

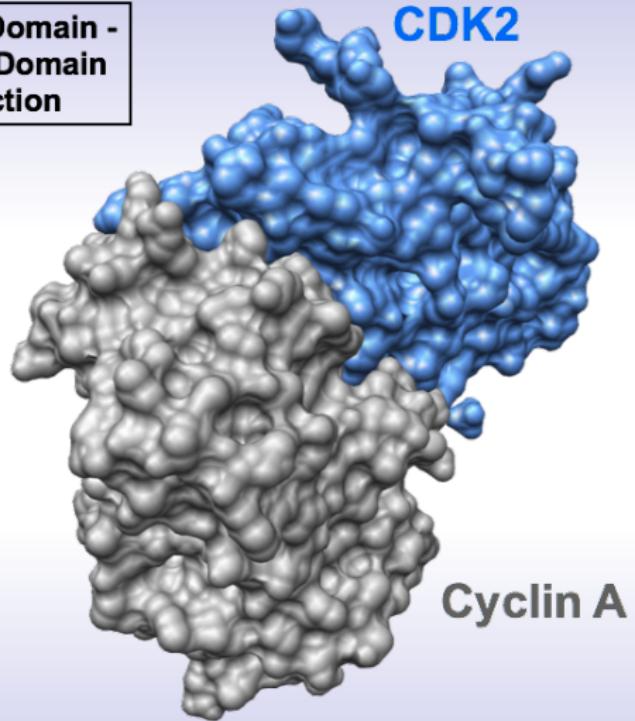
SHORT LINEAR MOTIFS

Holger Dinkel

EMBO Practical Course “Computational analysis of
protein-protein interactions – From sequences to
networks”

IMPORTANCE OF SHORT LINEAR MOTIFS

**Globular Domain -
Globular Domain
Interaction**

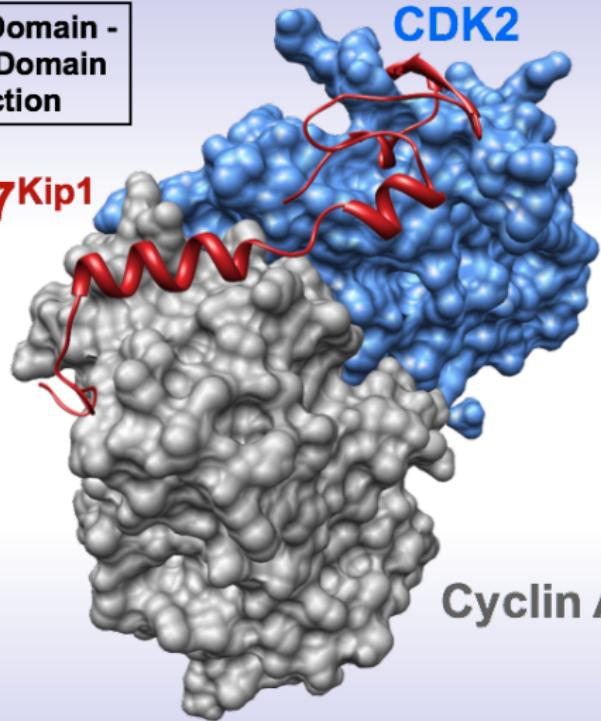


Cyclin A

IMPORTANCE OF SHORT LINEAR MOTIFS

Globular Domain -
Disordered Domain
Interaction

Globular Domain -
Globular Domain
Interaction

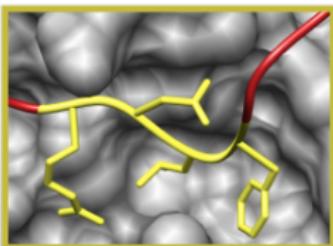


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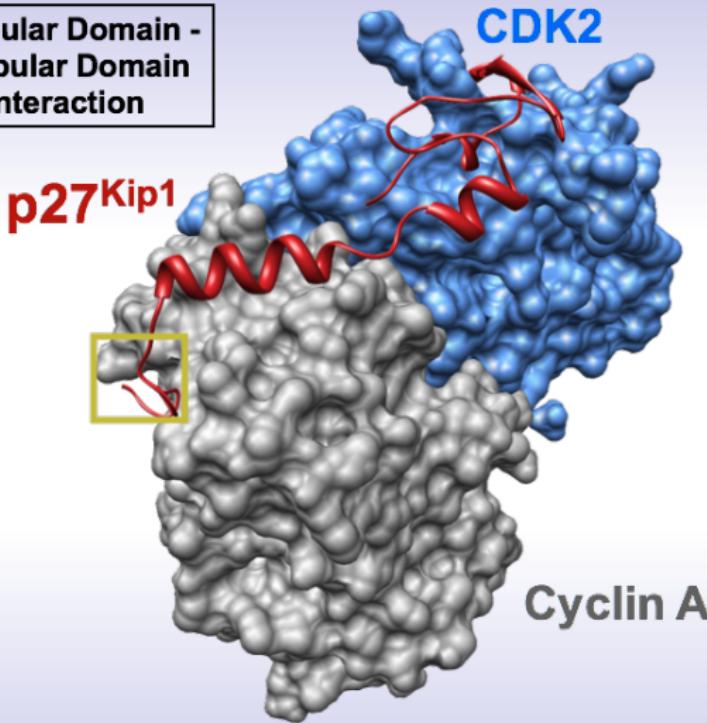
Globular Domain -
Globular Domain
Interaction

Globular Domain -
Disordered Domain
Interaction

Globular Domain -
Short Linear Motif
Interaction



RNLF



Cyclin A

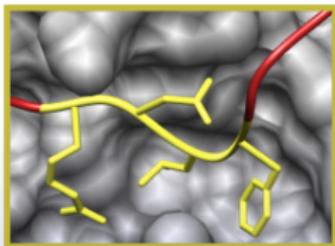
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Interaction

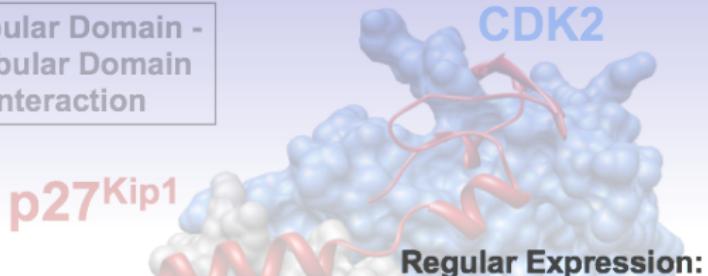
Globular Domain -
Globular Domain
Interaction

PDB 1JSU

Russo et al., Nature. 1996;
382: 325-331.



RNLF



LIG_CYCLIN_1

Regular Expression:
[RK]xLx{0,1}[FYLIVMP]

Defined positions

Fixed positions

Degenerate positions

Undefined positions

Fixed-length wildcard

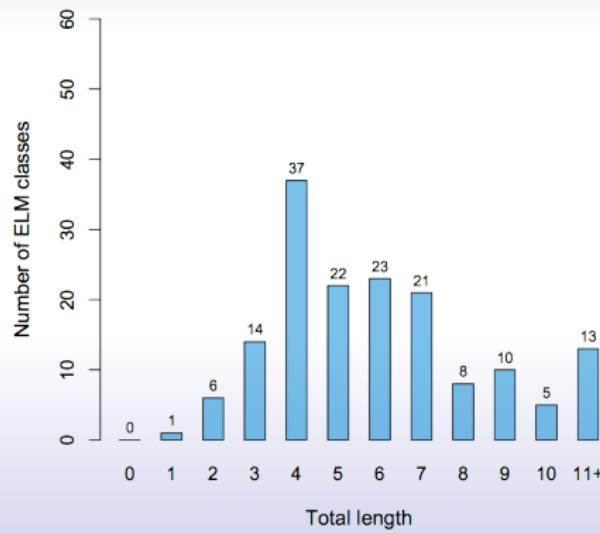
Flexible-length wildcard {min,max}

Cyclin A

ATTRIBUTES OF SHORT LINEAR MOTIFS

LINEAR MOTIFS

- are small.
- have few defined positions.
- mediate transient, low affinity interactions.

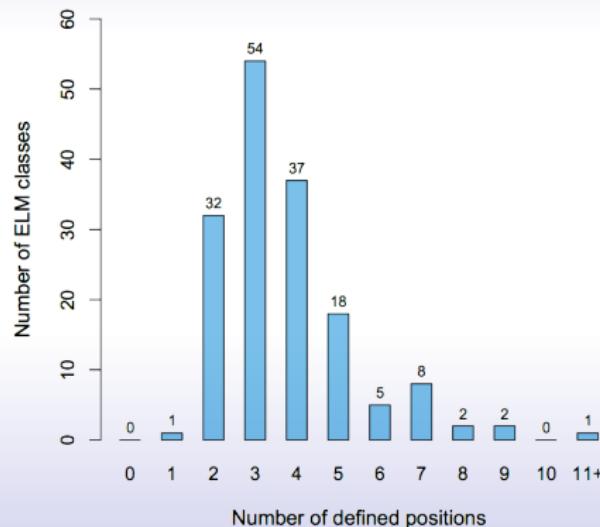


"Attributes of short linear motifs"; DAVEY, VAN ROEY, WEATHERITT, TOEDT, UYAR, ALtenBERG, BUDD, DIELLA, DINKEL & GIBSON; (MOL BIOSYST. 2011)

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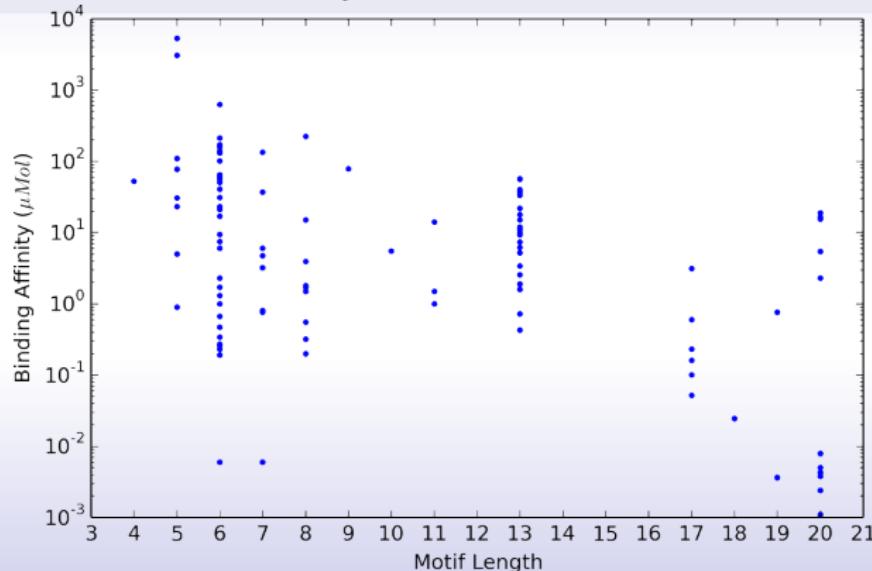


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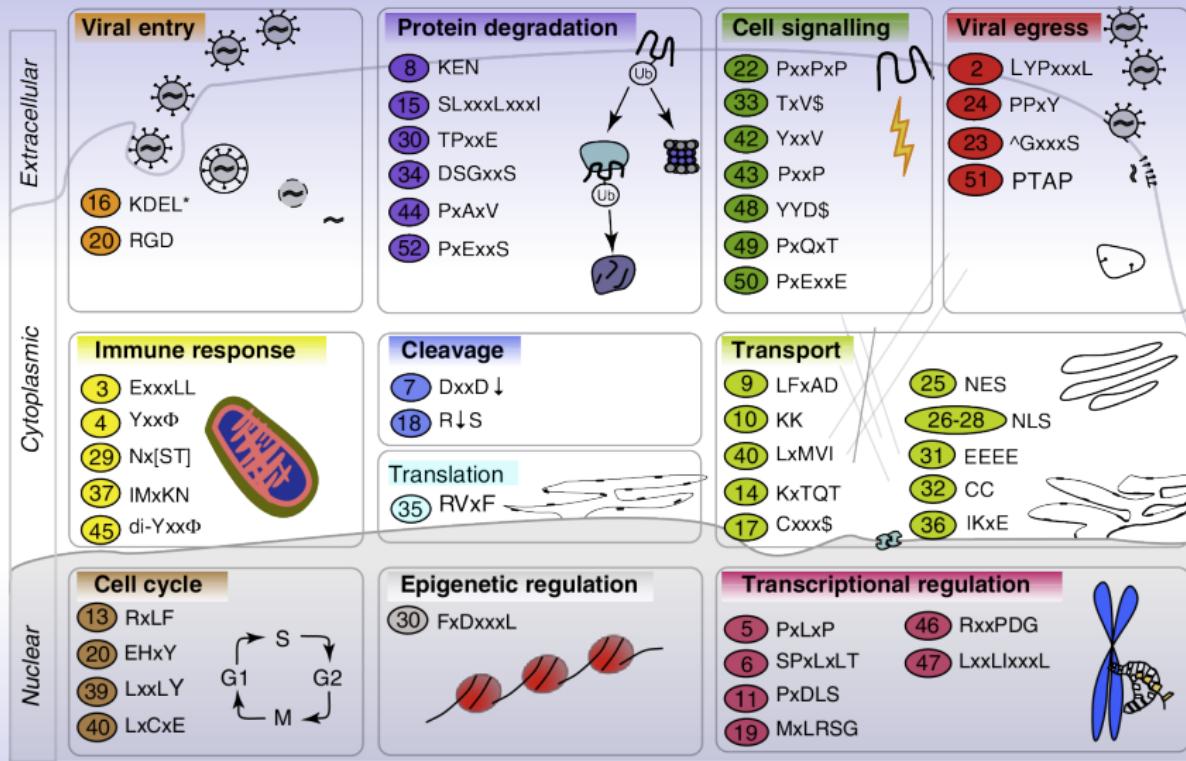


PREVALENCE OF SHORT LINEAR MOTIFS

DOMAIN FREQUENCIES FROM PFAM (HUMAN PROTEOME):

Domain Family	Frequency [Domains / Proteins]	Pattern of recognized motif
PDZ	573 / 342	$[ST]x[ACVILF]_{-COOH}$
SH3	451 / 382	$PxxP$
SH2	237 / 219	$_P Yxx[IV]$
WW	151 / 103	$PPxY$
PTB	142 / 133	NPx_pY

IMPORTANCE OF SHORT LINEAR MOTIFS: VIRUSES



"How viruses hijack cell regulation"; DAVEY, TRAVÉ & GIBSON; (TIBS 2010)

IMPORTANCE OF SHORT LINEAR MOTIFS: DISEASES

LIDDLE'S-SYNDROME: WW-INTERACTION MOTIF

has been implicated with autosomal dominant activating mutations in the WW interaction motif in the β - and γ -subunits of the epithelial sodium channel **ENAC**. These mutations abrogate the binding to the ubiquitin ligase **NEDD4-2**, ultimately resulting in increased Na^+ reabsorption, plasma volume extension and hypertension.

IMPORTANCE OF SHORT LINEAR MOTIFS: DISEASES

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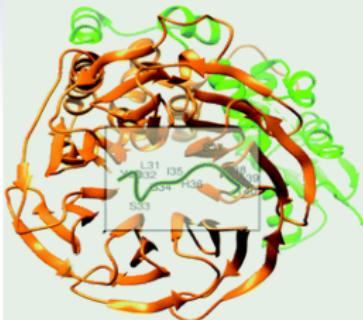
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BACILLUS ANTHRACIS “LETHAL FACTOR”

The protein **LEF_BACAN** is a metalloprotease (one of the three proteins composing the anthrax toxin) that specifically targets mitogen-activated protein kinase kinases (MKKs). which are important regulators of signal transduction as they phosphorylate and thus activate specific MAPKs (such as ERK1, ERK2, p38 or JNK). *Bacillus anthracis*’ “lethal factor” cleaves its MKK substrates within or close to the MAPK docking sites, thus effectively preventing the MKK to dock to its MAPK.

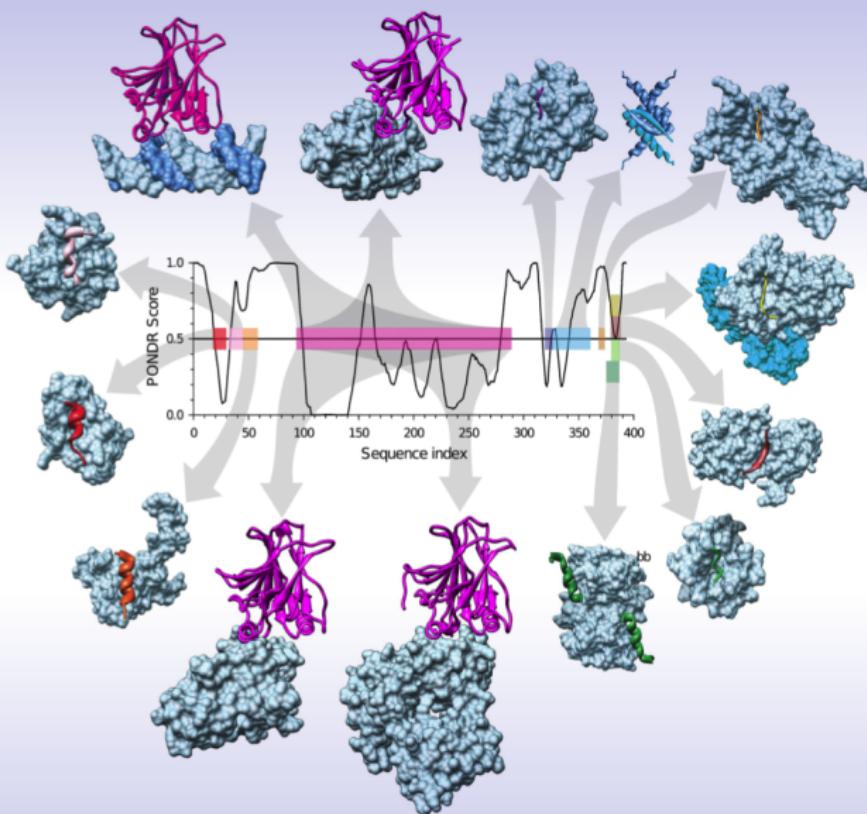
IMPORTANCE OF SHORT LINEAR MOTIFS: CANCER

β -CATENIN



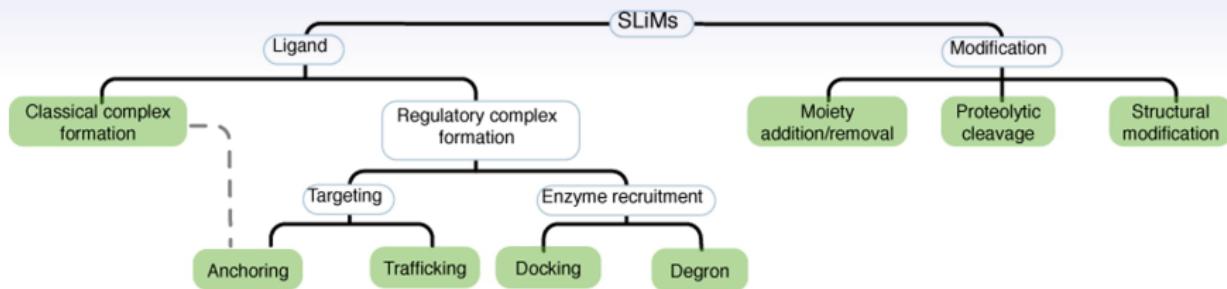
The most recurrently mutated experimentally validated motif in the COSMIC DB is the conserved proteasomal degradation motif (DEG_SCF_TRCP1_1) in the highly disordered N-terminal region of β -Catenin which mediates binding to the WD40 repeat domain of the β -TRCP subunit of the SCF-betaTRCP E3 ubiquitin ligase complex. (more than 1700 mutation entries for this motif derived from 1692 unique samples based on 256 different publications)

IMPORTANCE OF SHORT LINEAR MOTIFS: P53



"Understanding protein non-folding"; UVERSKY & DUNKER; (BIOCHIMICA ET BIOPHYSICA ACTA 2010)

CLASSIFICATION OF MOTIFS



MOTIF CLASSES: MODIFICATION SITES

DESCRIPTION:

Modification Motifs mediate specific binding to the active site of a modifying enzyme to allow subsequent catalytic post-translational modification of the target site.

EXAMPLE:

NAME MOD_CDK_1

REGEx $xxx([ST])Px[KR]$

Kinase domain

CDK site

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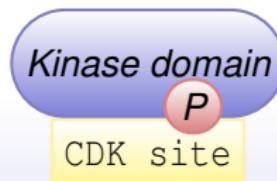
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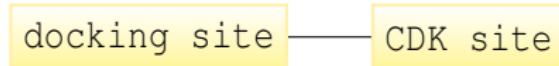
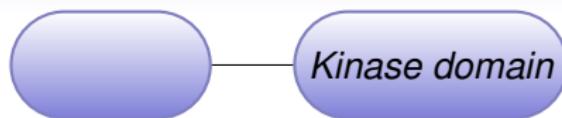
MOTIF CLASSES: DOCKING MOTIFS

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Docking motifs recruit enzymes via a surface that is distinct from the active site.

EXAMPLE:

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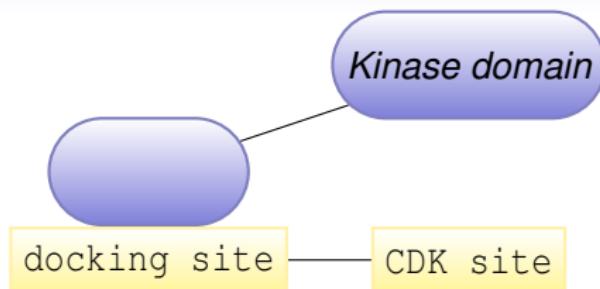
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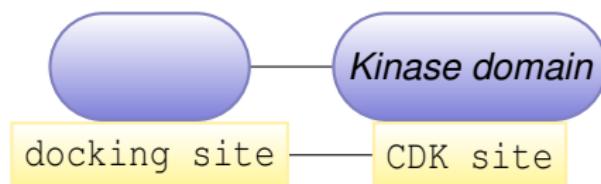
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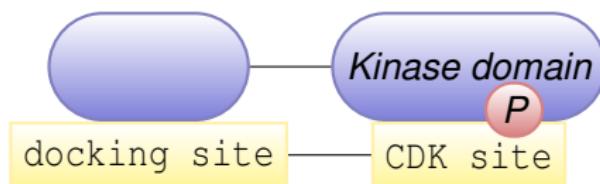
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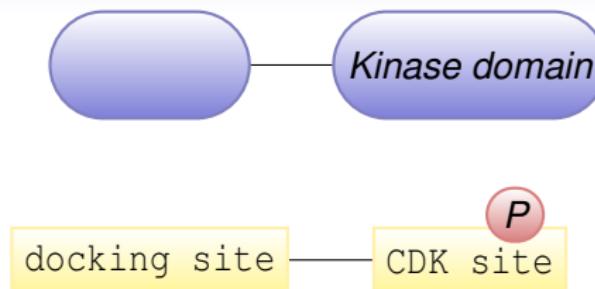
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MOTIF CLASSES: CLEAVAGE MOTIFS

DESCRIPTION:

Proteolytic processing of proteins into smaller polypeptides by protease-catalyzed hydrolysis of specific peptide bonds

EXAMPLE:

NAME CLV_Separin_Metazoa

REGEx $E[IMPVL][MLVP]Rx$

Separase

Cleavage site

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Separase



MOTIF CLASSES: DEGRADATION MOTIFS

DESCRIPTION:

Degradation motifs (Degrons) recognized by E3 Ubiquitin Ligase complexes priming proteins for degradation, regulating protein half-life.

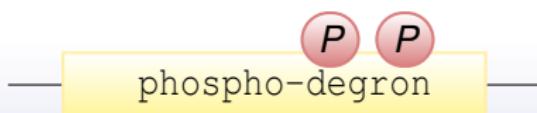
EXAMPLE:

NAME DEG_SCF_TRCP1_1

REGEx $D(S)GXX([ST])$

FBW7

SCF E3 Ligase



MOTIF CLASSES: DEGRADATION MOTIFS

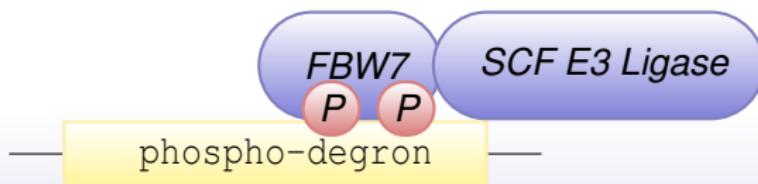
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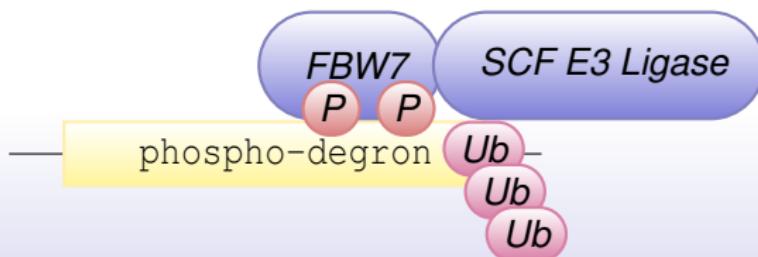
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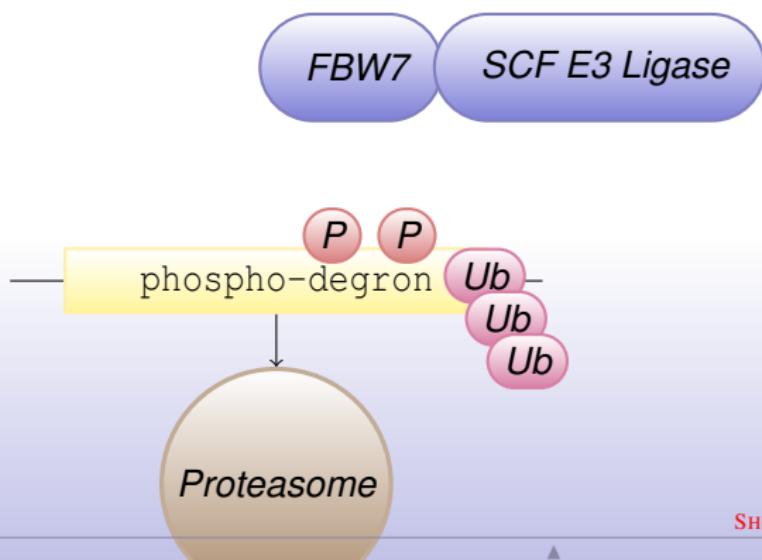
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MOTIF CLASSES: TARGETING/ANCHORING MOTIFS

DESCRIPTION:

TARGETING motifs allow a protein to bind to the transport machinery that relocalizes it to a particular sub-cellular location.

ANCHORING motifs are recognized by biomolecules specific to a sub-cellular location and thereby retain the motif-containing protein at that location.

EXAMPLE:

NAME TRG_NLS_MonoCore_2

REGEx [^DE](K[RK]|RK)[KRP][KR][^DE]

Importin α

NLS

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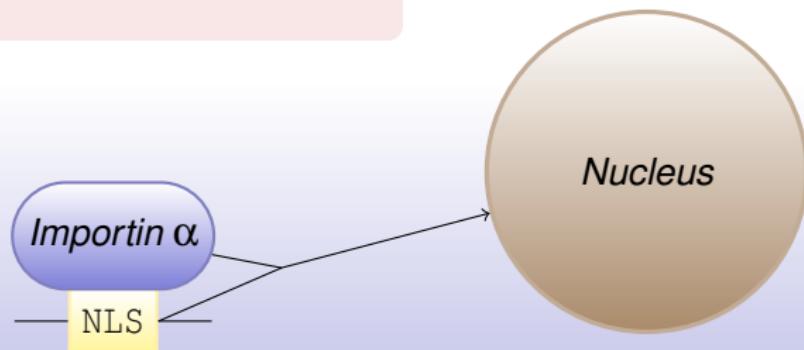
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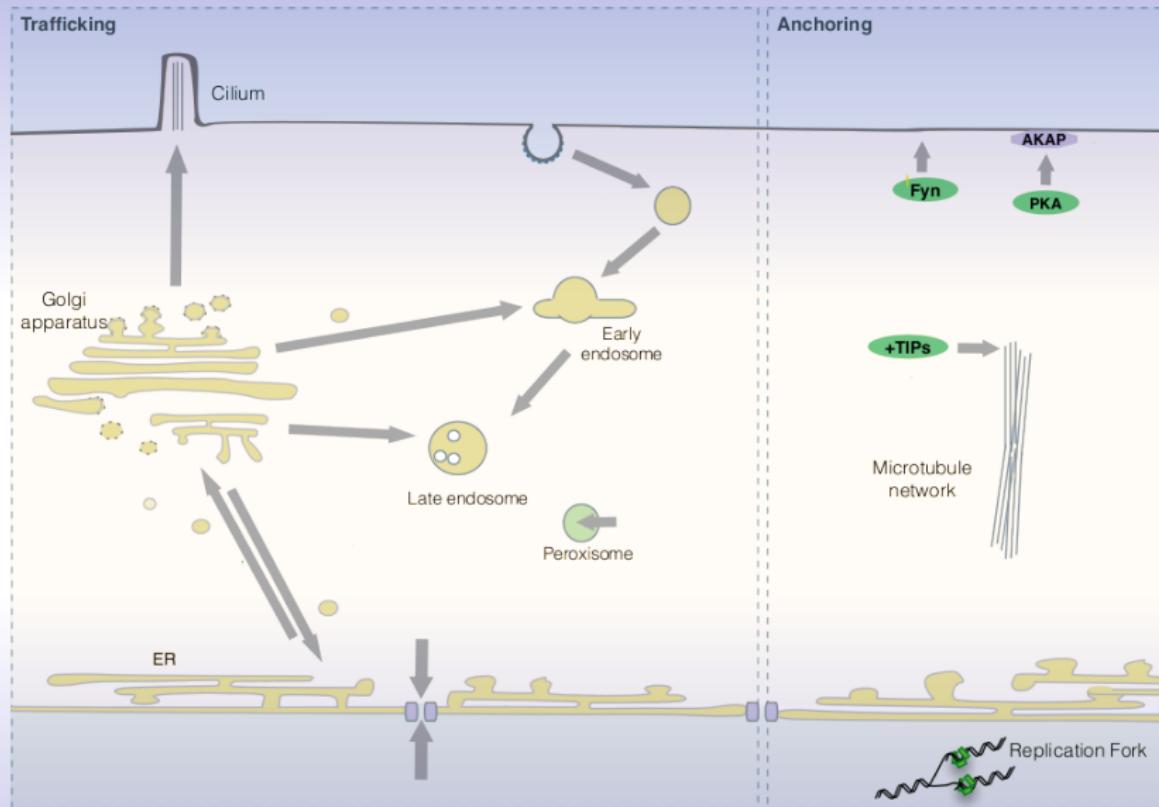
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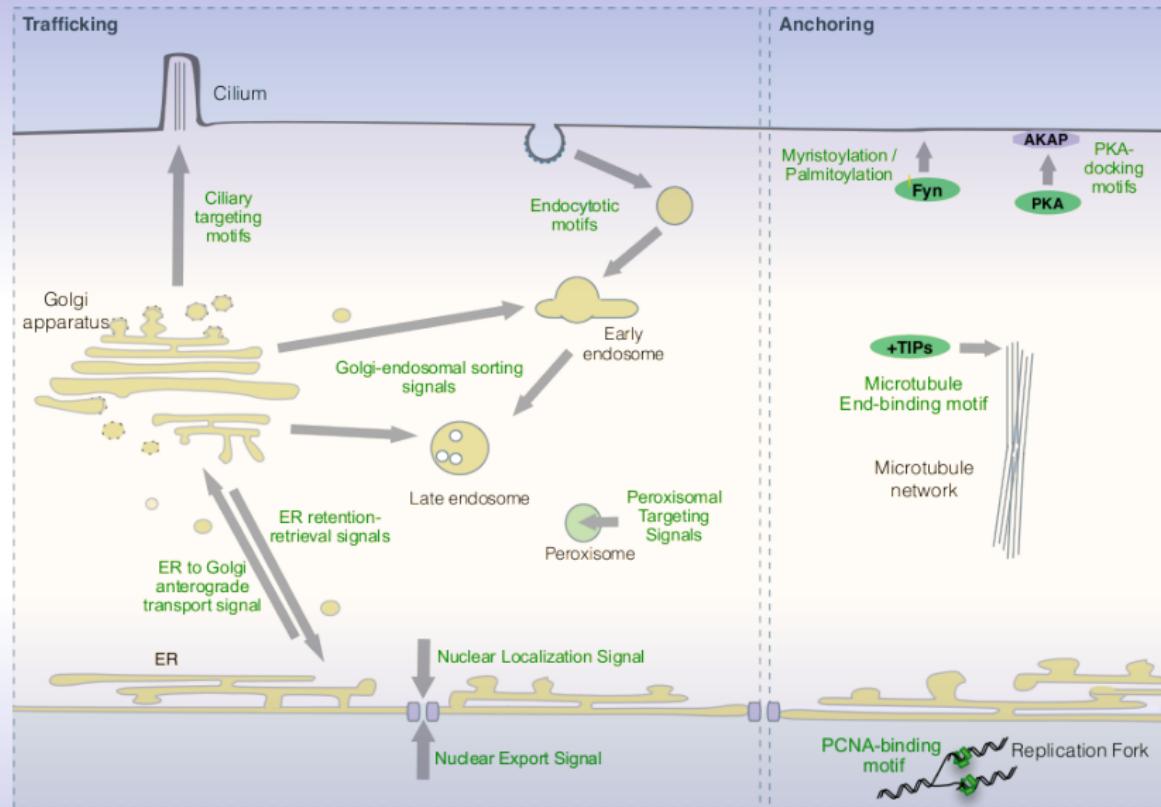


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"Short linear motifs: Ubiquitous and functionally diverse protein interaction modules directing cell regulation"; VAN ROEY, UYAR, WEATHERITT, DINKEL, SEILER, BUDD, GIBSON & DAVEY; (CHEM. REVIEWS; 2014)

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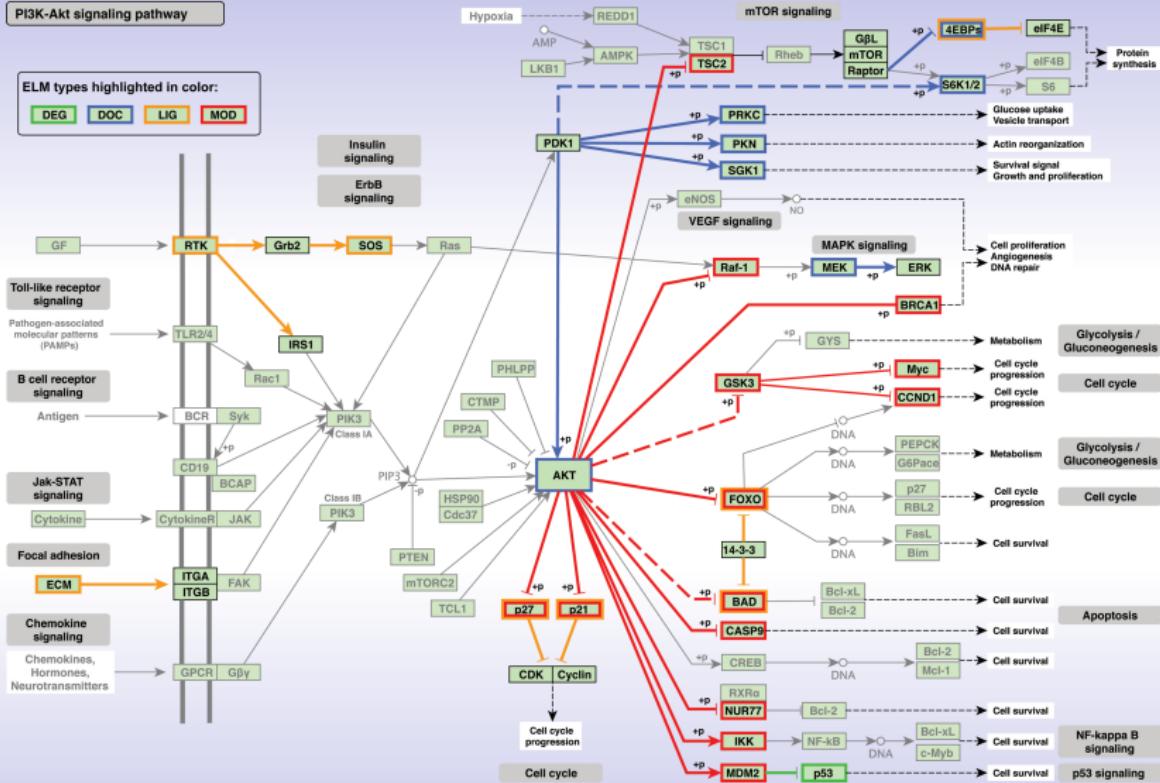


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VISUALIZING MOTIF-MEDIATED INTERACTIONS



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OUTLOOK: THE ELM RESOURCE

The Eukaryotic Linear Motif resource for *Functional Sites in Proteins*

ELM is a **REPOSITORY** of more than 240 thoroughly annotated motif classes with over 2700 annotated instances.

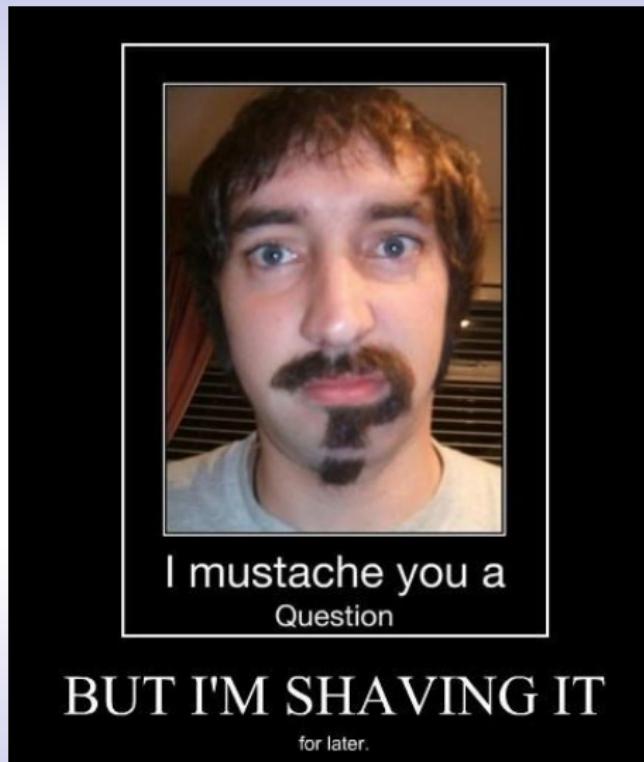
It is also a **PREDICTION TOOL** to detect these motifs in protein sequences employing different filters to distinguish between **functional** and **non-functional** motif instances.

SUMMARY

SHORT LINEAR MOTIFS

- small, versatile modules which mediate transient interactions
- important regulators of cellular processes.
- “kidnapped” by viruses
- play an important role in diseases
- collected in the Eukaryotic Linear Motif Resource (ELM)

QUESTIONS?



I mustache you a
Question

BUT I'M SHAVING IT

for later.