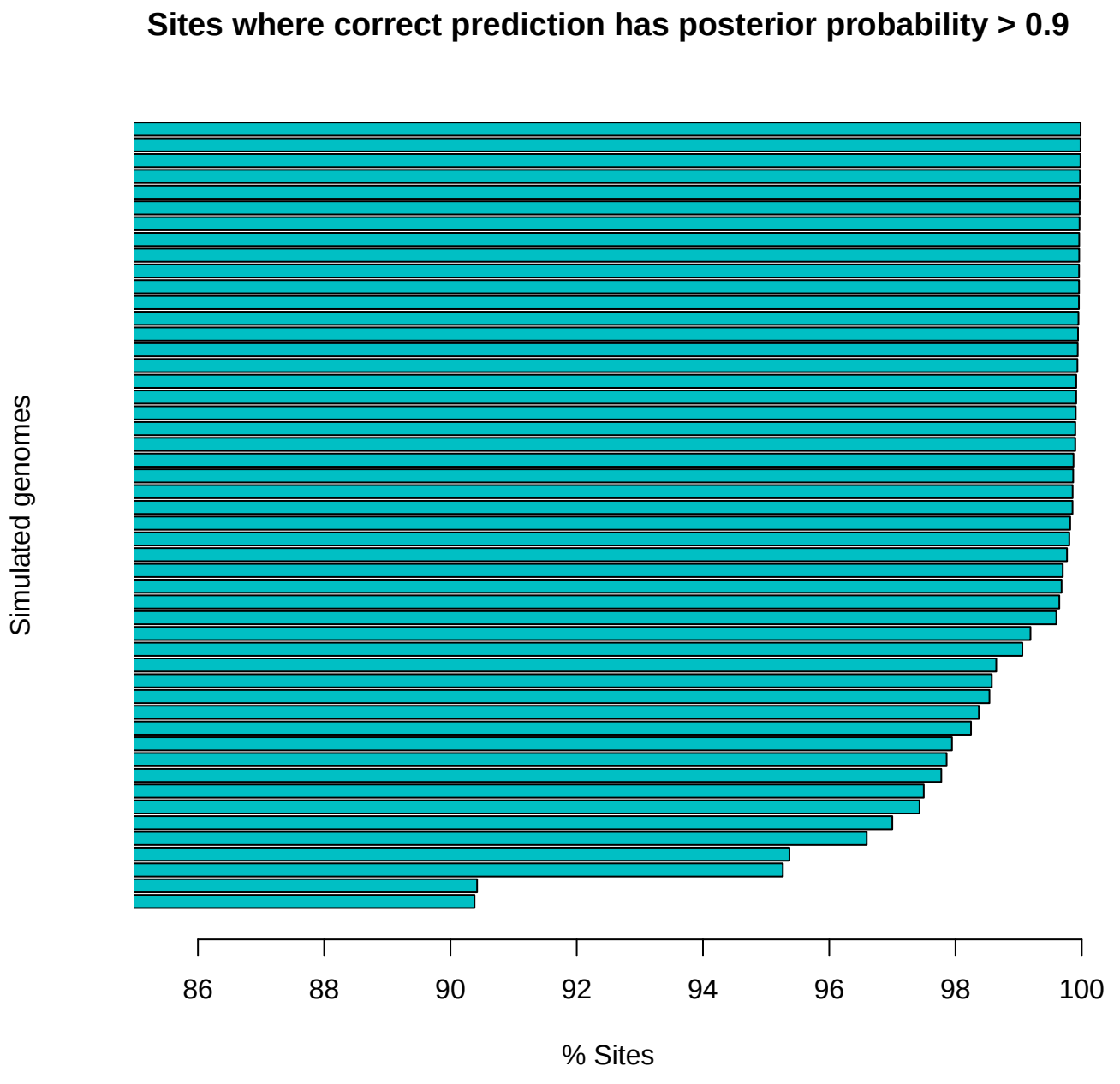
Sites where correct prediction has posterior probability > 0.9
Simulated genomes
86
88
90
92
94
96
98
100
% Sites