

Identification of diagnostic markers for tuberculosis by proteomic fingerprinting of serum

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	Sensitivity	Specificity	Accuracy
Mass Peaks			
All	0.89	0.77	0.83
10 positive correlated and 10 negative correlated	0.78	0.77	0.78
Markers			
Transthyretin	0.94	0.52	0.73
CRP	0.82	0.48	0.65
neopterin	0.71	0.74	0.72
SAA	0.82	0.52	0.68
neopterin-SAA	0.88	0.61	0.75
CRP-SAA	0.88	0.52	0.70
CRP-neopterin	0.82	0.65	0.74
Transthyretin-SAA	0.88	0.52	0.70
Transthyretin-neopterin	0.94	0.52	0.73
Transthyretin-CRP	0.82	0.52	0.67
Transthyretin-CRP-neopterin	0.82	0.65	0.74
Transthyretin-CRP-SAA	0.71	0.78	0.74
Transthyretin-neopterin-SAA	0.82	0.74	0.78
CRP-neopterin-SAA	0.82	0.52	0.67
Transthyretin-CRP-neopterin-SAA	0.88	0.74	0.81

Webtable 3: SVM classifiers performance on selected mass cluster peaks and biomarkers of second dataset