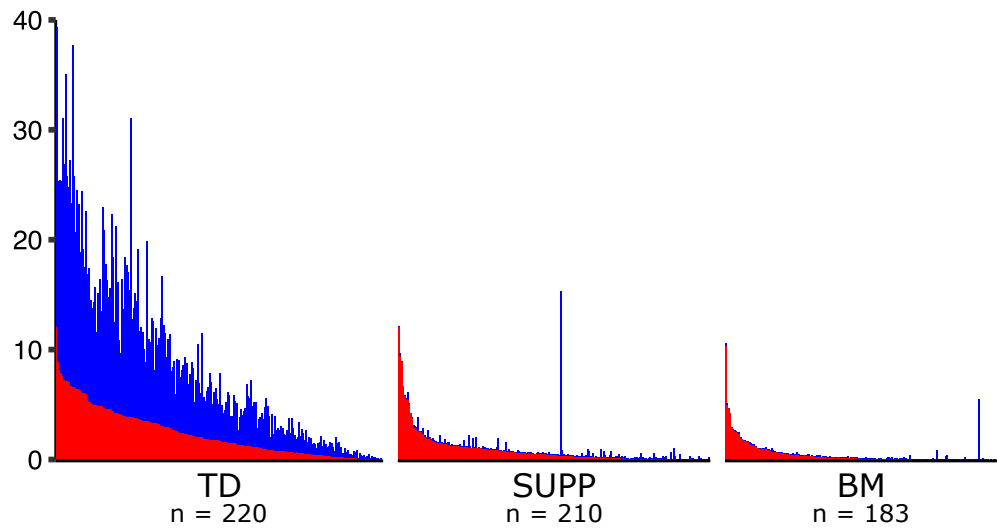


Mean depth of coverage (X)
of *H. parainfluenzae* &
OadG gene



Mean depth of coverage (X)
of *H. parainfluenzae* M1C142-1
(GCA_014931375.1) &
OadG gene

