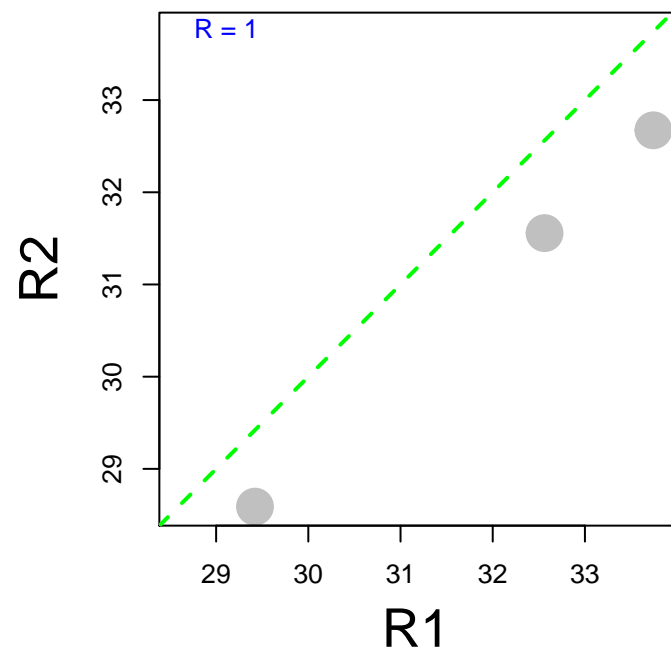
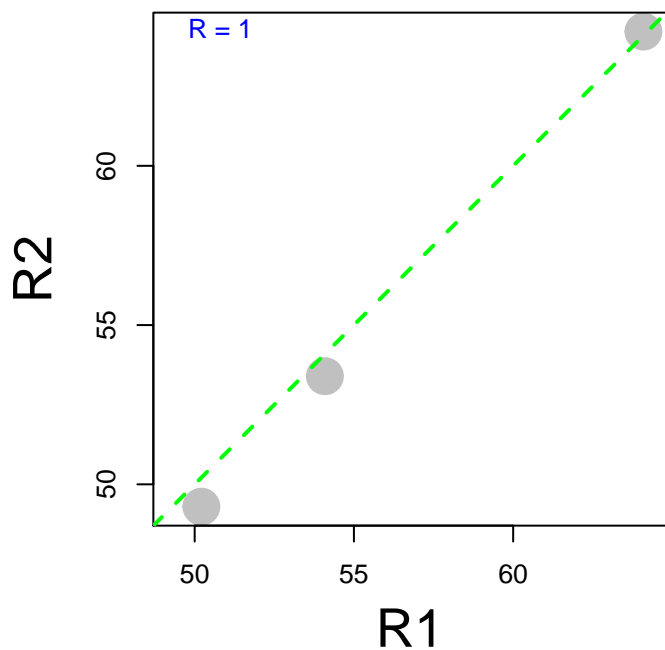


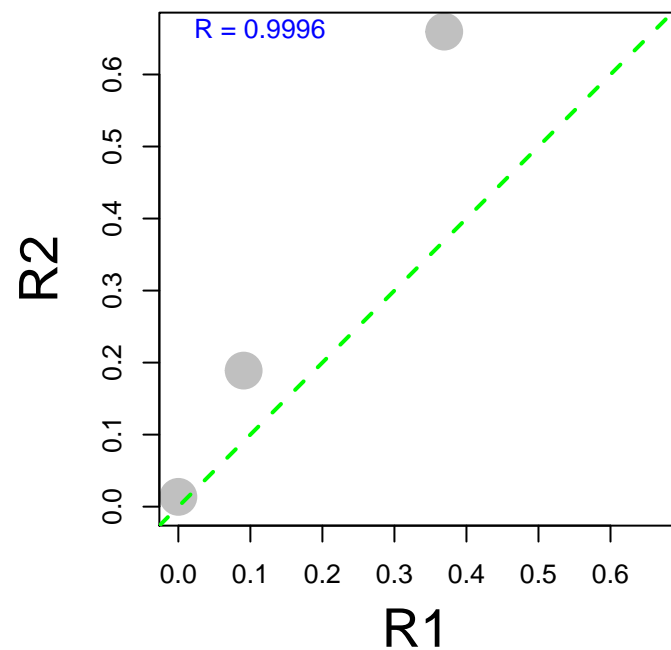
**Average score of sequence quality**



**Overall percentage of GC bases**



**Average percentage of Ns**



**Percentage of duplicated reads**

