prd

MNSGOGRVNOLGGVELNGRPLPNNLRLKLVEMAADGLRPCVLSROLRVSH

mel

2L: homeobox

50

| sim | MNSGQGRVNQLGGVFINGRPLPNNIRLKI VEMAADGI RPCVI SRQLRVSH | 50 50 |
|------------|---|------------|
| | Paired | |
| mel | GCVSKI LNRYQETGSI RPGVI GGSKPRI ATPEI ENRI EEYKRSSPGMFSW | 100 |
| sim | GCVSKI LNRYQETGSI RPGVI GGSKPRI ATPEI ENRI EEYKRSSPGMFSW | 100 |
| mel | EI REKLI REGVCDRSTAPSVSAI SRLVRGRDAPLDNDMSSASGSPAGDGT | 150 |
| sim | EI REKLI REGVCDRSTAPSVSAI SRLVRGRDAPLDNDMSSASGSPAGDGT | 150 |
| mel sim | KASSSCGSDVSGGHHNNGKPSDEDI SDCESEPGI ALKRKQRRCRTTFSAS KASSSCGSDVSGGHHNNGKPSDEDI SDCESEPGI ALKRKQRRCRTTFSAS Homeobox | 200 200 |
| mel sim | QLDELERAFERTQYPDI YTREELAQRTNLTEARI QVWFSNRRARLRKQHT QLDELERAFERTQYPDI YTREELAQRTNLTEARI QVWFSNRRARLRKQHT Double-click here to edit text. | 250 250 |
| mel | SVSGGAPGGAAASVSHVAASSSLPSVVSSVPSMAPLAMMPGSLDPATVYQ | 300 |
| sim | SVSGGAPGGAAASVSHVAASSSLPSVVSSVPSMAPLAMMPGSLDPATVYQ | 300 |
| mel | QQYDFYGSHANI SVSAAAPMASSNLSPGI TTTPPHHHQFYNPSANTASYI | 350 |
| sim | QQYDFYGSHANI SVSAAAPMASSNLSPGI TTTPPHHHQFYNPSANTASYI | 350 |
| mel | MPGENGNTTPTGNI I VSSYETQLGSVYGTETETHQTMPRNESPNESVSSA | 400 |
| sim | MPGENGNTTPTGNI I VSSYETQLGSVYGTETEAHQTMPRNESPNESVSSA | 400 |
| mel | FGQLPPTPNSLSAVVSGAGVTSSSGANSGADPSQSLANASAGSEELSAAL | 450 |
| sim | FGQLPPTPNSLSAVVSGAGVTSSSGANSAADPSHSLANASAGSEDLSAAL | 450 |

pro

2L: homeobox

| mel | KVESVDLI AASQSQLYGGWSSMQALRPNAPLSPEDSLNSTSSTSQALDVT | 500 |
|------------|---|-----|
| sim | KVESVDLI AASQSQLYGGWSSMQALRPNAPLSPEDSLNSTSSTSQALDVT | 500 |
| mel | AHQMFHPYQHTPQYASYPAPGHAHSHHGHPHAPHPHAHPHPQYAGAHPHY | 550 |
| sim | AHQMFHPYQHTPQYASYPAPGHAHSHHGHPHAPHPHAHPHPQYAGAHPHY | 550 |
| mel sim | PPPSSSAHFMPQNFNAAAFPSPSKVNYTTMPPQPFYPSWY 590 PPPSSSAHFMPQNFNAAAFPSPSKVNYTTMPPQPFYPSWY 590 | |