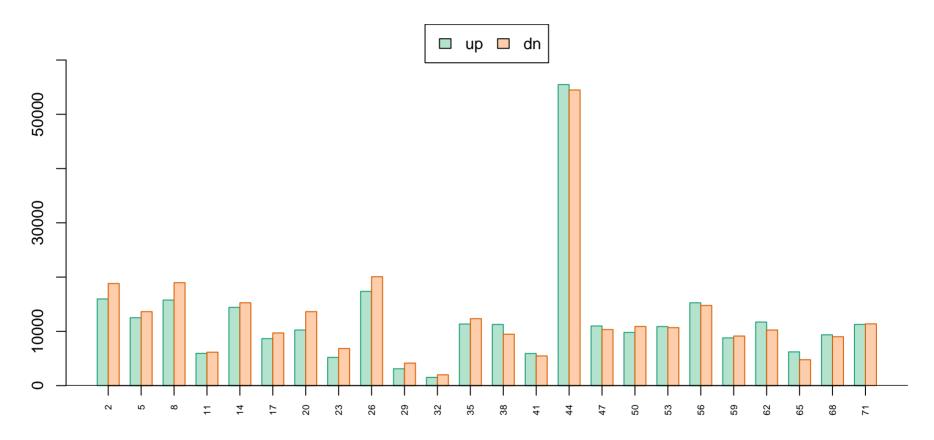
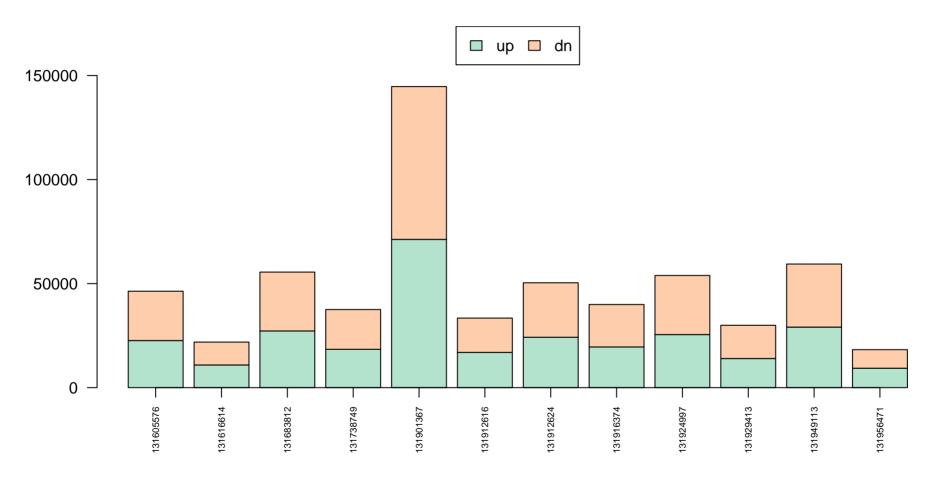
## Primer matches (# reads)



## Primer matches by patient (# reads)



## Primers identified on forward reads

## **Primers identified on reverse reads**

