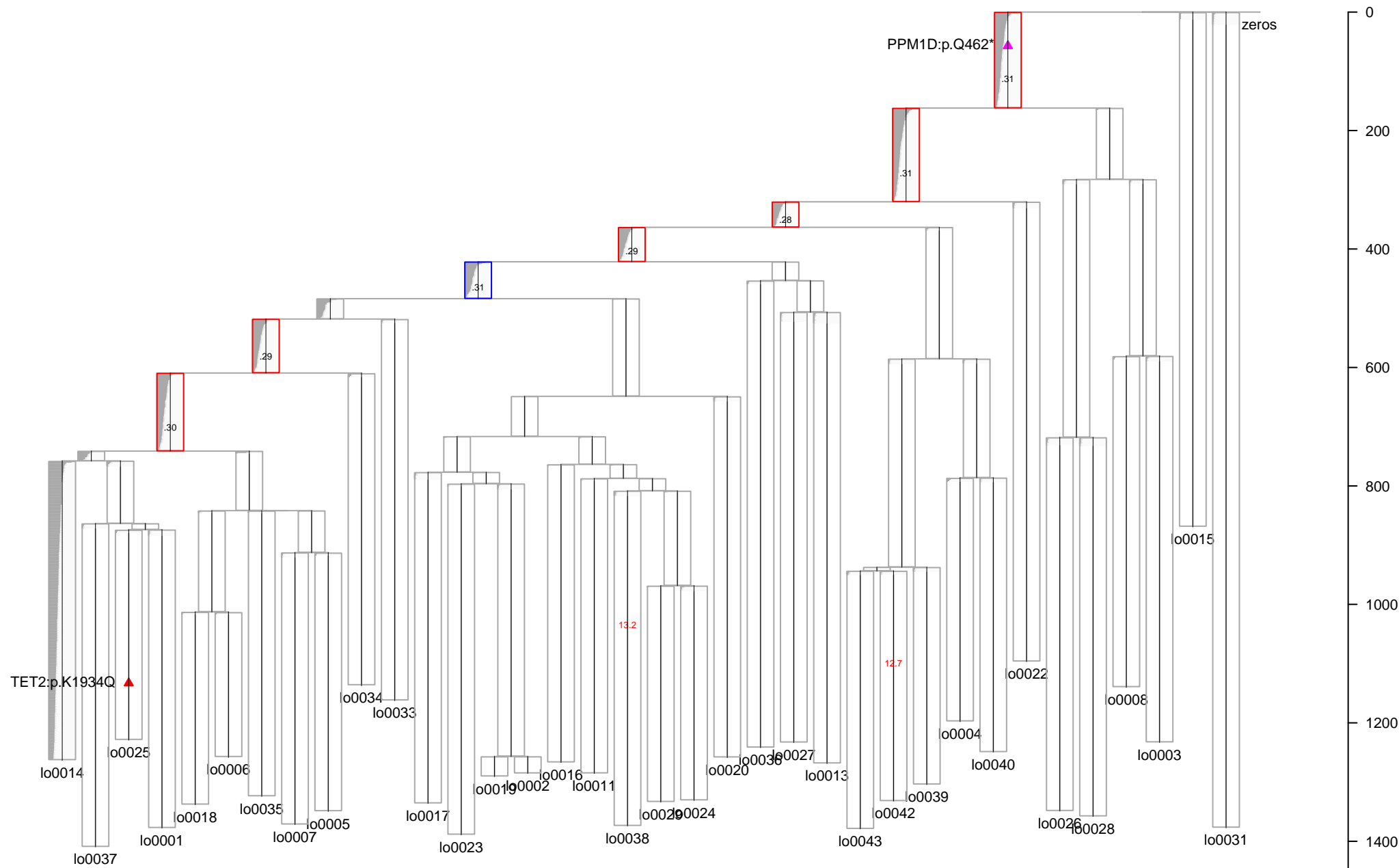


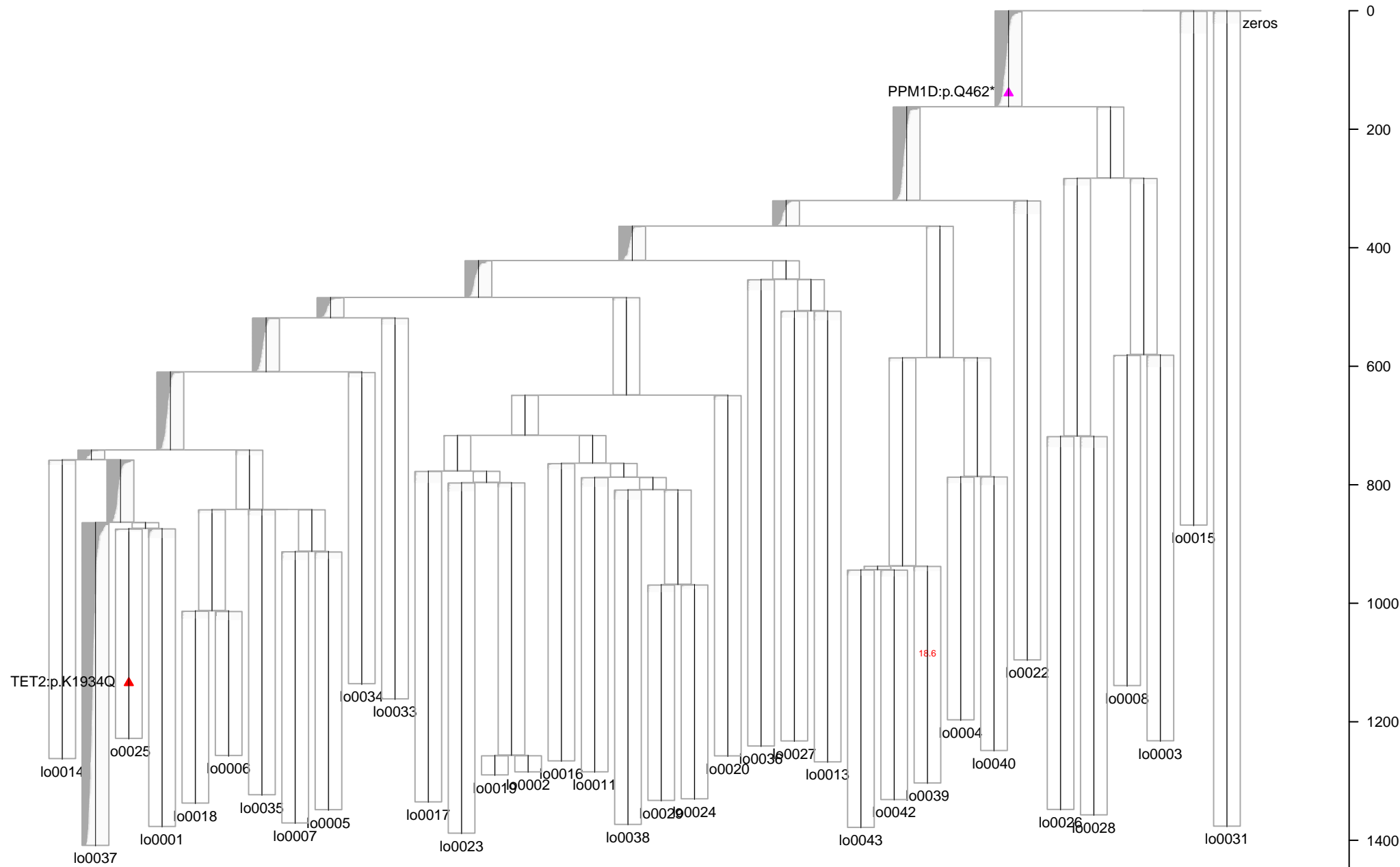
Tree By Colony Quality Assessment

This file reports the VAF distribution of the variants assigned to each branch on a per colony basis. This allows one to 'walk through' the trees on a per colony basis to visualise both the branch placement and VAF of all the variants present in that single colony with respect to the rest of the tree. This is particularly helpful to ensure that variants belonging to a single colony are not found in non-ancestral branches whilst also allowing one to assess if other branches in the tree suffer from a lack of sensitivity for picking up specific variants. The report includes all colonies - including those that are dropped from the final tree and also some additional samples of interest. For colonies that are in the final tree it is expected that the VAFs will be clonal on branches that are ancestral to the colony of interest and zero for those that are not ancestral. Branches are highlighted if they show significant deviation from this expectation ($VAF < 0.35$ and $VAF > 0.05$; Binomial test on aggregate mutant read count and aggregate depth; blue $p < 0.05$ and red $p < 0.05/\text{number of branches}$). Branches where the depth is significantly lower than the depth of variants across the whole tree are annotated with the branch depth shown in red.

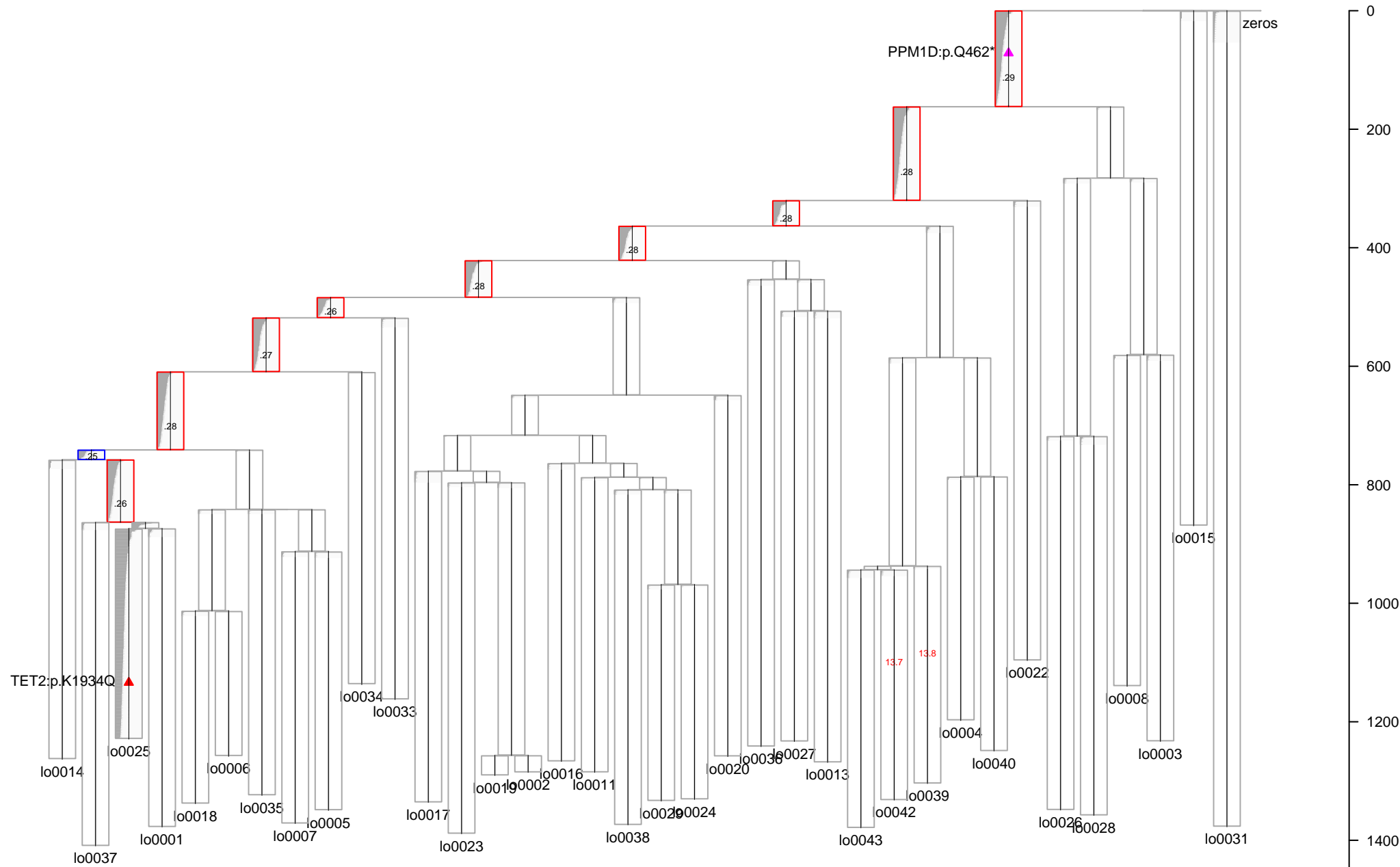
PD6634: Annotated with VAF from lo0014
Mean Depth=13.73



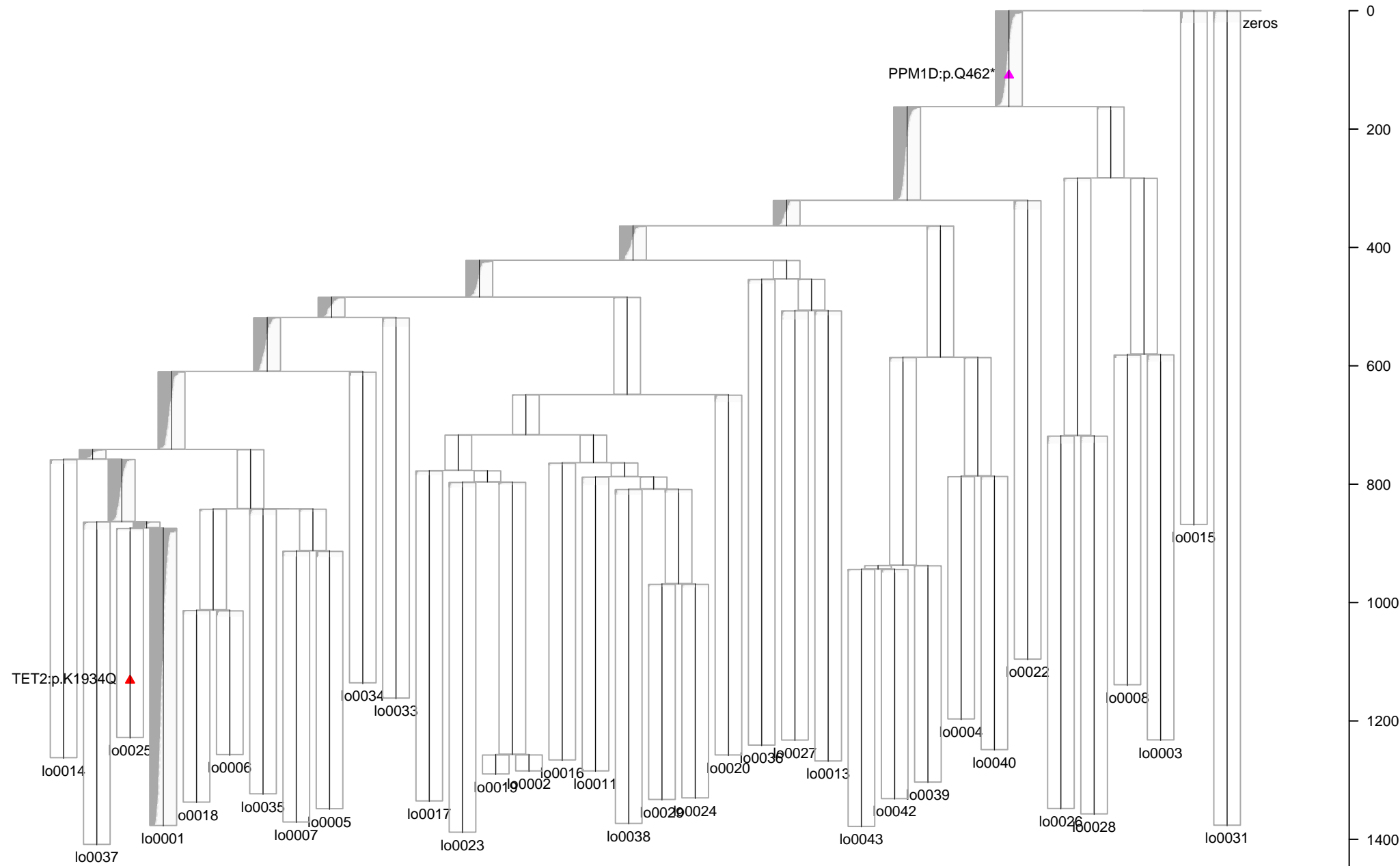
PD6634: Annotated with VAF from lo0037
Mean Depth=19.43



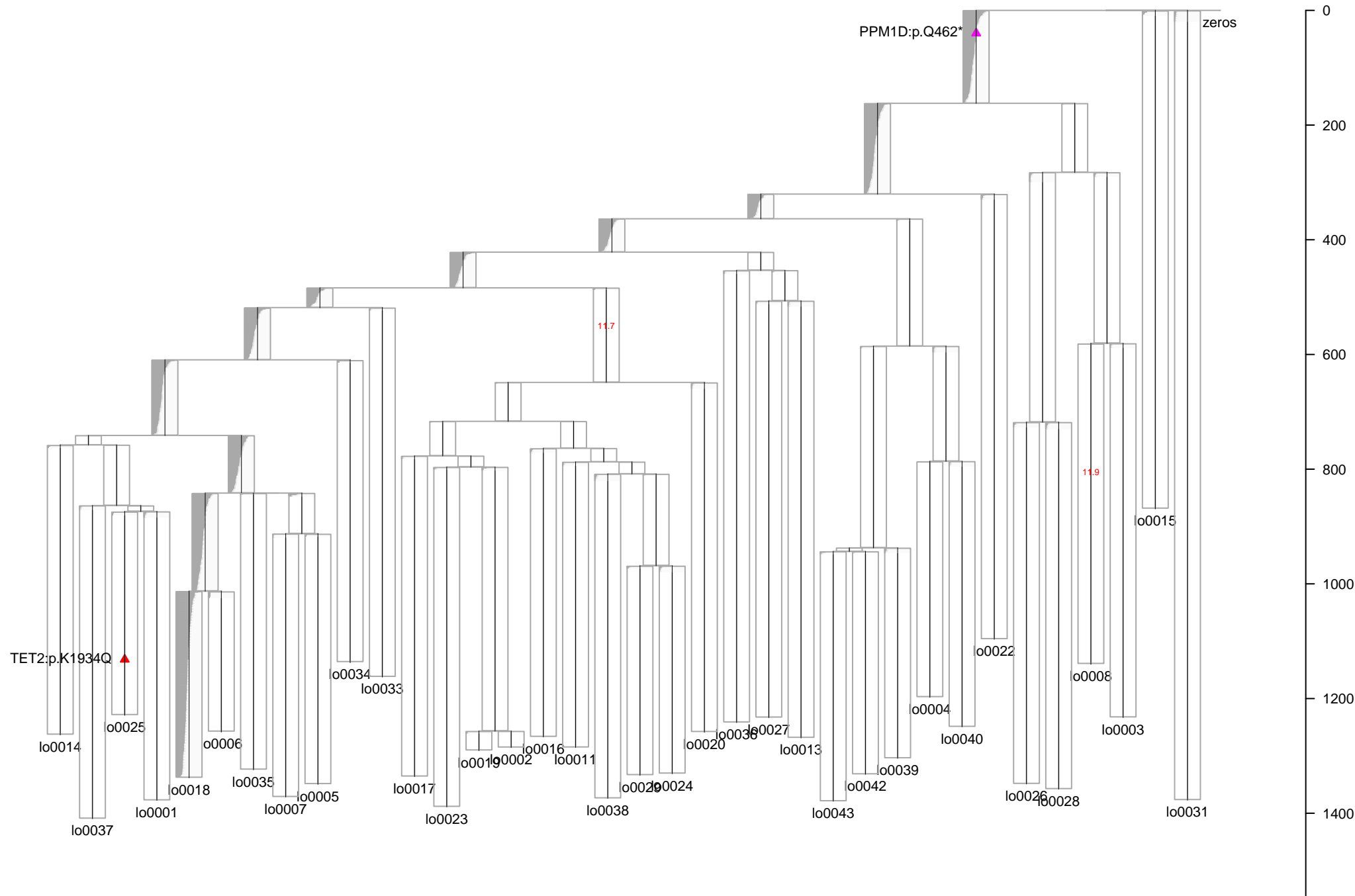
PD6634: Annotated with VAF from lo0025
Mean Depth=14.45



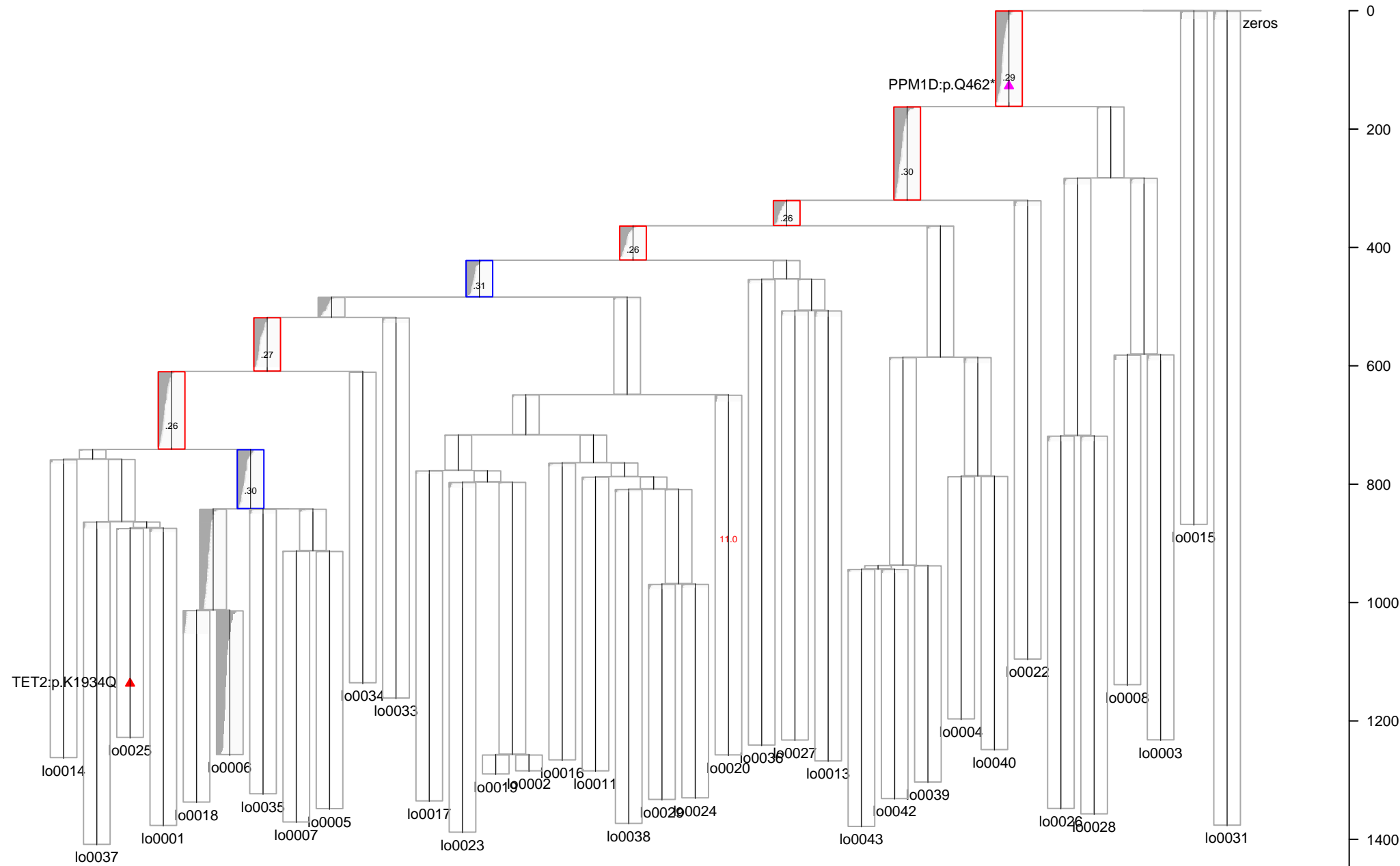
PD6634: Annotated with VAF from lo0001
Mean Depth=14.30



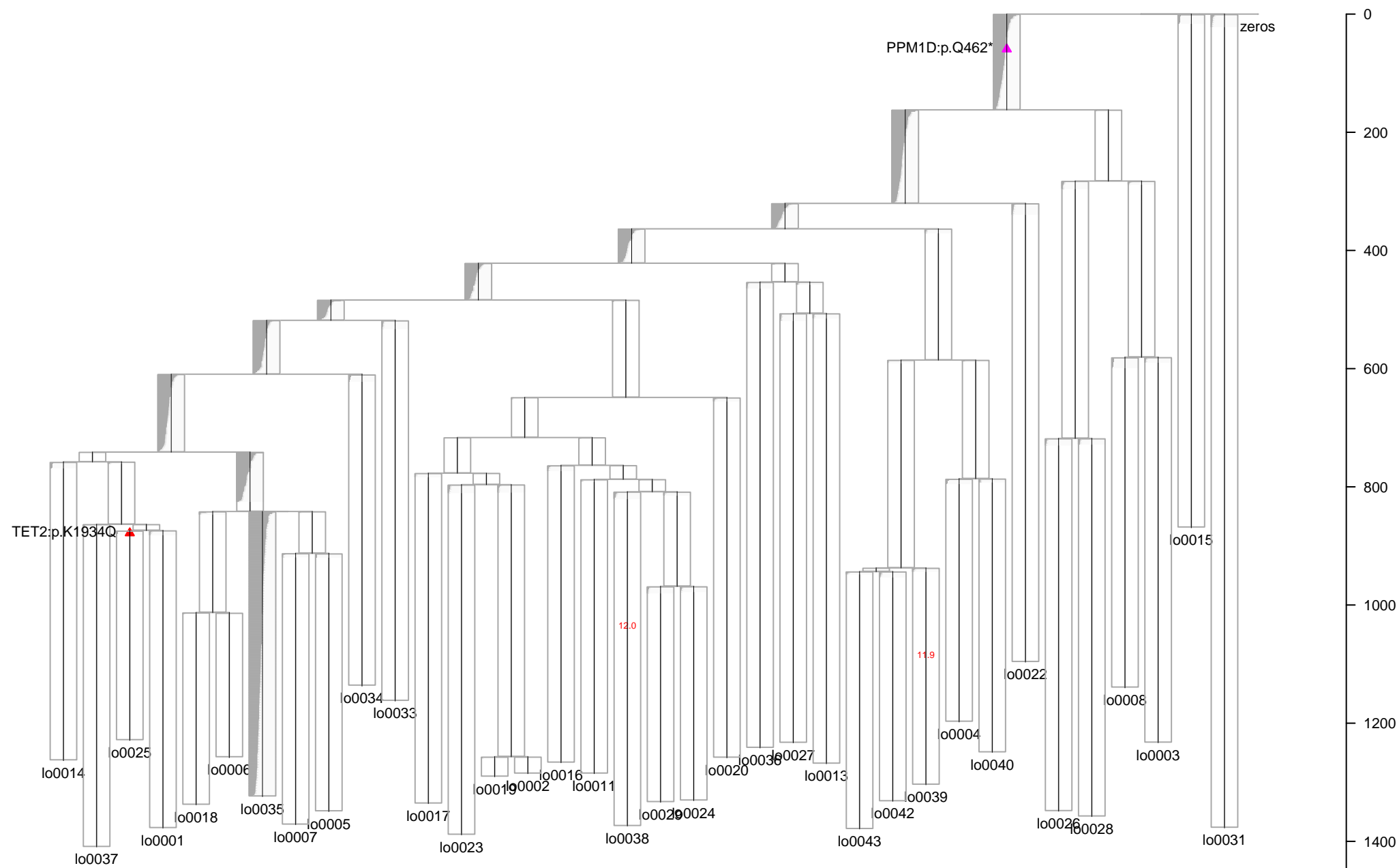
PD6634: Annotated with VAF from lo0018
Mean Depth=12.60



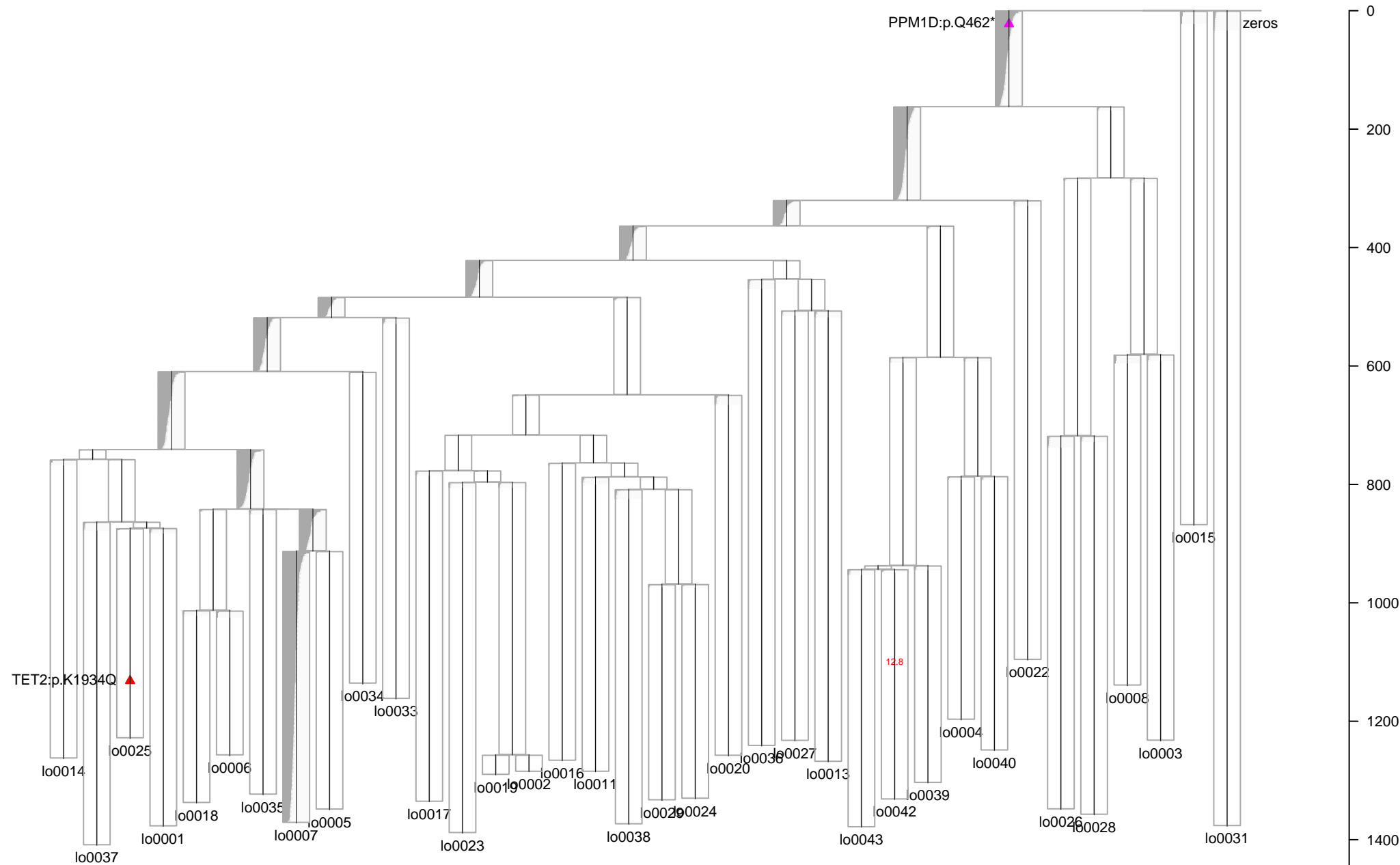
PD6634: Annotated with VAF from lo0006
Mean Depth=11.54



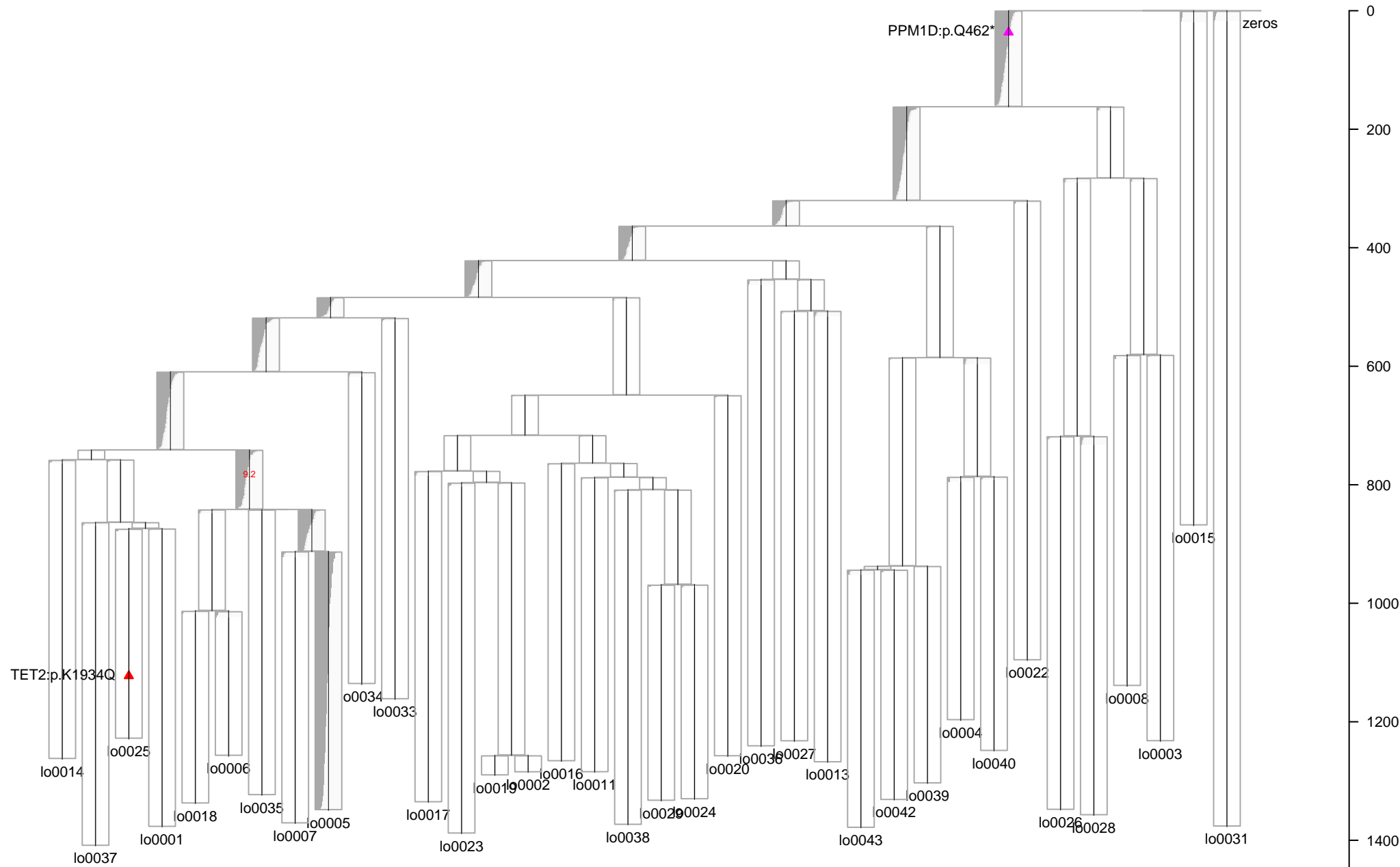
PD6634: Annotated with VAF from lo0035
Mean Depth=12.59



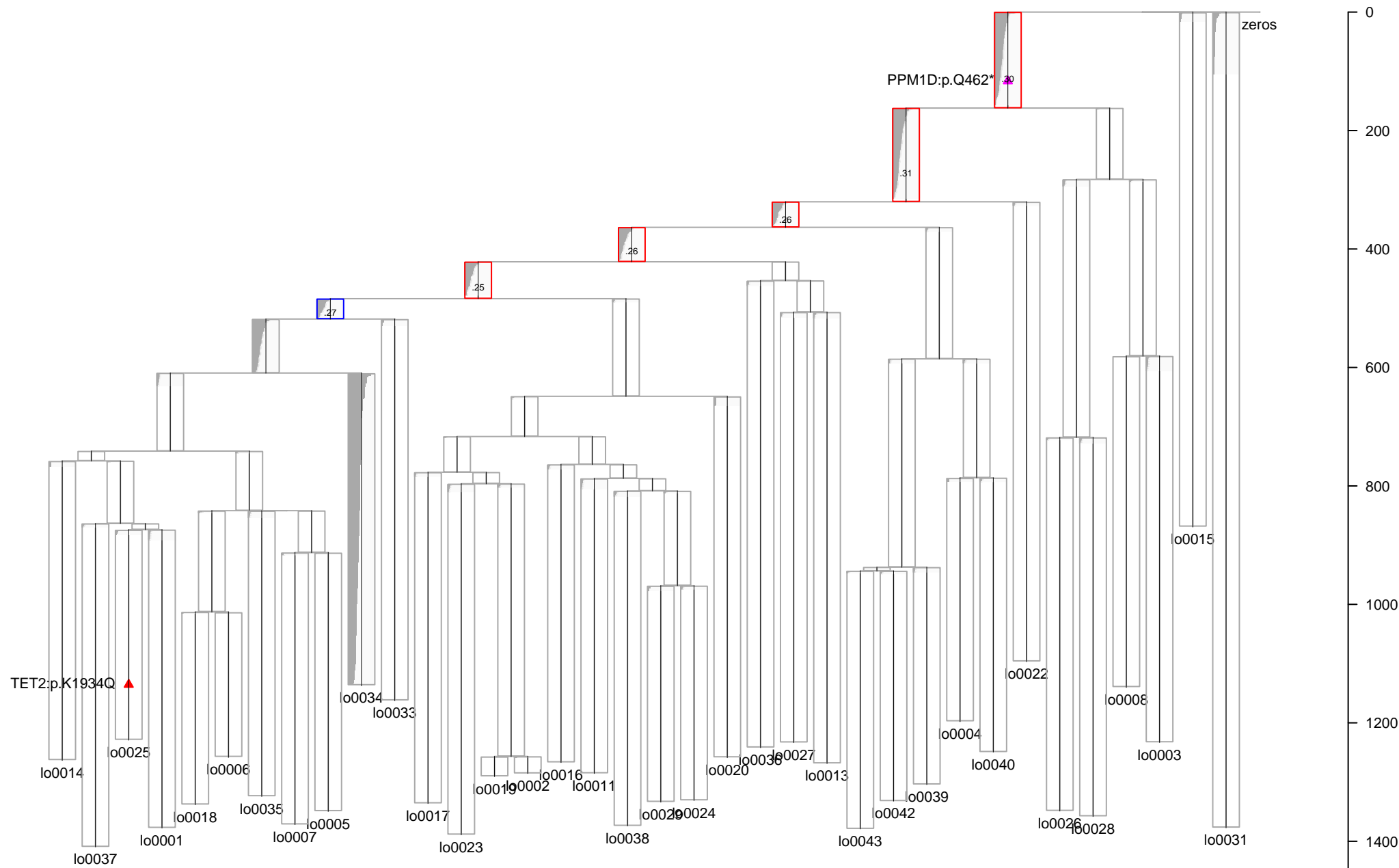
PD6634: Annotated with VAF from lo0007
Mean Depth=13.35



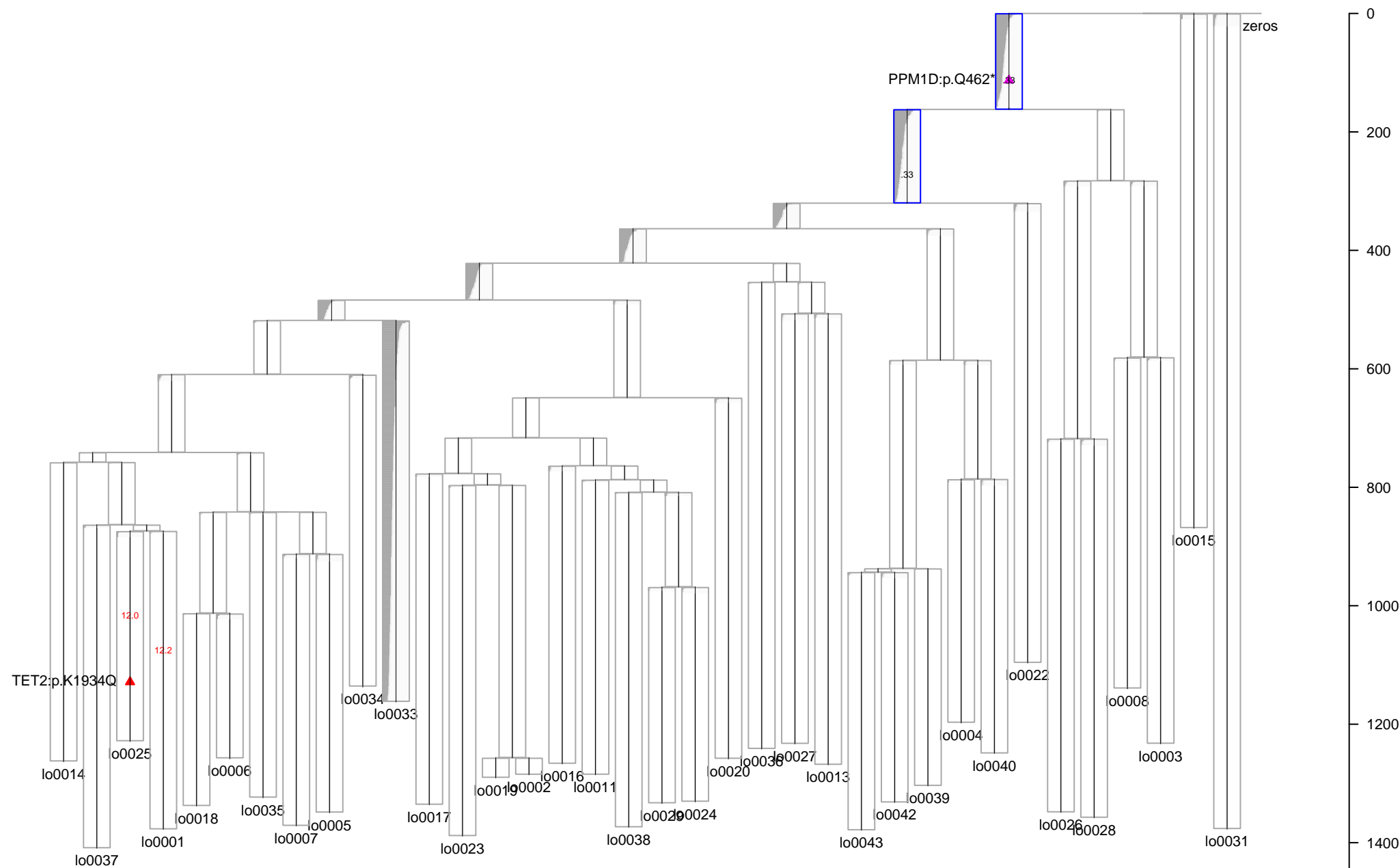
PD6634: Annotated with VAF from lo0005
Mean Depth=10.02



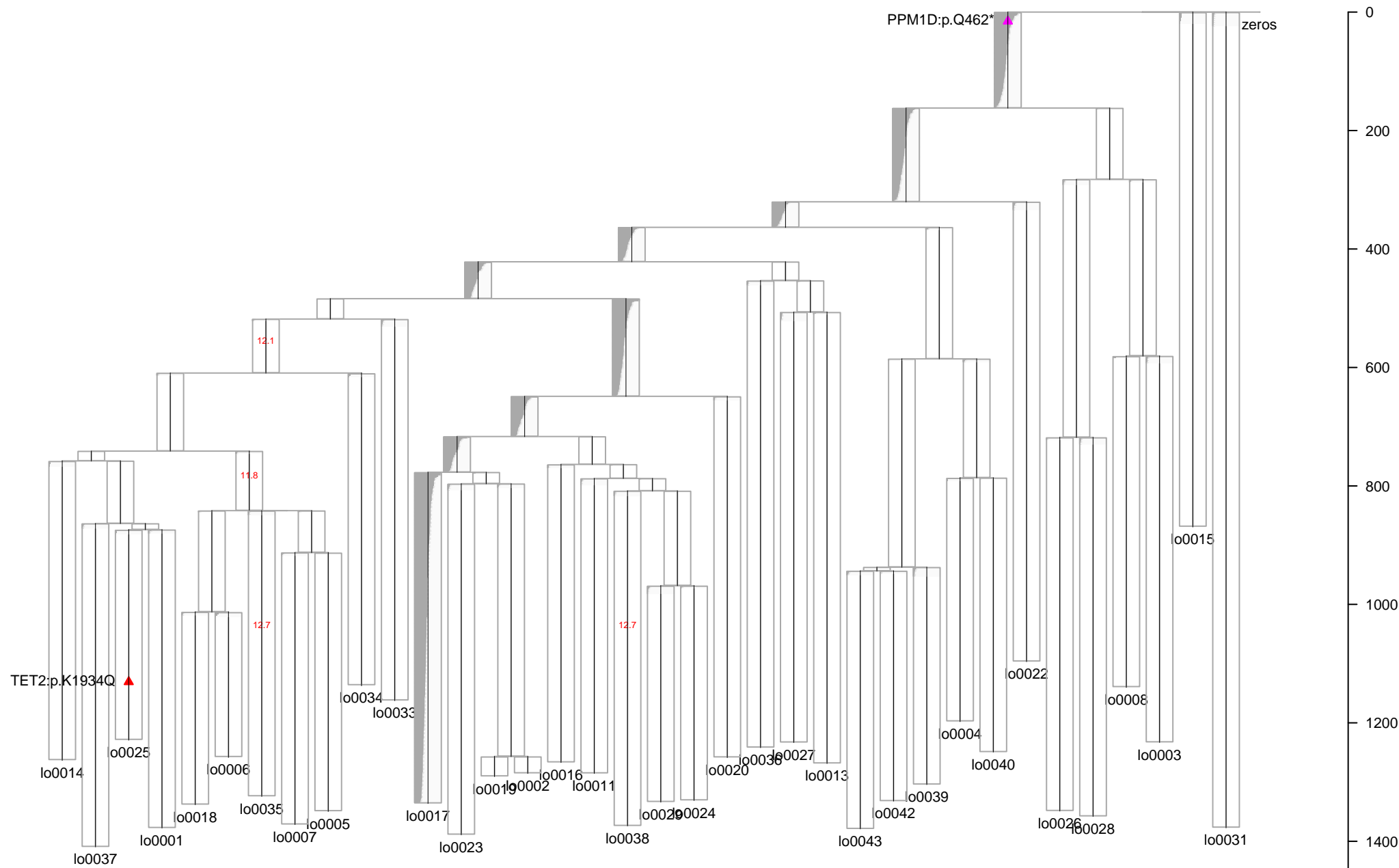
PD6634: Annotated with VAF from lo0034
Mean Depth=10.35



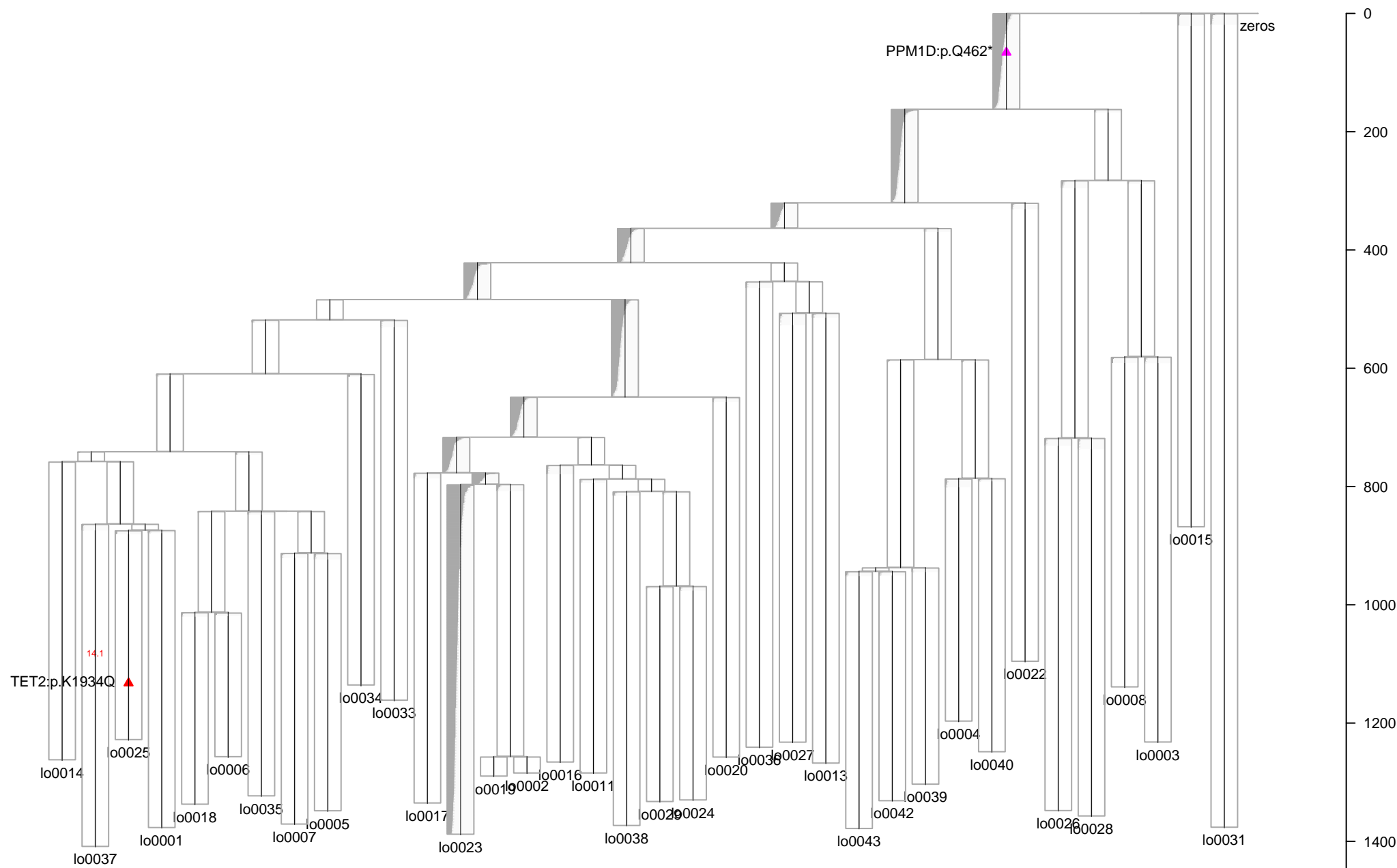
PD6634: Annotated with VAF from lo0033
Mean Depth=12.77



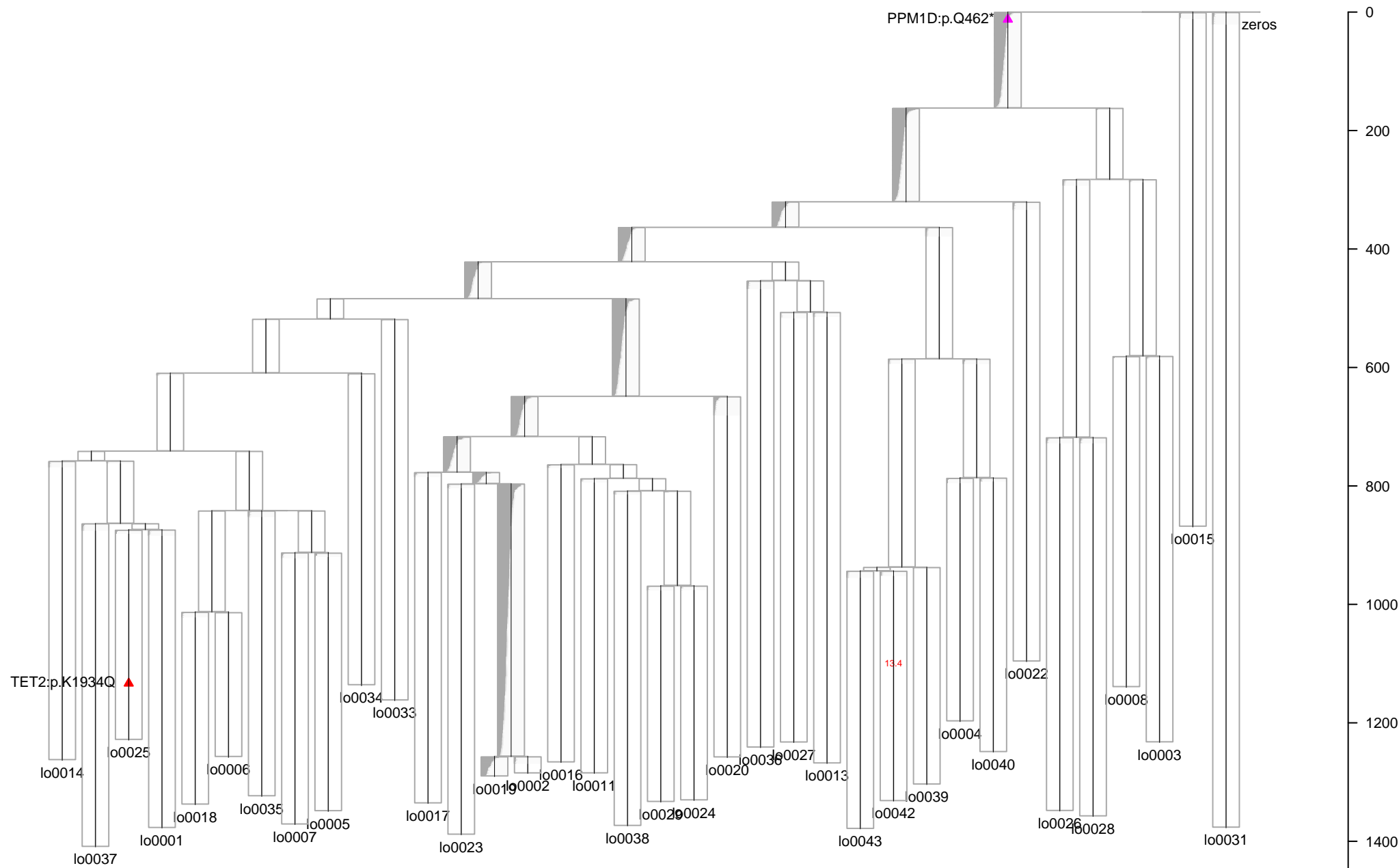
PD6634: Annotated with VAF from lo0017
Mean Depth=13.32



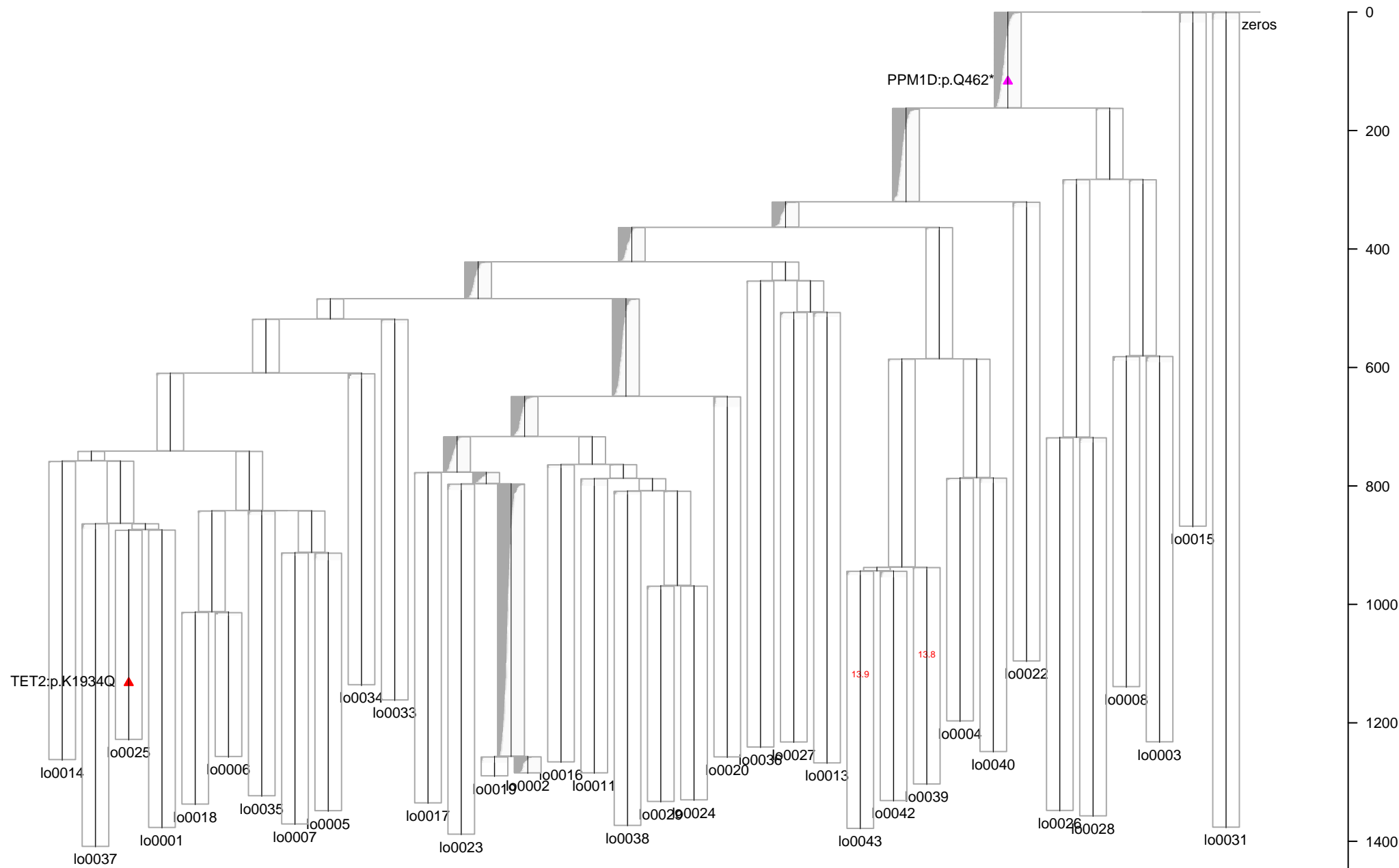
PD6634: Annotated with VAF from lo0023
Mean Depth=14.64



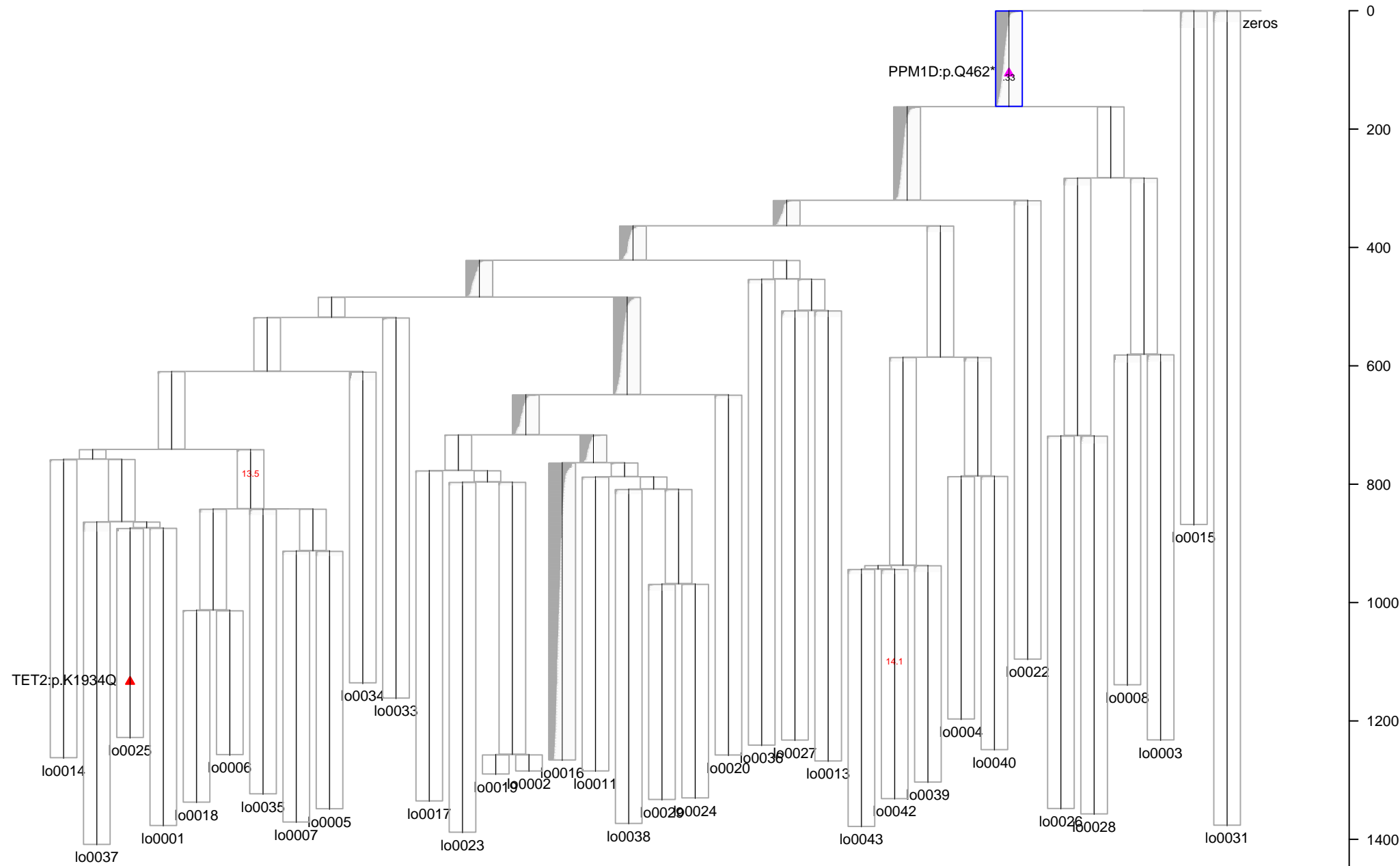
PD6634: Annotated with VAF from lo0019
Mean Depth=13.94



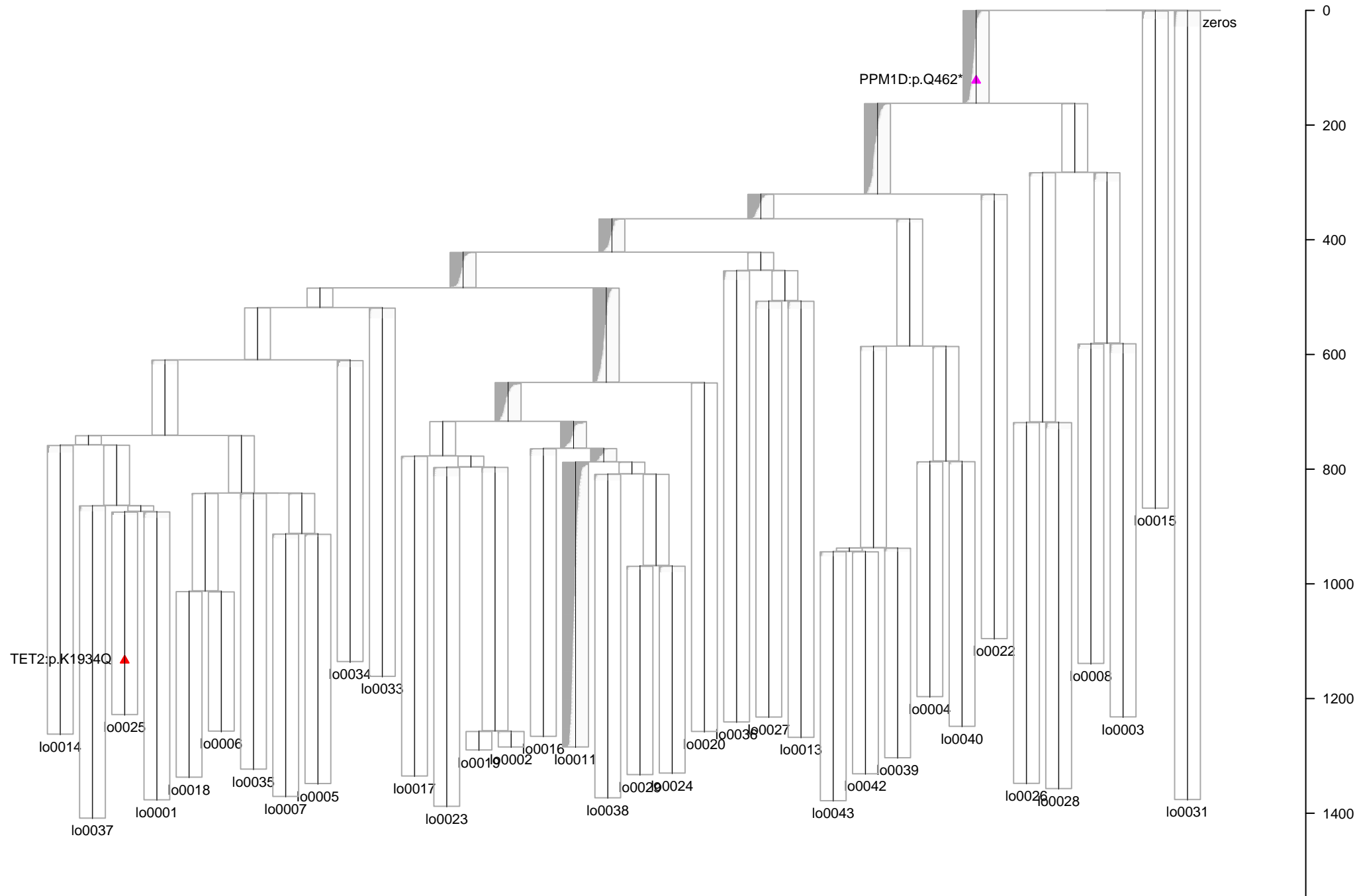
PD6634: Annotated with VAF from lo0002
Mean Depth=14.45



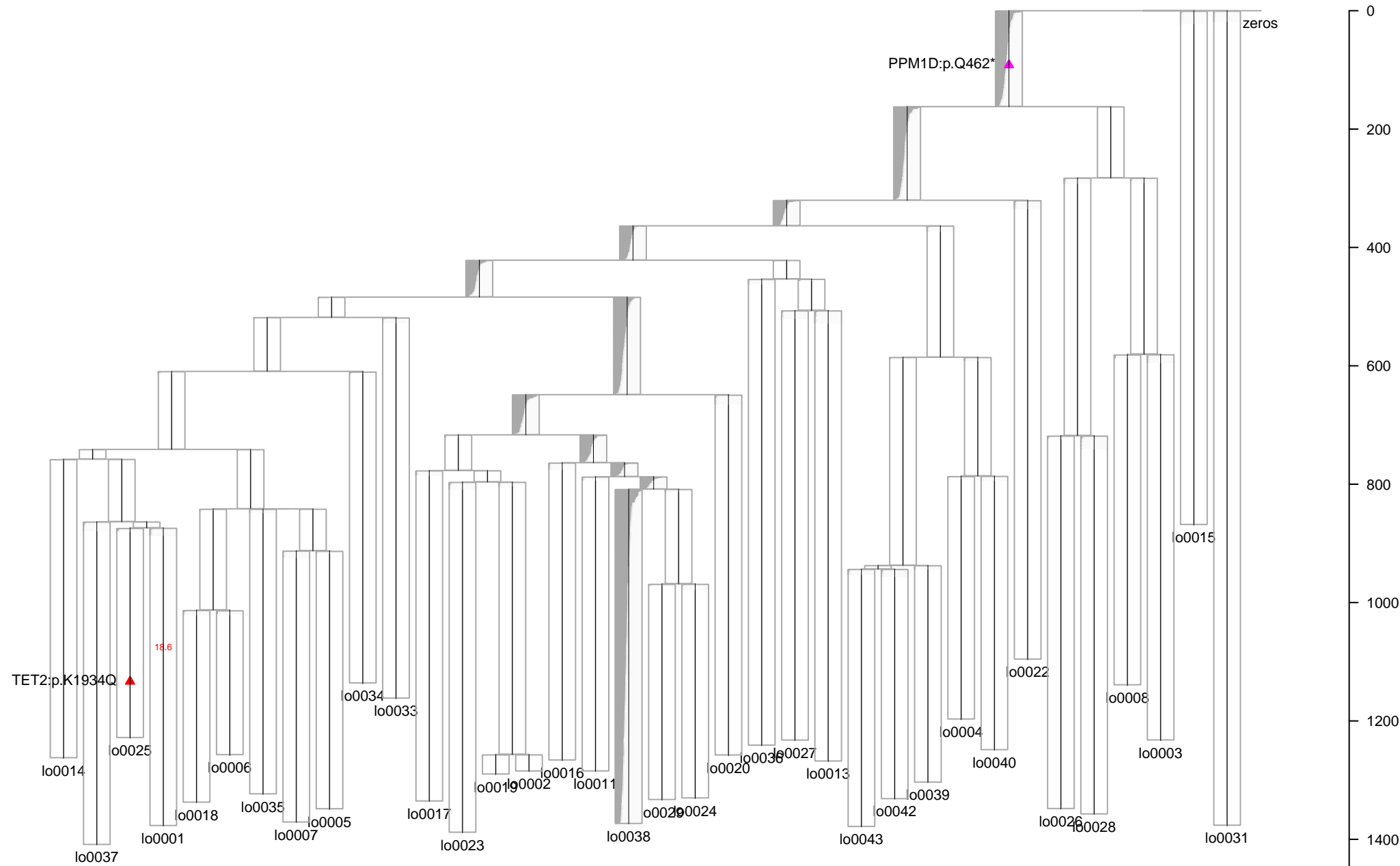
PD6634: Annotated with VAF from lo0016
Mean Depth=14.72



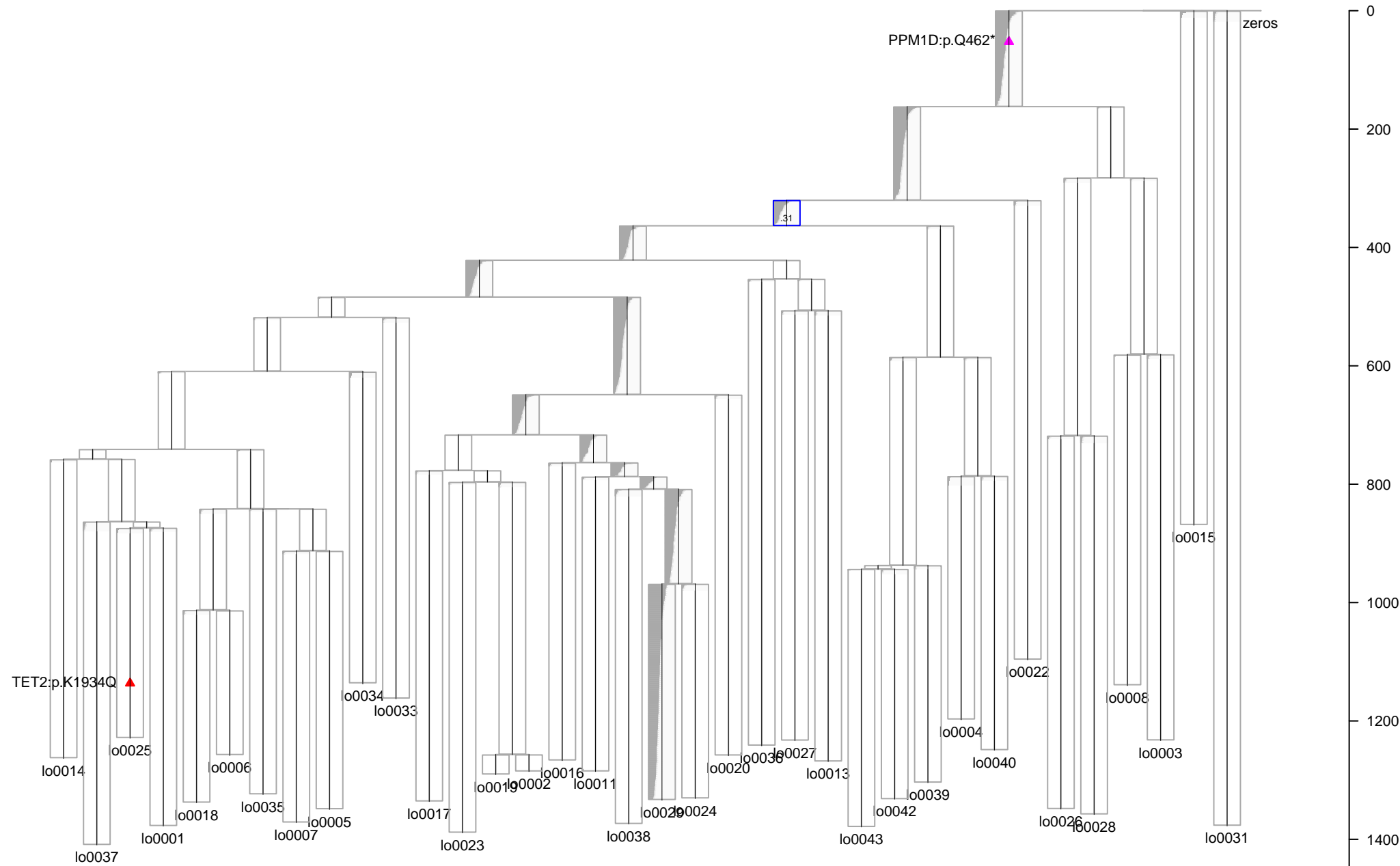
PD6634: Annotated with VAF from lo0011
Mean Depth=13.04



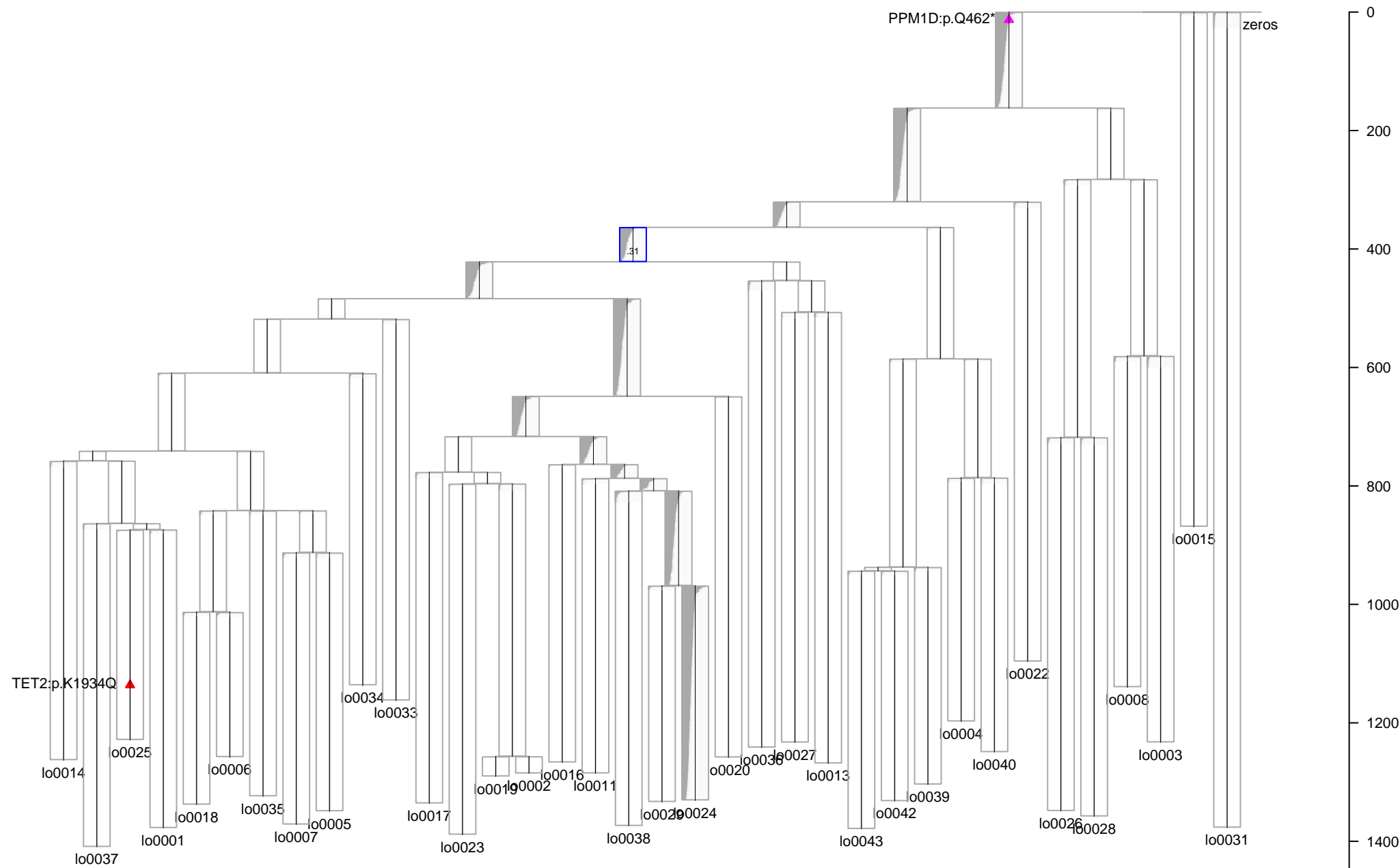
PD6634: Annotated with VAF from lo0038
Mean Depth=19.36



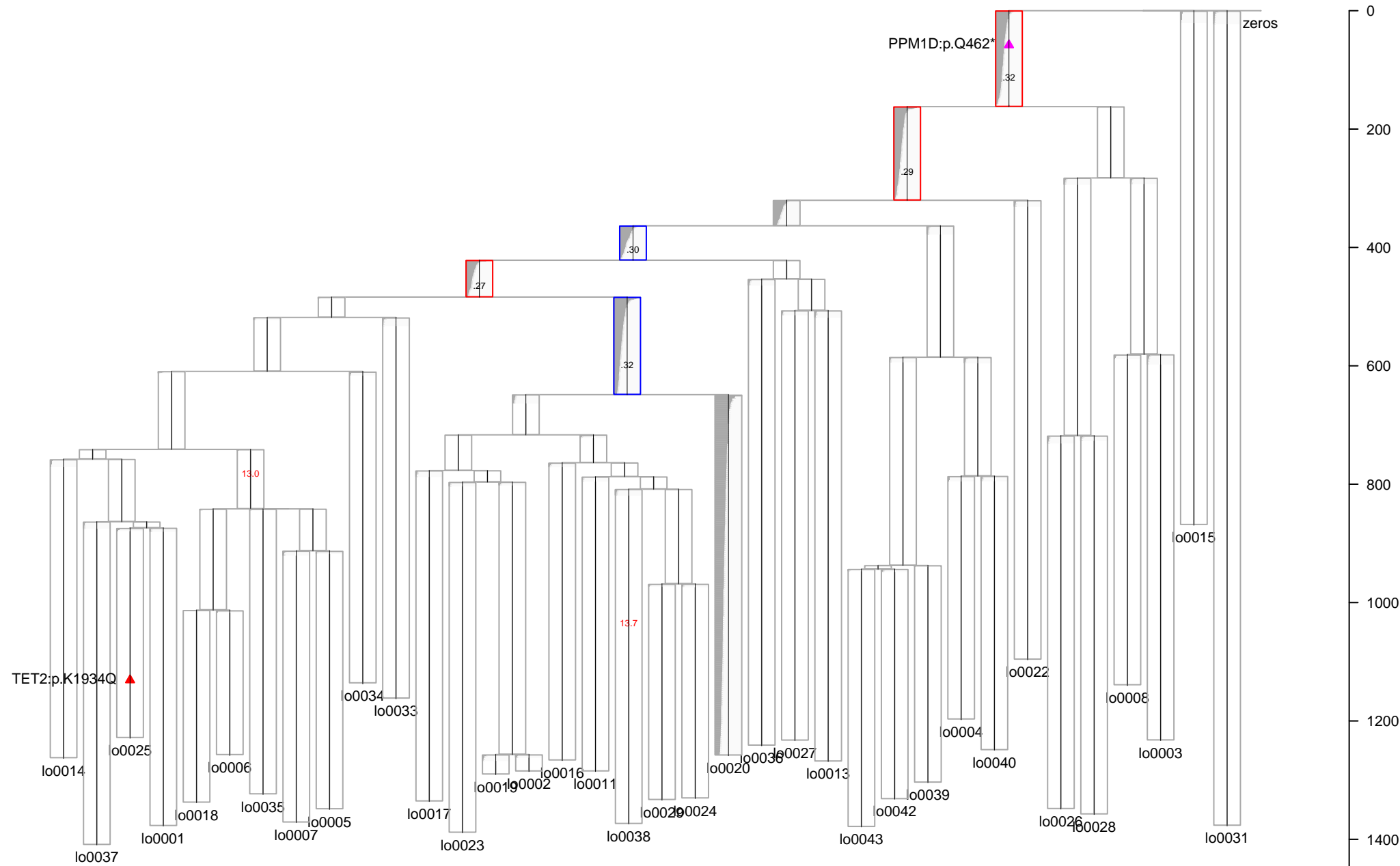
PD6634: Annotated with VAF from lo0029
Mean Depth=12.66



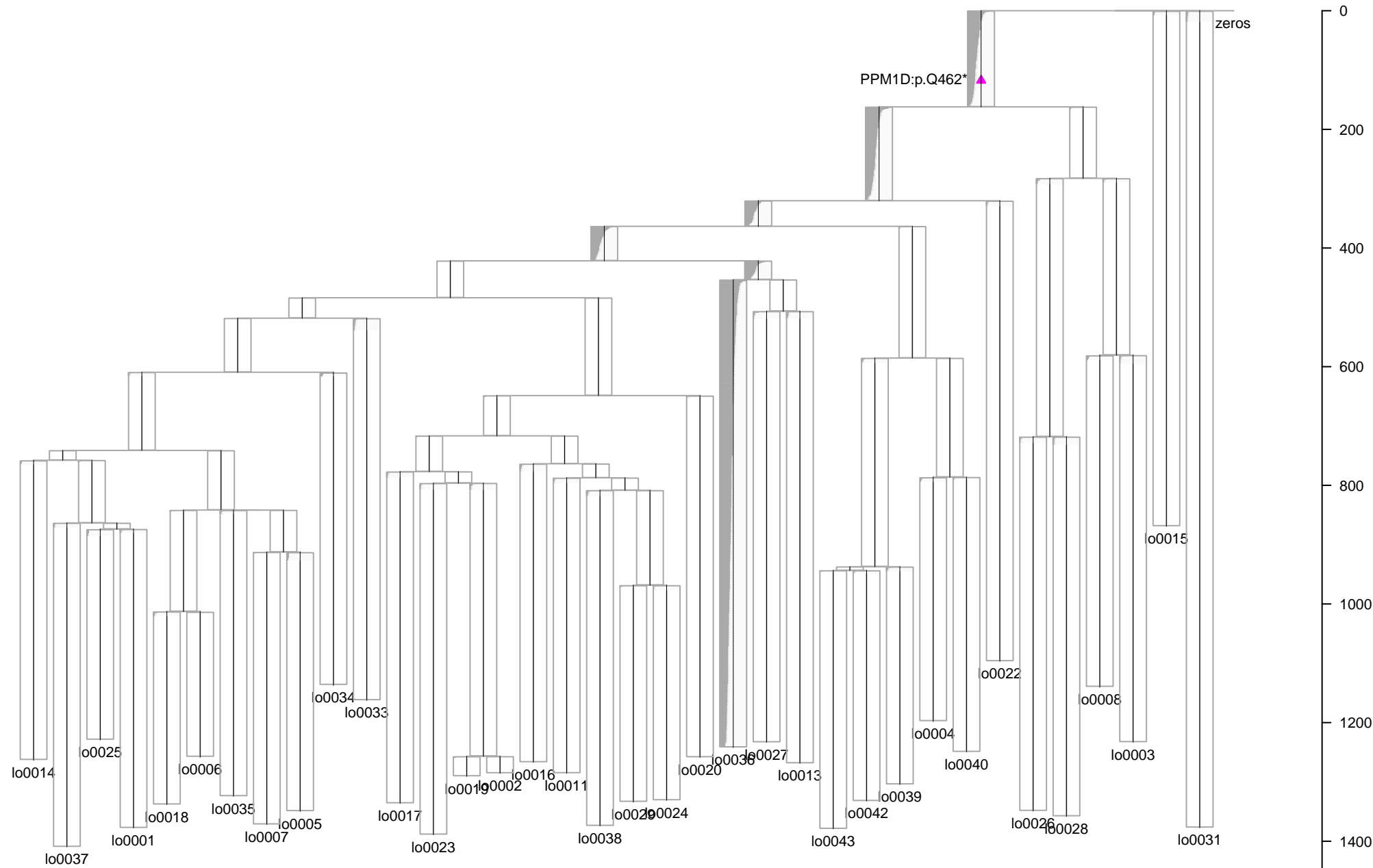
PD6634: Annotated with VAF from lo0024
Mean Depth=15.59



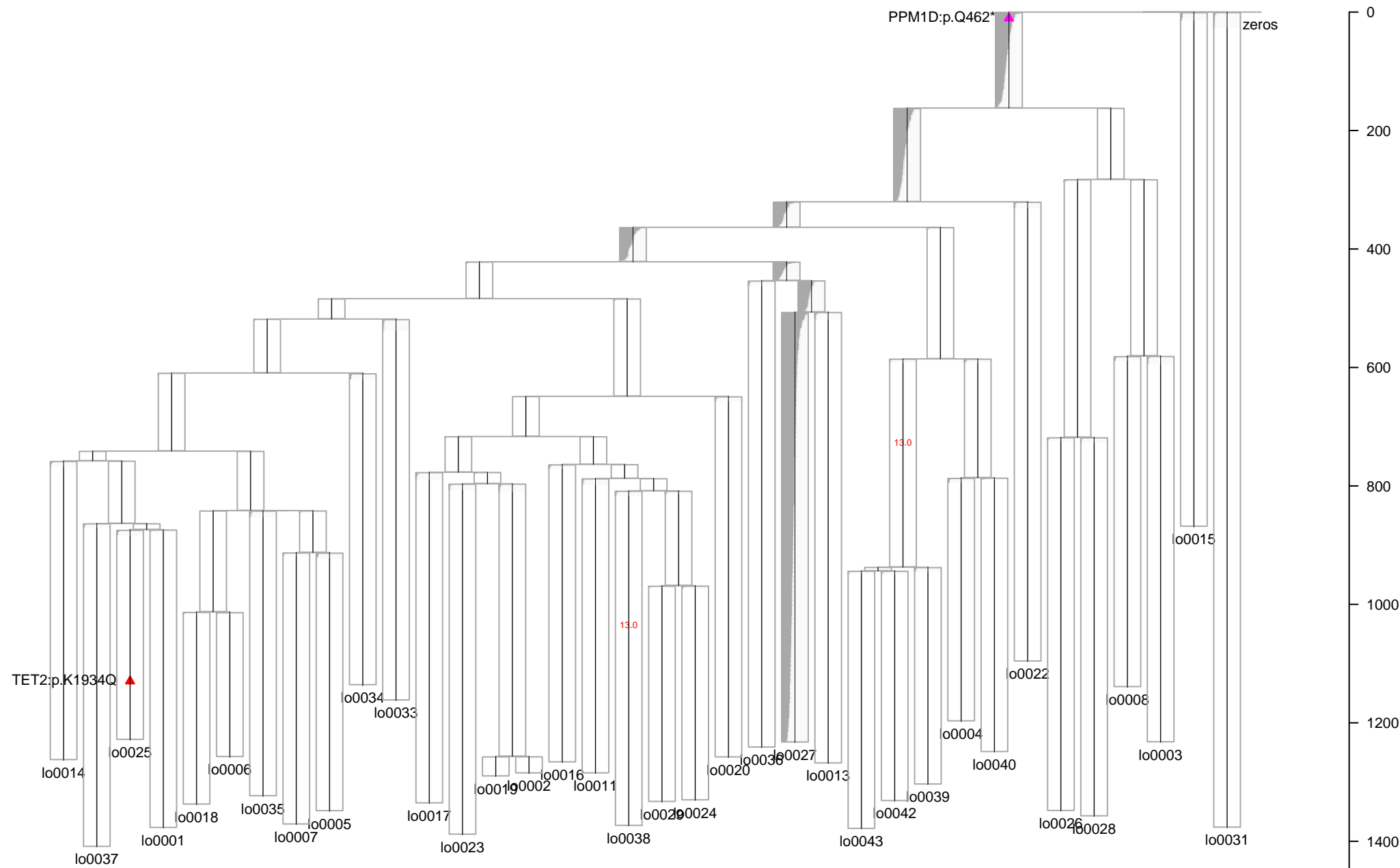
PD6634: Annotated with VAF from lo0020
Mean Depth=14.17



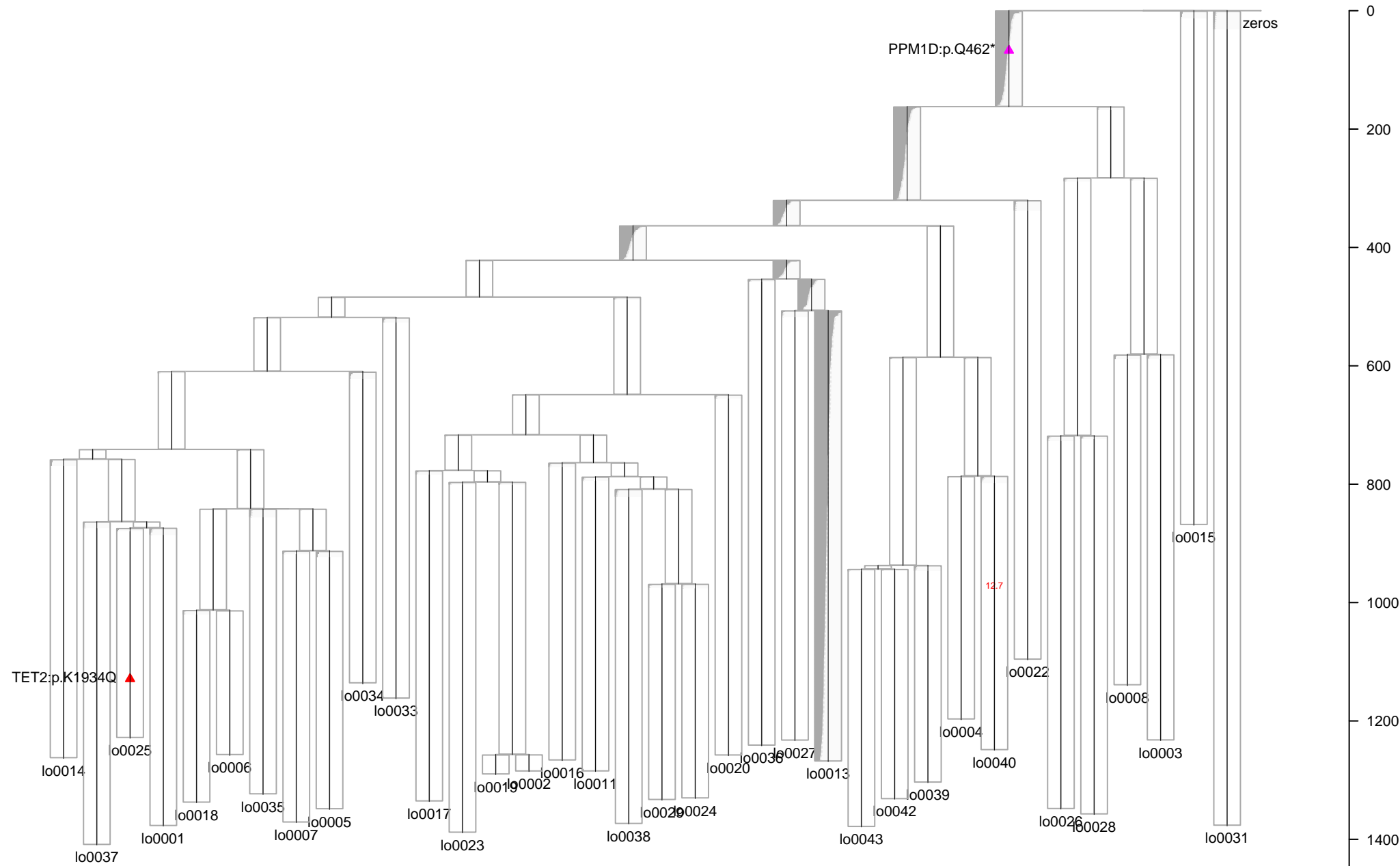
PD6634: Annotated with VAF from lo0036
Mean Depth=12.28



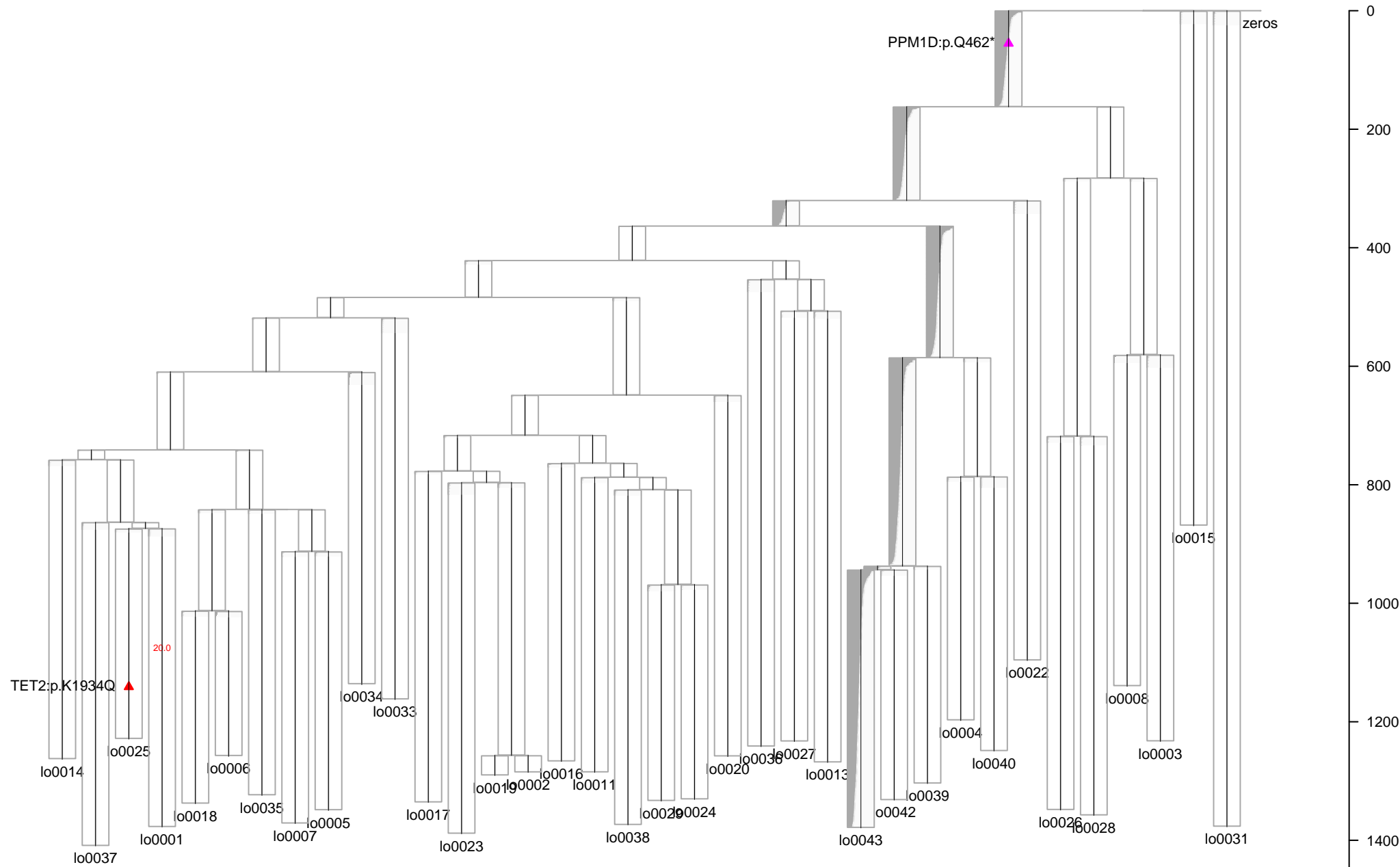
PD6634: Annotated with VAF from lo0027
Mean Depth=13.85



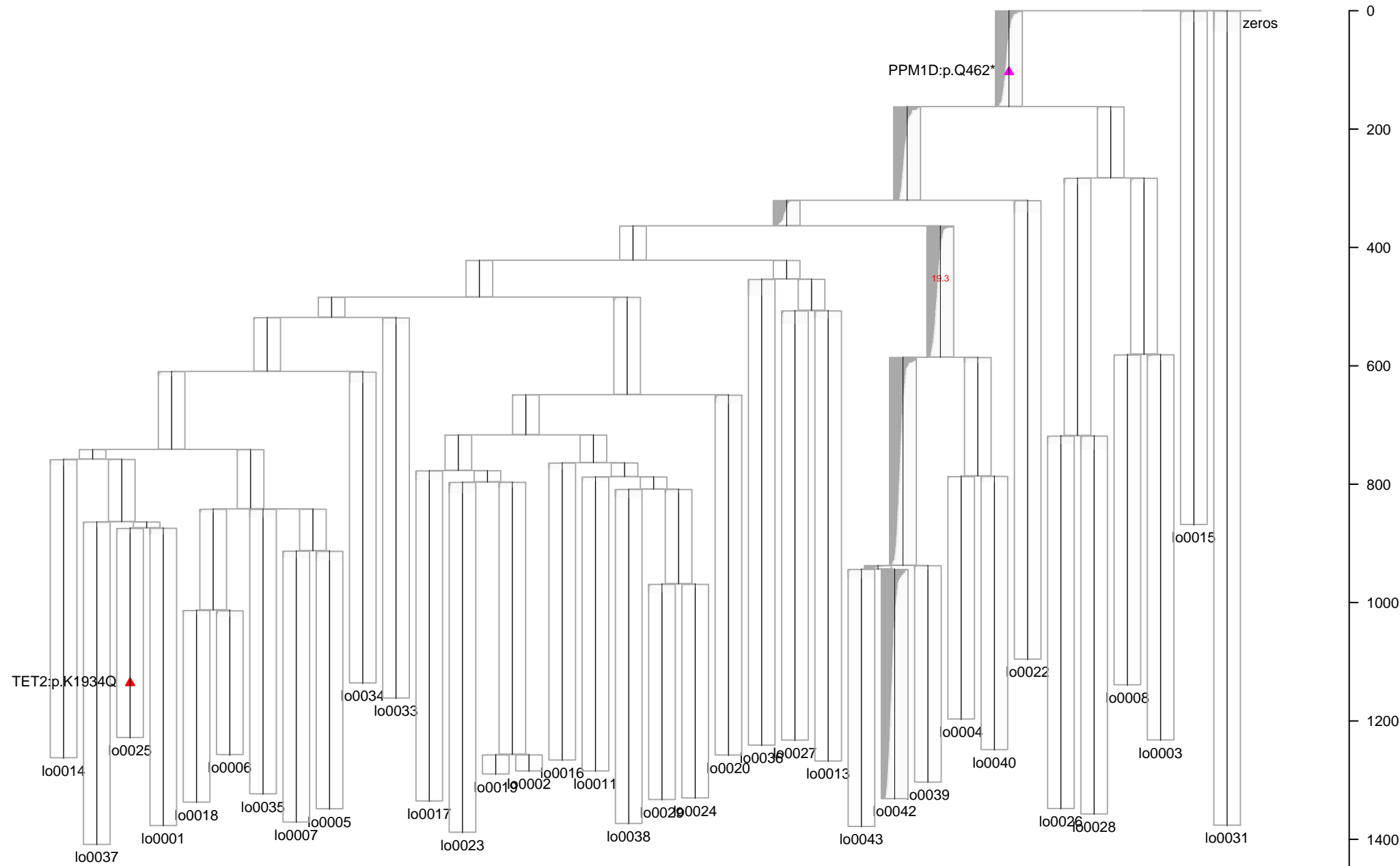
PD6634: Annotated with VAF from lo0013
Mean Depth=13.52



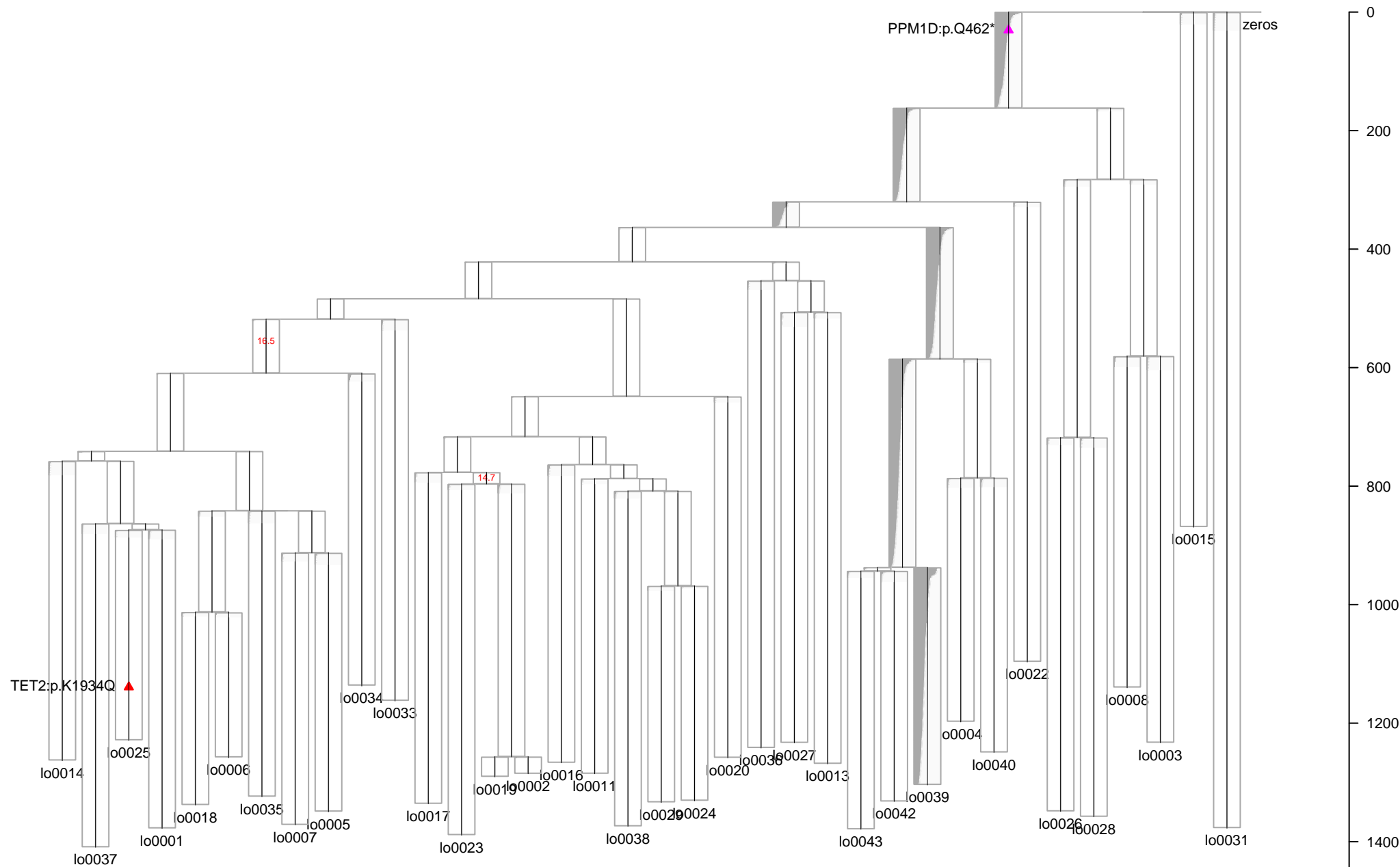
PD6634: Annotated with VAF from lo0043
Mean Depth=20.87



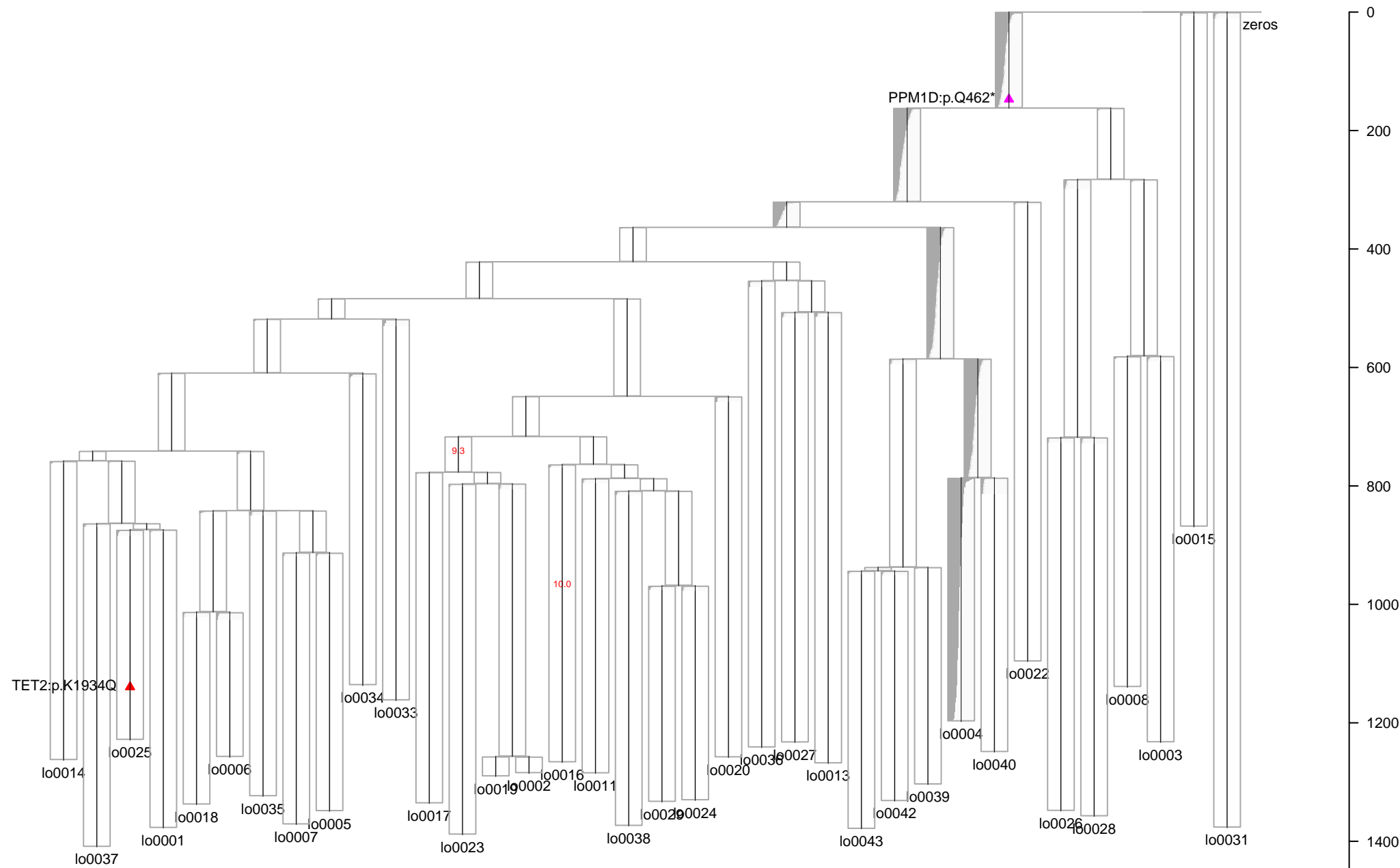
PD6634: Annotated with VAF from lo0042
Mean Depth=20.34



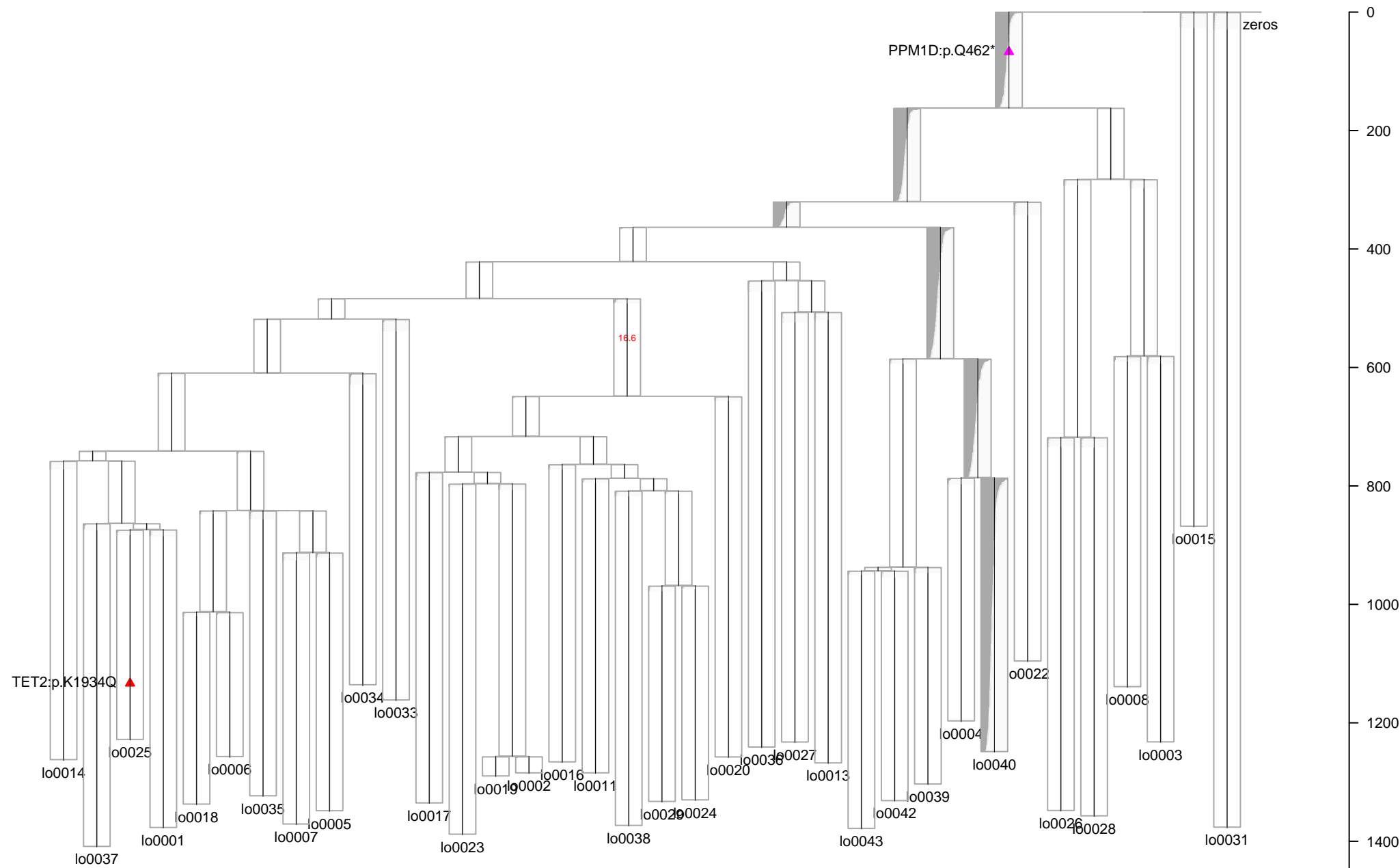
PD6634: Annotated with VAF from lo0039
Mean Depth=17.70



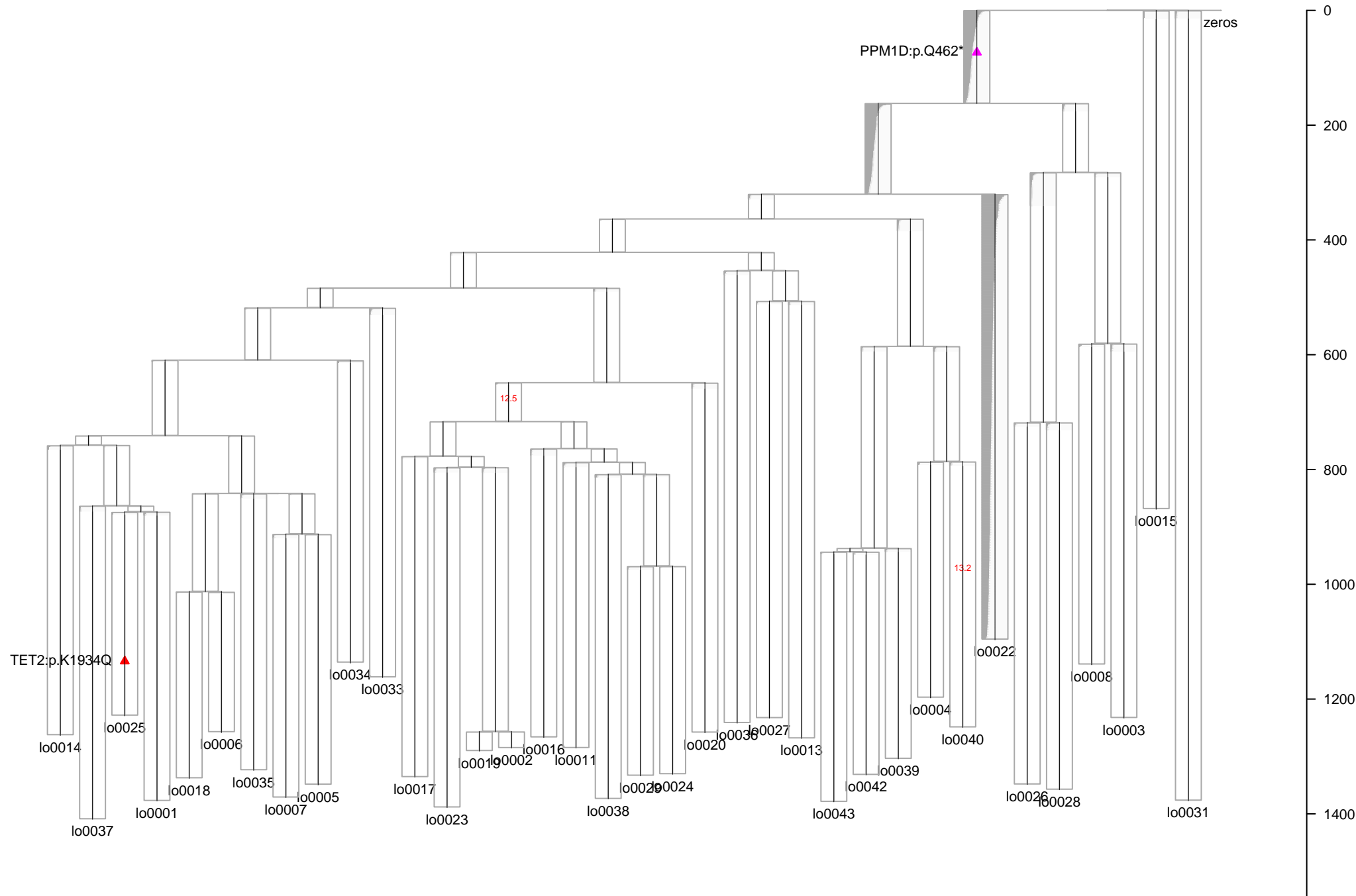
PD6634: Annotated with VAF from lo0004
Mean Depth=10.42



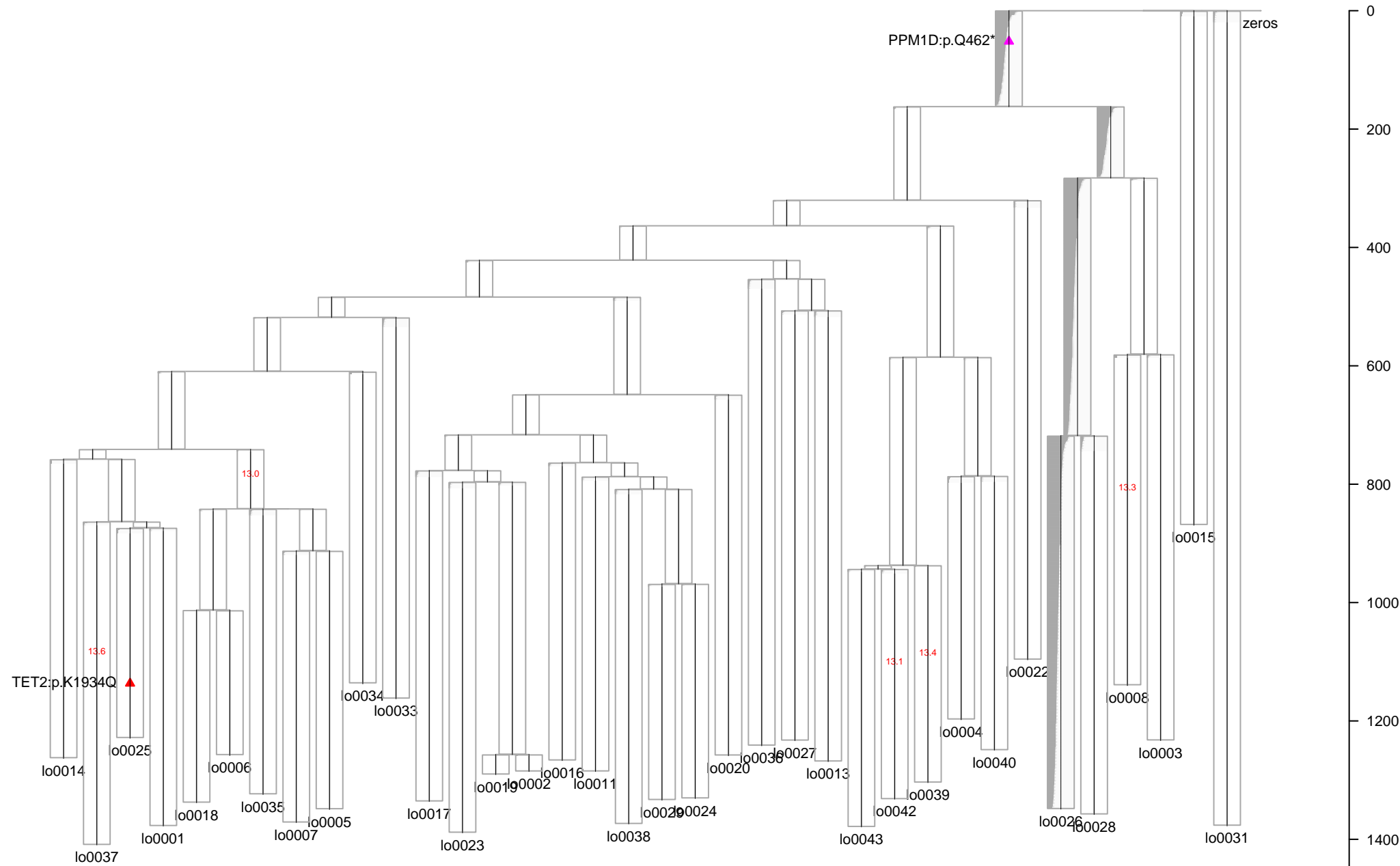
PD6634: Annotated with VAF from lo0040
Mean Depth=17.84



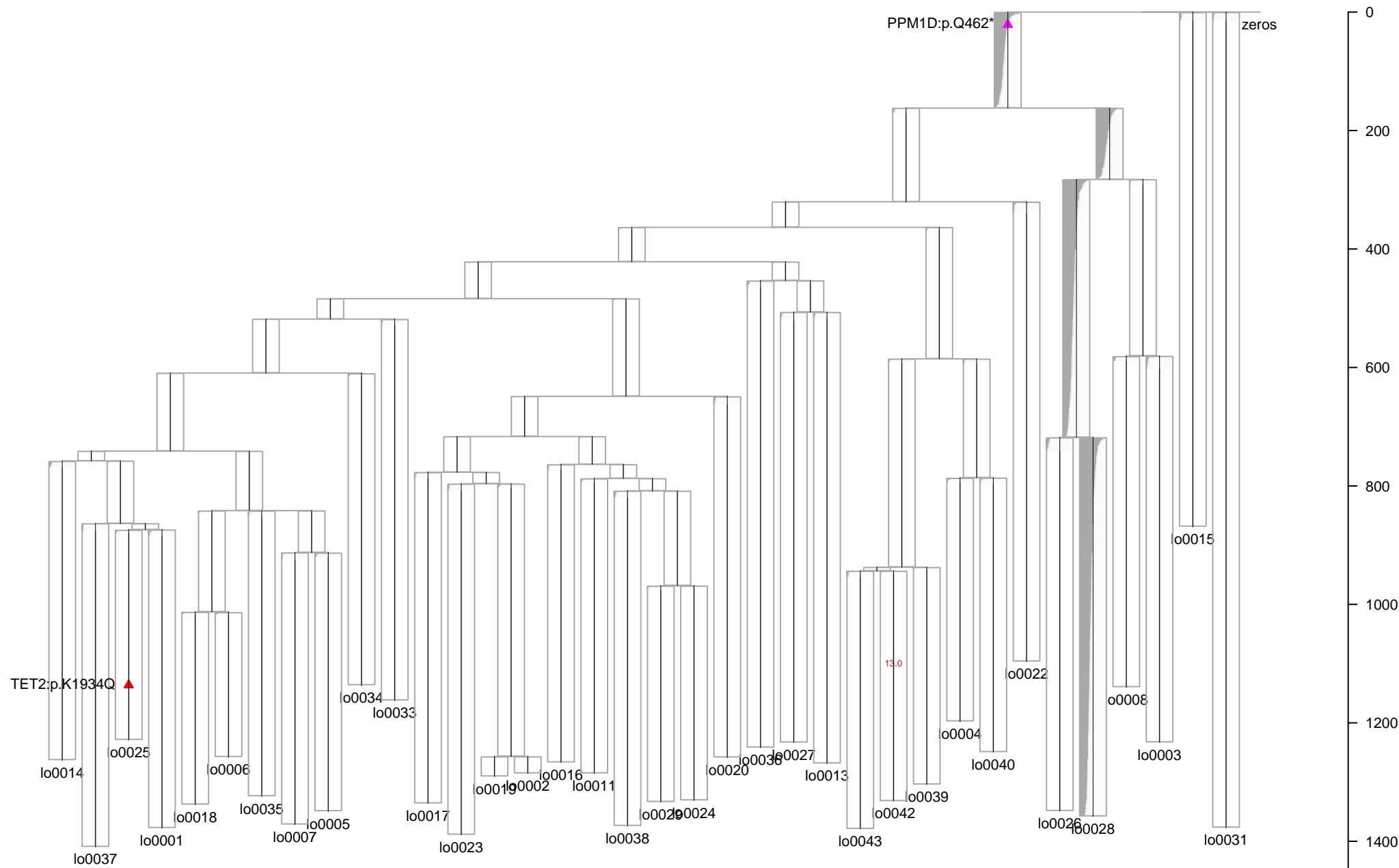
PD6634: Annotated with VAF from lo0022
Mean Depth=13.76



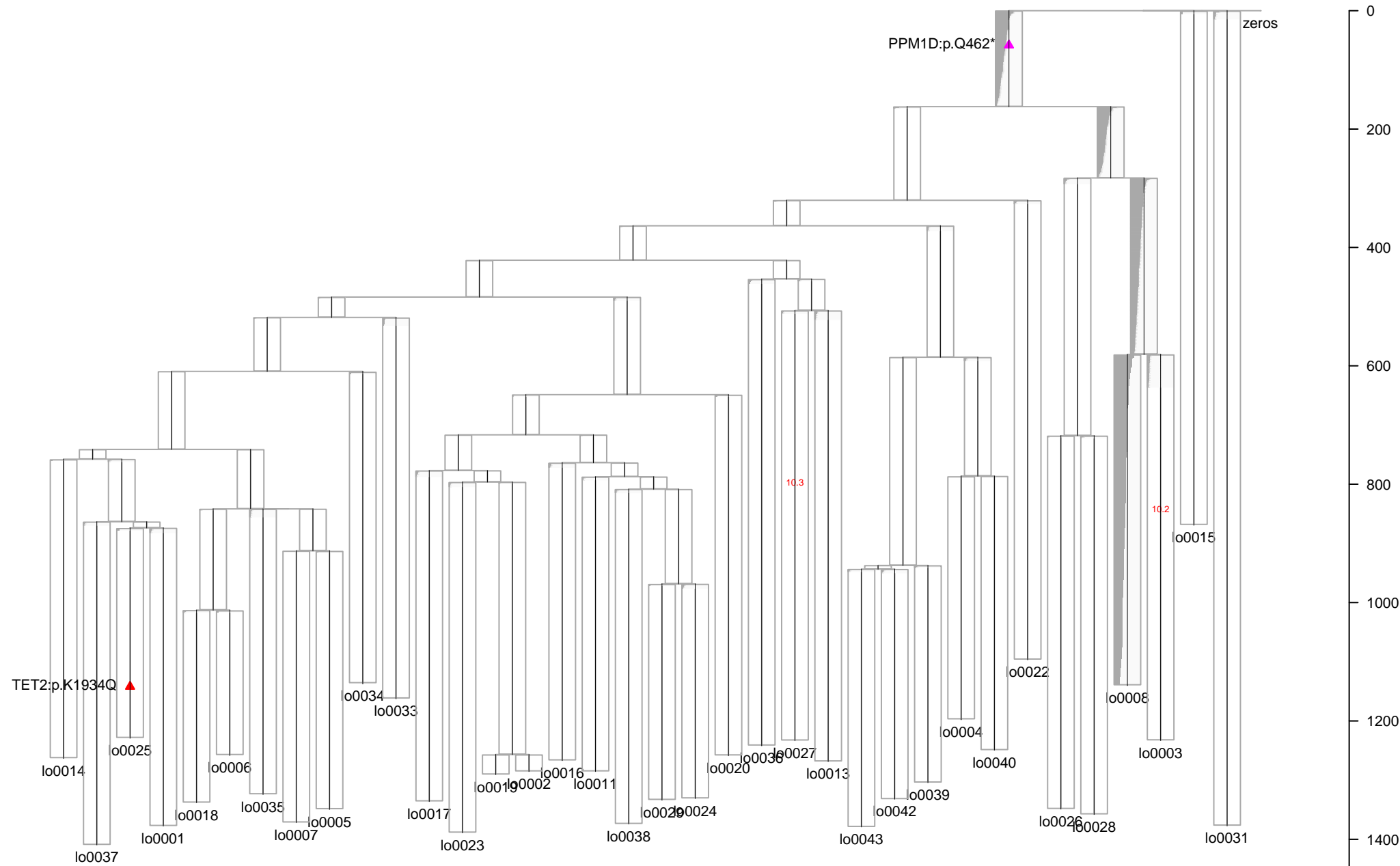
PD6634: Annotated with VAF from lo0026
Mean Depth=14.18



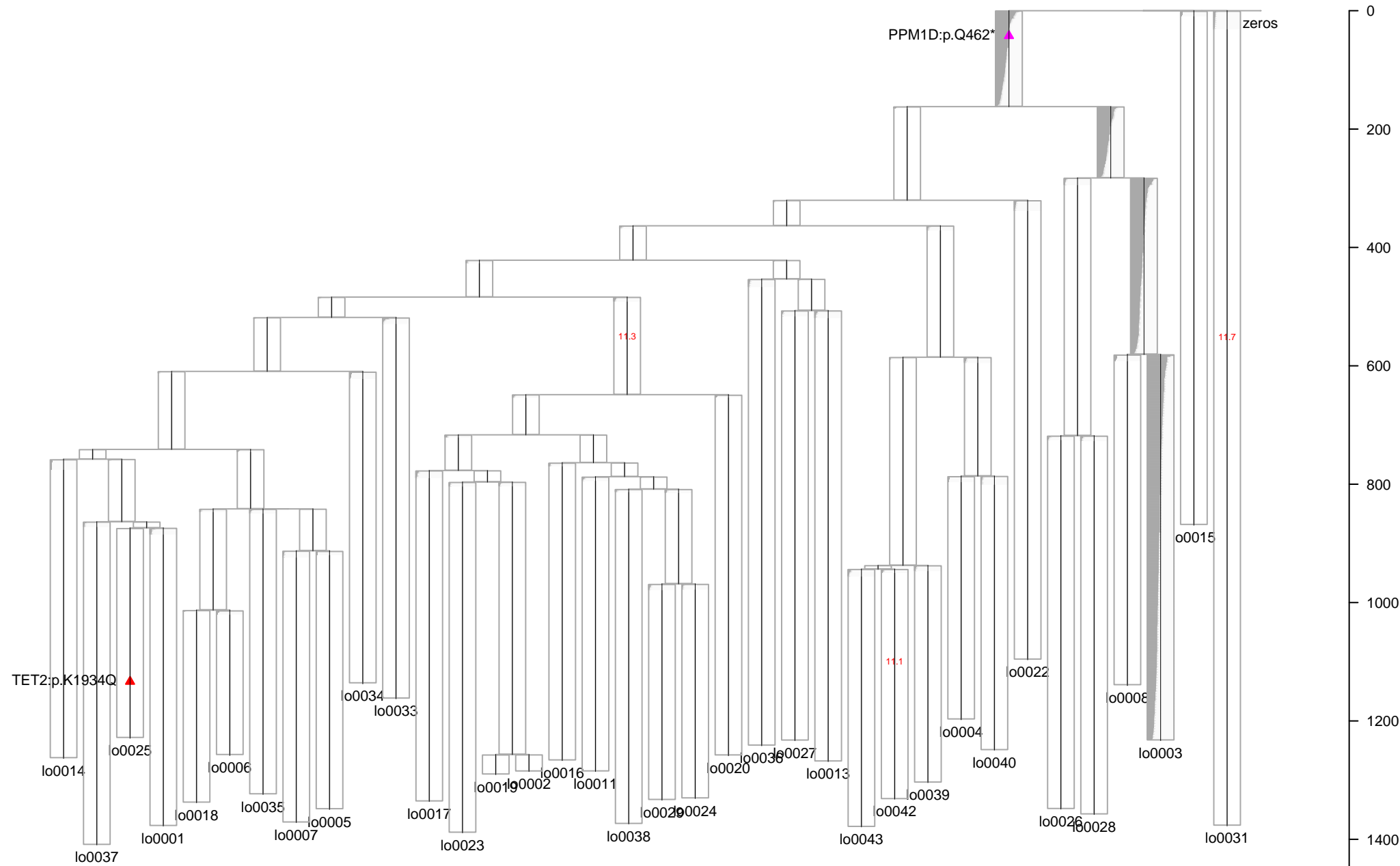
PD6634: Annotated with VAF from lo0028
Mean Depth=13.70



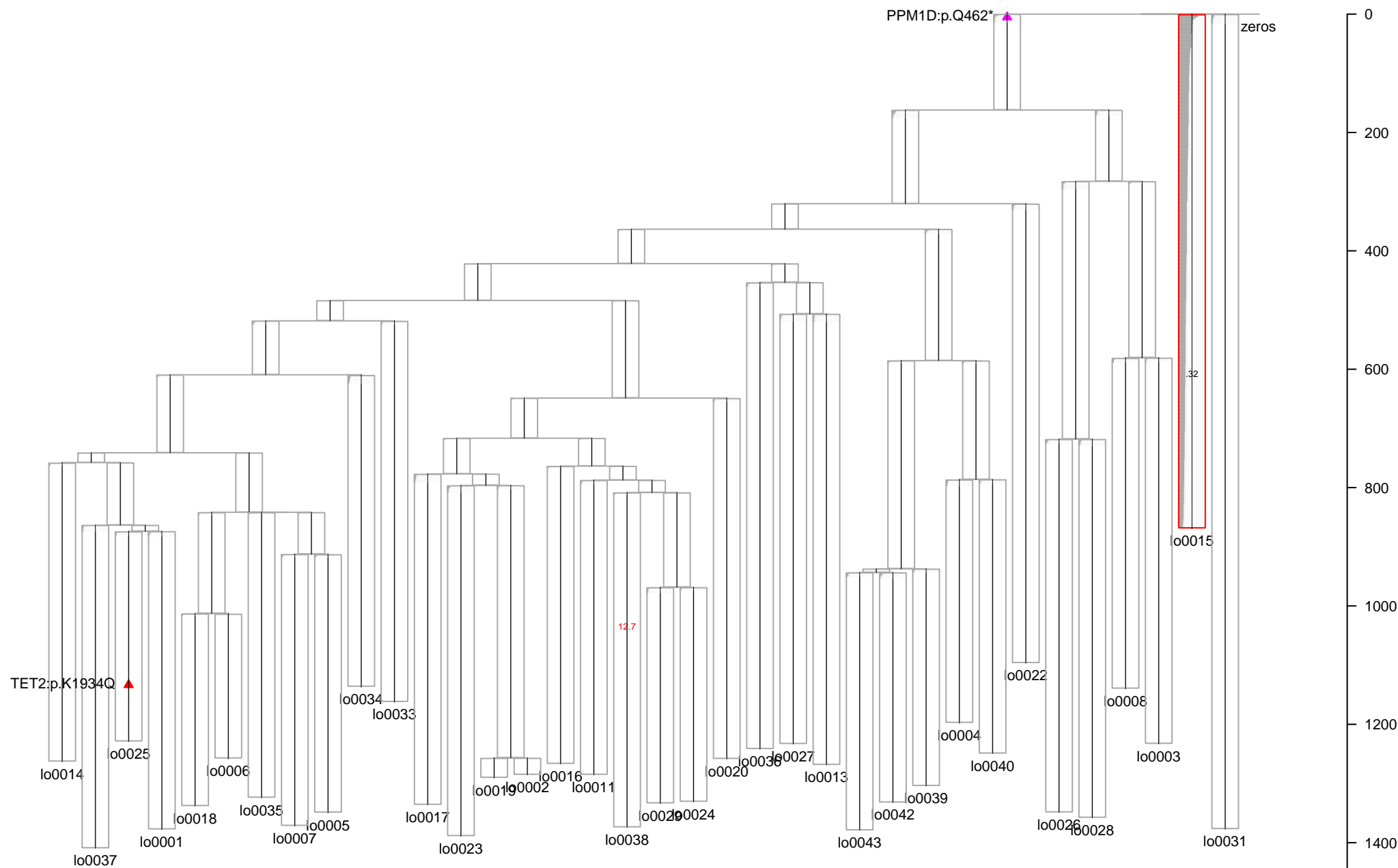
PD6634: Annotated with VAF from lo0008
Mean Depth=10.67



PD6634: Annotated with VAF from lo0003
Mean Depth=12.04



PD6634: Annotated with VAF from lo0015
Mean Depth=13.20



PD6634: Annotated with VAF from lo0031
Mean Depth=16.30

