

Figure S2

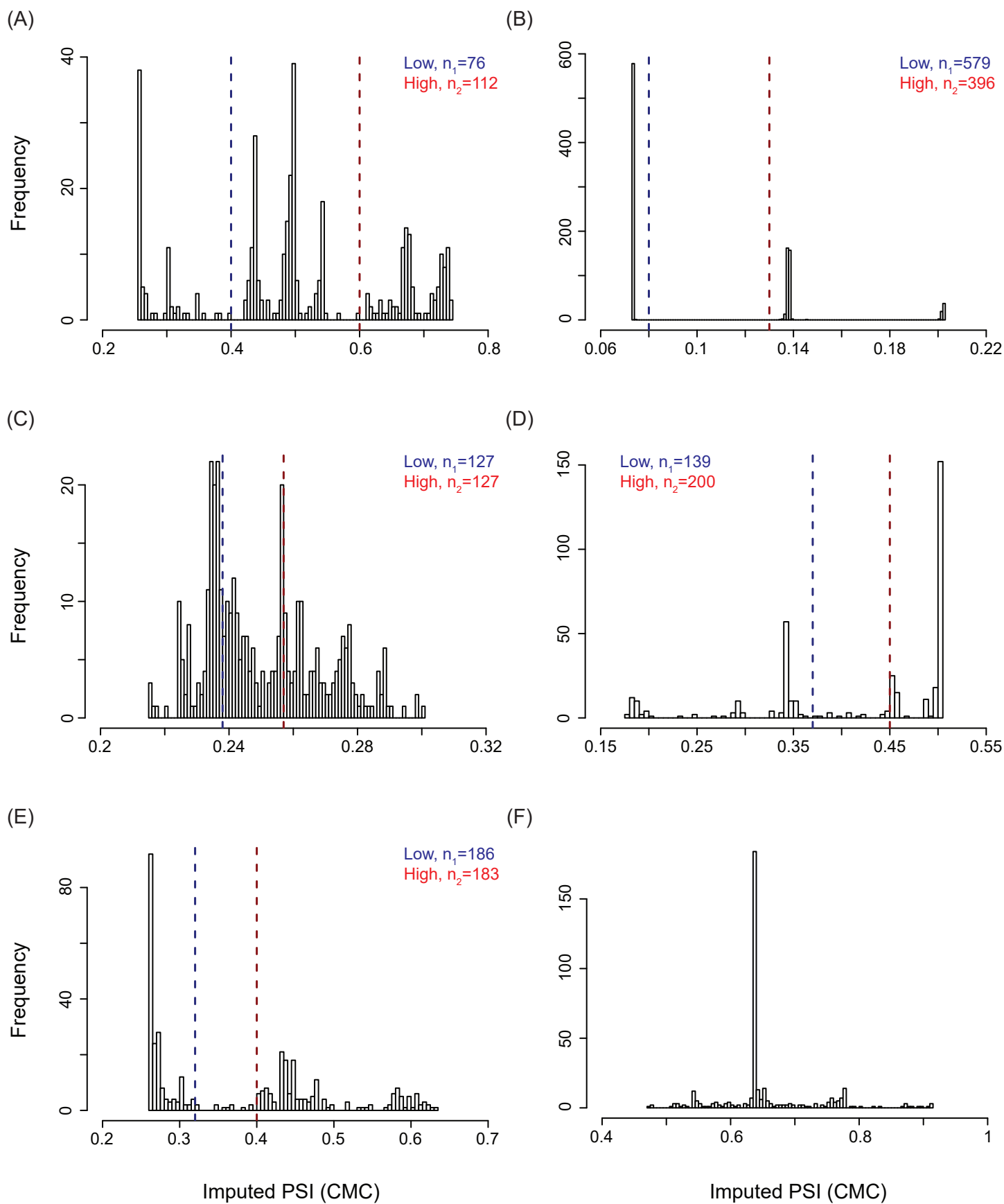


Figure S2. Sample stratification of the six causal splicing events based on cis-regulated PSI.

CMC samples with genetically imputed PSI values less than the level marked by the blue dashed line (n_1) were labeled as low, and those greater than the level marked by the red dashed line (n_2) were labeled as high. Intervening samples (n_0) with PSI values between the high and low were unused.

(A) ENSE00002438745 in *LINC00665*. $n_1 = 76$, $n_2 = 112$, $n_0 = 192$.

(B) ENSE00001875548 in *NSUN4*. $n_1 = 579$, $n_2 = 396$, $n_0 = 0$.

(C) ENSE00002674786 in *SRRM2*. $n_1 = 127$, $n_2 = 127$, $n_0 = 126$.

(D) ENSE00002079807 in *ELOVL7*. $n_1 = 139$, $n_2 = 200$, $n_0 = 41$.

(E) ENSE00003572542 in *DRC1*. $n_1 = 186$, $n_2 = 183$, $n_0 = 11$.

(F) ENSE00001693995 in *TBC1D5*. Samples on the high and low ends are not of sufficient numbers.