

CLUSTAL O(1.2.4) multiple sequence alignment

```
2JFQ      MNKPIGVIDSGVGGTLVAKIMRQLPNETIYYLGDIGRCPYGPRPGEQVKQYTVEIARKL      60
P63637|MURI_STAAM MNKPIGVIDSGVGGTLVAKIMRQLPNETIYYLGDIGRCPYGPRPGEQVKQYTVEIARKL      60
Q6GHT5|MURI_STAAR MNKPIGVIDSGVGGTLVAKIMRQLPNETIYYLGDIGRCPYGPRPGEQVKQYTVEIARKL      60
Q2YXC5|MURI_STAAB MNKPIGVIDSGVGGTLVAKIMRQLPNETIYYLGDIGRCPYGPRPGEQVKQYTVEIARKL      60
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```
2JFQ      MEFDIKMLVIACNTATAVALEYLQKTLISVIGVIEPGARTAIMTTRNQNVLVLGTEGTI      120
P63637|MURI_STAAM MEFDIKMLVIACNTATAVALEYLQKTLISIPVIGVIEPGARTAIMTTRNQNVLVLGTEGTI      120
Q6GHT5|MURI_STAAR MEFDIKMLVIACNTATAVALEYLQKTLISIPVIGVIEPGARTAIMTTRNQNVLVLGTEGTI      120
Q2YXC5|MURI_STAAB MGFDIKMLVIACNTATAVALEYLQKTLISIPVIGVIEPGARTAIMTTRNQNVLVLGTEGTI      120
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```
2JFQ      KSEAYRTHIKRINPHVEVHGACPGFVPLVEQMRYSPTTITSIVIHQTLKRWRNSESDTV      180
P63637|MURI_STAAM KSEAYRTHIKRINPHVEVHGACPGFVPLVEQMRYSPTTITSIVIHQTLKRWRNSESDTV      180
Q6GHT5|MURI_STAAR KSEAYRTHIKRINPHVEVHGACPGFVPLVEQMRYSPTTITSIVIHQTLKRWRNSESDTV      180
Q2YXC5|MURI_STAAB KSEAYRTHIKRINPHVEVHGACPGFVPLVEQMRYSPTTITSIVIHQTLKRWRNSESDTV      180
*****
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```
2JFQ      ILGCTHYPLLYKPIYDYFGGKKTVISSGLETAREVSALLTFSNEHASYTEHPDHRFFATG      240
P63637|MURI_STAAM ILGCTHYPLLYKPIYDYFGGKKTVISSGLETAREVSALLTFSNEHASYTEHPDHRFFATG      240
Q6GHT5|MURI_STAAR ILGCTHYPLLYKPIYDYFGGKKTVISSGLETAREVSALLTFSNEHASYTEHPDHRFFATG      240
Q2YXC5|MURI_STAAB ILGCTHYPLLYKPIYDYFGGKKTVISSGLETAREVSALLTFSNEHASYTEHPDHRFFATG      240
*****
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```
2JFQ      DTHITNIIKEWLNLSVNVERISVND      266
P63637|MURI_STAAM DPTHITNIIKEWLNLSVNVERISVND      266
Q6GHT5|MURI_STAAR DTHITNIIKEWLNLSVNVERISVND      266
Q2YXC5|MURI_STAAB DTHITNIIKEWLNLSVNVERISVND      266
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