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and if two ones are added, the given condition is false. The condition is false, thus false will be returned. The question in OP's example is, when the line takes too long, we don't need to add the second group, but just one. The selection is a selection between one or two. In your example you could do something like `if len(d) == len(data): data_list = data.sort(key=some_key_function, reverse=True) else: data_list = data` or `if len(d) == len(data): data_list = data.sort(key=some_key_function, reverse=True) elif len(d) != len(data): data_list = data` As other answer have said, the truth value of expression depends on context. Elevated expression of MDM2-FL, a spliced variant of MDM2, in human hepatocarcinogenesis. There is now evidence that MDM2 gene and/or its product, MDM2-FL, can function as an oncogene. In the current study, MDM2-FL was detected in 52 of 66 (79%) human liver tumors and 3 of 19 (16%) normal human livers, suggesting that this spliced variant is overexpressed in the hepatocarcinogenesis. MDM2-FL mRNA was also found in human hepatoma cell lines, which expressed a significant amount of MDM2-FL protein. Thus, overexpression of MDM2-FL might be a contributing factor to the etiology of human hepatocarcinogenesis. Selectivity of translocation of the Sec and Tat signal peptides across model and natural membranes. For efficient secretion, the signal peptide (SP) of a protein is recognized by the translocon (translocon), which is composed of SecY, SecE, and SecG complexed with the membrane, and SecA that catalyzes translocation. The efficiency of the translocon varies widely between different translocation substrates, probably due to differences in the physical properties of the membrane and in the composition and dynamics of the translocon. We have investigated the interaction of the model SP, SecM, and the natural SP, signal recognition particle (SRP), with the translocon and phospholipid bilayers 2d92ce491b