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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain C12_S_E7_2]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain C12_S_E7_4]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D08_H_8_6]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D09_K_2_3]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D09_K_4_1]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D09_K_4_2]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D10_B_F1_6]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D10_B_F3_5]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D10_B_S3_16]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D10_K_S11_6]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D11_T_E18]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis pseudocinerea, strain BpsVD110]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis pseudocinerea, strain BpsD11_KL_cal2]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis pseudocinerea, strain BpsD12_E_cal10]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D13_E_IF04]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D13_E_IL4]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D13_E_IL7]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D13_E_IL12]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain G09_S33]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D08_H_8_4]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain N11_K_W06]
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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain N11_S_E10]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain N11_S_E15]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain T4]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis calthae, strain BcalMUCL2830]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis cinerea, strain D08_H_8_3]

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>Bcin05g01790(mrr1), partial sequence [organism=Botrytis pseudocinerea, strain BpsVD256]
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