



Supplementary Figure S3 Control single nucleotide polymorphism analysis for HM fibroblasts and hiPSCs as well as single nucleotide polymorphism analysis of KS fibroblasts. (A) Number of transcripts with or without single nucleotide polymorphism (SNP) variation in HM fibroblast and hiPSC. **(B)** Number of transcripts with SNP variation, with or without Y chromosome homologs. **(C)** Number of transcripts with or without SNP variation in KS and HF fibroblasts. Results combined from three biological replicates. (See also Fig. 4, Supplementary Table SVIII)