

## Additional file 6

### 1. STAD data

**Table 1.** Statistics of gene expression profiling, DNA methylation and clinical cohort information of STAD under study.

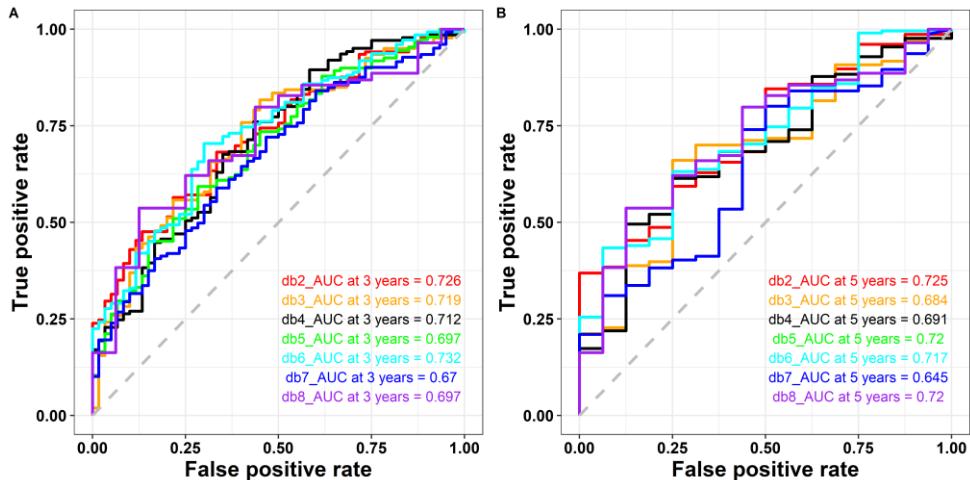
Item	Statistics
<b>#. STAD clinical cohort</b>	
Tumor	375
Normal	32
<b>Survival status (tumor)</b>	
Living	204
Deceased	171
<b>Race (tumor)</b>	
Asian	74
Black	11
White	238
Not reported	52
<b>Tumor stage</b>	
I	53
II	111
III	150
IV	38
Not reported	23
<b>Follow-up (months)</b>	0.03-125.8
<b>Age (years)</b>	
Range	35-90
Median	63
<b>Gender (tumor)</b>	
Male	241
Female	134
<b>#. (Epi) genomic data</b>	
mRNA profiling (DEGs)	16486
DNA methylation (DMPs)	6197

## 2. The results of our proposed methods

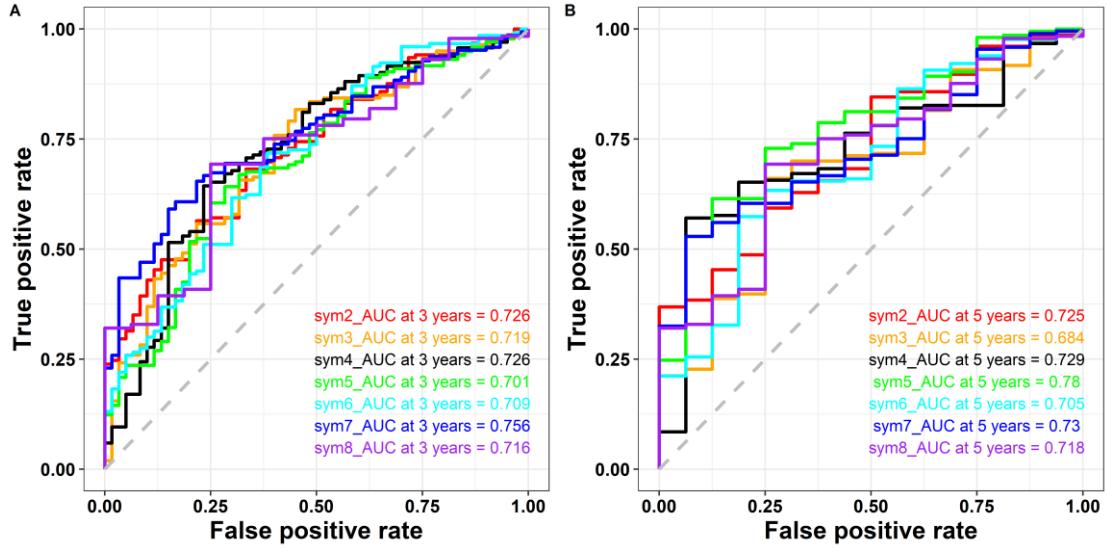
**Table 1.** The detailed analysis results of STAD dataset about comparisons on the adopted wavelet-based, SWT-CNN methods and classic LASSO methods with diverse predictors

	<b>Wavelet function</b>	<b>AUC at 3 years</b>	<b>AUC at 5 years</b>	<b>gene number</b>
<b>db</b>	db2	0.726	0.725	8
	db3	0.719	0.684	7
	db4	0.712	0.691	5
	db5	0.697	0.720	7
	db6	0.732	0.717	7
	db7	0.670	0.645	4
	db8	0.697	0.720	7
	bior1.1	0.656	0.655	7
<b>bior</b>	bior1.3	0.725	0.718	6
	bior1.5	0.710	0.706	6
	bior2.2	0.725	0.718	6
	bior2.4	0.707	0.690	6
	bior2.6	0.700	0.719	7
	bior2.8	0.725	0.718	6
	bior3.1	0.698	0.714	5
	bior3.3	0.701	0.697	5
	bior3.5	0.701	0.697	5
	bior3.7	0.700	0.719	7
<b>sym</b>	bior3.9	0.714	0.682	7
	bior4.4	0.710	0.706	6
	bior5.5	0.725	0.718	6
	bior6.8	0.710	0.706	6
	sym2	0.726	0.725	8
	sym3	0.719	0.684	7
	sym4	0.726	0.729	5
	sym5	0.701	0.780	7
<b>coif</b>	sym6	0.709	0.705	5
	sym7	0.756	0.730	12
	sym8	0.716	0.718	8
	coif2	0.709	0.705	5
	coif3	0.709	0.705	5
<b>haar</b>	coif4	0.712	0.691	5
	coif5	0.712	0.691	5
<b>dmeyer</b>		0.692	0.681	9
<b>dmeyer</b>		0.710	0.706	6
	rbio1.1	0.672	0.771	5
	rbio1.3	0.685	0.666	6

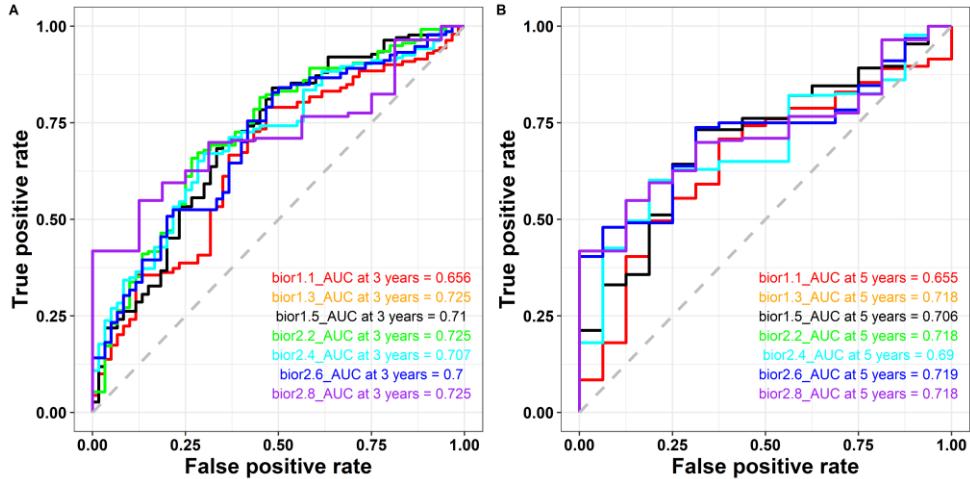
	rbio1.5	0.670	0.623	7
	rbio2.2	0.673	0.695	8
	rbio2.4	0.670	0.623	7
	rbio2.6	0.688	0.636	5
	rbio2.8	0.708	0.648	8
<b>rbio</b>	rbio3.1	0.656	0.655	6
	rbio3.3	0.656	0.655	6
	rbio3.5	0.673	0.695	8
	rbio3.7	0.655	0.623	4
	rbio3.9	0.673	0.695	8
	rbio4.4	0.710	0.706	6
	rbio5.5	0.700	0.719	7
	rbio6.8	0.710	0.706	6
	fk4	0.709	0.703	6
	fk6	0.712	0.691	5
<b>fk</b>	fk8	0.716	0.718	8
	fk14	0.715	0.705	7
	fk18	0.697	0.720	7
	fk22	0.714	0.693	11
<b>SWT-CNN</b>		0.673	0.742	7
<b>LASSO</b>		0.682	0.709	14



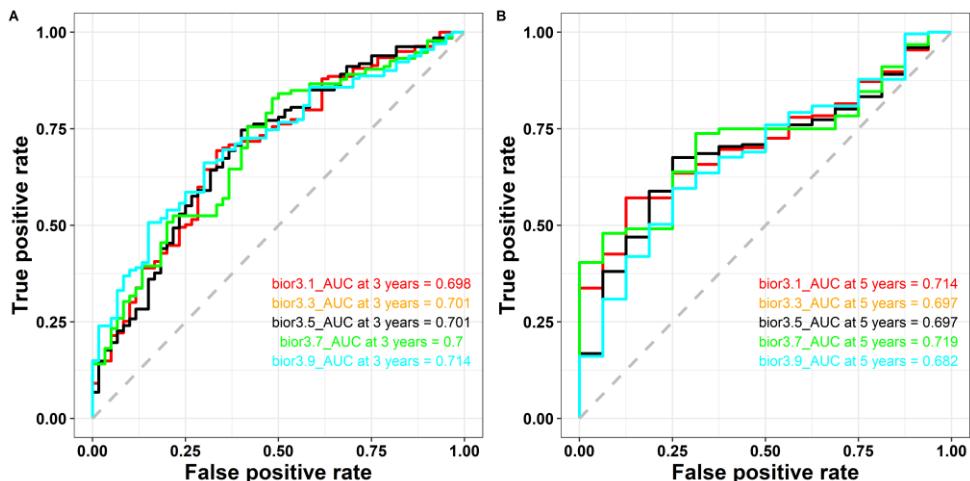
**Figure 1.** The performance of STAD dataset on db basis function.



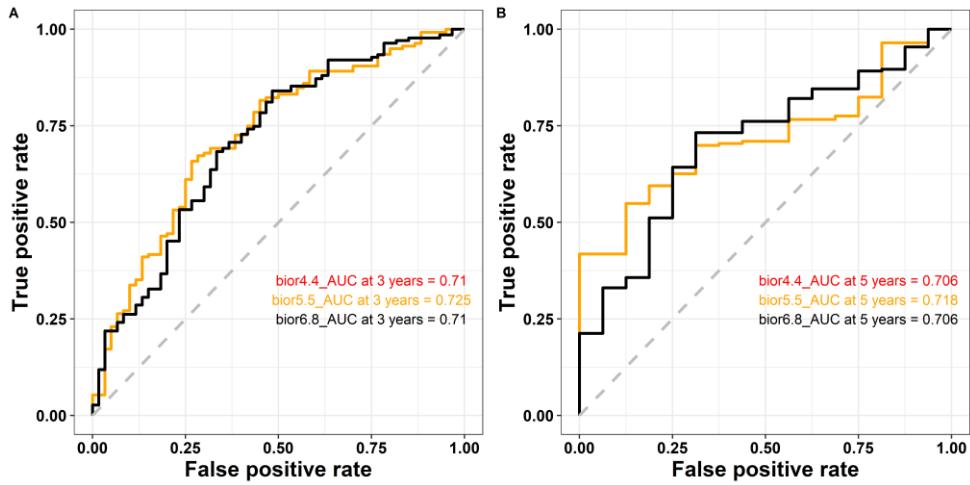
**Figure 2.** The performance of STAD dataset sym basis function.



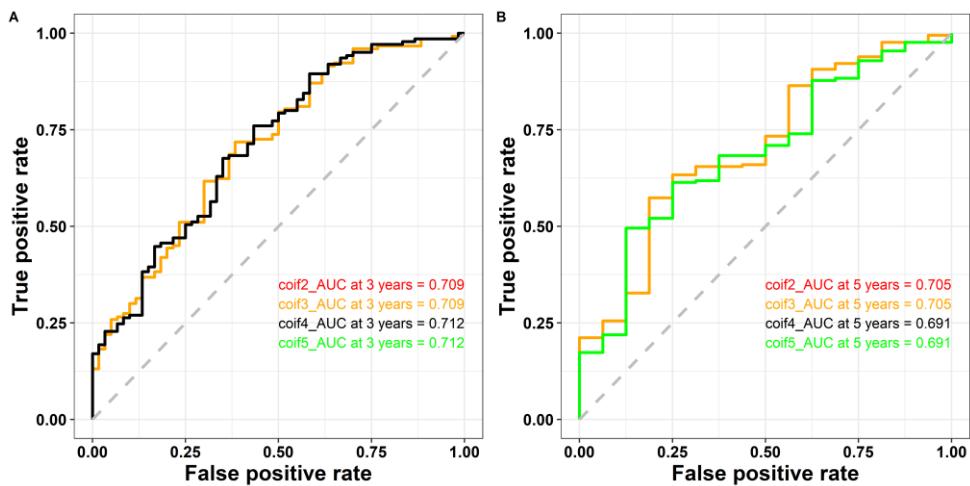
**Figure 3.** The performance of STAD dataset on bior basis function (bior1.1~bior2.8).



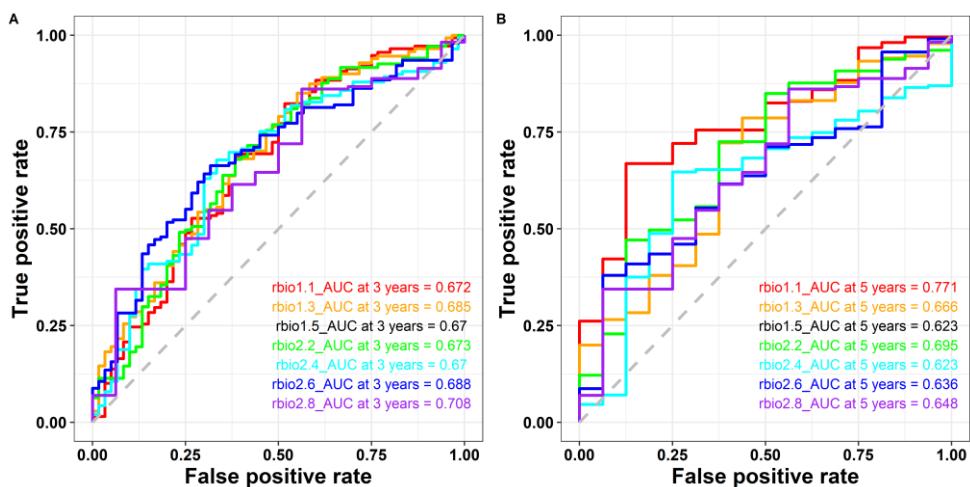
**Figure 4.** The performance of STAD dataset on bior basis function (bior3.1~bior3.9).



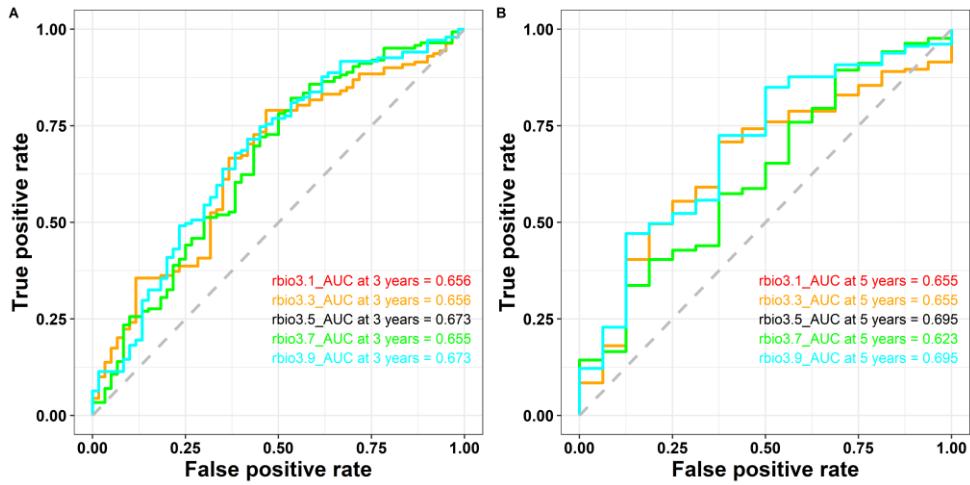
**Figure 5.** The performance of STAD dataset on bior basis function (bior4.4~bior6.8).



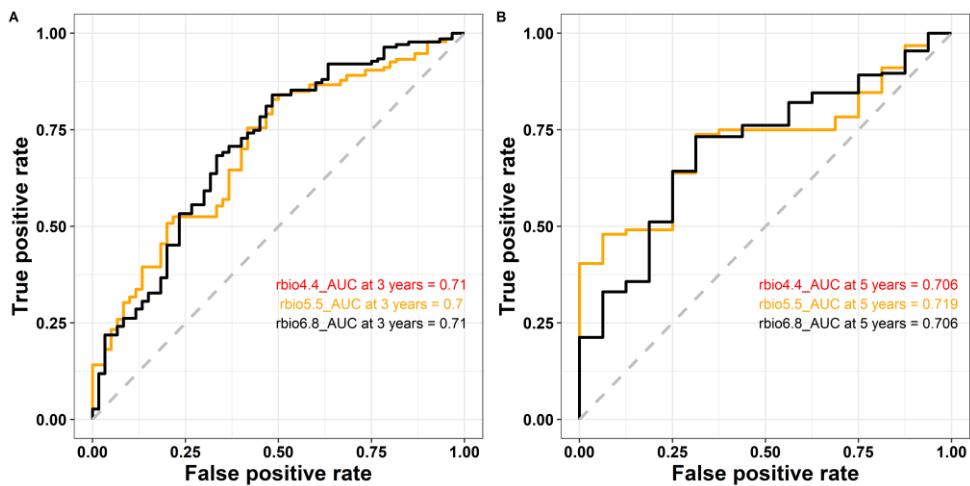
**Figure 6.** The performance of STAD dataset on coif basis function.



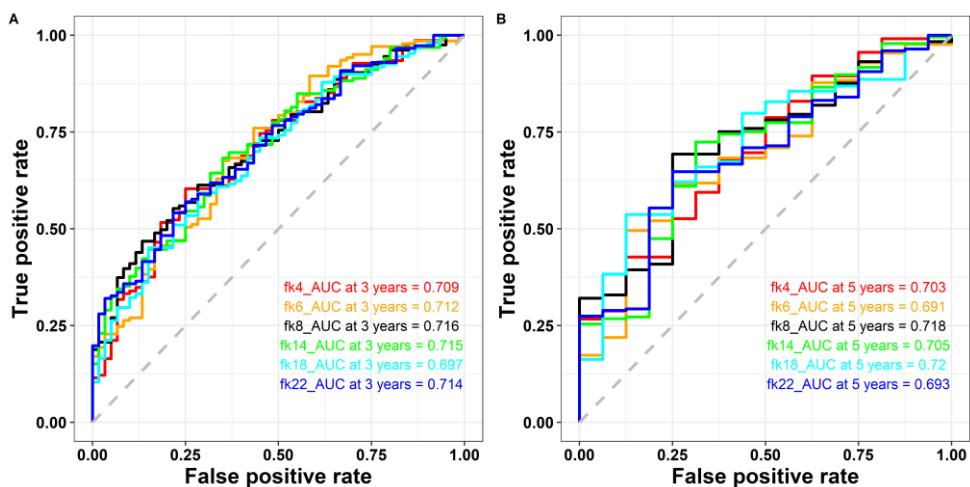
**Figure 7.** The performance of STAD dataset on rbio basis function (rbio1.1~rbio2.8).



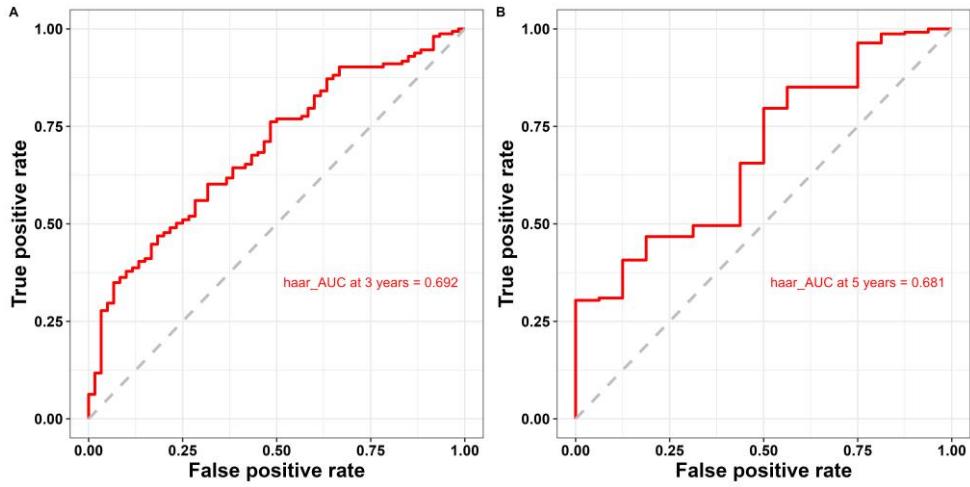
**Figure 8.** The performance of STAD dataset on rbio basis function (rbio3.1~rbio3.9).



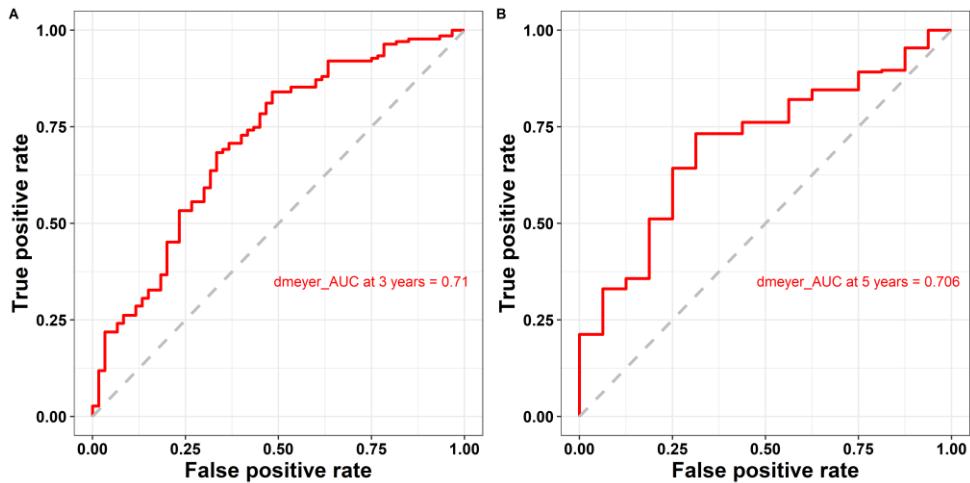
**Figure 9.** The performance of STAD dataset on rbio basis function (rbio4.4~rbio6.8).



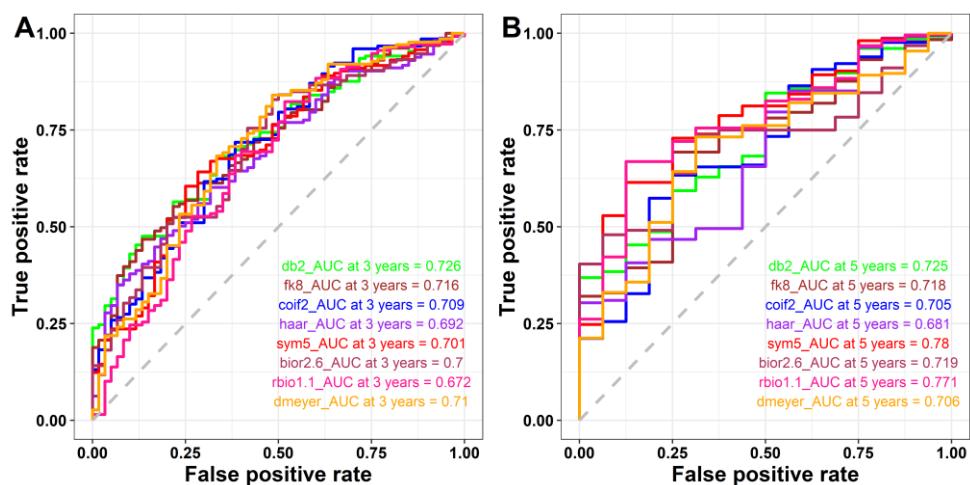
**Figure 10.** The performance of STAD dataset on fk basis function.



**Figure 11.** The performance of STAD dataset on haar basis function.

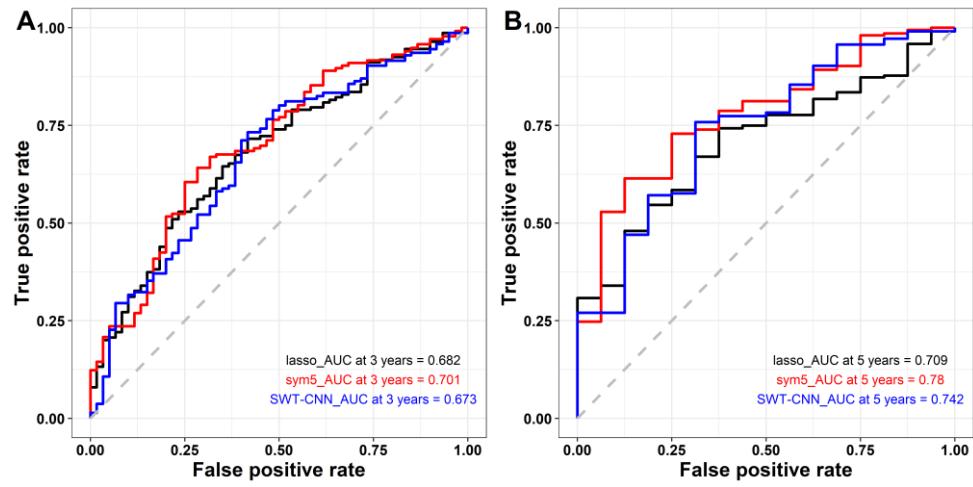


**Figure 12.** The performance of STAD dataset on dmeyer basis function.



**Figure 13.** The performance of STAD dataset on all basis functions (Take the smoothness corresponding to the best result for each basis function).

### 3. Comparison of the three methods



**Figure 16.** Comparison of the three methods on the STAD dataset. (A) AUC at 3 years: sym5\_AUC at 3 years = 0.701, lasso\_AUC at 3 years = 0.682, SWT-CNN at 3 years = 0.673; (B) AUC at 5 years: sym5\_AUC at 5 years = 0.780, lasso\_AUC at 5 years = 0.709, SWT-CNN at 5 years = 0.742.