

37 Participants enrolled

D0 samples, Pv (+)*: n = 37
ssPCR(+): n = 35

2-year follow up after treatment

Month 1: weekly
Month 2-4: monthly

No Pv recurrences: 9 participants

Pv recurrences: 28 participants

Total episodes, Pv (+): n = 142
ssPCR(+): n = 140

1st year (n=66)
2nd year (n=76)

1 Pv sample was not
analysed by MS genotyping

MS genotyping

D0 + recurrences
n = 174 Pv samples

Overall genetic diversity analysis (n=136)

Monoclonal/complete allele data (n=125)

Monoclonal/incomplete allele data (n=11)

Polyclonal/complete+incomplete (n=28)

Negative PCR amplification (n=10)

Identified
haplotypes
(n=162)