

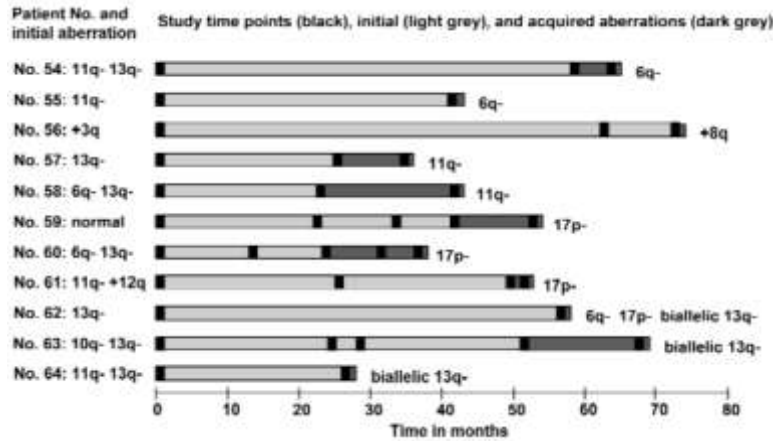
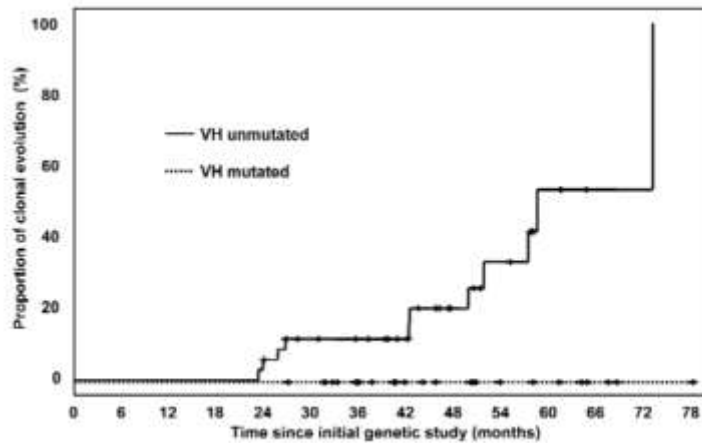
A**B**

Figure 1 a-b. Clonal evolution of genomic aberrations in CLL. **(A)** The 11 patients in whom clonal evolution was observed are illustrated with the genomic aberrations present at initial analysis (time point 0) on the left and the acquired aberrations over time on the right. Time points of genetic analyses are indicated by black boxes. The duration of the presence of the initial and the acquired aberrations are indicated in light grey and dark grey, respectively. **(B)** Risk of clonal evolution depending on the VH mutation status. The proportion of patients with clonal evolution is illustrated over time according to the presence of mutated or unmutated VH genes. Clonal evolution was only observed among cases with unmutated VH. The median time to clonal evolution among cases with unmutated VH was 73 months (95% confidence interval: 58.4 months – Inf).