

Title: Supplementary Data 1:

Description: Details of all rare variants curated to be pathogenic or likely pathogenic in ACMG SF 2.0 genes in the MGRB cohort.

Title: Supplementary Data 2:

Description: Common variant score analysis supplementary data.

Title: Supplementary Data 3:

Description: Putative incidental somatic variants identified in the MGRB.

Title: Supplementary Data 4:

Description: Resources used in this work.

Title: Supplementary Data 5:

Description: Development and derivation of the somatic SNV detector.