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**S3 Fig. Prostate cancer CSPRS distributions before (A) and after (B – F) defining global risk variants.** Five sets of global variants were defined as variant whose allele frequency differences between the four ancestry groups within the 1000 Genomes Project reference were below 25% (B), 20% (C), 15% (D), 10 % (E) and 5 % (F). Red lines indicate 10% quantiles of the corresponding UKB PRS distribution in all controls. Abbreviations: AFR: African; EAS: East Asian; EUR: European, SAS: South Asian