

## Supplementary Files

Table S1| The true positive rate (TP), the false positive rate (FP), the false discovery rate (FDR), the average number of false positive in the final model (ANFP), model size (Size), and the area under the curve (AUROC), the number of filtered truly important variable from FDR and RF (# filter) under the low correlation coefficients,  $\rho=0$  for the 500 simulations.

screening	method	TP	FP	FDR	ANFP	Size	AUROC	# filter
$FDR_{.05}$ – ISIS	LASSO	0.2348 (0.00233)	2e-06 (0)	0.02917 (0.00277)	0.204 (0.01912)	6.074 (0.063)	0.81701 (0.00165)	6.074 (0.063)
	SCAD	0.2348 (0.00233)	2e-06 (0)	0.02917 (0.00277)	0.204 (0.01912)	6.074 (0.063)	0.81701 (0.00165)	
	MCP	0.2348 (0.00233)	2e-06 (0)	0.0283 (0.00274)	0.198 (0.01893)	6.068 (0.06292)	0.81701 (0.00165)	
$FDR_{.20}$ – ISIS	LASSO	0.28056 (0.00259)	1.28e-05 (0)	0.13122 (0.00566)	1.284 (0.06609)	8.298 (0.10205)	0.82968 (0.00165)	8.298 (0.10205)
	SCAD	0.28056 (0.00259)	1.28e-05 (0)	0.13092 (0.00565)	1.28 (0.066)	8.294 (0.10184)	0.82965 (0.00165)	
	MCP	0.28056 (0.00259)	1.23e-05 (0)	0.12636 (0.00568)	1.234 (0.06602)	8.248 (0.10092)	0.82978 (0.00165)	
$RF_{1000}$ – ISIS	LASSO	0.45512 (0.00257)	0.000116 (0)	0.5053 (0.00279)	11.622 (0.06418)	23 (0)	0.84898 (0.00158)	14.392 (0.06546)
	SCAD	0.45728 (0.00261)	0.000115 (0)	0.50296 (0.00283)	11.568 (0.06514)	23 (0)	0.84888 (0.00157)	
	MCP	0.46192 (0.00269)	0.000114 (0)	0.49791 (0.00293)	11.452 (0.06735)	23 (0)	0.84956 (0.00157)	

(): standard deviation

Table S2| The true positive rate (TP), the false positive rate (FP), the false discovery rate (FDR), the average number of false positive in the final model (ANFP), model size (Size), and the area under receiver operating characteristic curve (AUROC), the number of filtered truly important variable from FDR and RF (# filter) under the low correlation coefficients,  $\rho= 0.4$  for the 500 simulations.

screening	method	TP	FP	FDR	ANFP	Size	AUROC	# filter
<i>FDR</i> <sub>.05</sub> – <i>ISIS</i>	LASSO	0.19448 (0.00149)	1.8e-06 (0)	0.02953 (0.00327)	0.184 (0.0211)	5.046 (0.04413)	0.76712 (0.00159)	5.052 (0.0448)
	SCAD	0.1936 (0.00149)	1.8e-06 (0)	0.02953 (0.00327)	0.184 (0.0211)	5.024 (0.0443)	0.76702 (0.00159)	
	MCP	0.19088 (0.00154)	1.8e-06 (0)	0.02973 (0.00329)	0.184 (0.0211)	4.956 (0.04561)	0.76658 (0.00159)	
<i>FDR</i> <sub>.20</sub> – <i>ISIS</i>	LASSO	0.22728 (0.00188)	9.9e-06 (0)	0.12385 (0.00588)	0.99 (0.05595)	6.672 (0.07893)	0.77586 (0.00165)	6.690 (0.07975)
	SCAD	0.2248 (0.00179)	9.9e-06 (0)	0.12493 (0.00591)	0.99 (0.05595)	6.61 (0.07741)	0.77574 (0.00165)	
	MCP	0.22152 (0.00185)	9.9e-06 (0)	0.12579 (0.00597)	0.988 (0.05591)	6.526 (0.07863)	0.77527 (0.00165)	
<i>RS</i> <sub>1000</sub> – <i>ISIS</i>	LASSO	0.3428 (0.00246)	0.0001442 (0)	0.62705 (0.00269)	14.412 (0.06232)	22.982 (0.00595)	0.76657 (0.00184)	11.876 (0.07313)
	SCAD	0.33832 (0.0024)	0.0001452 (0)	0.63175 (0.00263)	14.514 (0.06108)	22.972 (0.00739)	0.76576 (0.00185)	
	MCP	0.33496 (0.00236)	0.000146 (0)	0.63542 (0.00258)	14.598 (0.06007)	22.972 (0.00739)	0.76553 (0.00184)	

( ): standard deviation

Table S3| Average true regression coefficients for the 25 variables

Variable name	True regression coefficient
g01	1.12939
g02	0.28505
g03	-1.45882
g04	0.12632
g05	-0.35443
g06	-1.01840
g07	1.65116
g08	-0.76401
g09	0.84540
g010	-0.69254
g11	-0.07449
g12	0.96375
g13	0.23261
g14	-0.42302
g15	0.47535
g16	-0.12398
g17	-0.33047
g18	-0.87834
g19	1.57071
g20	-1.23074
g21	0.73234
g022	-0.55067
g23	1.17932
g24	0.96817
g25	-0.72762

Table S4| AUROC scores, the corresponding confidence interval, and standard error for LASSO, SCAD, and MCP methods on six different screening scenarios with prostate cancer data.

Method		AUROC	Confidence Interval		S.E
Screening	Variable Selection		lower	upper	
$FDR_{0.01}$ – <i>ISIS</i>	LASSO	0.74622	0.67467	0.81777	0.03650
	SCAD	0.74585	0.67429	0.8174	0.03651
	MCP	0.76437	0.69508	0.83366	0.03535
$FDR_{0.05}$ – <i>ISIS</i>	LASSO	0.69712	0.61929	0.77495	0.03971
	SCAD	0.71405	0.63739	0.79072	0.03911
	MCP	0.68335	0.60335	0.76335	0.04081
$FDR_{0.20}$ – <i>ISIS</i>	LASSO	0.70906	0.63178	0.78634	0.03943
	SCAD	0.68944	0.61178	0.76711	0.03962
	MCP	0.71539	0.63801	0.79278	0.03948
$RF_{1000}$	LASSO	0.73586	0.66149	0.81024	0.03794
	SCAD	0.73757	0.66269	0.81245	0.03820
	MCP	0.73757	0.66269	0.81245	0.03820
$RF_{2000}$	LASSO	0.73684	0.66227	0.81140	0.03804
	SCAD	0.70626	0.62894	0.78357	0.03944
	MCP	0.71929	0.64044	0.79815	0.04023
$RF_{4000}$	LASSO	0.69273	0.61330	0.77216	0.04052
	SCAD	0.72100	0.64440	0.79759	0.03907
	MCP	0.69480	0.61450	0.77511	0.04097

S.E was estimated by bootstrapping approach.

Figure S1| Selection frequencies of each of the 25 variables across the LASSO, the SCAD, and the MCP during 500 simulations with  $\rho = 0$ . The x-axis depicts the names of the variables, and the y-axis is the frequency of variables selected out of 500 simulations. The variables not depicted on the x-axis in Figure 2 did not have any counts. Each of the three methods is identified by color in legend.

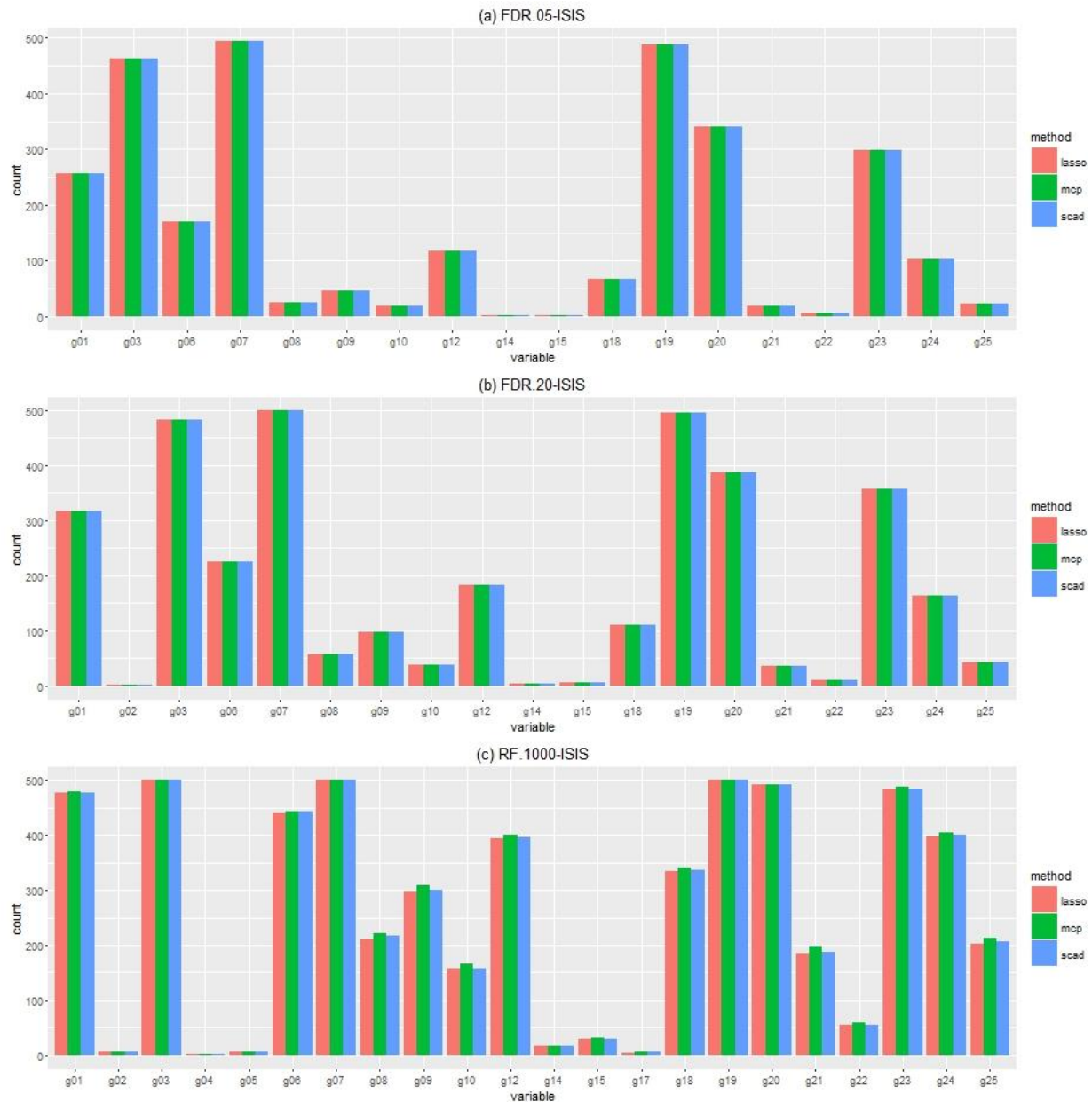


Figure S2| Selection frequencies of each of the 25 variables across the LASSO, the SCAD, and the MCP during 500 simulations with  $\rho = 0.4$ . The x-axis depicts the names of the variables, and the y-axis is the frequency of variables selected out of 500 simulations. The variables not depicted on the x-axis in Figure 2 did not have any counts. Each of the three methods is identified by color in legend.

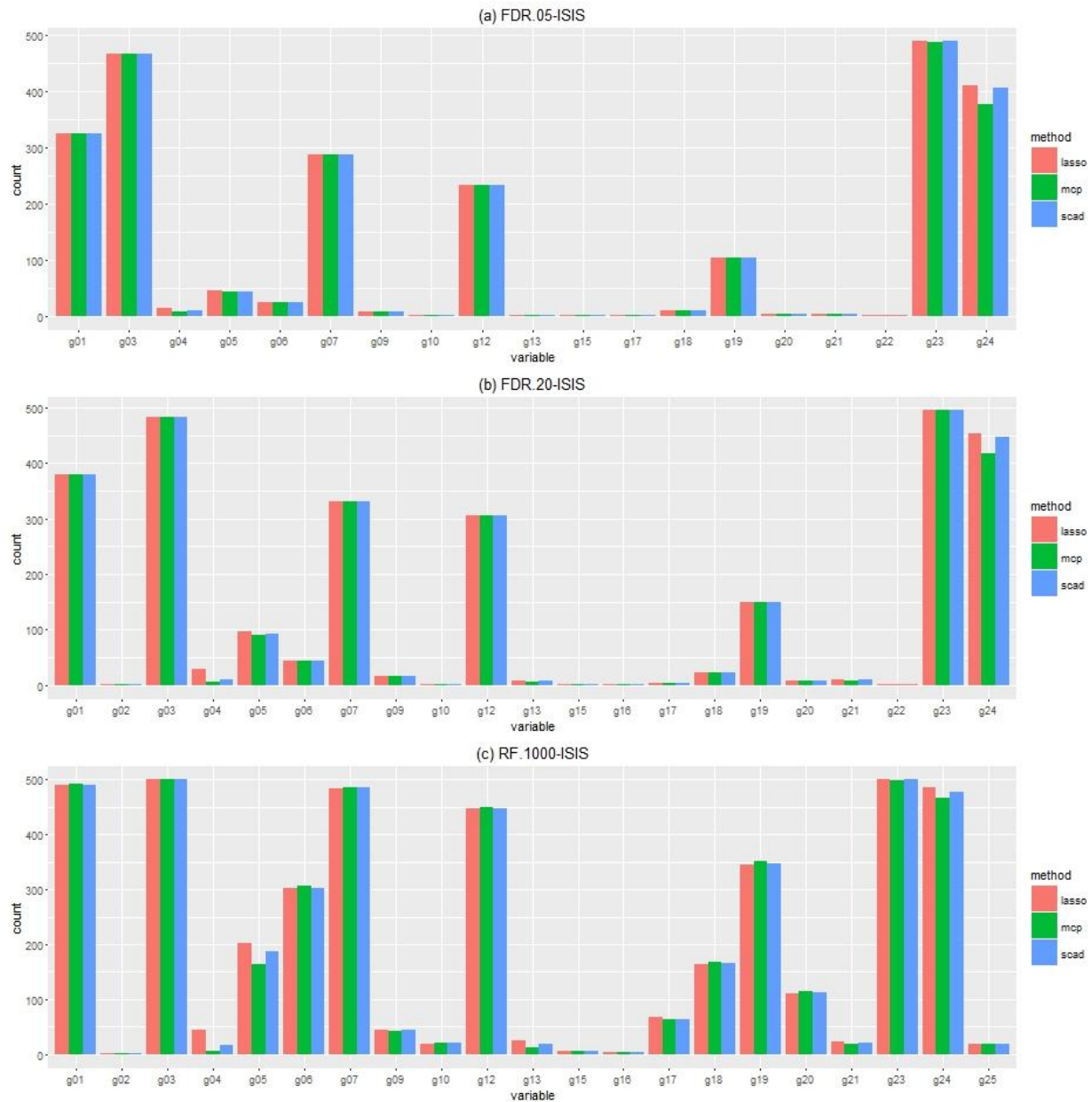


Figure S3| First column is the box plots of the AUROC scores under  $\rho = 0$ . x-axis is the name of variable selection methods ,and y-axis is the AUROC scores. Second column is the box plots of the corresponding false discovery rate, and y-axis the false discovery rate during 500 simulations.

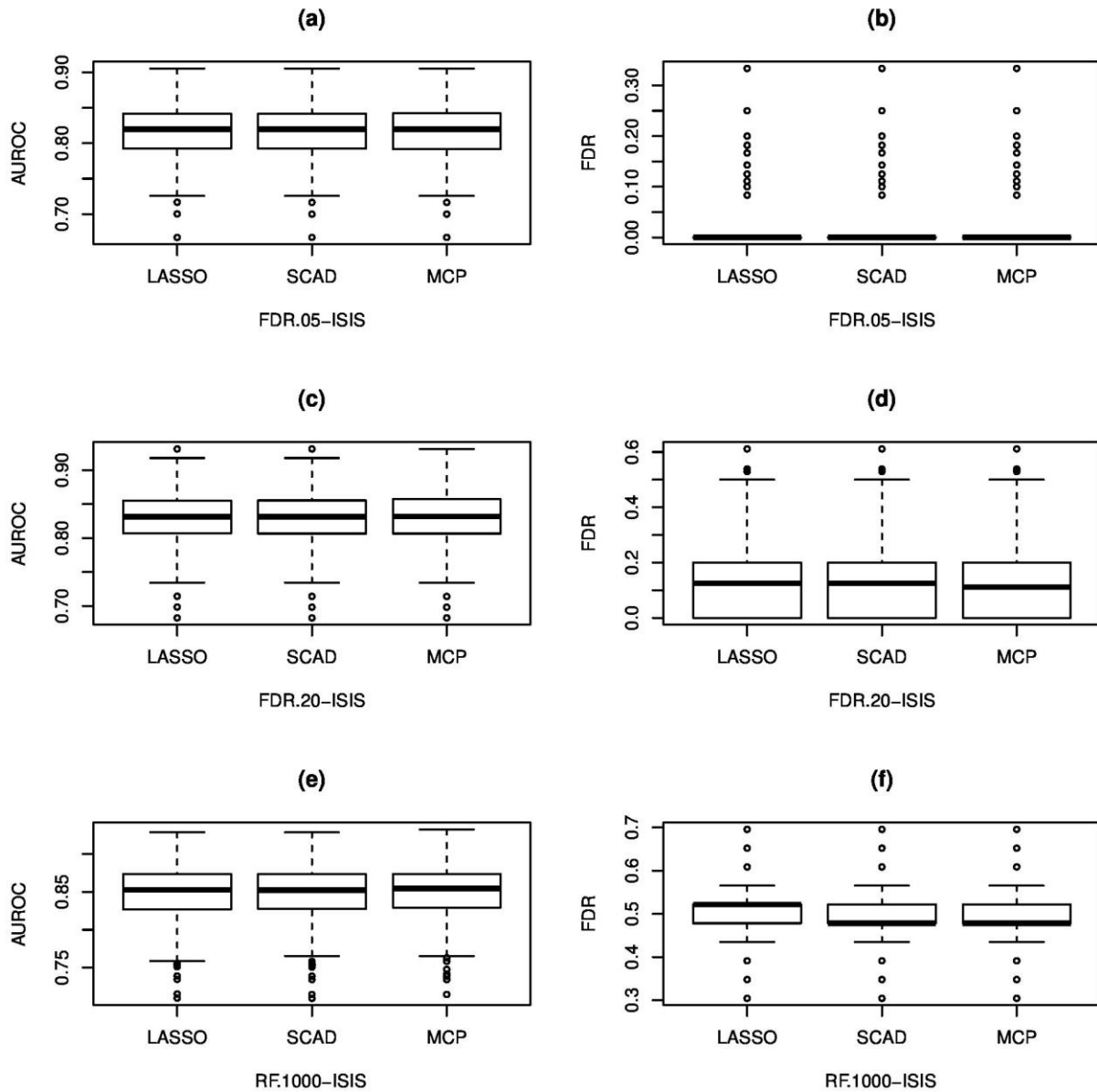


Figure S4| First column is the box plots of the AUROC scores under  $\rho = 0.4$ . x-axis is the name of variable selection methods ,and y –axis is the AUROC scores. Second column is the box plots of the corresponding false discovery rate, and y-axis the false discovery rate during 500 simulations.

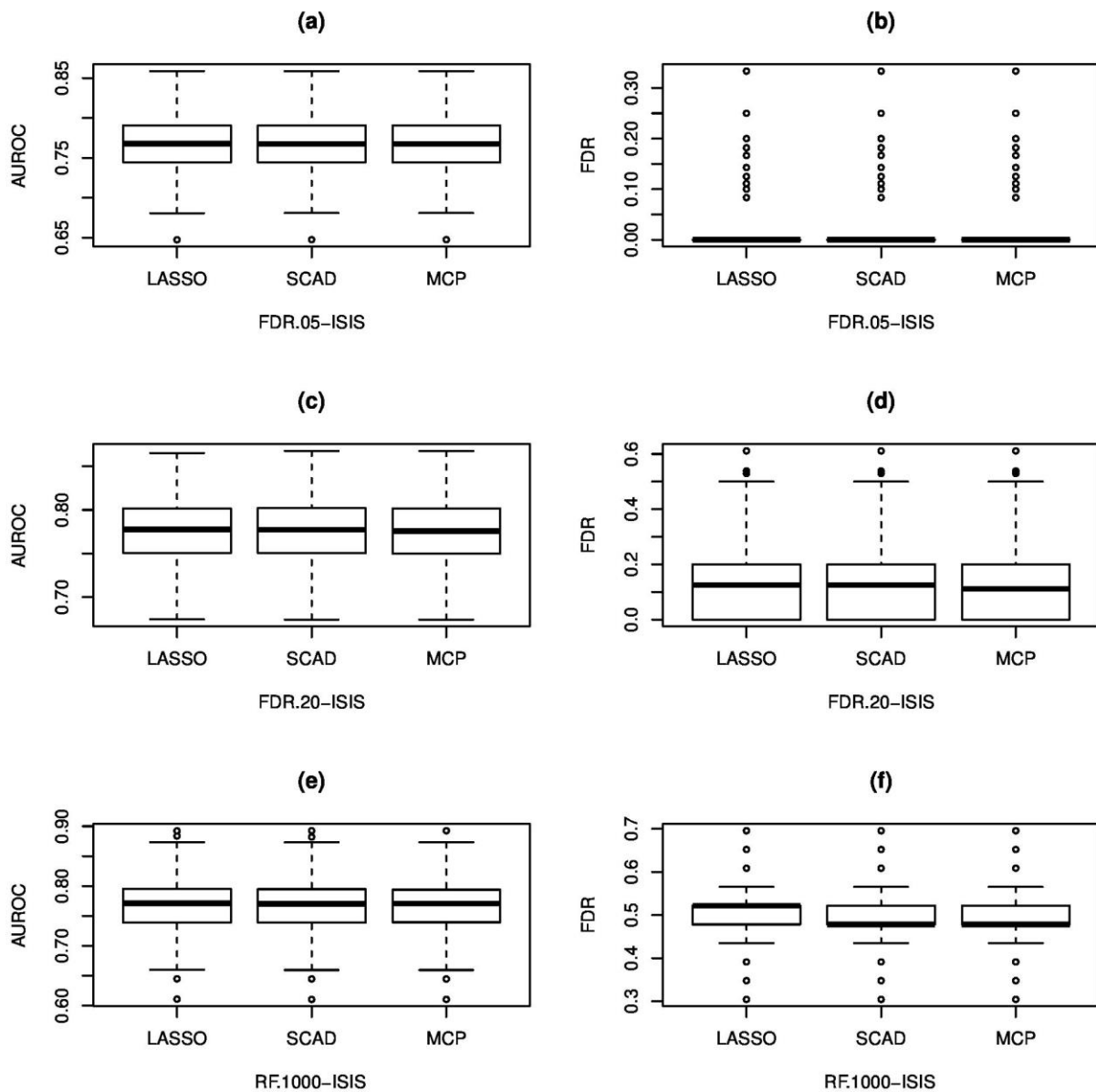




Figure S5| Boxplots of the expressions of 12 detected genes on the prostate cancer. Both AN07 and MYBPC1 are the top down-regulated whereas the rest of them are the top up-regulated. The red color is no metastasis and another is metastasis.

