

The AraC-type transcriptional regulator GliR (PA3027) activates genes of glycerolipid metabolism in *Pseudomonas aeruginosa*

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Supplementary Materials

Text S1: Sequences of PA3027 ChIP-seq peak summits used for motif searching (peaks marked by numbers and gene ID identified in the vicinity, see Table 2).

>1 PA0235

CCGCGAGCGGTTATGCGCCCTACGCGATGCCGACCCCTGGGGTAGGGCGAATAACGCCACCGGCGTTATCCGCCG
ATACACCGAAGCCCGGCGGCGGATAACCGCAAGCGGTTATGCGCCCTACGCGATGCCGACCCCTGGGGTAGGGCG
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>2 PA4045

TAACGCCGGTGGCGTTATTGCGCCCTACGGCAATTGGGGGGTAGGGCGAATAACCGCAACGCGGTTATCGCTGCCA
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GGGTAGGGCGAATAACCGCAACGCGGTTATCCGCCGTTGTGCTGGGTTT

>3 PA3572

CGCCTGAGCGCGGATCAGGCCGTTTCGCTGTTTTAGTTTCGGGCAGCACATGGGGCGCGCGCAGGCGCGCCTCGAGG
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>4 PA3464

ATGTCGGGAAAGTCTAGCCGCTGGCGCCTGGGCGCCATTGATGGCGGTCATGGGGCCGGCCCGGCGTGGATAGA
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CTTGCCGGCACCGAGTCGAGACAAGAGCGAAAAGCCGATGAAGTTGCTGT

>5 PA3464

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>6 PA3397

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>7 PA3391

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ATGTCGCGGAACCTGGCGCAGTTGGACCCGATCGAAGATGCCACCGCGCACGTAGCCGGAACCCCTGAAGGAATAC
TCGCCGTCGCCGAGCACGACGAAGGCGTATTGCCTGGCTTCAGCGACGC

>8 PA3342

AGCGTCTCTACGATCTCTACGAGCATGCCGATCGCGCGCTTTACGAAGCCAAGGCGTTTCGGCCGTAACCAGGTGG
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GGTTCGACAGGAAAAAATGCCAGGCACAGCCGCTCTTCGACAGAGCTGCCG

>9 PA3341

CACATTTTCGGCACCTTGCTAGCCCAGACTTCCCGTGCTTGGCGCGCTGAACTCGACCGCCGCCCTCAGTCACCTC
GGCCTGTCCCAGGCGCGCTGGTTGGTGTGTTGCATCTGGCGCGCCATCGCGATTTCGCCACCCAGCGCGAGTTG
GCCAATCGGTTCGGGGTCGAAGGCCCCACCTGGCGCGGCTGCTCGACGG

>10 PA3309

CGGCGATTTCCTCGCTGGCGCGCTCCTGGCAGACGCGATCGAAGGTGGCGCCTTCGACGTTGCTCGCCTCGACCT
TCTCGCGGGCCTTGGCCAGGACGTCATCGGCGGCTTTCAGCGCGCTTCGCGCAGCGGCTCGGGTTGTAGTACA
CCGCGTAGTCGGGAAGGTGGCGCAGCGGGCTGTTCGACGATGGTGATGACG

>11 PA3083

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>12 PA2742

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>13 PA2601

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>14 PA2601

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>15 PA2147

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>16 PA1414

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>17 PA1196

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>18 PA0928

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>19 PA4610

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TGCCCATATGAAAAAGGGCGCCGAGGCGCCCTTTTTTCATGCCGTTGTTCG

>20 PA4772

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TGGCTGCTCCGGCGTCGTGCGCCCGCGCCGGGTTCGGGCAAAAAATCCGTC

>21 PA5208

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>22 PA5243

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>23 PA5294

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>24 PA3026

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CACGCTCCAGGAGCCGGGCTGGACGAAACGCGCCGAATGCAGGCCGAACAGGGGATCGCCGGAATGCTCGCAGAA
ATAGCCGAGCAGGTGCTCGTGGGCTTCGCCCCGCGAGGCGCTGGCGATTGTCGTTGAGCTGGTCGGCCTGGAGGCC
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CGAGGTGGAGCCAAGGGTCTGCATGGTCGTTGTTCTTGTCGTGGCGGGTGGAAACAGGCCGTCCGGCGTGCGTGCC
GGCTGTCTTGATCGGTACAGGGAGTGTCTCGATCCGACAGTGACAGGCGAGGGCGGCTGGCTAGTATAGGCAGC
CAGAAAGGTTCCGATAAACAGGTTGCCGCGACAAGCGAGGGGAGAGGTTTC