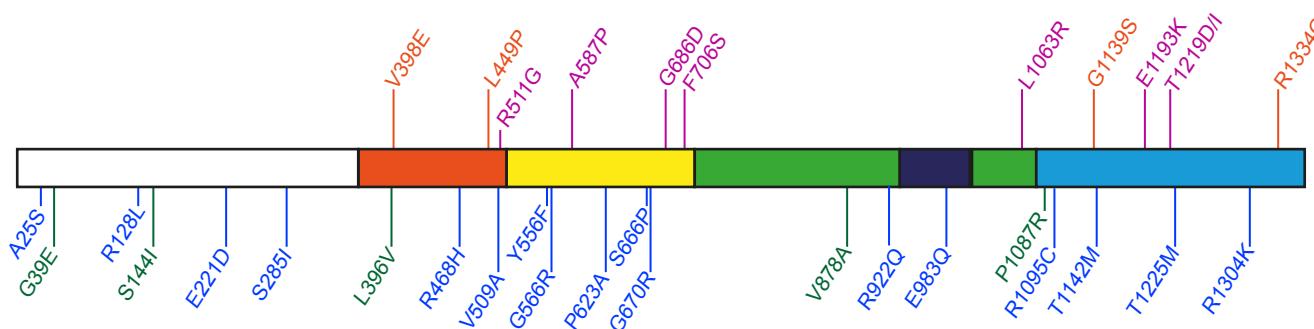


Detected (Pathogenic)



- Mismatch binding domain (aa 362-518)
- Connector domain (aa 519-717)
- Lever domain (aa 718-934 and 1009-1076)
- Clamp domain (aa 935-1008)
- ATPase domain (aa 1077-1360)