Table 1. Points awarded per in *trans* occurrence

Classification/Zygosity of other variant[1]	Points per Proband	
	Confirmed in *trans*	Phase unknown
Pathogenic or Likely pathogenic variant	1.0	0.5 (P) 0.25 (LP)
Homozygous occurrence *(max point 1.0)*	0.5	N/A
Uncertain significance variant on other allele *(max point 0.5)*	0.25	0.0

[1]All variants should be sufficiently rare (meet PM2 specification)

Table 2. Recommendation for determining the appropriate ACMG/AMP evidence strength level for PM3

PM3_Supporting	PM3	PM3_Strong	PM3_VeryStrong
0.5	1.0	2.0	4.0