H.W.#3

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Question #4.

In terms of accuracy (TP+TN/TP+TN+FP+FN), it turned out that the second model(I.e. model based on **PIK3CA_GRCh37_3:178952085-178952085_Missense-Mutation_SNP_A-A-T_A-A-G)** is more accurate.

Question #5.

When compared by "TP-FP", the model based on **RNF43_GRCh37_17:56435161-56435161_Frame-Shift-Del_DEL_C-C--** is the most useful for classifying samples. Maximum value for the quantity "TP-FP" is 11.

Question #6.

When compared by "%TP-%FP", the model based on **RNF43_GRCh37_17:56435161-56435161_Frame-Shift-Del_DEL_C-C-- i**s the most useful for classifying samples. Maximum value for the quantity "%TP-%FP" is 0.04.