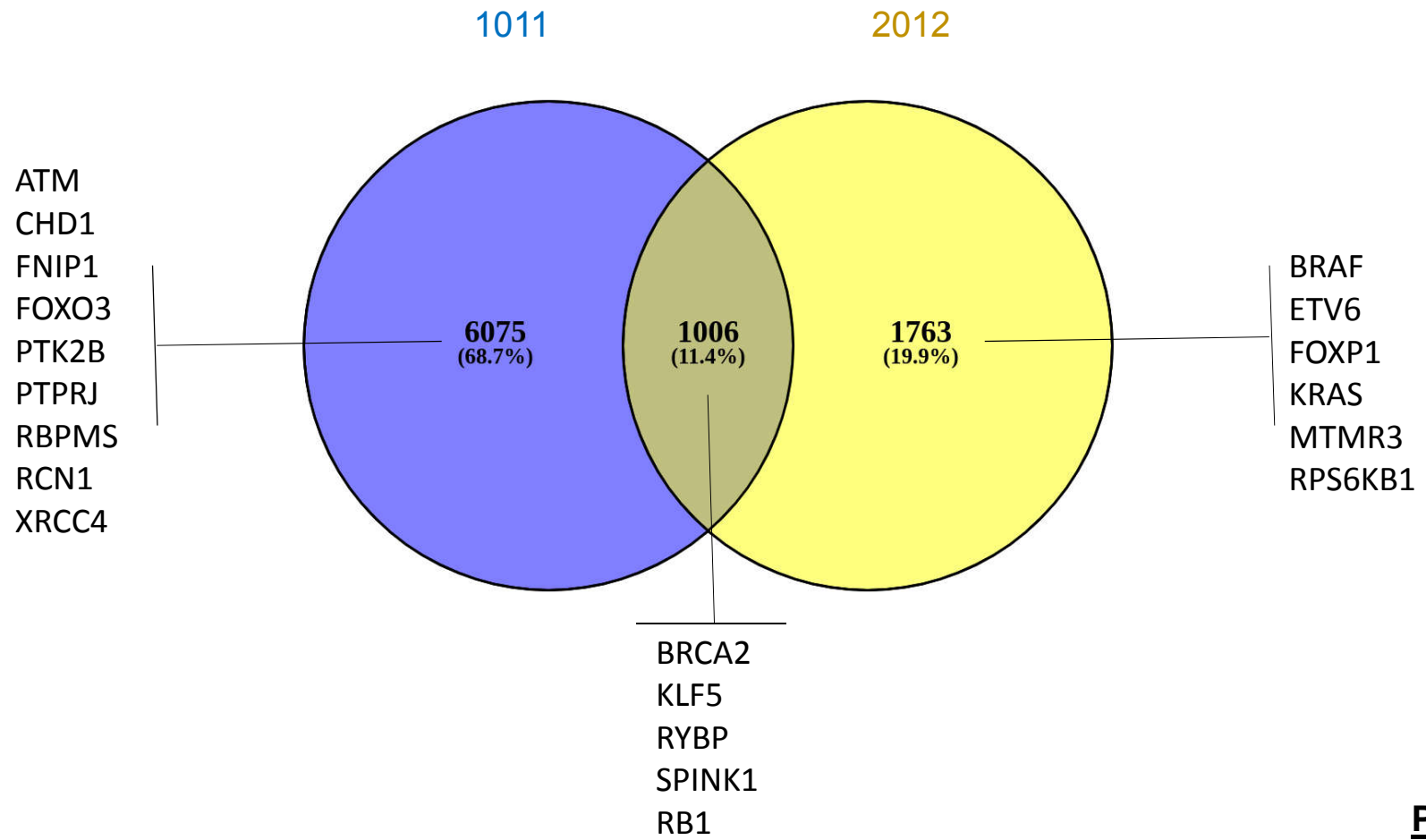
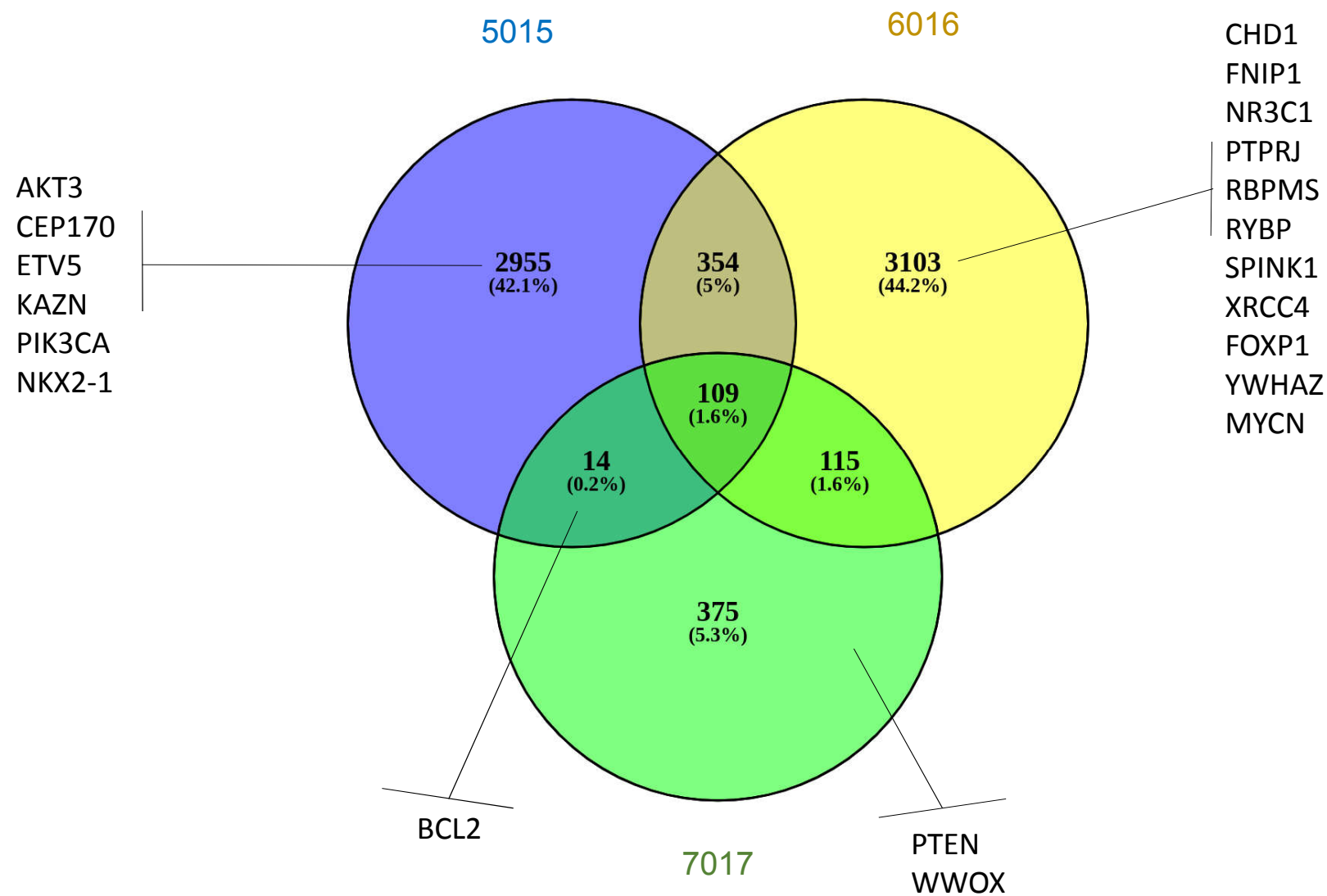


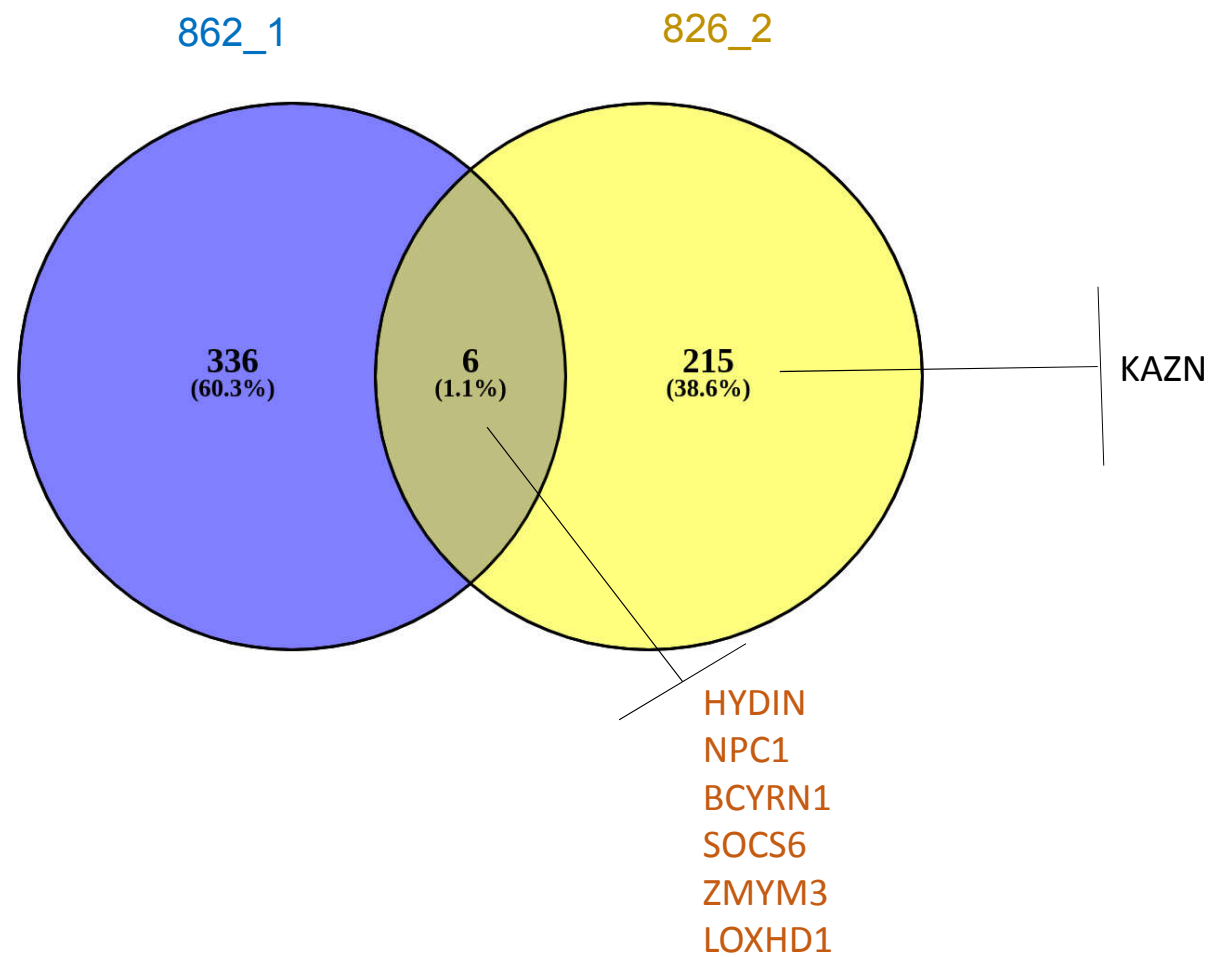
Supplementary Figures 2: Venn diagram representing CTC-shared SNVs/indels and CNA per patient. Genes highlight in the CNA section were also found in Friedlander *et al.* 2019 and genes highlight in the SNVs/indels were associated with prostate cancer in bold and telomere maintenance in italic. We also emphasize in red CTC-shared genes.

Copy Number Alterations (CNA)

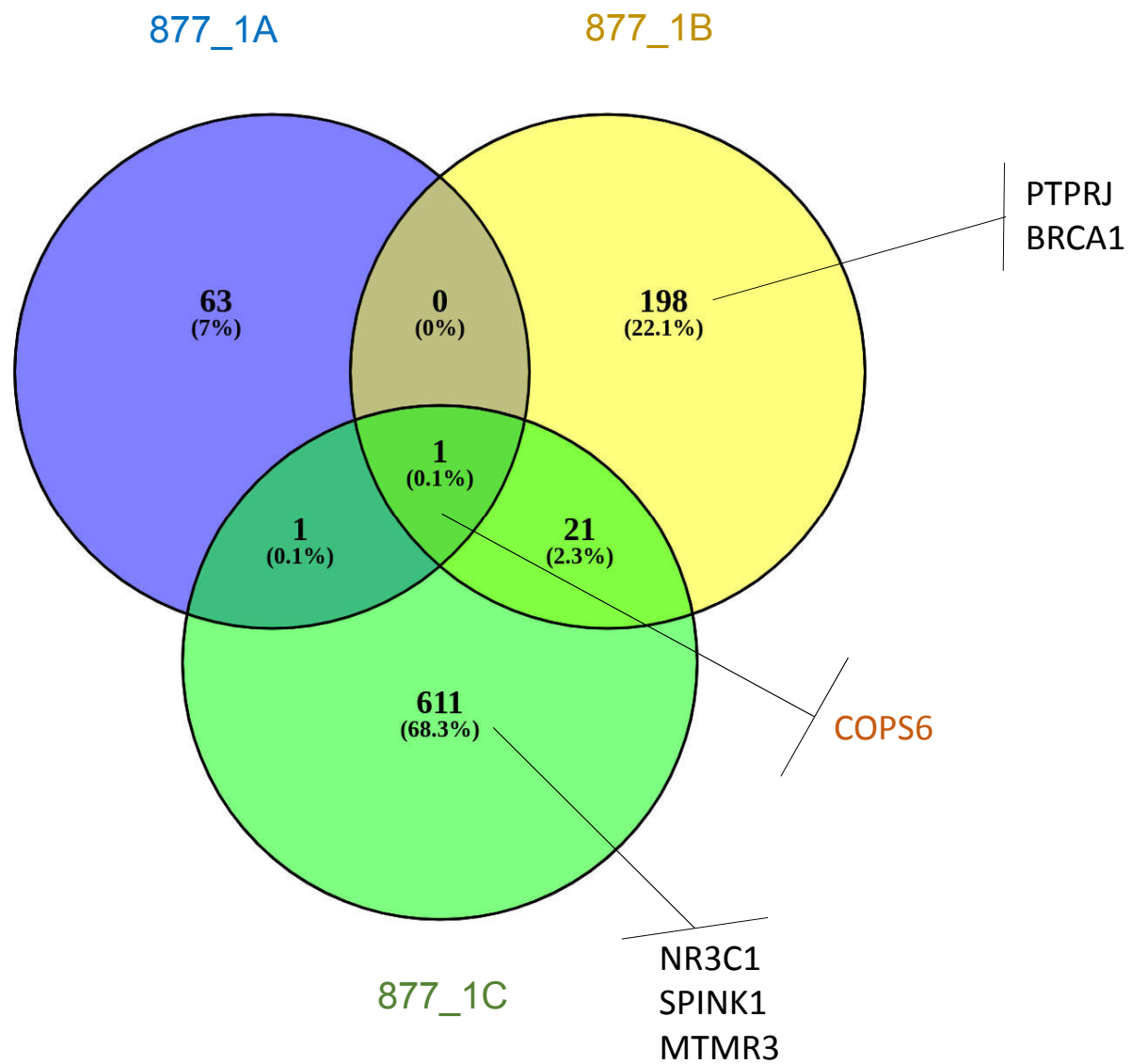




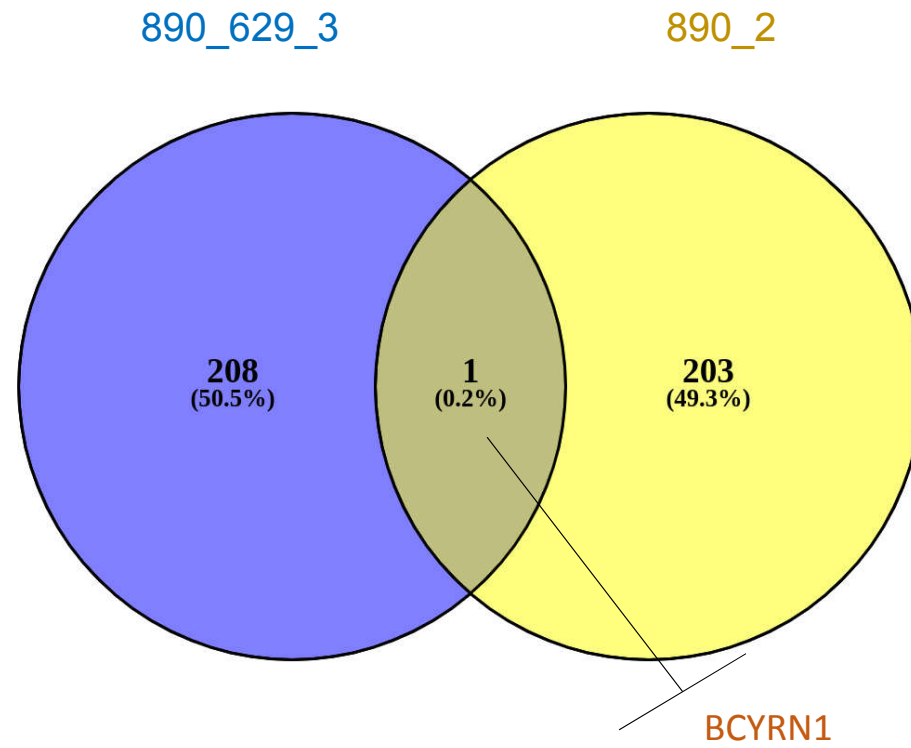
Patient 3



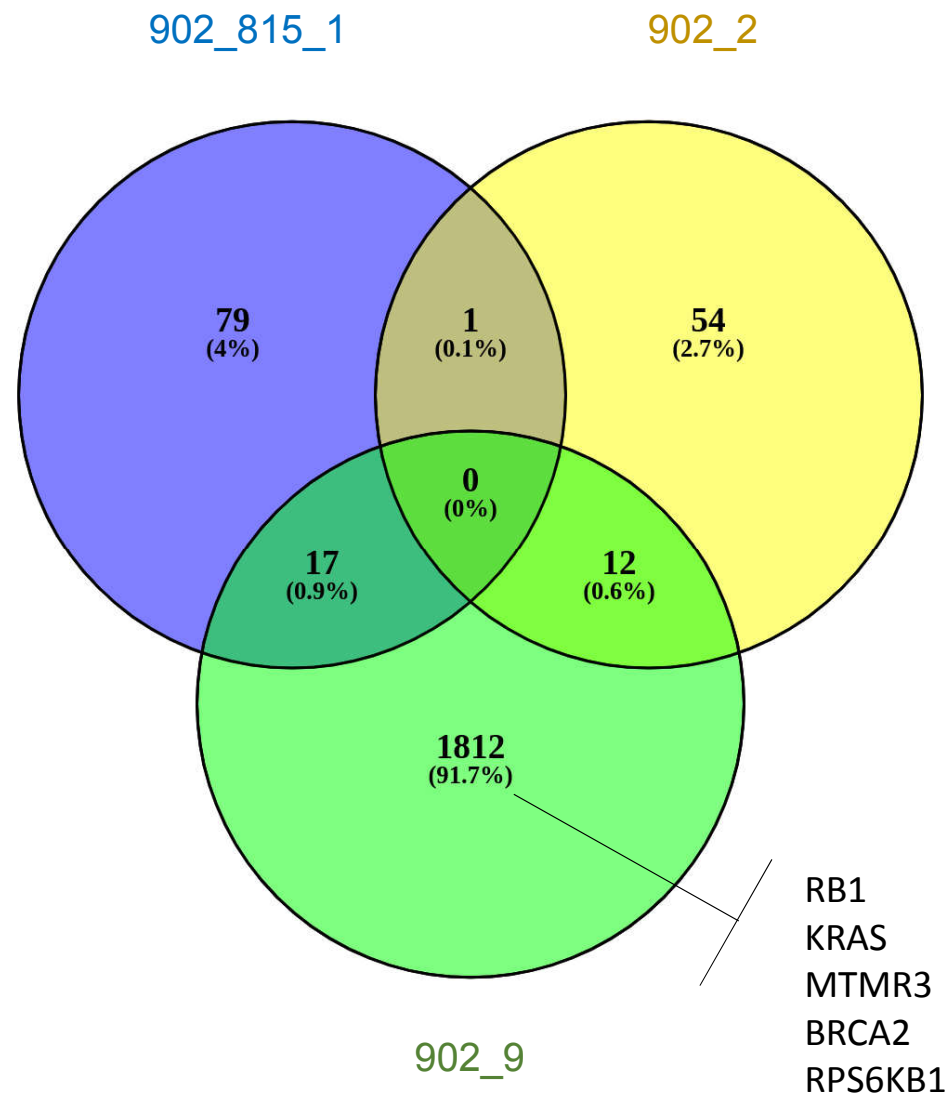
Patient 5



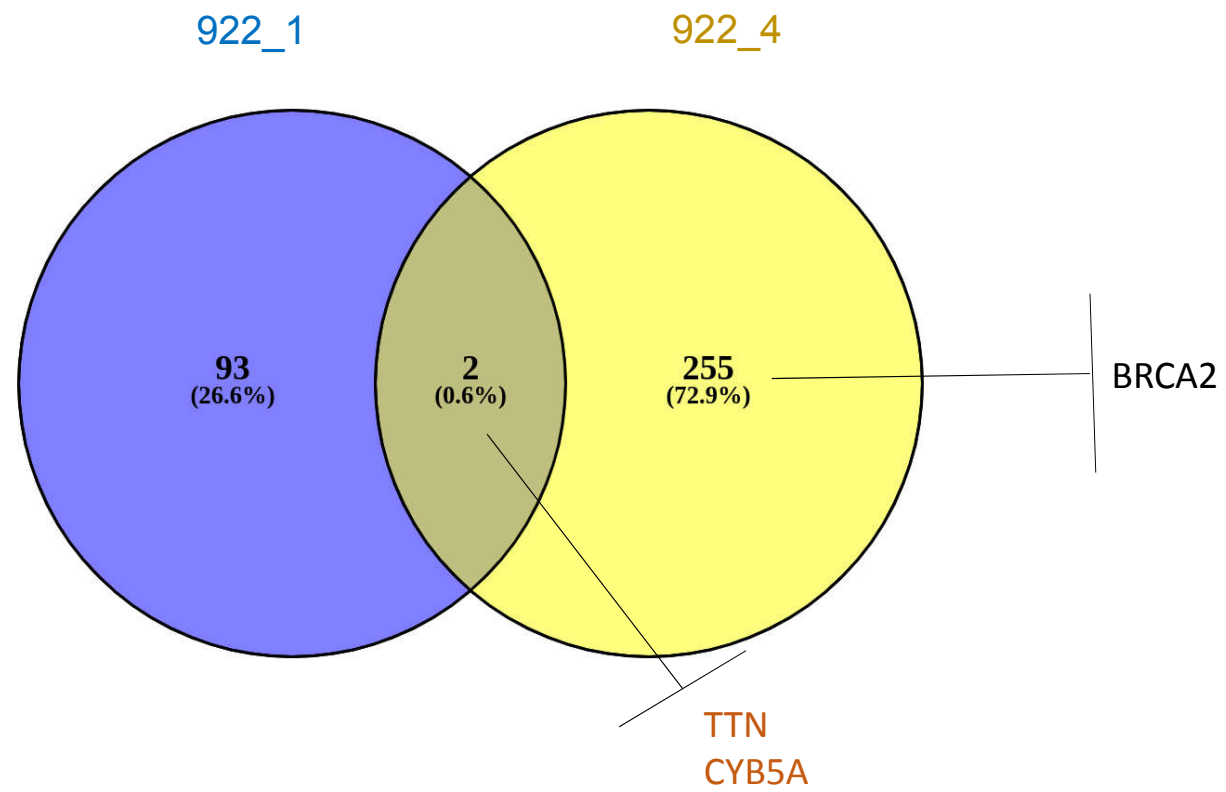
Patient 6



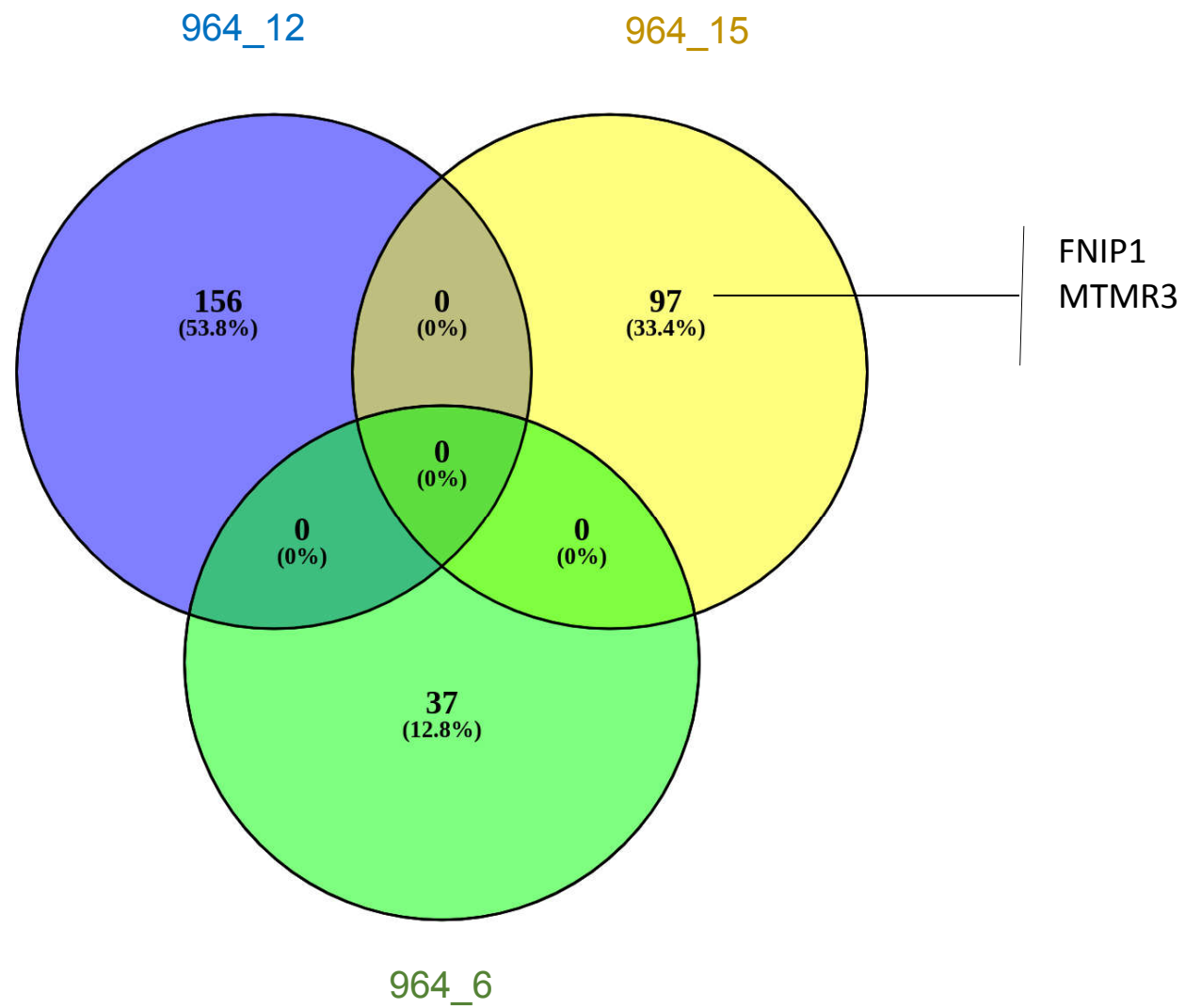
Patient 7



Patient 8

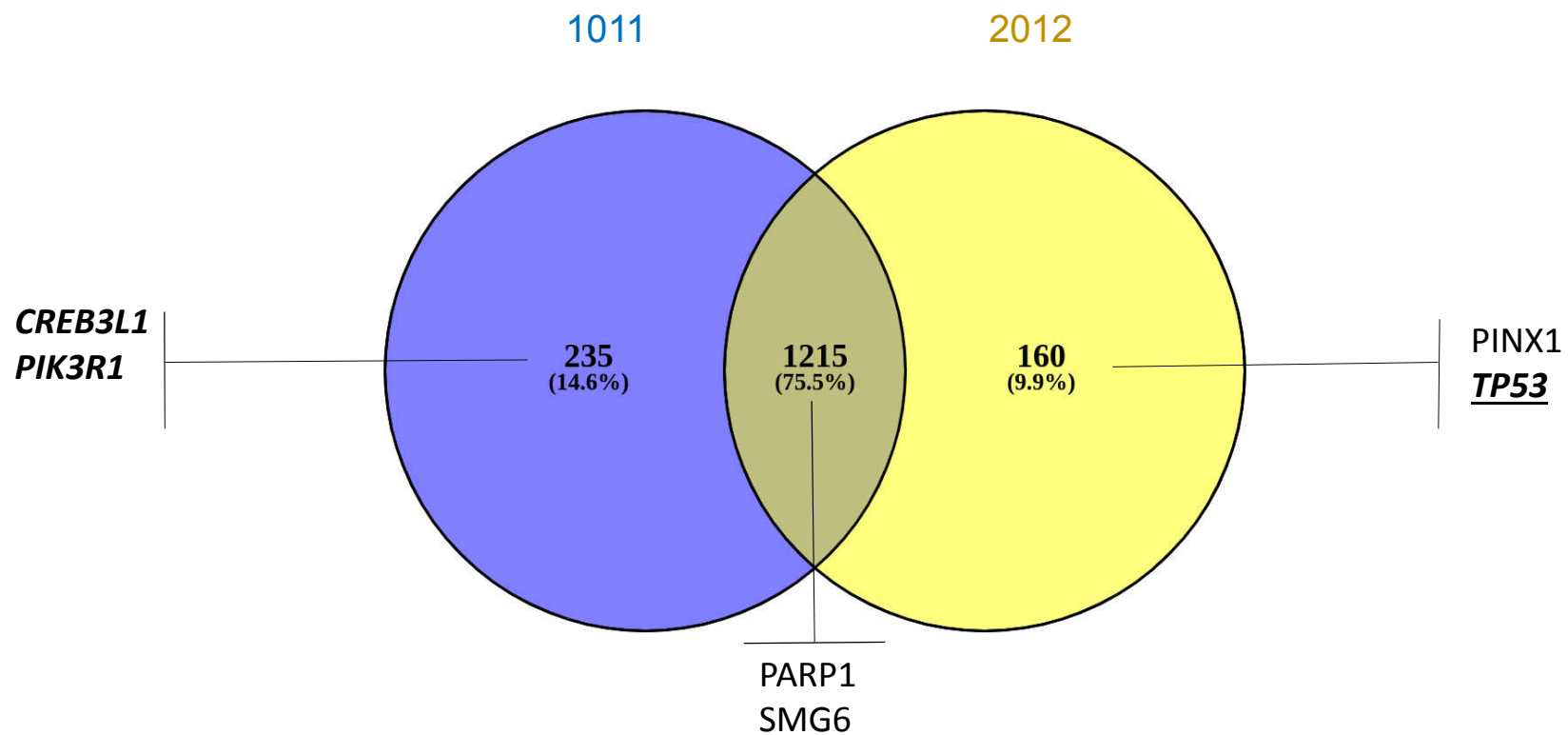


Patient 9

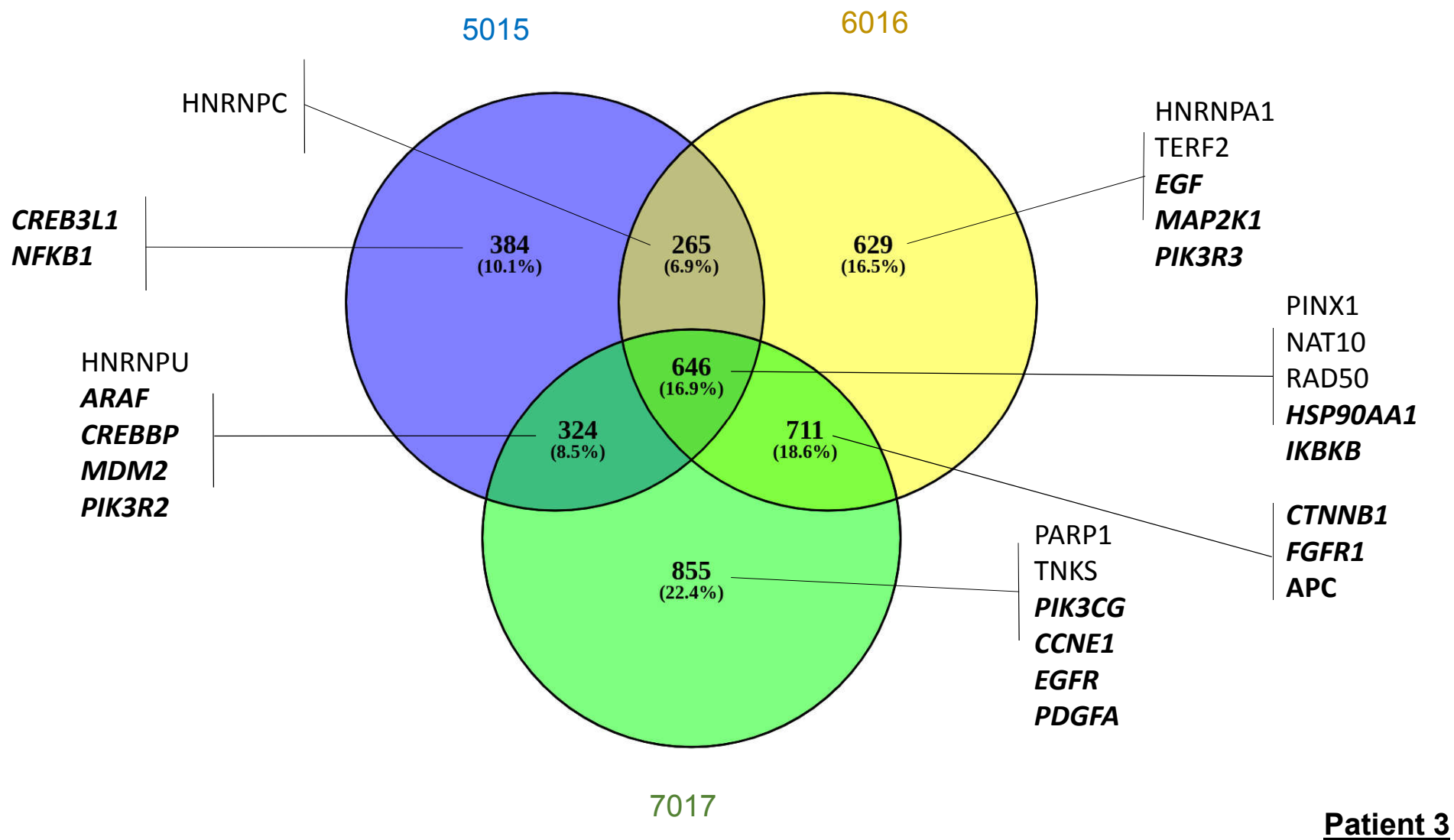


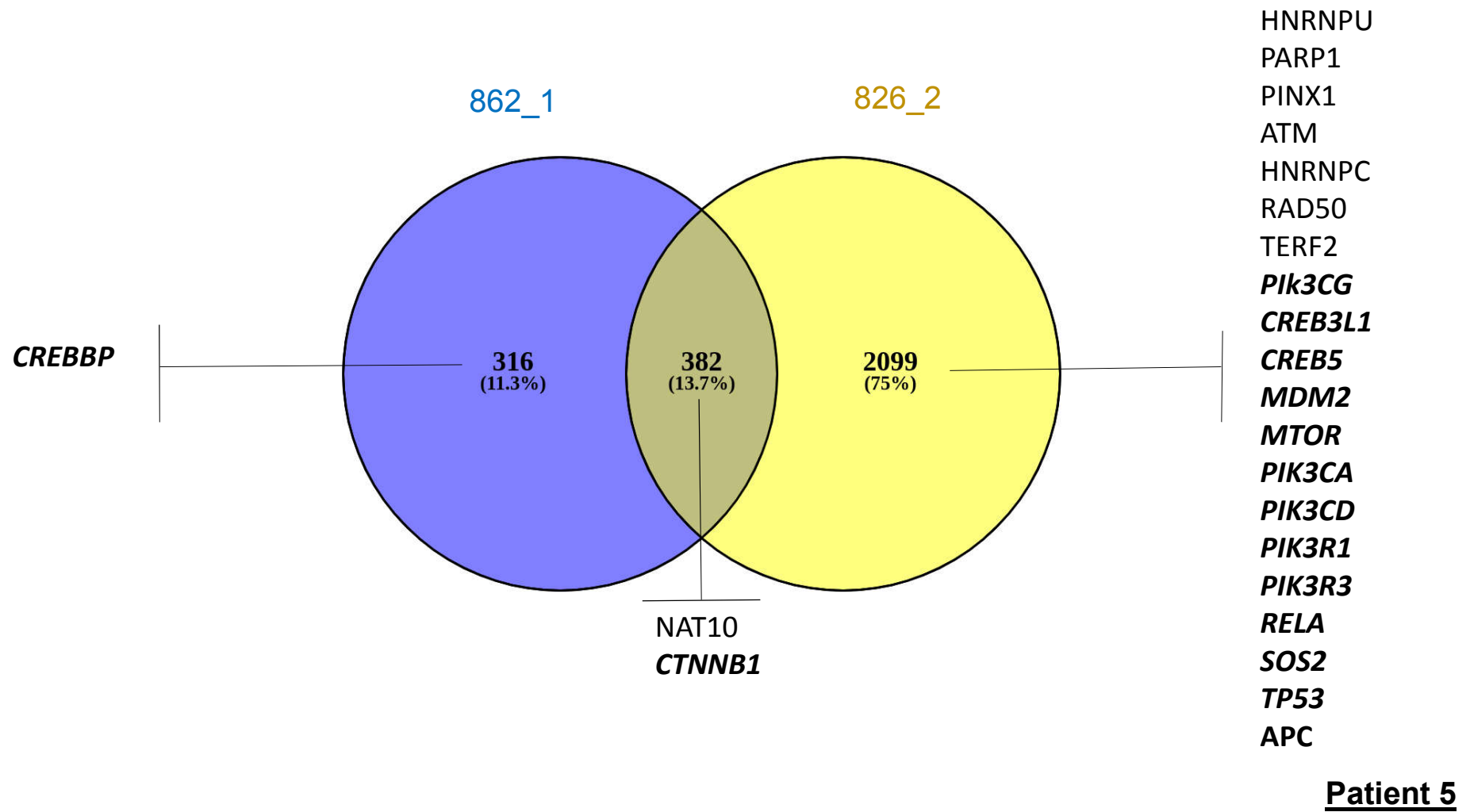
Patient 10

Genetic variations (SNP + INDELs)

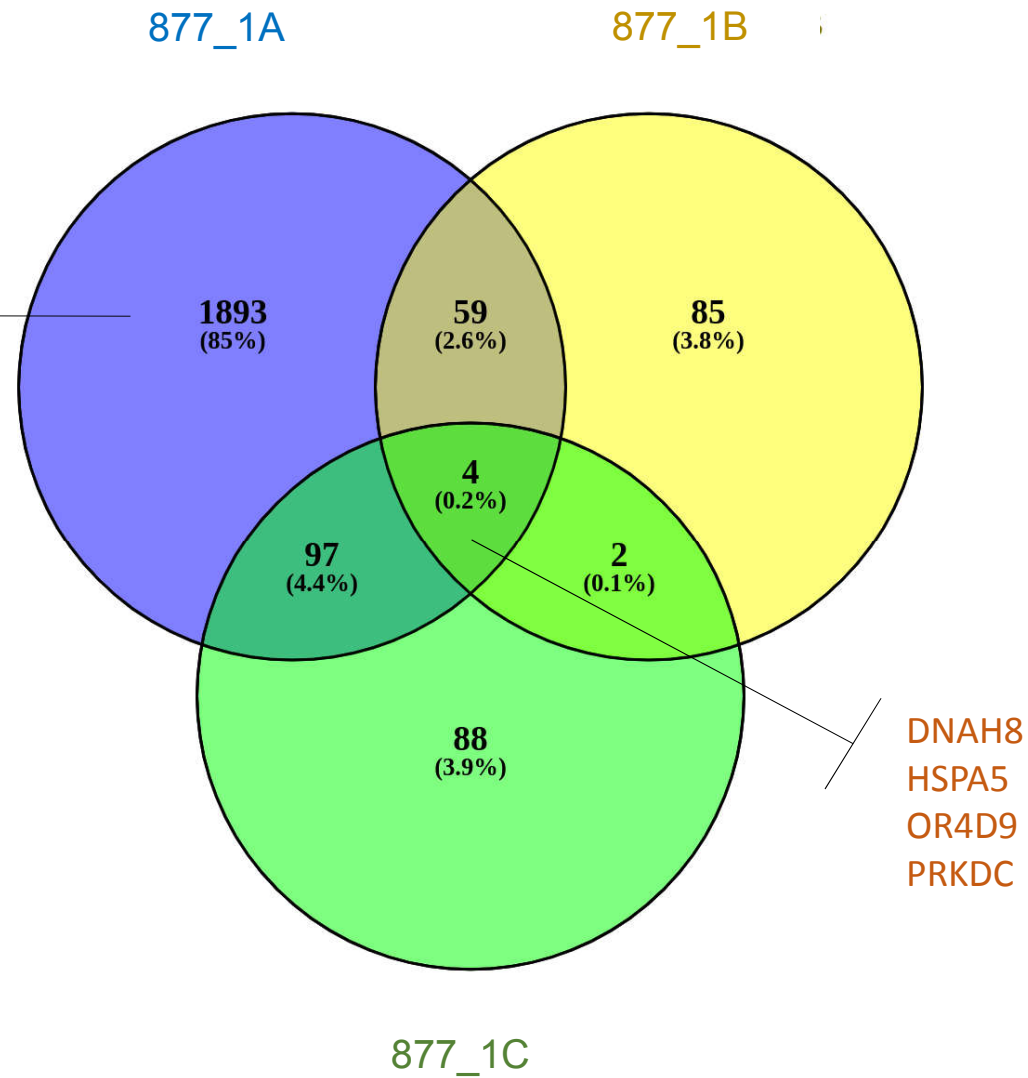


Patient 1

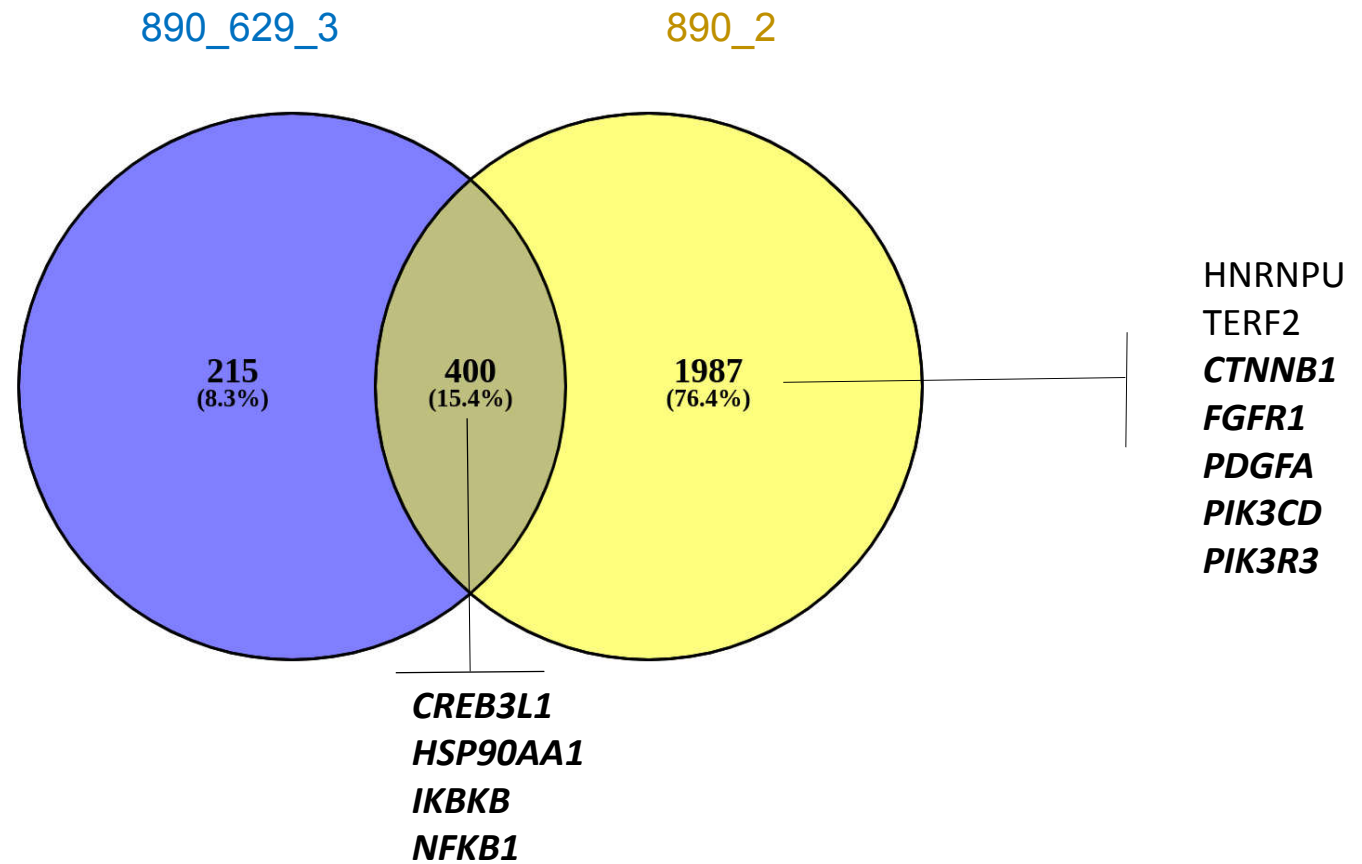




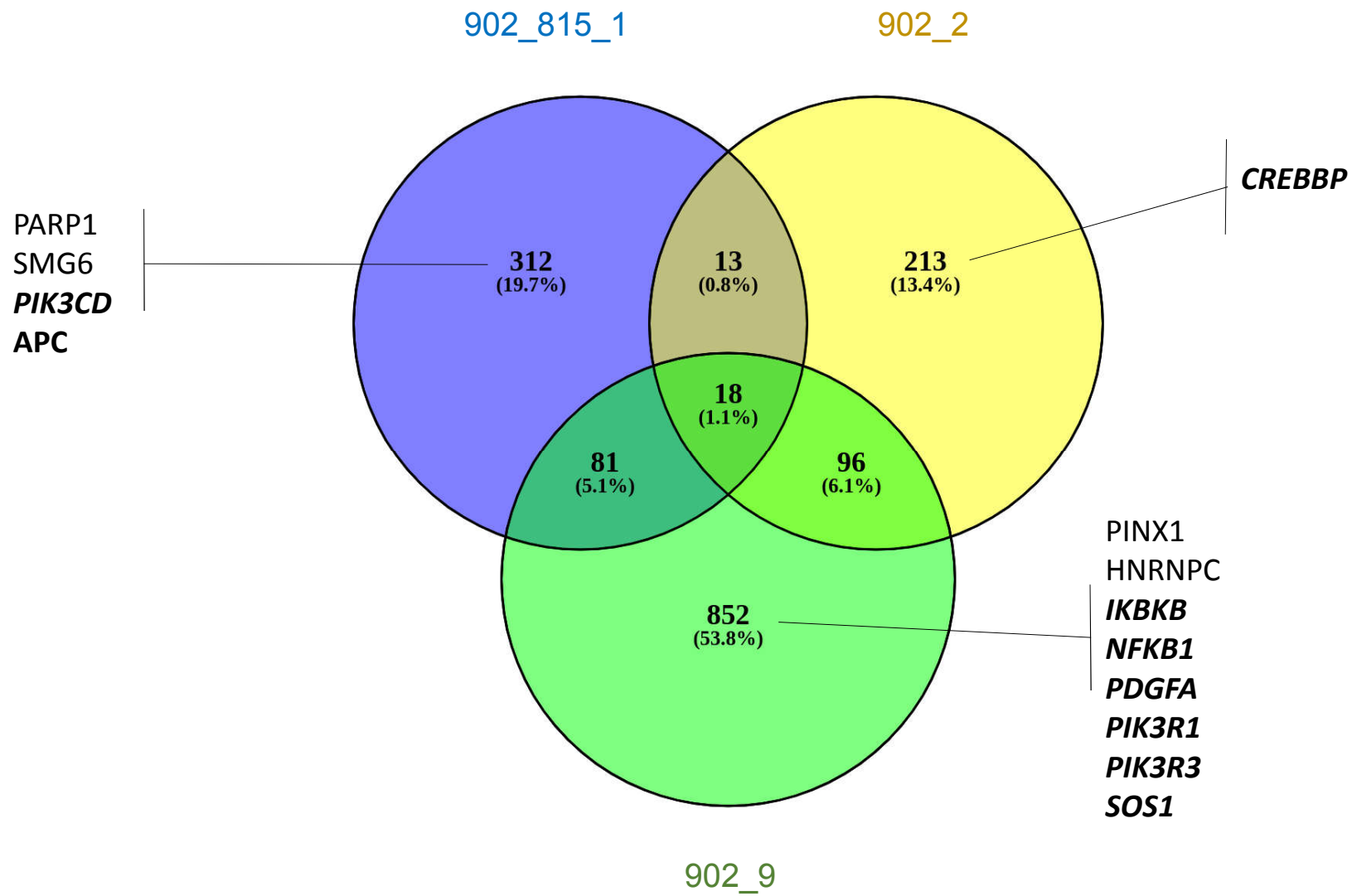
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NAT10
TERF2
PIK3CG
TP53
CREB3L1
CREB3L2
CREBBP
CTNNB1
FGFR1
HSP90A
A1
IKBKB
NFKB1
PDGFA
PIK3R3
SOS1
SOS2
APC



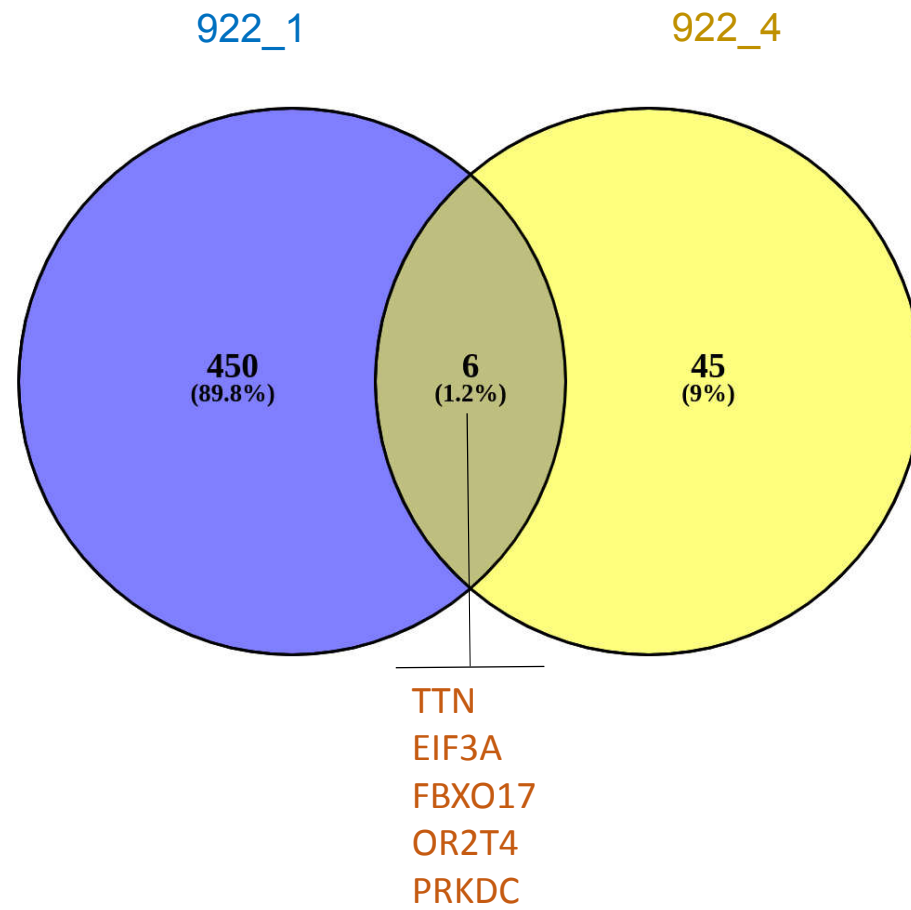
Patient 6



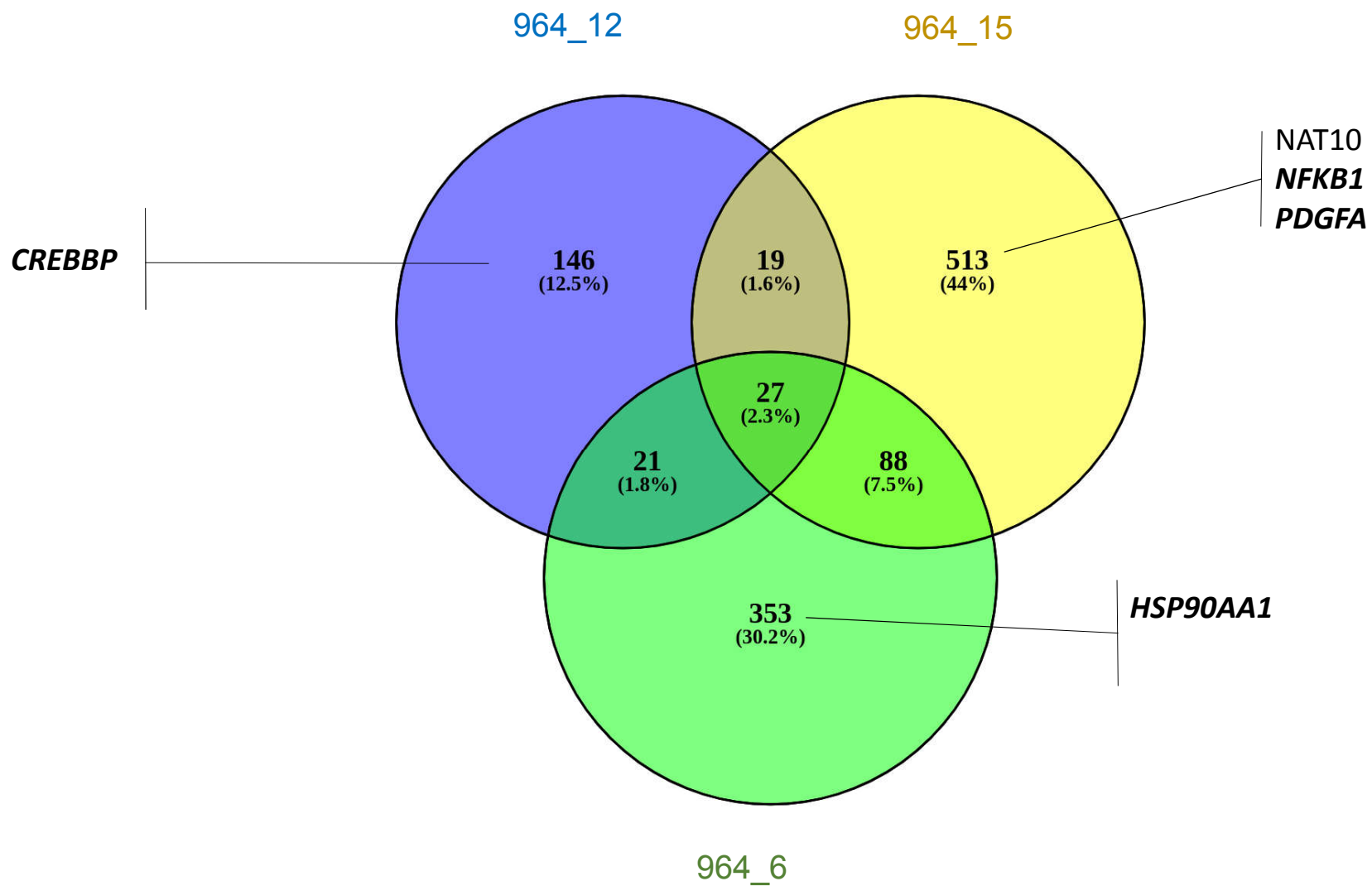
Patient 7



Patient 8



Patient 9



Patient 10