

Protein Kinase Sequence Motifs

Downloaded from https://esbl.nhlbi.nih.gov/Databases/Kinase_Logos/

Kinase data curated from Sugiyama et al. (PMID: 31324866). Sequence motifs for each protein kinase showing amino acid preferences flanking the post-translational modified site were identified by PTMLogo. All logos were generated from a minimum of 30 amino acid sequences, using Chi-squared filtering alpha = 0.0001. Exact number of reference sequences used for each kinase can be found at the link above. PTMLogo published by Saethang et al. 2019, PMID: 31318409.

Amino acids are color coded based on their properties. **L, I, M, V, A** - green. **K, R, H** – blue, **Q, N** - purple, **D, E** - red, **F, W, Y, P** - dark grey, **S, T** - light grey, **C** - pink, **G** - orange.

B represents the N-terminus and **J** represents the C-terminus.

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