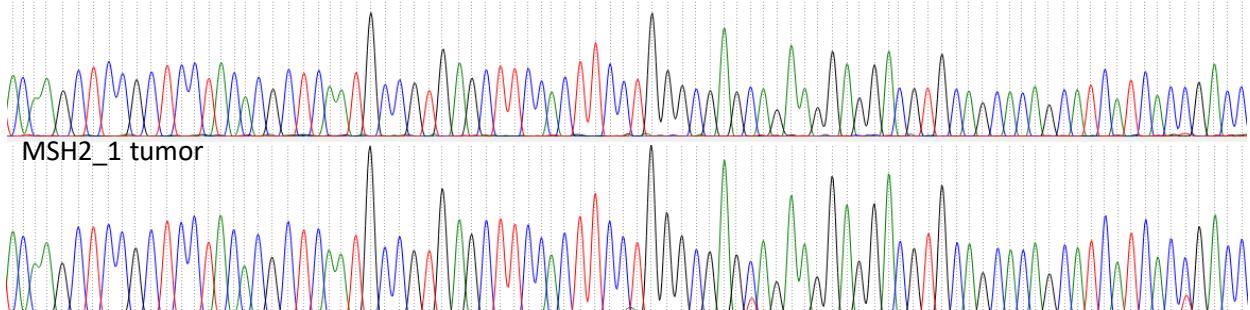


A**Healthy control**

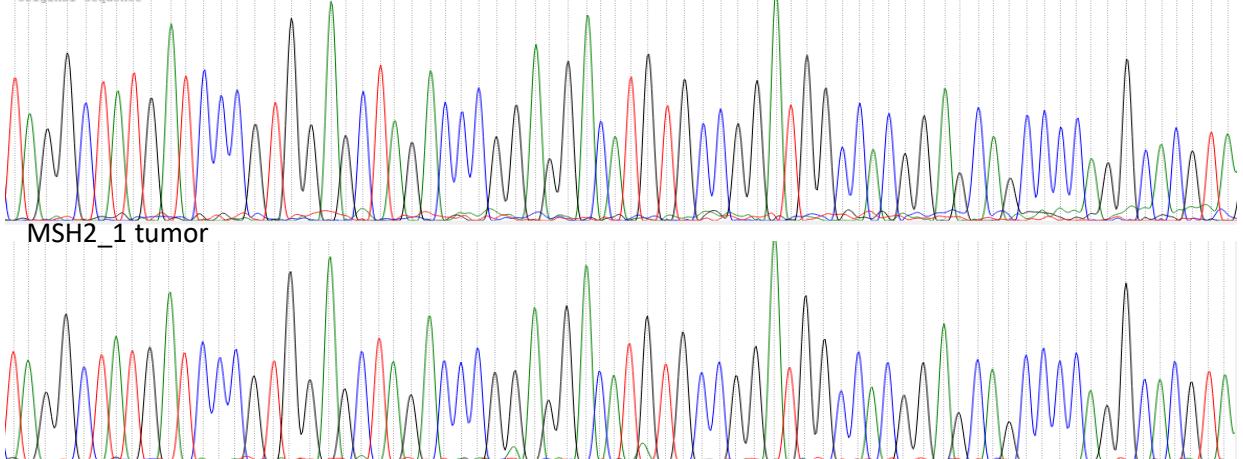
ACAA G C T C C G C T C C T A C A C G C T C A A T G C C G T G A G C T T C C A C T T C C T G G G C G A G C A G A A G G A G G A C G T G C A G C A C A G C A T C A T C A C C G A C C
Original sequence:



POLD1
p.Gln485Ter
c.1453C>T
gnomAD: -

B**Healthy control**

T A G G C T A T G A T C C C G T G G G A G C T A G A C C C G G A G G A C A T G T G C C C G G A T G G C C A C G G A G C A G C C C C A G G C A C G T A
Original sequence:



POLE
p.Glu442Lys
c.1324G>A
gnomAD: -

POLE
p.Met444Ile
c.1332G>A
gnomAD: -