

## Figure legends

Supplemental Figure 1. **Absolute neutrophil counts of the patient over time.** Grey area indicates reference values, red asterisks indicate infection episodes. Blue line indicates administration of G-CSF.

Supplemental Figure 2. **FNIP1 –disease causing mutations.** A) Shown is the Integrative Genomics Viewer (1) analysis mapping the read depth of the trio-whole exome sequencing data across the genomic locus surrounding the *FNIP1* gene. Violet line indicates the location of the de novo deletion. B) Representation of described FNIP1 - disease causing variants. Bold indicates homozygosity; compound heterozygous mutations are marked in the same colours (blue and red), mutation associated with UPD is represented in *italics*.

## References

1. Robinson J, Thorvaldsdóttir H, Winckler W, Guttman M, Lander E, Getz G, Mesirov JP. Integrative genomics viewer. *Nat Biotechnol.* 2011 Jan;29(1):24-26.