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ScanProsite Res

Output format: Graphical view - this view shows ScanProsite results together with ProRule-based predicted intra-domain feat

Hits for USERPAT1 "[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]" on LGri

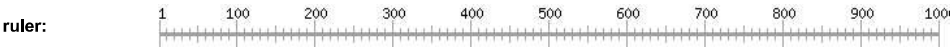
found: 94 hits in 70 sequences

Legend:



Please note that the graphical representations of domains displayed hereafter are for illustrative purposes only, and that they are not intended to be used for any other purpose. For more information about how these graphical representations are constructed, go to <https://prosite.expasy.org/mydomain>

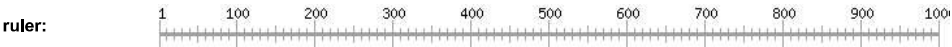
hits by patterns: [94 hits (by 1 pattern) on 70 sequences]



USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?>26784

195 - 202: FvGAaYkF



USERPAT1 :

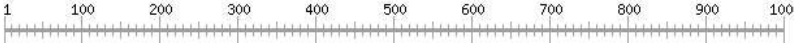
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)

26784

321 - 328: QtGVqVgI

ruler:



1fi1_A



[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>1fi1_A%0A&output=nice](http://prosite.expasy.org/cgi-bin/prosite/scanprosite/ScanView.cgi?seq=>1fi1_A%0A&output=nice))

USERPAT1 :

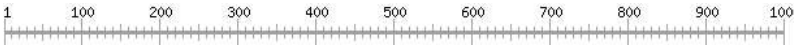
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)

26784

472 - 479: RgGVnYlF

ruler:



1im0_A



[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>1im0_A%0A&output=nice](http://prosite.expasy.org/cgi-bin/prosite/scanprosite/ScanView.cgi?seq=>1im0_A%0A&output=nice))

USERPAT1 :

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

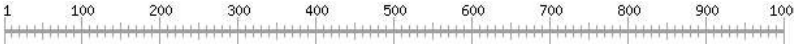
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)

26784

217 - 224: ElGLsYpI

251 - 258: RvGVgVmL

ruler:



1k24_A



[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>1k24_A%0A&output=nice](http://prosite.expasy.org/cgi-bin/prosite/scanprosite/ScanView.cgi?seq=>1k24_A%0A&output=nice))

USERPAT1 :

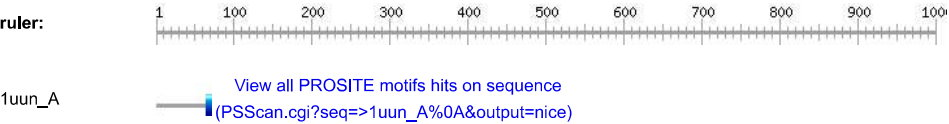
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)

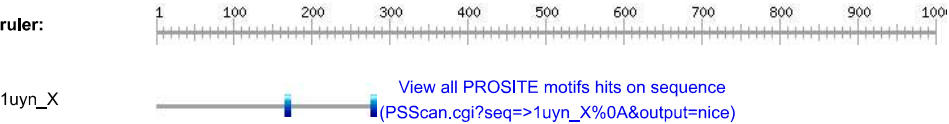
26784

105 - 112: RvGAgLgF

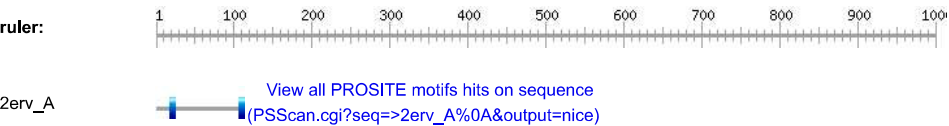
240 - 247: EyGLrVgI



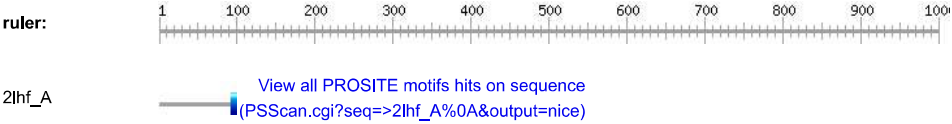
USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]
67 - 74: QiGFpWsl



USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]
164 - 171: EgGLrYdL
272 - 279: RvGVgYrF



USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]
18 - 25: RIGLsWdW
107 - 114: RiGAgLkF

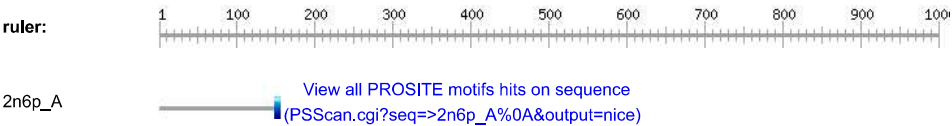


USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]

93 - 100: FgGAtLgL

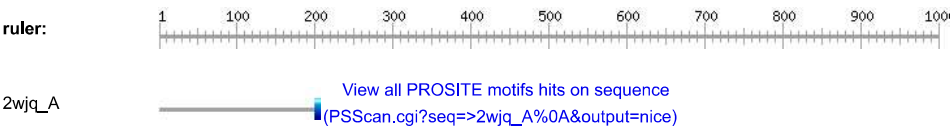


USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]

149 - 156: EIGFdYmL

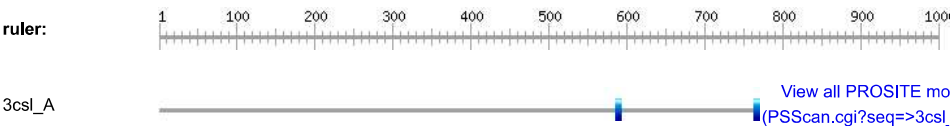


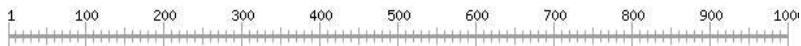
USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]

198 - 205: RiGVsFkL

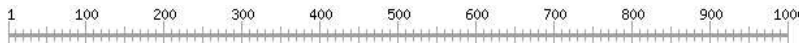


USERPAT1 :Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)]**573 - 580:** FrGLeYqL**746 - 753:** QgGVeYrF**ruler:**

3dzm_A

[View all PROSITE motifs hits on sequence](#)

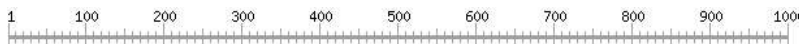
(PSScan.cgi?seq=>3dzm_A%0A&output=nice)

USERPAT1 :Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)]**128 - 135:** QlGLgLgV**201 - 208:** RlGAaYrF**ruler:**

3emo_C

[View all PROSITE motifs hits on sequence](#)

(PSScan.cgi?seq=>3emo_C%0A&output=nice)

USERPAT1 :Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)]**86 - 93:** QnGLaIgV**ruler:**

3fh_A

[View all PROSITE motifs hits on sequence](#)

(PSScan.cgi?seq=>3fh_A%0A&output=nice)

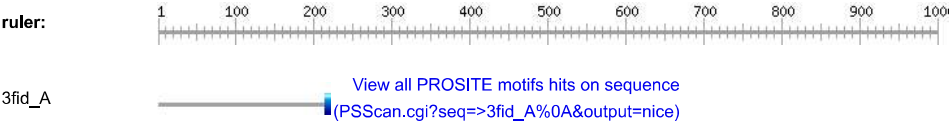
USERPAT1 :Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)]

26784

68 - 75: HrGVlVlV

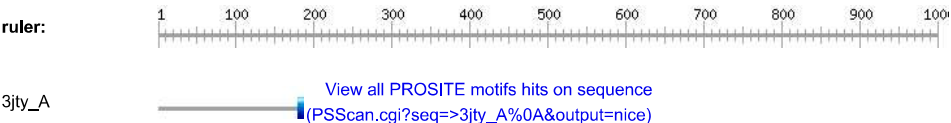
438 - 445: EyGFgLrF

543 - 550: HsGFsVgW



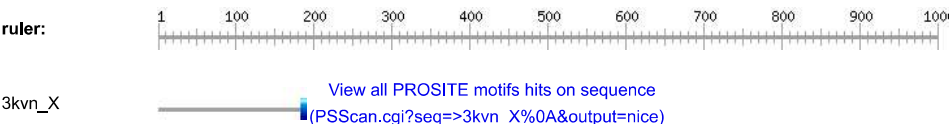
USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175>)]
26784

212 - 219: KeGLiYkV



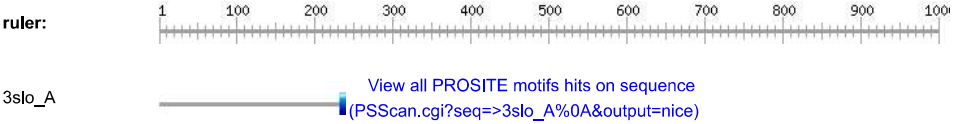
USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175>)]
26784

178 - 185: FqGAeYaF

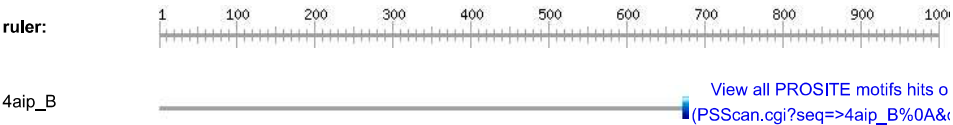


USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175>)]
26784

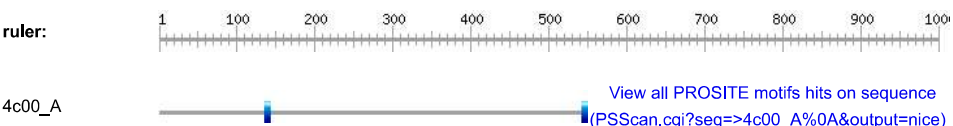
182 - 189: QaGARyViV



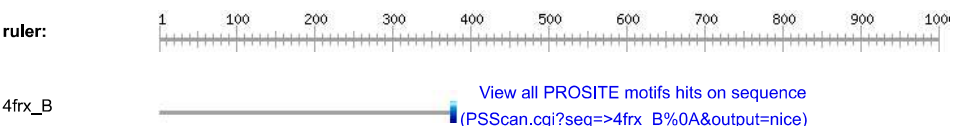
USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?> 26784
230 - 237: RaGLgVqF



USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?> 26784
658 - 665: RIGVnYkF



USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?> 26784
134 - 141: QLGIaLgL
531 - 538: EhGLqFyI

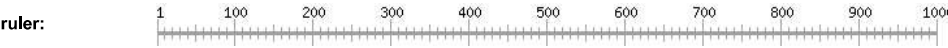


USERPAT1 :

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175)]
26784

369 - 376: FkGLgFtW

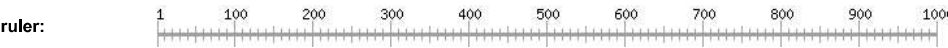


USERPAT1 :

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175)]
26784

163 - 170: RvGVqYsF

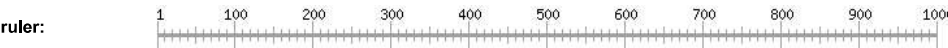


USERPAT1 :

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175)]
26784

206 - 213: KvGVnFyW

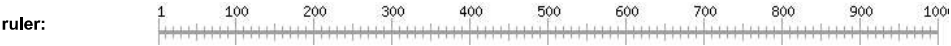


USERPAT1 :


Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175)]
26784

142 - 149: KyGVsWaf



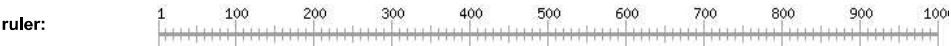
5bun_C



[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>5bun_C%0A&output=nice](#))

USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?> 26784

280 - 287: KiGLnFsL



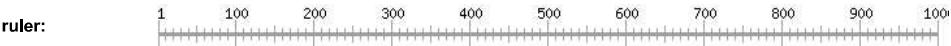
5dl8_B




[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>5dl8_B%0A&output=nice](#))

USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?> 26784

98 - 105: KyGAtLkL



5fq6_E



[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>5fq6_E%0A&output=nice](#))

USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?> 26784

126 - 133: EdGLtYkV



5fq6_M



[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>5fq6_M%0A&output=nice](#))

USERPAT1 :

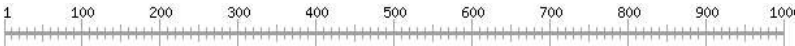
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175](https://pubmed.ncbi.nlm.nih.gov/11535175/) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?>26784

613 - 620: EvGLnMaF

939 - 946: RFGFnLmV

ruler:



5jzt_A



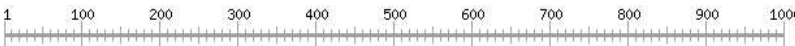
USERPAT1 :

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175](https://pubmed.ncbi.nlm.nih.gov/11535175/) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?>26784

414 - 421: EiGApVpL

ruler:



5mwv_A



USERPAT1 :

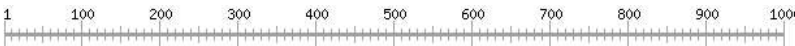
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175](https://pubmed.ncbi.nlm.nih.gov/11535175/) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?>26784

155 - 162: EtGLqYtF

246 - 253: QnGLsVsL

ruler:



5o65_B

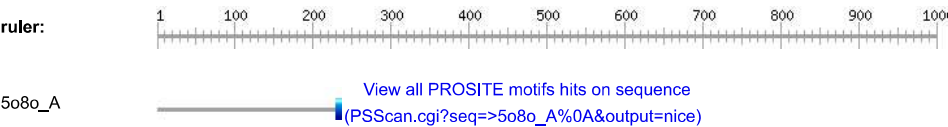


USERPAT1 :

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?PMID=11535175)]
26784

208 - 215: QfGVGVaF

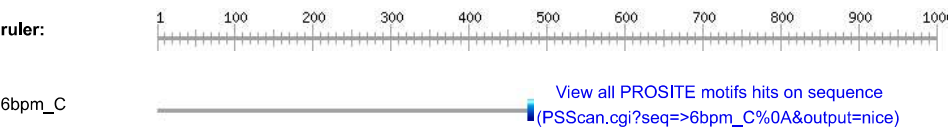


USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?PMID=11535175)]
26784

227 - 234: QaGVdMtL

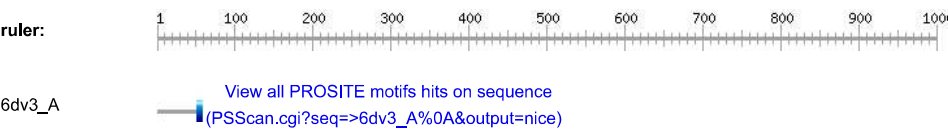


USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?PMID=11535175)]
26784

467 - 474: KaGAlYhL

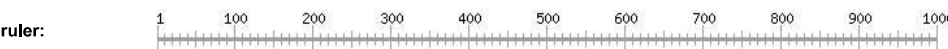


USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?PMID=11535175)]
26784

54 - 61: QlGLiWYF



6eus_A

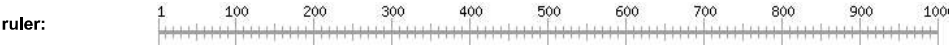
[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>6eus_A%0A&output=nice](#))

USERPAT1 :

Pattern: [KRH~~Q~~FE]-x-G-[IVL~~F~~AC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?>26784

7 - 14: KaGAdVnL



6h04_A

[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>6h04_A%0A&output=nice](#))

USERPAT1 :

Pattern: [KRH~~Q~~FE]-x-G-[IVL~~F~~AC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?>26784

339 - 346: KpGIPyPI



6h04_B

[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>6h04_B%0A&output=nice](#))

USERPAT1 :

Pattern: [KRH~~Q~~FE]-x-G-[IVL~~F~~AC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?>26784

408 - 415: EyGAaLaW



6h04_C

[View all PROSITE motifs hits on sequence](#)
([PSScan.cgi?seq=>6h04_C%0A&output=nice](#))

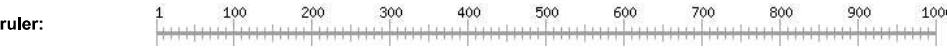
USERPAT1 :

Pattern: [KRH~~Q~~FE]-x-G-[IVL~~F~~AC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)]

26784

187 - 194: KsGFsFgF

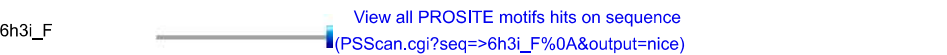
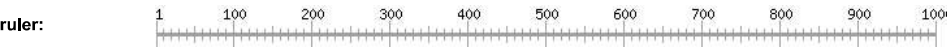


USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)]

26784

1412 - 1419: KwGInLpF

1564 - 1571: QyGFgFnL



USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)]

26784

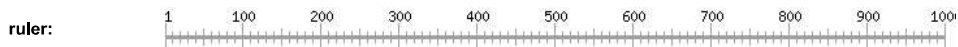
216 - 223: KIGVsLeL



USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>)]

26784

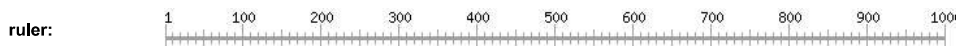
480 - 487: EIGLkVdL

**USERPAT1 :**

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175)]

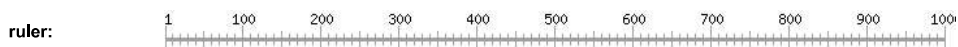
528 - 535: RiGALVLF

**USERPAT1 :**

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175)]

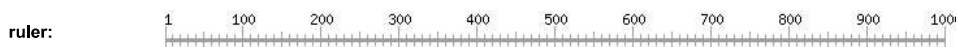
229 - 236: KKGIyvtV

**USERPAT1 :**

Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**

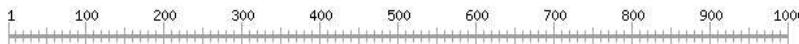
Approximate number of expected random matches [Ref: [PMID 11535175 \(http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175\)](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175)]

150 - 157: QvGLrYdL



USERPAT1 :

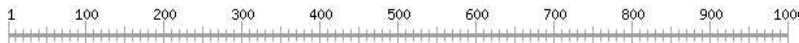
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>) 26784**277 - 284:** RaGLpLhF**488 - 495:** ELGfKFsV**ruler:**

6v81_A

[View all PROSITE motifs hits on sequence \(PSScan.cgi?seq=>6v81_A%0A&output=nice\)](http://prosite.expasy.org/cgi-bin/prosite/scanprosite/ScanView.cgi?seq=>6v81_A%0A&output=nice)**USERPAT1 :**

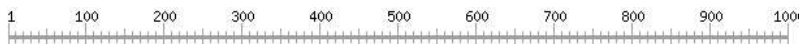
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>) 26784**304 - 311:** ELGvpVtF**500 - 507:** RqGAeLaW**ruler:**

6ve2_O

[View all PROSITE motifs hits on sequence \(PSScan.cgi?seq=>6ve2_O%0A&output=nice\)](http://prosite.expasy.org/cgi-bin/prosite/scanprosite/ScanView.cgi?seq=>6ve2_O%0A&output=nice)**USERPAT1 :**

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

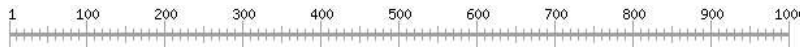
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>) 26784**192 - 199:** RaGLIMvF**ruler:**

6w6m_A

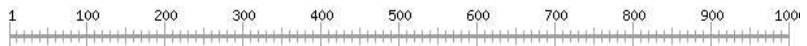
[View all PROSITE motifs hits on sequence \(PSScan.cgi?seq=>6w6m_A%0A&output=nice\)](http://prosite.expasy.org/cgi-bin/prosite/scanprosite/ScanView.cgi?seq=>6w6m_A%0A&output=nice)**USERPAT1 :**

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

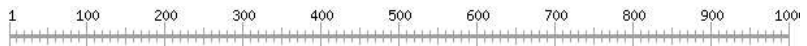
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?PMID=11535175>) 26784

178 - 185: EIGVrwgV**ruler:**

6wum_a

**USERPAT1 :**Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]**52 - 59:** EeGFgVfL**273 - 280:** RIGVlYpL**433 - 440:** RkGLqVgV**437 - 444:** QvGVgIsF**ruler:**

6x1k_A

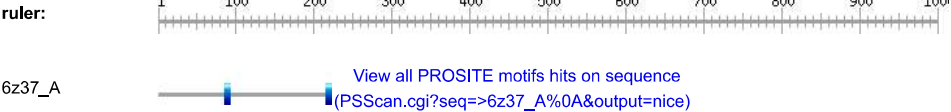
**USERPAT1 :**Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]**115 - 122:** QaGLsYrI**ruler:**

6z34_A

**USERPAT1 :**Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]**43 - 50:** HIGLgMdI**187 - 194:** RIGLtYqL

199 - 206: RvGIvYnF

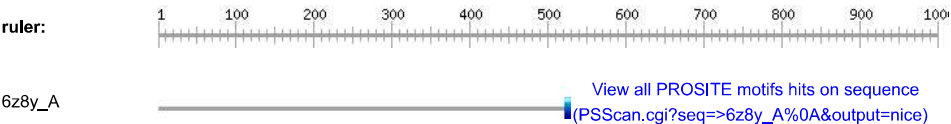
310 - 317: RyGAnkVvF



USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175>)]
26784

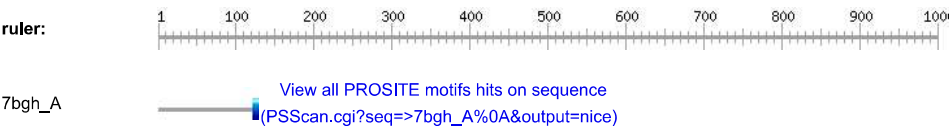
85 - 92: RfGAgLfV

213 - 220: RIGLtYkL



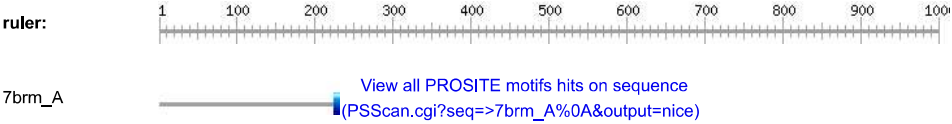
USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175>)]
26784

511 - 518: ELGAsYqL

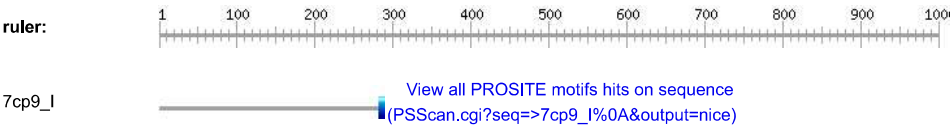


USERPAT1 :
Pattern: **[KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]**
Approximate number of expected random matches [Ref: [PMID 11535175](http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175) (<http://www.ncbi.nlm.nih.gov/80/entrez/query.fcgi?PMID=11535175>)]
26784

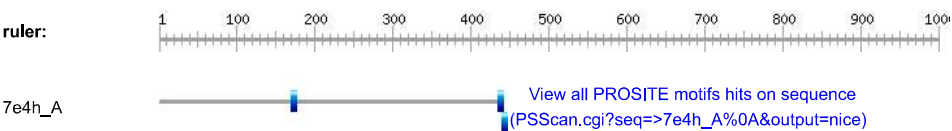
121 - 128: RgGAeFsw



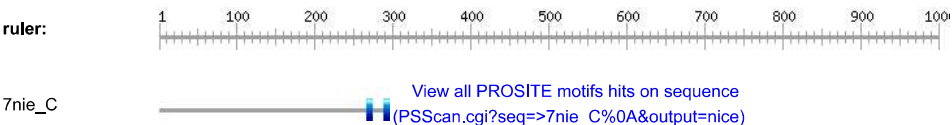
USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]
221 - 228: EtGViFlI



USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]
278 - 285: QcGFgLtI



USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]
167 - 174: QrGARLsL
427 - 434: RkGFqFgL
431 - 438: QfGLgLaf



USERPAT1 :

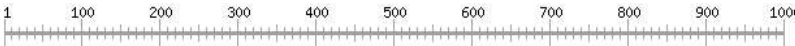
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?>26784

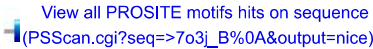
263 - 270: RpGVkLtL

285 - 292: KLGLaLeL

ruler:



7o3j_B



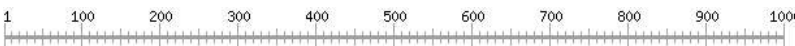
USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?>26784

16 - 23: RvGAYnL

ruler:



7omm_D



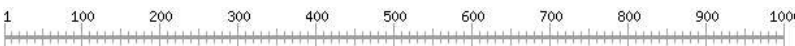
USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?>26784

312 - 319: KaGLtFlV

ruler:



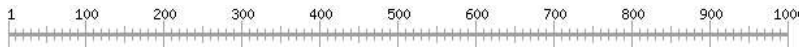
7rj5_D



USERPAT1 :

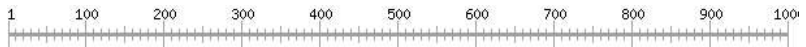
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?>26784

168 - 175: ErGAwVaV**ruler:**

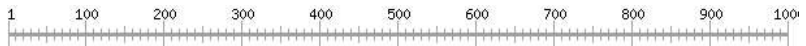
7vku_X

[View all PROSITE motifs hits on sequence](#)
(PSScan.cgi?seq=>7vku_X%0A&output=nice)

USERPAT1 :Pattern: [KRH~~Q~~FE]-x-G-[IVL~~F~~AC]-x-[IVL~~F~~MYW]-x-[IVL~~F~~W]Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]**146 - 153:** KiGCgLqF**ruler:**

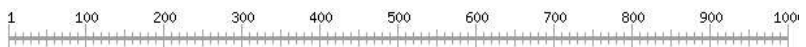
7vtz_A

[View all PROSITE motifs hits on sequence](#)
(PSScan.cgi?seq=>7vtz_A%0A&output=nice)

USERPAT1 :Pattern: [KRH~~Q~~FE]-x-G-[IVL~~F~~AC]-x-[IVL~~F~~MYW]-x-[IVL~~F~~W]Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]**325 - 332:** FfGAmYdL**ruler:**

7xzi_4

[View all PROSITE motifs hits on sequence](#)
(PSScan.cgi?seq=>7xzi_4%0A&output=nice)

USERPAT1 :Pattern: [KRH~~Q~~FE]-x-G-[IVL~~F~~AC]-x-[IVL~~F~~MYW]-x-[IVL~~F~~W]Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)]**145 - 152:** RiGCrIdL**ruler:**

7xzi_E

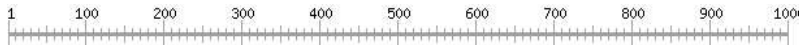
[View all PROSITE motifs hits on sequence](#)
(PSScan.cgi?seq=>7xzi_E%0A&output=nice)

USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)**326 - 333:** RhGcGmTL

ruler:



7xzi_9

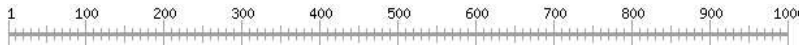
[View all PROSITE motifs hits on sequence](#)
(PSScan.cgi?seq=>7xzi_9%0A&output=nice)

USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)**242 - 249:** HaGcGLmL

ruler:



8bmxC

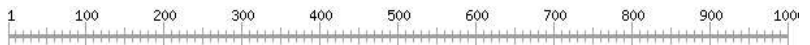
[View all PROSITE motifs hits on sequence](#)
(PSScan.cgi?seq=>8bmxC%0A&output=nice)

USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)**430 - 437:** EhGFtYdV

ruler:



8bmxD

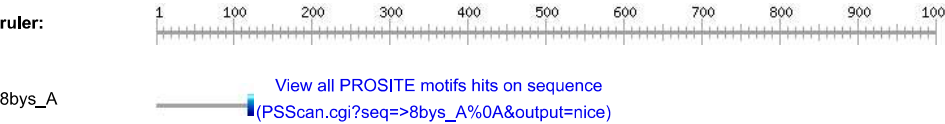
[View all PROSITE motifs hits on sequence](#)
(PSScan.cgi?seq=>8bmxD%0A&output=nice)

USERPAT1 :

Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]

Approximate number of expected random matches [Ref: [PMID 11535175](#) (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)

67 - 74: RdGIGWiV



8bys_A

USERPAT1 :
Pattern: [KRHQFE]-x-G-[IVLFAC]-x-[IVLFMYW]-x-[IVLFW]
Approximate number of expected random matches [Ref: PMID 11535175 (<http://www.ncbi.nlm.nih.gov:80/entrez/query.fcgi?26784>)

118 - 125: FdGAqLnV

horizontal scaling:

do not show text labels: ☐

do not show sites in hits: ☐

do not show ranges in hits: ☐

[Shaded Alignment of hits \(/cgi-bin/aligner?color=1&zgff=264838152952.scan.gz\)](#)