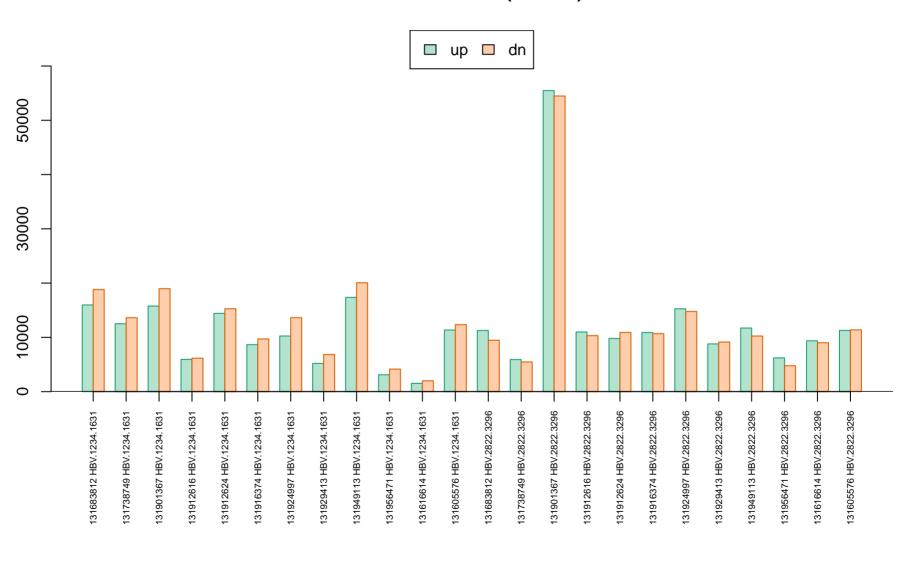
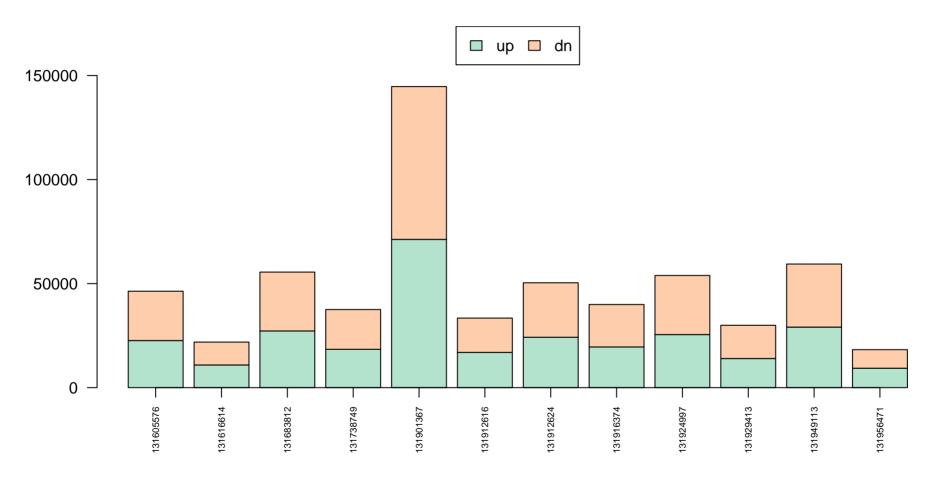
## Primer matches (# reads)



## Primer matches by patient (# reads)



## **Primers identified on forward reads**

## Primers identified on reverse reads

