

Genotyping efficiency (%)

100.0  
10.0  
1.0  
0.1

RHP targeted sequencing

Mutation identification

WES/WGS

U2AF1.S34F

IDH2.R140Q

TET2.Q1537\*

EZH2.P132L

ETV6.R360W

TET2.Q876\*

TET2.Q1547\*

TET2.R1216\*

TET2.S792

TET2.H1380Y

ASXL1.R693\*

ASXL1.V507fs

TET2.R1216\*

ASXL1.G642fs

ZRSR2.R126\*

TET2.Q1537\*

ASXL1.L697fs\*

ASXL1.L697fs

RPS24.chr10:79795273:T/C

RAB9A.3pUTR

MTAP.rearr

HNRNPUL1.F559F

MTAP.rearr

MALAT1.chr11:65270399:G/A

NFIC.chr19:3462800:G/A

FAM138A.chr10:133133347:G/A

HNRNPUL1.F559F:G/A

PRKDC.chr8:48713410:C/T

CEP350.chr10:133133347:G/A

SF3A1.chr19:3462800:G/A

SMARCC1.chr15:47627735:G/A

NOTCH1.chr9:139389423:C/T

MAP4K5.P667S

SMARCD1.chr12:50490729:C/T

DOLPP1.R227S

MALAT1.chr11:65270399:G/A