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Gmet_R3001  (+)  158167   158284   GATAGACGATAATACTAAACCATCCGCGAGGGTGGGACGGAAAGCCTACAGGGTCTCCCC-GAGACAGCCGGGTCGCC-AAATATCTCAACAG--GATATGC-CGGC--CCCGGCTTTTTTTCATG
Gmet_R3002  (-)  264958   265068   AATAGACGACAATACTAAACCATCCGCGAGGATGGGGCGGAAAGCCCATAGGGTCTCACC-GAGACAGCCGGGTTGCCGAAATATTTTTT-------------CGGCGACCCGGCTTTTTCGTTT
Gmet_R3003  (-)  271381   271497   GATACACGACAATACTAAACCATCCGCGAGGATGGGGCGGAAAGCCCACAGGGTCTCACC-GAGACAGCCGGGTTGCCGAAATGTCACCA-----CATTTT--CGACAGCCCGGTTTTTTTTGTGC
Gmet_R3004  (+)  325752   325867   GATAGACGATAATACTCAACCATCCGCGAGGATGGGGCGGAAAGCCTACAGGGTCTCTTC-GAGACAGCCGGGTTGCCGAAATATCTCACA----GATATG--CGGC--CCCGGCTTTTTTTGTGT
Gmet_R3005  (+)  1079463  1079581  GATAT-CGACAATACTAAACCATCCGCGAGGGTGGGACGGAAAGCCTACAGGGTCTCTCT-GAGACAGCCGGGATGCCGAAATATCACAATTCG-TGATGCT-CGGT--CCCGGCATTTCTTTTTT
Gmet_R3006  (-)  1211932  1212051  GATAG-CGATACTACTAAACCATCCGCGAGGATGGGGCGGAAAGCCTACAGGGTCTCATC-GAGACAGCCGGGTTGCCGAACTATCACACCACA-GATAGG--CGGCGACCCGGCTTTTTATTTT
Gmet_R3007  (-)  1212114  1212231  GATAG-CGATACTACTAAACCATCCGCGAGGATGGGGCGGAAAGCCTACAGGGTCTCATC-GAGACAGCCGGGTTGCCGAACTATCACAATA---GATAGG--CGGCGACCCGGCTTTTTGTTT
Gmet_R3008  (-)  1324366  1324481  GATAGTCGATAATACTAAACCATTCGTGAGAATGGGGCGGAAAGCCTACAGGGTCTTACT-GAGACAGCCGGGTTGCCGAAATATCACCA-----GATATT--CG-CAACCCGGCTTTTTTGTTG
Gmet_R3009  (-)  1324505  1324620  GATAGACGACAATACTAAACCATTCGCGAGAATGGGGCGGAAAGCCTACAGGGTCTTATT-GAGACAGCCGGGTTGCCGAAATATCATTAT----GATAA-T-CGGC--CCCGGCTGTTCTGTTT
Gmet_R3010  (+)  1334236  1334351  GATAGACGACAATACTAAACCATCCGCGAGGGTGGGACGGAAAGCCTACAGGGTCTCCCC-GAGACAGCCGGGTCGCCGAAATATCGCAT----CGATATTC-CGA---CCCGGCTTTTTTTGTG
Gmet_R3011  (+)  1911269  1911387  GATAGACGACAATACTAAACCATCCGCGAGGATGGGGCGGAAAACCCACAGGGTCTCCCT-GAGACAGCCGGGTTGCCGAAATATCAATT-----GATATATATGGCAACCCGGCTGTTTTATTA
Gmet_R3012  (+)  1977790  1977908  GATAGACGATAATACTCAACCATTCGCGAGAATGGGGCGGAAAGCCTATAGGGTCTCATG-CAGACAGCCGGGTTGCCGAAATATCATTTT---AGAATGATGCGGC--CCCGGCTTTTTTGCGT
Gmet_R3013  (+)  2280689  2280806  GATAGACGATAATACTCAACCATCCGTGAGGATGGGGCGGAAAGCCTATTGGGTCTCACC-GAGACAGCCGGGTTGCCGAAATATCTGACCAA-AGATATA---GGC--CCCGGCTTTTTTTGCT
Gmet_R3014  (+)  2447616  2447732  GATACACGACAATACTCAACCATCCGCGAGGATGGGGCGGAAAGCCTACAGGGTCTCCAG-CAGACAGCCGGGTTGCCGAAATATCACT-----CGATGATAGCGGT--CCCGGCTTTTTGTGC
Gmet_R3015  (+)  3913061  3913175  GATAGACGACAATACTTAACCATTCGCGAGAATGGGGCGGAAAGCCTATAGGGTCTCATG-GAGACAGCCGGGTTGCCGAAATATCACA------CGATATT--TGGT--CCCGGCTTTTTTATTG
Gmet_R3016  (+)  3913269  3913383  GGTAGACGATAATACTAAACCATTCGCGAGAATGGGGCGGAAAGCCTATAGGGTCTCCCT-GAGACAGCCGGGTTGCCGAAATATCACG-----CGATATT--TGGT--CCCGGCTTTTTTGTGT
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