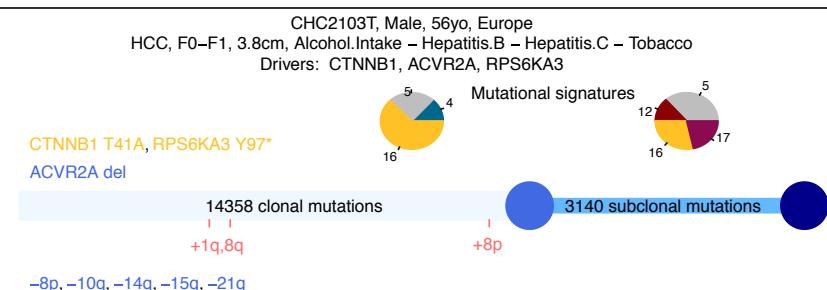
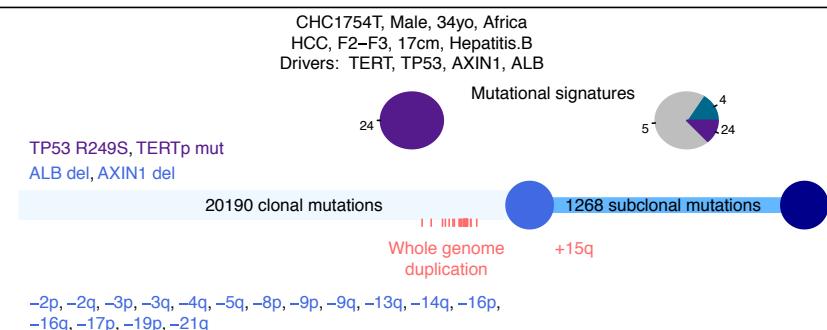
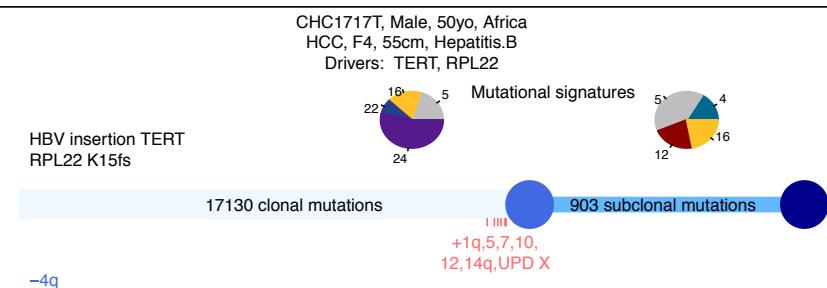
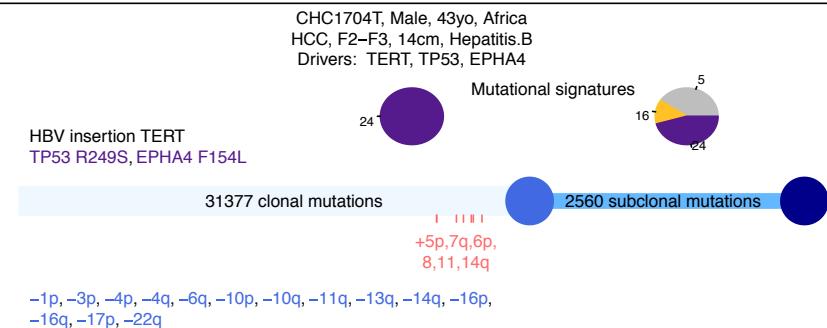
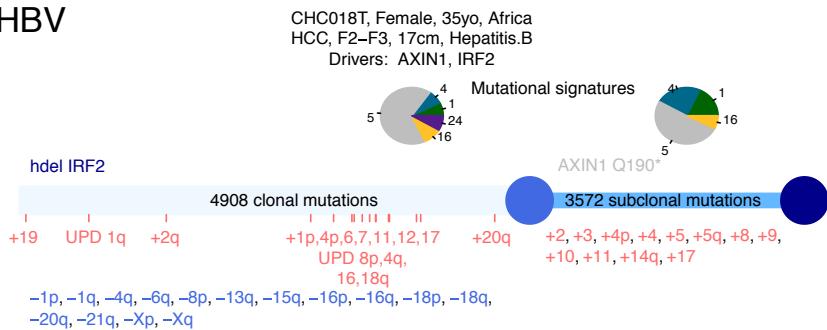
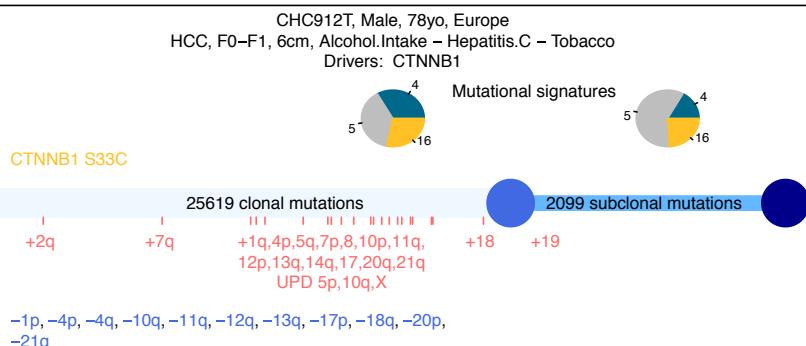
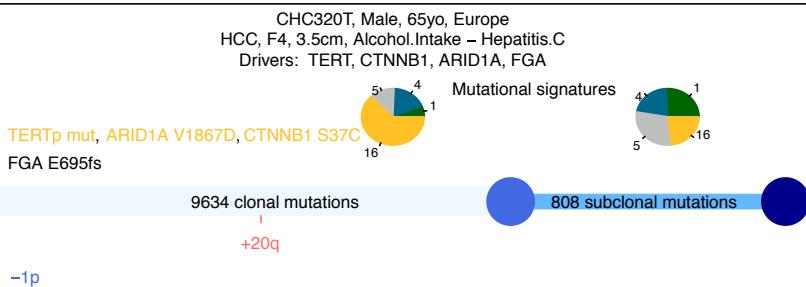
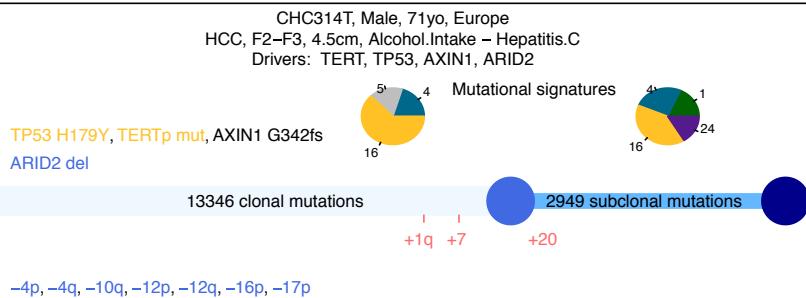
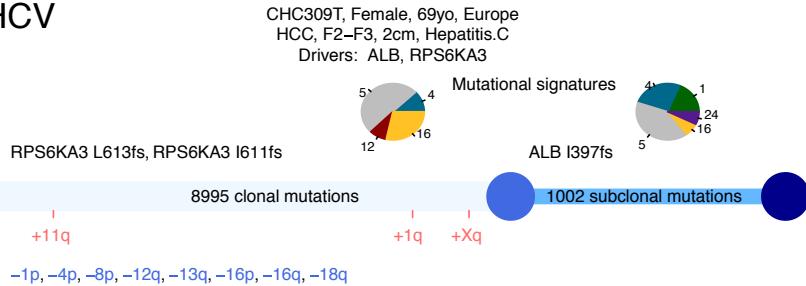


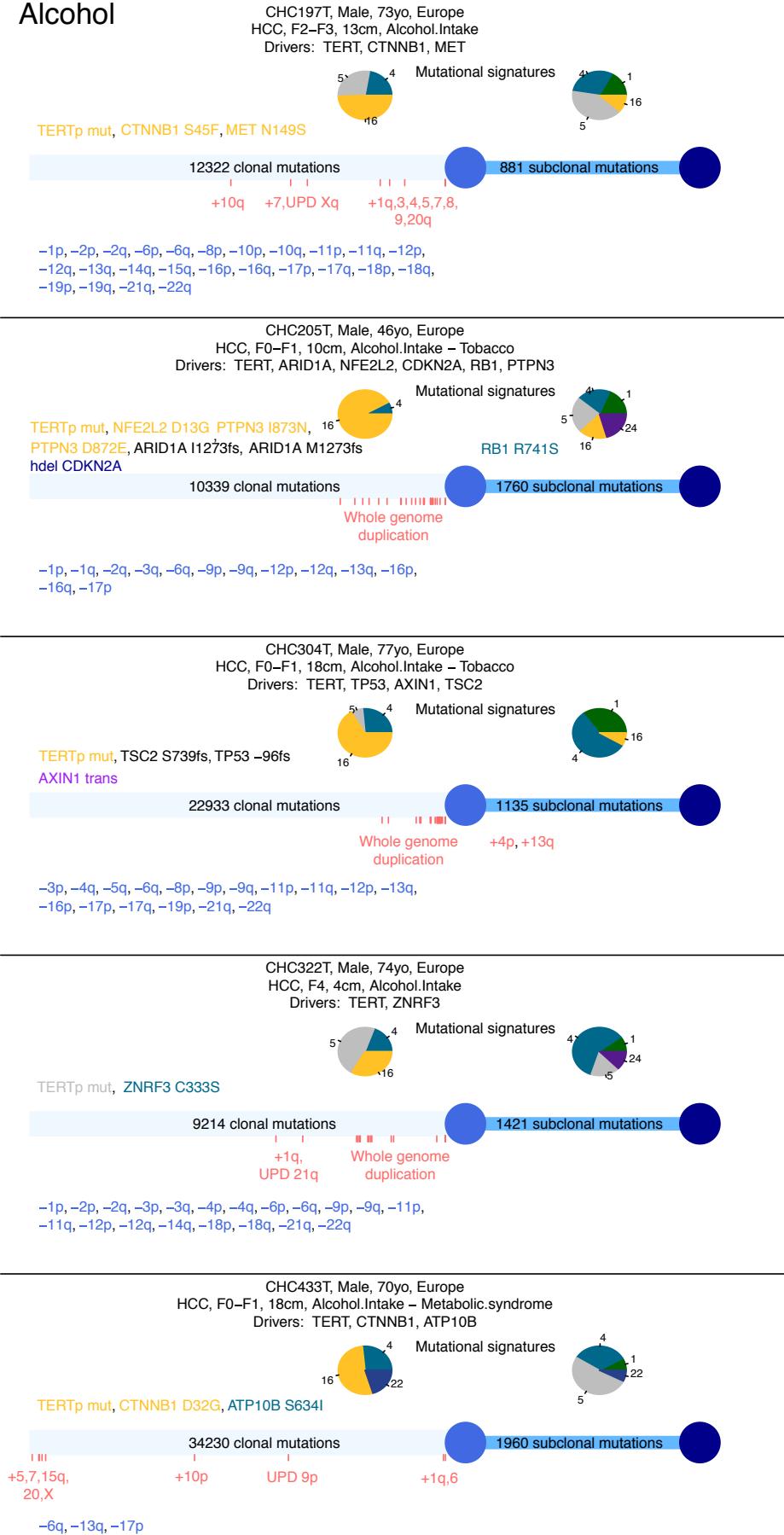
## HBV

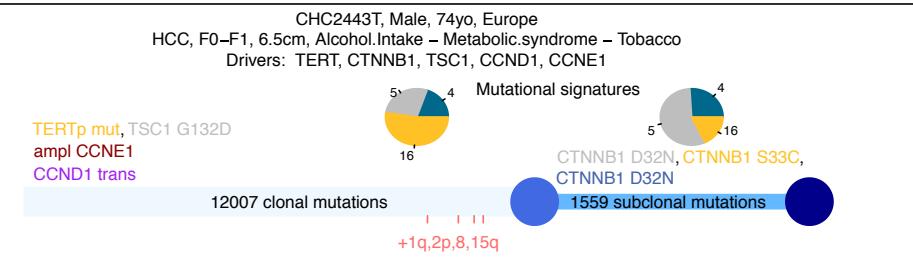
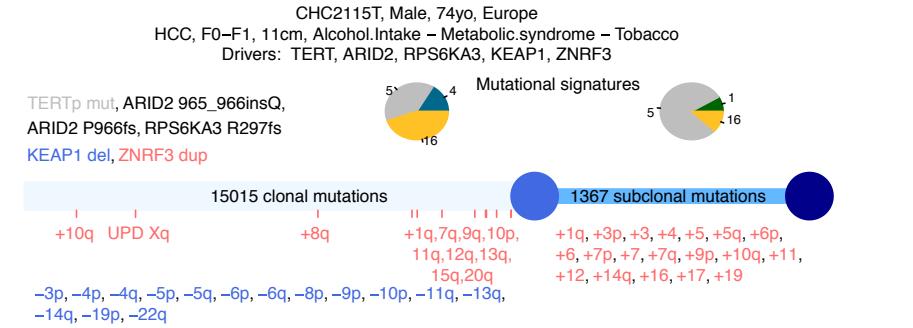
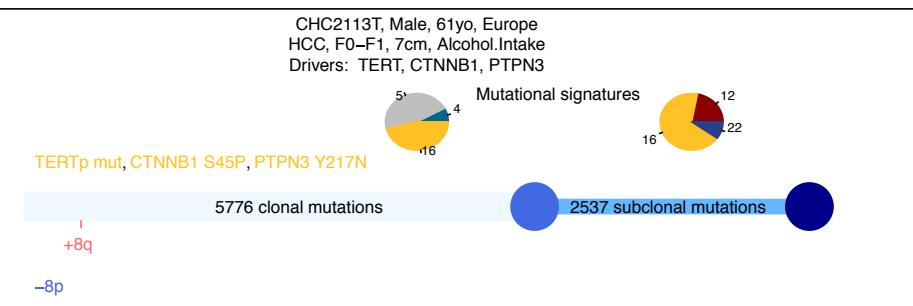
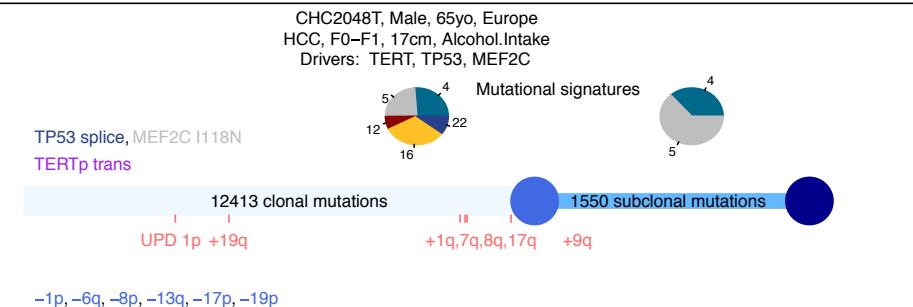
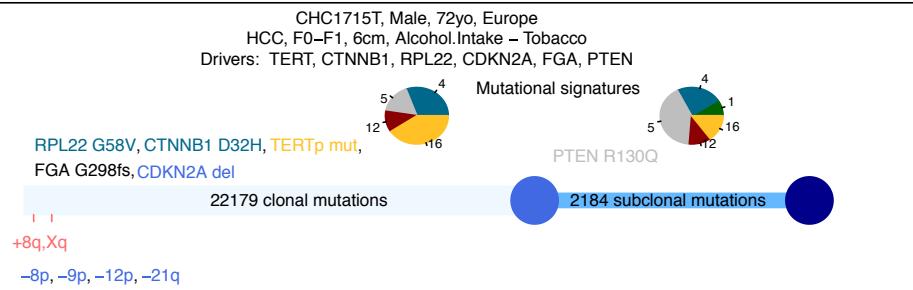


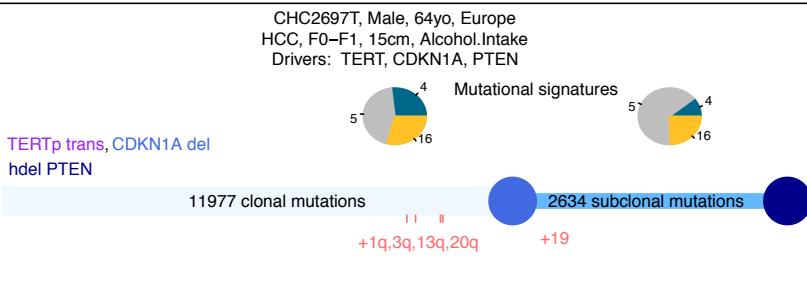
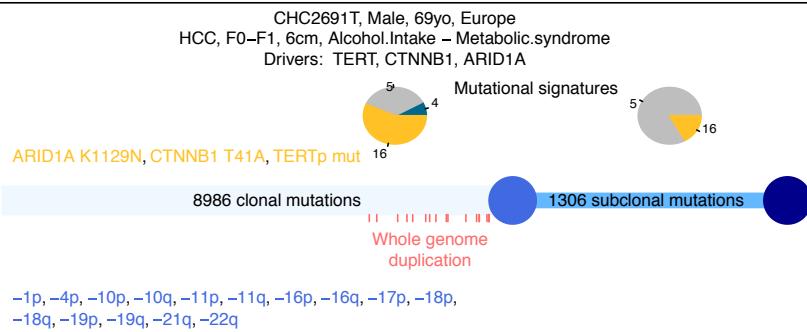
## HCV



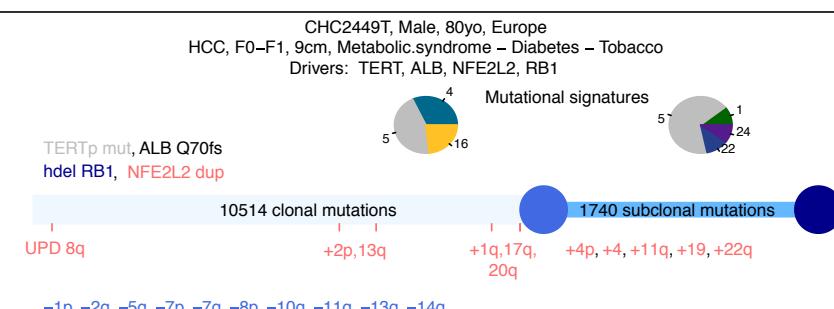
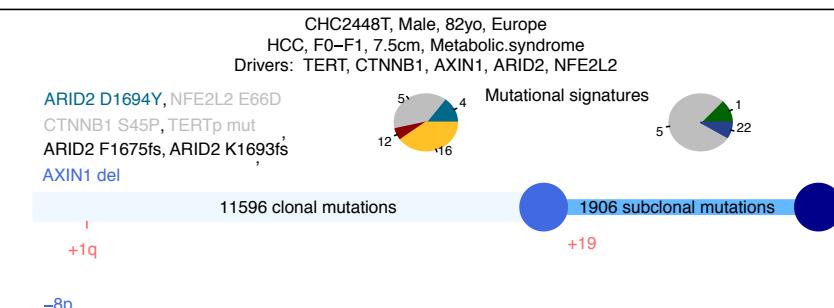
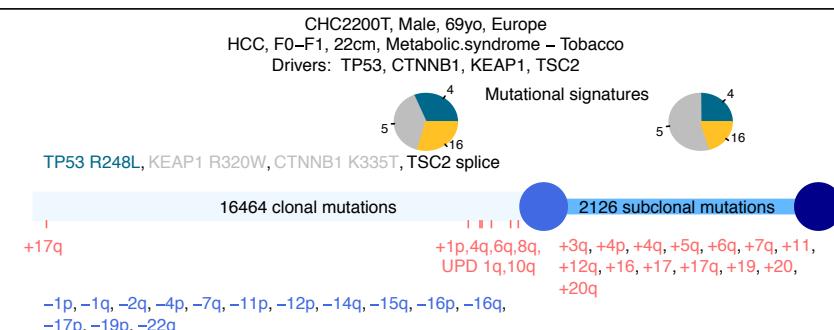
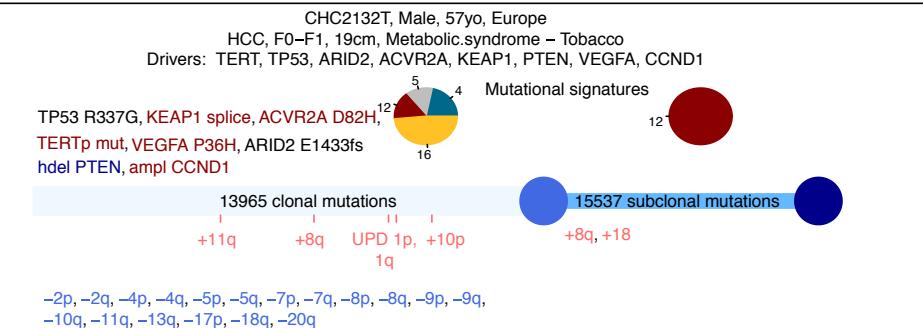
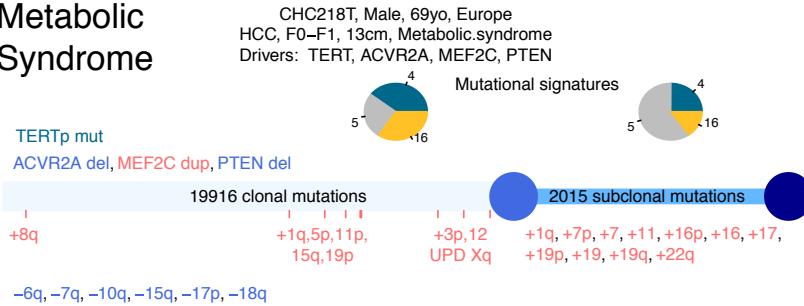
## Alcohol

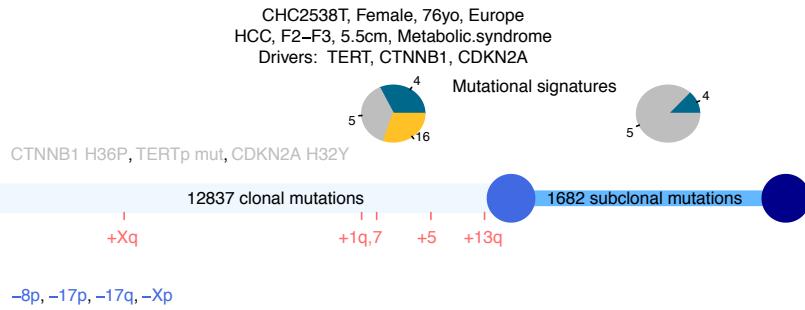




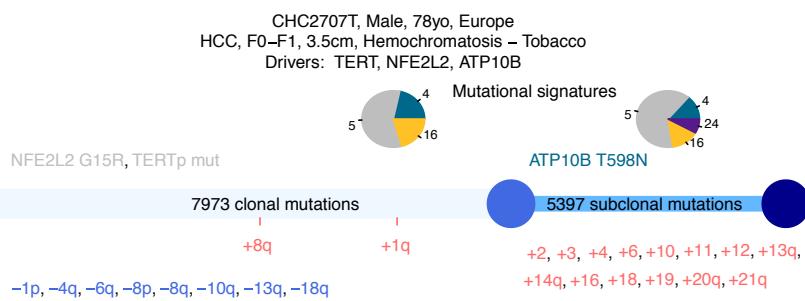


## Metabolic Syndrome

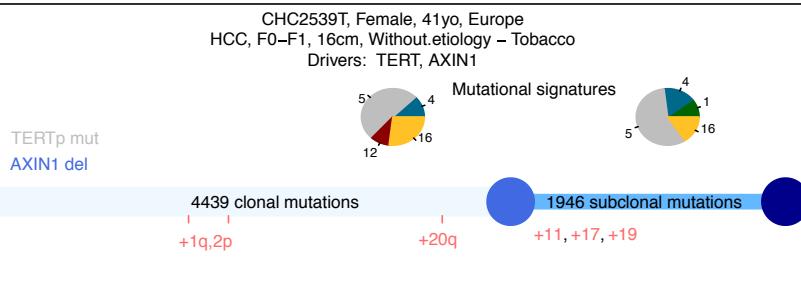
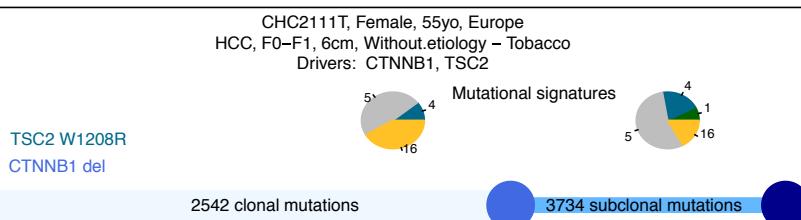
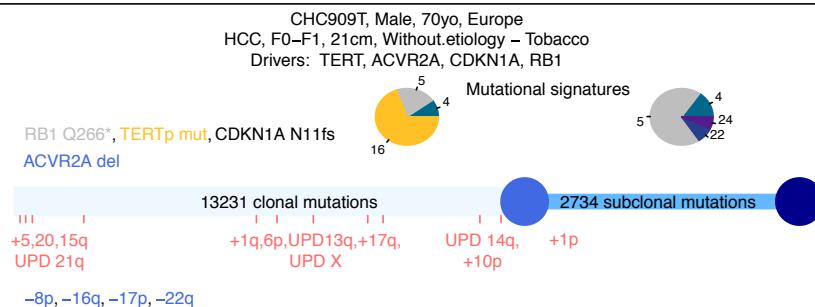
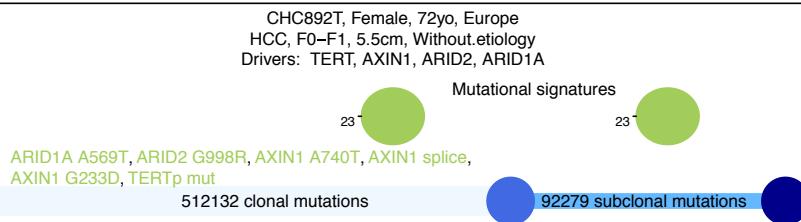
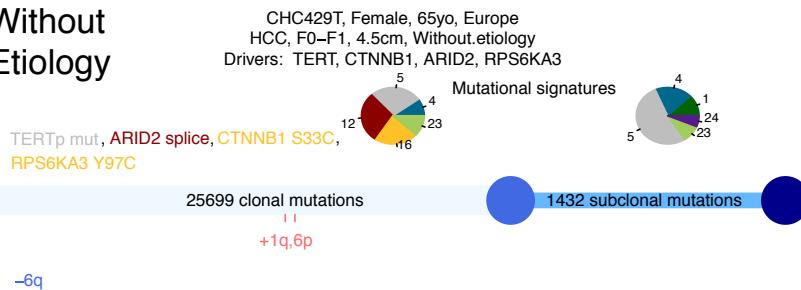




## Hemochromatosis



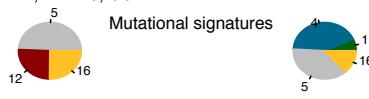
## Without Etiology



## Fibrolamellar Carcinomas

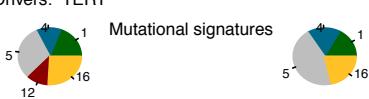
CHC255T, Female, 39yo, Europe  
FLC, F0–F1, 14cm, Without.etiology – Tobacco  
Drivers: ALB, ZNRF3, CCNE1

ampl CCNE1  
ALB del, ZNRF3 del



CHC334T, Female, 24yo, Europe  
FLC, F0–F1, 9cm, Without.etiology  
Drivers: TERT

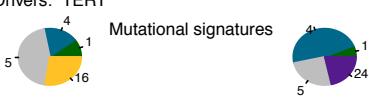
TERTp mut



CHC442T, Female, 27yo, Europe  
FLC, F0–F1, 10cm, Without.etiology – Tobacco  
Drivers: TERT

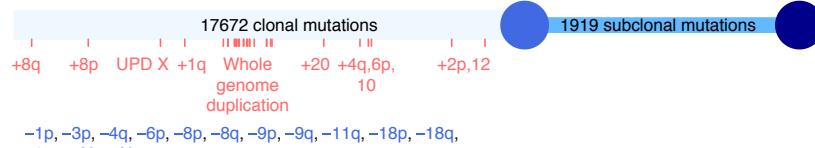
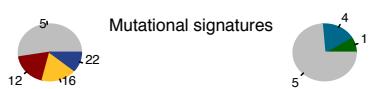
TERTp mut

PRKACA-DNAJB1 fusion



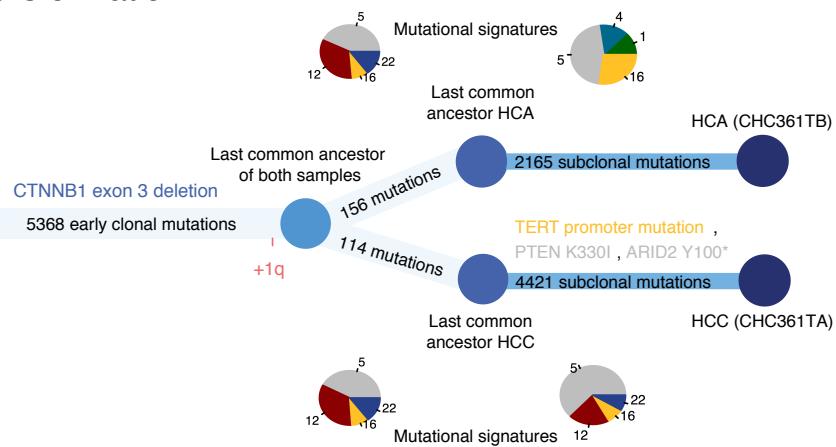
\* This tumor was too contaminated to distinguish duplicated from non-duplicated somatic mutations.

CHC906T, Female, 50yo, Europe  
FLC, F0–F1, 8cm, Without.etiology

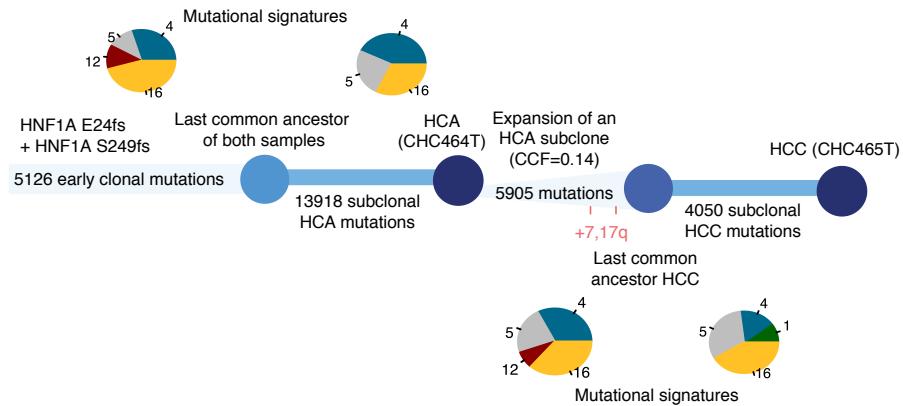


## HCA → HCC transformation

CHC361TA/B, Female, 67yo, Europe  
HCA, F0–F1, 6cm, Metabolic.syndrome – Diabetes  
Drivers: TERT, CTNNB1, ARID2, PTEN



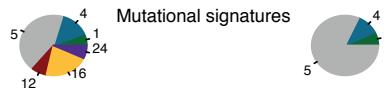
CHC464T, Female, 42yo, Europe  
HCA, F0–F1, 12cm, Without.etiology – Tobacco  
Drivers: HNF1A



## HCA

CHC605T, Female, 47yo, Europe  
HCA, F0–F1, 6cm, Metabolic.syndrome – Diabetes

Drivers:

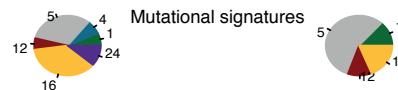


2079 clonal mutations

3781 subclonal mutations

CHC2432T, Female, 43yo, Europe  
HCA, F0–F1, 1.5cm, Without. etiology

Drivers:

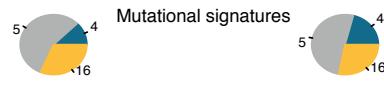


2299 clonal mutations

780 subclonal mutations

CHC2446T, Female, 45yo, Europe  
HCA, F0–F1, 5cm, Without. etiology – Tobacco

Drivers:



2050 clonal mutations

3349 subclonal mutations

-6p

## Legend

### Tumor progression tree

- Last common ancestor
- Final tumor

Clonal alterations

Subclonal alterations

### Mutational signatures

- 1
- 4
- 5
- 6
- 12
- 16
- 17
- 22
- 23
- 24

### Copy-number and structural variations (SVs):

Deletions, Gains, Translocations