Alteration **Mutation Taster** SIFT PolvPhen-2 Probably Damaging (.999)

NT*

NT

NT

NT

NT

Probably Damaging (1.000)

Probably Damaging (1.000)

Probably Damaging (.999)

Probably Damaging (.999)

Supplementary Table 1. Predicted Effects of the Alterations in *HUWE1* on Protein Function

Disease-causing

Disease-causing

Disease-causing

**Three missense mutations reported by Froyen et al.[22] and Isrie et al.[23]

p.R4063Q c.12188G>A Disease-causing Disease-causing p.G4310R c.12928G>C

c.12037C>T

c.12559C>T

c.8942G>A

p.R4013W**

p.R4187L**

p.R2981H**

*NT- Not Tolerated