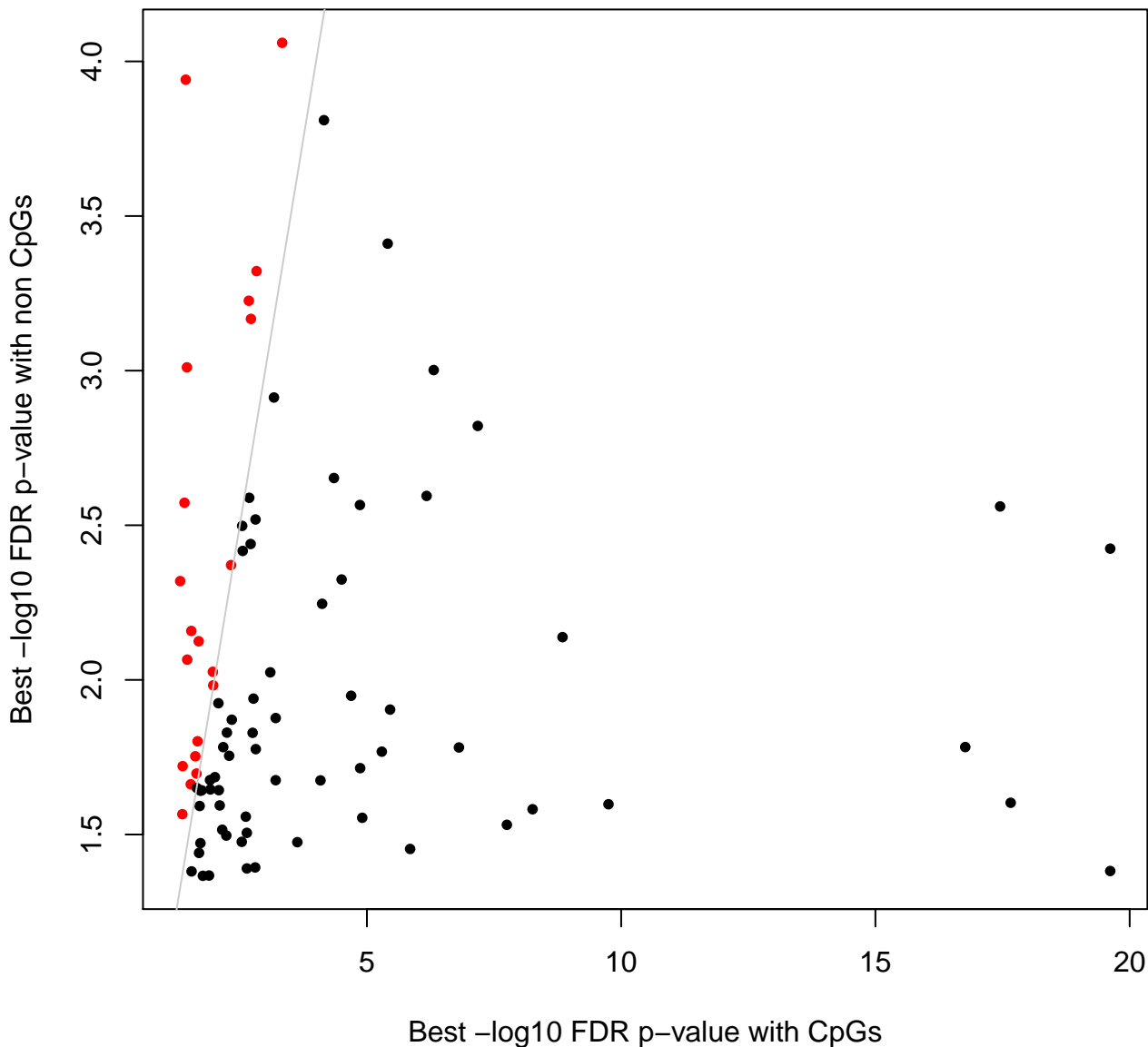
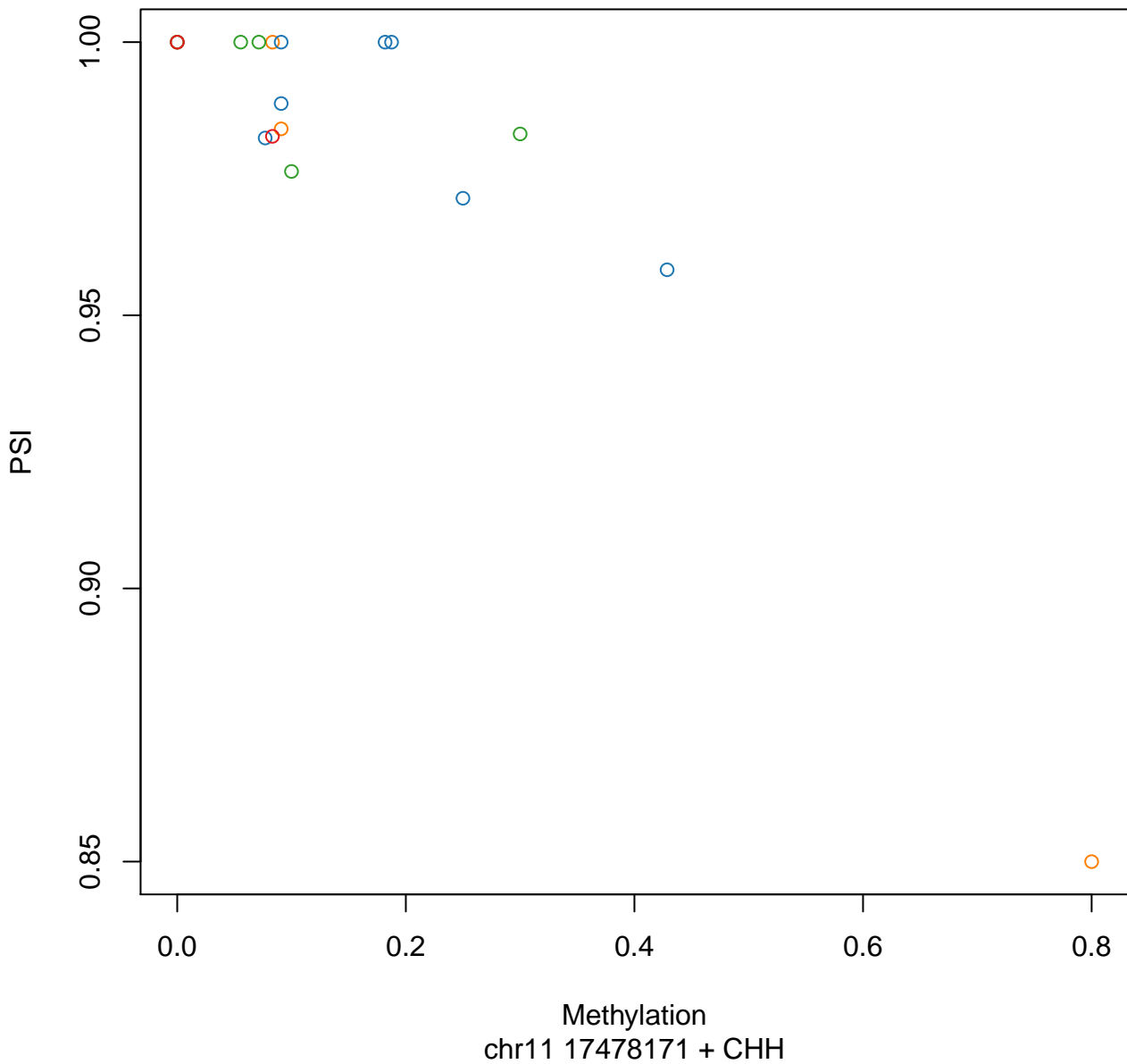


85 common psi meQTLs
Based on top 5000 psis expressed in Glia



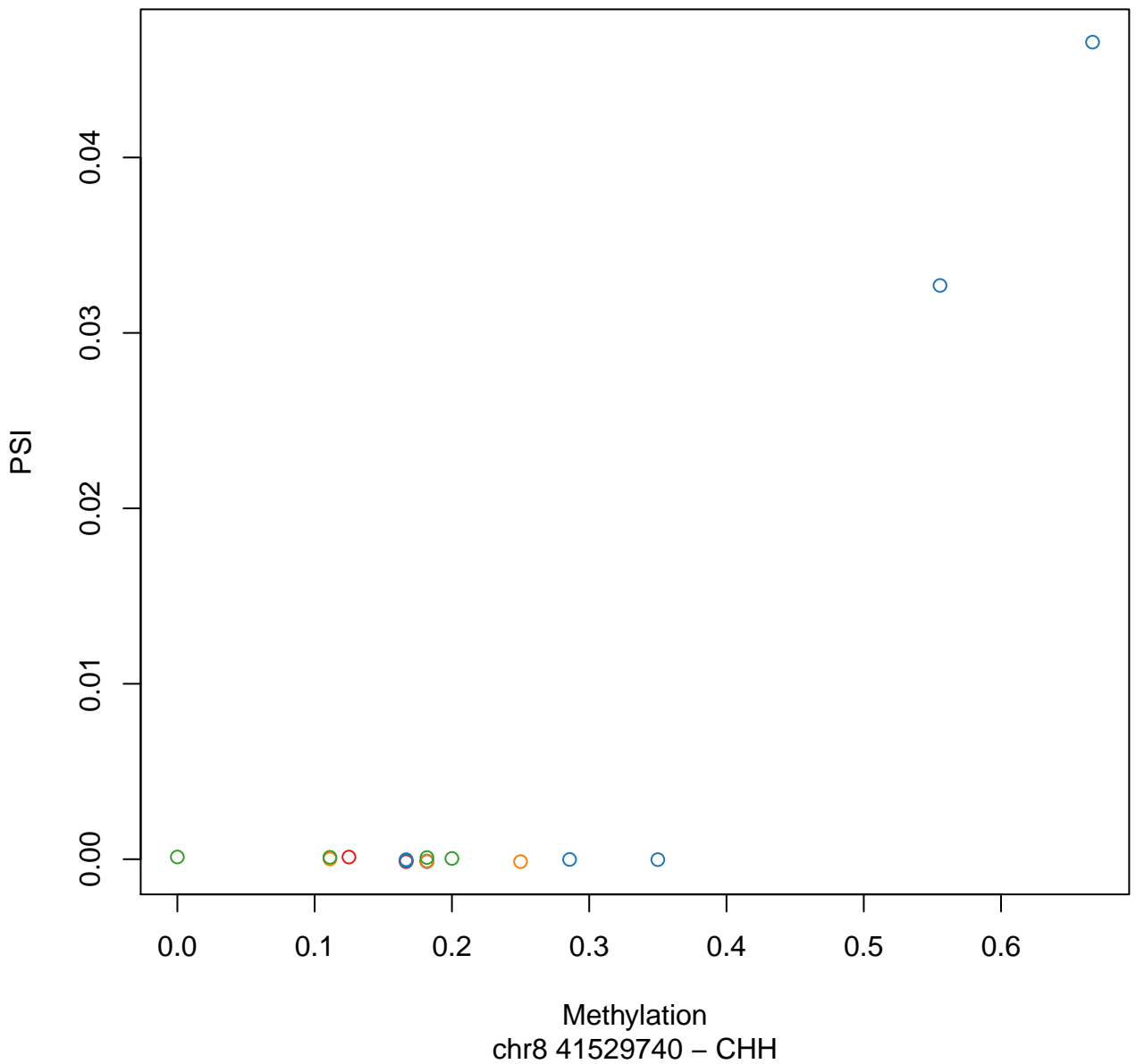
psi ENSG00000006071.12_2 FDR 0.000115

ABCC8



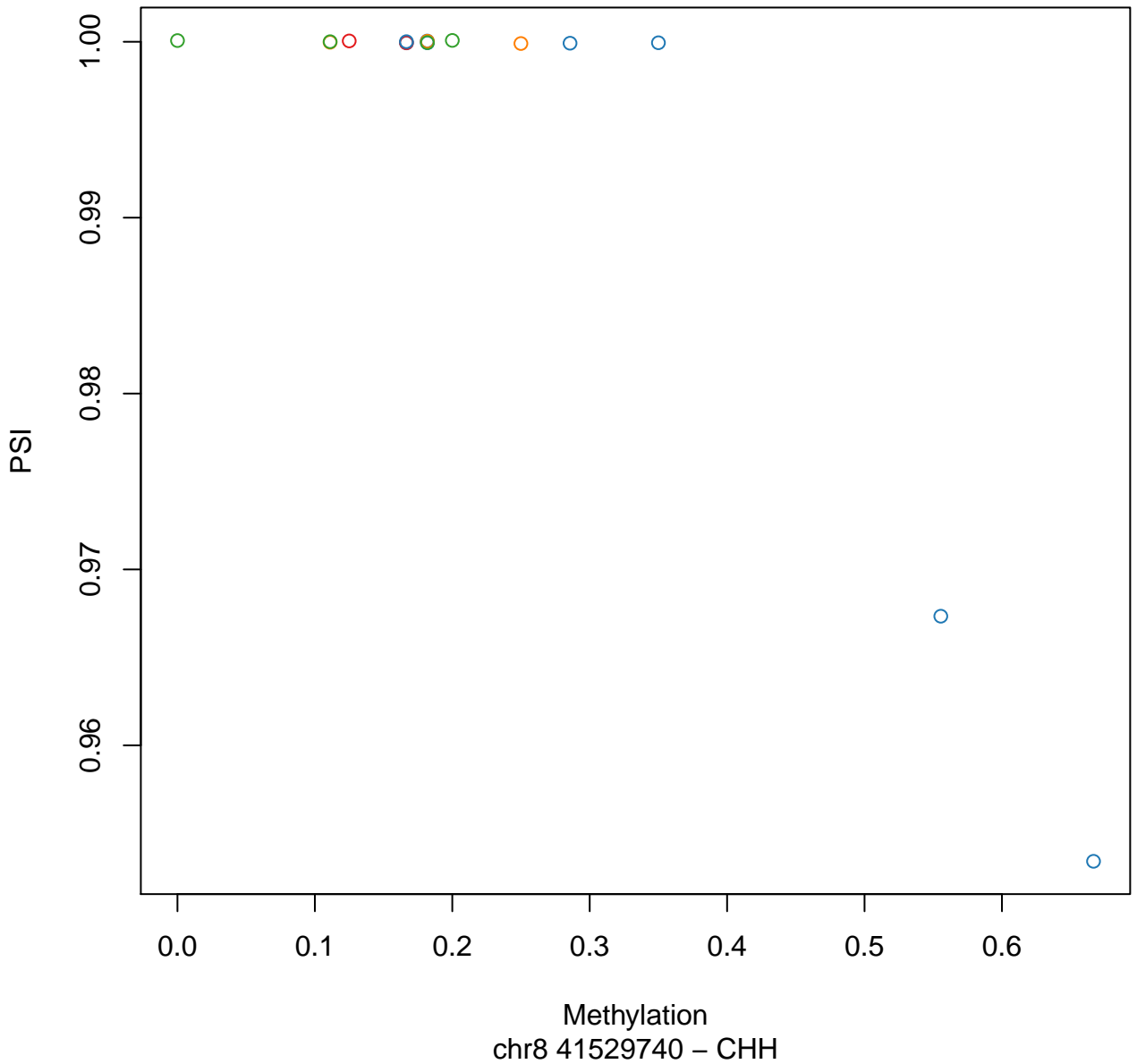
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ANK1



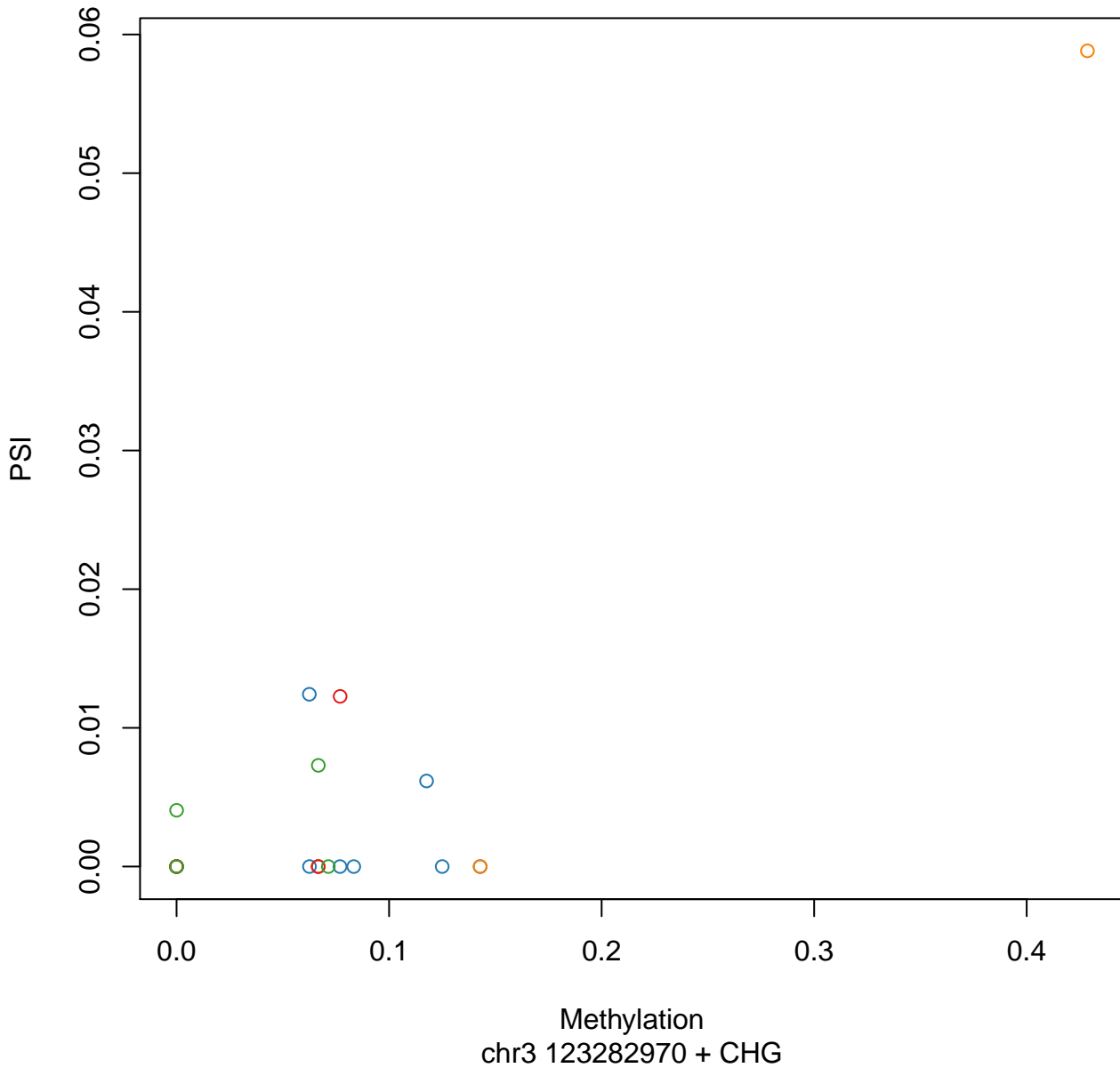
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ANK1



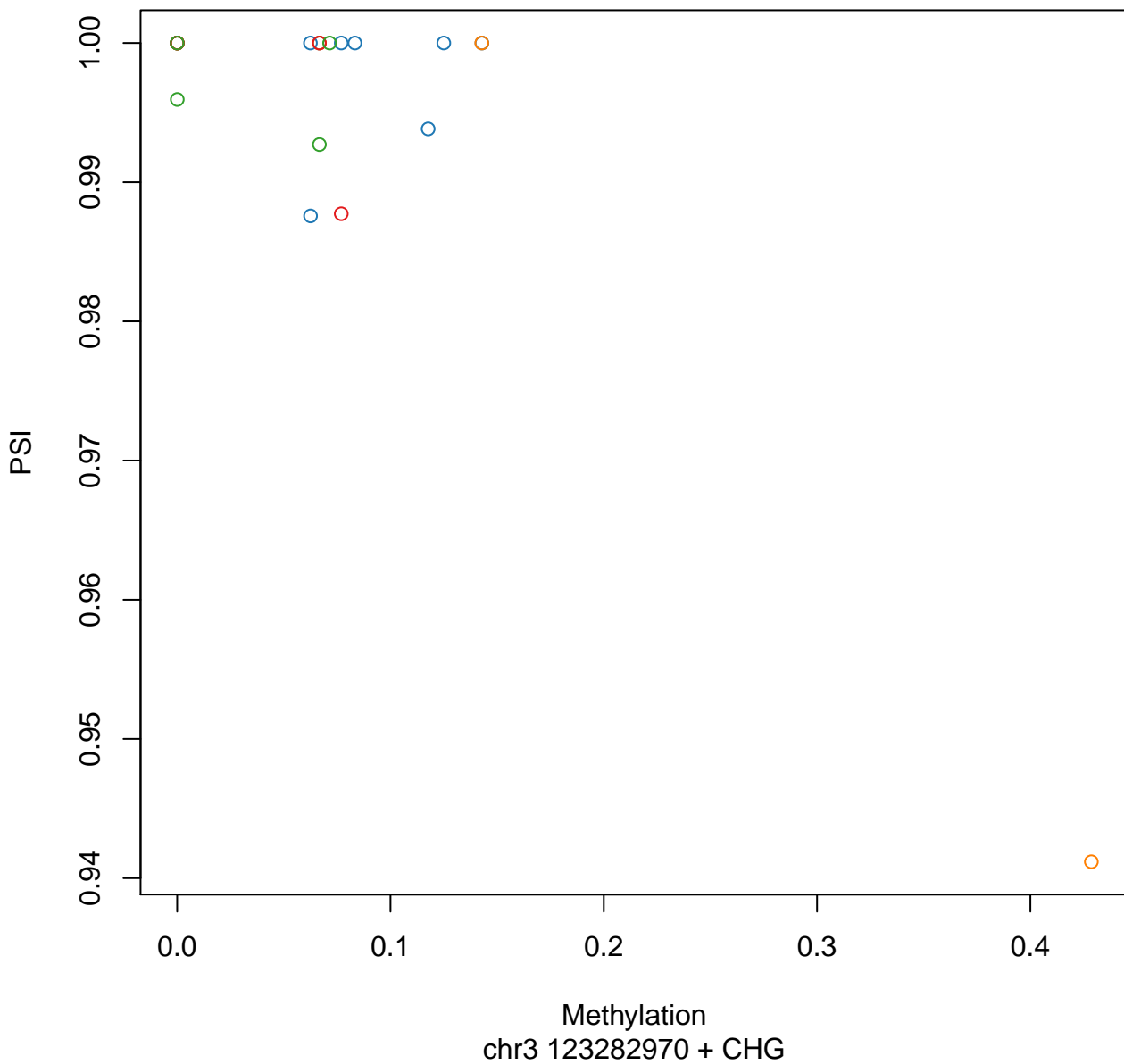
psi ENSG00000206527.9_1 FDR 0.00267

HACD2



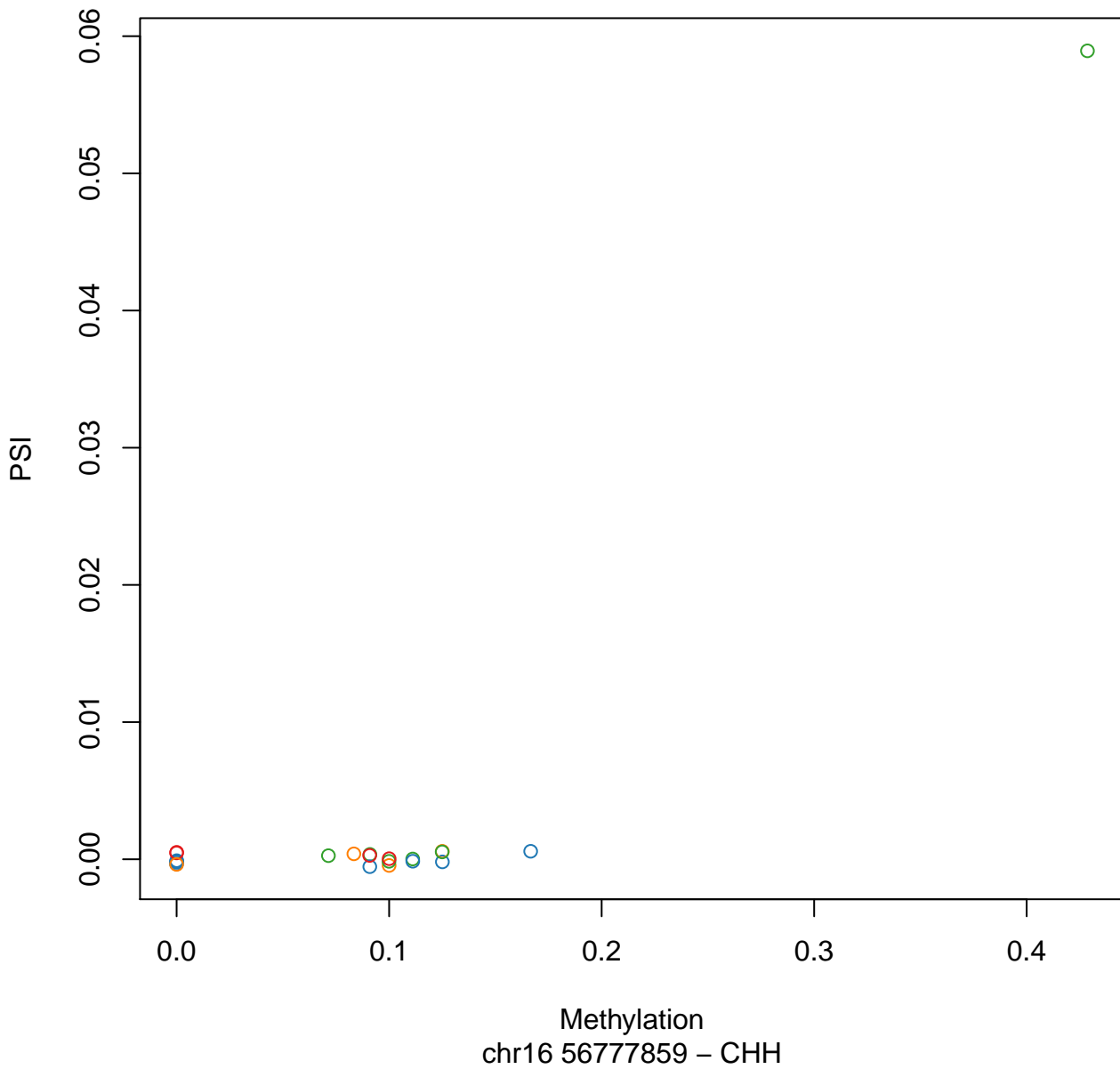
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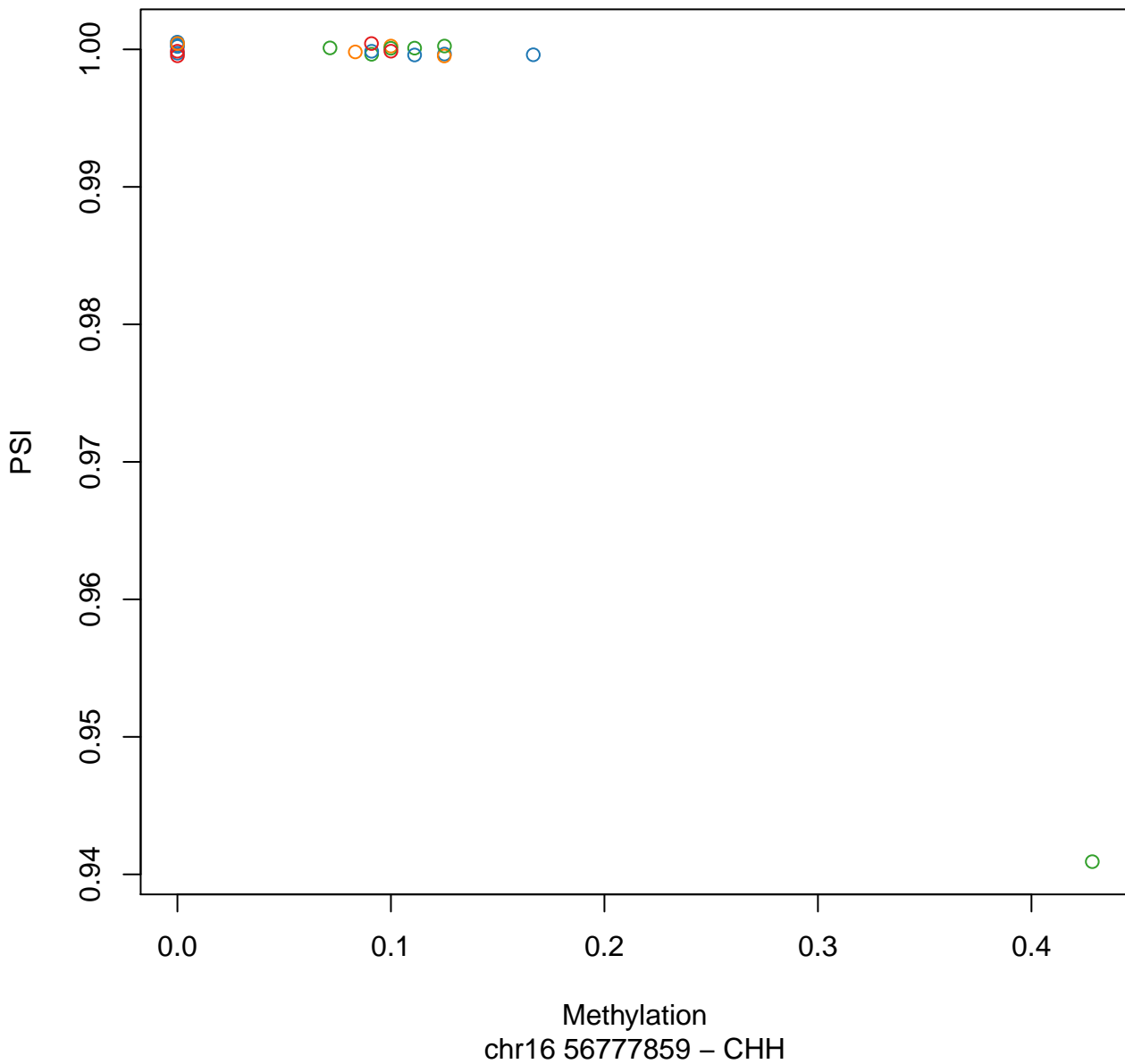
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NUP93

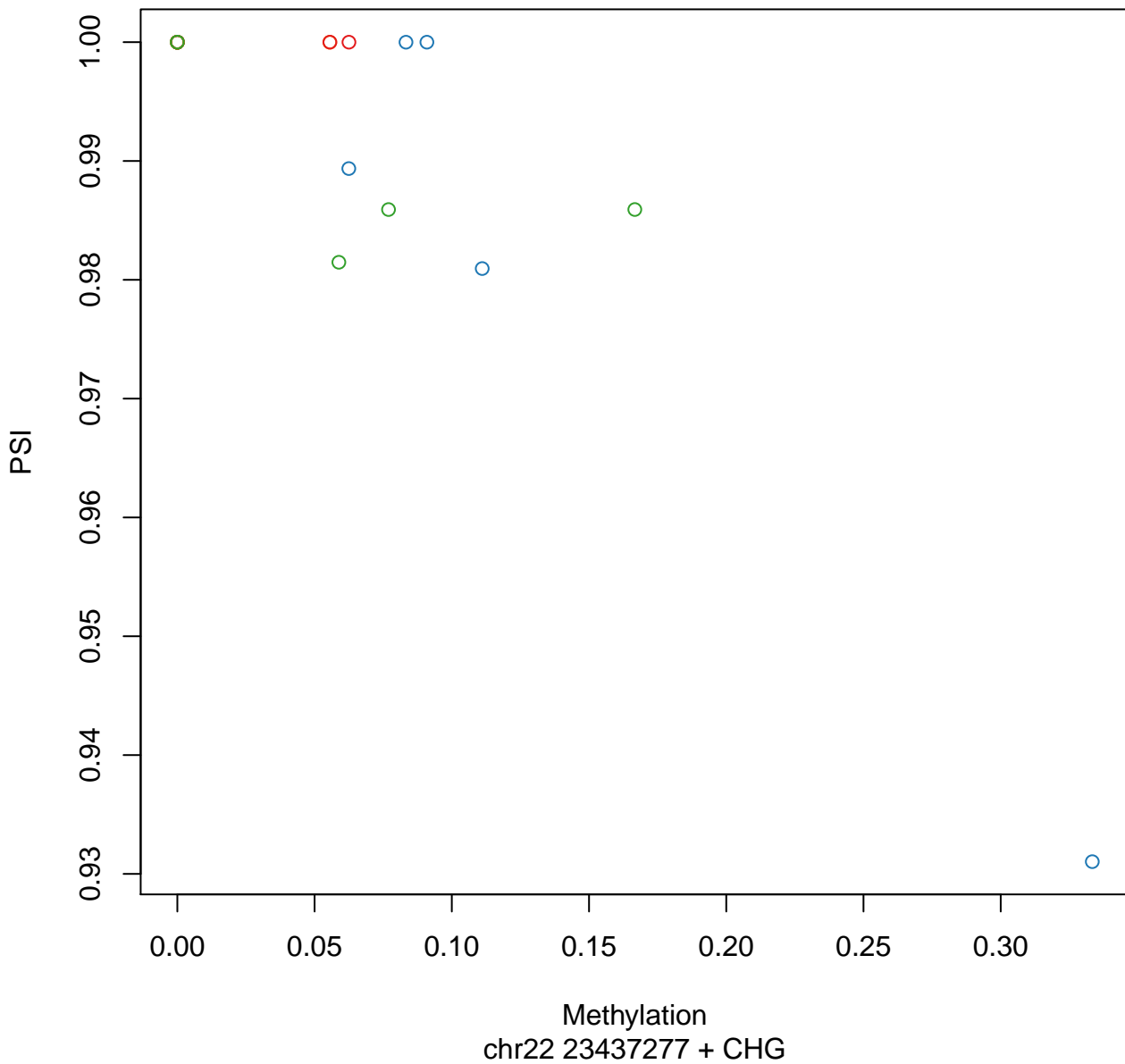


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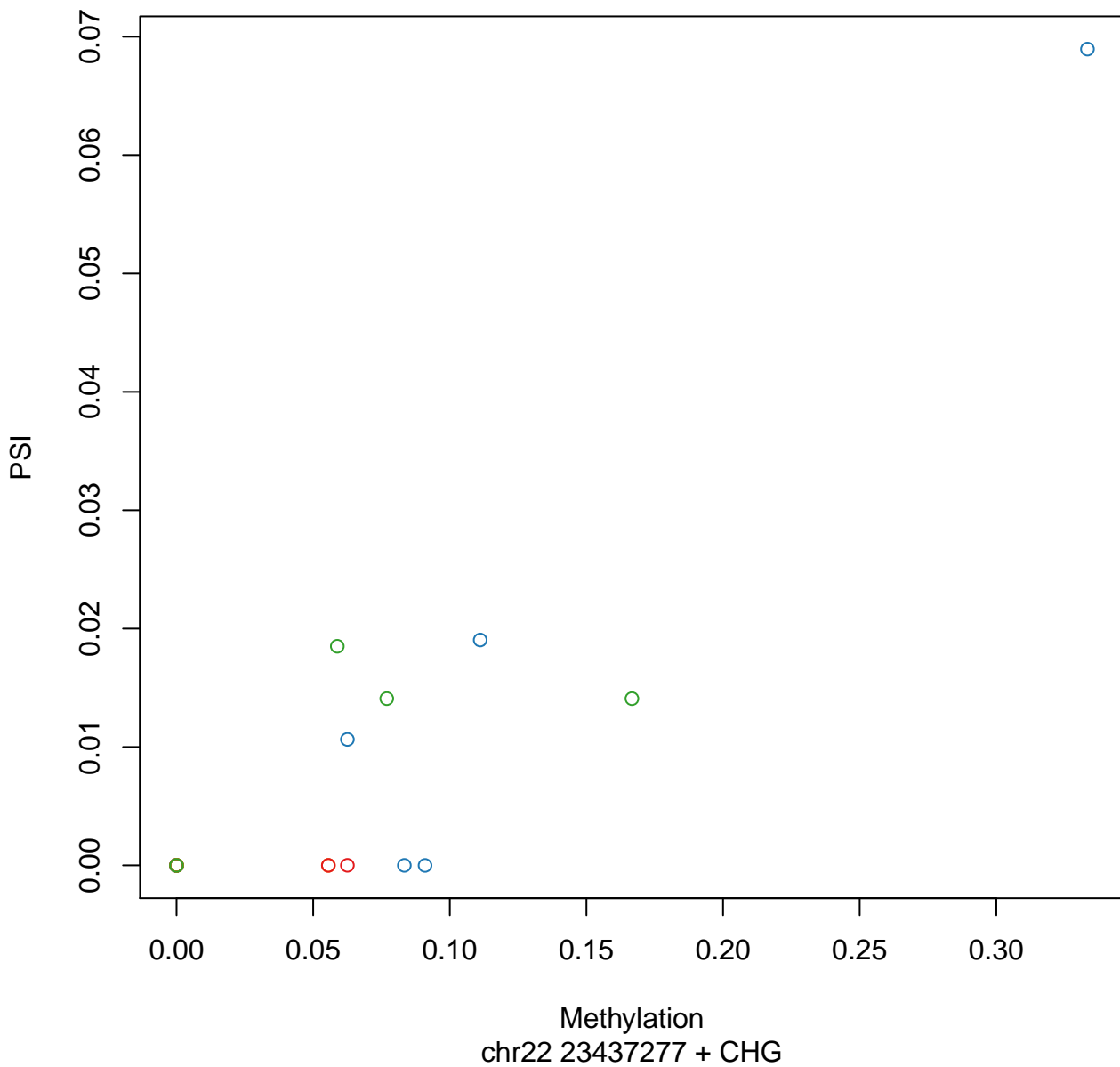
NUP93



psi ENSG00000128266.8_1 FDR 8.71e-05
GNAZ

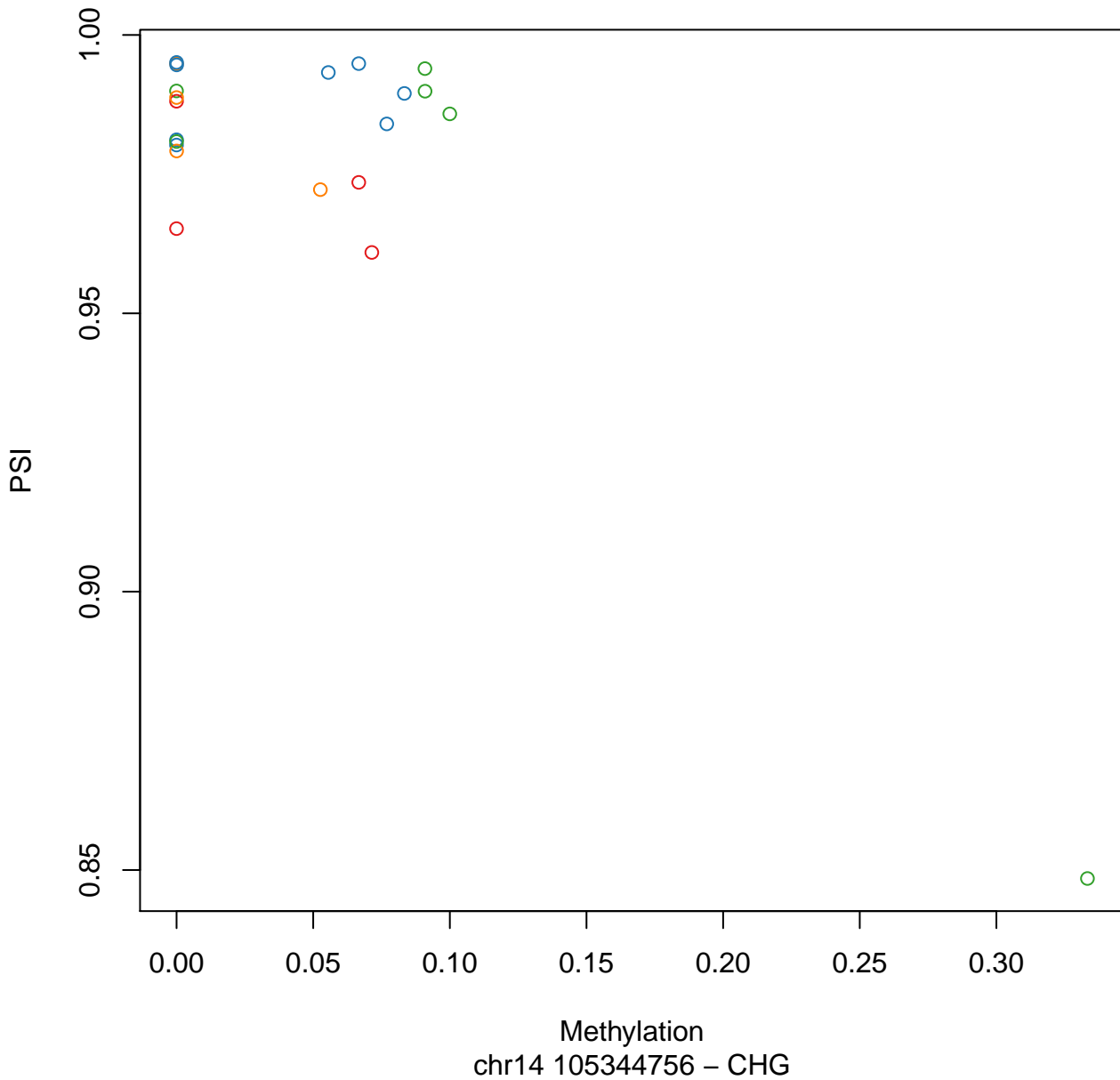


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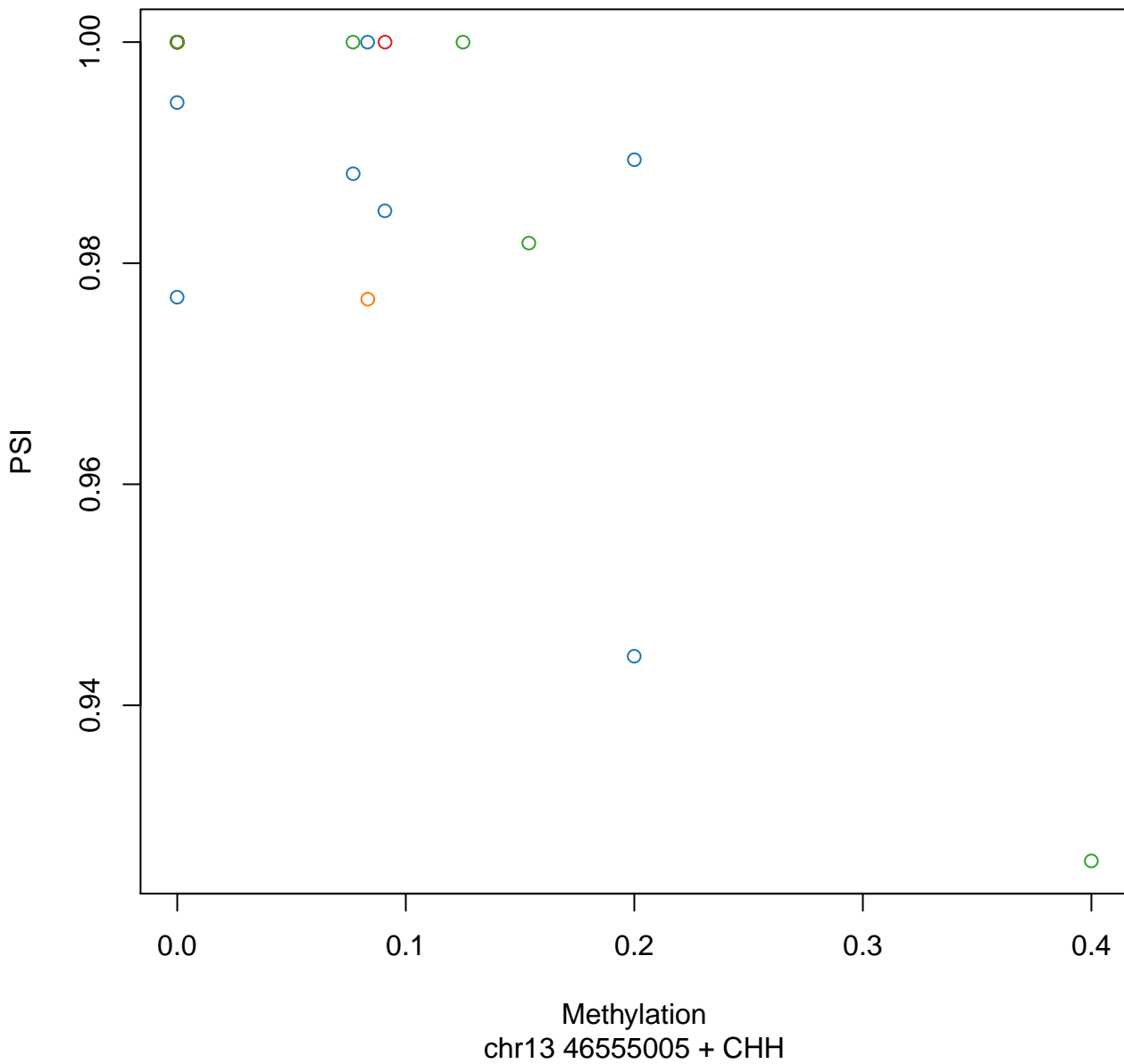


psi ENSG00000099814.15_1 FDR 0.00694

CEP170B

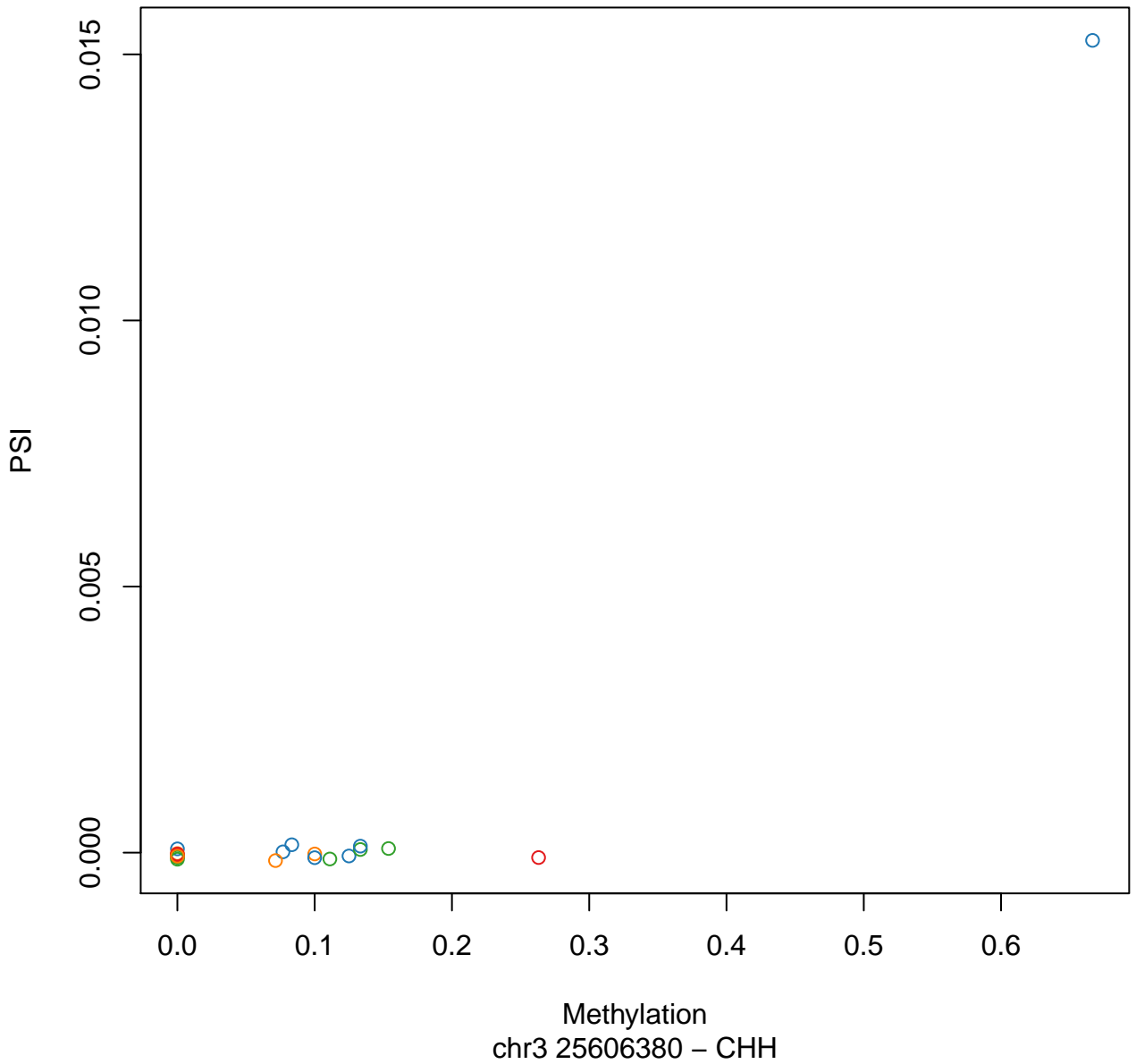


psi ENSG00000123200.16_2 FDR 0.0086
ZC3H13



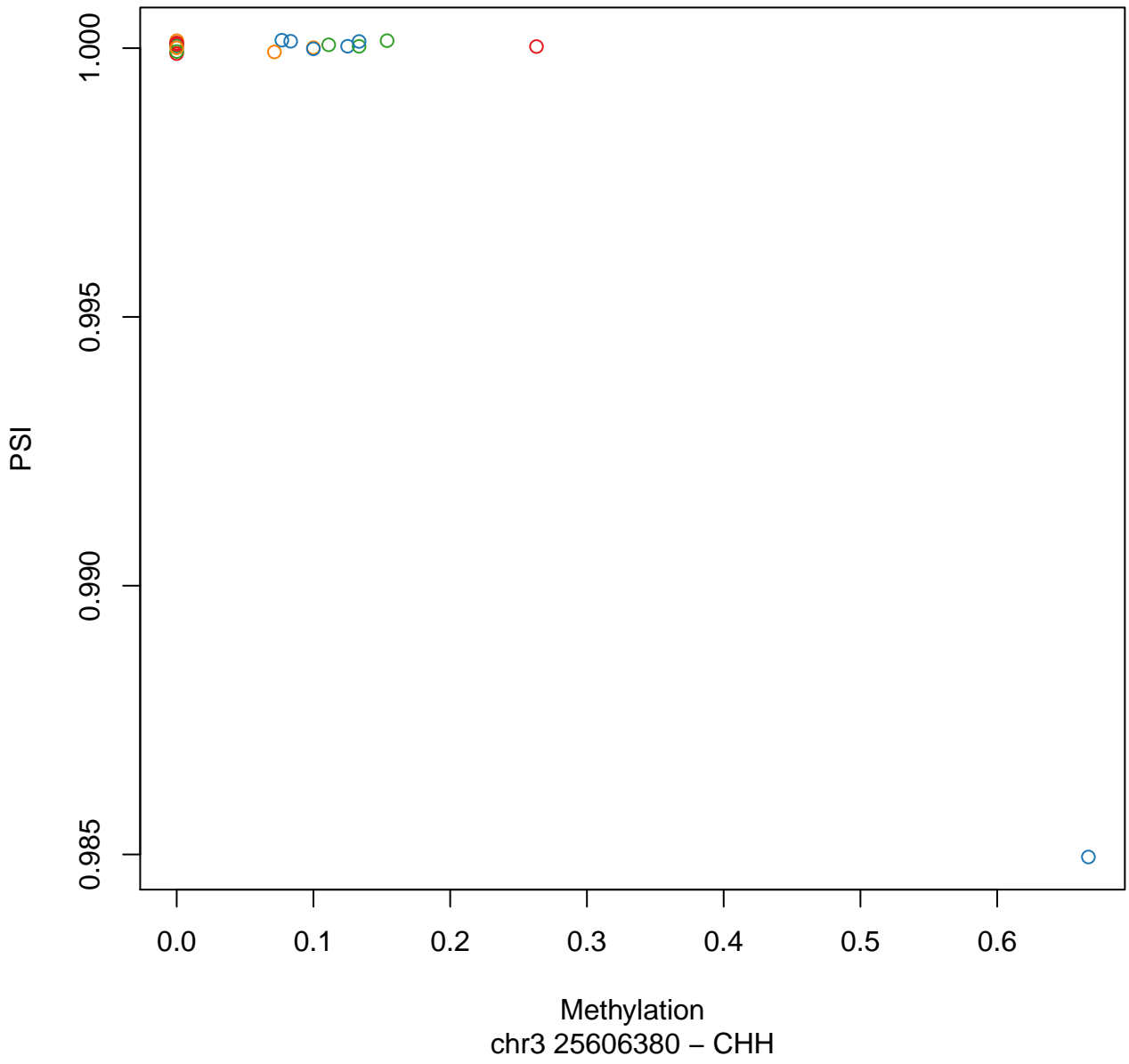
psi ENSG00000077092.18_2 FDR 0.000594

RARB

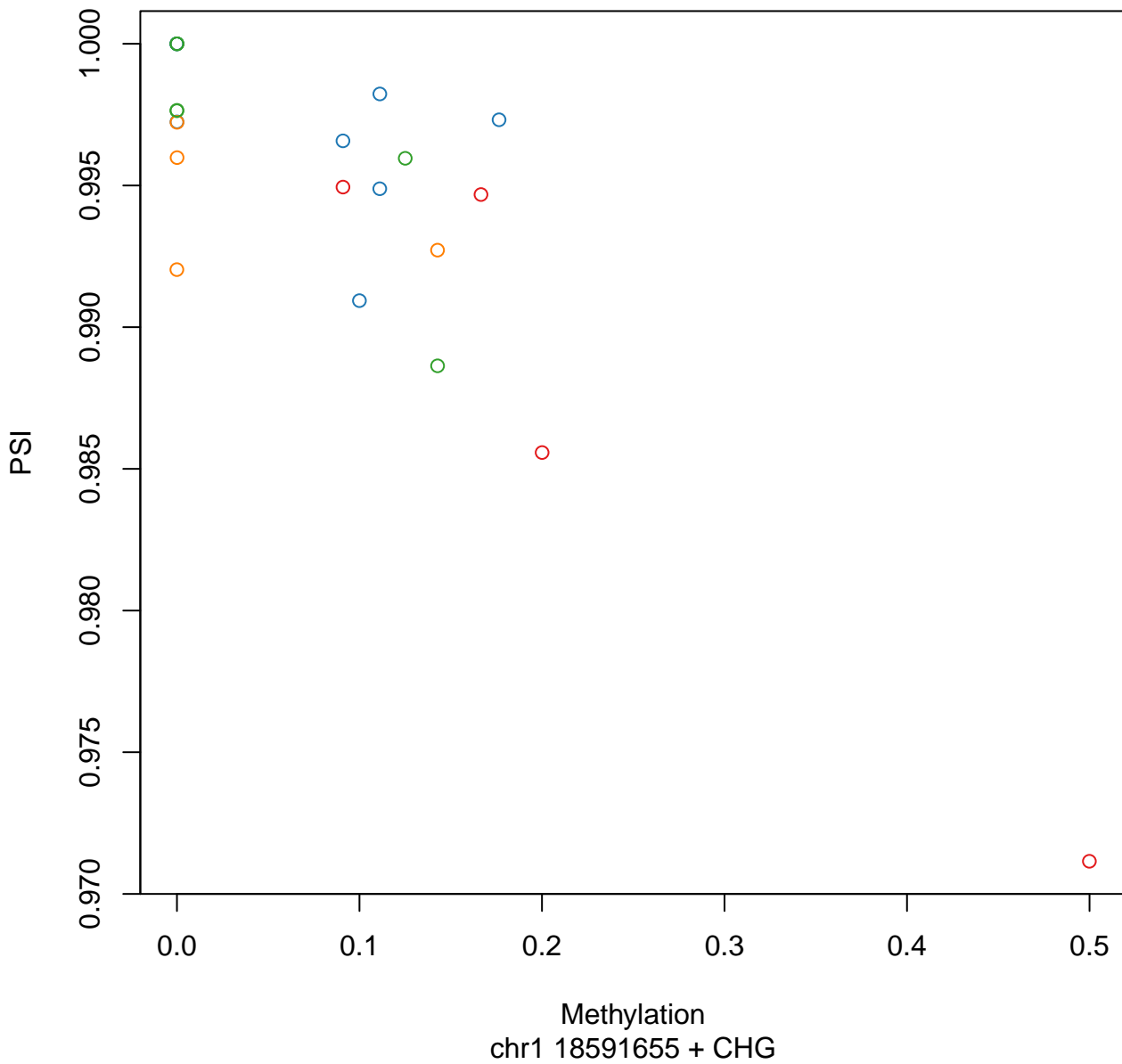


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RARB

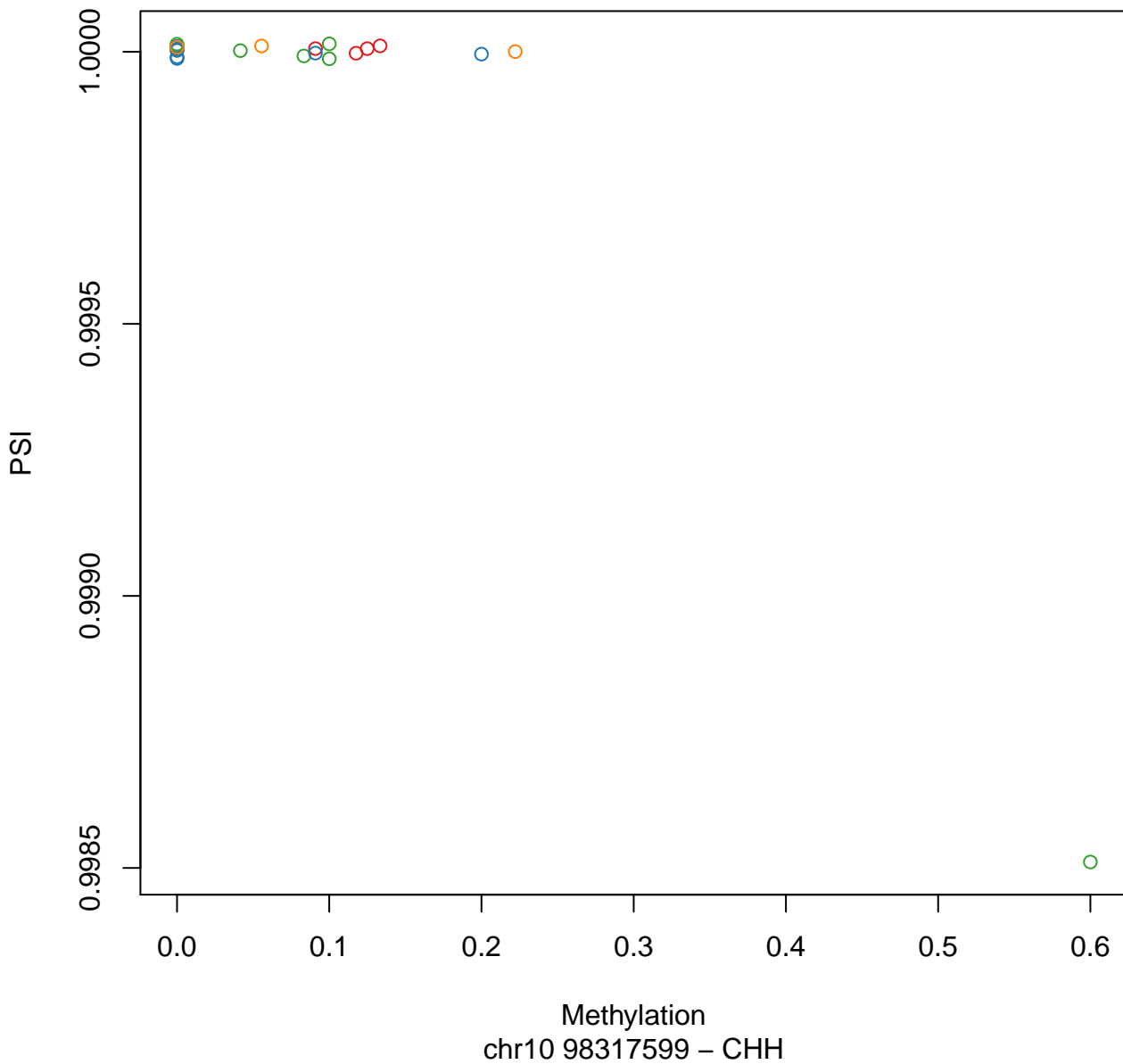


psi ENSG00000117154.11_2 FDR 0.000477
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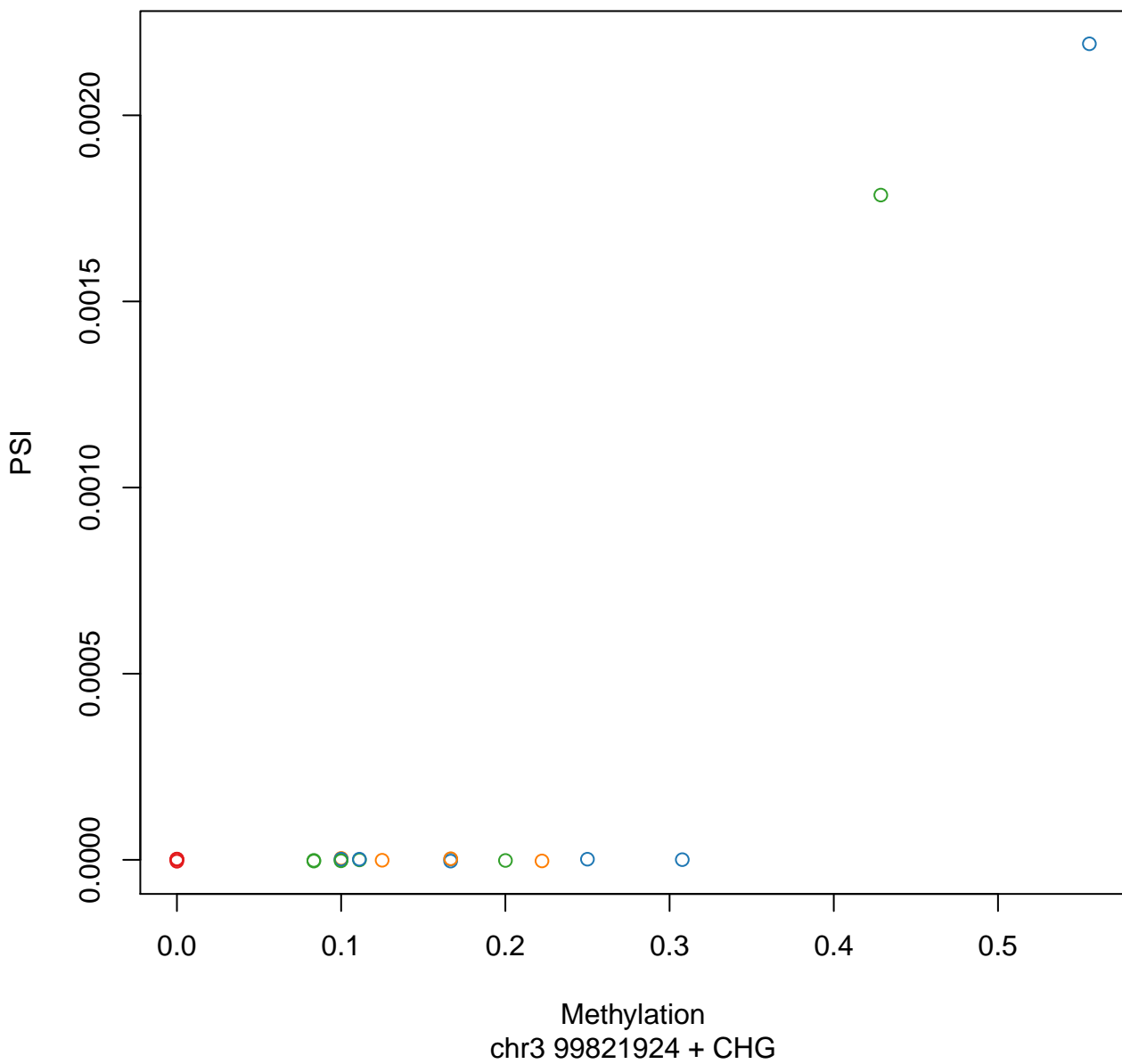


psi ENSG00000077147.14_1 FDR 0.00068

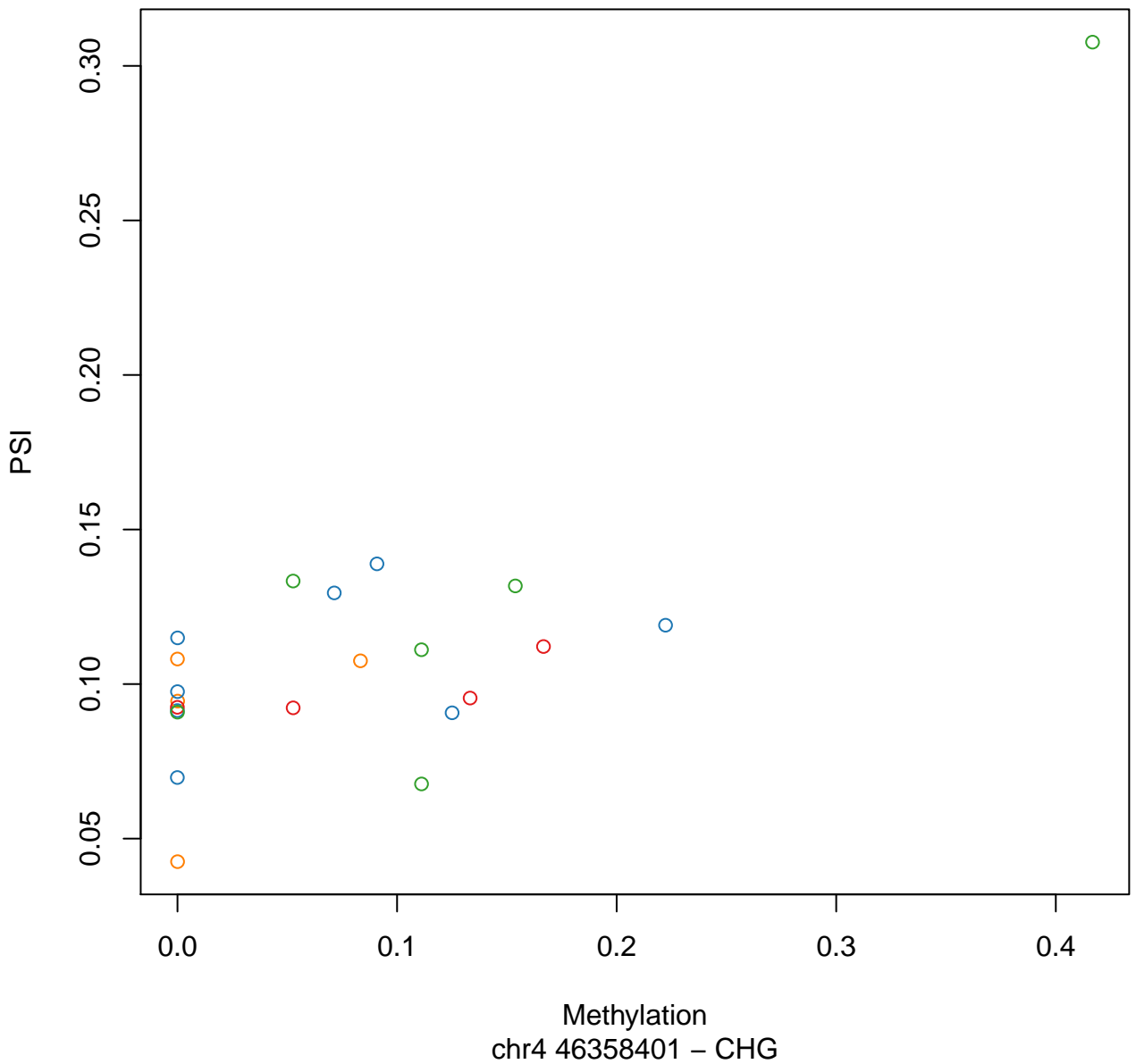
TM9SF3



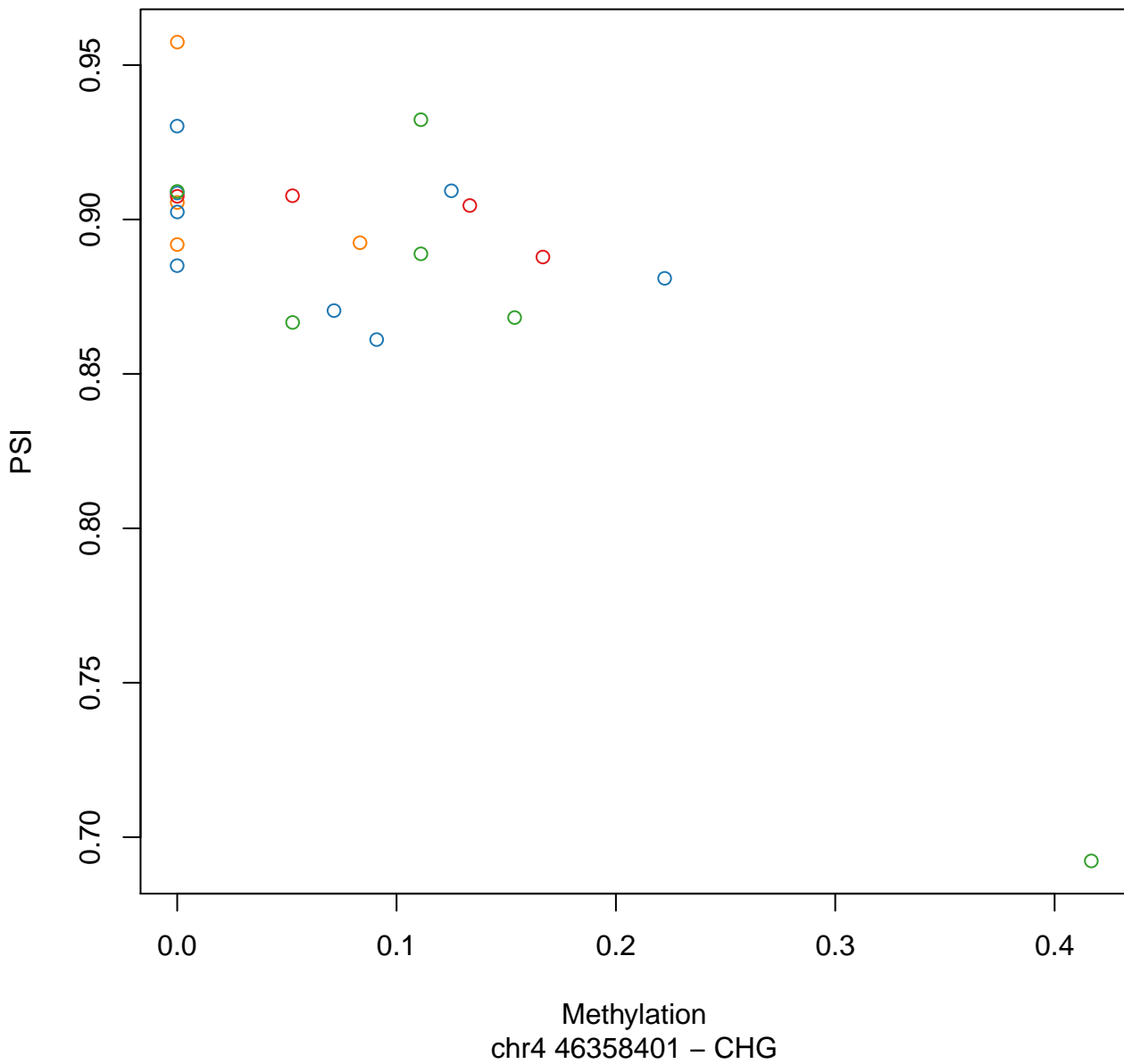
psi ENSG00000184220.11_2 FDR 0.0075
CMSS1



psi ENSG00000151834.15_1 FDR 0.019
GABRA2

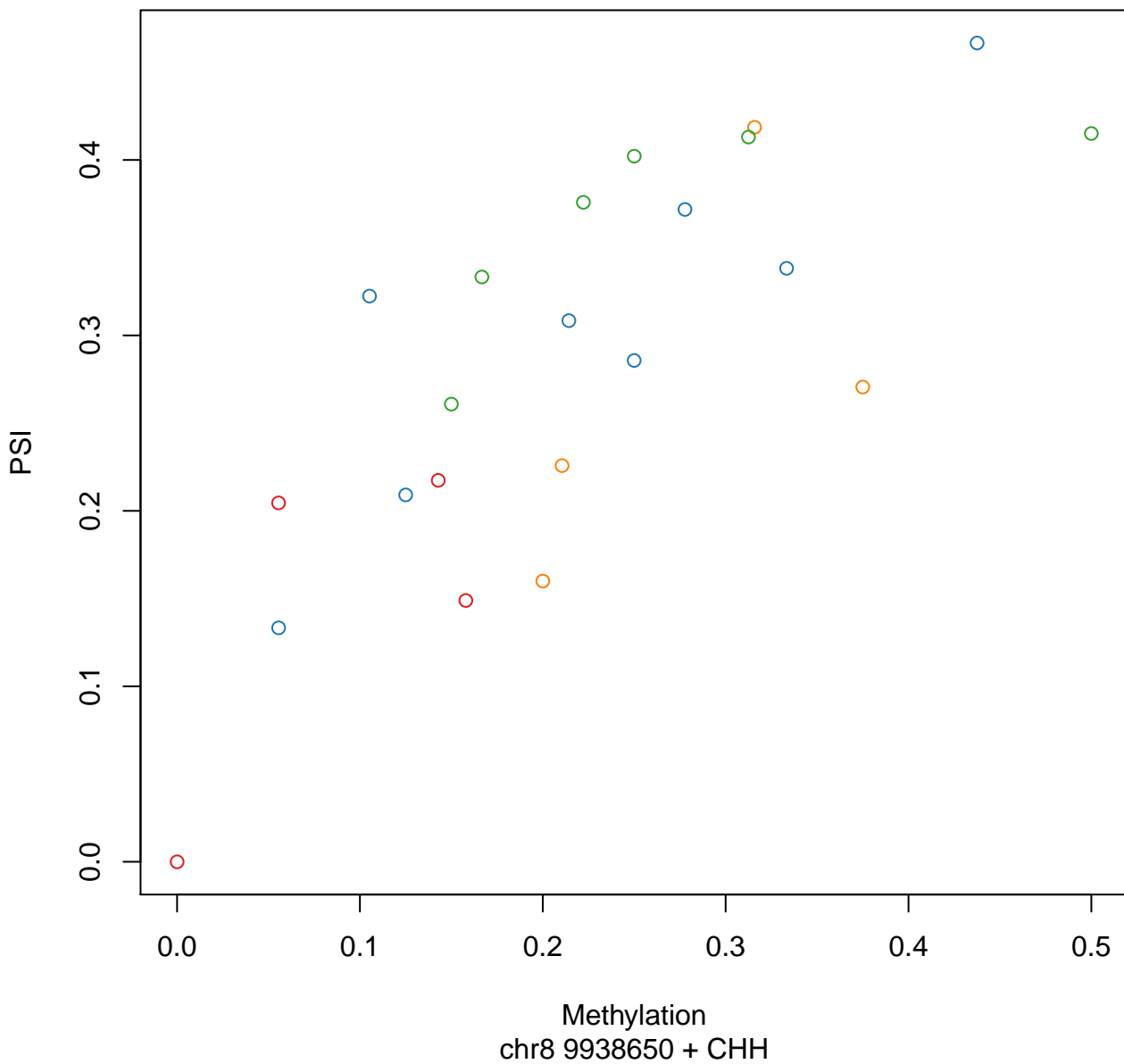


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GABRA2



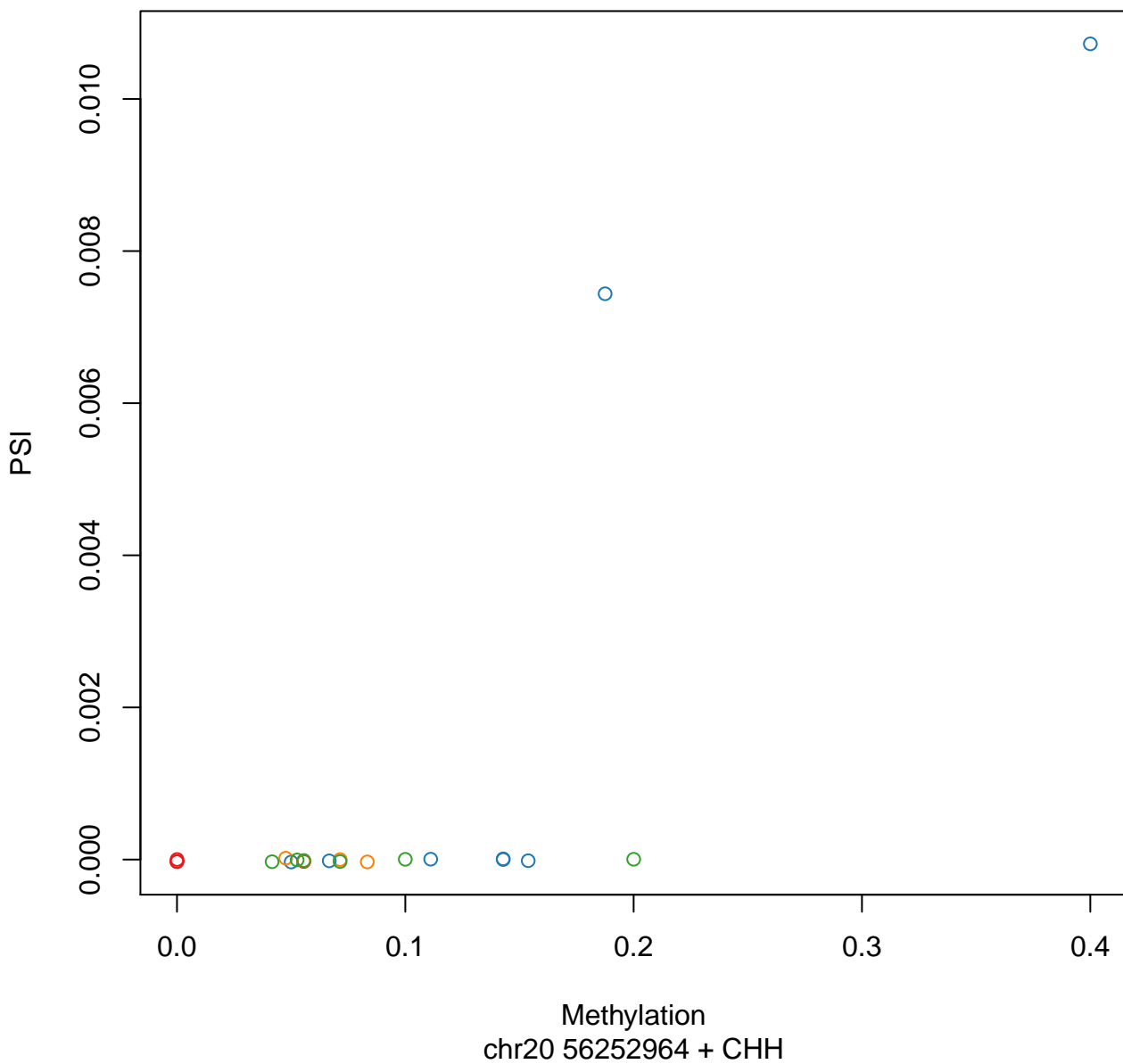
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MSRA



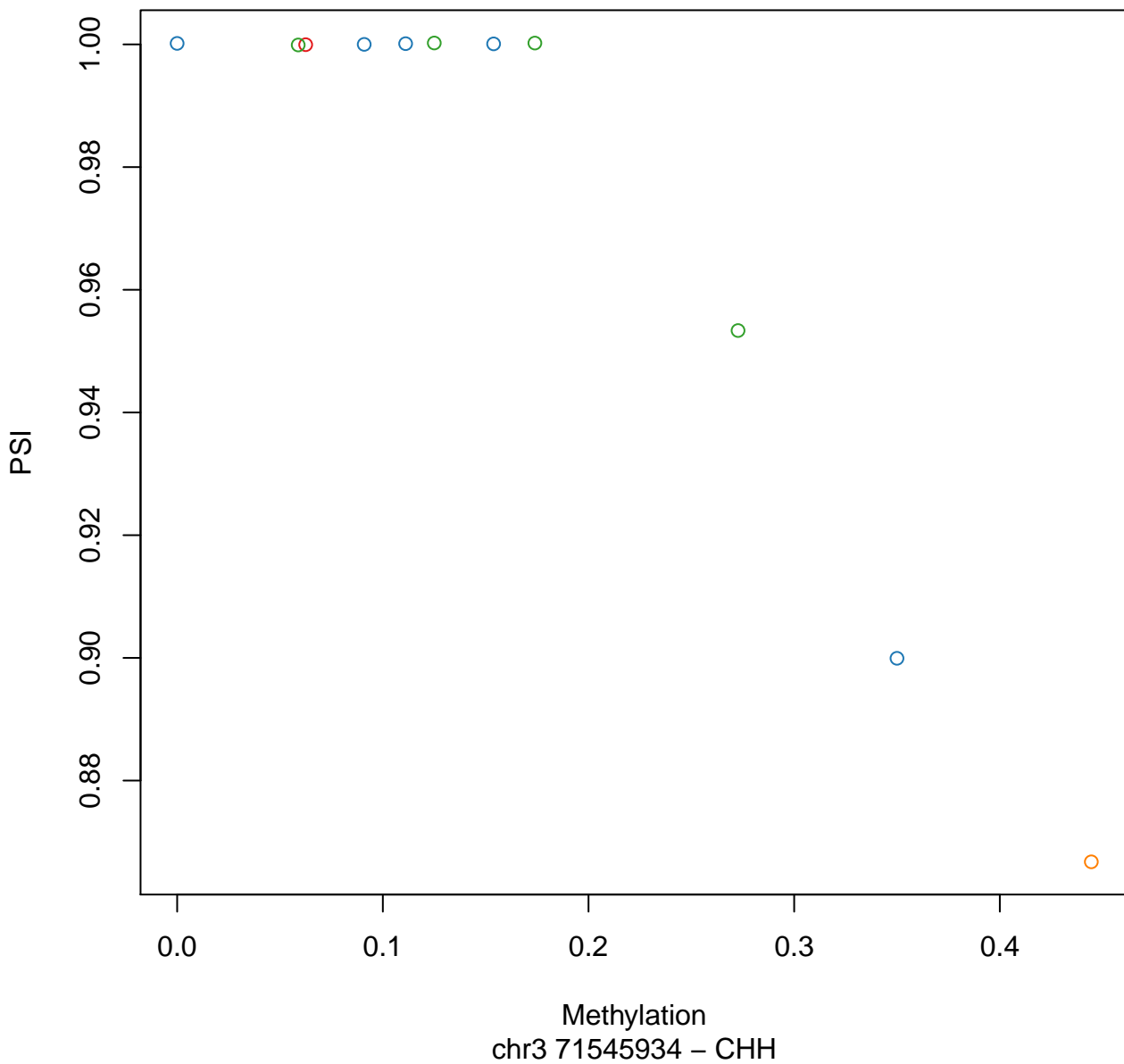
psi ENSG00000124225.15_2 FDR 0.0158

PMEPA1



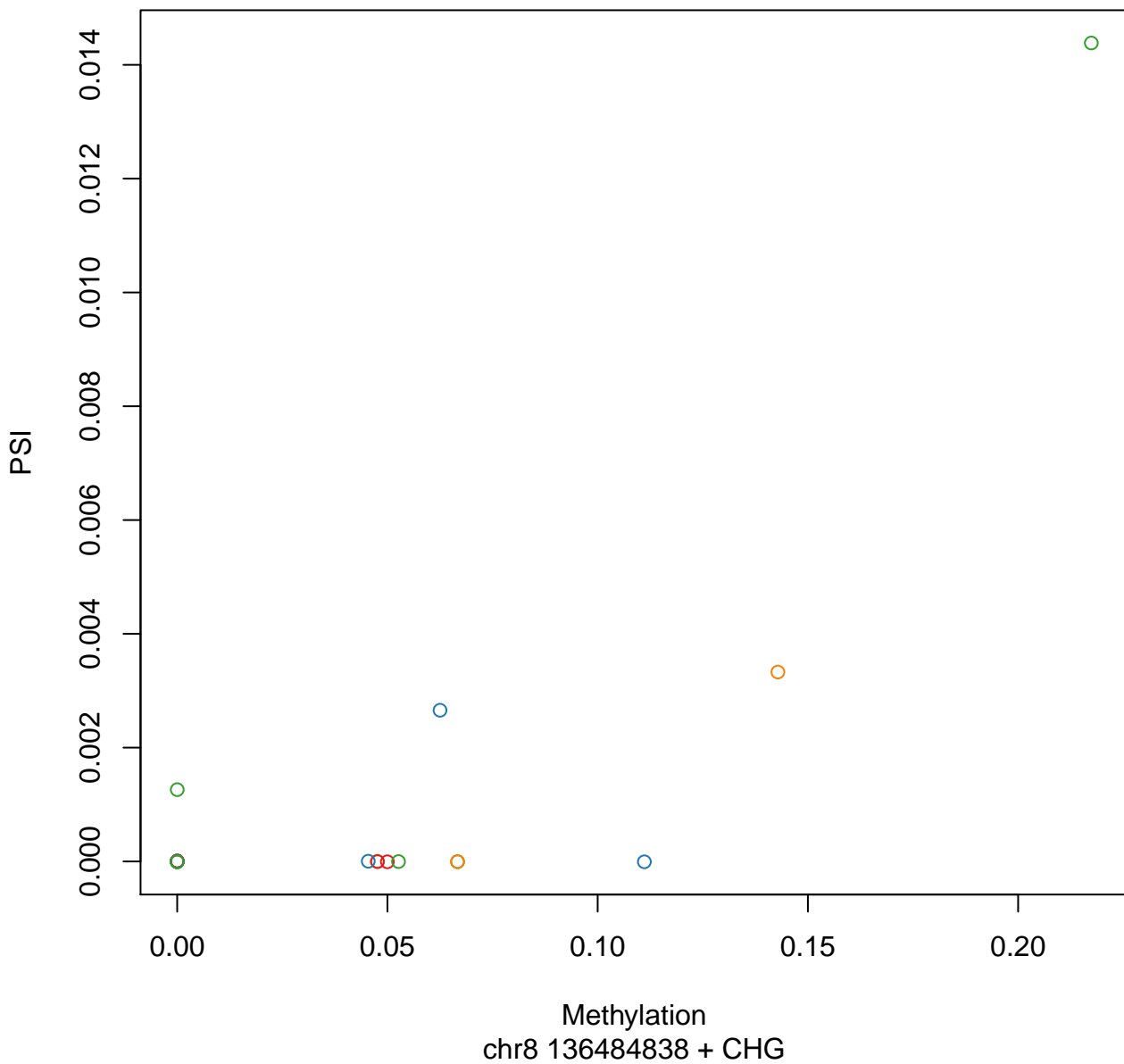
psi ENSG00000114861.18_2 FDR 0.0177

FOXP1



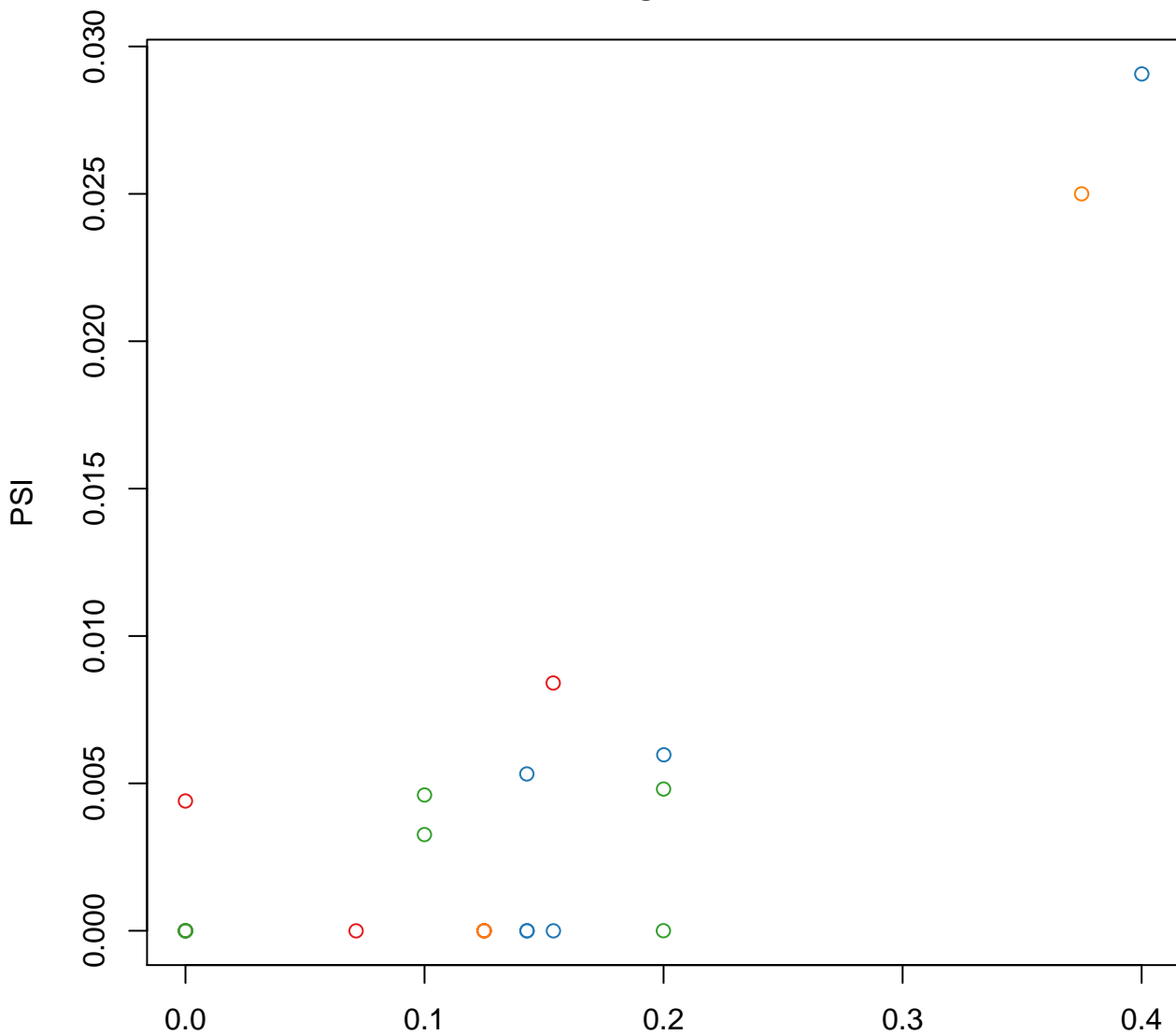
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KHDRBS3



psi ENSG00000154978.12_1 FDR 0.00942

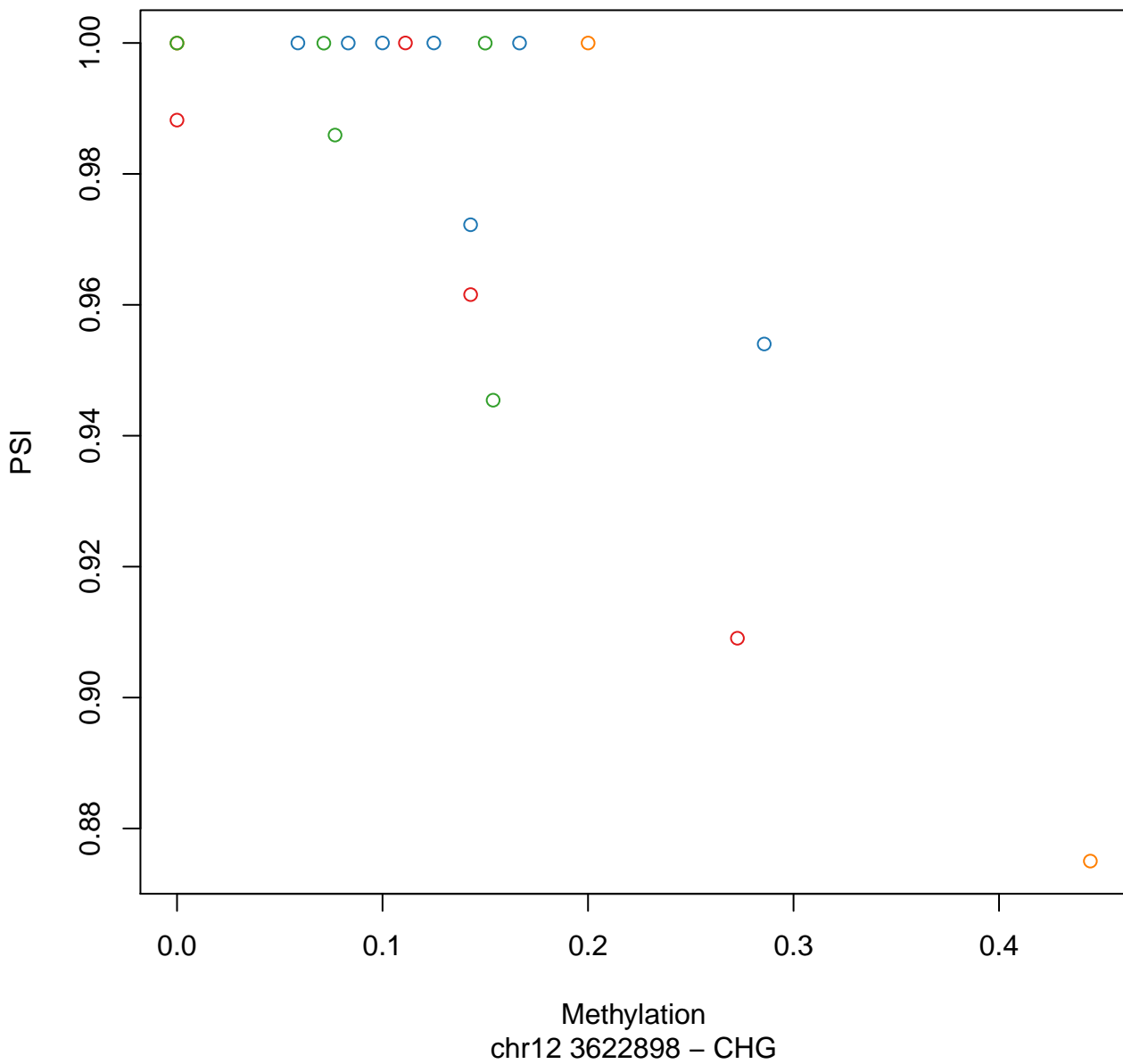
VOPP1



Methylation
chr7 55520841 - CHG

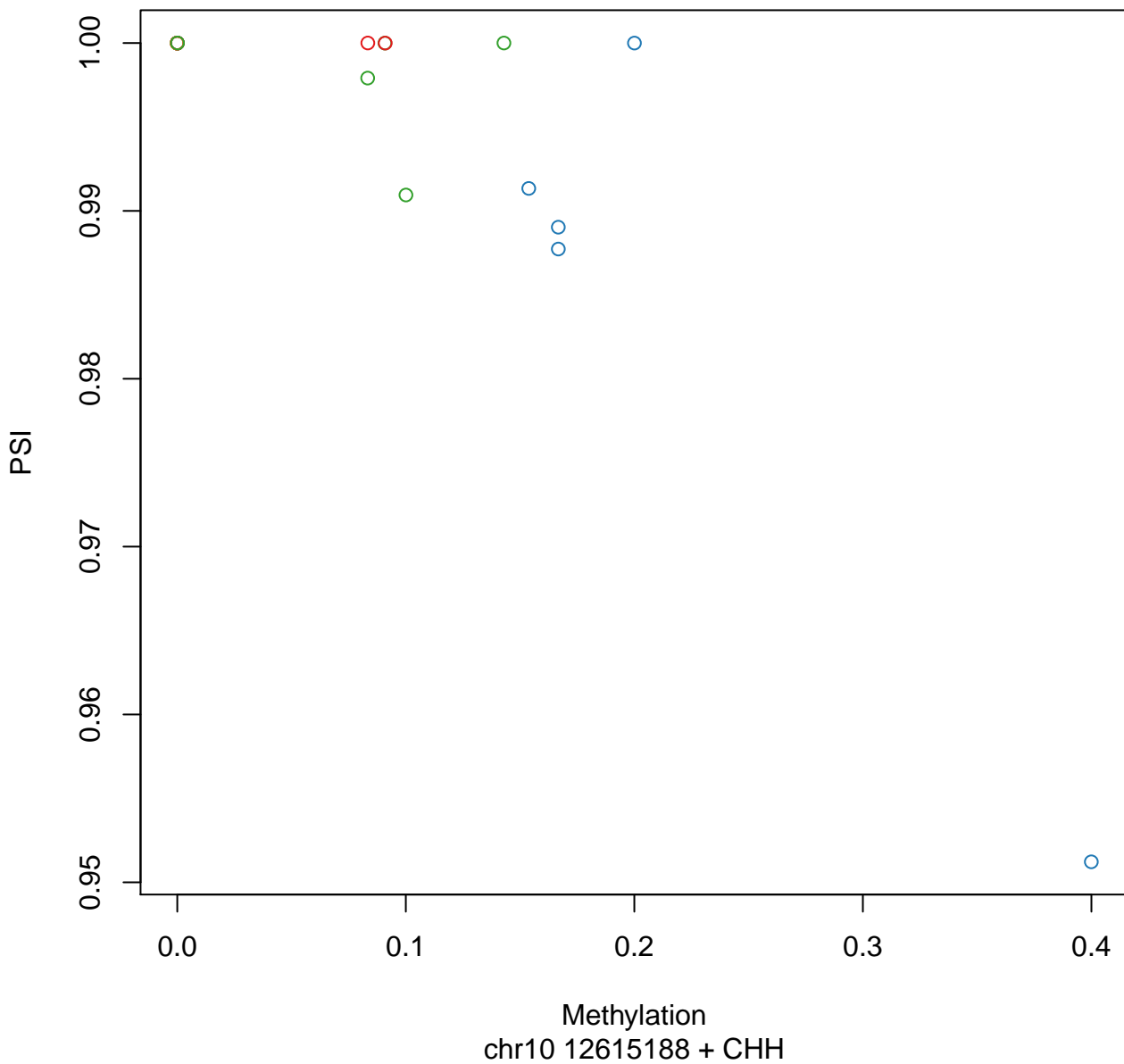
psi ENSG00000111218.11_2 FDR 0.0201

PRMT8



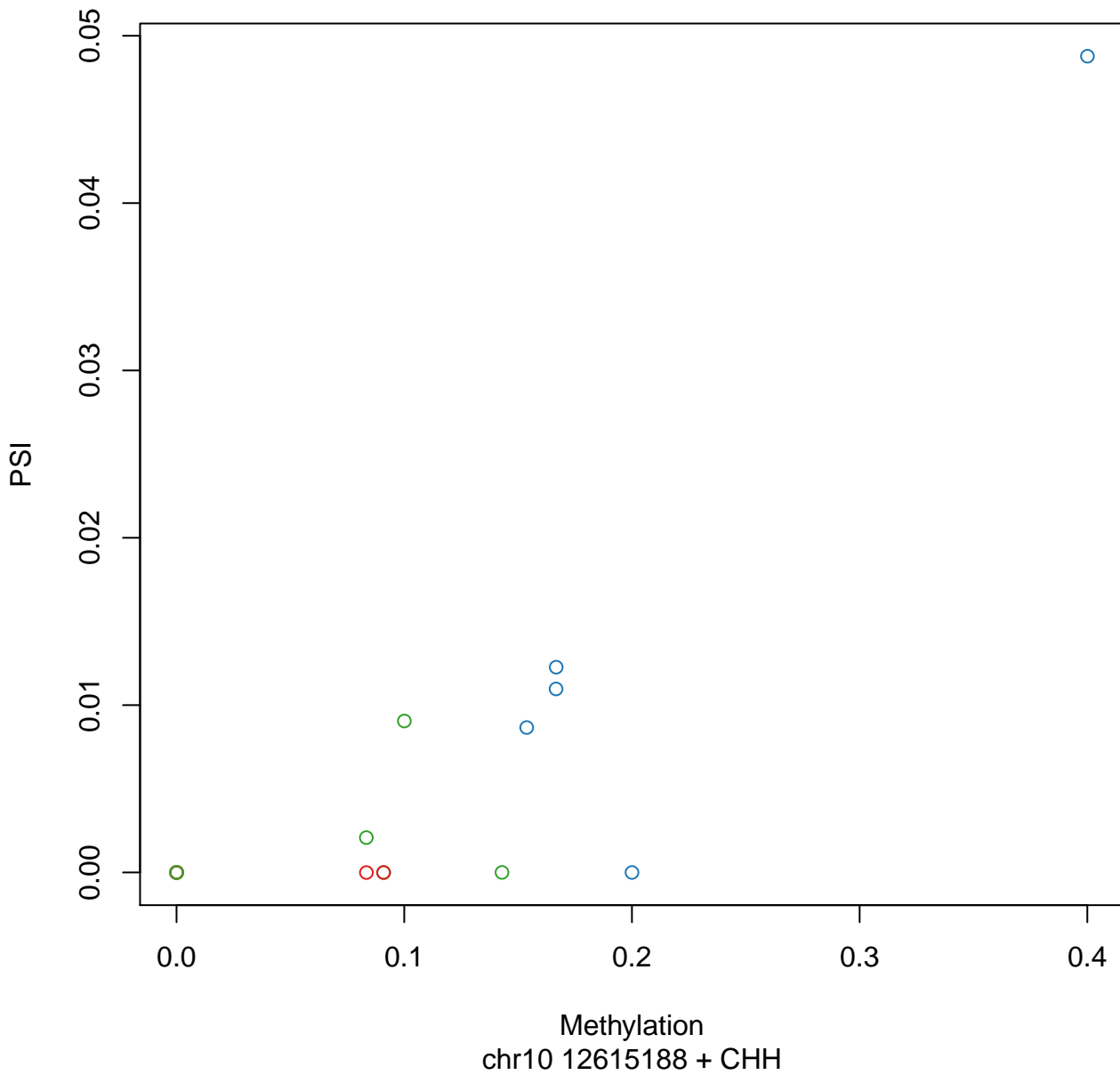
psi ENSG00000183049.12_1 FDR 0.00425

CAMK1D



psi ENSG00000183049.12_1 FDR 0.00425

CAMK1D



psi ENSG00000143499.13_1 FDR 0.0104

SMYD2

