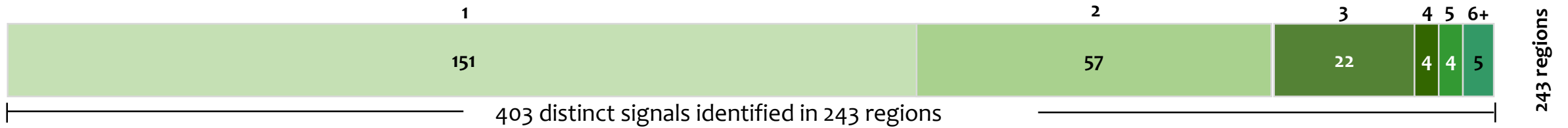
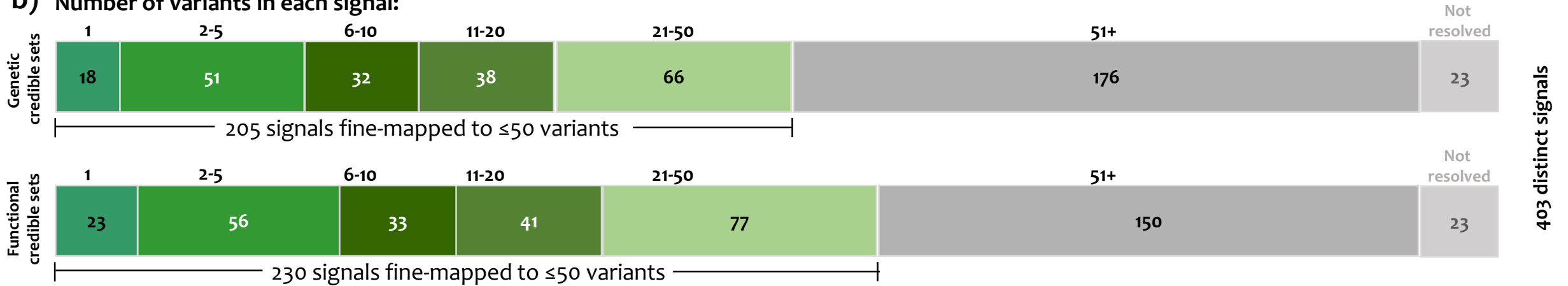


a) Number of signals/regions:



b) Number of variants in each signal:



c) Posterior probability for variant to be causal:

