

CDK12 Pathogenic Mutations in African American and White Patients with Prostate Cancer

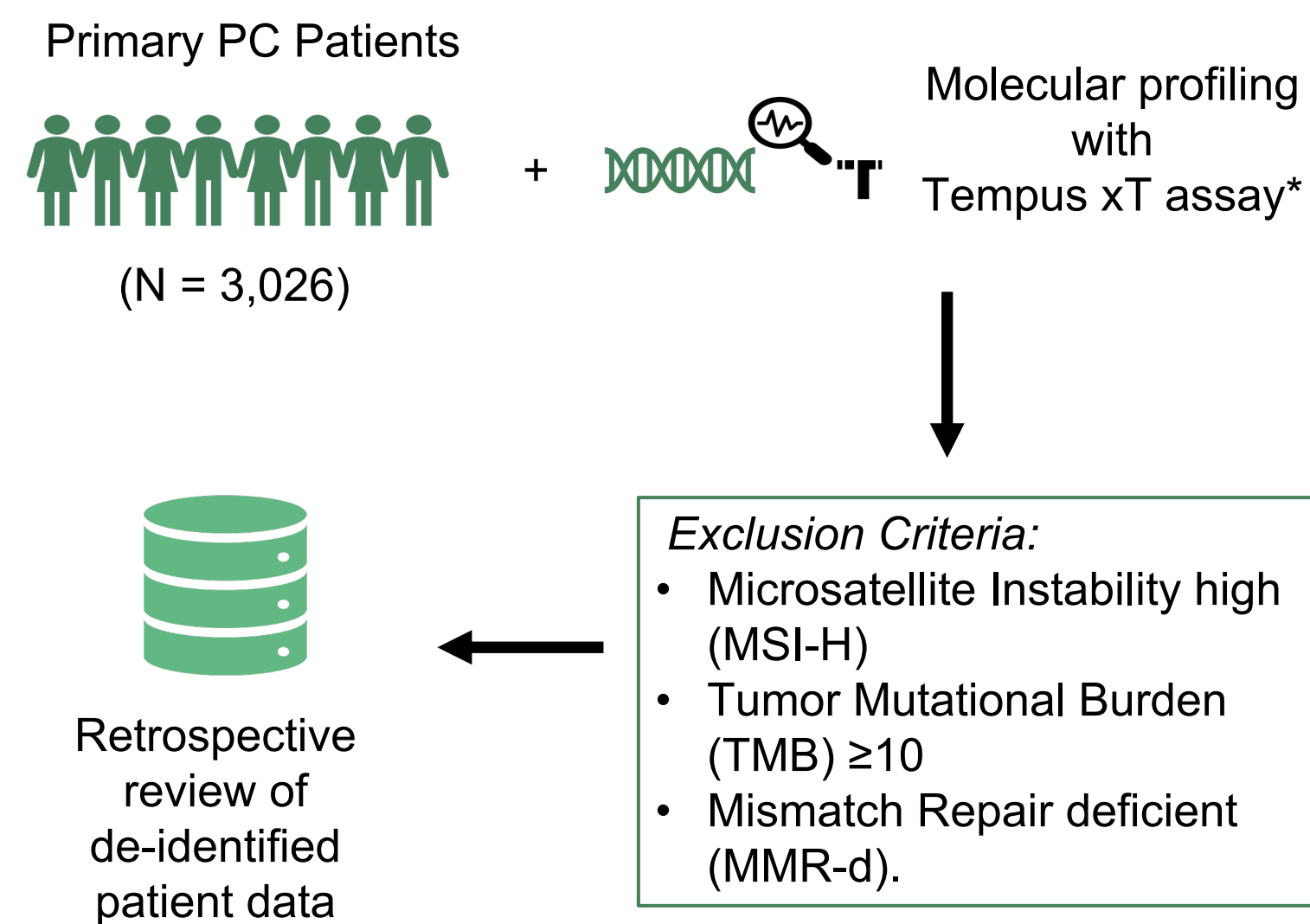
Elisa Ledet, PhD¹, Nicholas Mitsiades, MD, PHD², Ellen Jaeger, MSPH³, Calvin Chao, MD³, A Oliver Sartor, MD¹

¹School of Medicine, Tulane Medical School, New Orleans, LA, ²UC Davis Comprehensive Cancer Center, UC Davis, ³Tempus Labs, Inc.

INTRODUCTION

- African American men are underrepresented in prostate cancer (PC) trials but have a greater probability of developing and dying from PC than other race populations.
- CDK12 mutations in PC are associated with aggressive disease, increased metastasis, and decreased overall survival.
- This study reports data from next-generation sequencing (NGS) of tissue from a real-world dataset of patients with PC to investigate the frequency of pathogenic CDK12 mutations in African American men and White men.

METHODS



Chi-square and Fischer's Exact tests were used where appropriate to compare alterations between AA and White patients

- Frequency and types of pathogenic CDK12 mutations
- Co-mutation in other assessed genes

Corrections for multiple testing were also applied.

*Tempus xT assay - solid tumor, 648 genes, 500x coverage, full transcriptome RNAseq for a subset of patients

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Correspondence: eledet@Tulane.edu

SUMMARY

- African American men with prostate cancer show a **higher frequency of pathogenic CDK12 mutations** than White men in a large real-world population.
- Co-mutations did not vary by race, and the alterations were most likely to be truncating mutations in both races.

RESULTS

Table 1. Cohort Characteristics.

	Overall, N = 3,026 ¹	White, N = 2,444 ¹	Black or African American, N = 582 ¹
Age at Diagnosis	65 (60, 72)	66 (60, 73)	63 (58, 68)
Unknown	439	354	85
Ethnicity			
Not Hispanic or Latino	1,365 (97%)	1,147 (97%)	218 (98%)
Hispanic or Latino	45 (3.2%)	41 (3.5%)	4 (1.8%)
Unknown	1,616	1,256	360
Stage			
Stage 4	1,511 (92%)	1,207 (92%)	304 (94%)
Stage 3	88 (5.4%)	73 (5.6%)	15 (4.6%)
Stage 2	38 (2.3%)	32 (2.4%)	6 (1.8%)
Stage 1	3 (0.2%)	3 (0.2%)	0 (0%)
Unknown	1,386	1,129	257
Advanced stage	1,921 (63%)	1,541 (63%)	380 (65%)
Tissue site			
Prostate	1,935 (64%)	1,558 (64%)	377 (65%)
Other site	1,091 (36%)	886 (36%)	205 (35%)

¹ Median (IQR); n (%)

² Wilcoxon rank sum test; Pearson's Chi-squared test; Fisher's exact test

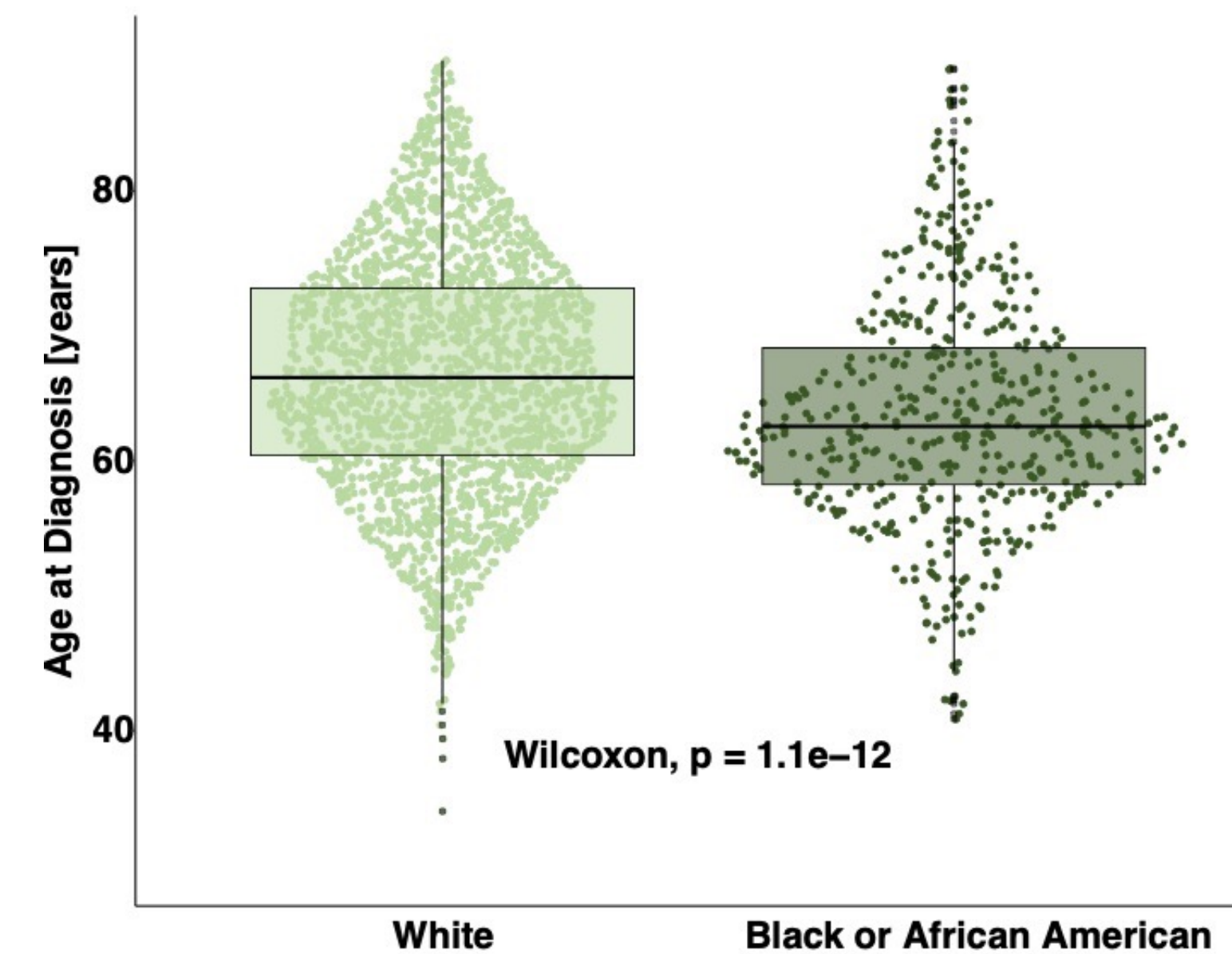


Figure 1. Age at Diagnosis.

582 African American and 2,444 White patients were included. African American were younger at diagnosis than Whites (median 63 vs 66 years; $p < 0.001$).

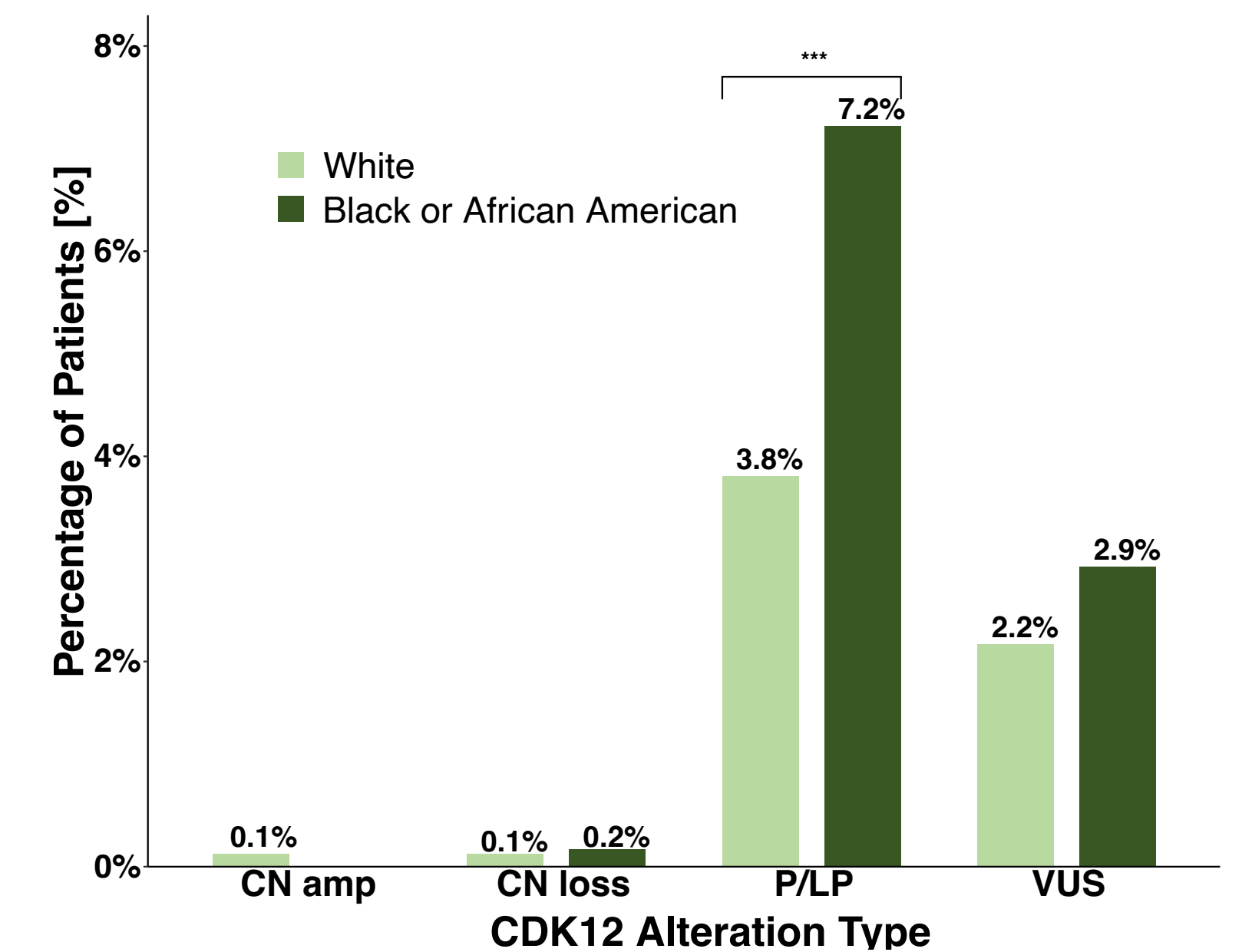


Figure 2. CDK12 Alteration Types.

Tissue samples from African American men were significantly more likely than those from Whites to express a pathogenic CDK12 mutation (42 [7.2%] vs 93 [3.8%], $p < 0.001$).

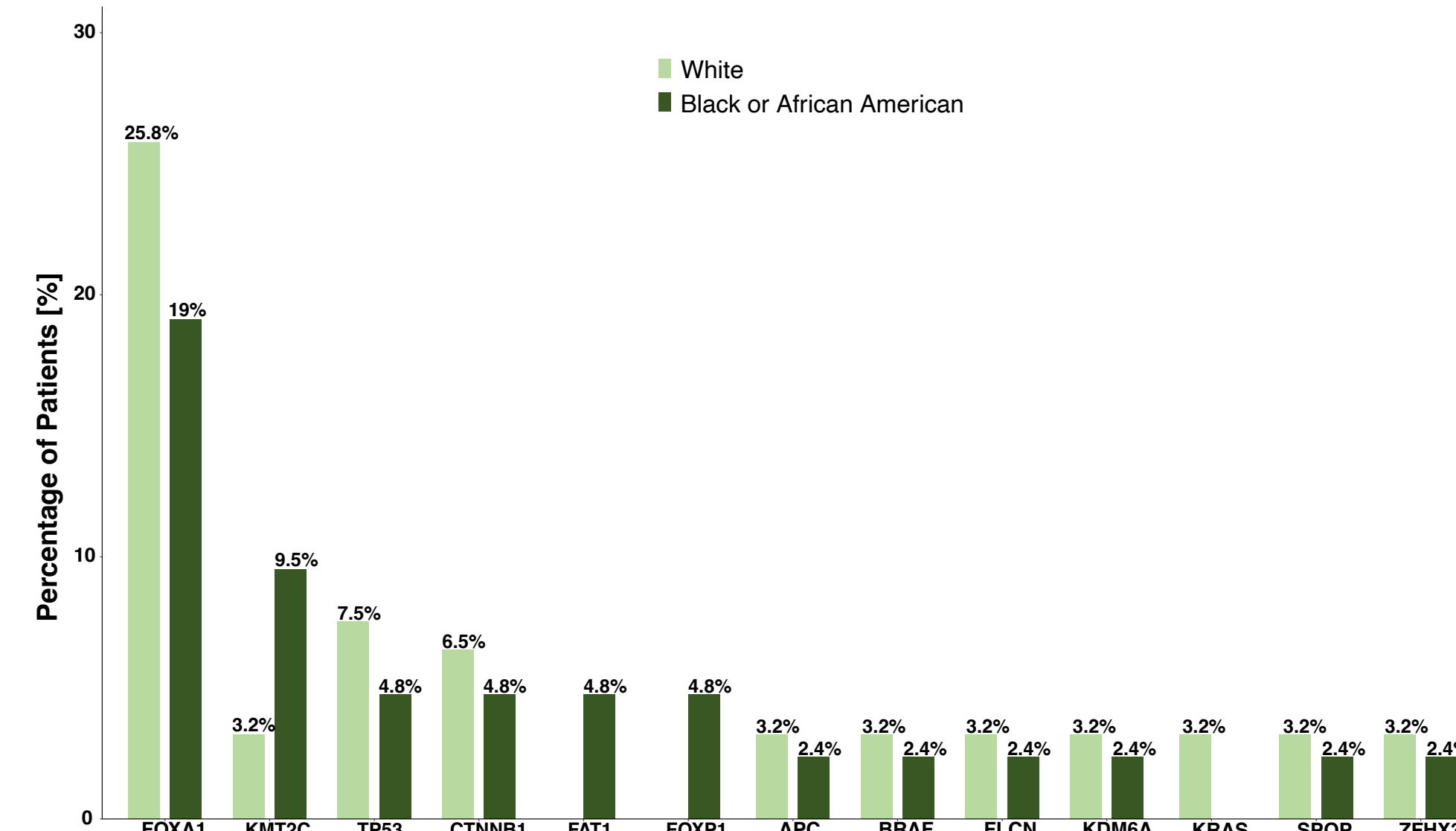


Figure 3. Short variant P/LP alterations found in 3% or more of the CDK12 altered population.

Most of the variants identified were truncating mutations. There were no differences in truncating vs. non-truncating alterations between White and African American men. There were no differences in the prevalence or distribution of co-mutations between the two populations.