

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Clinical information of the cohort. A total of 191 CMT in the study cohort are shown with available clinical information.

File Name: Supplementary Data 2

Description: Sequencing information of WES. The number of total reads sequenced and the mapping rates are shown.

File Name: Supplementary Data 3

Description: Sequencing information of RNA-seq.

File Name: Supplementary Data 4

Description: Exonic mutations of 183 CMT. A total of 10,855 exonic SNVs and indels are listed with genomic coordinates based on CanFam3.1.

File Name: Supplementary Data 5

Description: Germline variants of CMT genomes.

File Name: Supplementary Data 6

Description: Mutation frequencies of genes in CMT and human breast cancers. The frequency (%) of non-silent mutations in benign-malignant CMTs are shown with those of TCGA breast cancers.