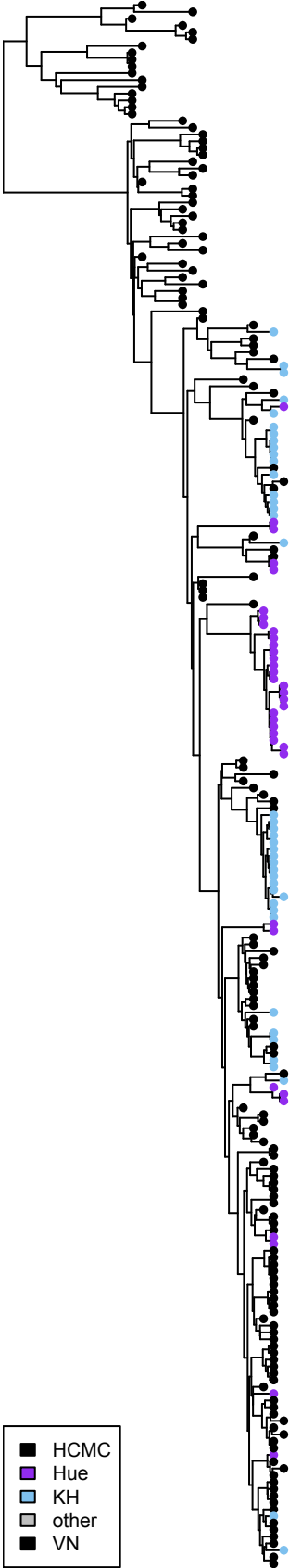


- HCMC
- Hue
- KH
- other
- VN



0 2.8 6.6 10.3 14.1



Genomic data visualization showing sequence alignment and coverage across a genomic region. The top track displays sequence alignment with various symbols (dots, lines) indicating mismatches or indels. Below the alignment, a series of colored bars (green, red, orange, blue, light blue, brown) represent different genomic features or annotations. The bottom track shows coverage or depth across the region, with a scale from 0 to 14.1 indicated on the left. The data is organized into two main sections, each with a corresponding color scale bar on the left.

