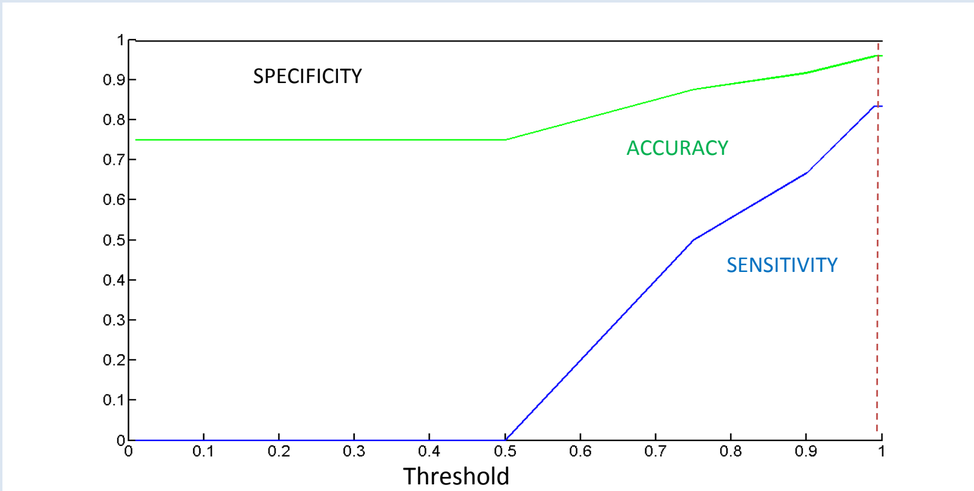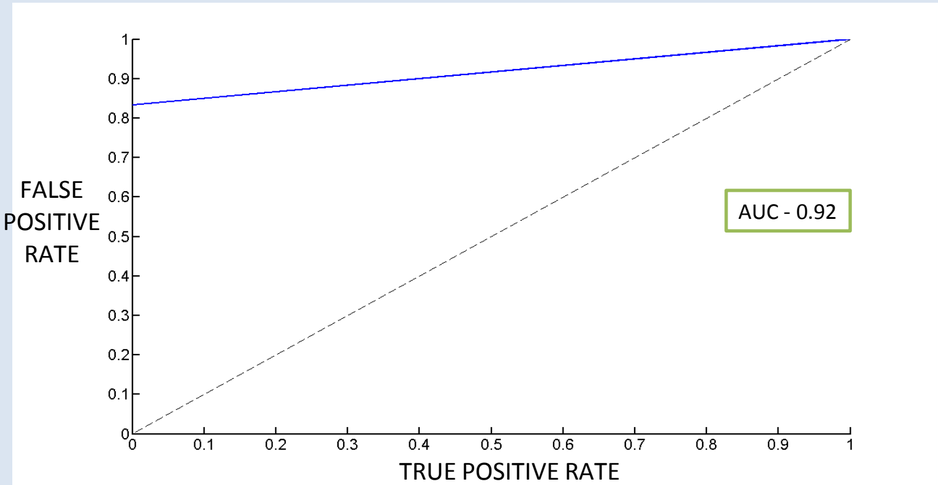Effect of changing the threshold for essential gene prediction in M. tuberculosis