

	CN2				QUESTGilles				inTrees			
	acc	balacc	nodes	time	acc	balacc	nodes	time	acc	balacc	nodes	time
wisconsinBreast	0.9498+0.02 $\sigma$	0.943+0.02 $\sigma$	16.0+2.94 $\sigma$	0.2478+0.17 $\sigma$	0.9484+0.01 $\sigma$	0.941+0.01 $\sigma$	23.6667+2.49 $\sigma$	0.7522+0.19 $\sigma$	0.9527+0.0 $\sigma$	0.9452+0.01 $\sigma$	16.0+2.94 $\sigma$	25.8307+0.26 $\sigma$

	Genetic				ISM_pruned				QUESTLoh			
	acc	balacc	nodes	time	acc	balacc	nodes	time	acc	balacc	nodes	time
wisconsinBreast	0.9355+0.01 $\sigma$	0.937+0.01 $\sigma$	43.0+23.72 $\sigma$	754.7636+189.11 $\sigma$	0.9484+0.01 $\sigma$	0.9458+0.02 $\sigma$	45.0+16.57 $\sigma$	172.5185+20.76 $\sigma$	0.9499+0.01 $\sigma$	0.9499+0.0 $\sigma$	8.3333+0.94 $\sigma$	0.2996+0.02 $\sigma$

	C4.5				ISM				CART			
	acc	balacc	nodes	time	acc	balacc	nodes	time	acc	balacc	nodes	time
wisconsinBreast	0.9441+0.01 $\sigma$	0.9337+0.02 $\sigma$	18.3333+4.99 $\sigma$	0.0105+0.0 $\sigma$	0.9427+0.01 $\sigma$	0.9336+0.02 $\sigma$	100.3333+7.54 $\sigma$	198.1263+18.2 $\sigma$	0.9413+0.01 $\sigma$	0.9345+0.02 $\sigma$	23.6667+15.08 $\sigma$	0.002+0.0 $\sigma$

	GUIDE			
	acc	balacc	nodes	time
wisconsinBreast	0.937+0.01 $\sigma$	0.9274+0.01 $\sigma$	11.6667+0.94 $\sigma$	0.2917+0.01 $\sigma$

Accuracy on wisconsinBreast dataset using 3 folds

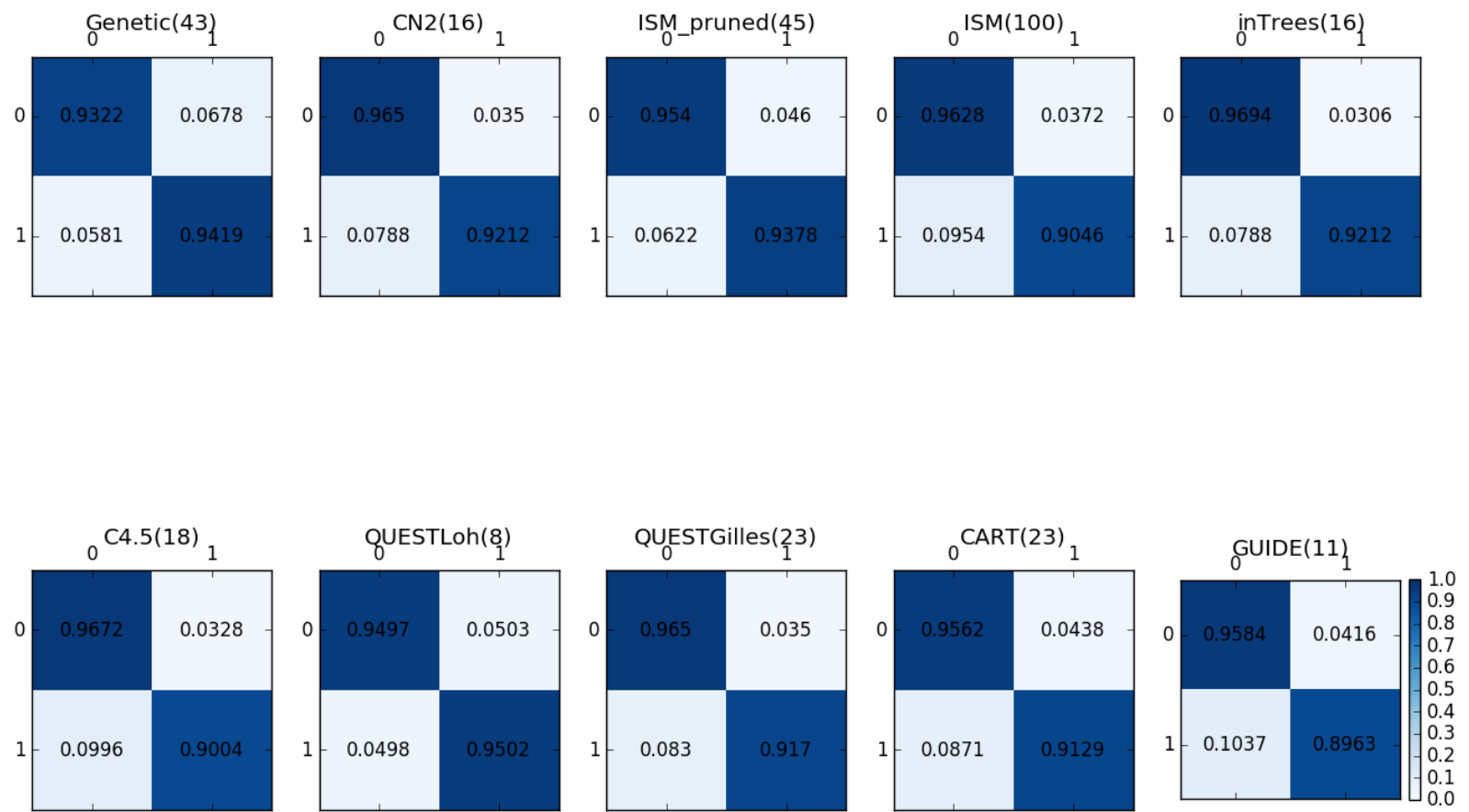


Figure 1: Confusion matrix for wisconsinBreast