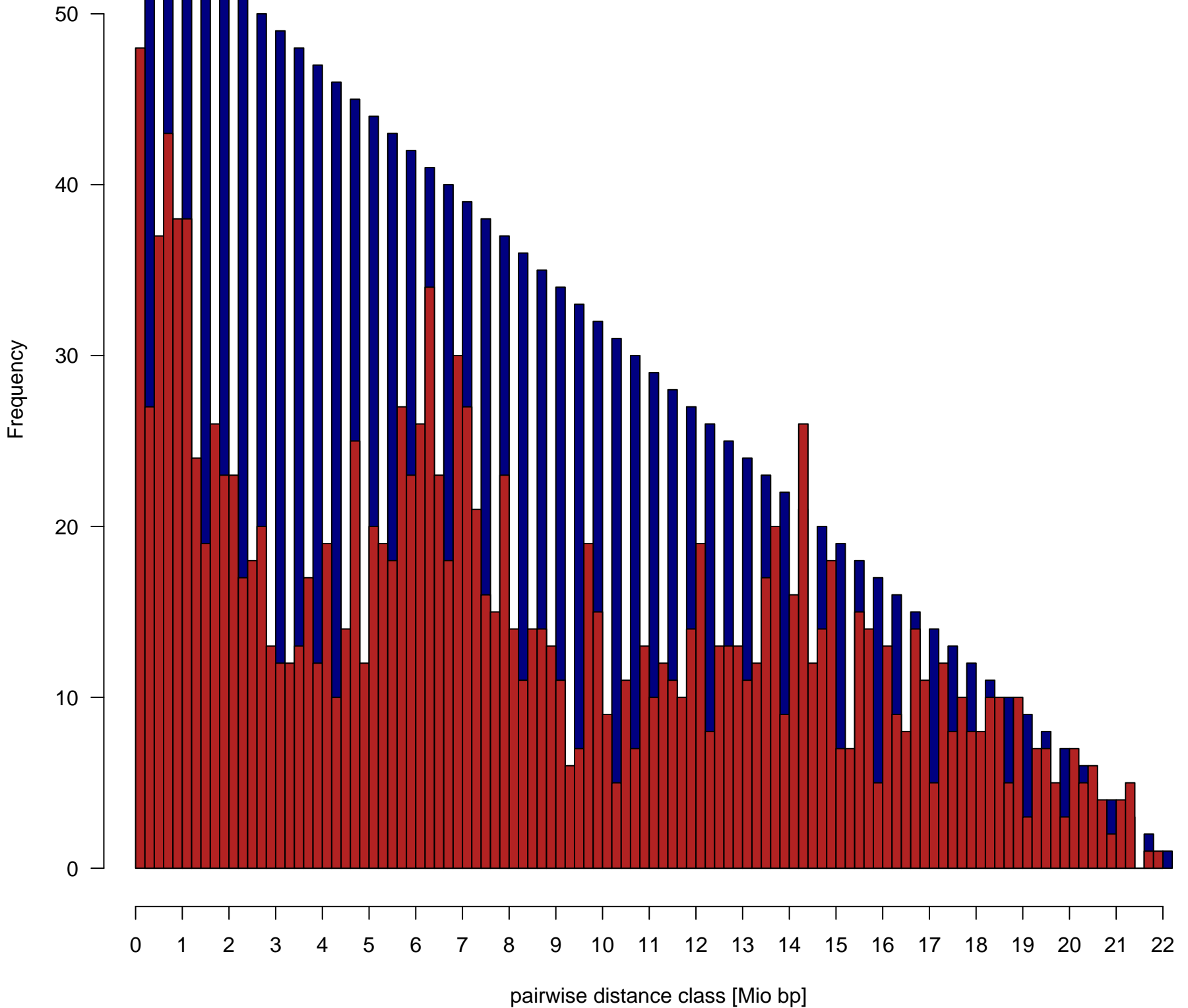
# Observed and expected distances between no-fixation regions on chromosome 9 in Italy

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.152 p-value = 0

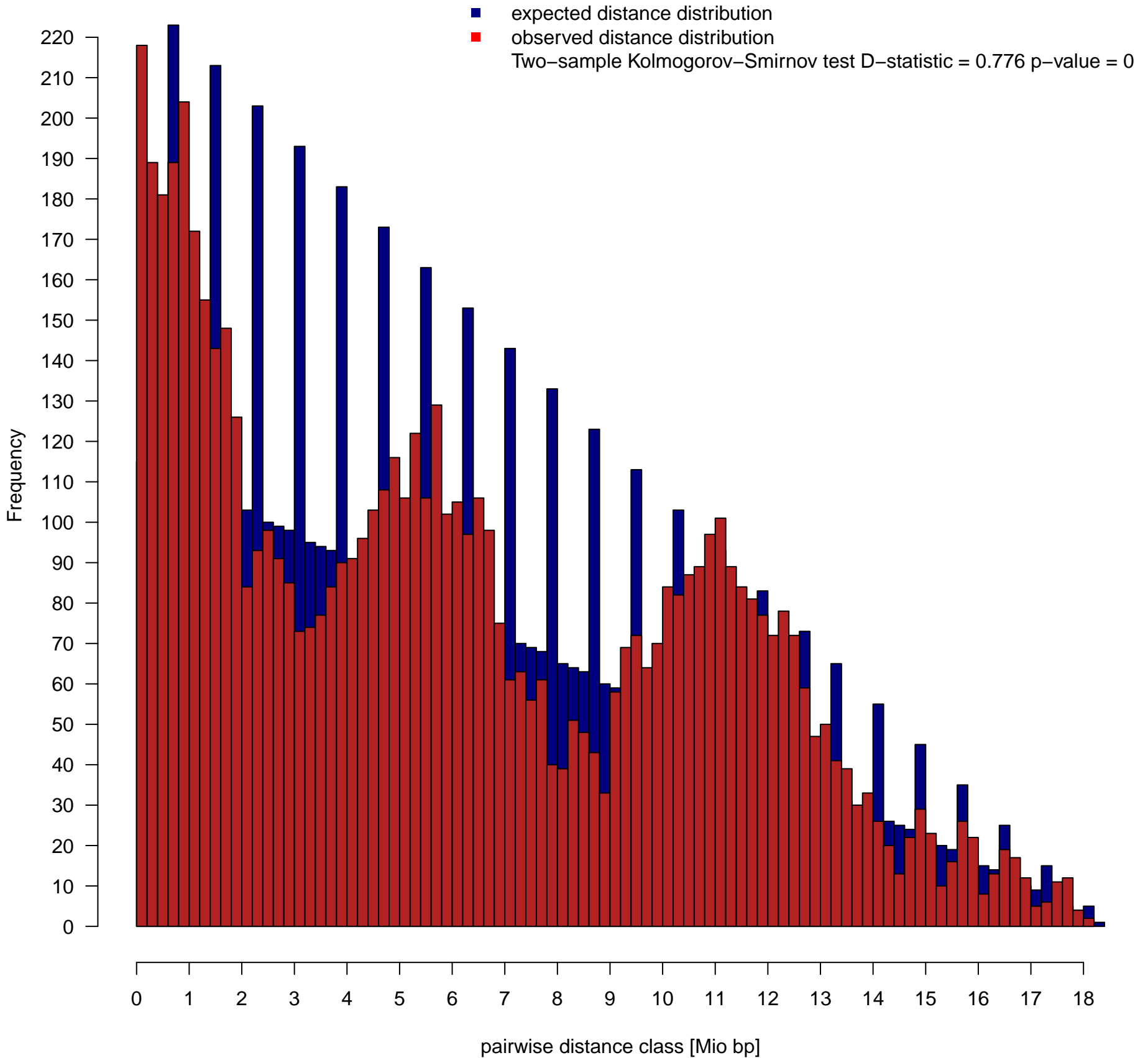


# Observed and expected distances between no-fixation regions on chromosome 10 in Italy

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.388 p-value = 0

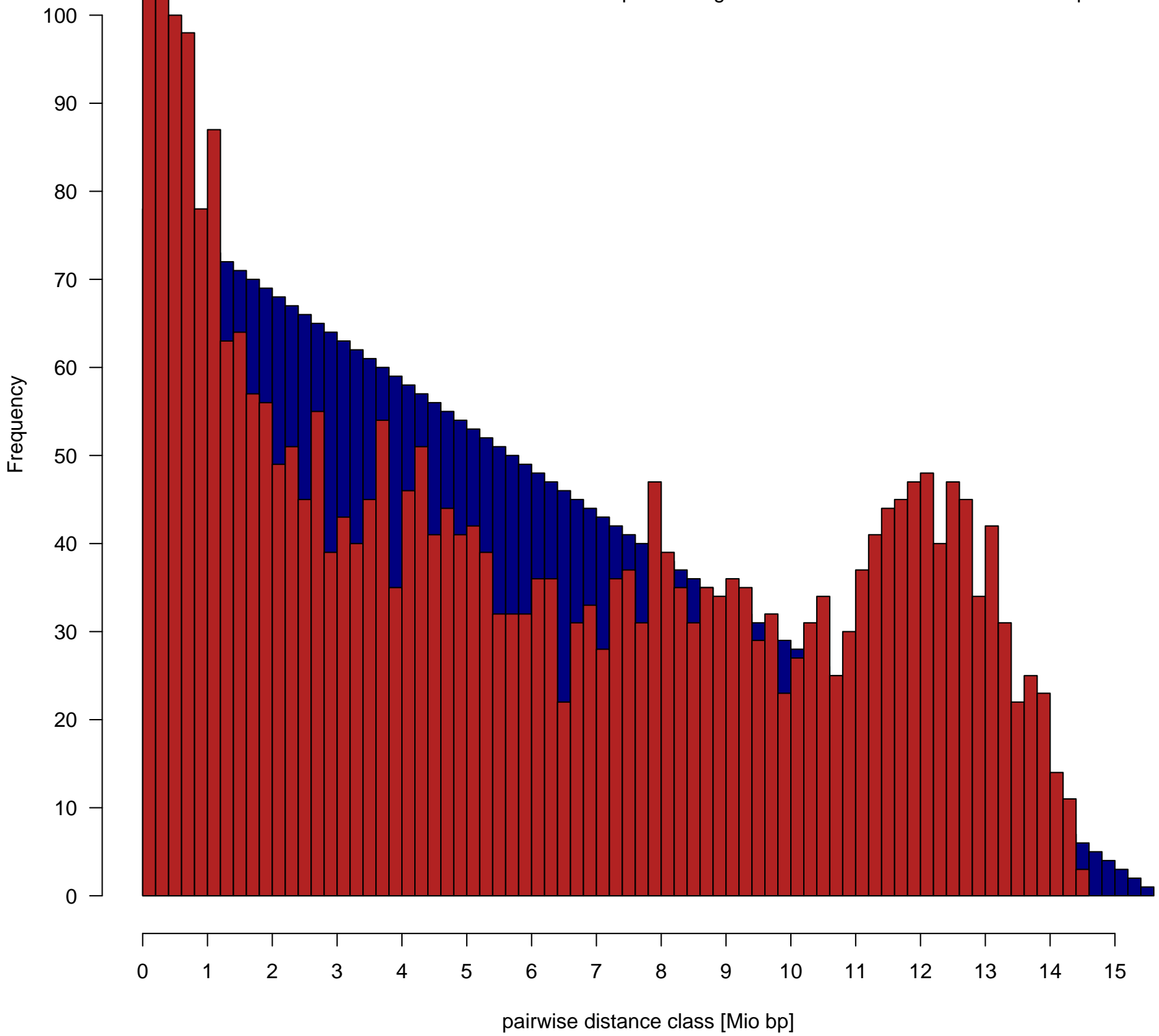


# Observed and expected distances between no-fixation regions on chromosome 11 in Italy



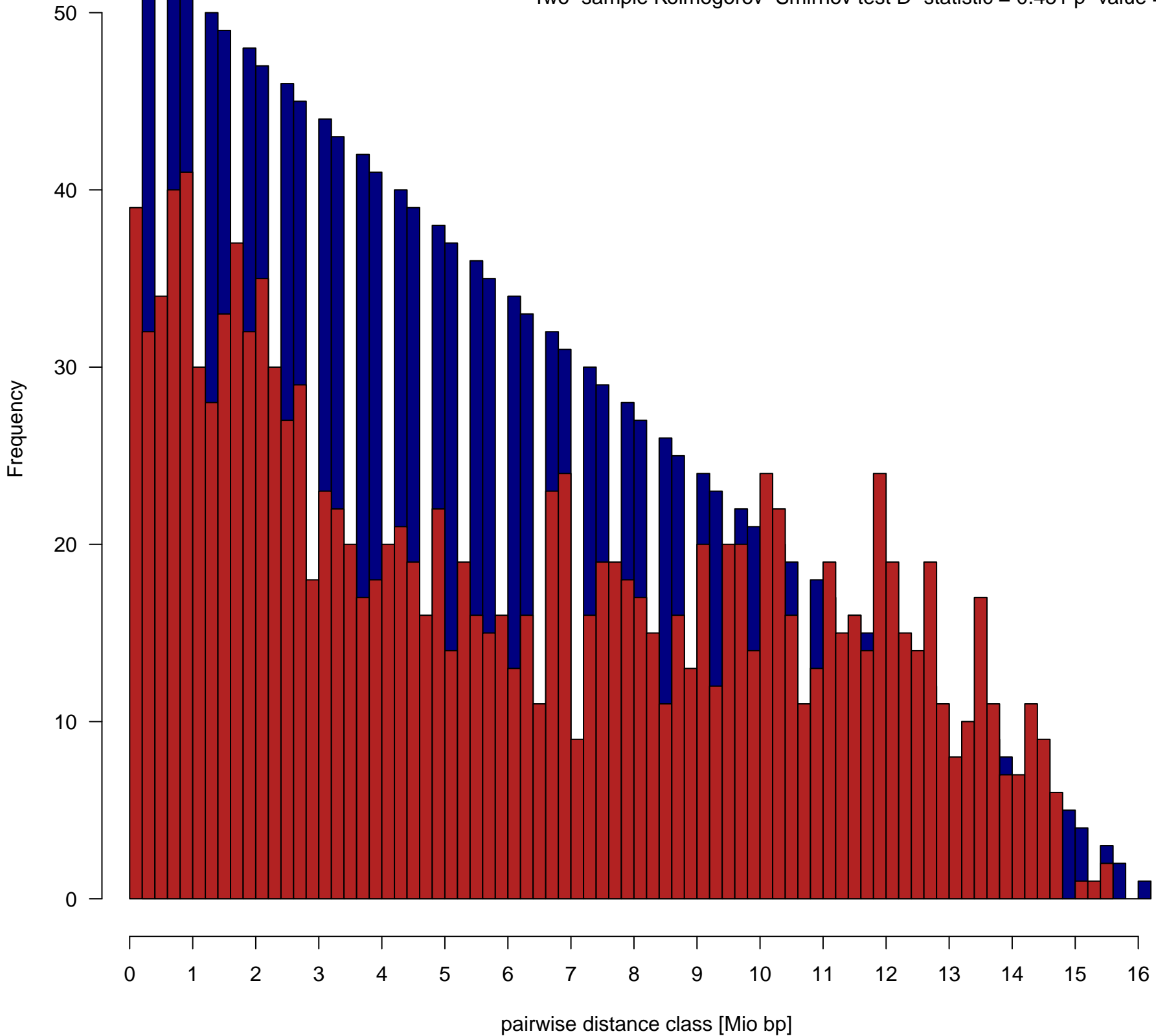
# Observed and expected distances between no-fixation regions on chromosome 12 in Italy

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.679 p-value = 0



# Observed and expected distances between no-fixation regions on chromosome 13 in Italy

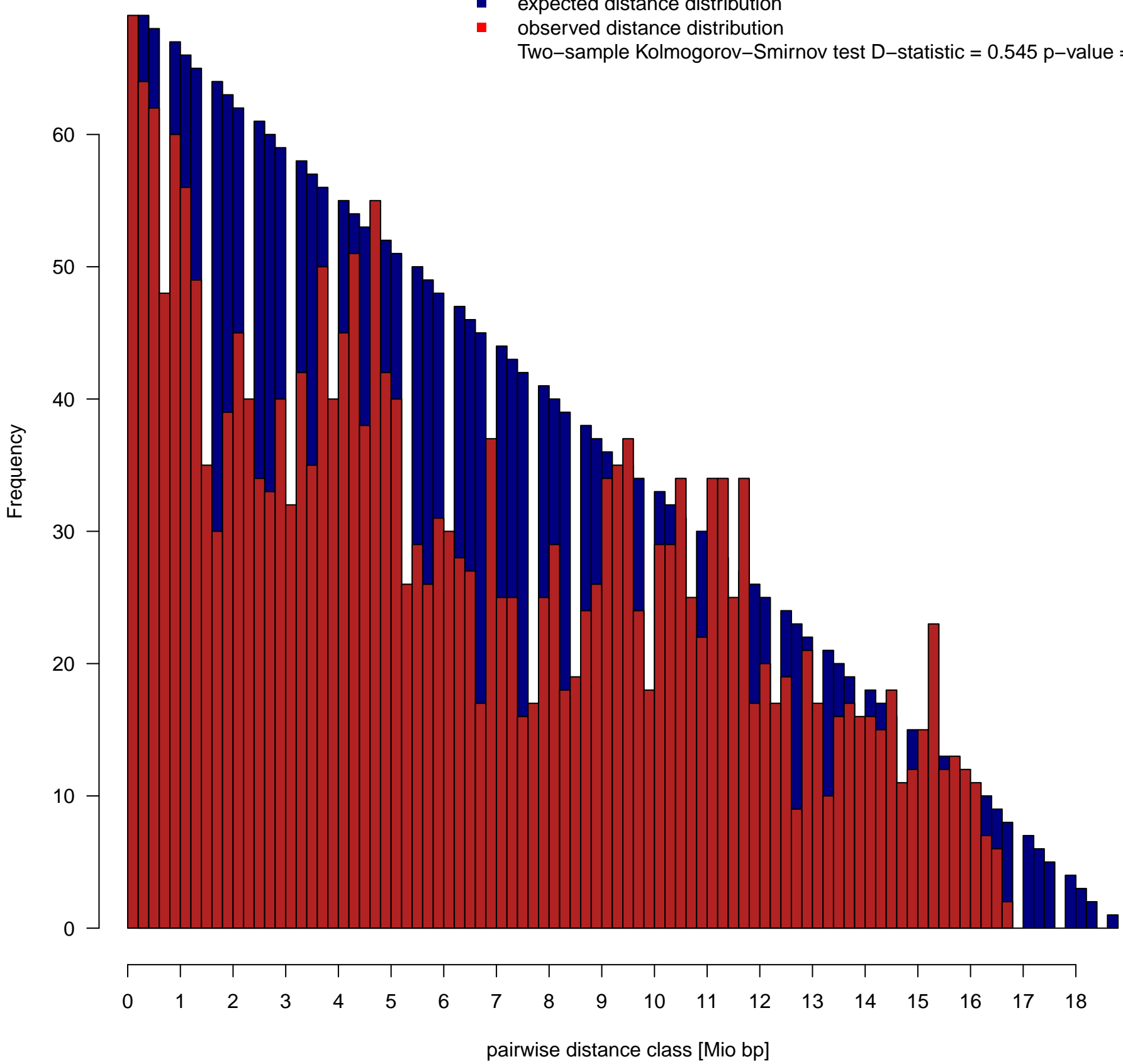
■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.451 p-value = 0



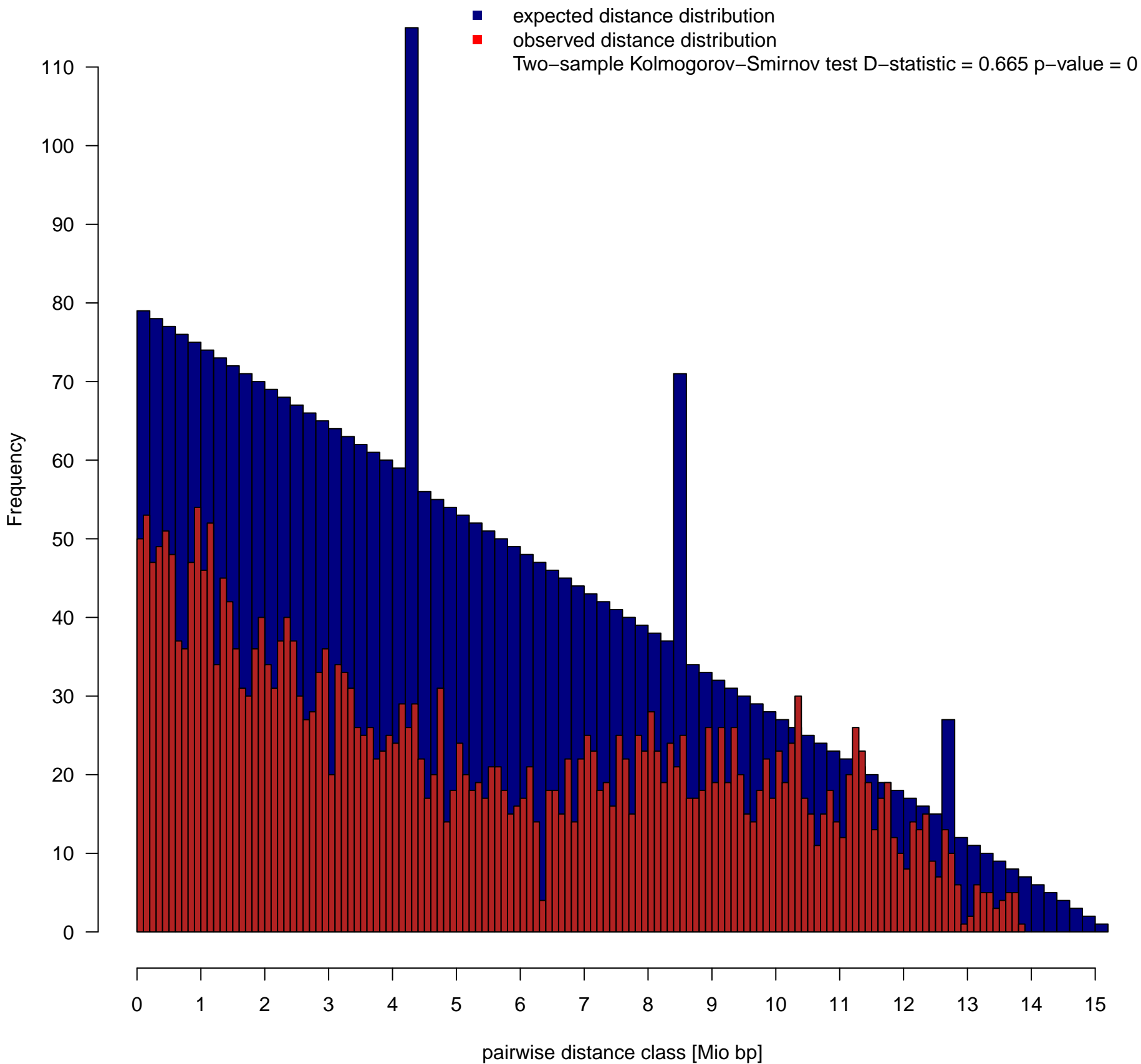


# Observed and expected distances between no-fixation regions on chromosome 14 in Italy

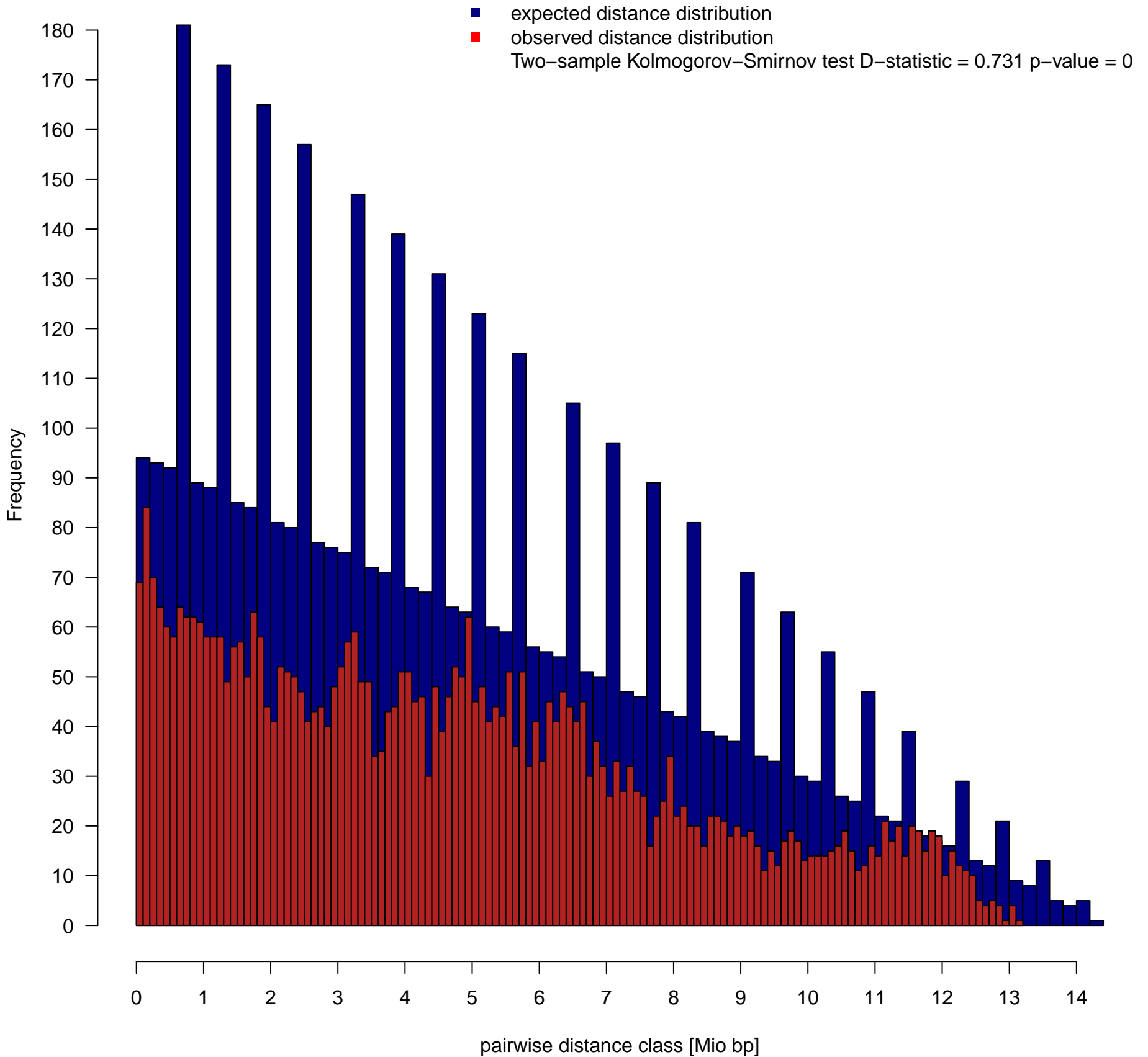
■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.545 p-value = 0



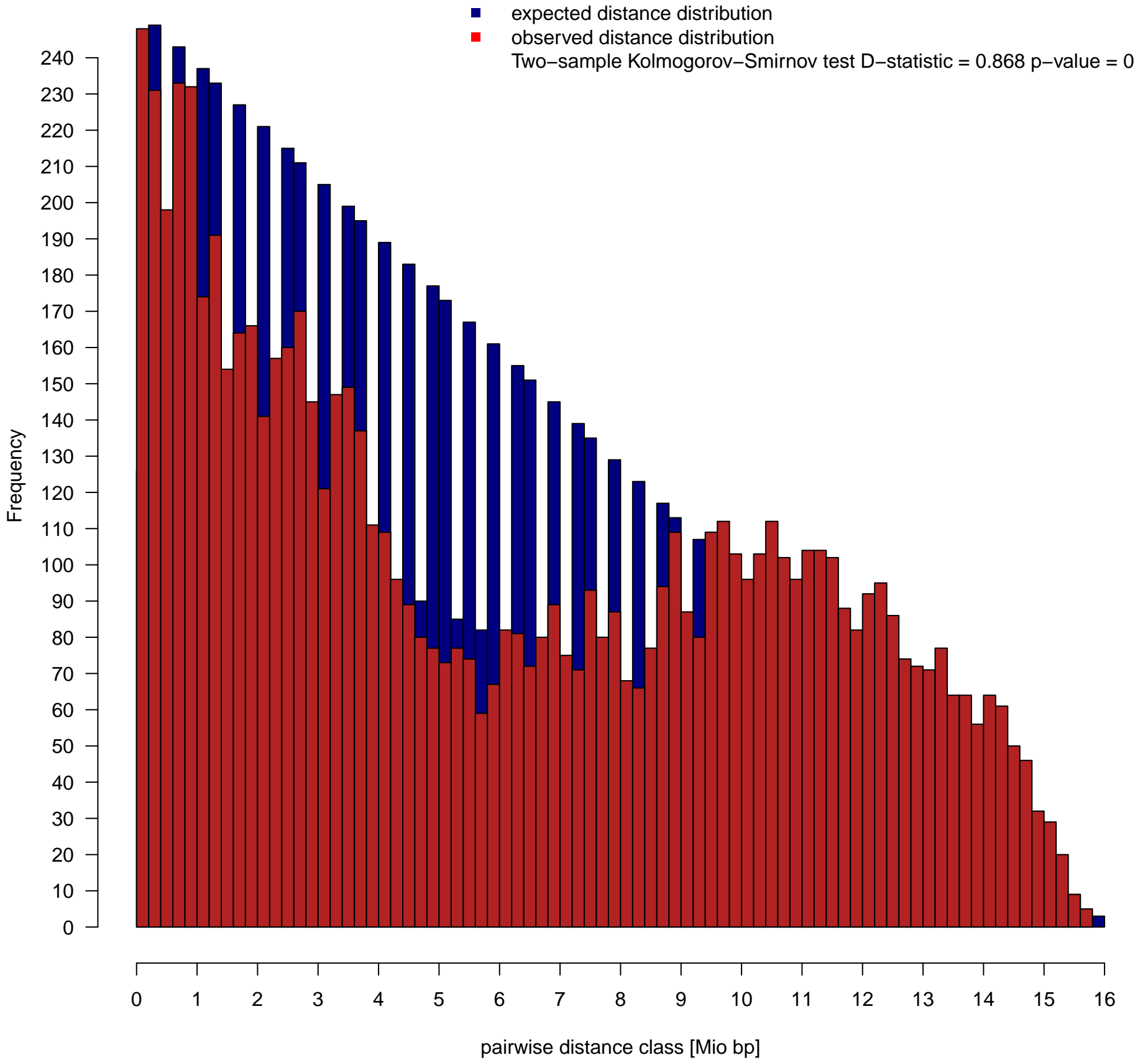
# Observed and expected distances between no-fixation regions on chromosome 15 in Italy



# Observed and expected distances between no-fixation regions on chromosome 16 in Italy

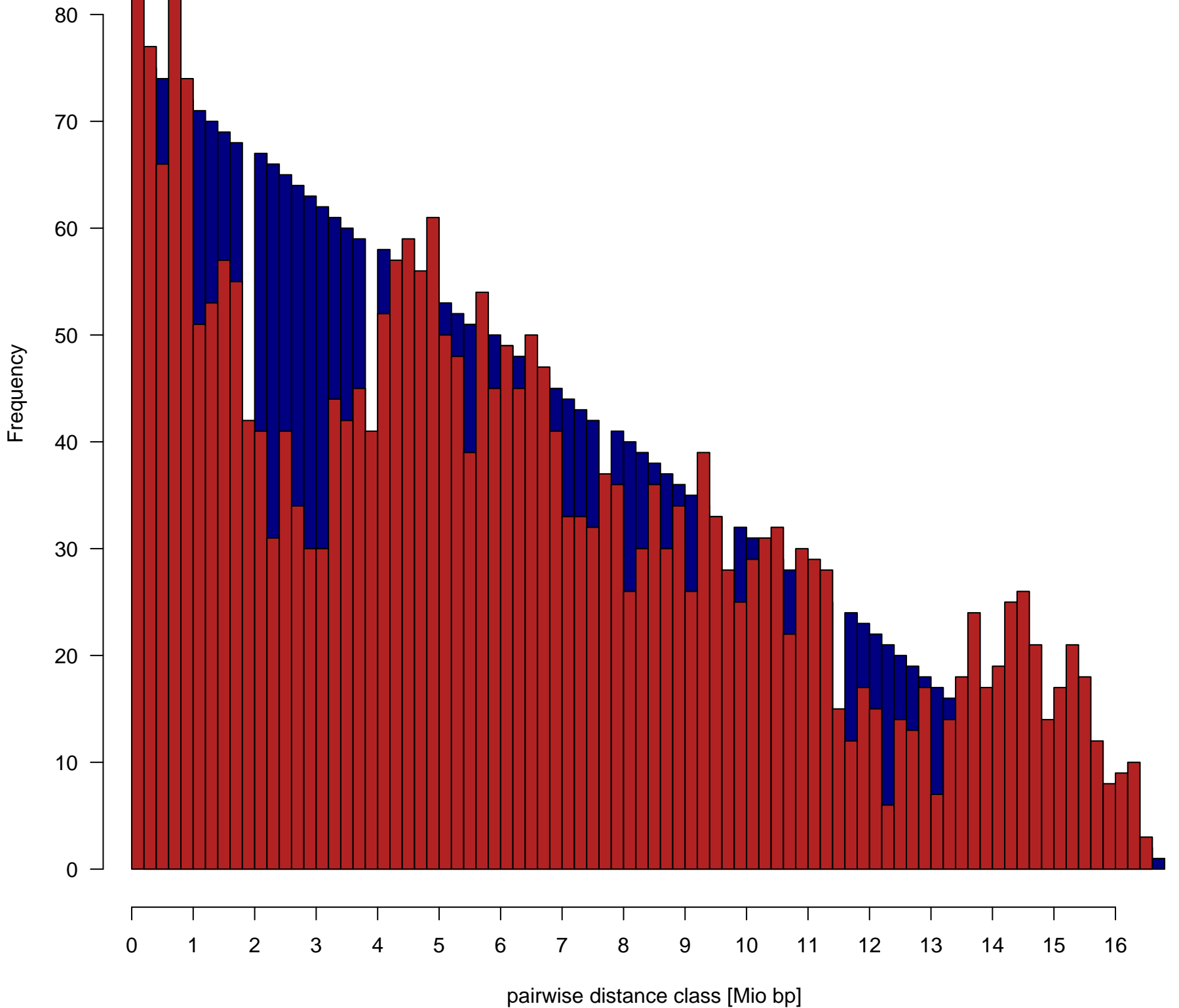


# Observed and expected distances between no-fixation regions on chromosome 17 in Italy



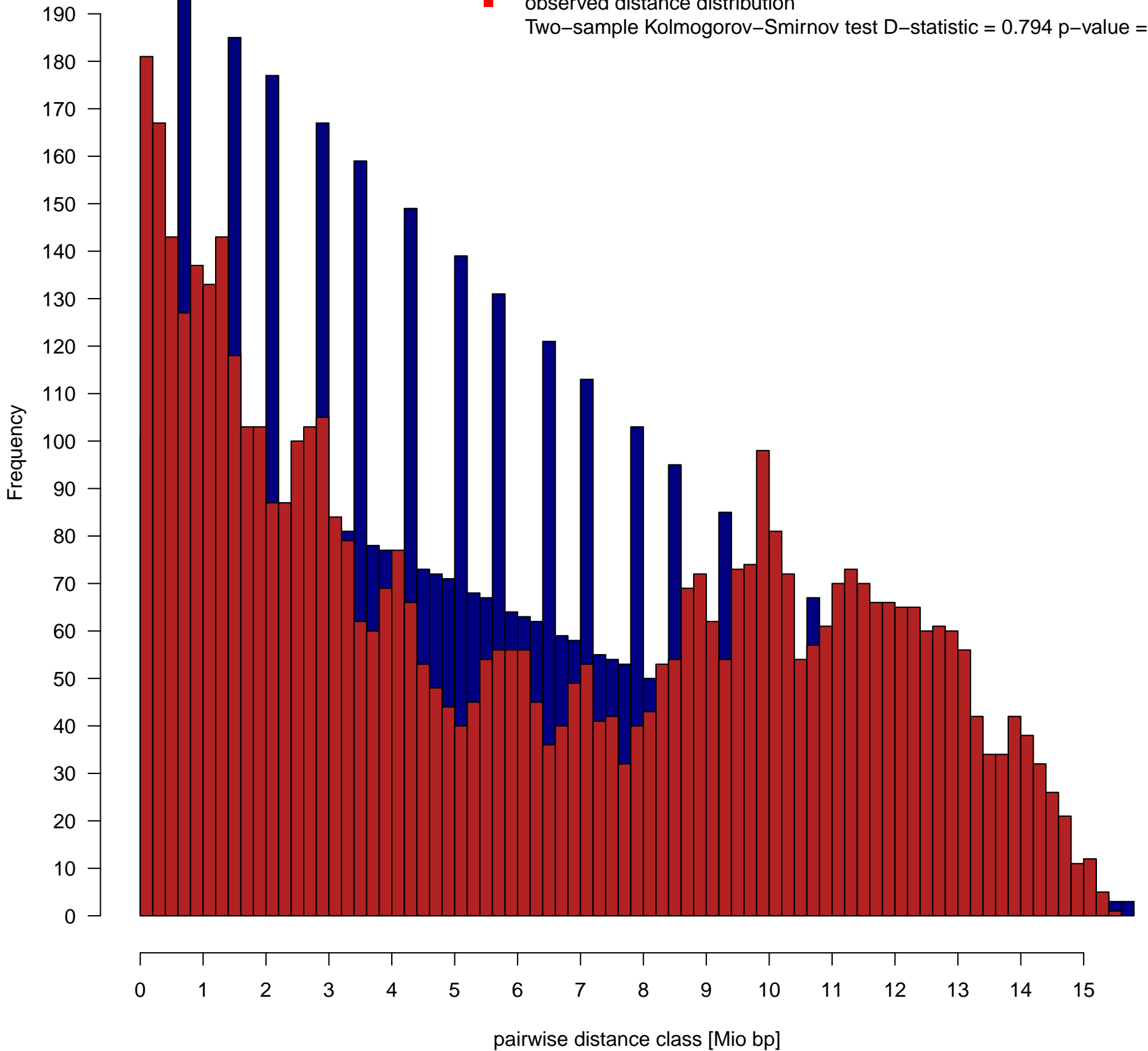
# Observed and expected distances between no-fixation regions on chromosome 18 in Italy

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.635 p-value = 0



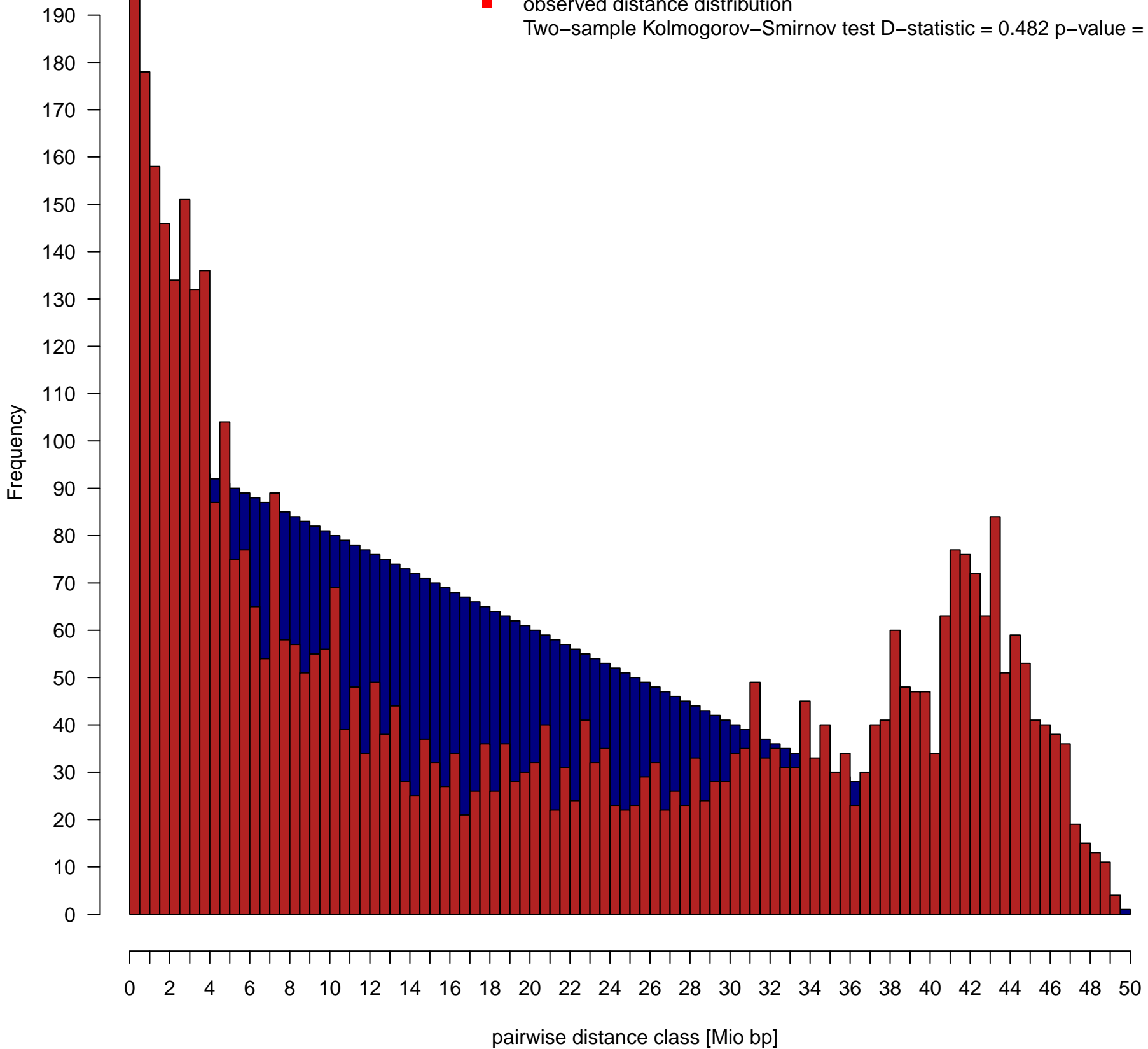
# Observed and expected distances between no-fixation regions on chromosome 19 in Italy

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.794 p-value = 0



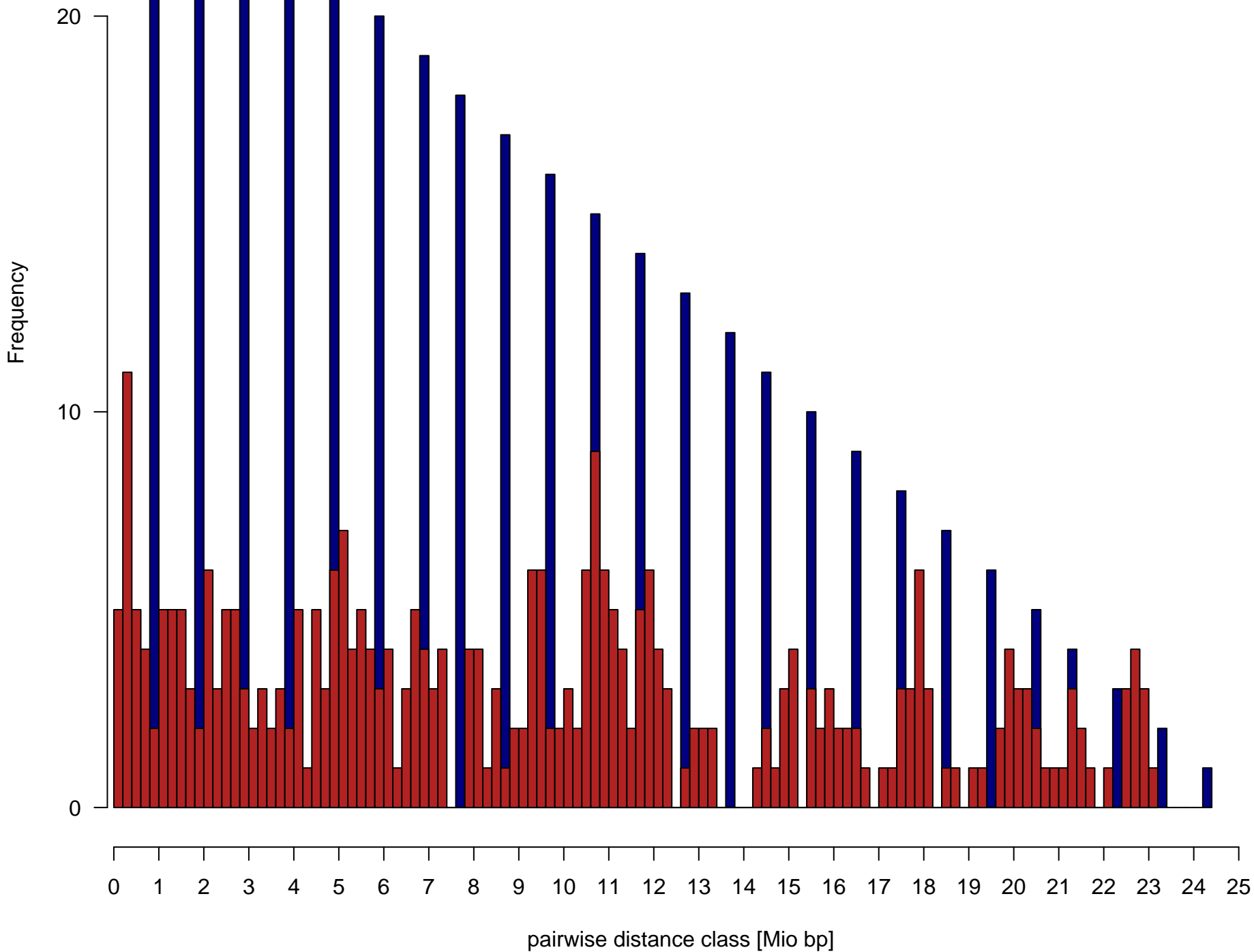
# Observed and expected distances between no-fixation regions on chromosome 1 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.482 p-value = 0



# Observed and expected distances between no-fixation regions on chromosome 2 in Hungary

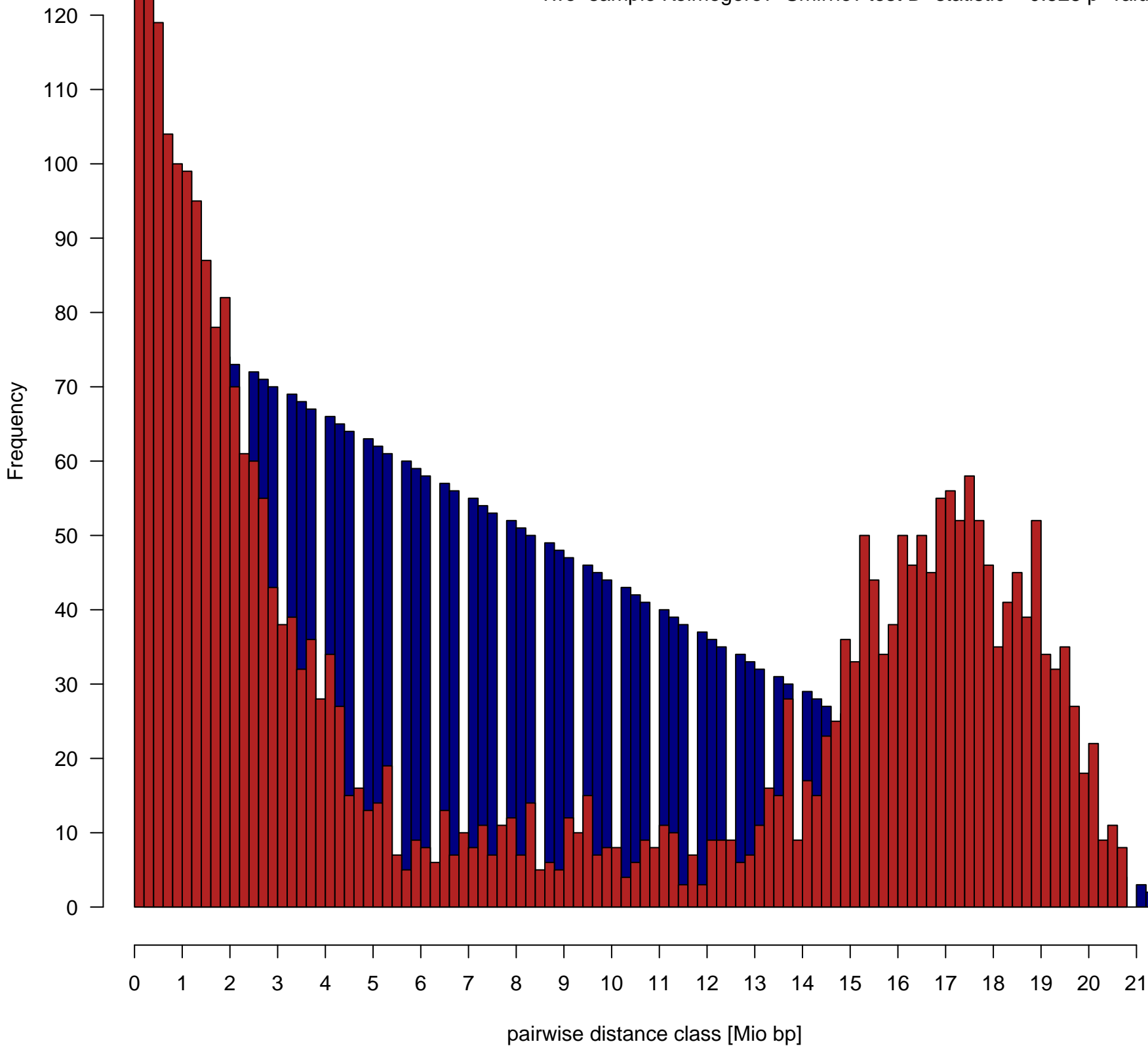
■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.091 p-value = 0





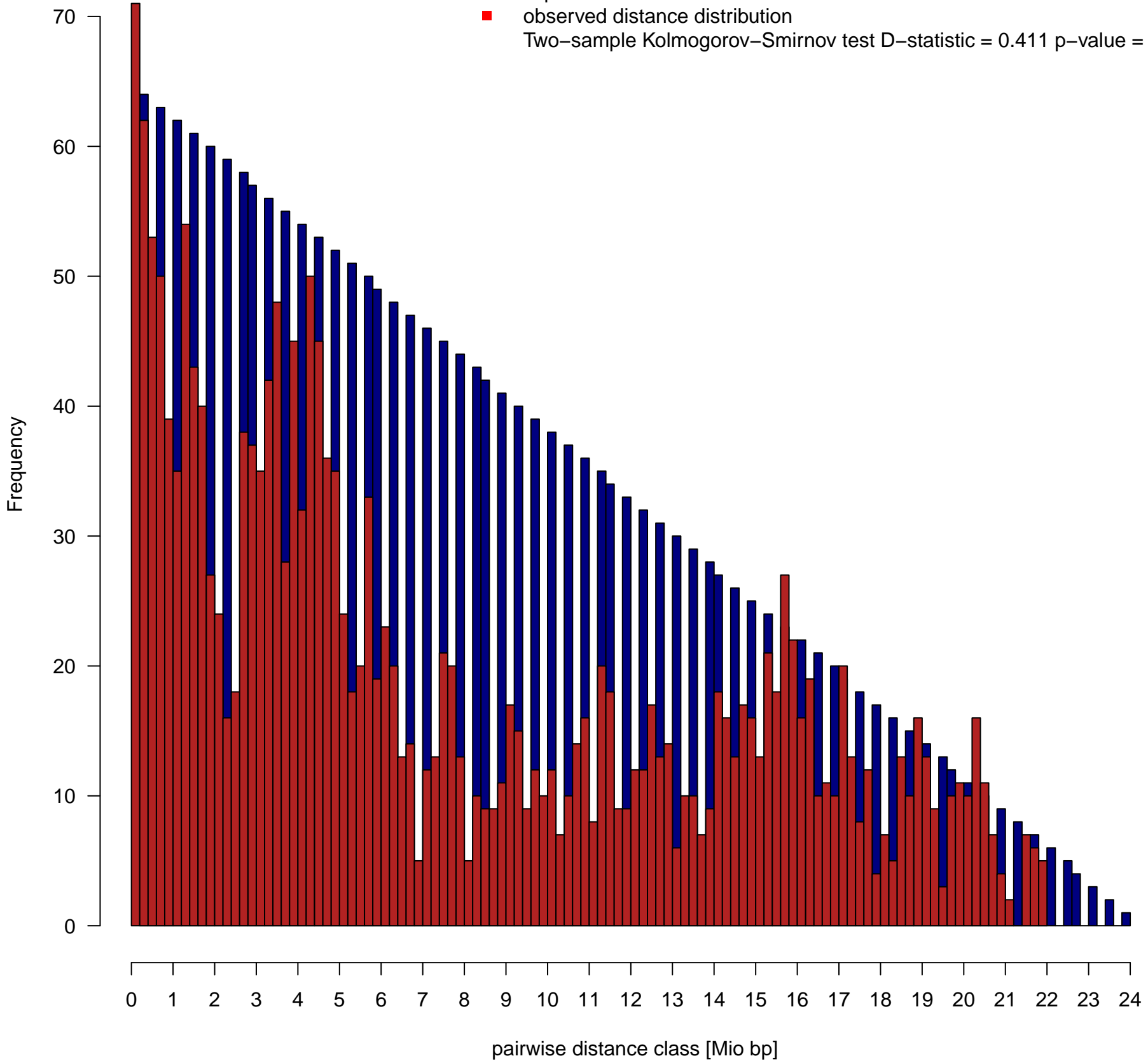
# Observed and expected distances between no-fixation regions on chromosome 3 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.528 p-value = 0



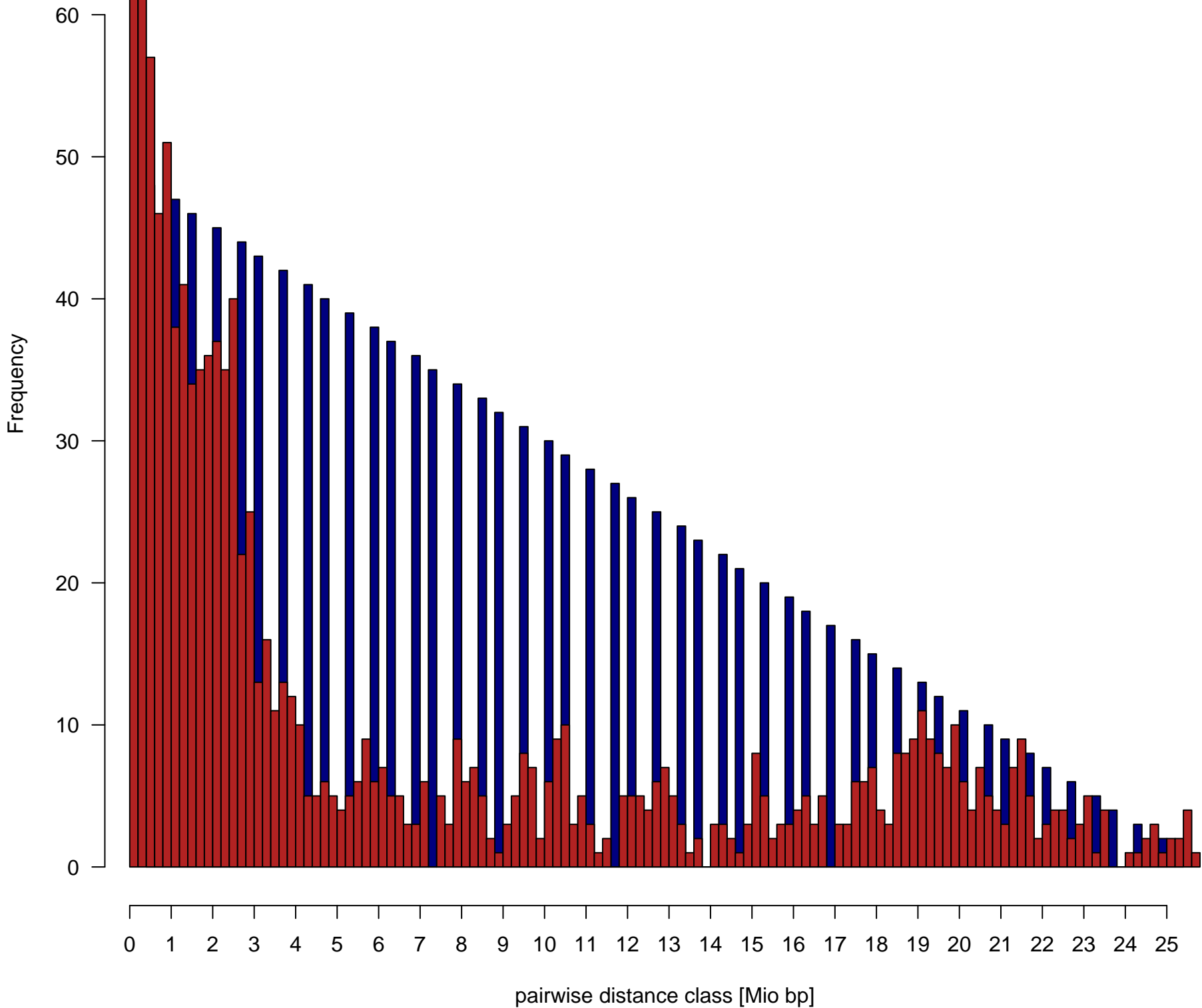
# Observed and expected distances between no-fixation regions on chromosome 4 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.411 p-value = 0



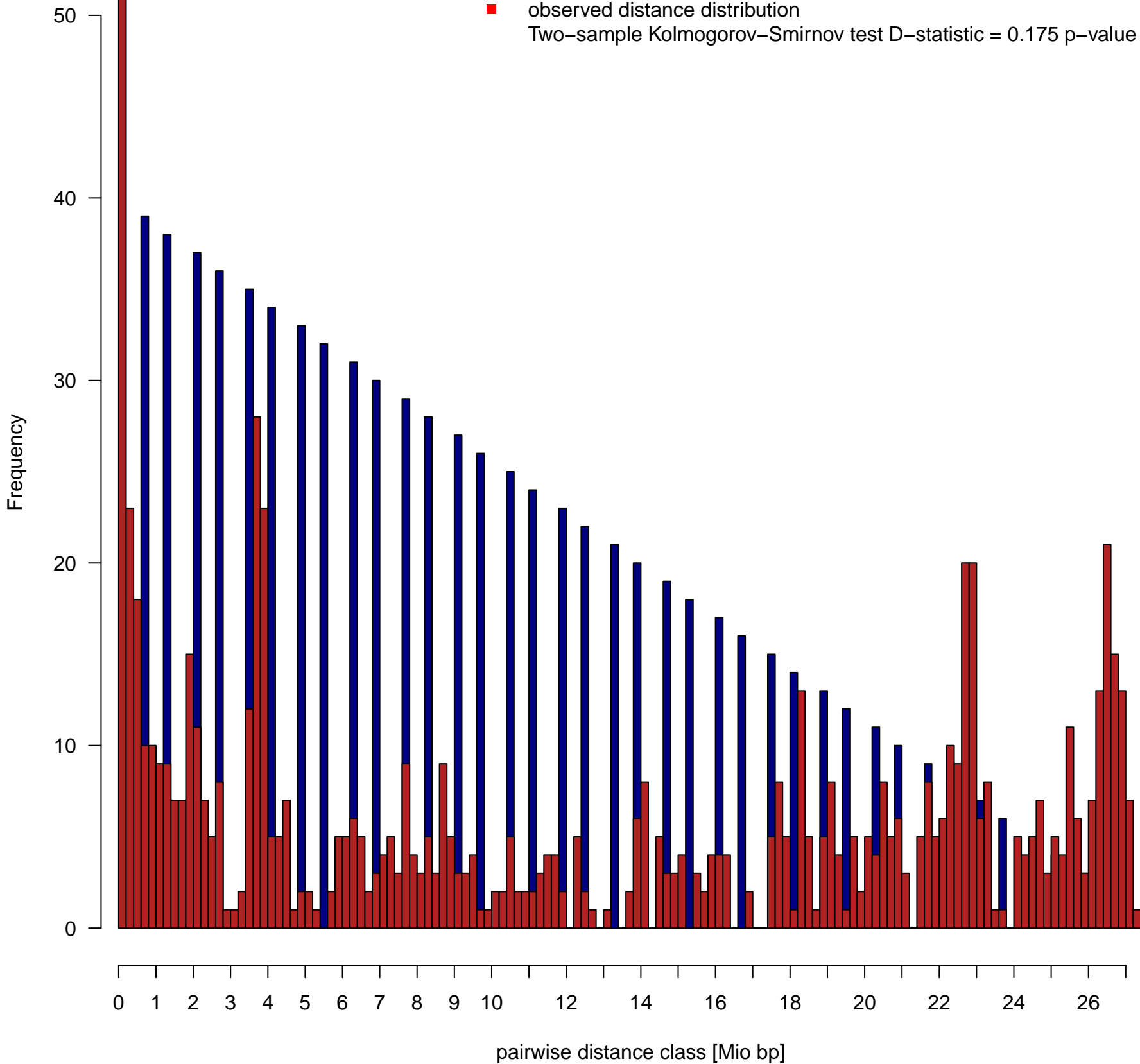
# Observed and expected distances between no-fixation regions on chromosome 5 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.231 p-value = 0



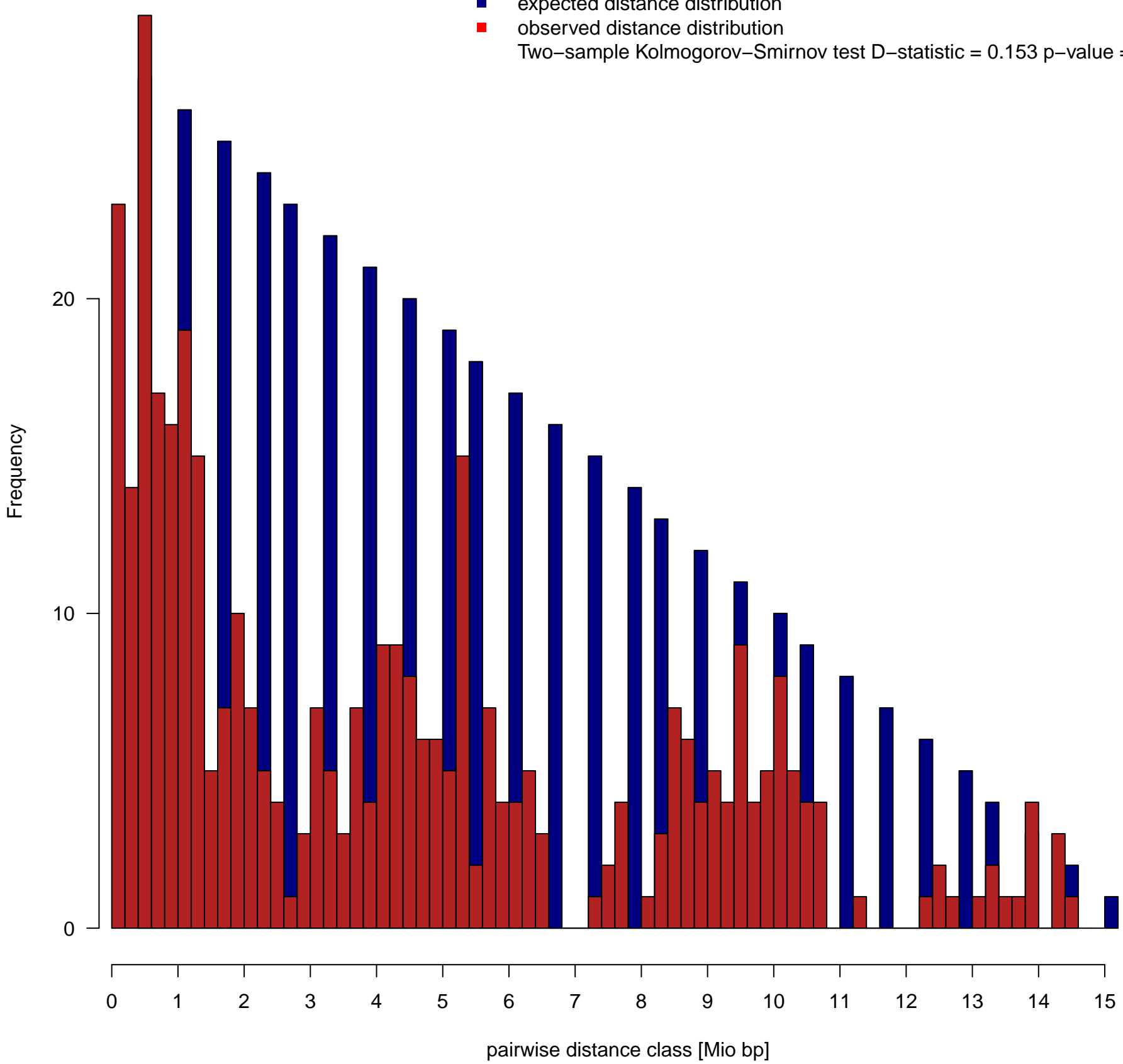
# Observed and expected distances between no-fixation regions on chromosome 6 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.175 p-value = 0



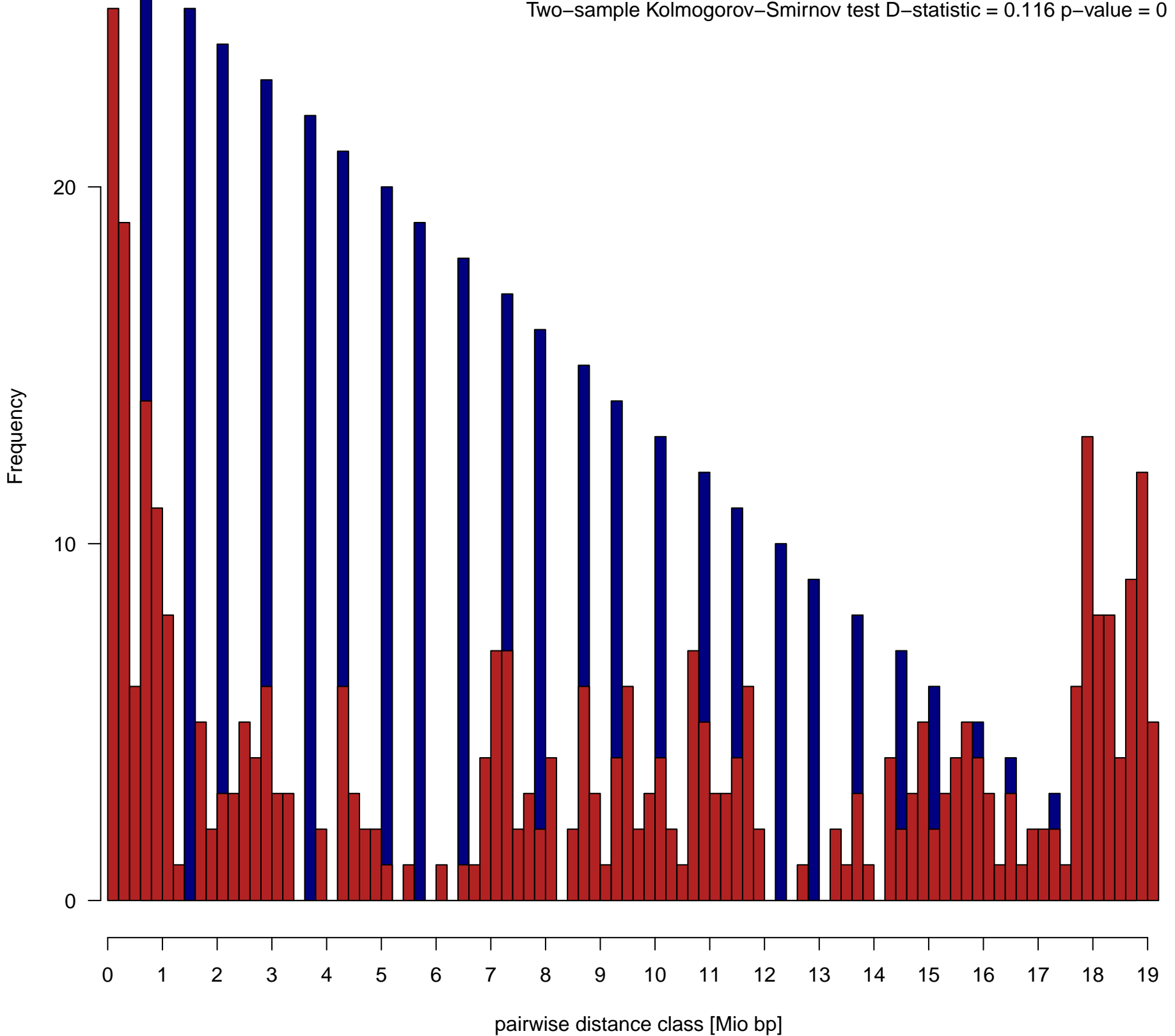
# Observed and expected distances between no-fixation regions on chromosome 7 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.153 p-value = 0



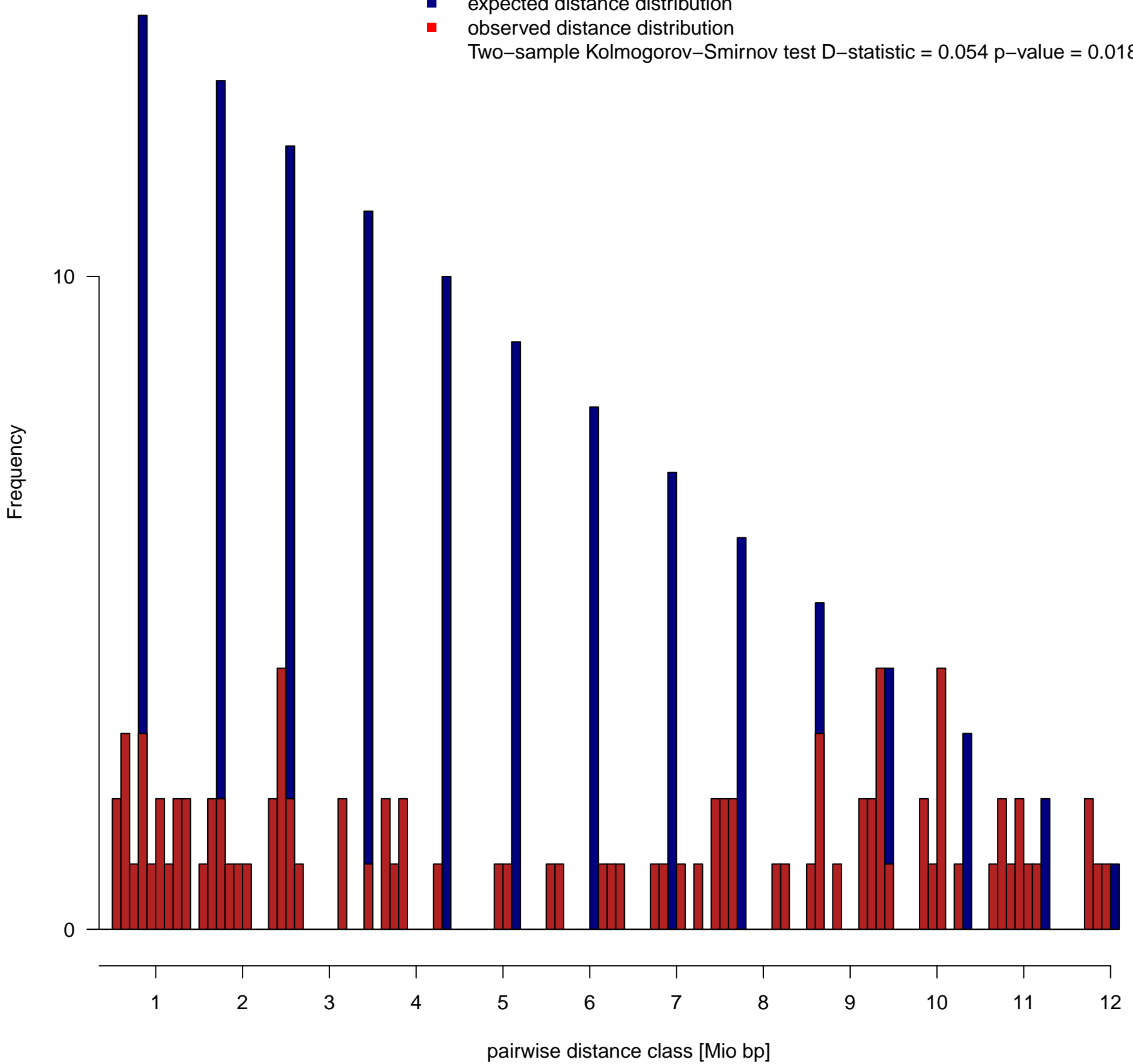
# Observed and expected distances between no-fixation regions on chromosome 8 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.116 p-value = 0



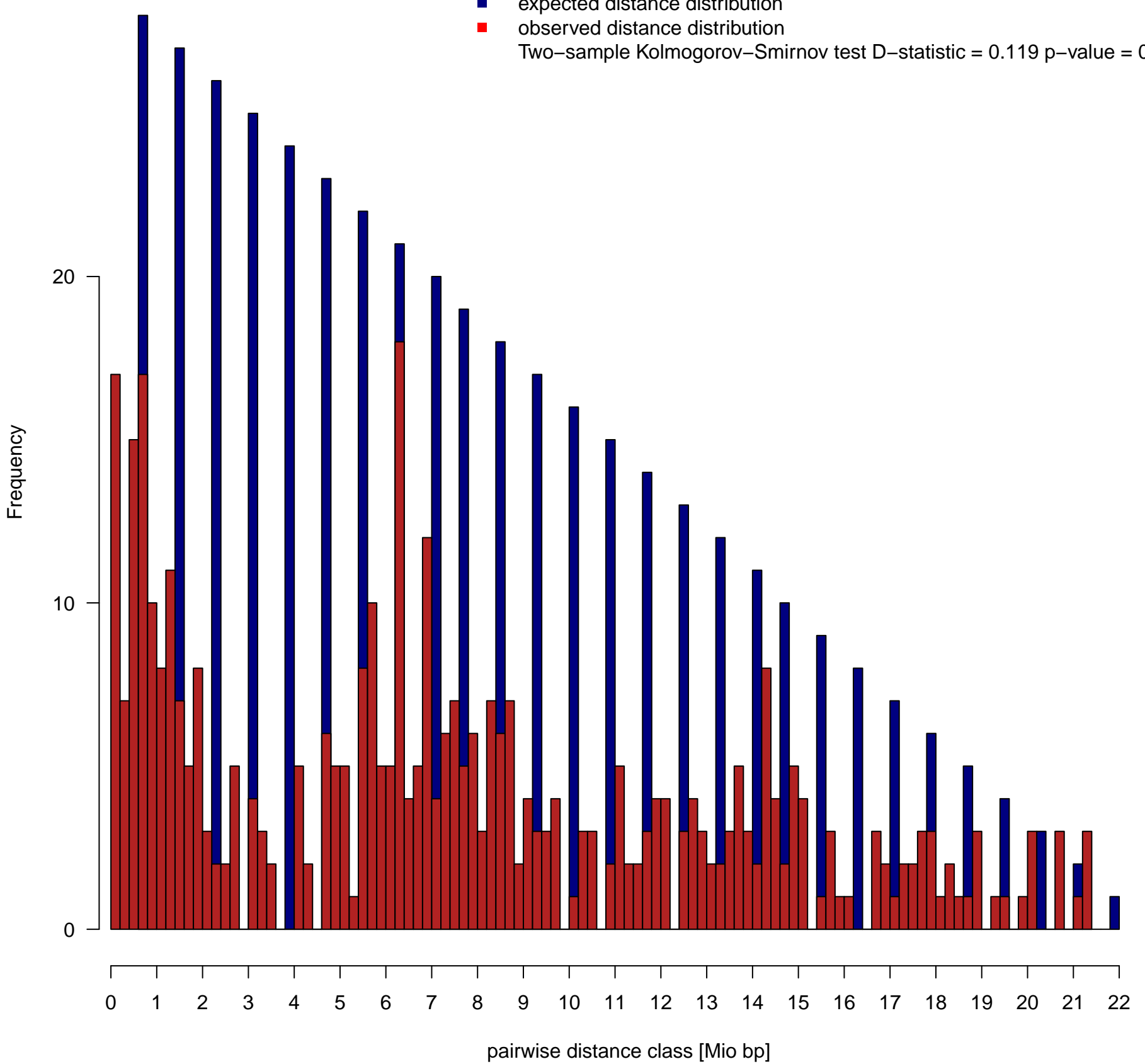
# Observed and expected distances between no-fixation regions on chromosome 9 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.054 p-value = 0.0186



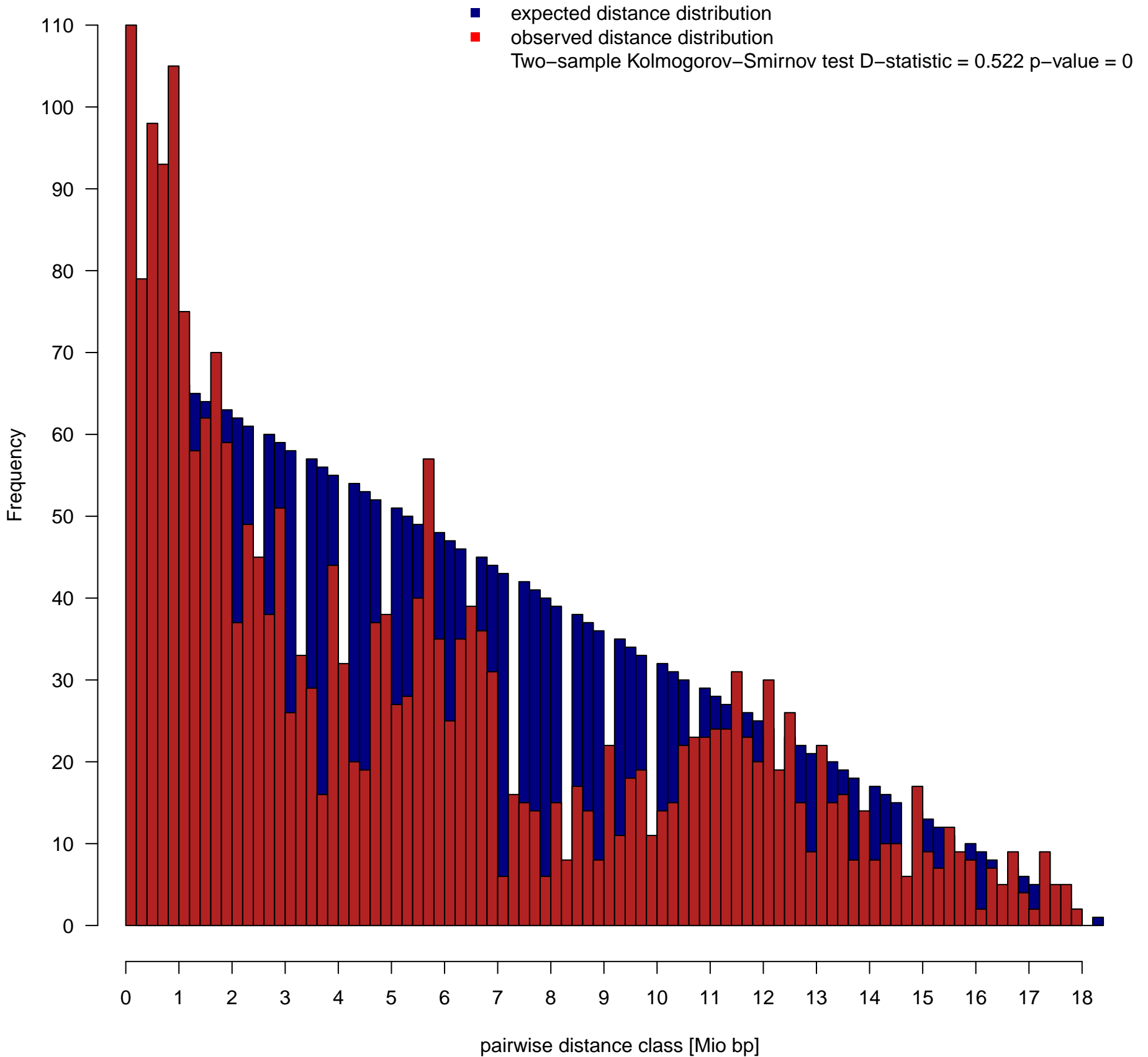
# Observed and expected distances between no-fixation regions on chromosome 10 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.119 p-value = 0

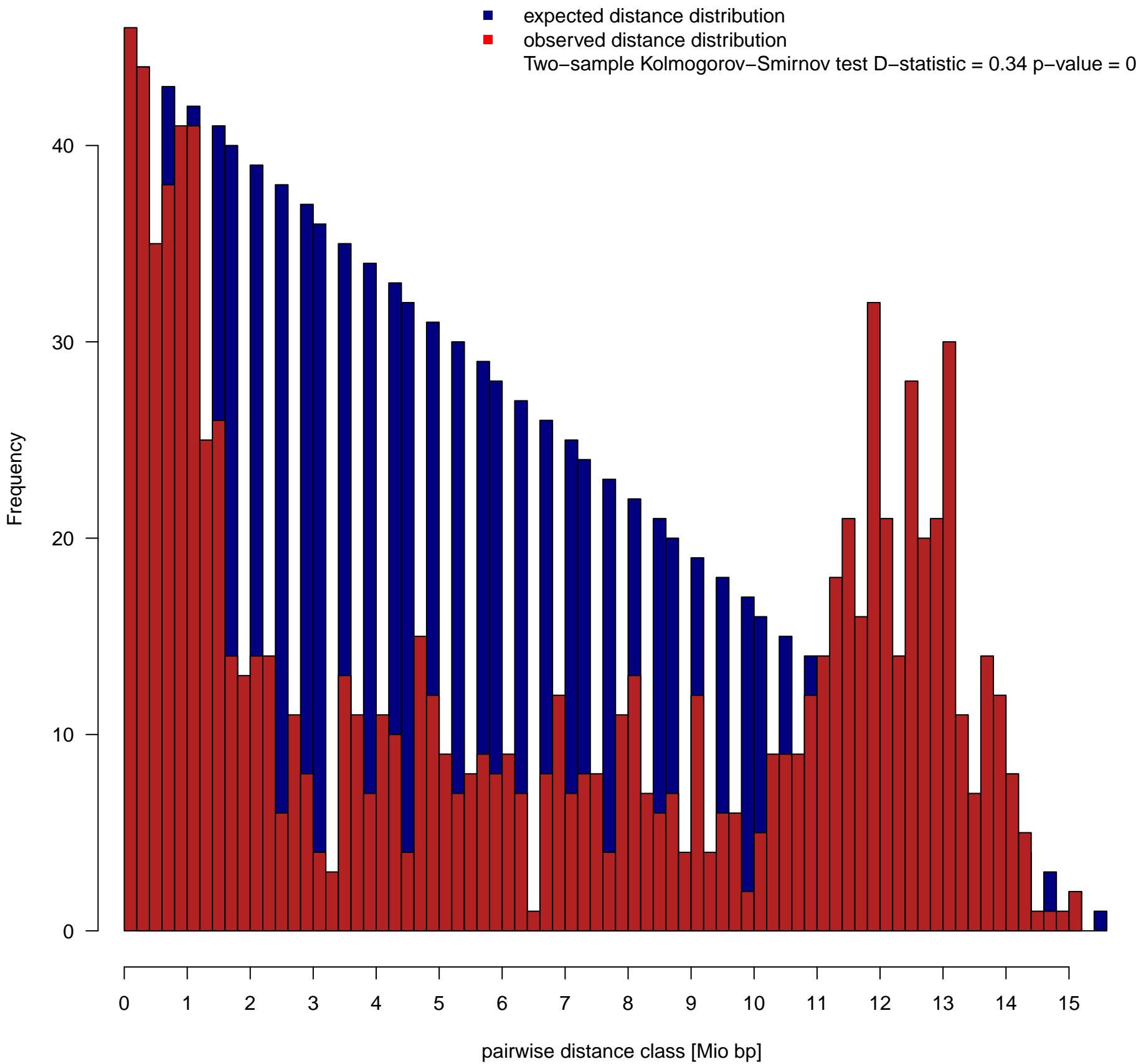




# Observed and expected distances between no-fixation regions on chromosome 11 in Hungary

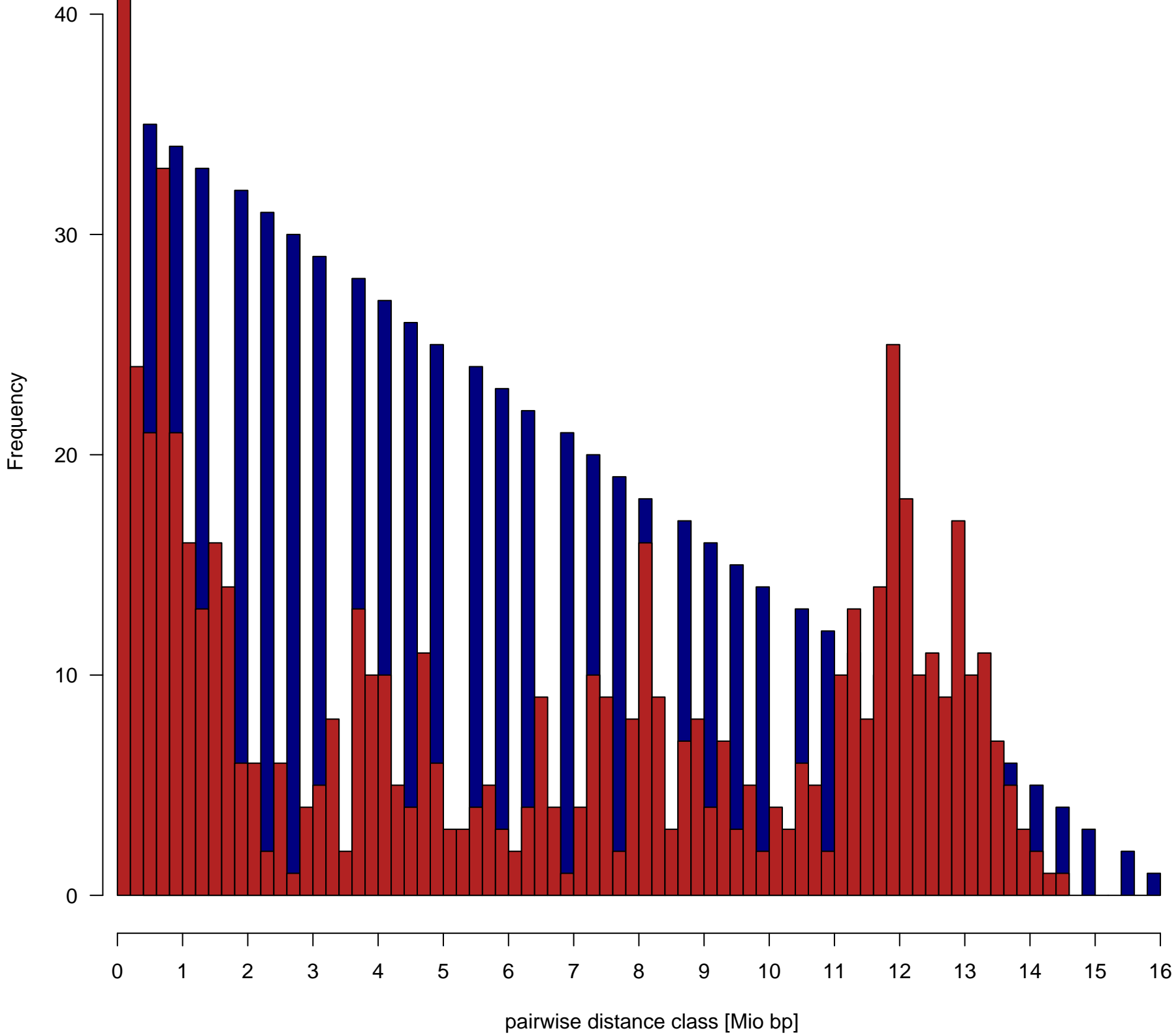


# Observed and expected distances between no-fixation regions on chromosome 12 in Hungary



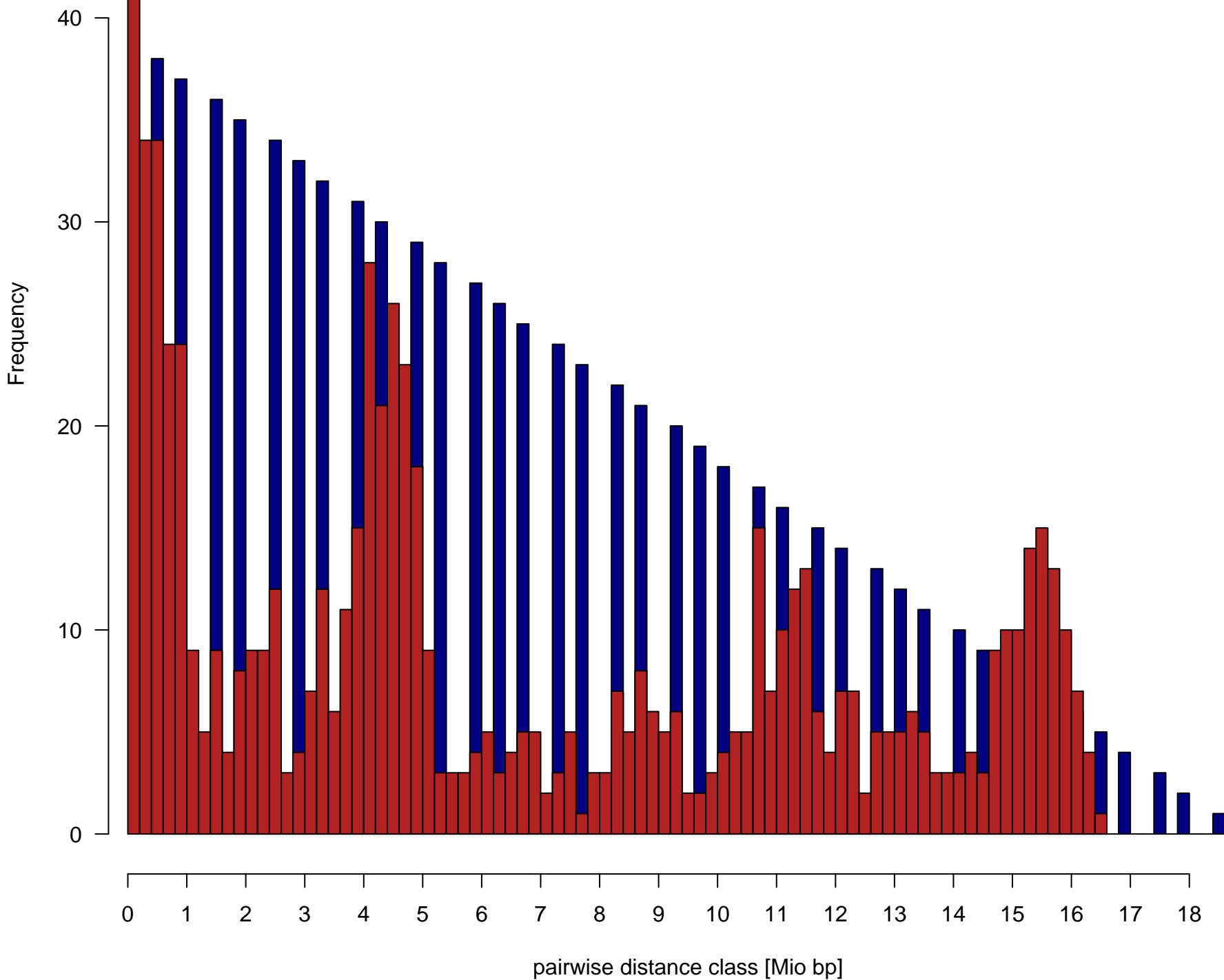
# Observed and expected distances between no-fixation regions on chromosome 13 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.222 p-value = 0



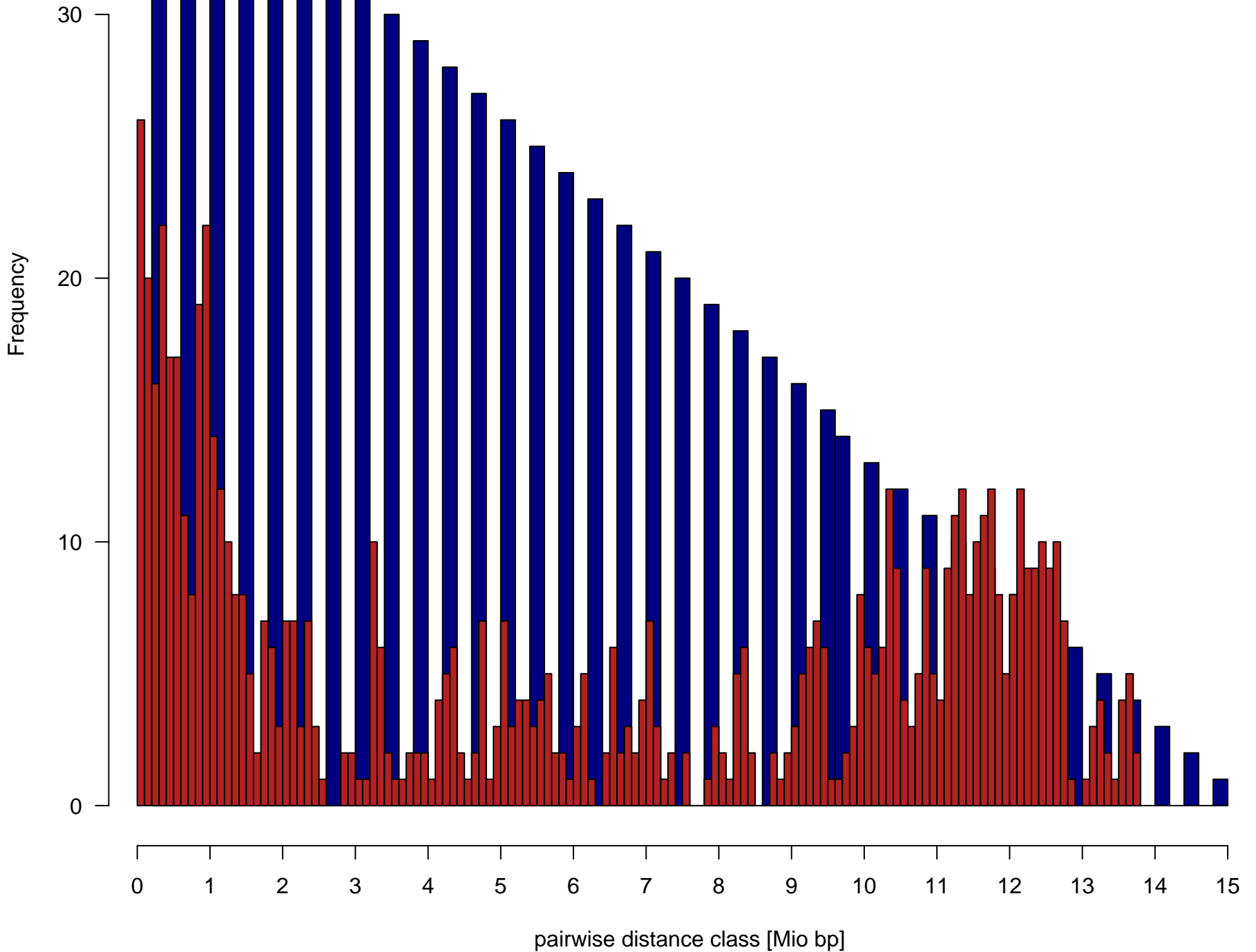
# Observed and expected distances between no-fixation regions on chromosome 14 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.23 p-value = 0



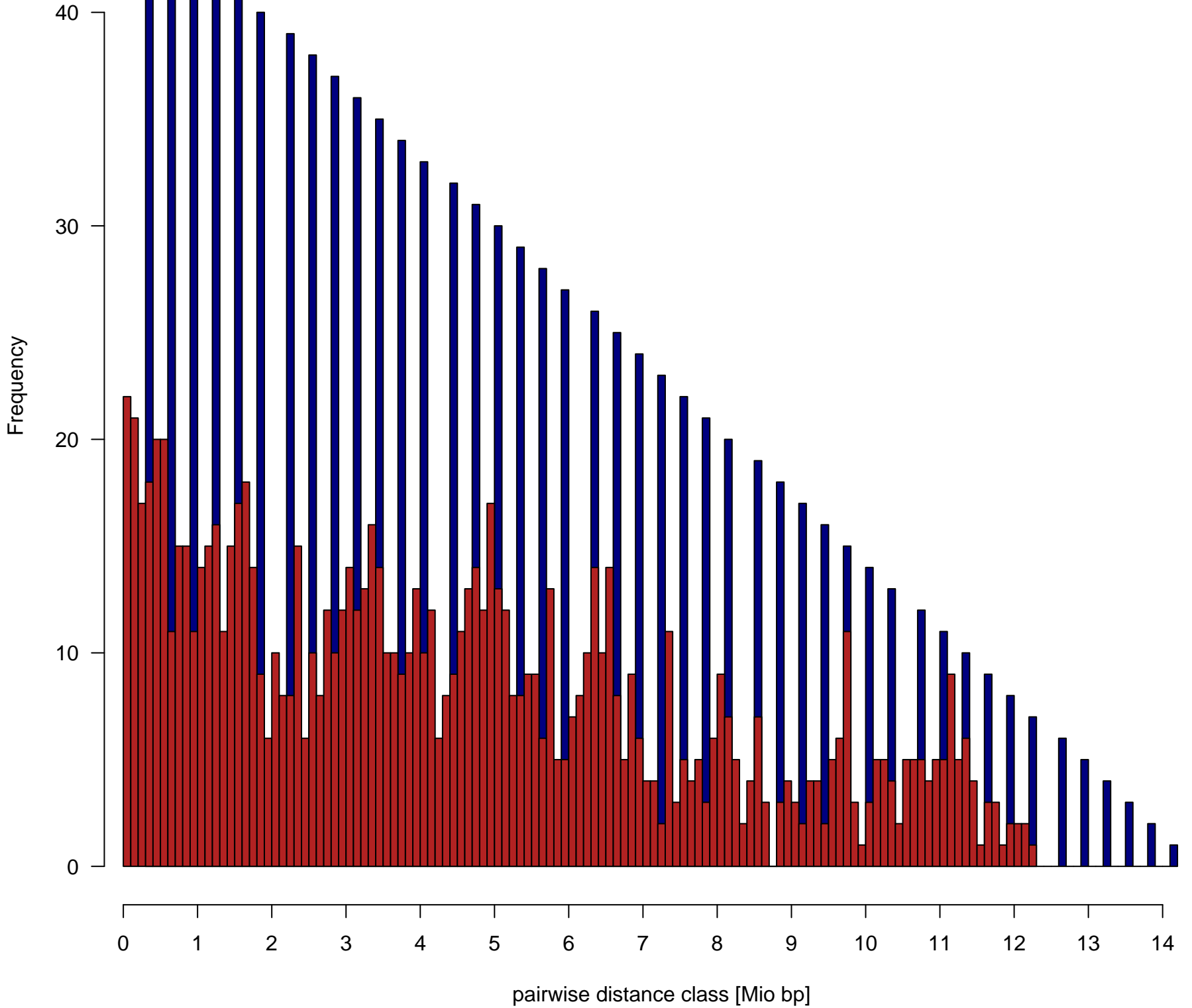
# Observed and expected distances between no-fixation regions on chromosome 15 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.271 p-value = 0



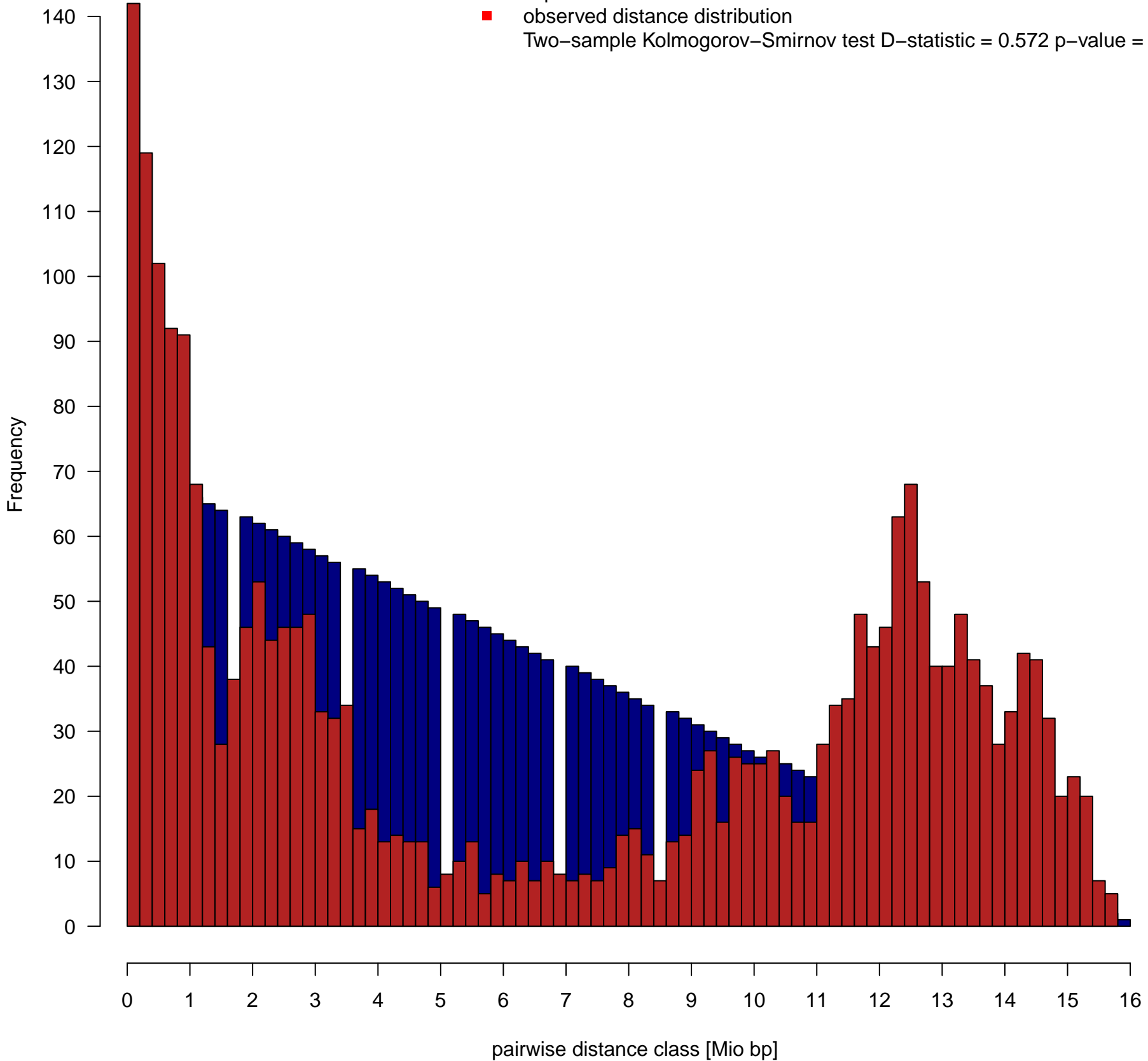
# Observed and expected distances between no-fixation regions on chromosome 16 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.374 p-value = 0



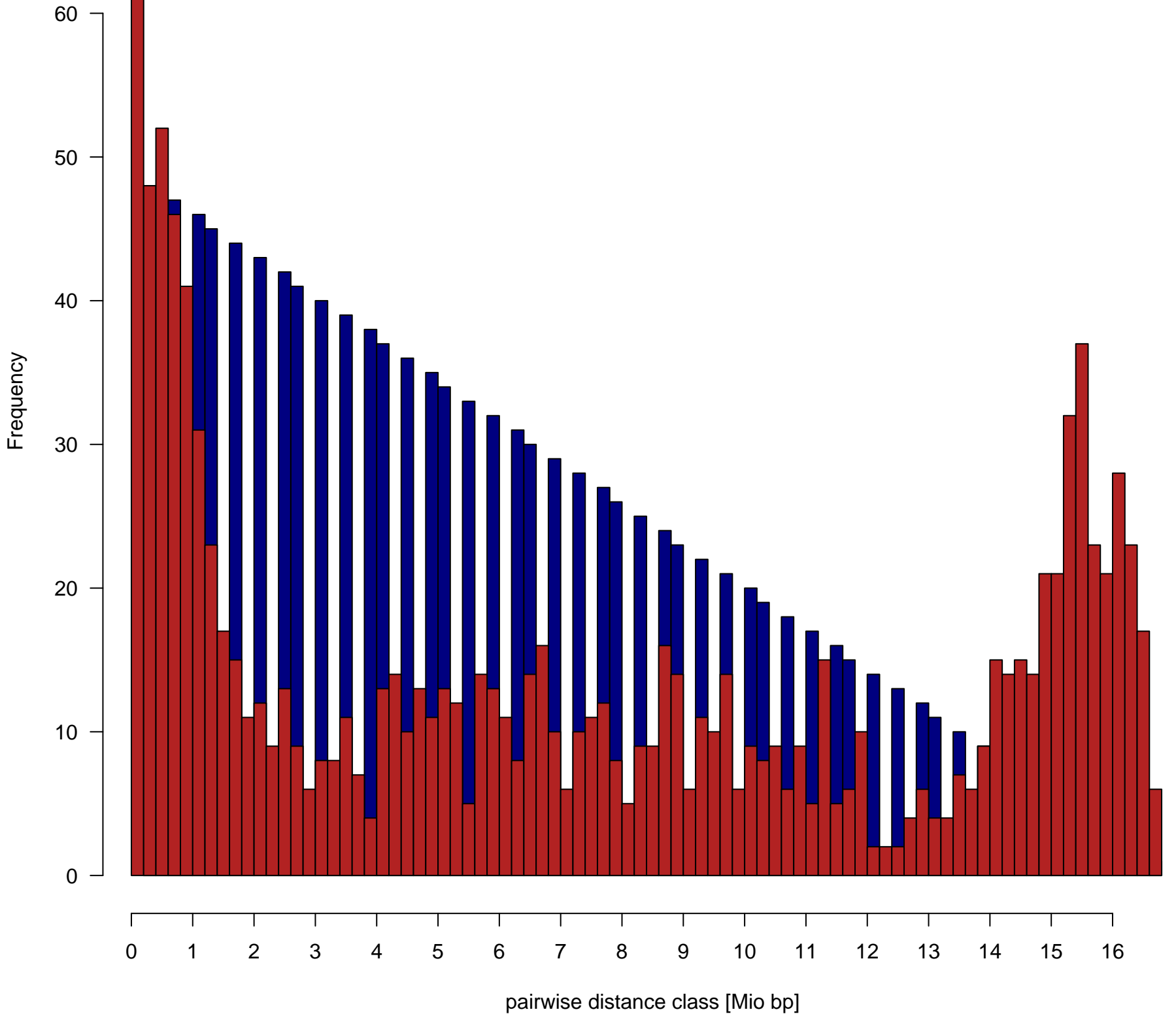
# Observed and expected distances between no-fixation regions on chromosome 17 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.572 p-value = 0



# Observed and expected distances between no-fixation regions on chromosome 18 in Hungary

■ expected distance distribution  
■ observed distance distribution  
Two-sample Kolmogorov-Smirnov test D-statistic = 0.367 p-value = 0





# Observed and expected distances between no-fixation regions on chromosome 19 in Hungary

