

```

GSNLLRPPQQFPEALRECPQQESDIVFLIDGSGSINNIDFQKMKEFVSTVMEQFKKSKTL
GSNLRQQPQKFPEALRGCPQEDSDIAFLIDGSGSIIPHDFRRMKEFVSTVMEQLKSKTL
**** . ** .***** ** .*** ***** ** .***** *****
FSLMQYSDEFRIHFTFNDFKRNPSRSHVSPKQLNGRKTASGIRKVVRELFHKTNGAR
FSLMQYSEEFRIHFTFKEFQNNPFRSLVKPITQLLGRTHATGIRKVVRELFNITNGAR
***** .***** .* ** ** * ** * ** * ** * .***** . *****
ENAAKILVVITDGEKFGDPLDYKDVIPEADRAGVIRYVIGVGNAFNKPQSRRELDTIASK
KNAFKILVVITDGEKFGDPLGYEDVIPEADREGVIRYVIGVDADRSEKSRQELNTIASK
** ***** ** ***** ** ***** ** .*** ** *****
PAGEHVFQVDNFEALNTIQNLQEKIFA Murine I domain
PPRDHVFQVNNFEALKTIQNLREKIFA Human I domain
* .***** ***** ***** .***** 77 % Identity

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