Gene expression differences between matched pairs of ovarian cancer patient tumors and patient-derived xenografts

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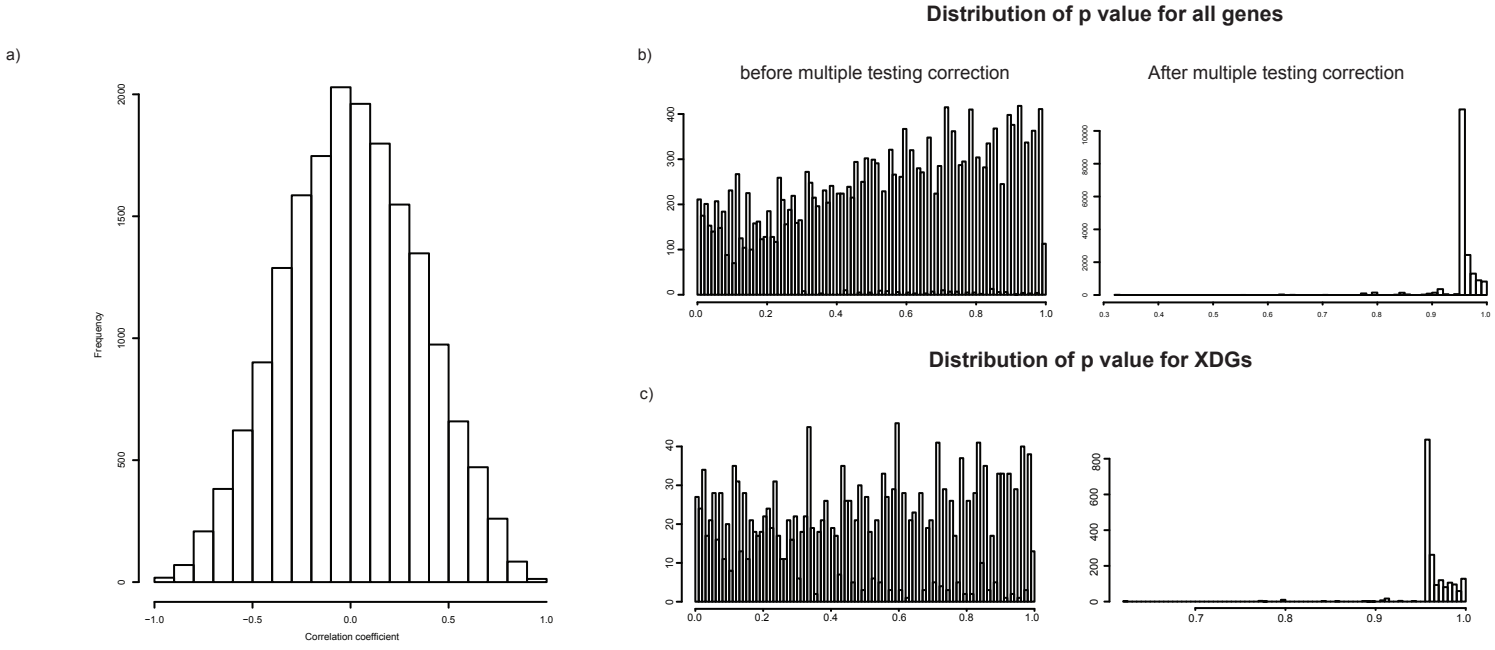

Figure S1. (a) Distribution of correlation coefficients between PDX passage and gene expression. (b) Distribution of p-values associated with correlation coefficients for all genes. (c) Distribution of p-values associated with correlation coefficients for XDGs.

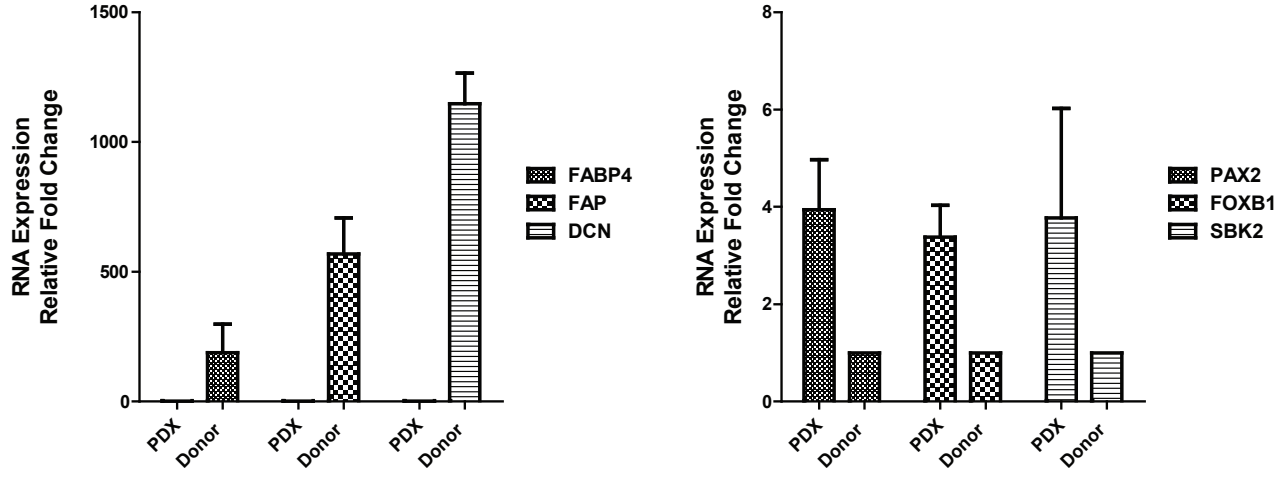

Figure S2. RT-qPCR validation of (a) three up-XDGs and (b) three down-XDGs. Data presented as relative expression fold change between PDX and Donor tumor. Error bars represent standard error of the mean (n = 3). Our qRT-PCR results indicate that expression of these 6 genes is consistent with our RNA-seq results.

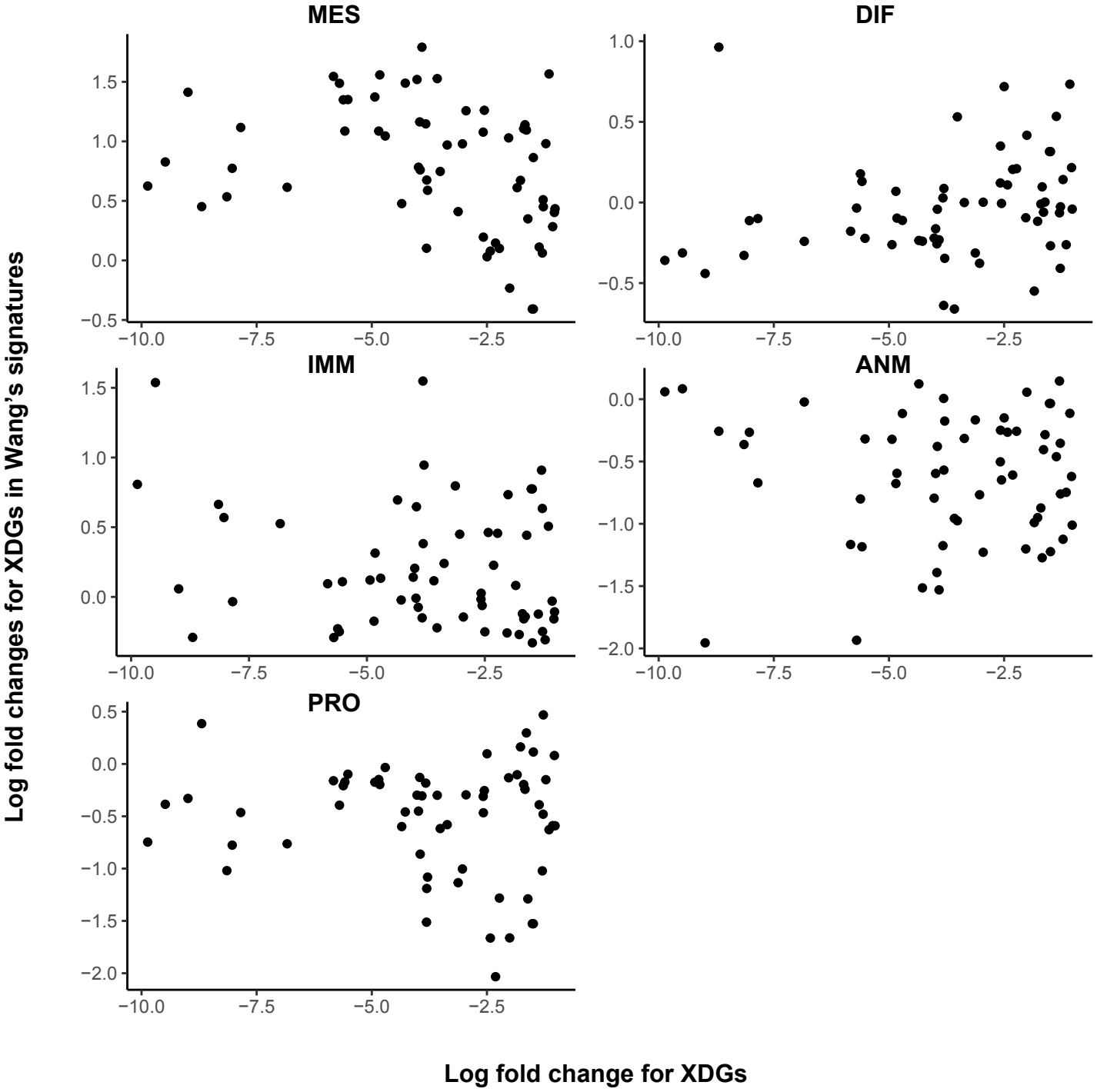

Figure S3. Comparison of XDGs with ovarian cancer subtype signatures defined by Wang. x axis indicates log2 fold change of XDGs for PDX/donor comparison. y axis indicates log2 fold change of XDGs for ovarian cancer subtype signatures defined by Wang.

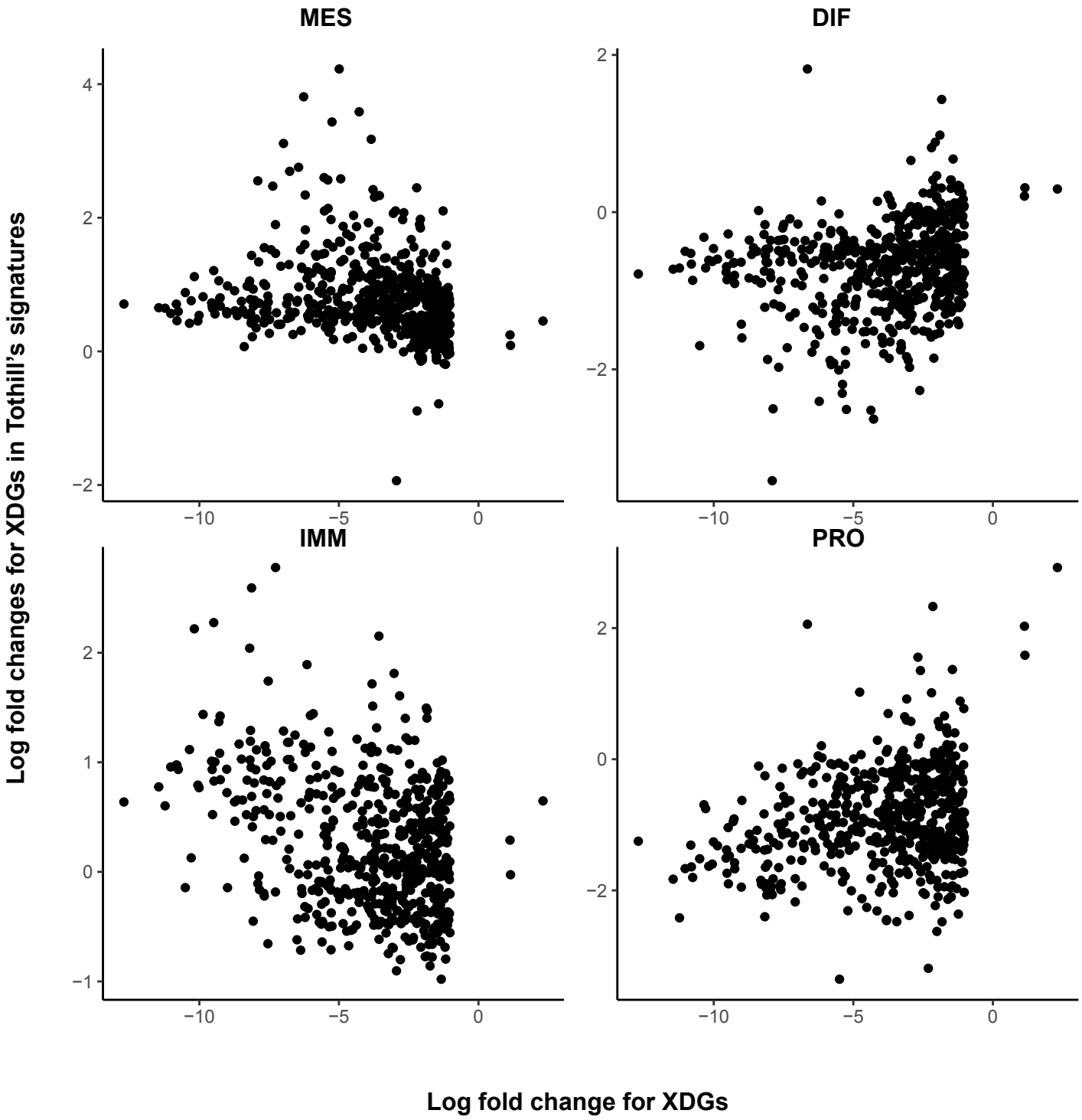

Figure S4. Comparison of XDGs with ovarian cancer subtype signatures defined by Tothill. x axis indicates log2 fold change of XDGs for PDX/donor comparison. y axis indicates log2 fold change of XDGs for ovarian cancer subtype signatures defined by Tothill.

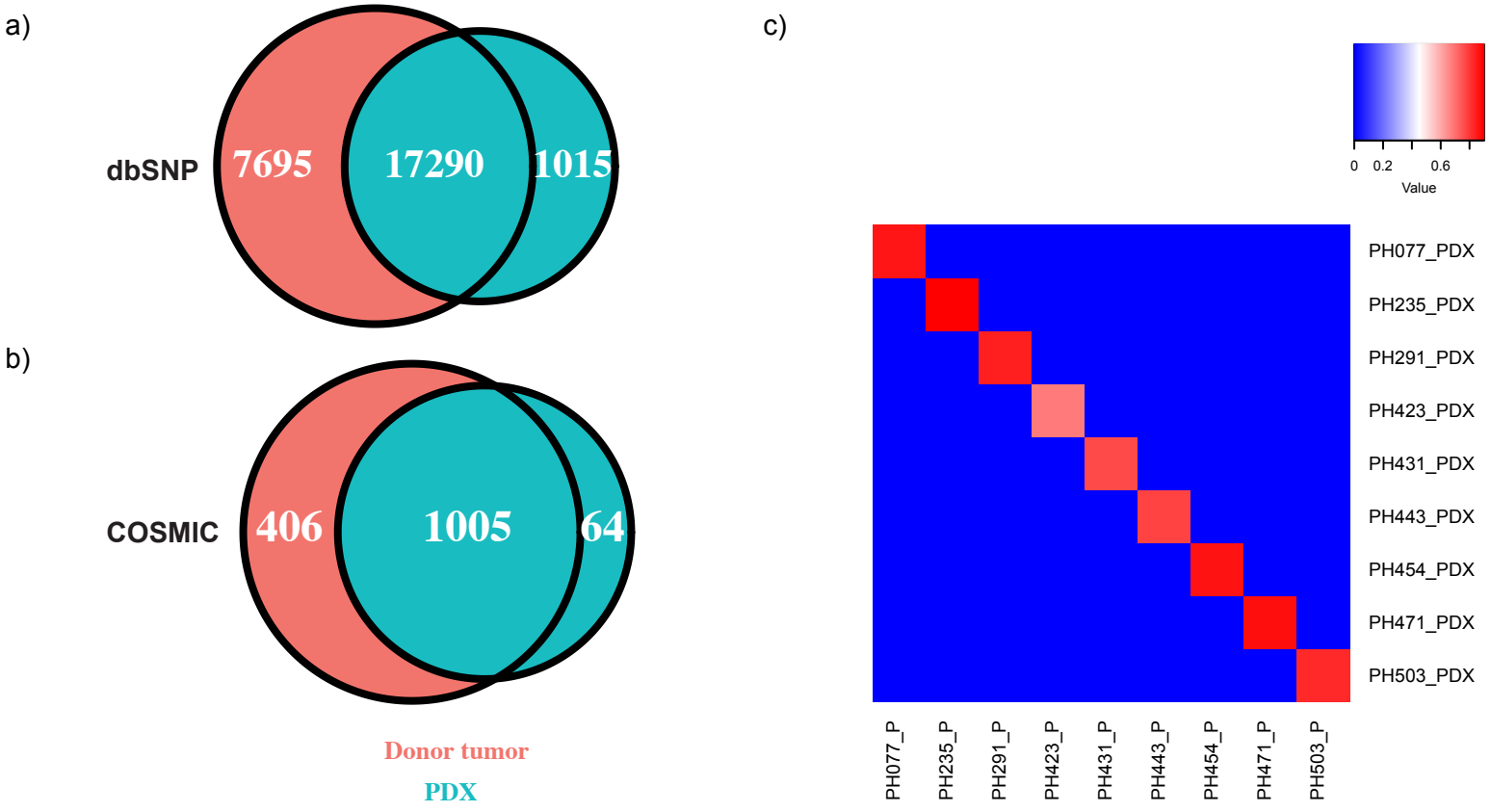

Figure S5: Venn diagram comparing overlap of a) dbSNP annotated; b) COSMIC annotated variants between donor tumor and PDX for PH503; c) Genotype concordance between donor tumor/PDX pairs examined using NGSCheckMate.