Assignment 2

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Objective

This report presents the results of the analysis conducted to determine whether a localization sequence is present in specific genes. Our team has been assigned to find whether the given sequence **SKL(serine-lysine-leucine)** is a tripeptide sequence found at the C-terminus of proteins targeted to the peroxisomes within cells which serves as a peroxisome localization signal **PTS1**, is present within various genes. Below is a table summarizing the presence of this sequence within each gene.

Code

We downloaded sequences for the given genes and saved them as text files. Then, we wrote a Python script to check whether the localization sequence is present in the gene. Finally, we saved the outputs in a excel file. You can run the script.py directly with all other files downloaded locally, or you can run the IPython notebook file in Google Colab.

Results

Gene	Localization Sequence Present	Gene	Localization Sequence Present
ALDH18A1	YES	CATIP	NO
ALG12	NO	CBY1	NO
ALG9	YES	CC2D2A	YES
ALMS1	YES	CCDC103	NO
AMER1	NO	CCDC112	NO
AMMECR1	NO	CCDC13	YES
ANK1	YES	CCDC14	NO
ANKRD11	YES	CEP83	NO
ANKRD26	YES	CEP89	YES
ANKS3	NO	CEP97	YES
ANKS6	NO	CERKL	NO
ANO2	NO	CETN1	NO
ANOS1	NO	CETN2	NO
BUB3	NO	CETN3	NO
C2CD3	NO	CFAP100	NO
CA4	NO	CFAP126	NO
CABCOCO1	NO	CFAP161	NO

Observation

We observed that the localization sequence is present in the genes ALDH18A1, ALG9, ALMS1, ANK1, ANKRD11, ANKRD26, CC2D2A, CCDC13, CEP89, and CEP97.

Reference

UniProt