

Table 1. Points awarded per in *trans* occurrence

Classification/Zygotity of other variant ¹	Points per Proband	
	Confirmed in <i>trans</i>	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

¹All 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