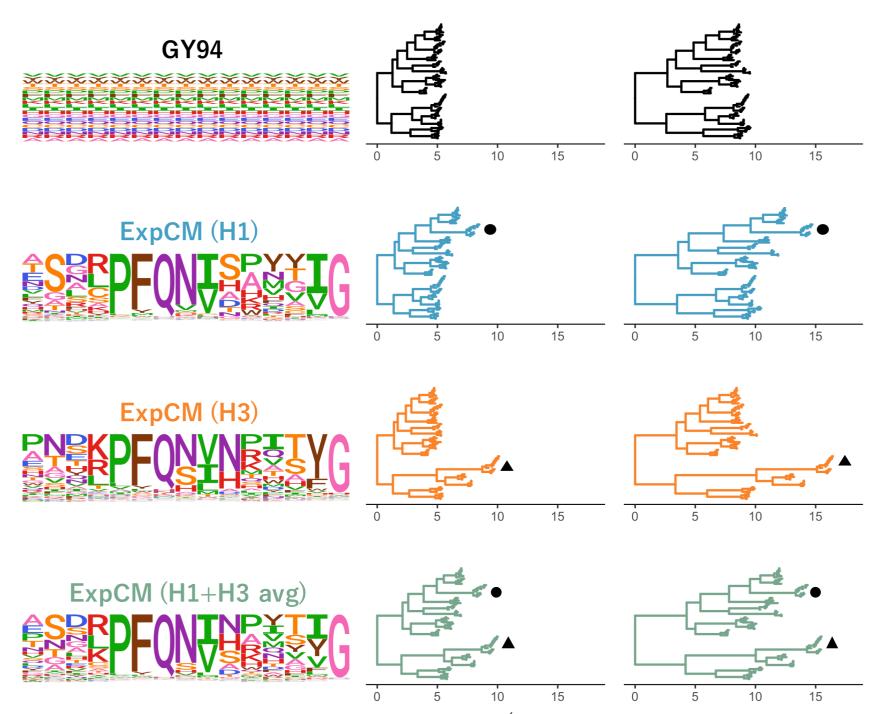
No rate variation

$\Gamma\omega$ rate variation



Branch length (average codon substitutions per site)