

Figure S6-A. Sensitivity and specificity of a predictive model based on association of *LINC00667* gene expression levels in metastatic melanoma (61 non-responders and 55 responders) with response to inhibitor of CTLA-4 (ipilimumab). AUC-value = 0.67, Mann-Whitney test p-value = 0.0019, fold change of median gene expression values (non-responders vs responders) = 2.2. Analysis was performed using the web-based tool ROC-ploter.

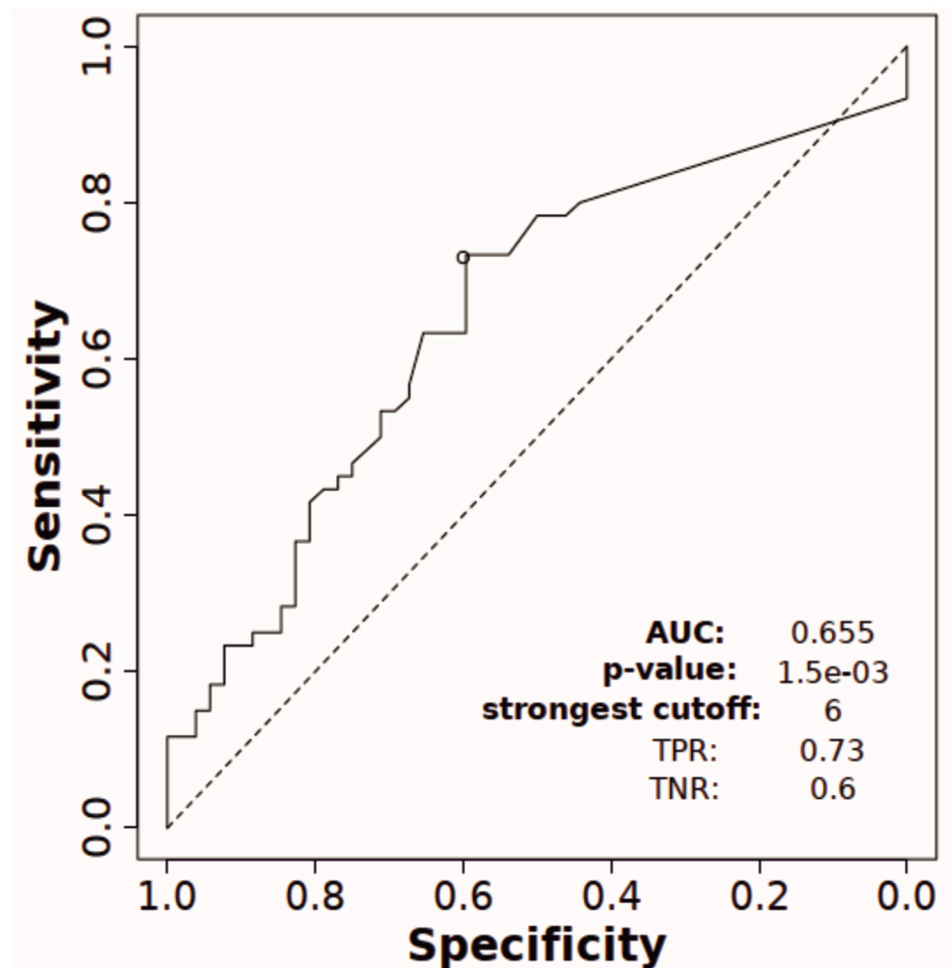


Figure S6-B. Sensitivity and specificity of a predictive model based on association of *LINC00526* gene expression levels in metastatic melanoma (61 non-responders and 55 responders) with response to inhibitor of CTLA-4 (ipilimumab). AUC-value = 0.65, Mann-Whitney test p-value = 0.0045, fold change of median gene expression values (non-responders vs responders) = 2.6. Analysis was performed using the web-based tool ROC-plottter.