

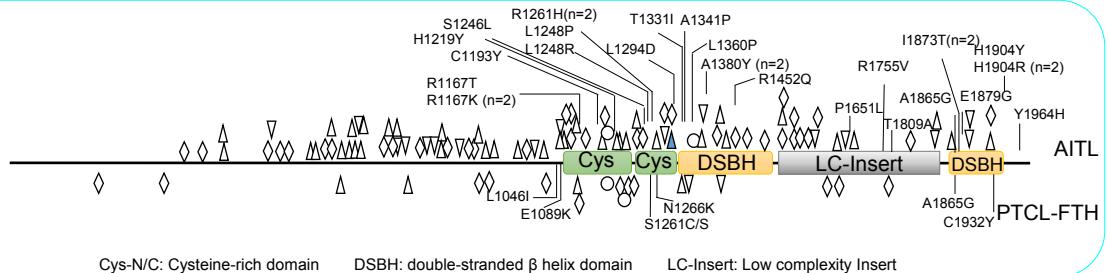
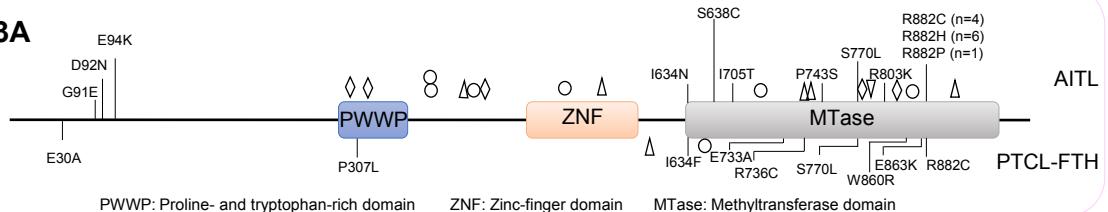
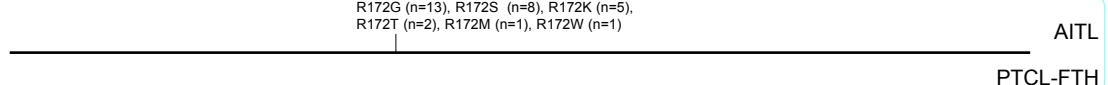
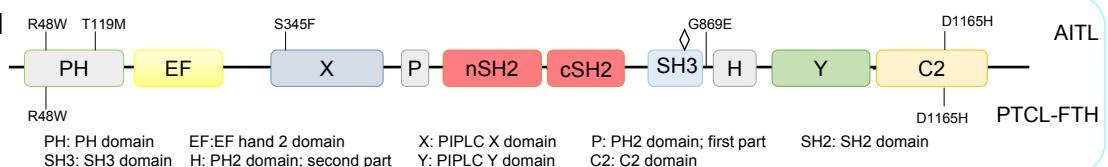
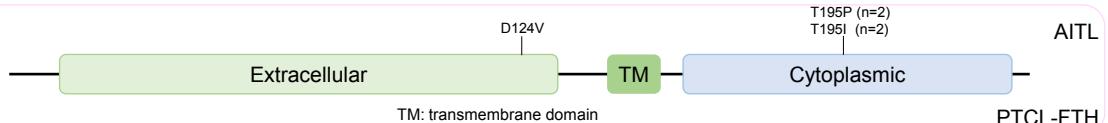
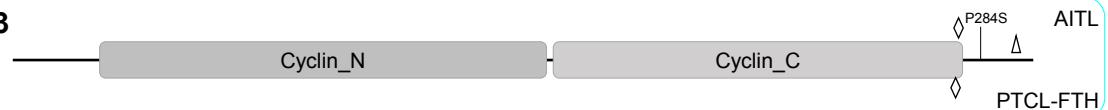
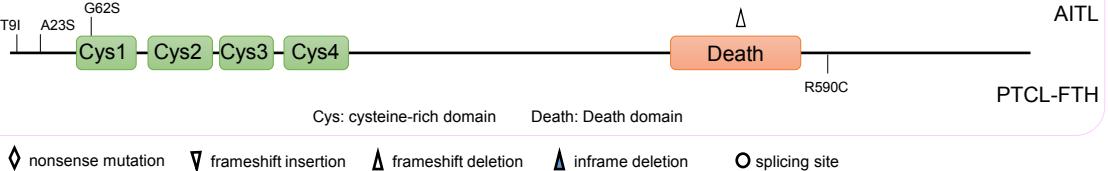
TET2**DNMT3A****IDH2****RHOA****PLCG1****CD28****CCND3****TNFRSF21**

Figure S1. Nature and distribution of mutations in *TET2*, *DNMT3A*, *IDH2*, *RHOA*, *PLCG1*, *CCND3* and *TNFRSF21* in AIHL and PTCL-FTH.