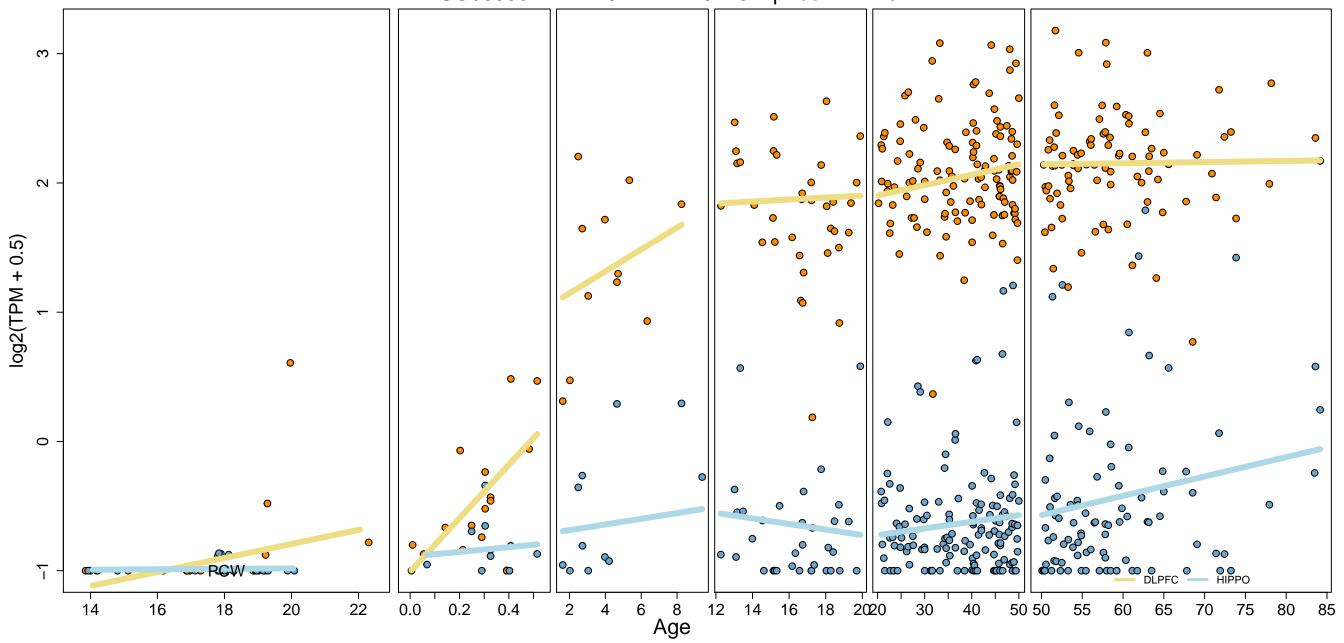
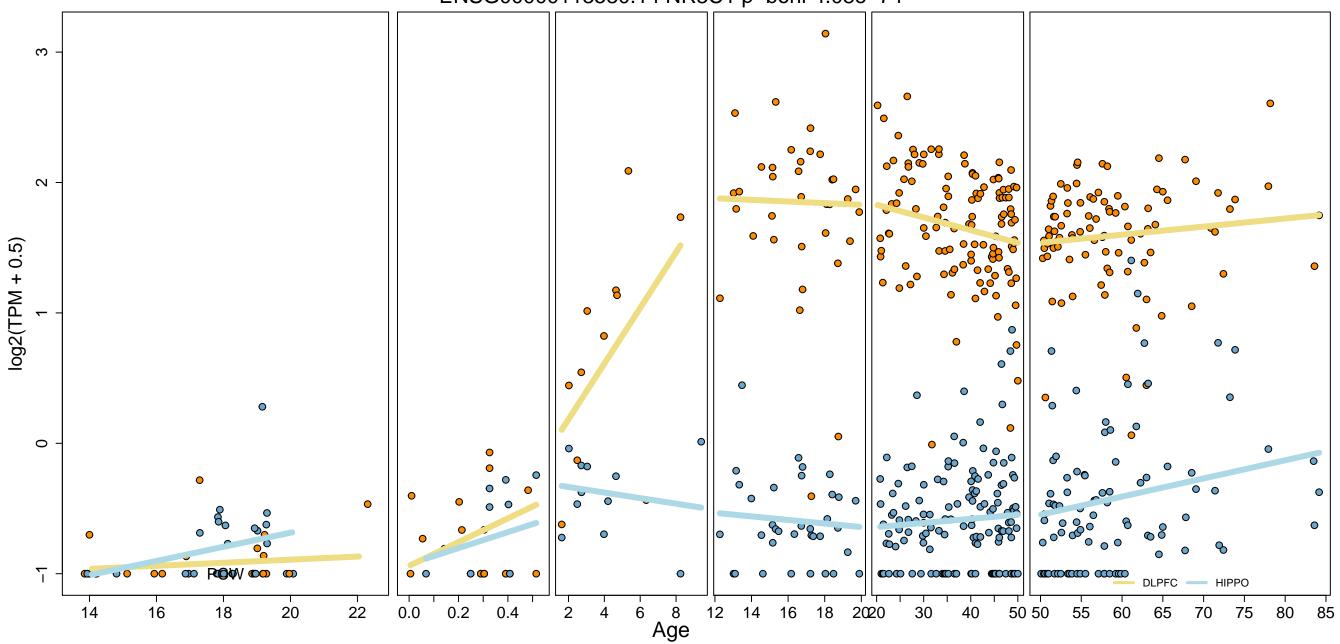
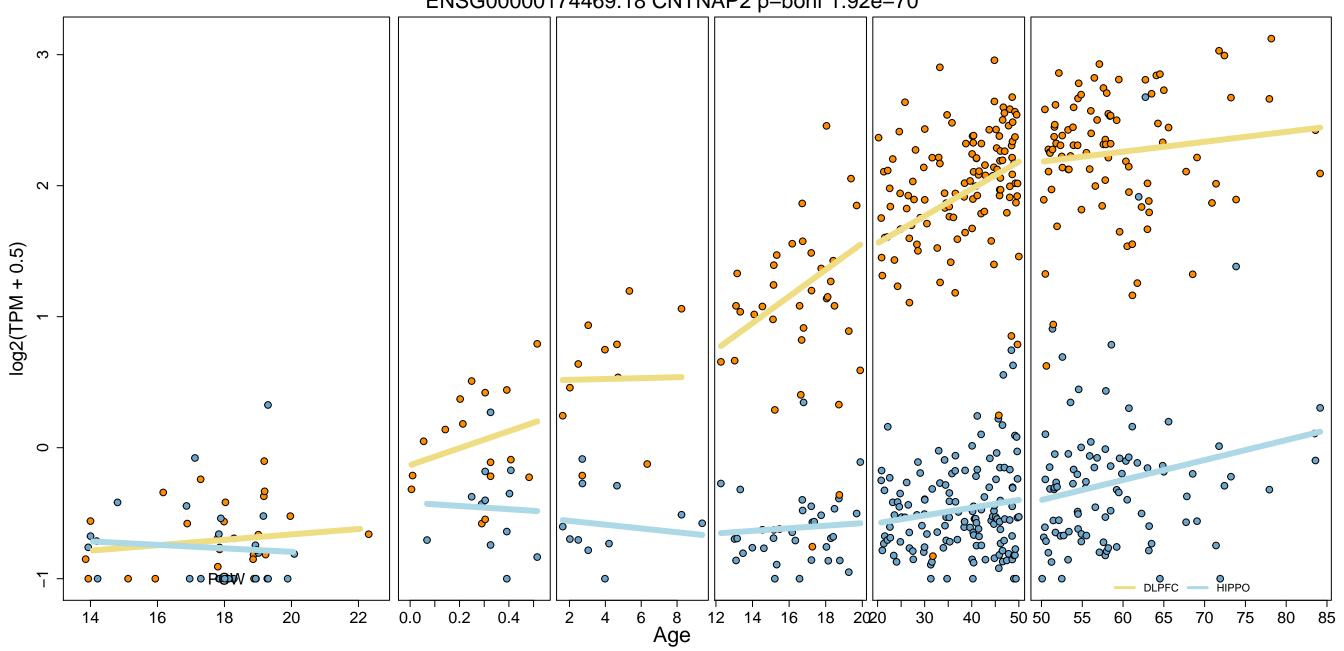
ENSG00000224271.5 RP11-191L9.4 p-bonf 1.47e-74



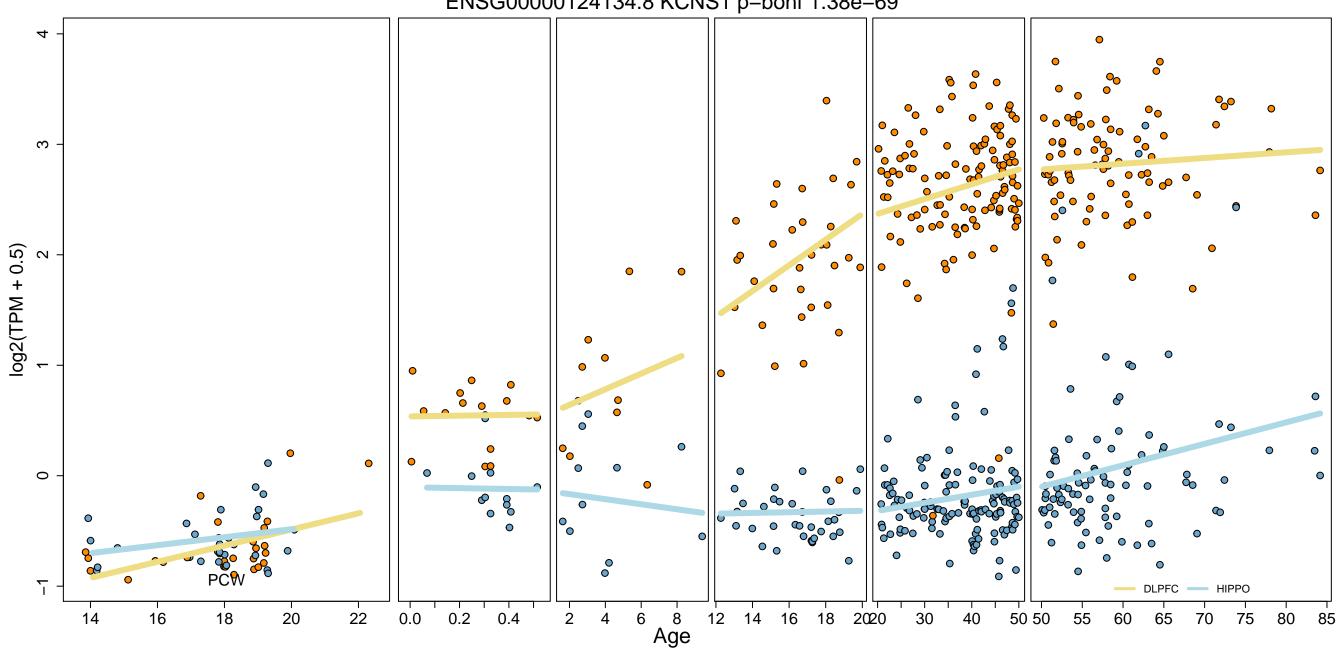
ENSG00000113580.14 NR3C1 p-bonf 4.08e-74



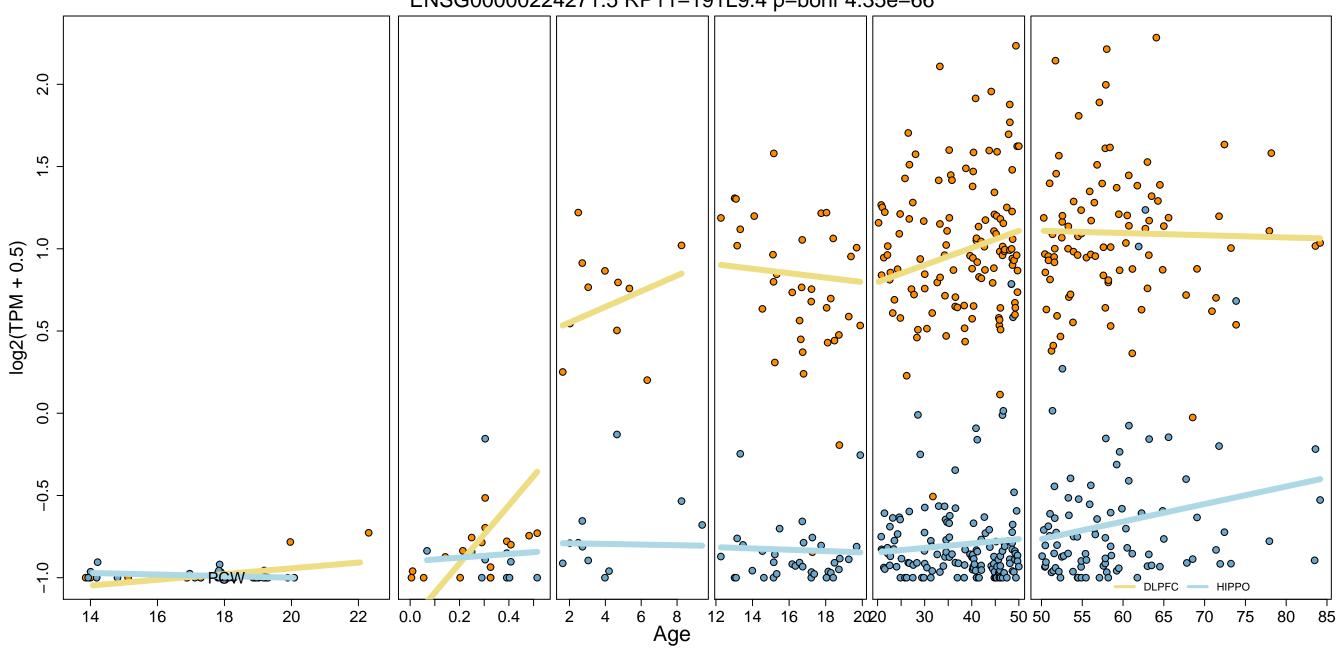
ENSG00000174469.18 CNTNAP2 p-bonf 1.92e-70



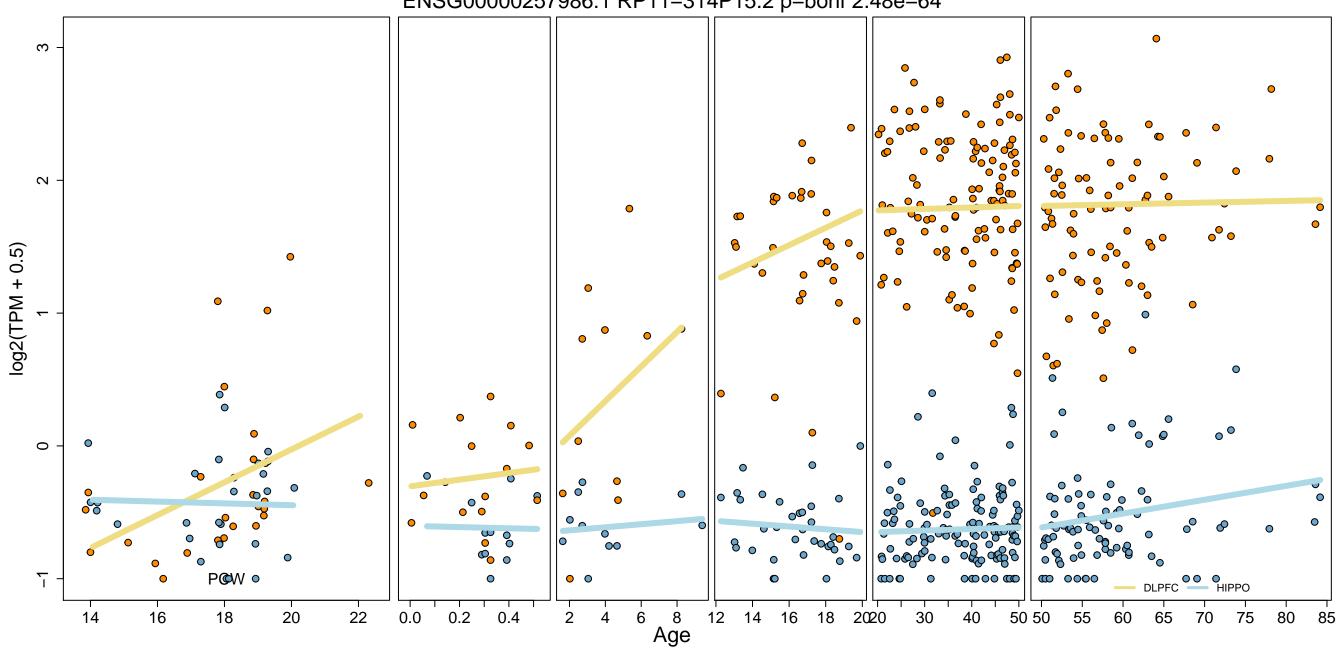
ENSG00000124134.8 KCNS1 p-bonf 1.38e-69



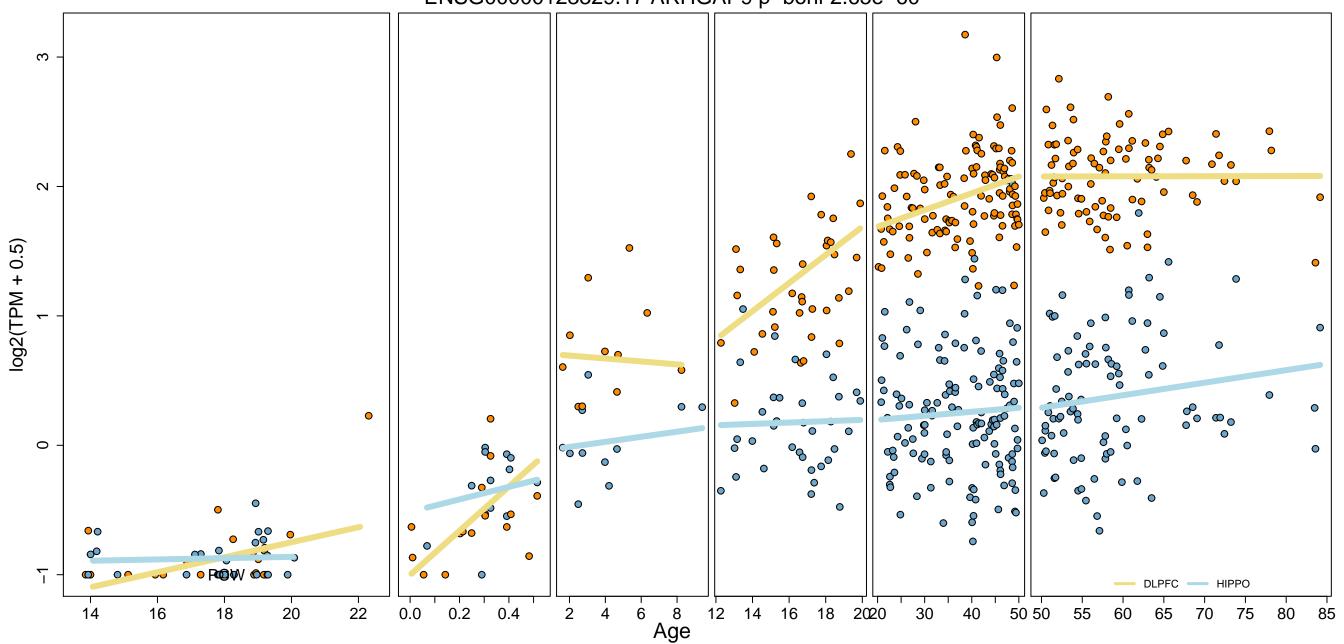
ENSG00000224271.5 RP11-191L9.4 p-bonf 4.35e-66



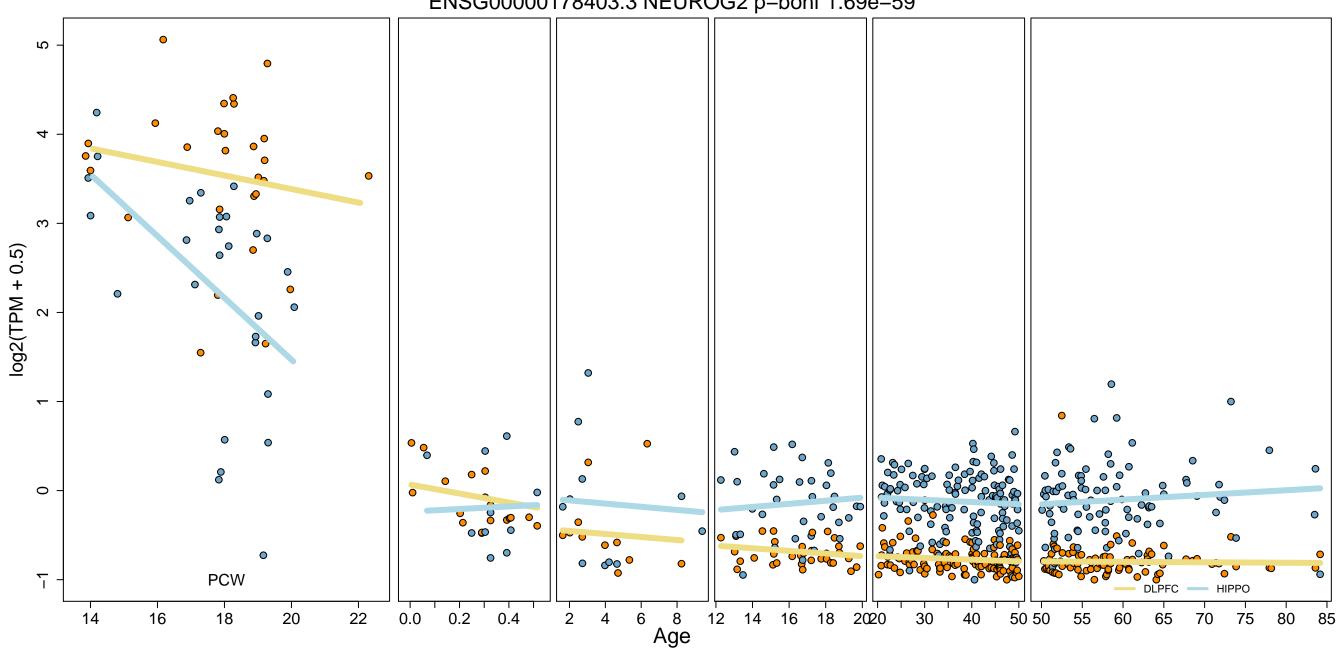
ENSG00000257986.1 RP11-314P15.2 p-bonf 2.48e-64



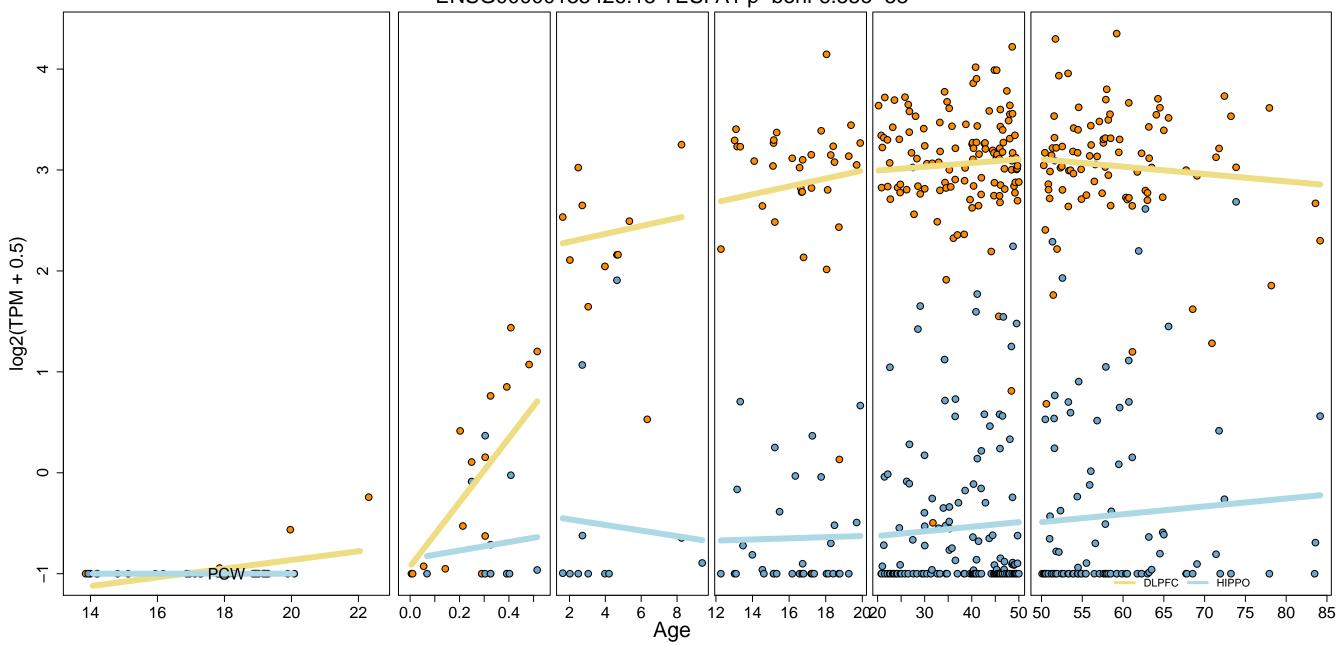
ENSG00000123329.17 ARHGAP9 p-bonf 2.65e-60



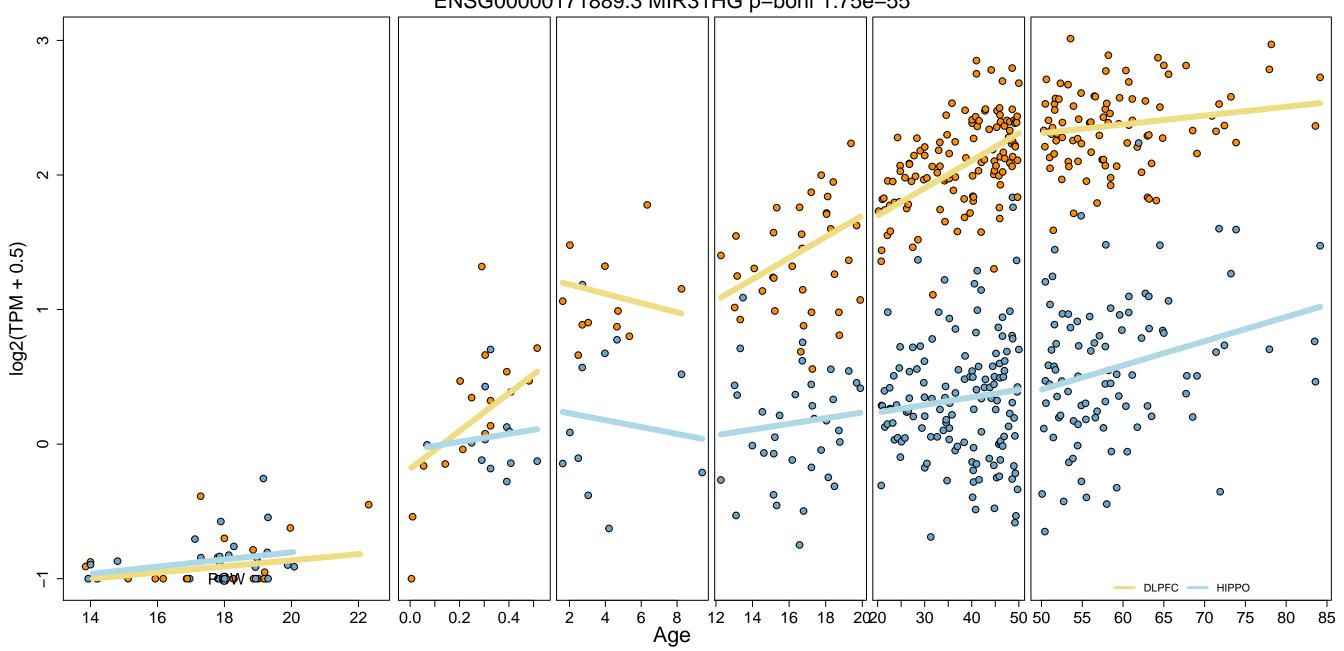
ENSG00000178403.3 NEUROG2 p-bonf 1.69e-59



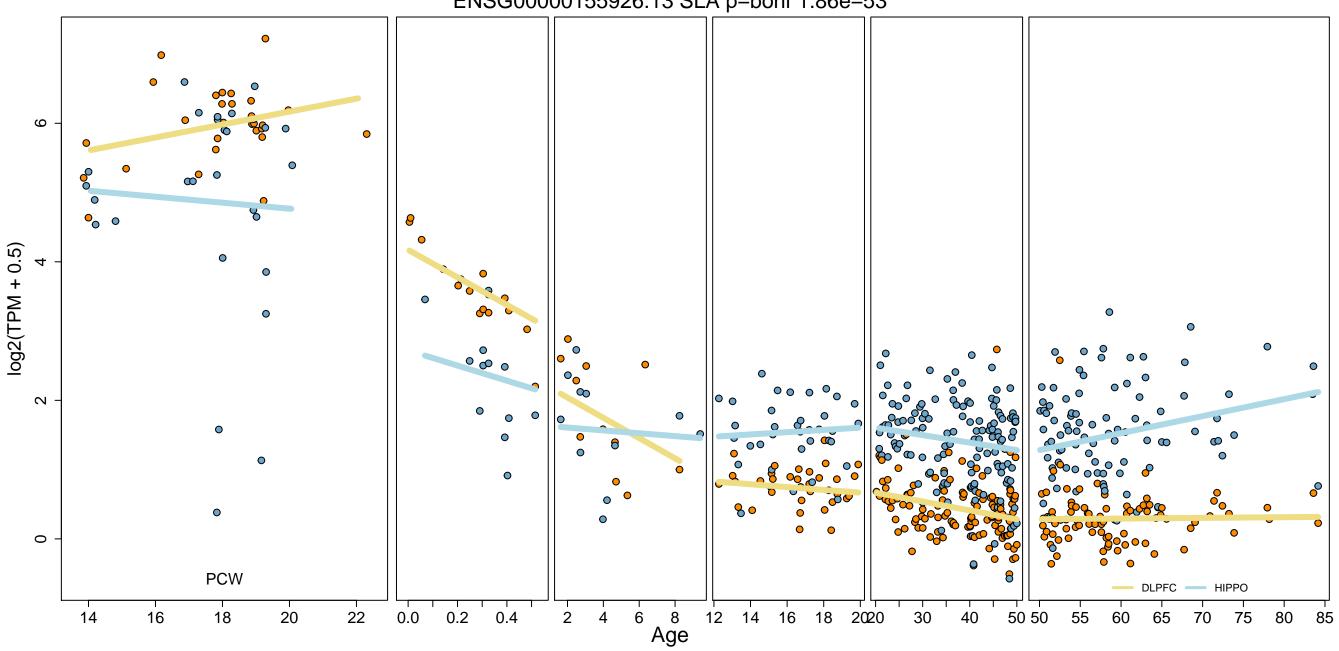
ENSG00000135426.15 TESPA1 p-bonf 6.55e-58



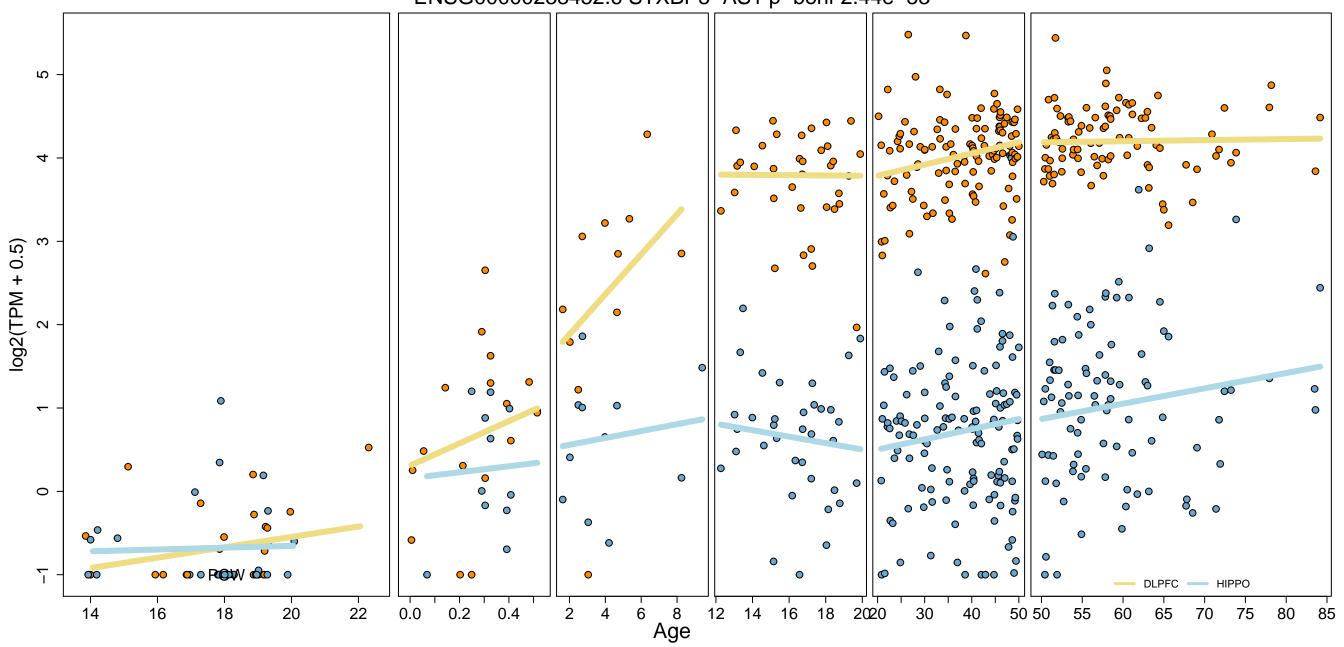
ENSG00000171889.3 MIR31HG p-bonf 1.75e-55



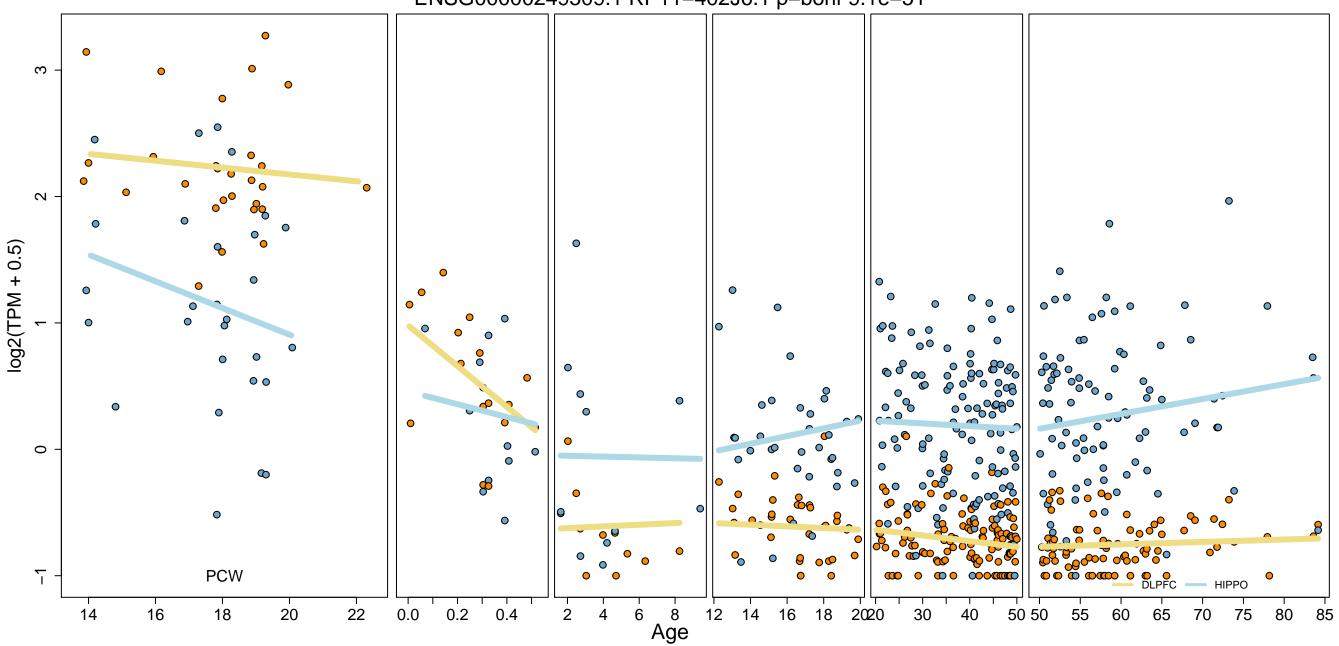
ENSG00000155926.13 SLA p-bonf 1.86e-53



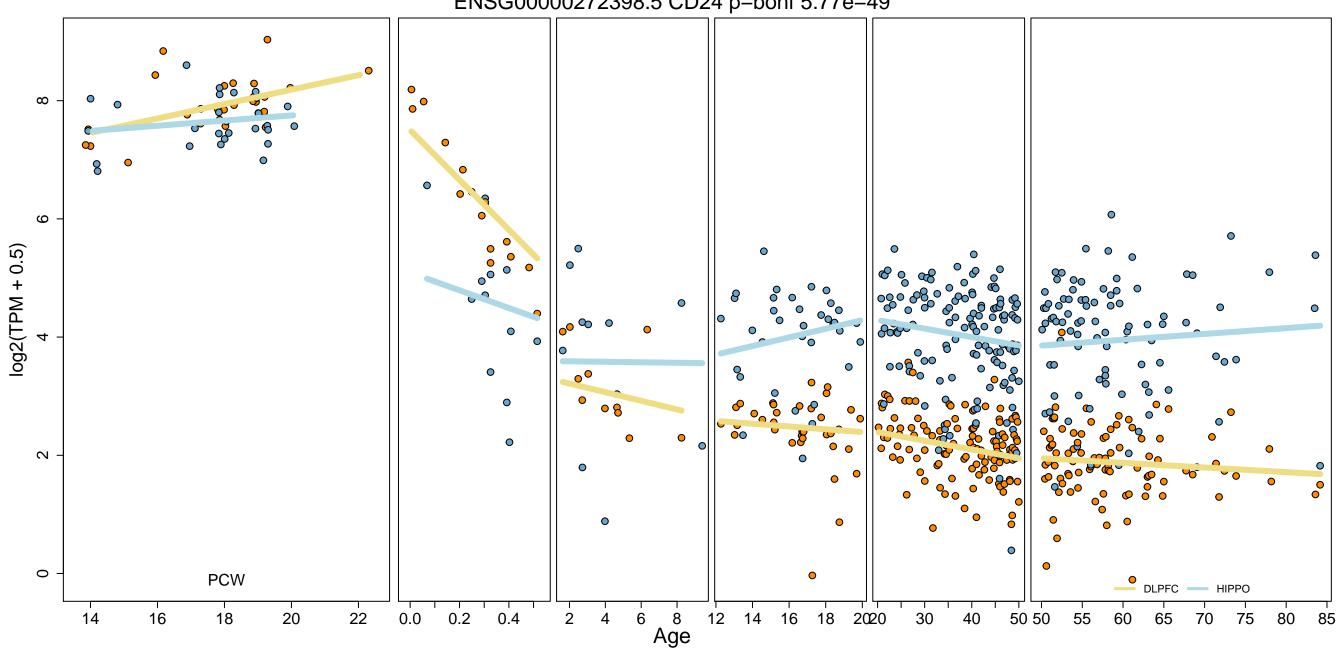
ENSG00000233452.6 STXBP5-AS1 p-bonf 2.44e-53



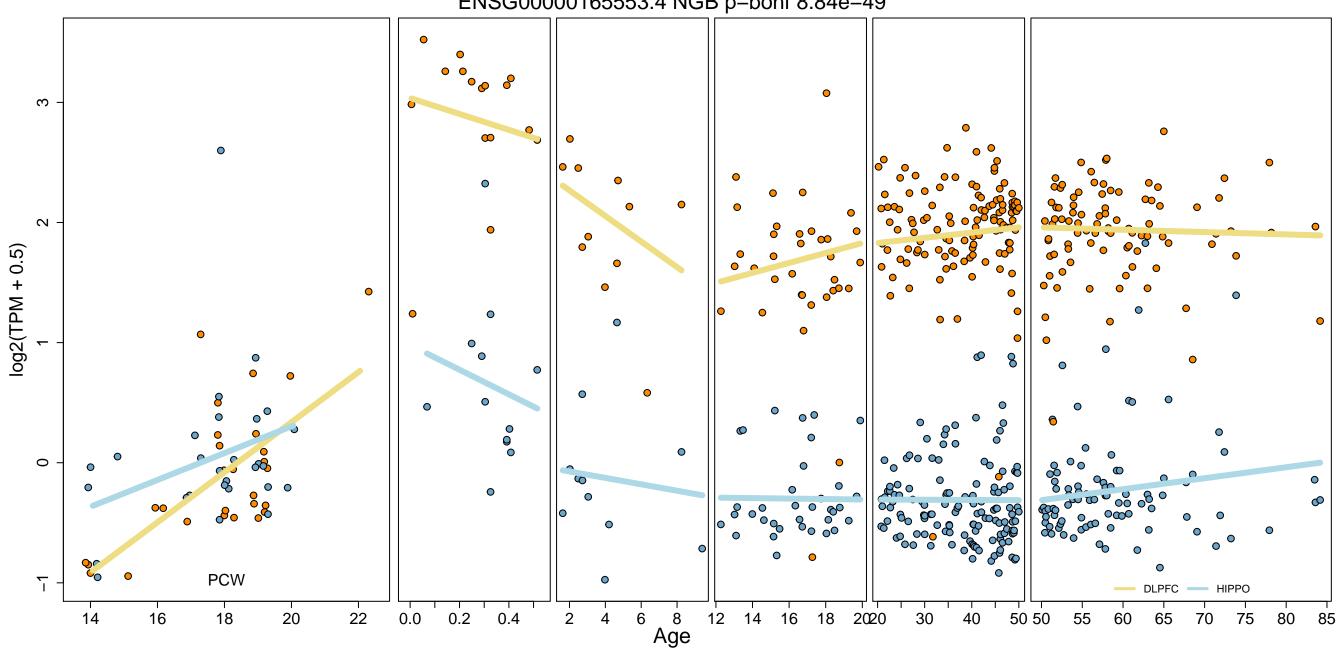
ENSG00000249509.1 RP11-402J6.1 p-bonf 9.1e-51



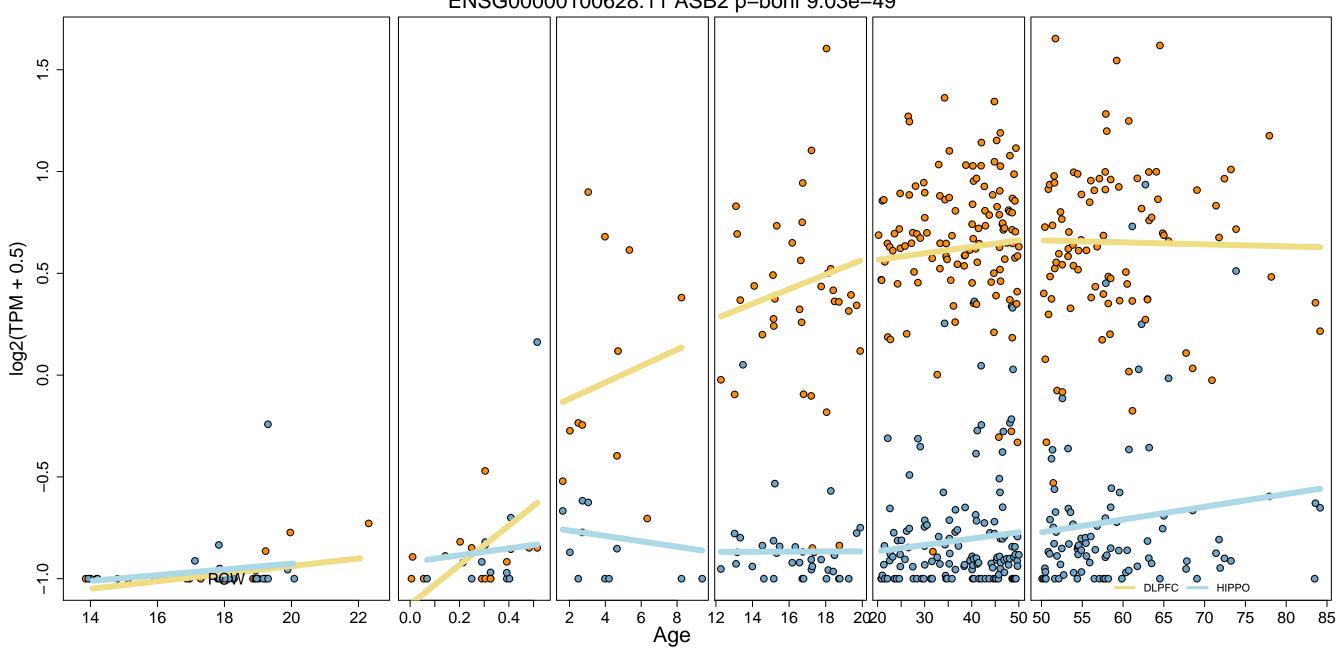
ENSG00000272398.5 CD24 p-bonf 5.77e-49



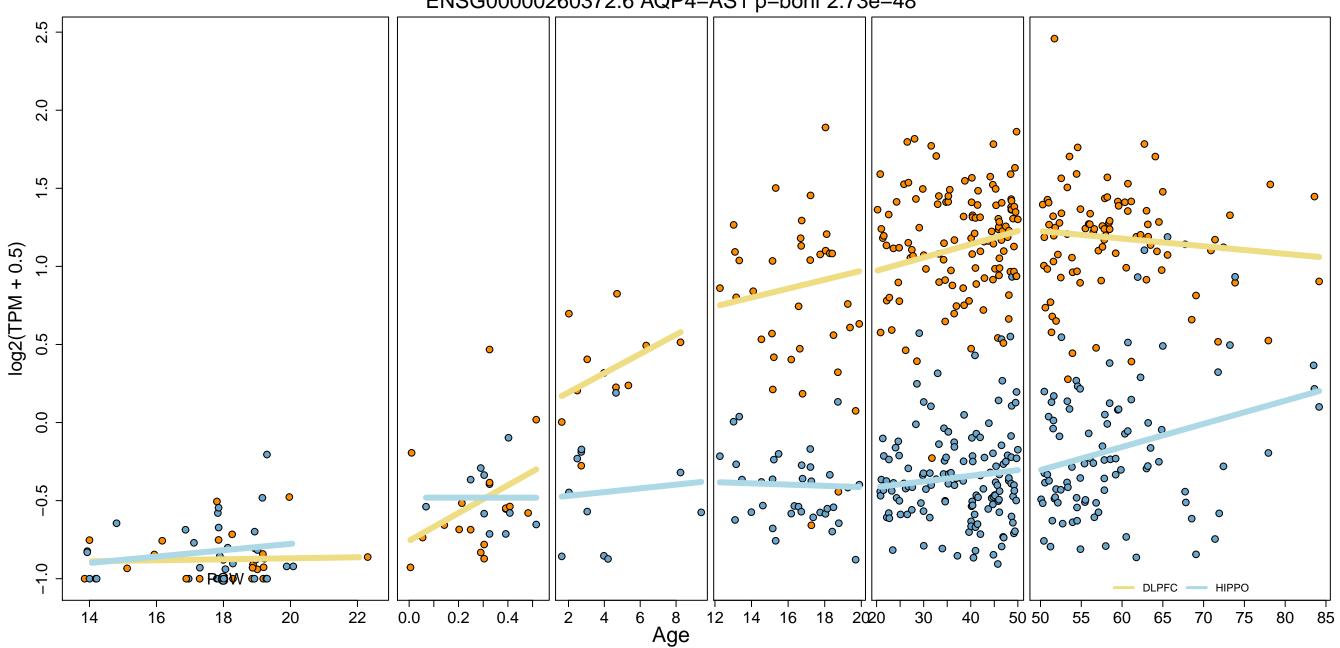
ENSG00000165553.4 NGB p-bonf 8.84e-49



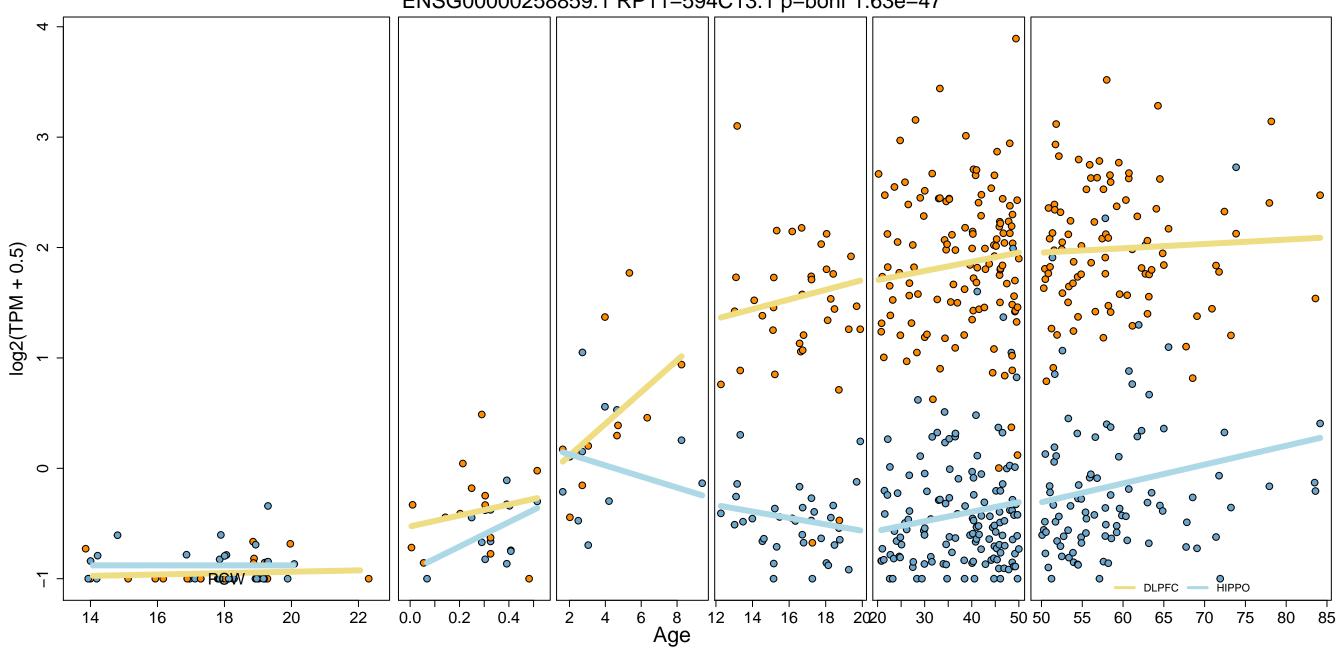
ENSG00000100628.11 ASB2 p-bonf 9.03e-49



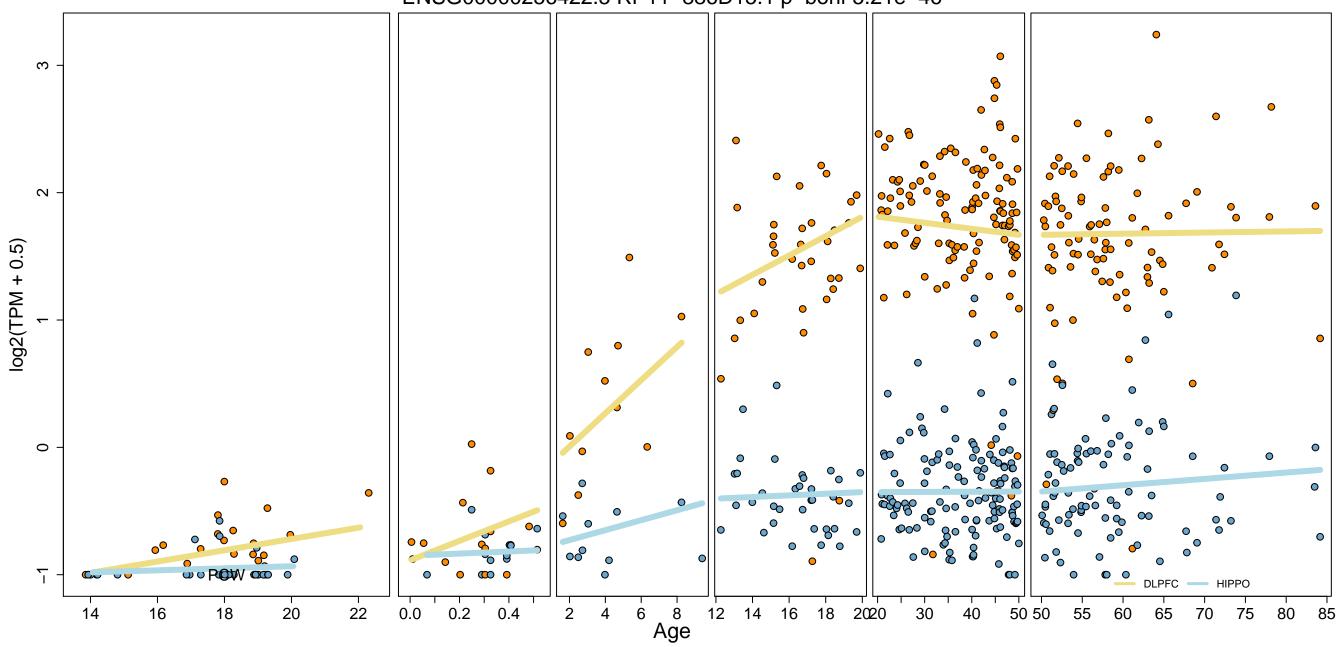
ENSG00000260372.6 AQP4-AS1 p-bonf 2.73e-48



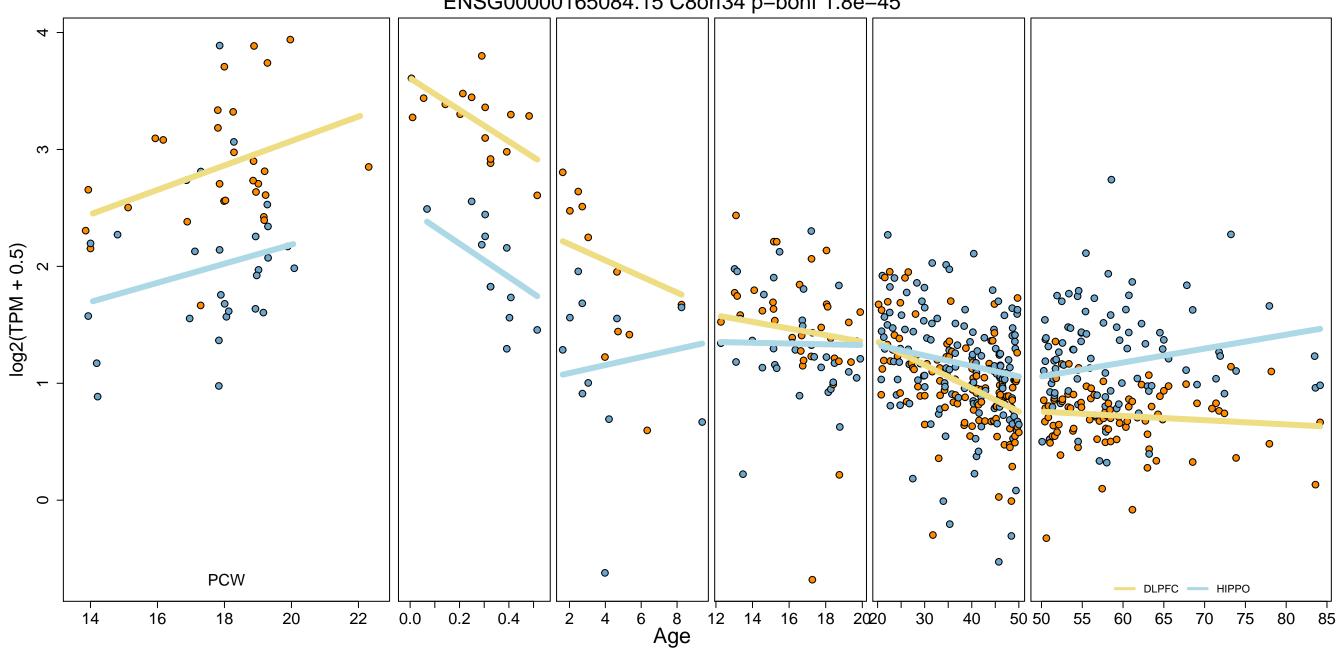
ENSG00000258859.1 RP11-594C13.1 p-bonf 1.63e-47



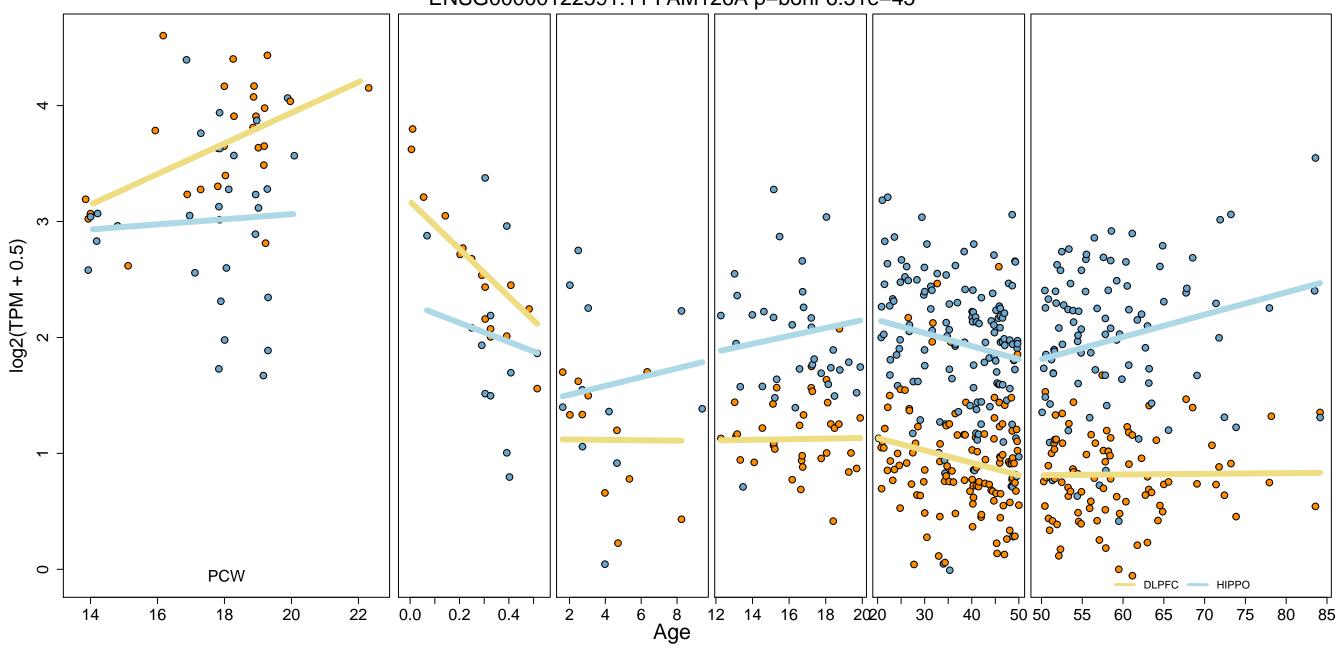
ENSG00000256422.5 RP11-886D15.1 p-bonf 5.21e-46



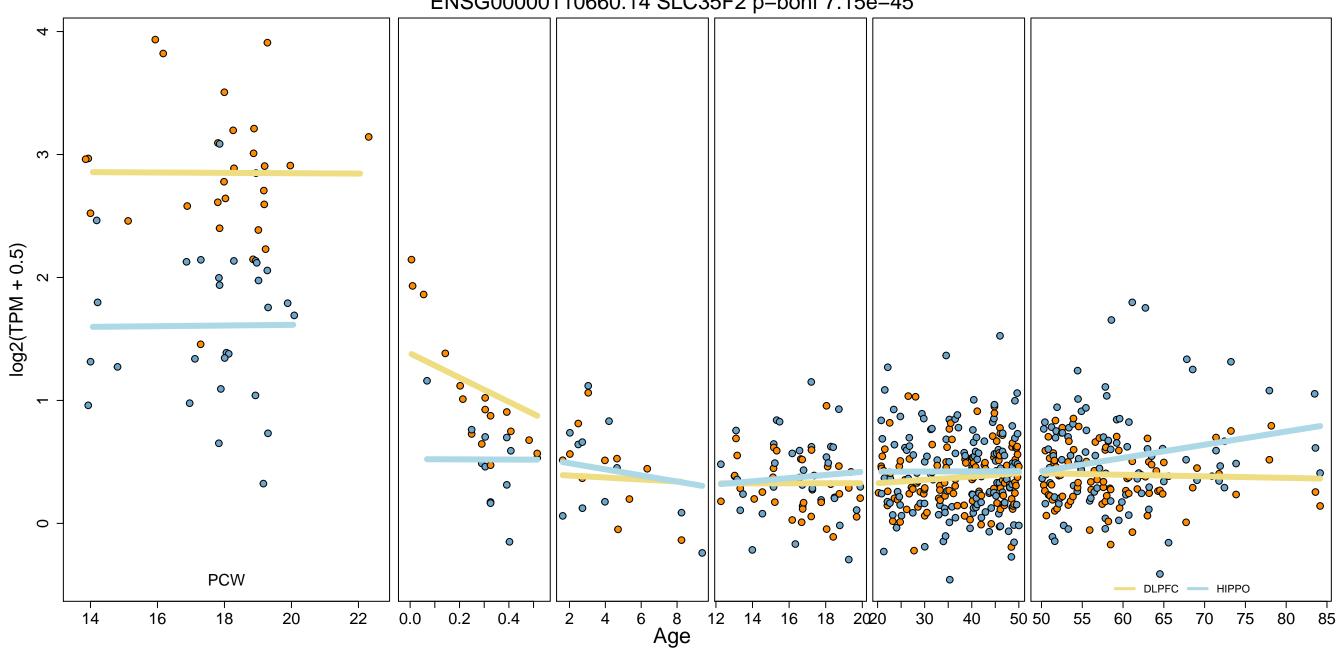
ENSG00000165084.15 C8orf34 p-bonf 1.8e-45



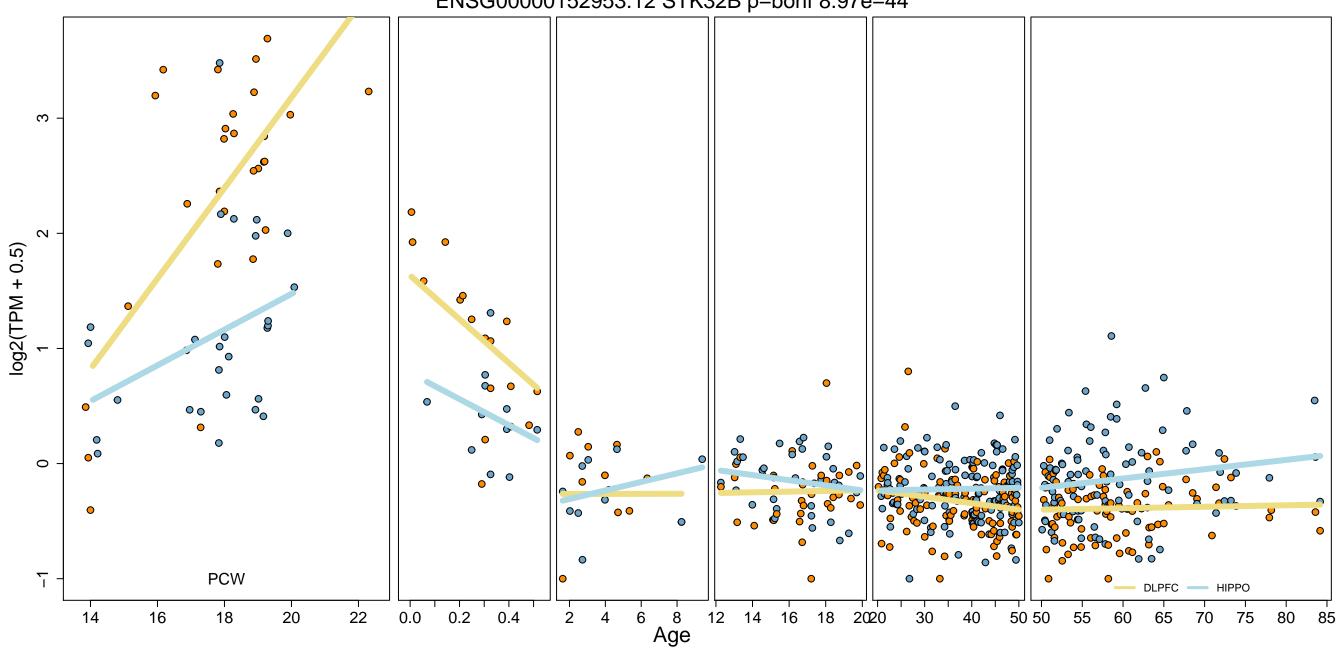
ENSG00000122591.11 FAM126A p-bonf 6.51e-45



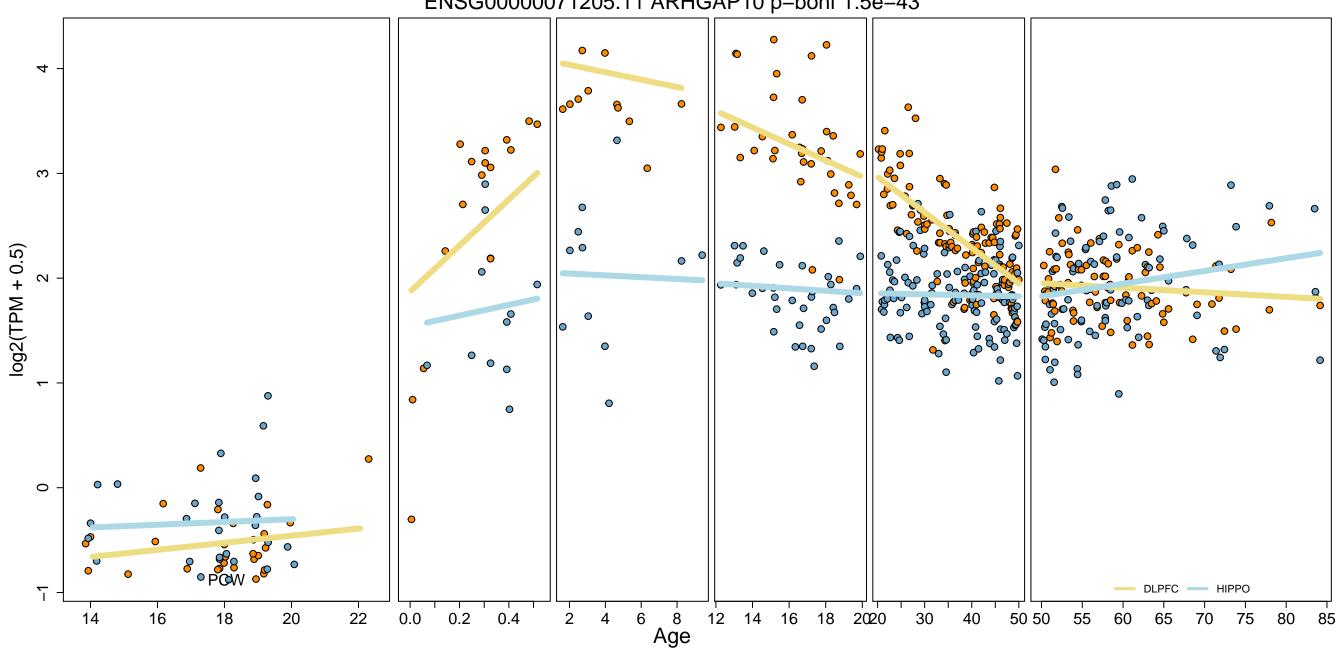
ENSG00000110660.14 SLC35F2 p-bonf 7.15e-45



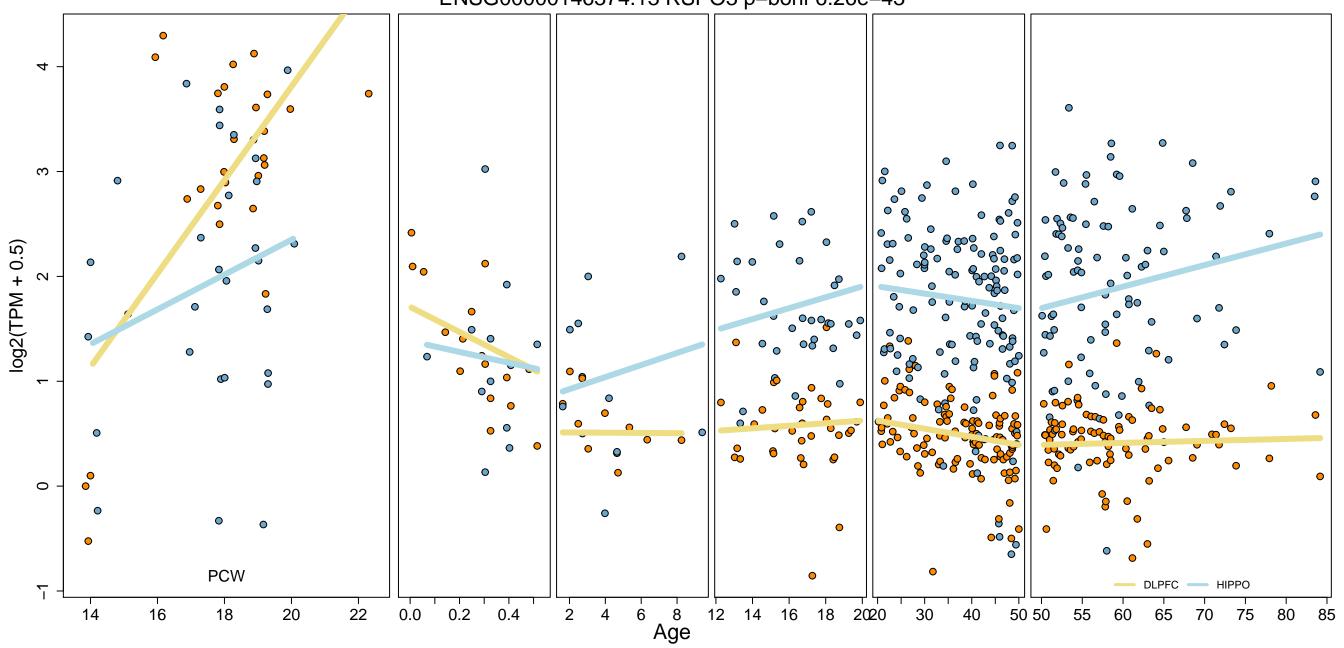
ENSG00000152953.12 STK32B p-bonf 8.97e-44



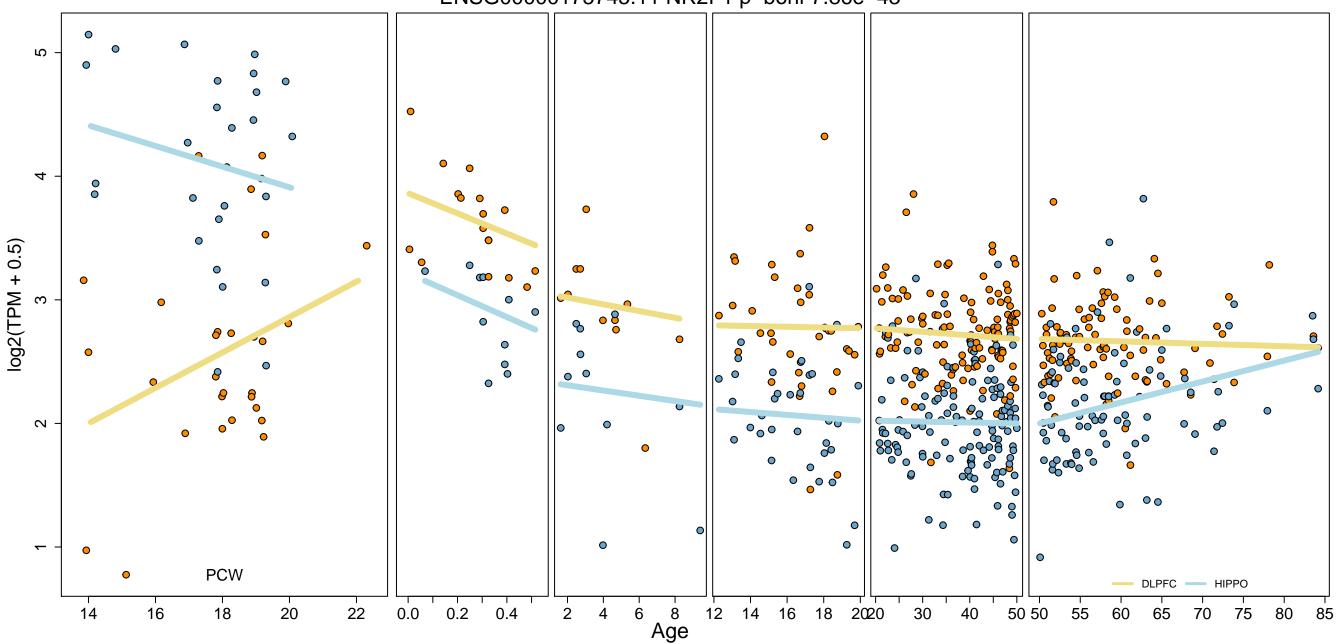
ENSG00000071205.11 ARHGAP10 p-bonf 1.5e-43



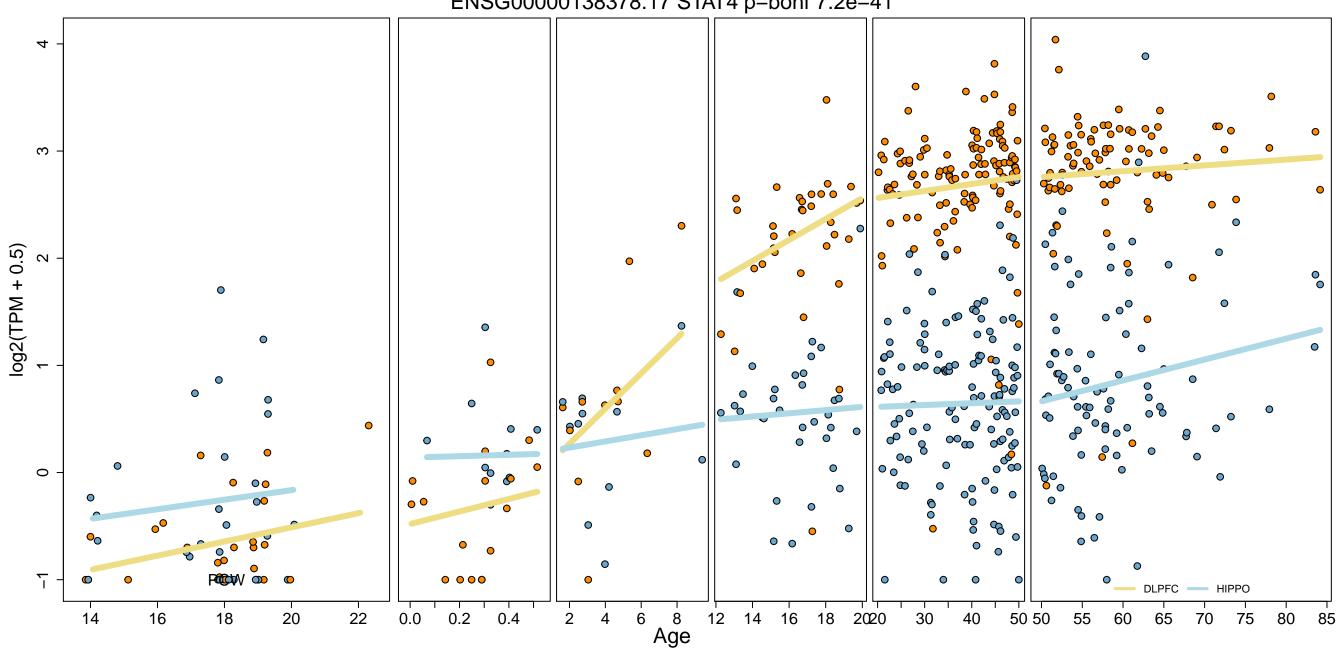
ENSG00000146374.13 RSPO3 p-bonf 6.26e-43



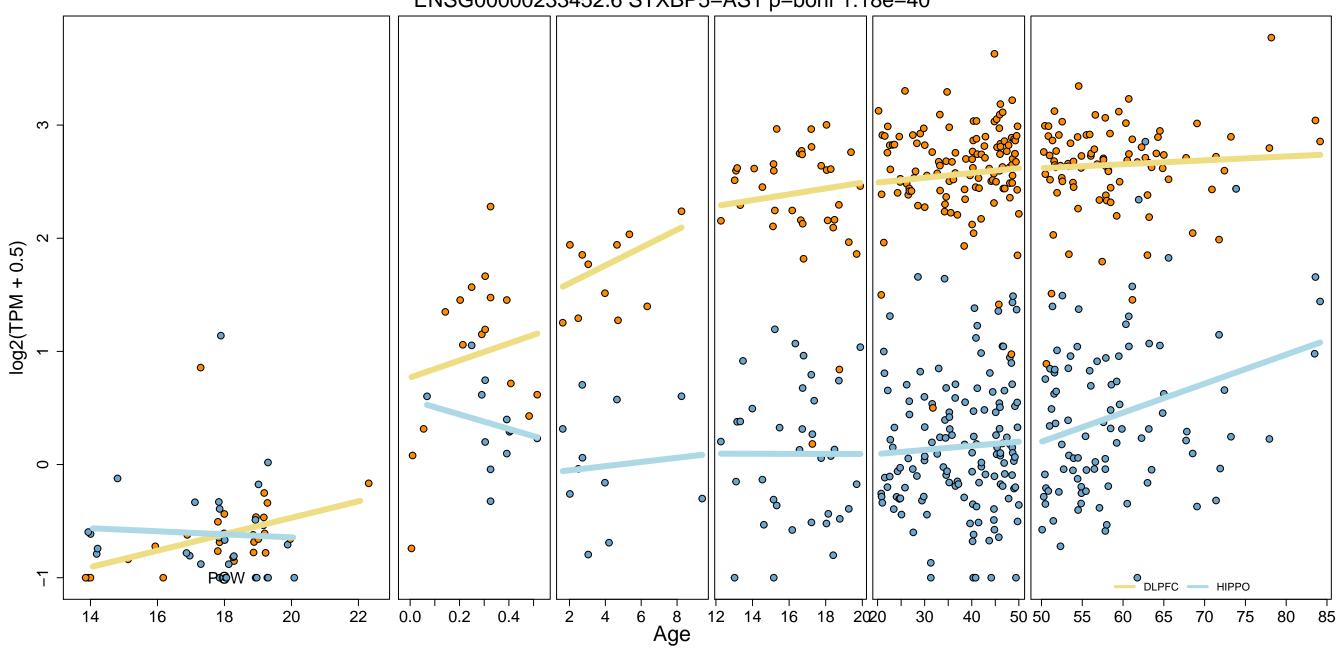
ENSG00000175745.11 NR2F1 p-bonf 7.86e-43



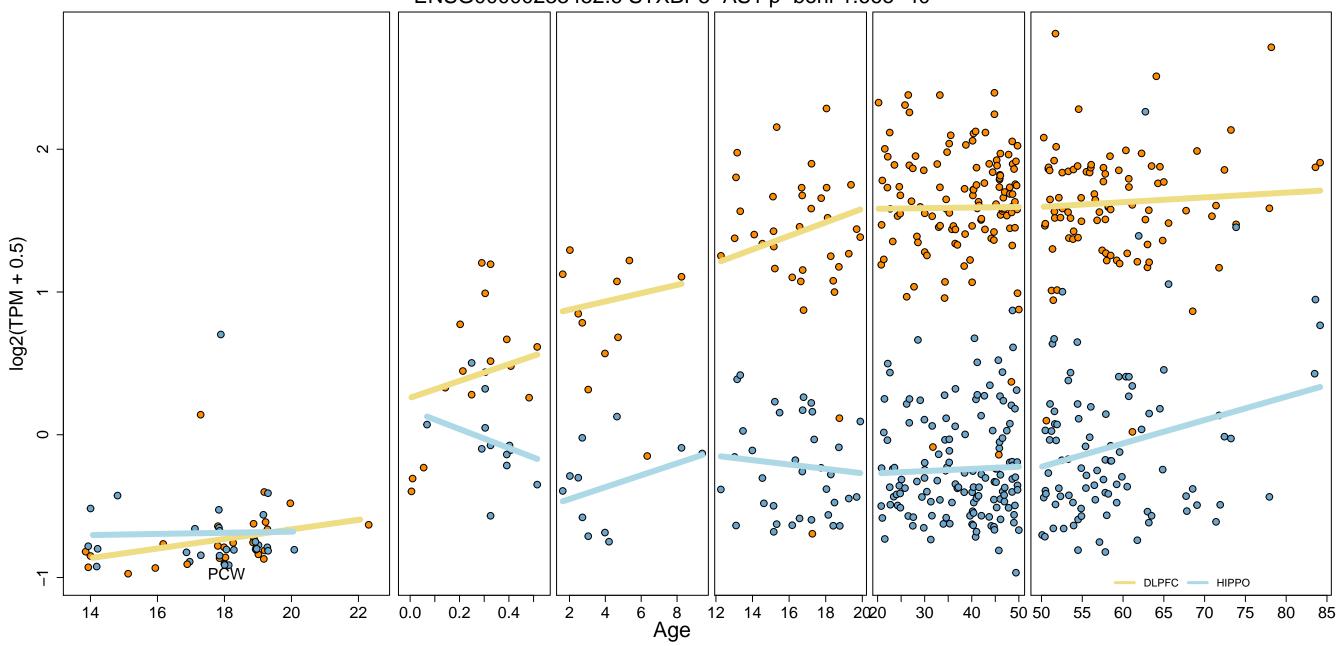
ENSG00000138378.17 STAT4 p-bonf 7.2e-41



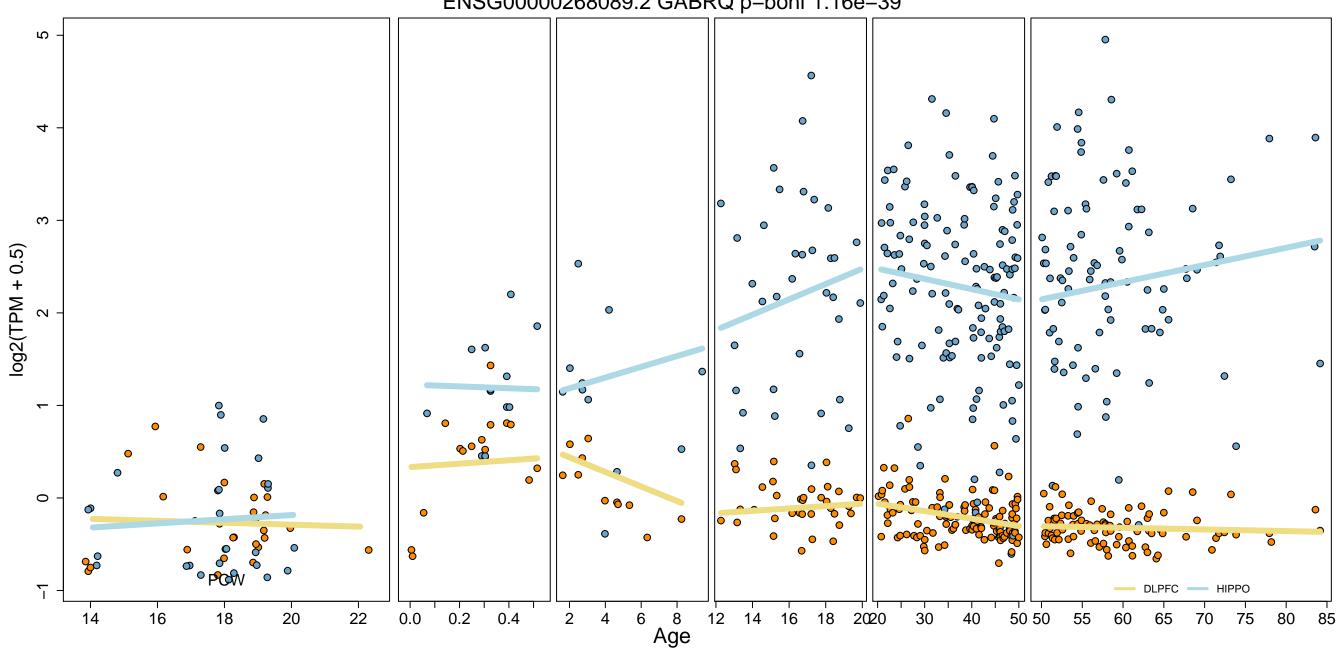
ENSG00000233452.6 STXBP5-AS1 p-bonf 1.18e-40



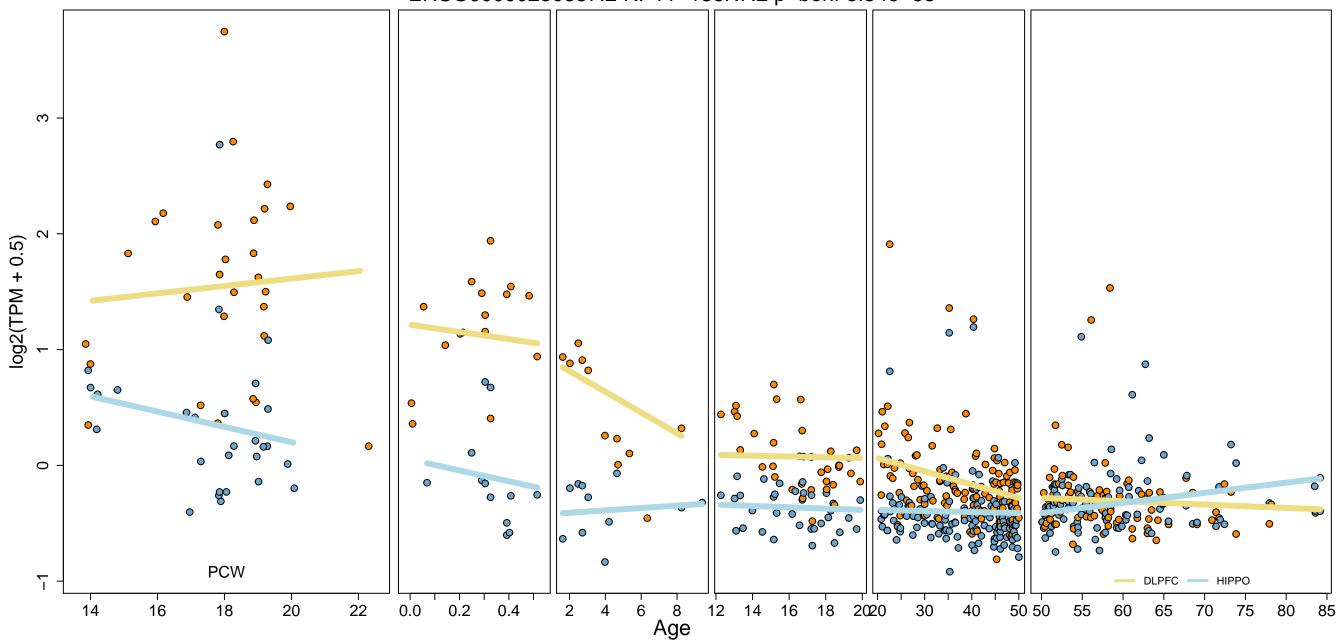
ENSG00000233452.6 STXBP5-AS1 p-bonf 1.66e-40



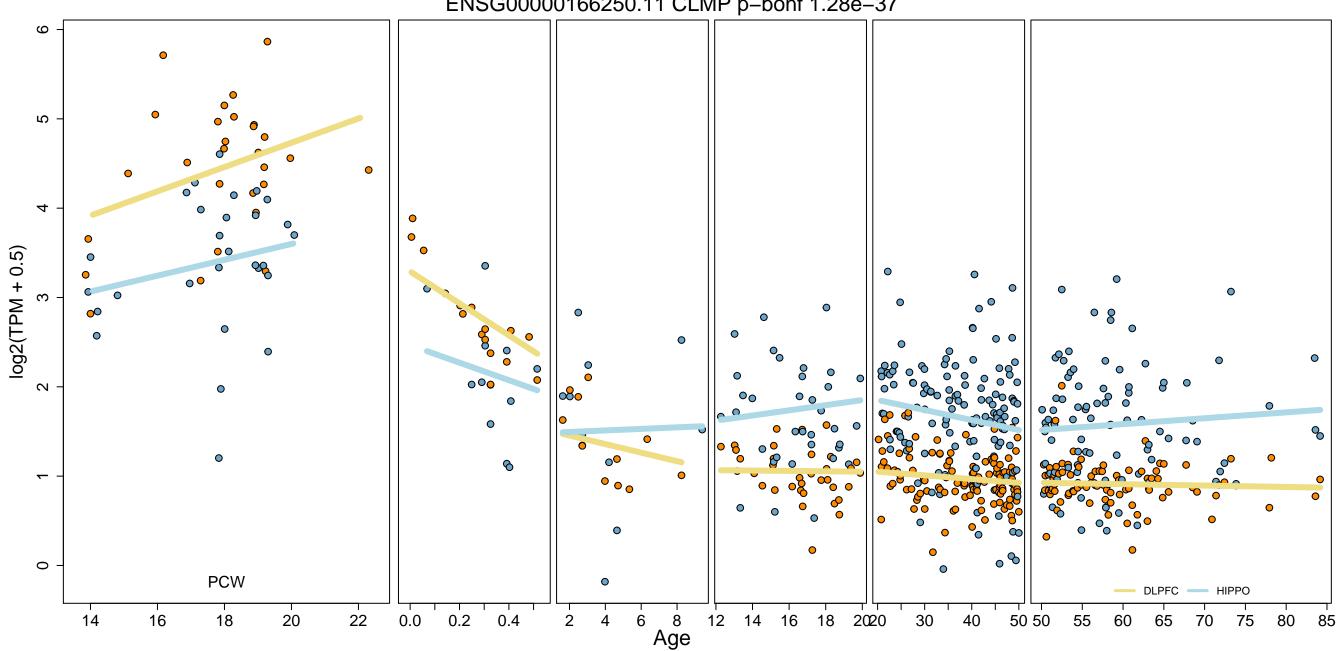
ENSG00000268089.2 GABRQ p-bonf 1.16e-39



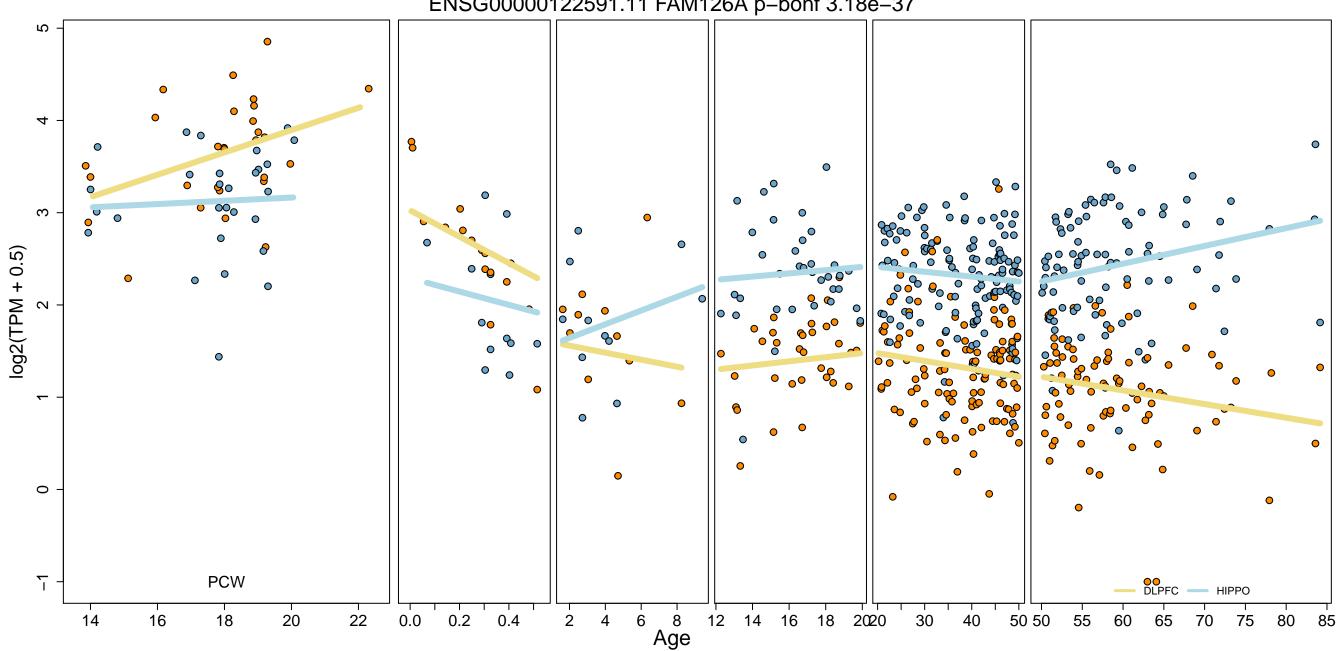
ENSG00000250387.2 RP11-136K7.2 p-bonf 8.34e-38



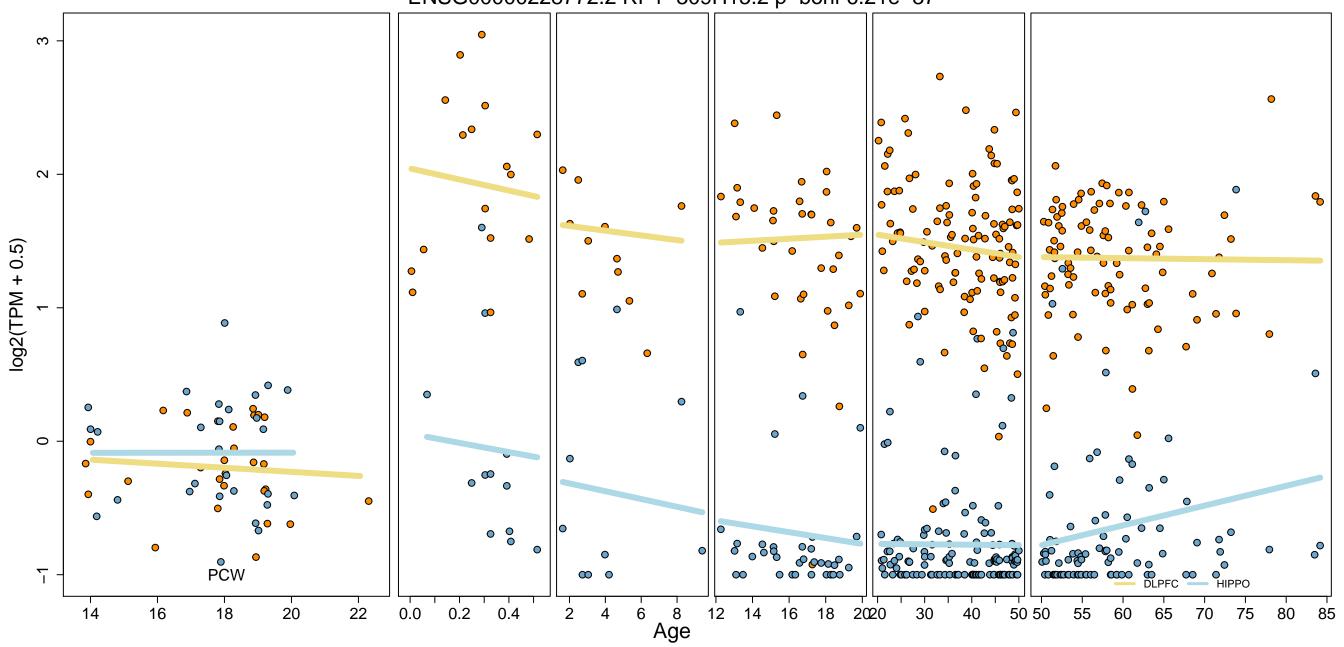
ENSG00000166250.11 CLMP p-bonf 1.28e-37



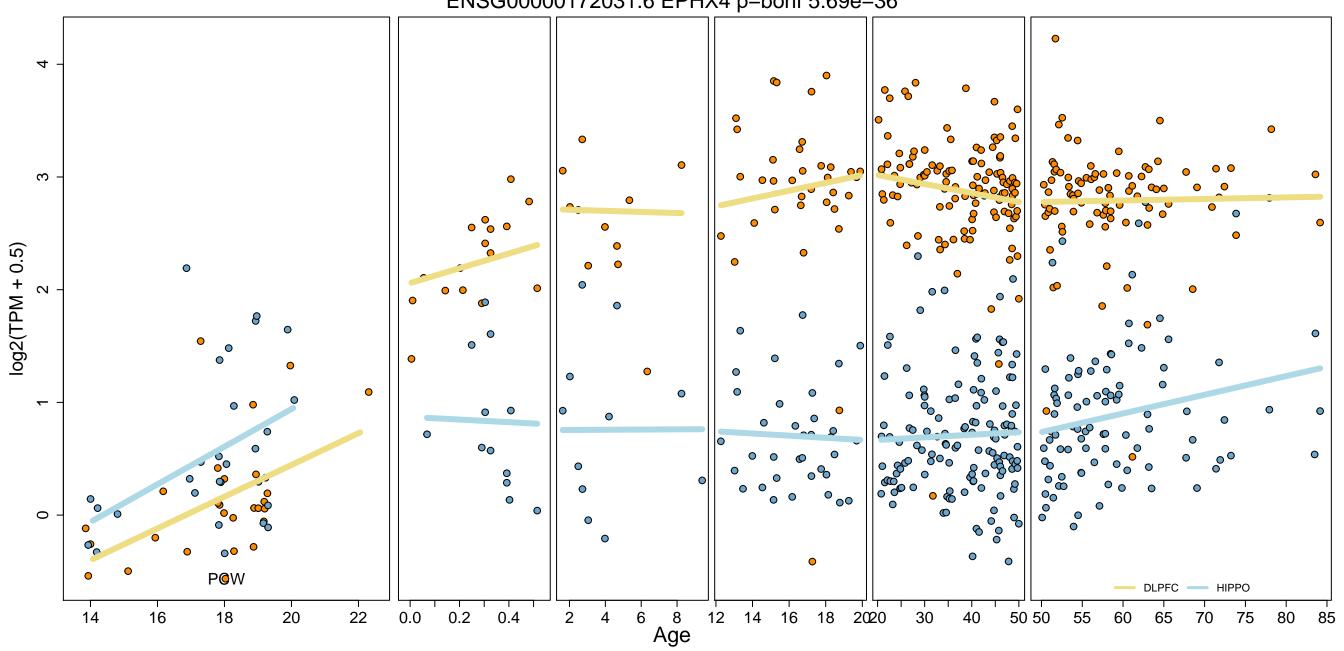
ENSG00000122591.11 FAM126A p-bonf 3.18e-37



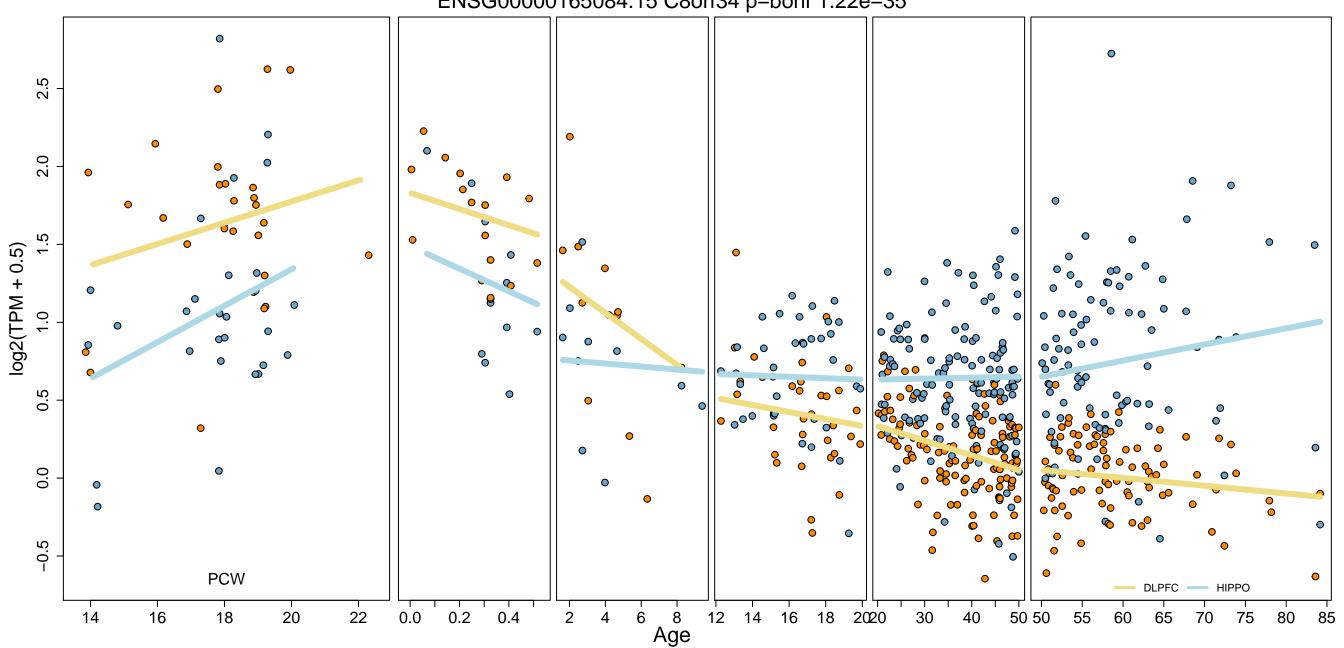
ENSG00000228772.2 RP1-309H15.2 p-bonf 6.21e-37



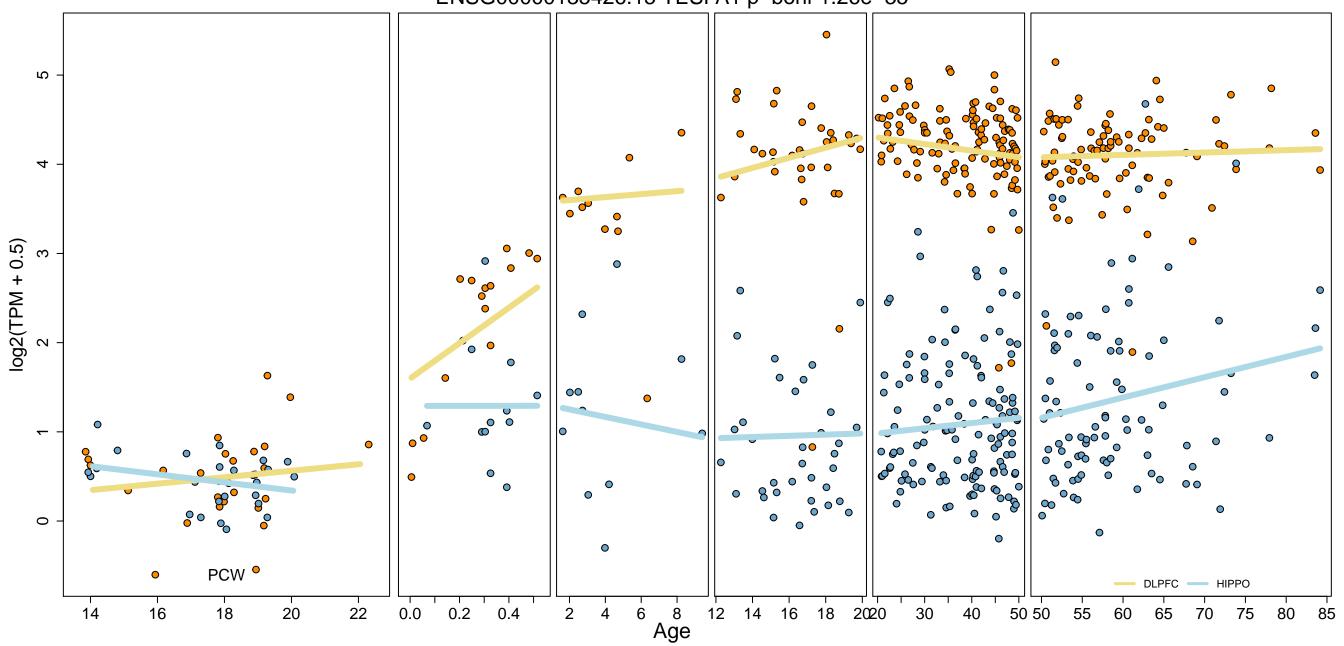
ENSG00000172031.6 EPHX4 p-bonf 5.69e-36



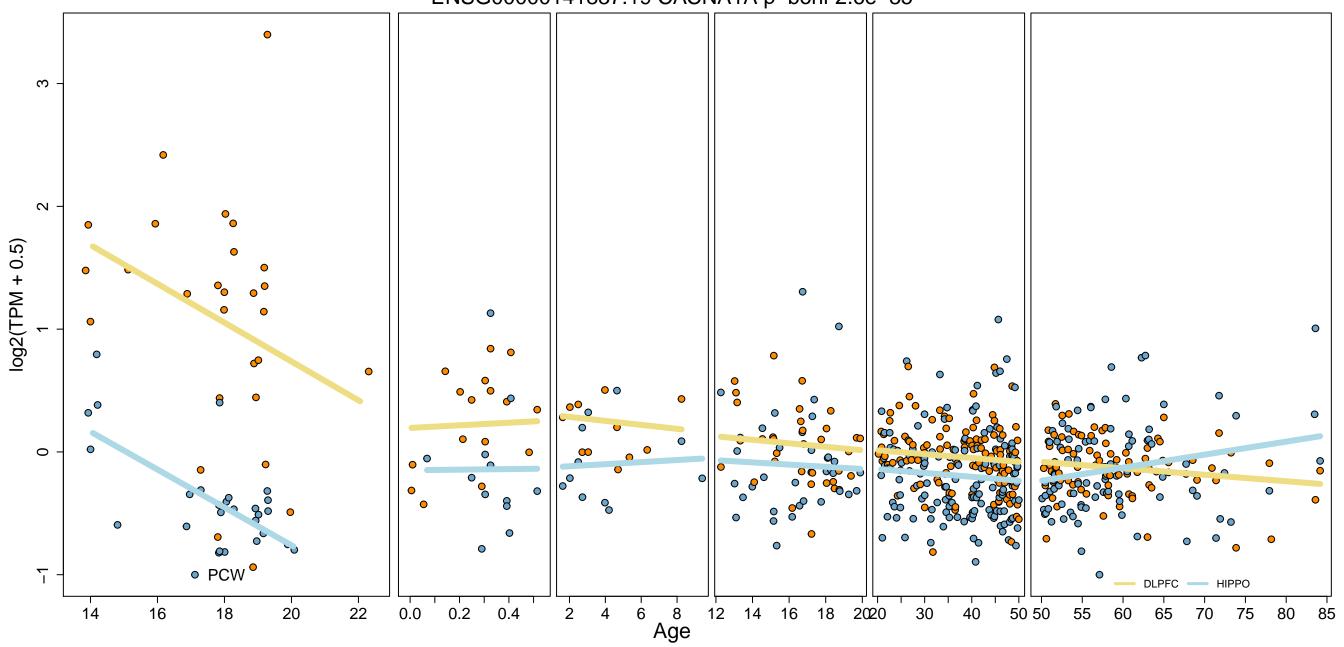
ENSG00000165084.15 C8orf34 p-bonf 1.22e-35



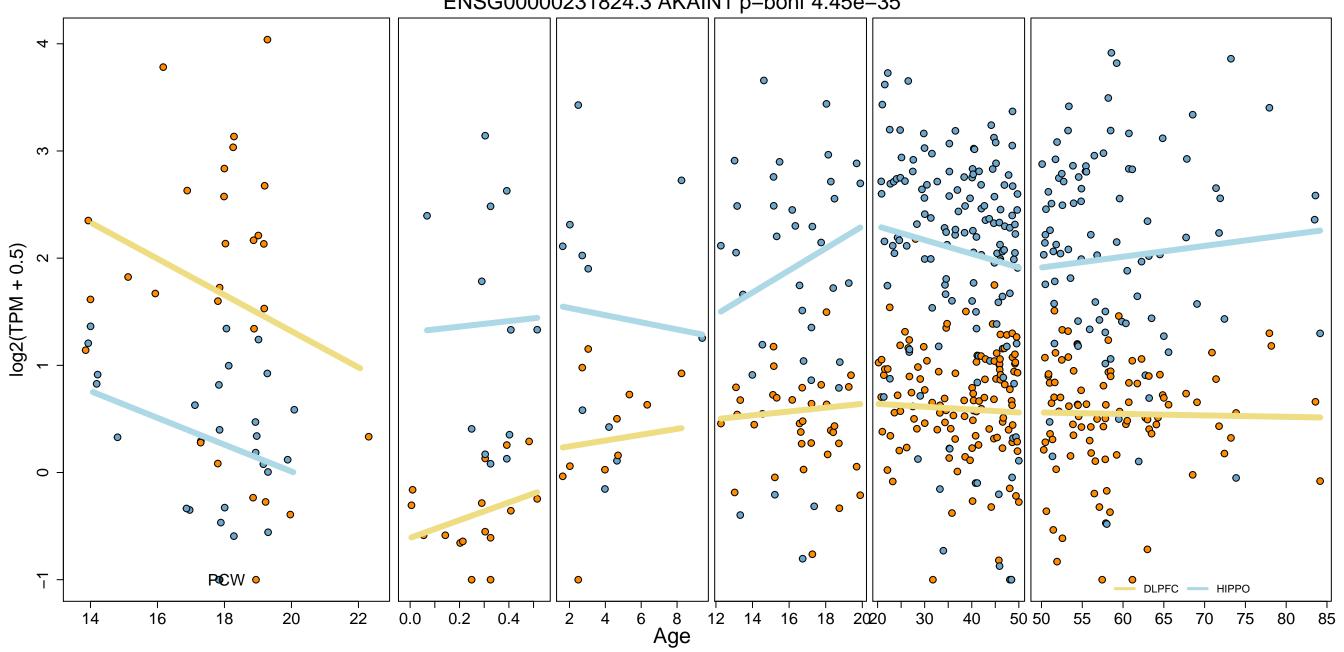
ENSG00000135426.15 TESPA1 p-bonf 1.26e-35



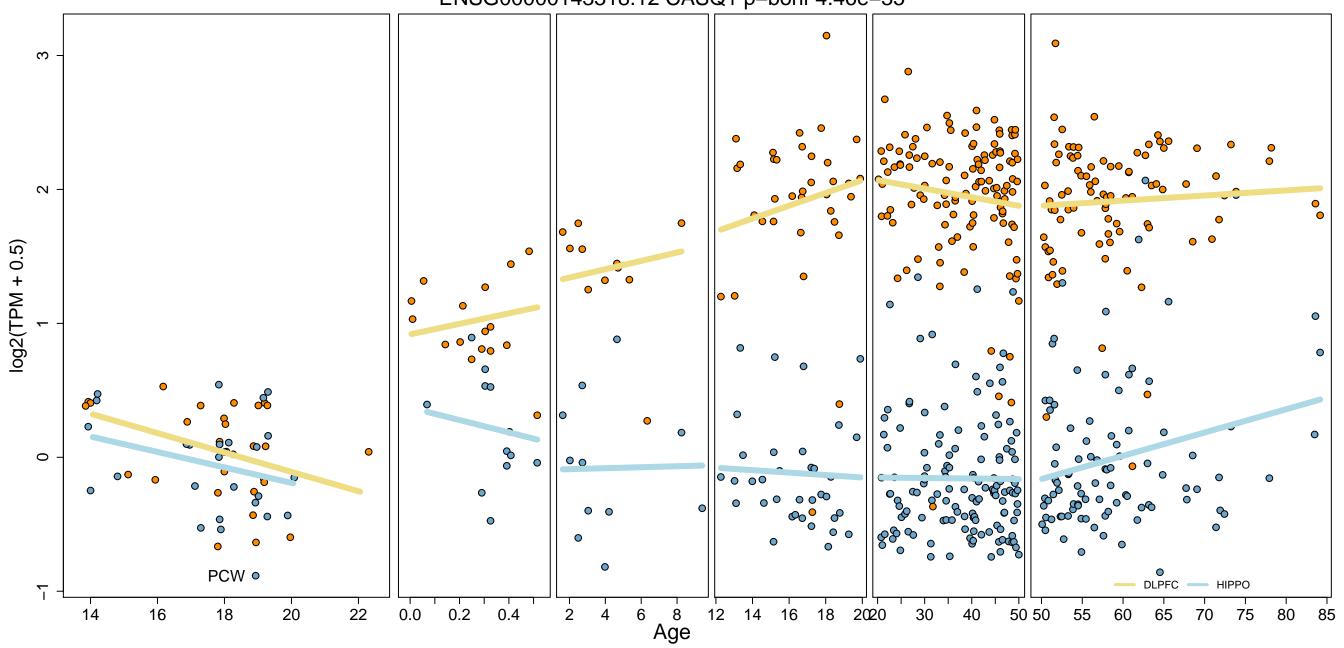
ENSG00000141837.19 CACNA1A p-bonf 2.6e-35



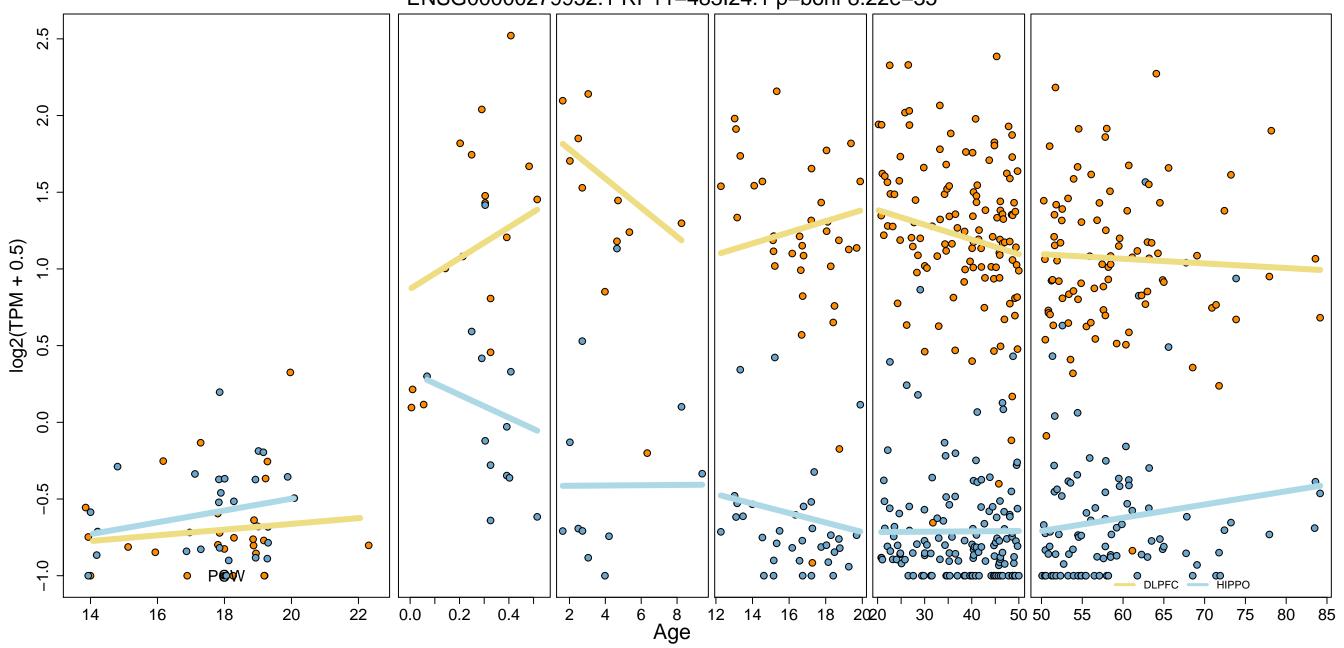
ENSG00000231824.3 AKAIN1 p-bonf 4.45e-35



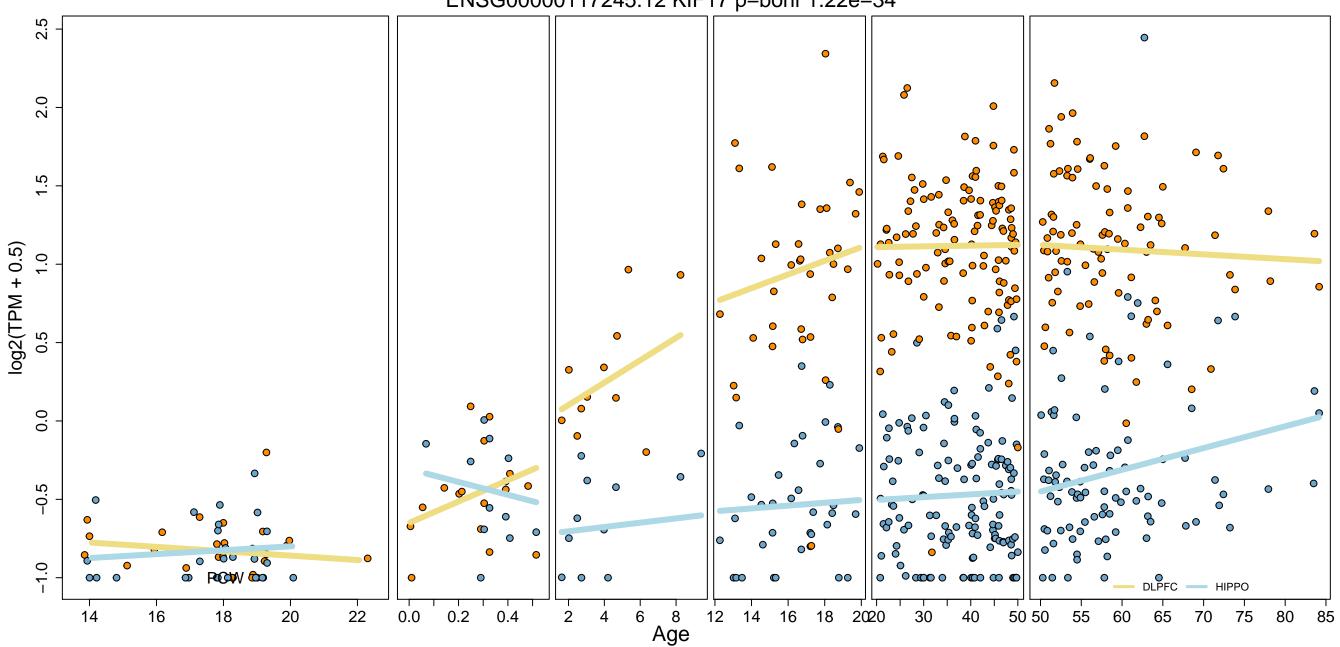
ENSG00000143318.12 CASQ1 p-bonf 4.46e-35



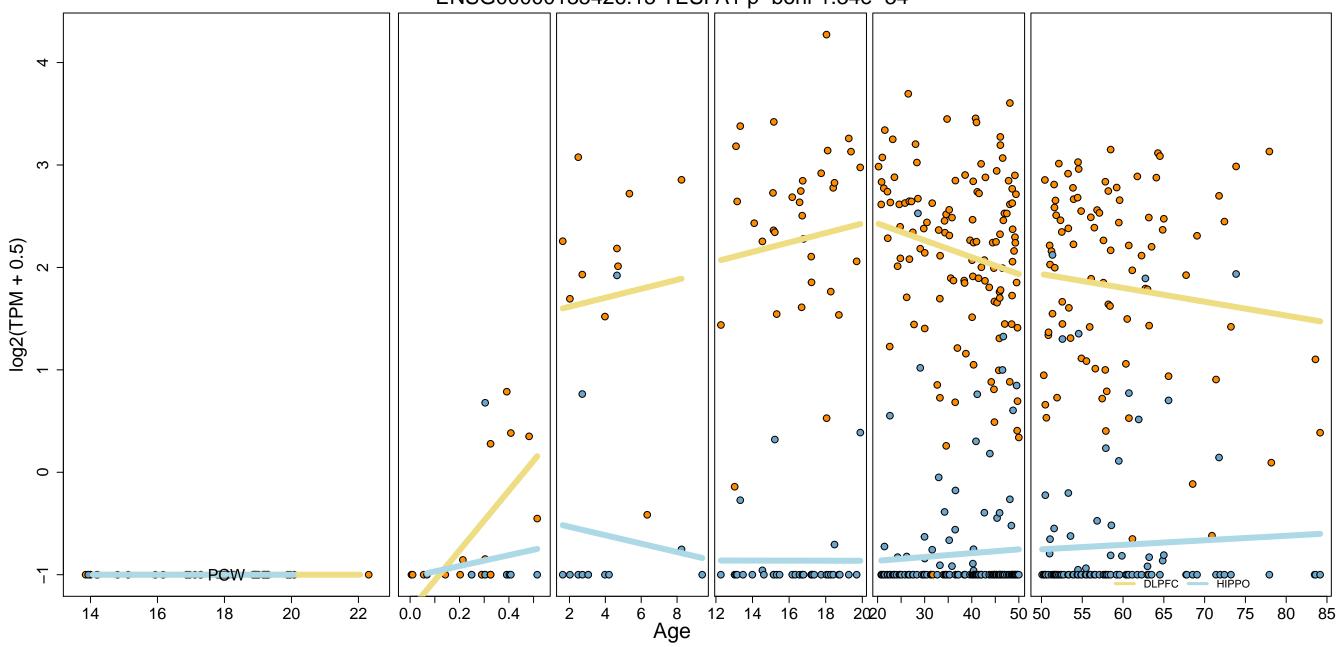
ENSG00000279952.1 RP11-483I24.1 p-bonf 8.22e-35



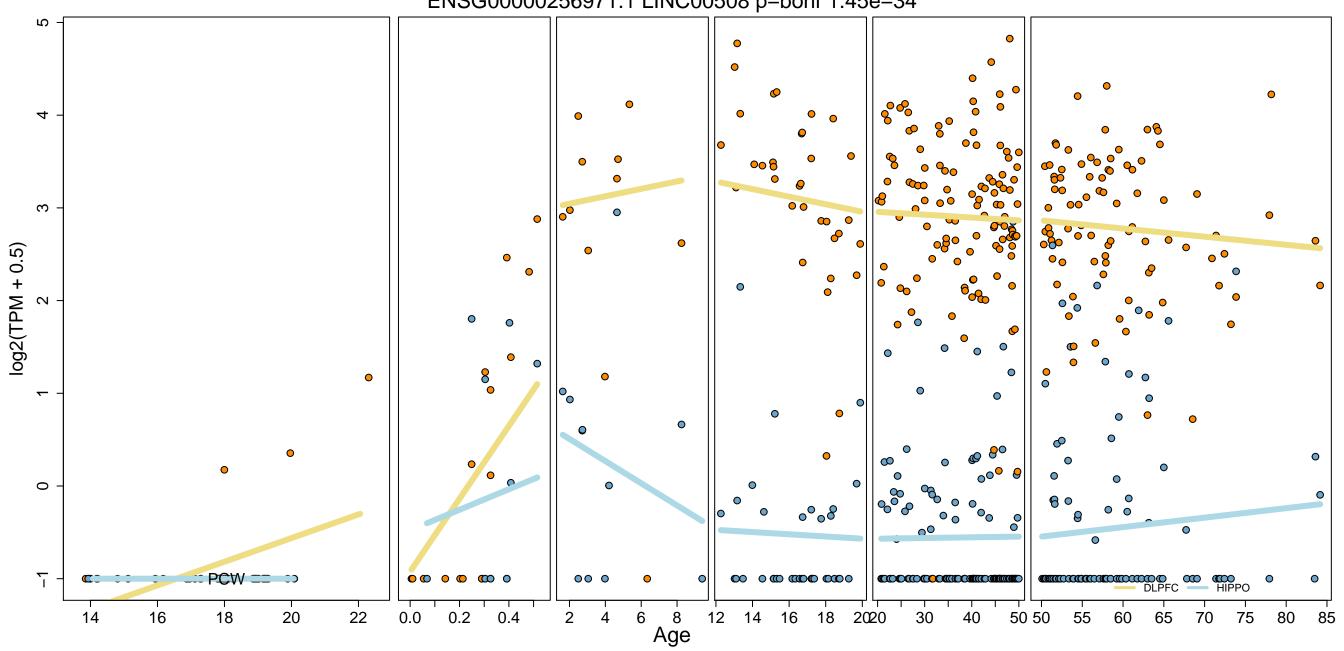
ENSG00000117245.12 KIF17 p-bonf 1.22e-34



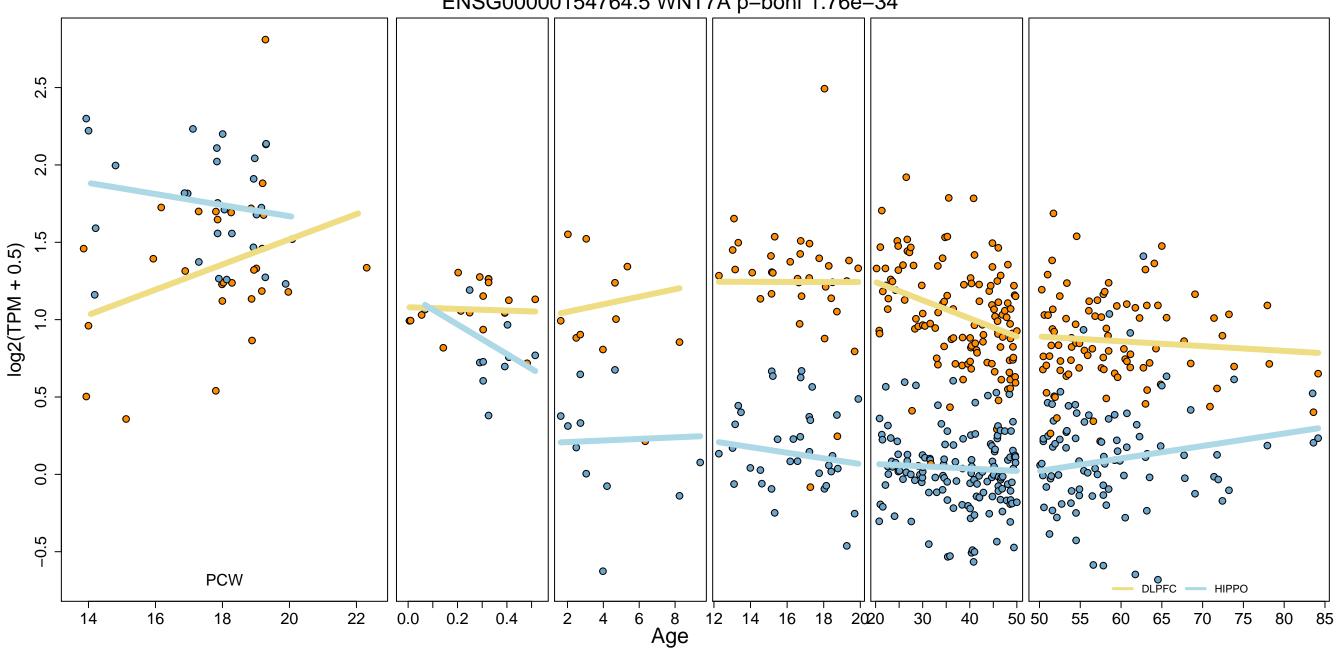
ENSG00000135426.15 TESPA1 p-bonf 1.34e-34



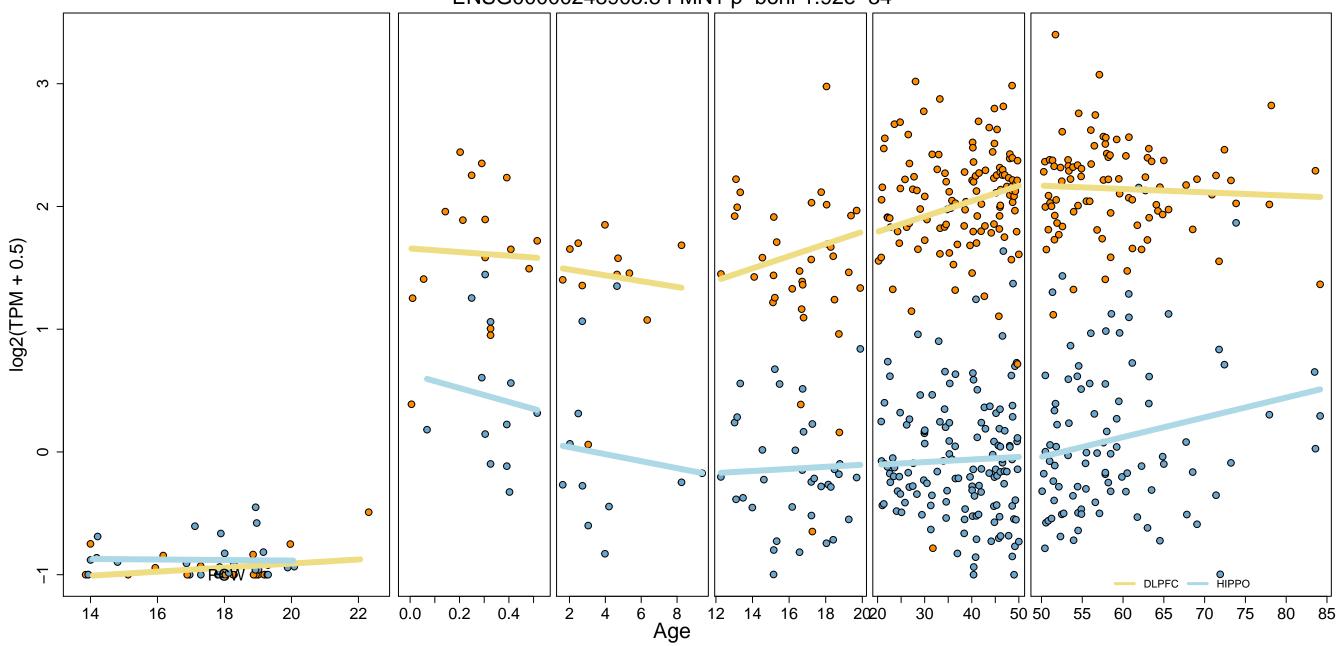
ENSG00000256971.1 LINC00508 p-bonf 1.45e-34



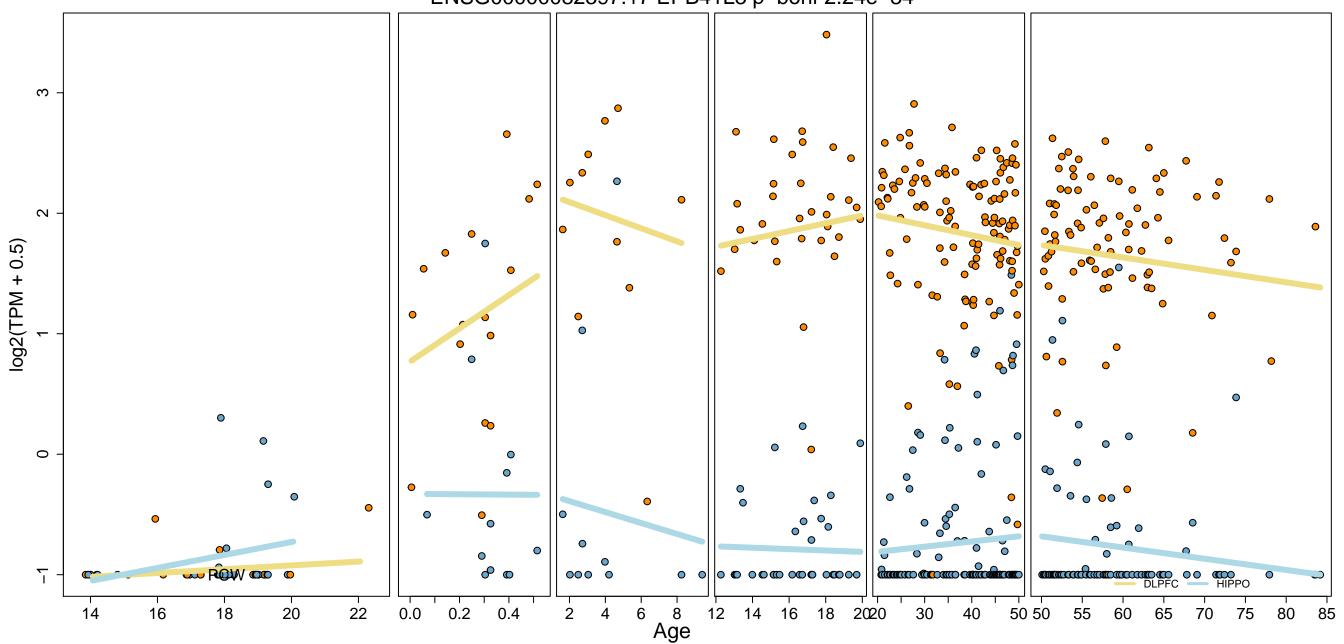
ENSG00000154764.5 WNT7A p-bonf 1.76e-34



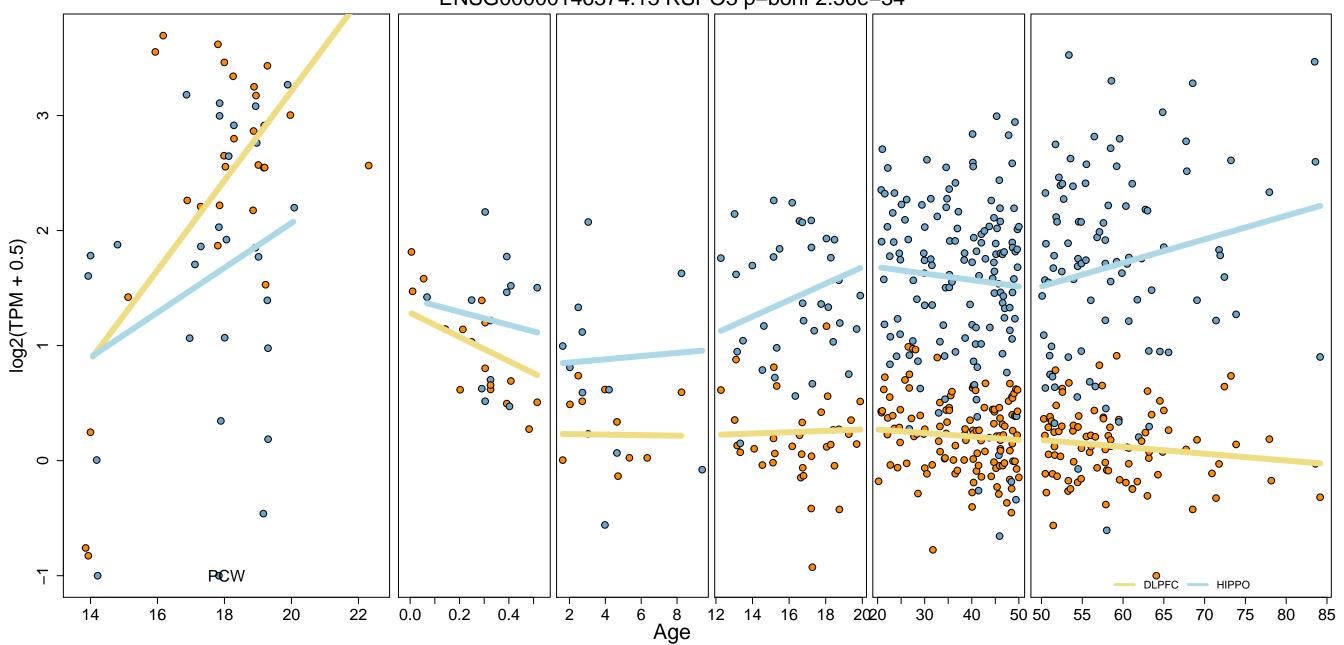
ENSG00000248905.8 FMN1 p-bonf 1.92e-34



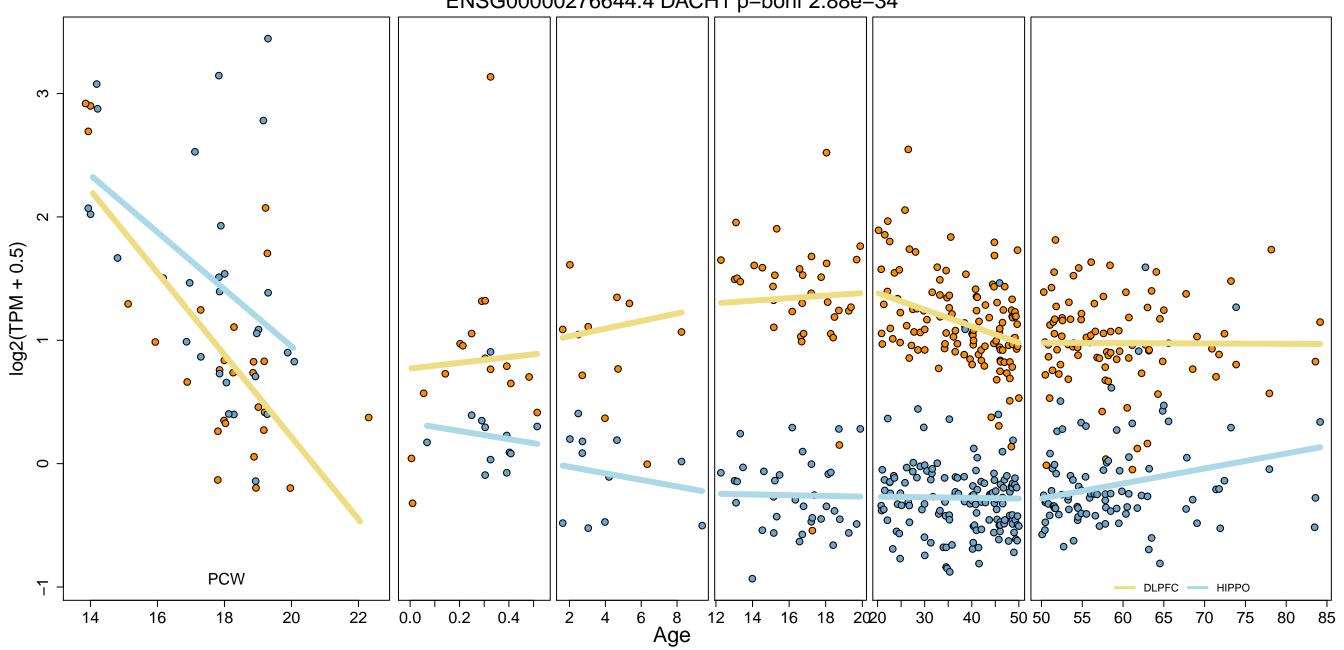
ENSG00000082397.17 EPB41L3 p-bonf 2.24e-34



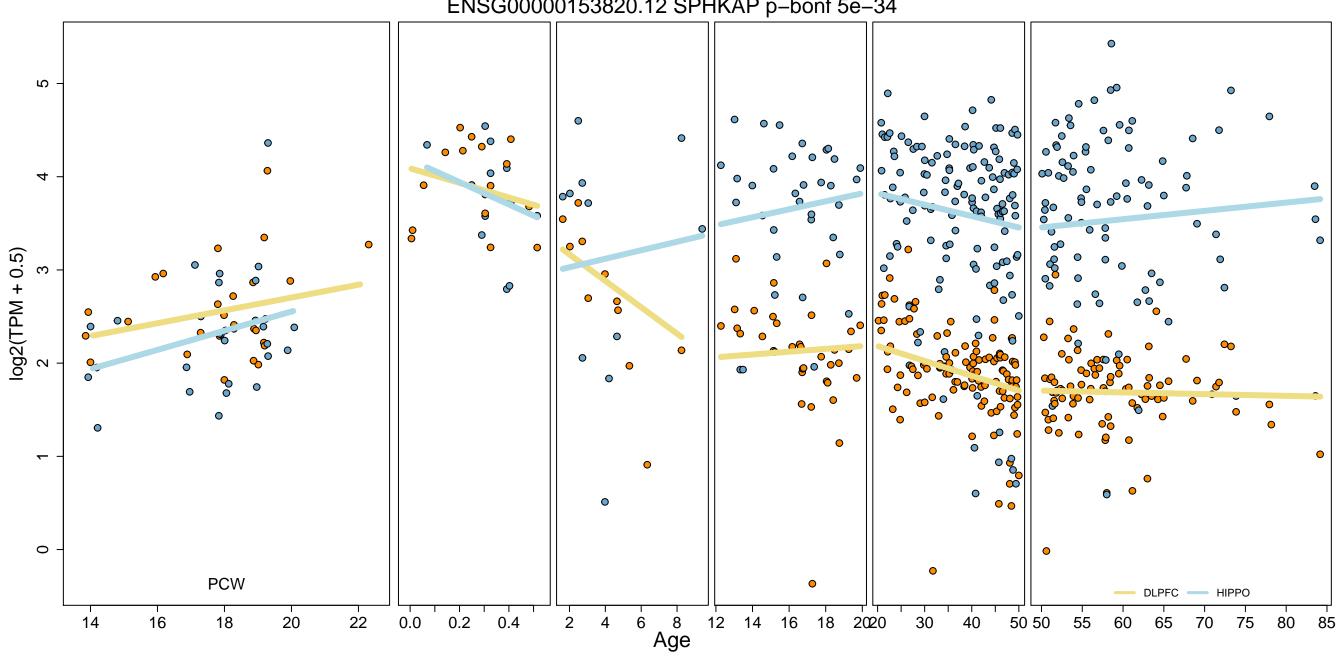
ENSG00000146374.13 RSPO3 p-bonf 2.56e-34



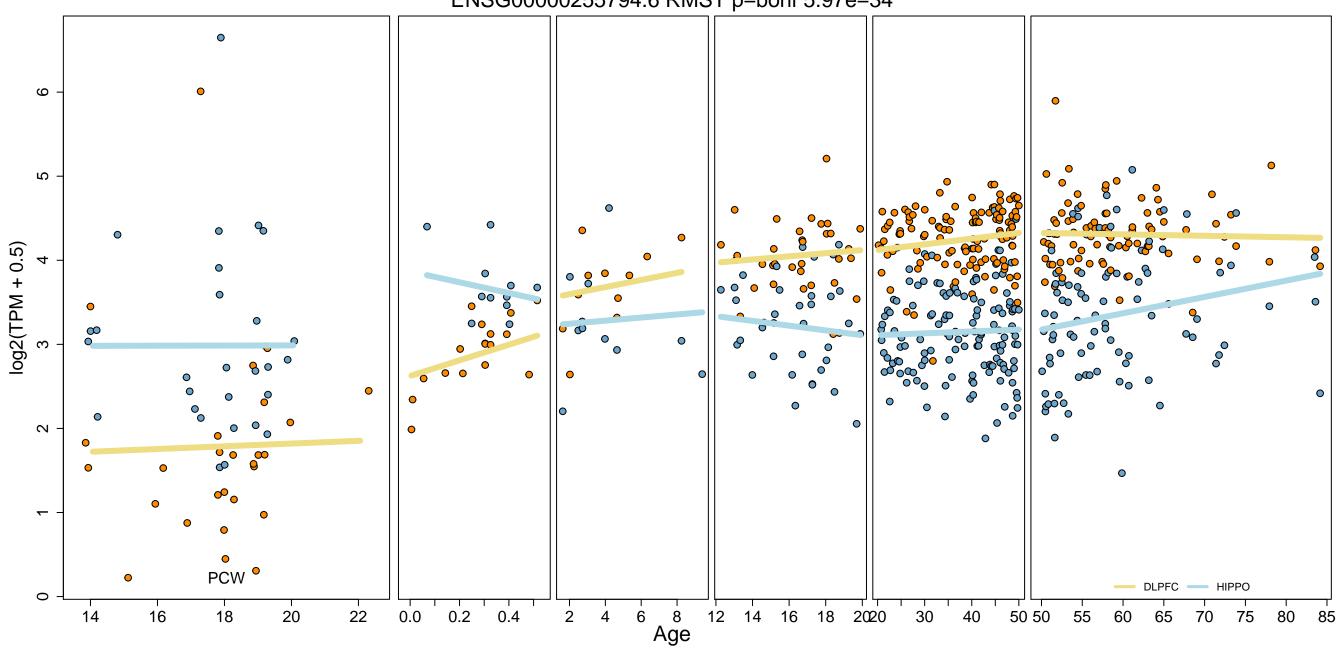
ENSG00000276644.4 DACH1 p-bonf 2.88e-34



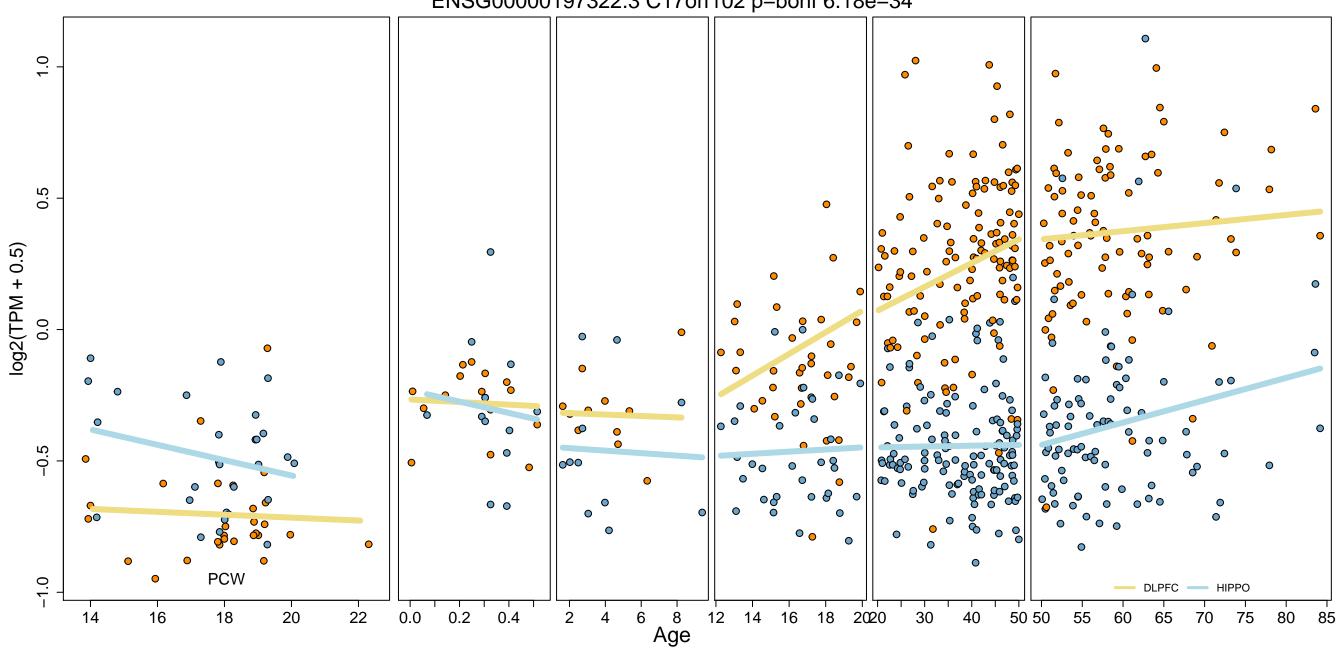
ENSG00000153820.12 SPHKAP p-bonf 5e-34



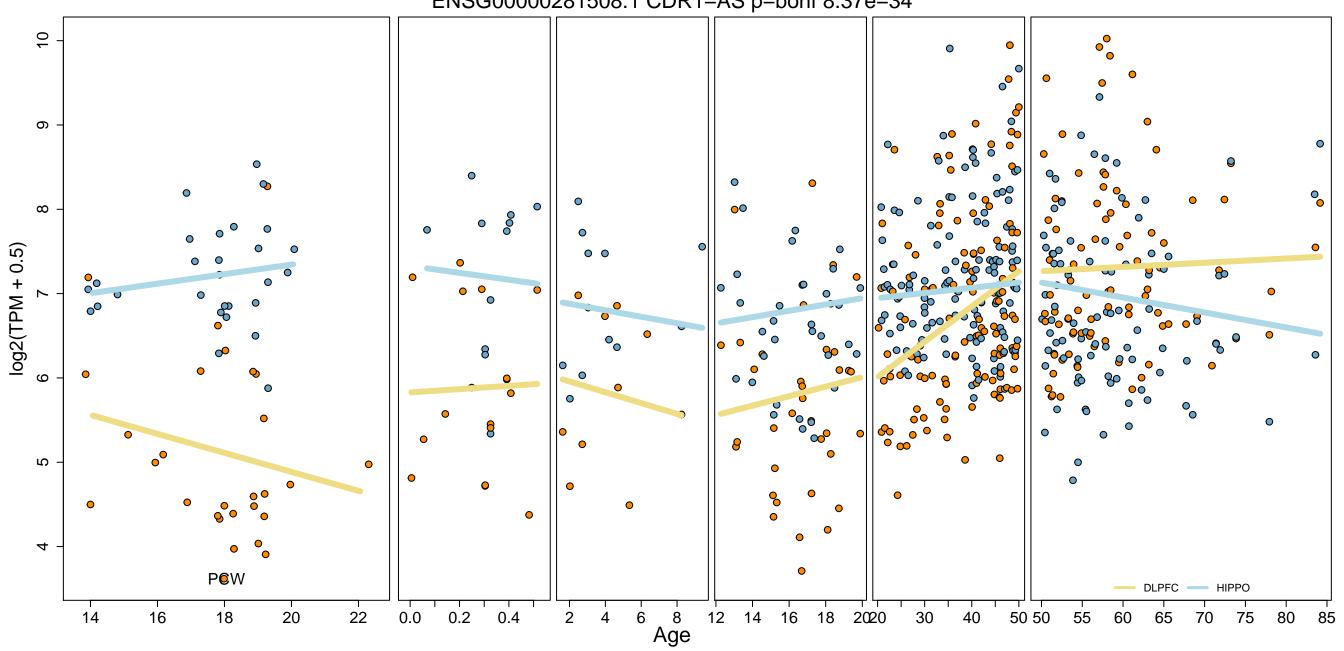
ENSG00000255794.6 RMST p-bonf 5.97e-34



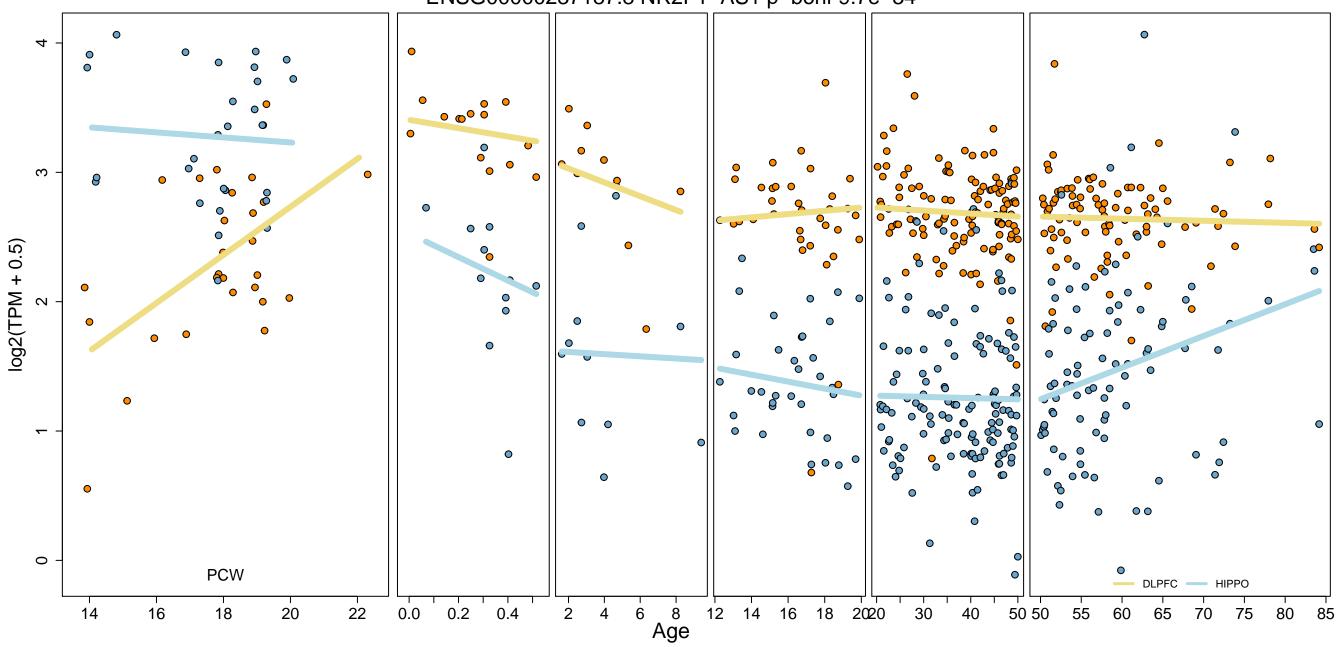
ENSG00000197322.3 C17orf102 p-bonf 6.18e-34



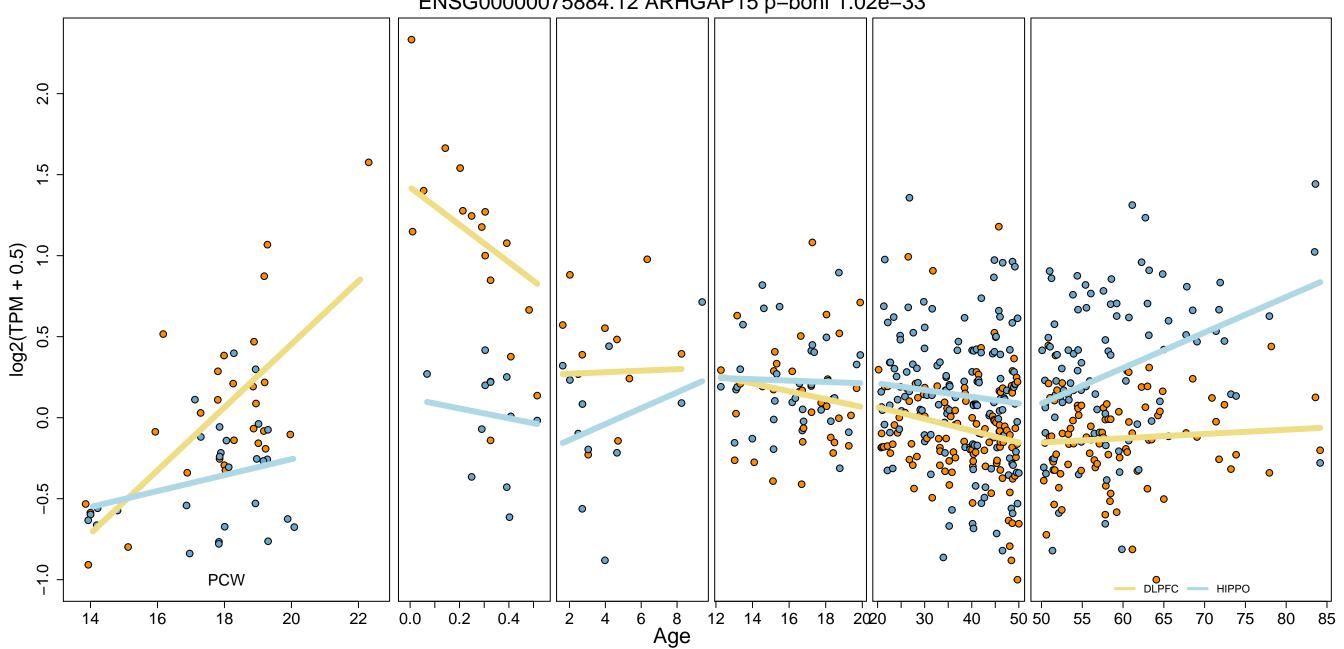
ENSG00000281508.1 CDR1-AS p-bonf 8.37e-34



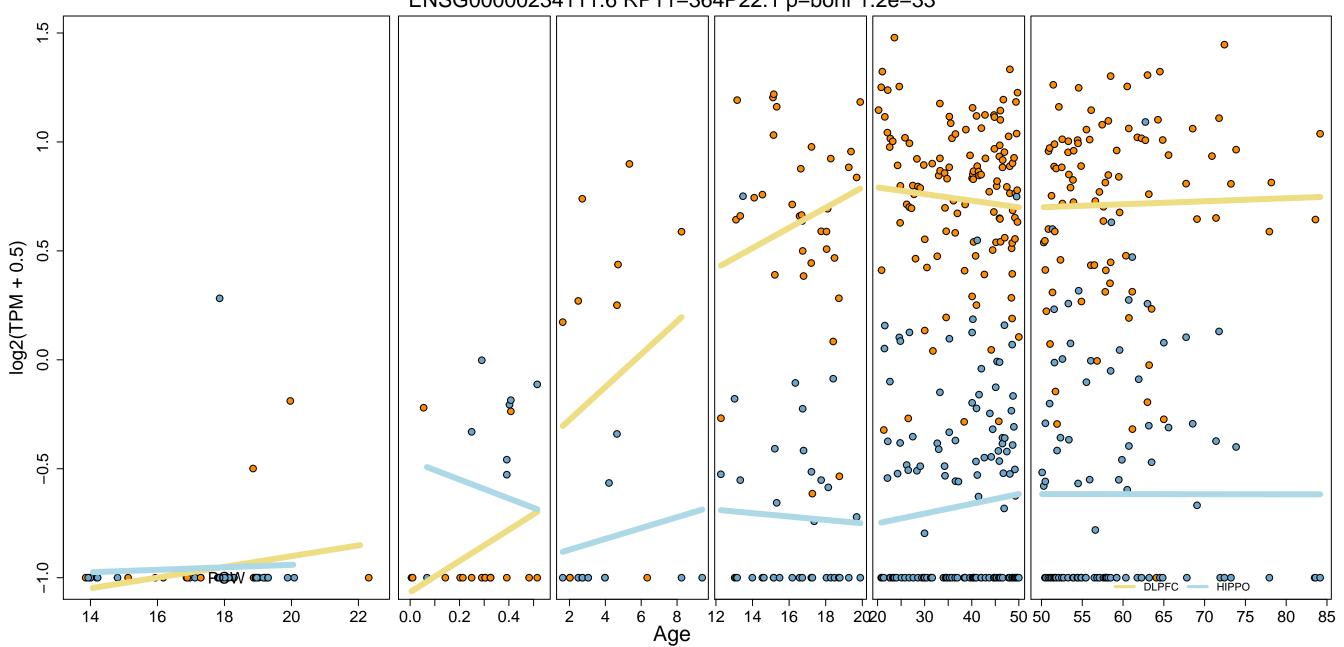
ENSG00000237187.8 NR2F1-AS1 p-bonf 9.7e-34



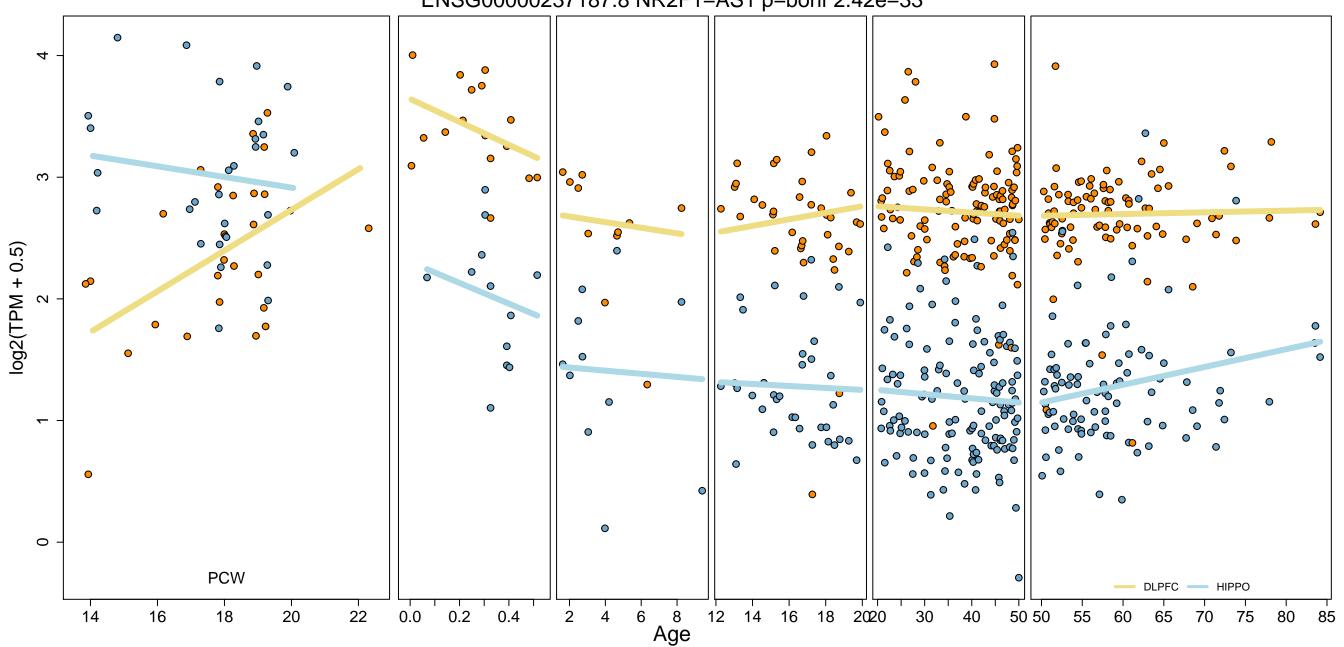
ENSG00000075884.12 ARHGAP15 p-bonf 1.02e-33



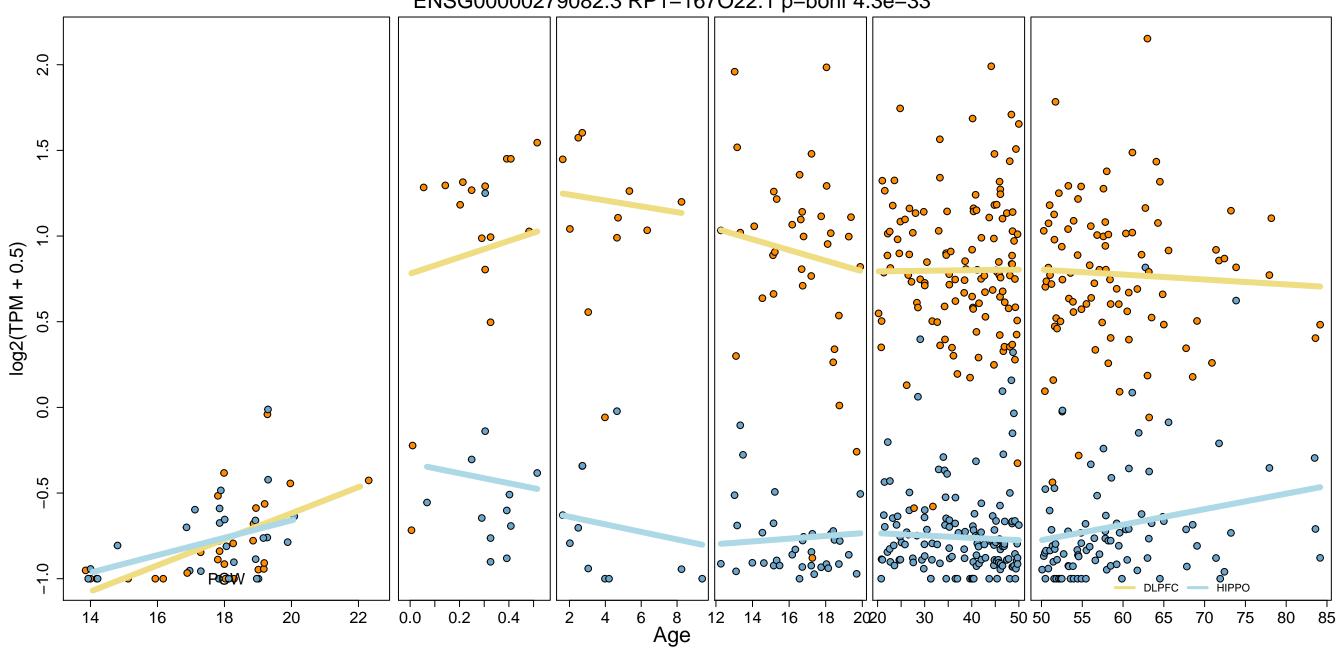
ENSG00000234111.6 RP11-364P22.1 p-bonf 1.2e-33



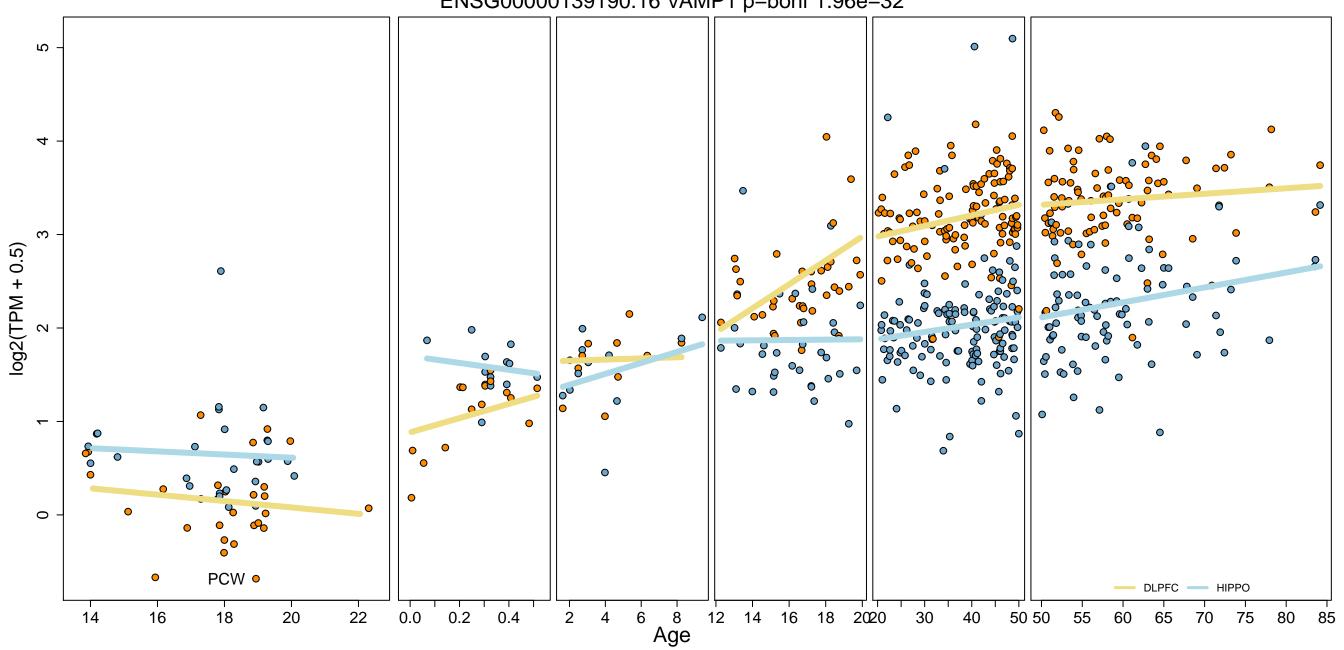
ENSG00000237187.8 NR2F1-AS1 p-bonf 2.42e-33



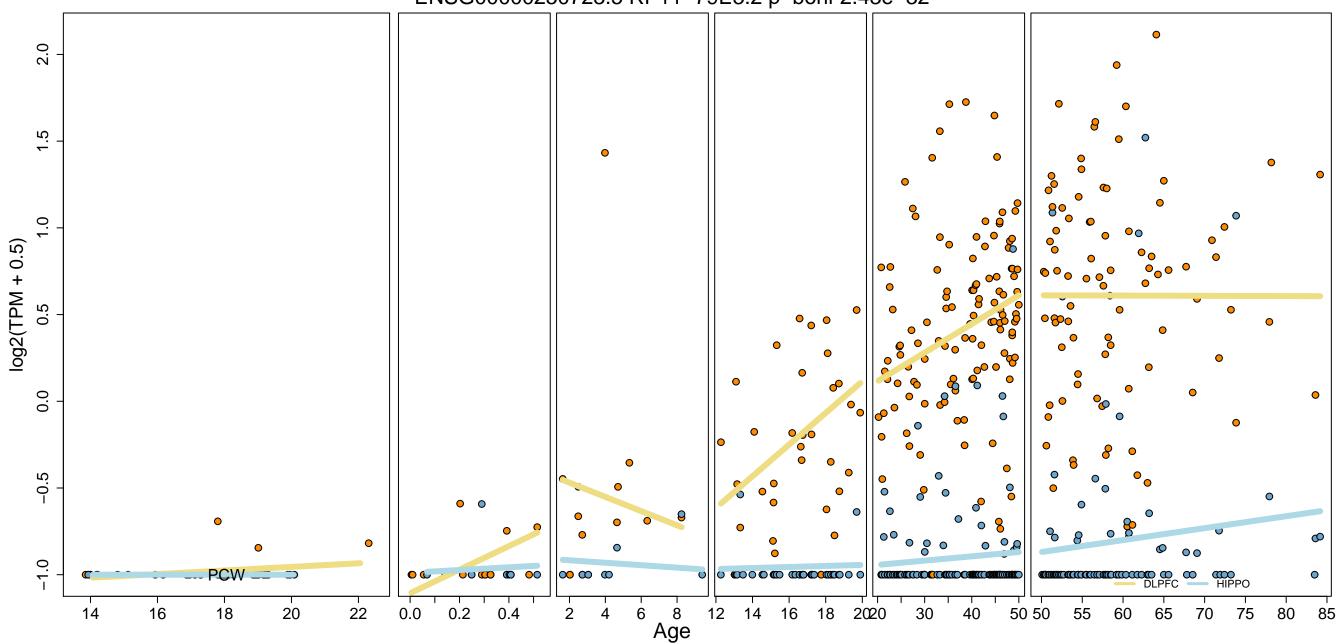
ENSG00000279082.3 RP1-167O22.1 p-bonf 4.3e-33



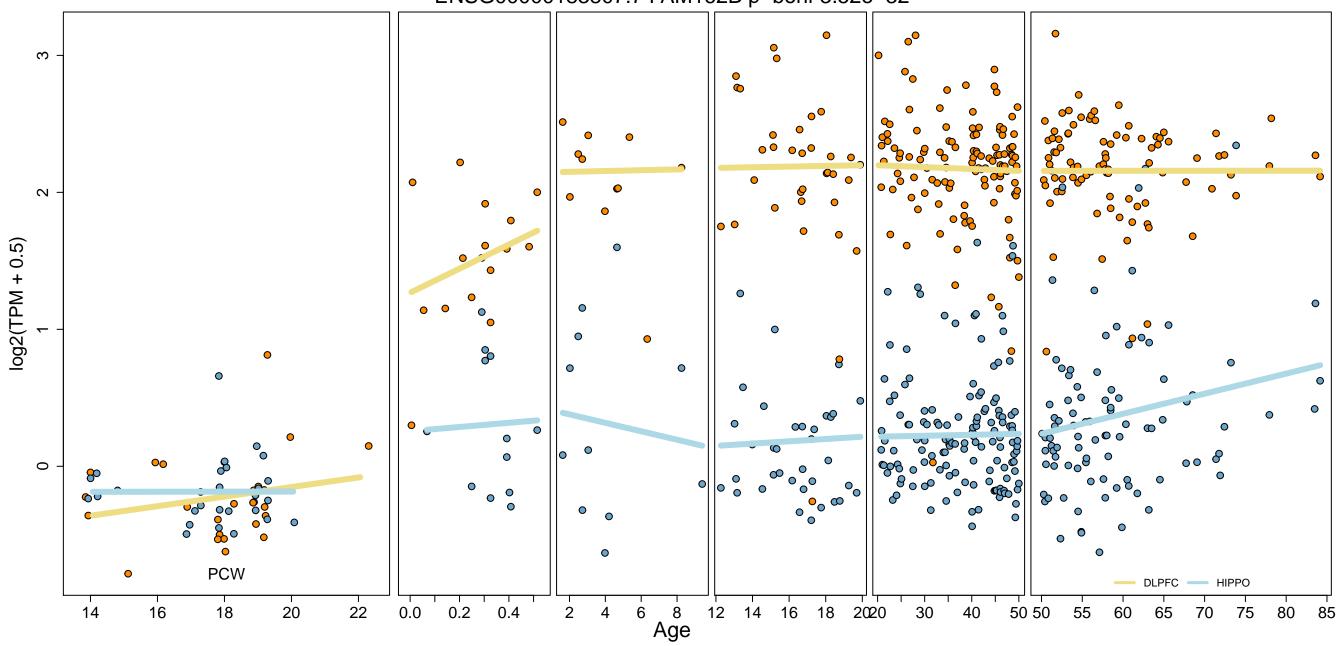
ENSG00000139190.16 VAMP1 p-bonf 1.96e-32



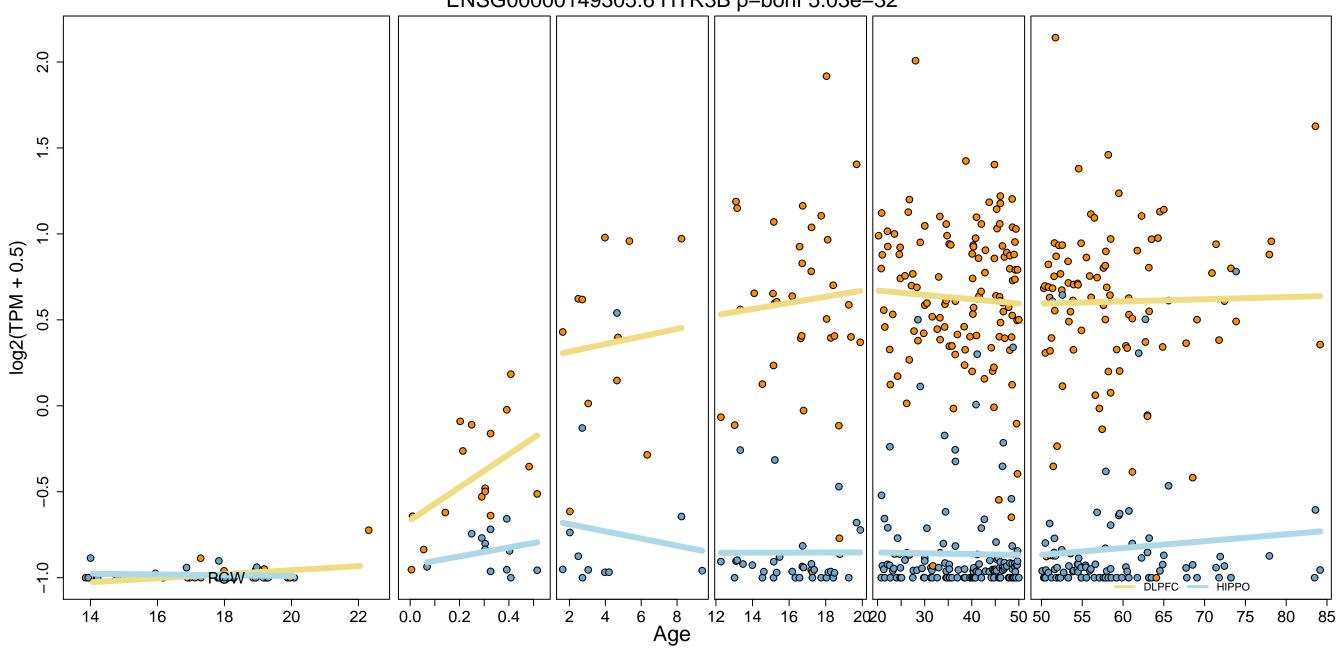
ENSG00000250723.5 RP11-79E3.2 p-bonf 2.43e-32



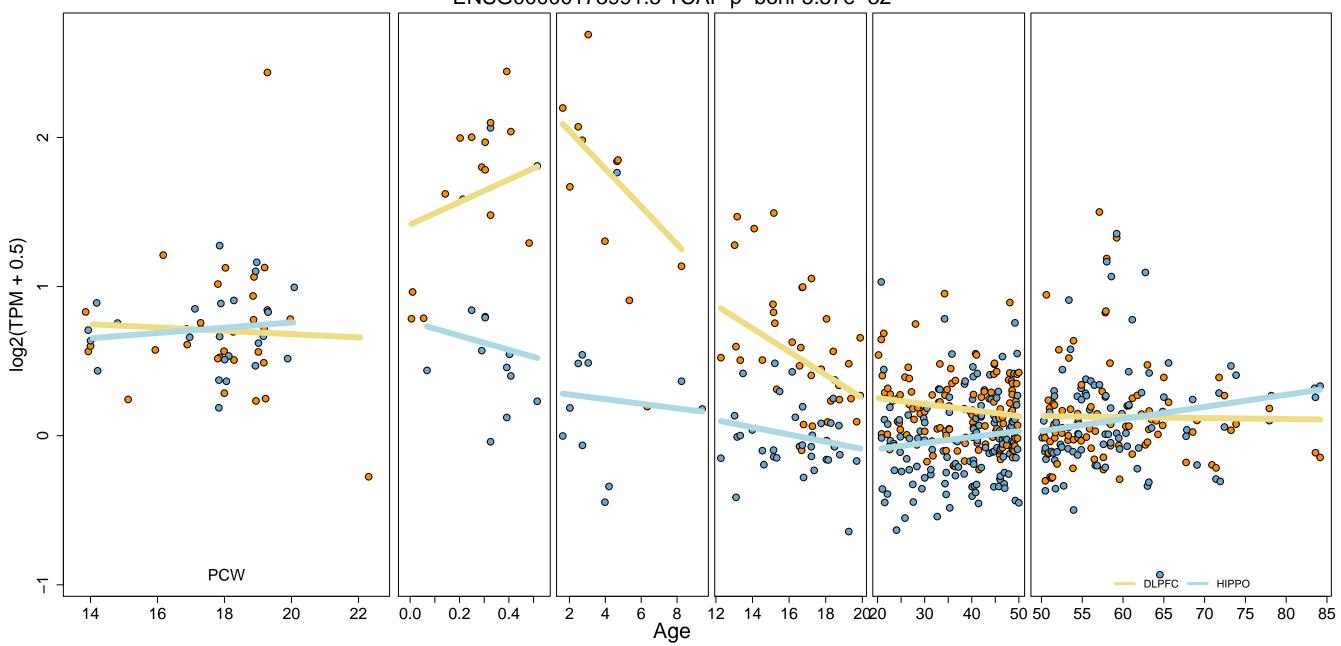
ENSG00000183807.7 FAM162B p-bonf 3.52e-32



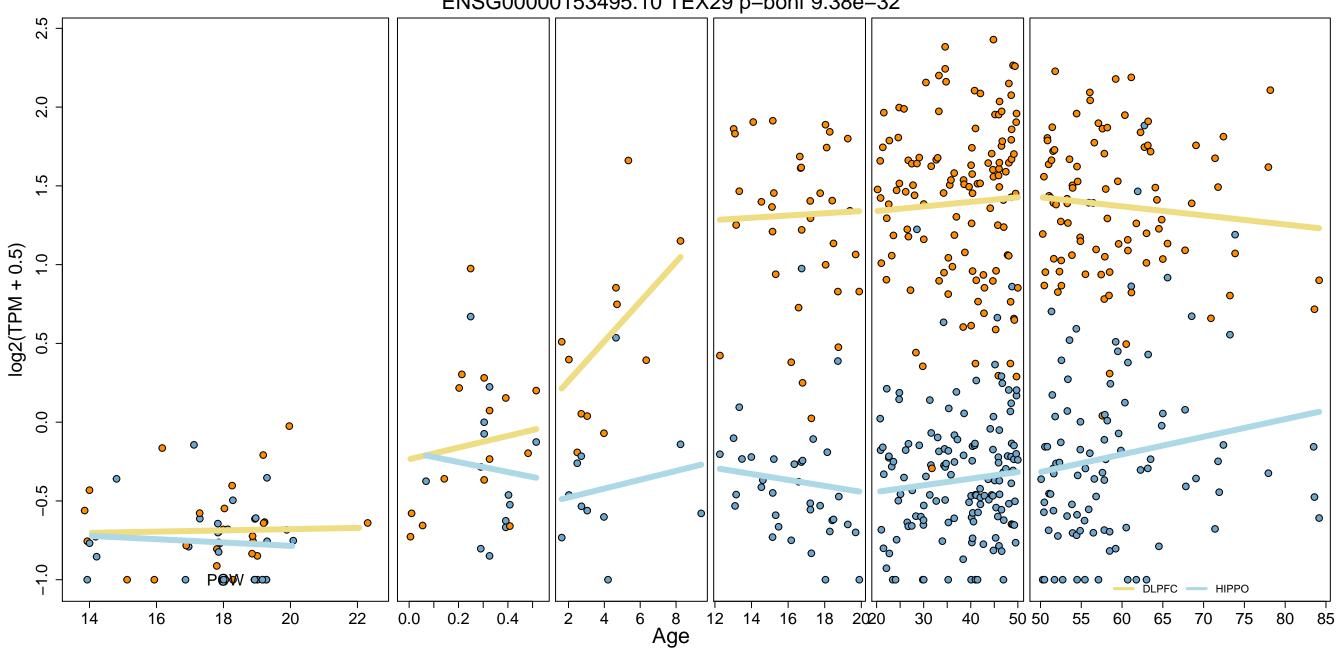
ENSG00000149305.6 HTR3B p-bonf 5.03e-32



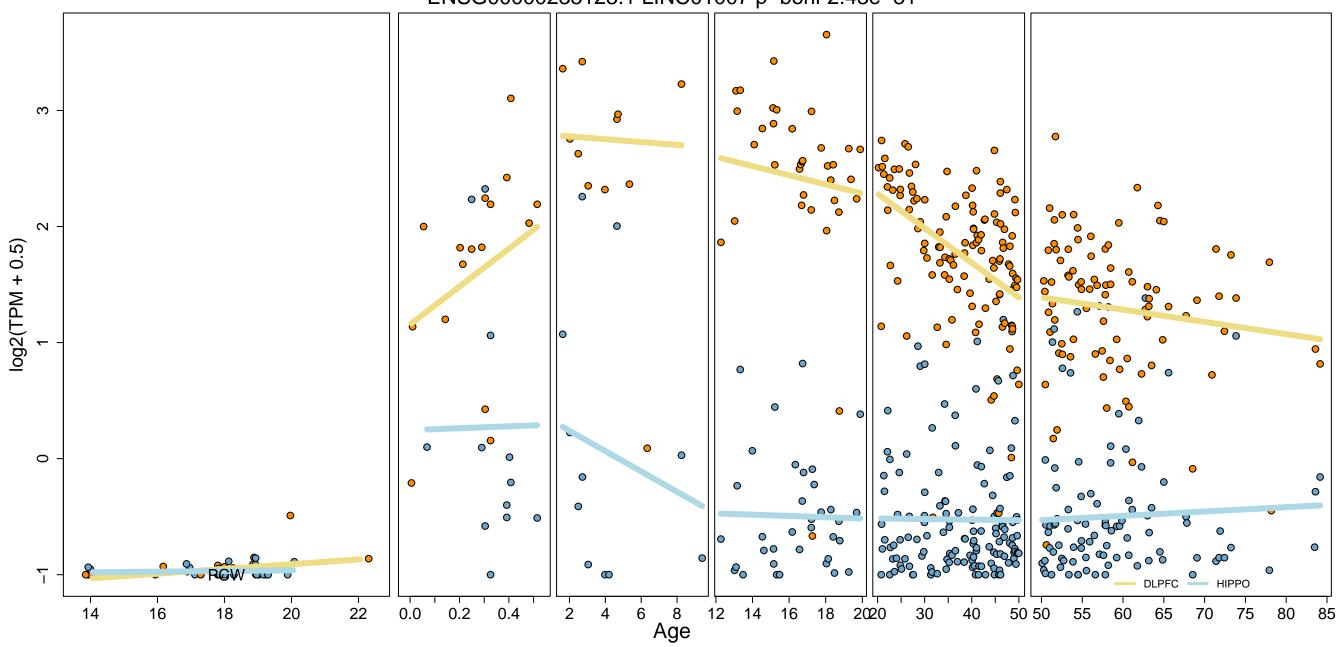
ENSG00000173991.5 TCAP p-bonf 5.57e-32



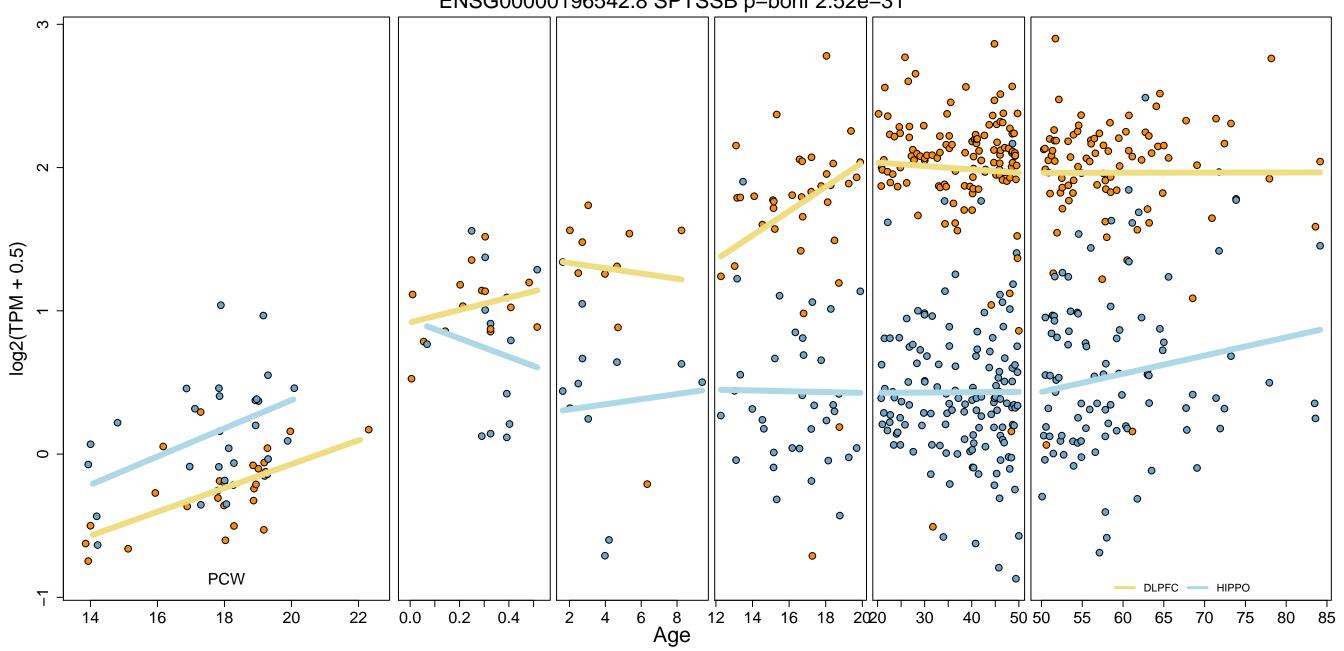
ENSG00000153495.10 TEX29 p-bonf 9.38e-32



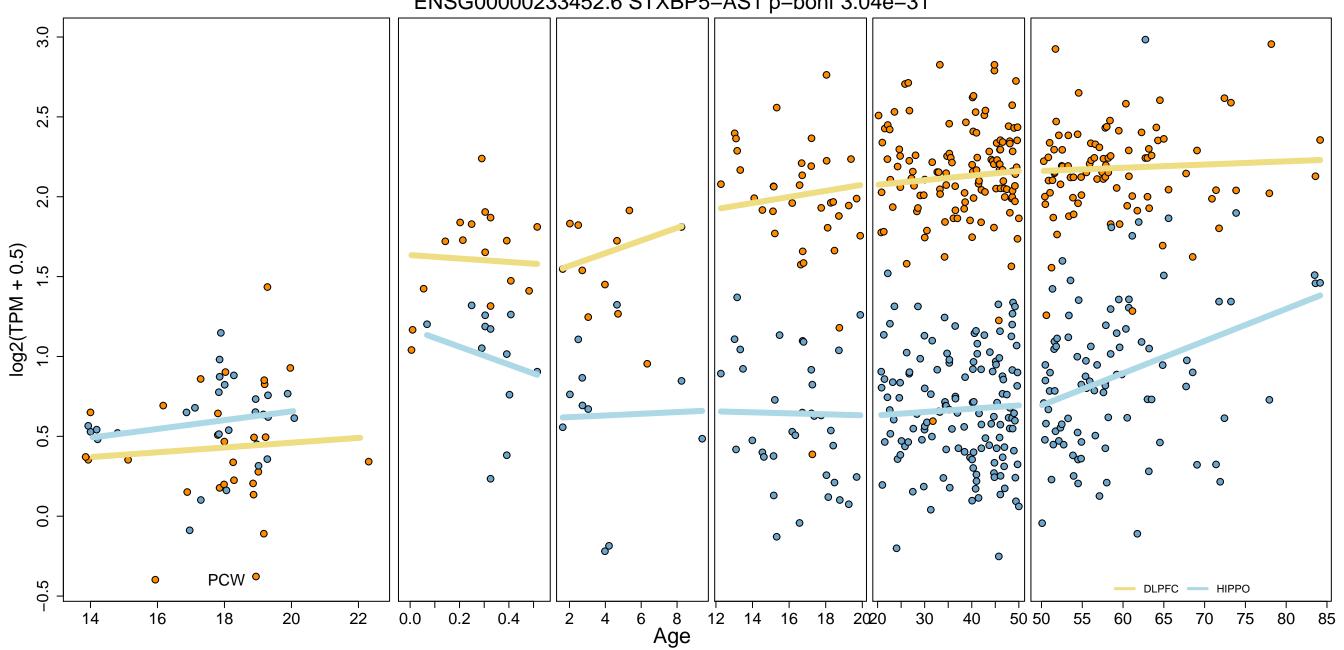
ENSG00000233123.1 LINC01007 p-bonf 2.43e-31



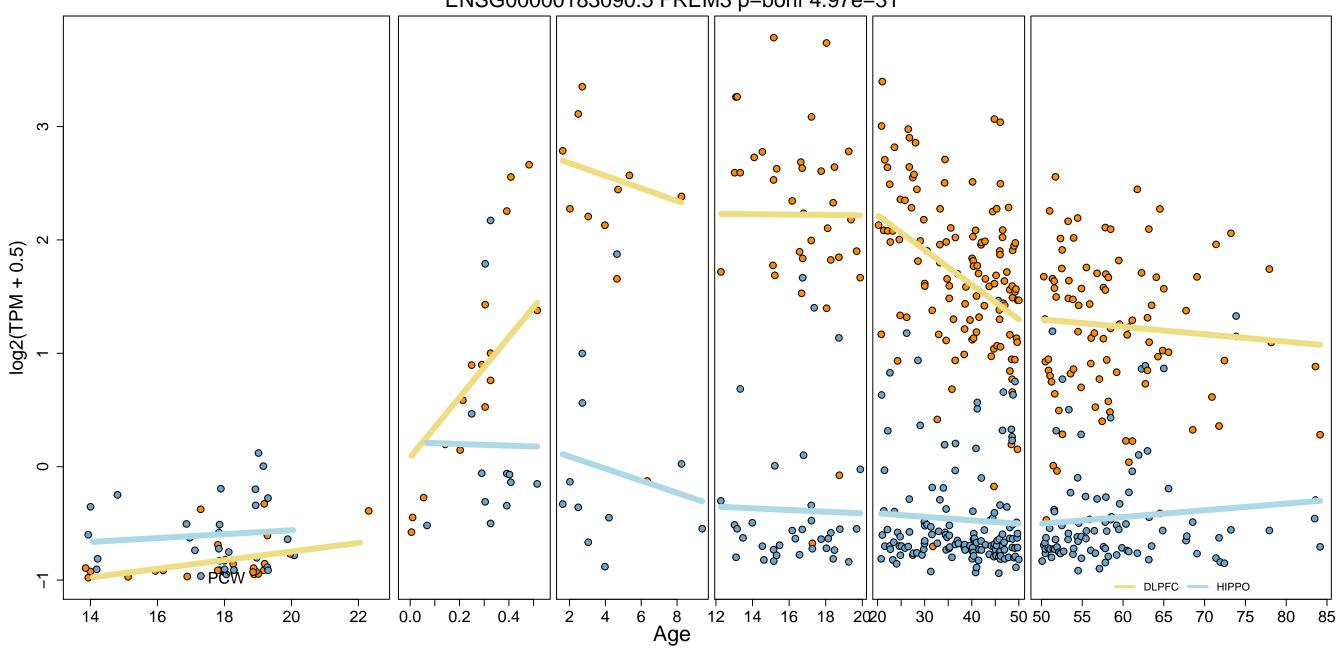
ENSG00000196542.8 SPTSSB p-bonf 2.52e-31



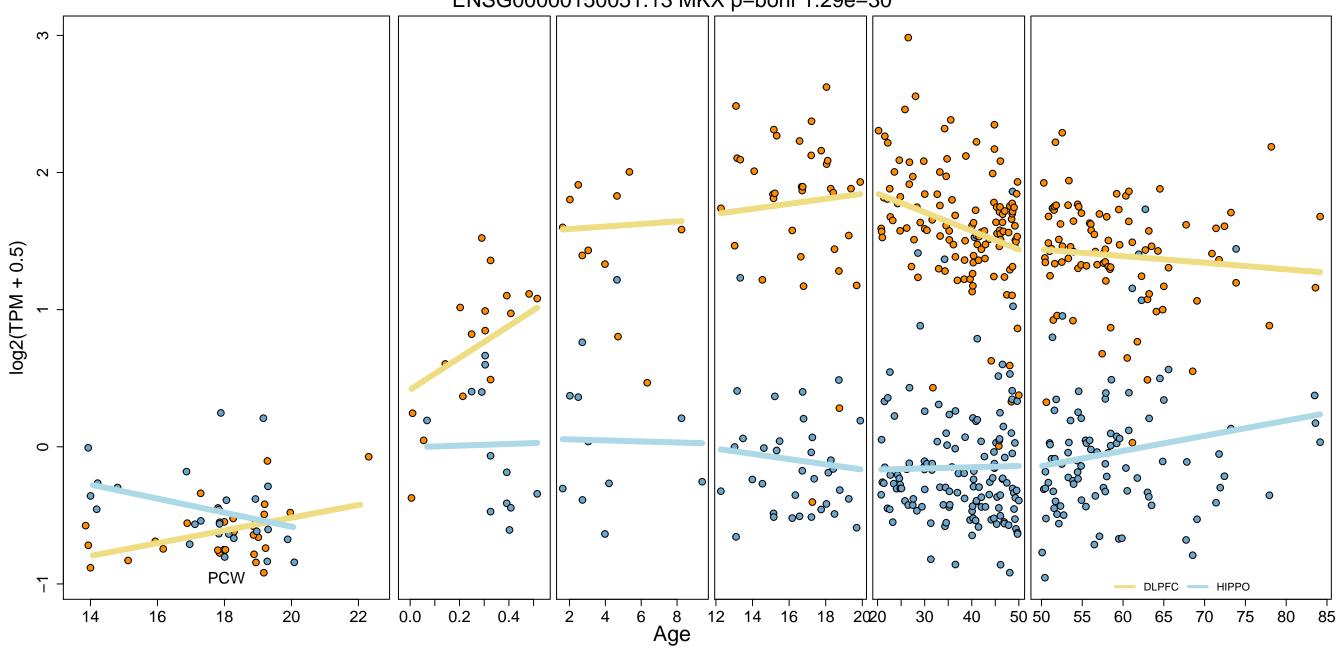
ENSG00000233452.6 STXBP5-AS1 p-bonf 3.04e-31



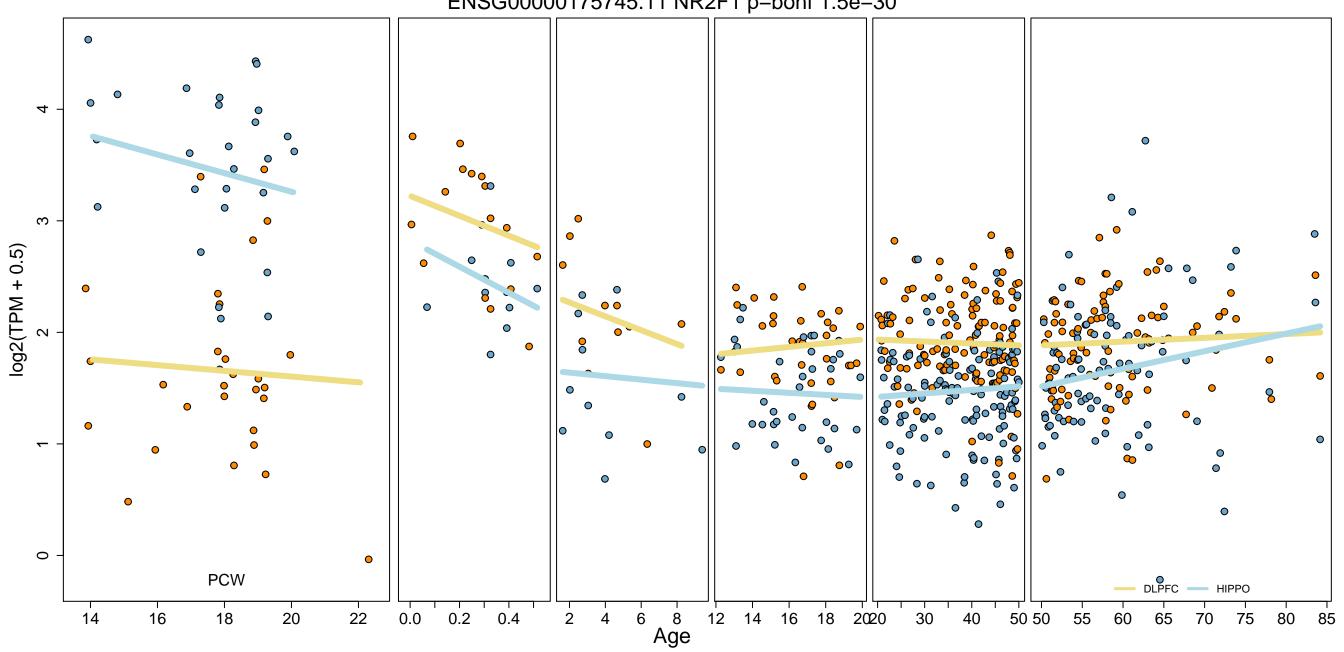
ENSG00000183090.5 FREM3 p-bonf 4.97e-31



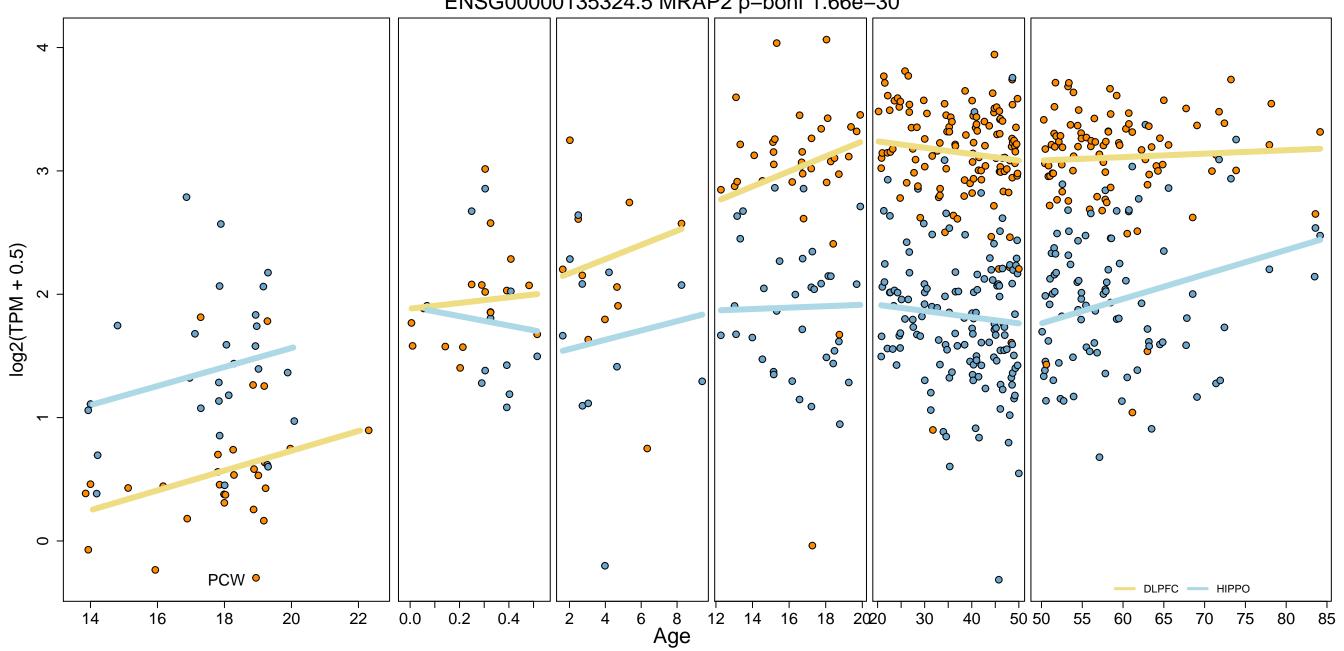
ENSG00000150051.13 MKX p-bonf 1.29e-30



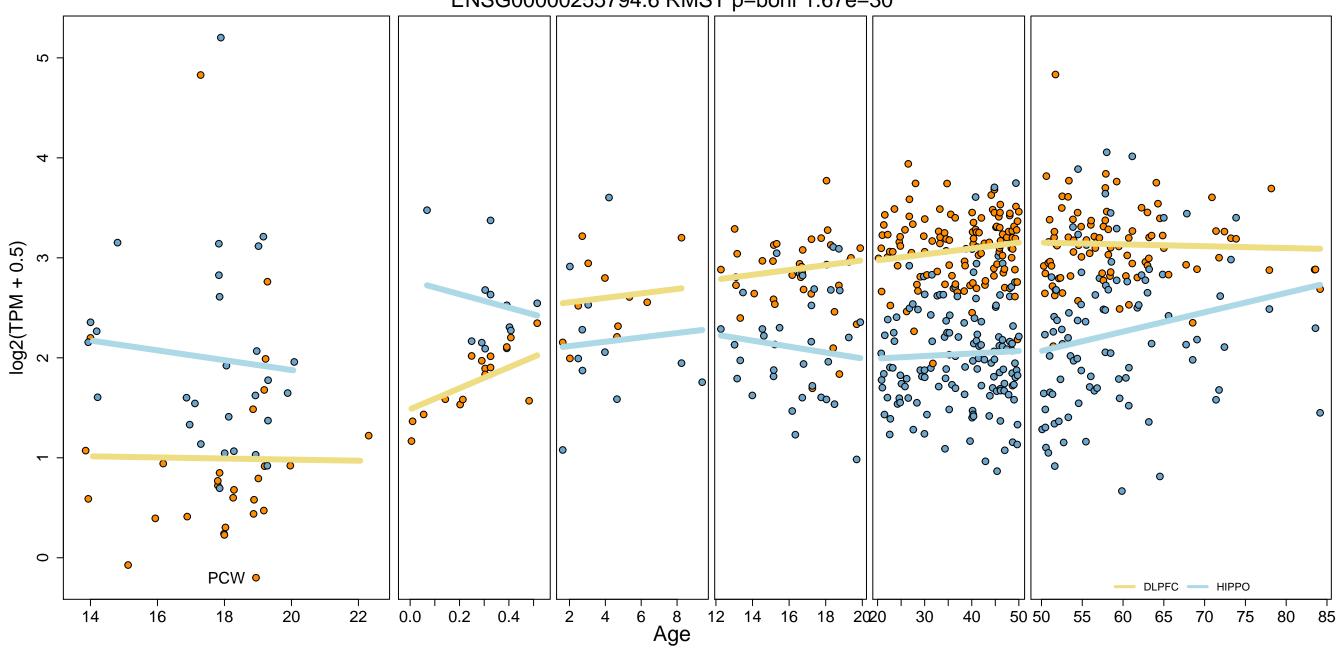
ENSG00000175745.11 NR2F1 p-bonf 1.5e-30



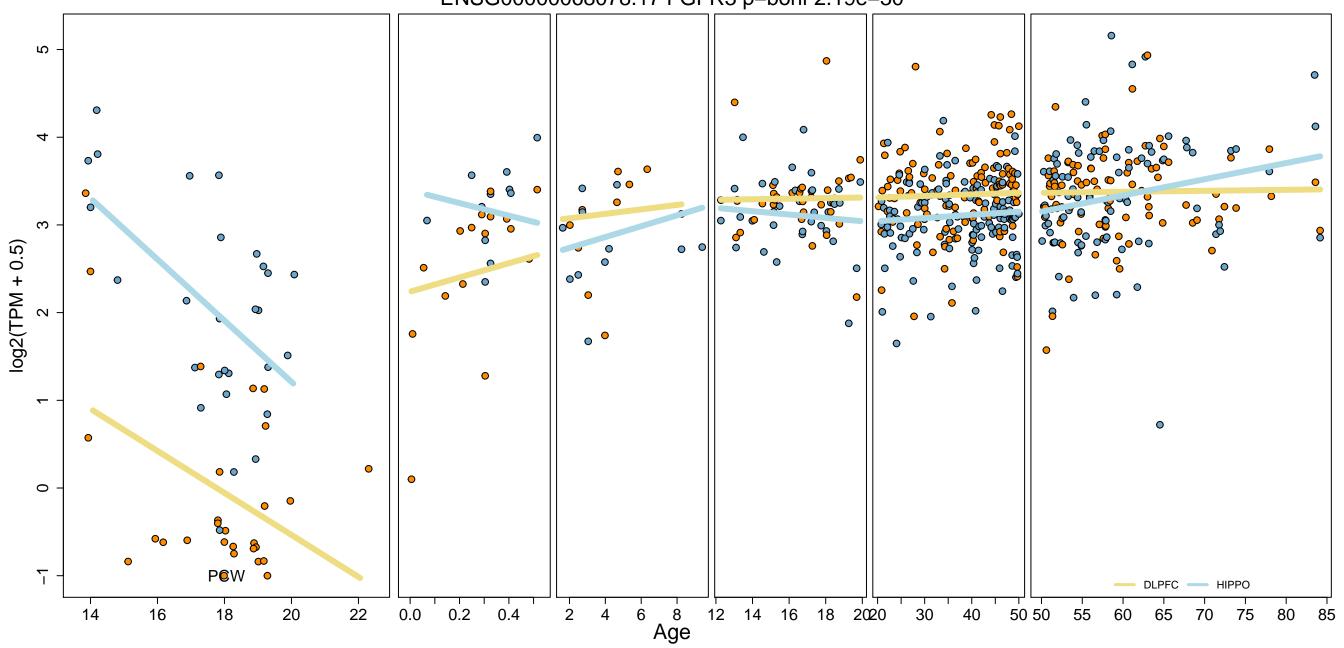
ENSG00000135324.5 MRAP2 p-bonf 1.66e-30



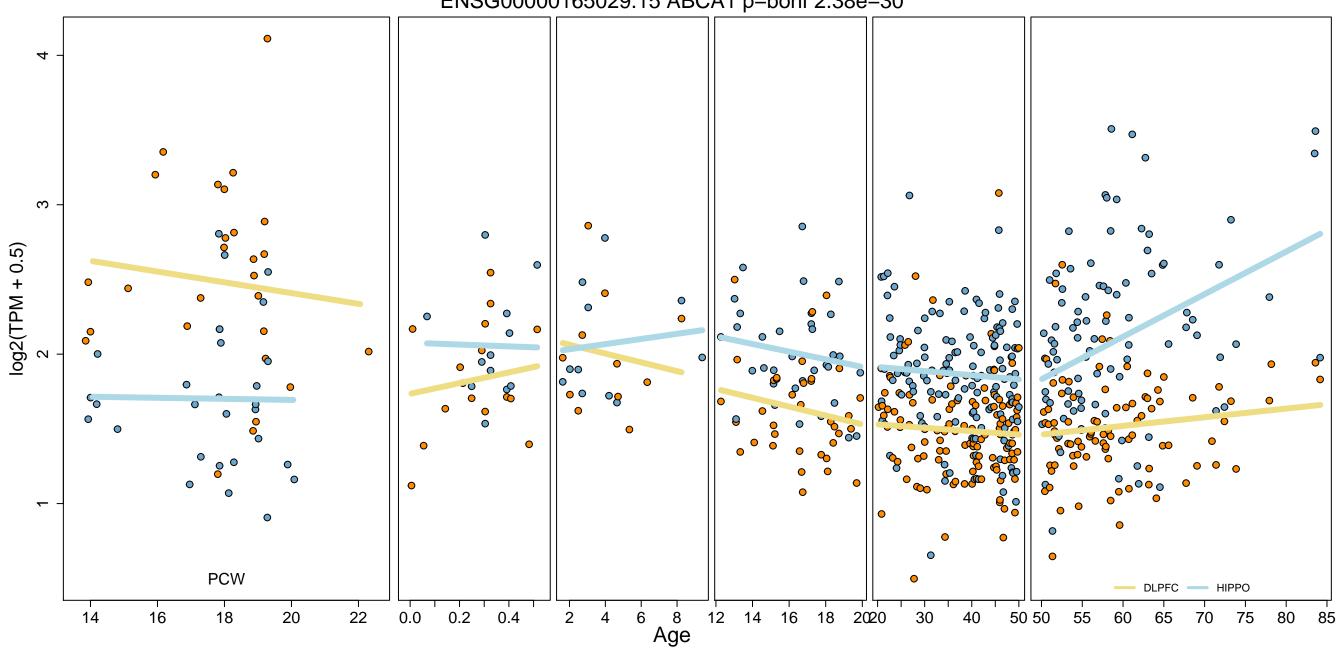
ENSG00000255794.6 RMST p-bonf 1.67e-30



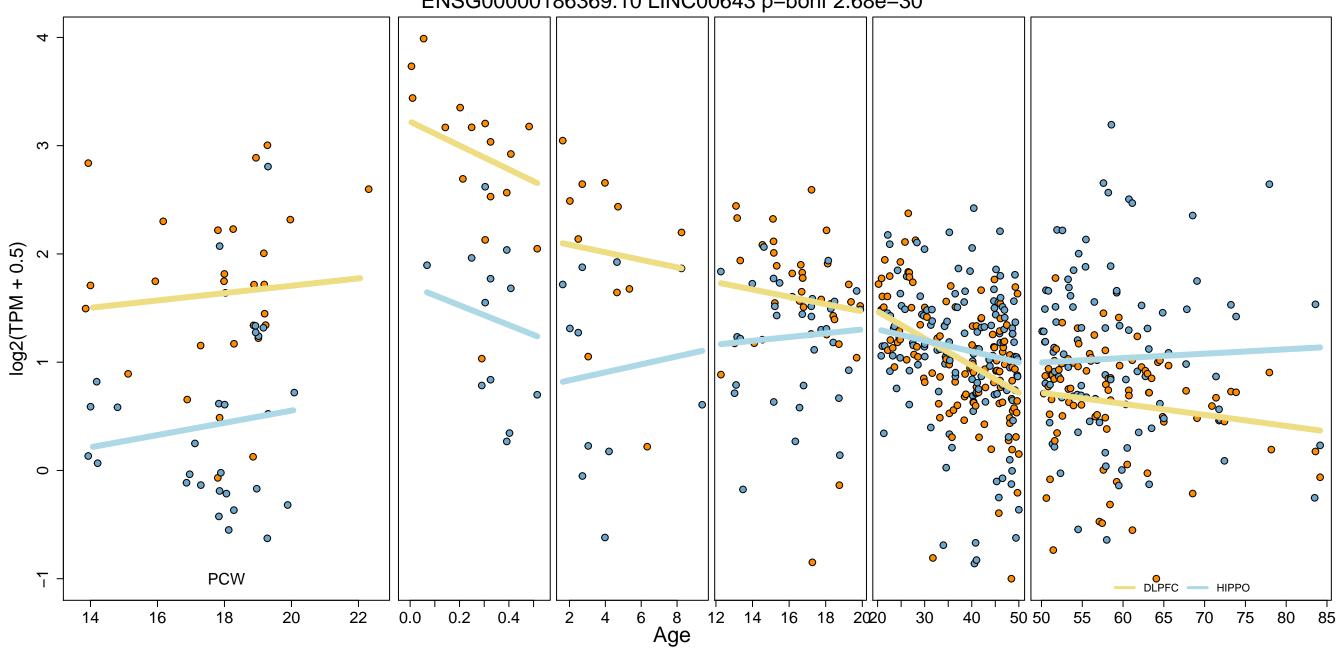
ENSG00000068078.17 FGFR3 p-bonf 2.19e-30



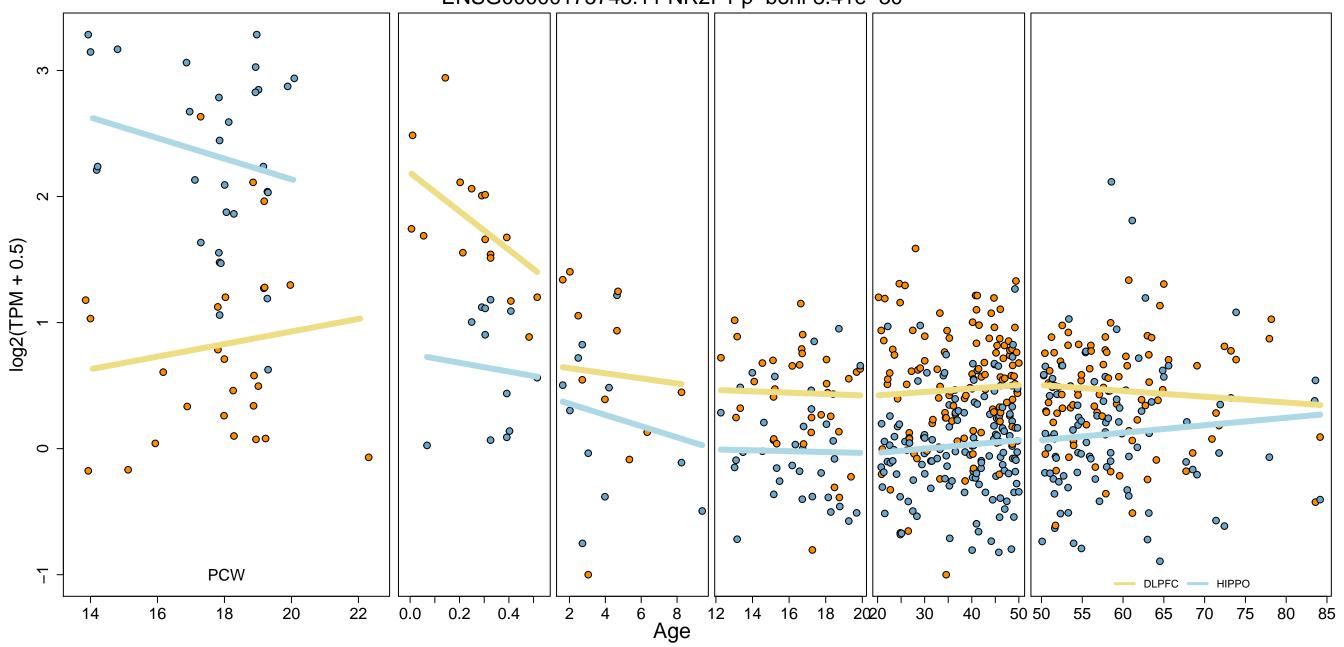
ENSG00000165029.15 ABCA1 p-bonf 2.38e-30



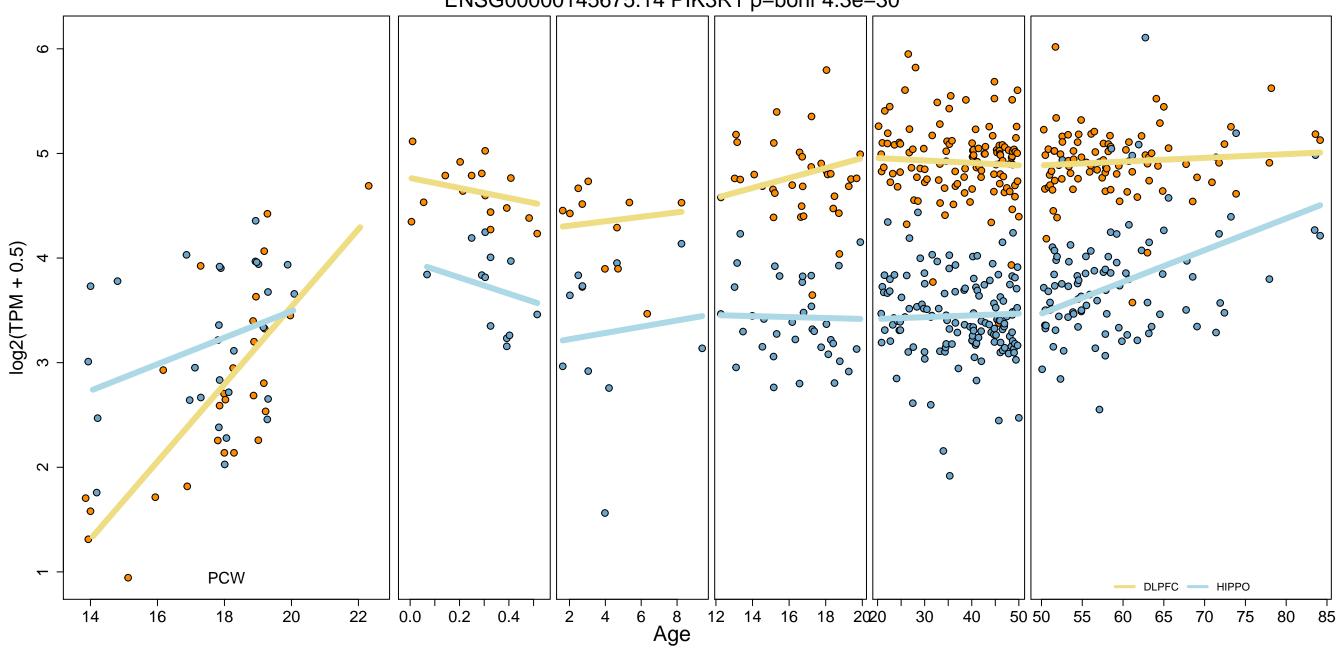
ENSG00000186369.10 LINC00643 p-bonf 2.68e-30



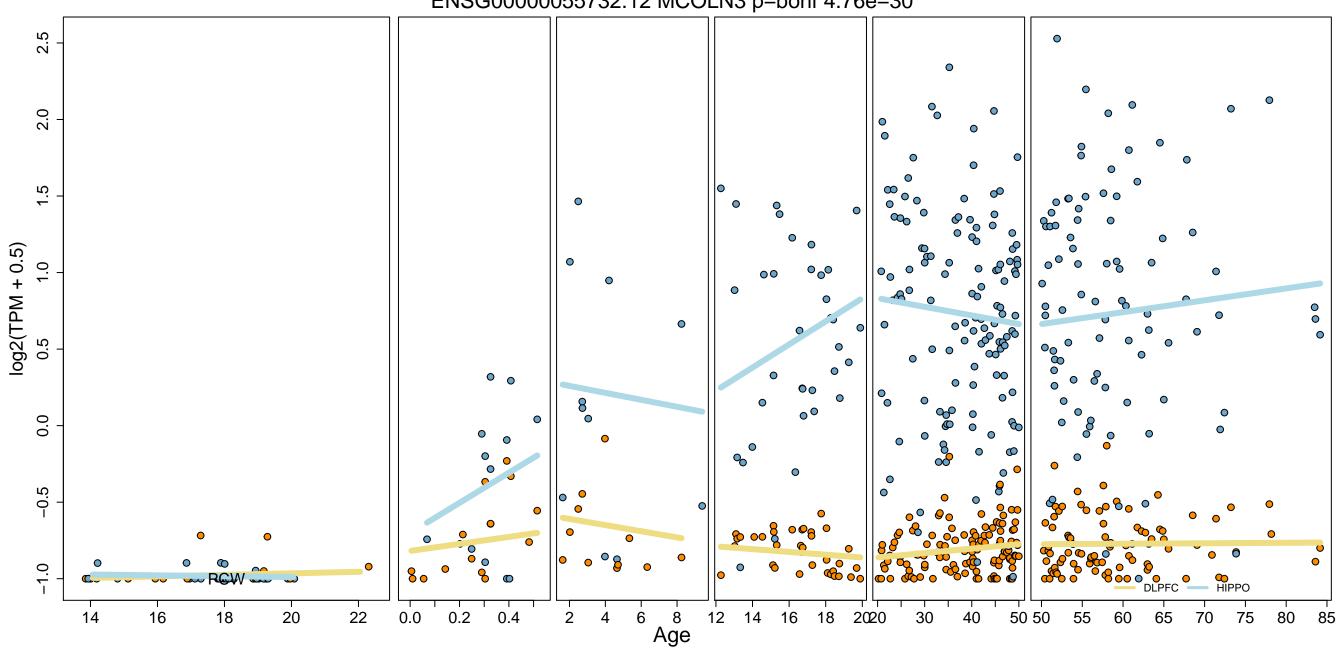
ENSG00000175745.11 NR2F1 p-bonf 3.41e-30



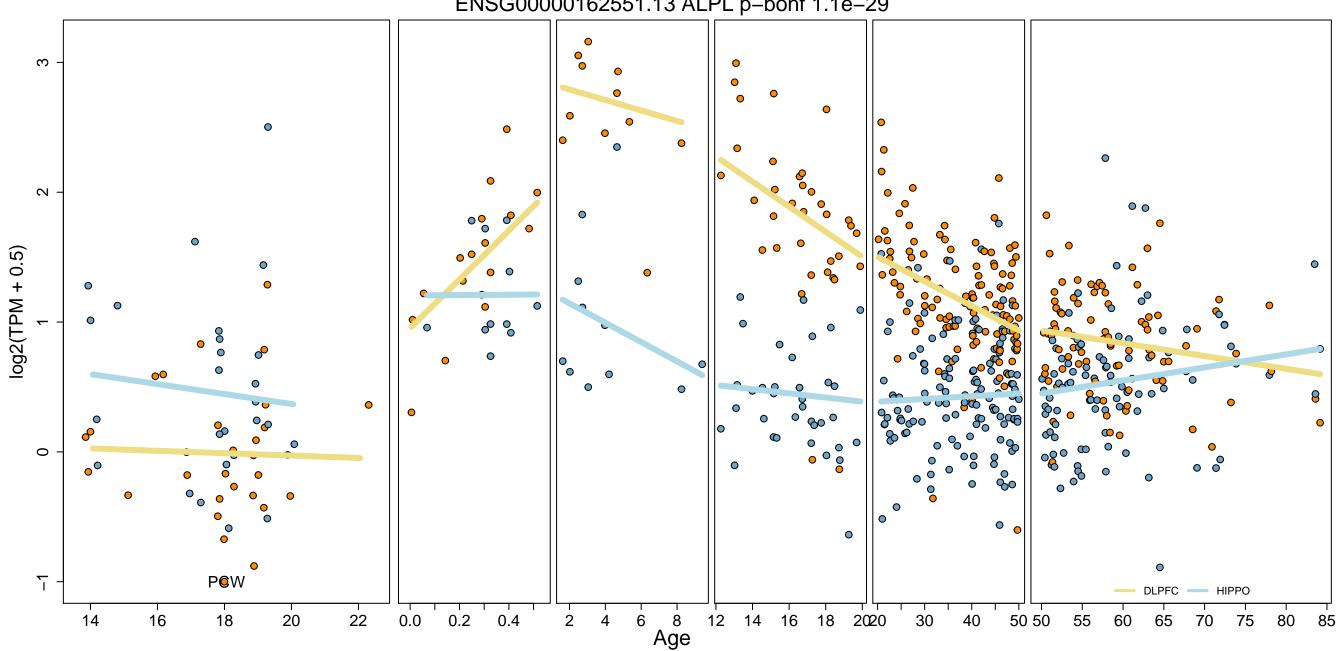
ENSG00000145675.14 PIK3R1 p-bonf 4.3e-30



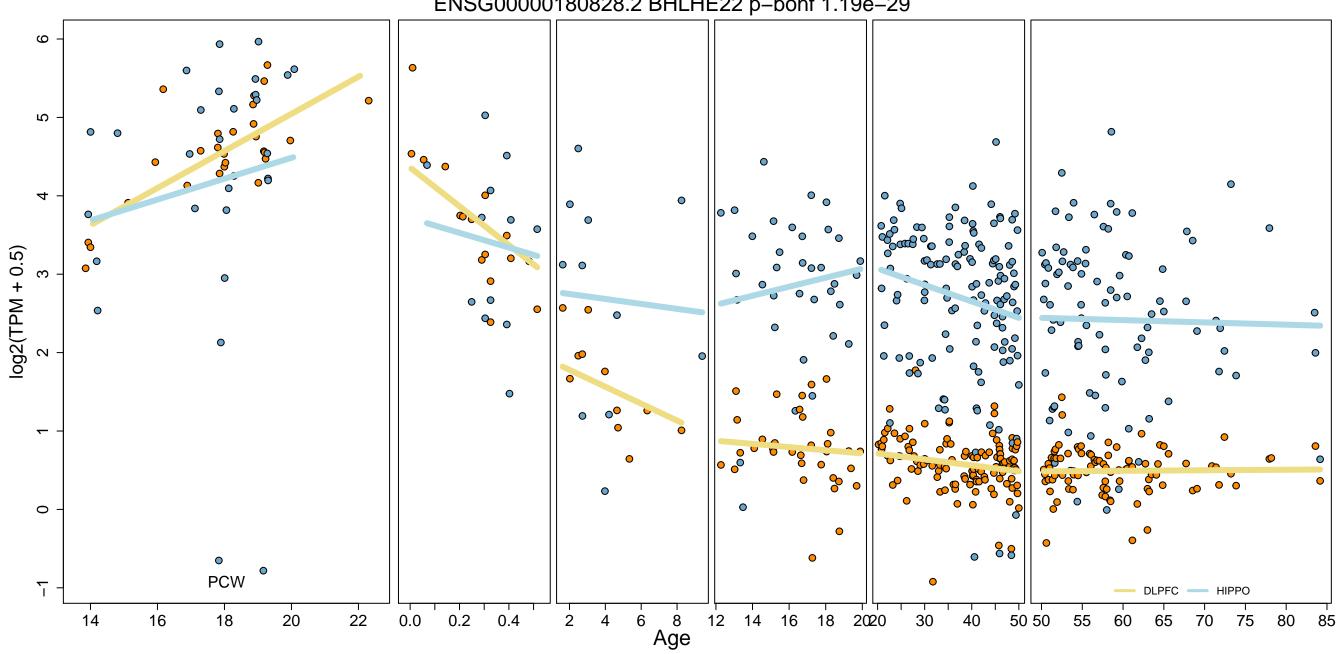
ENSG00000055732.12 MCOLN3 p-bonf 4.76e-30



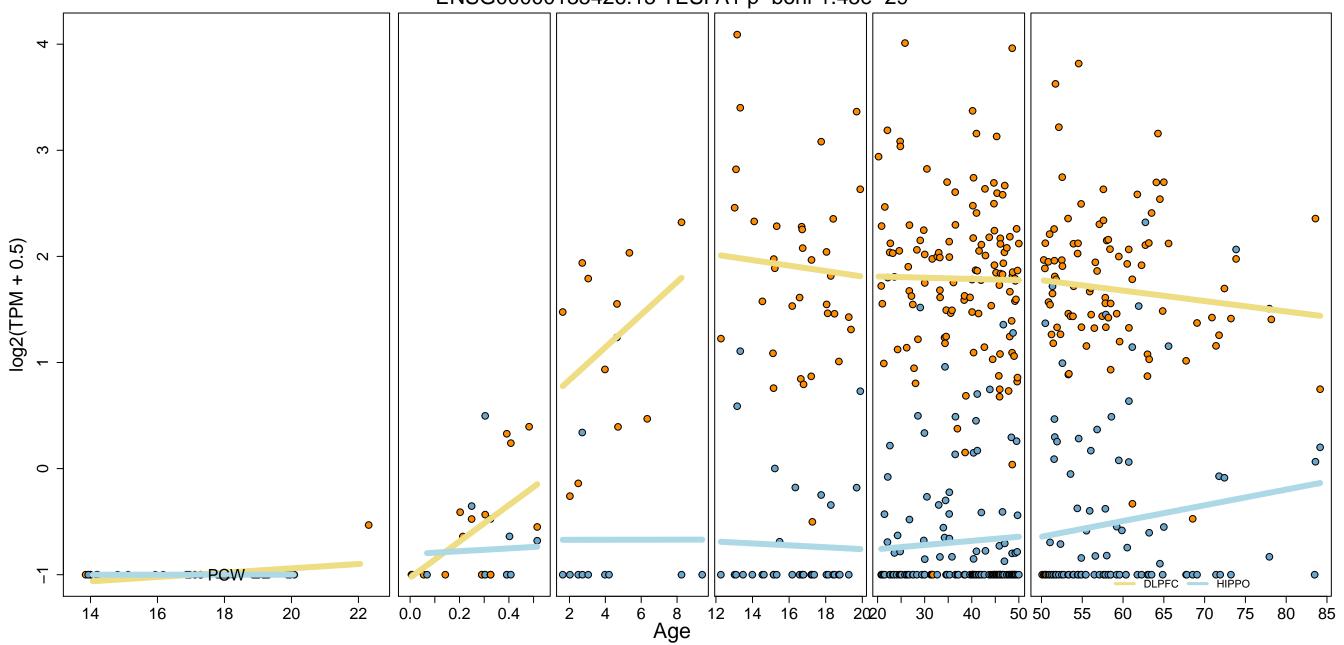
ENSG00000162551.13 ALPL p-bonf 1.1e-29



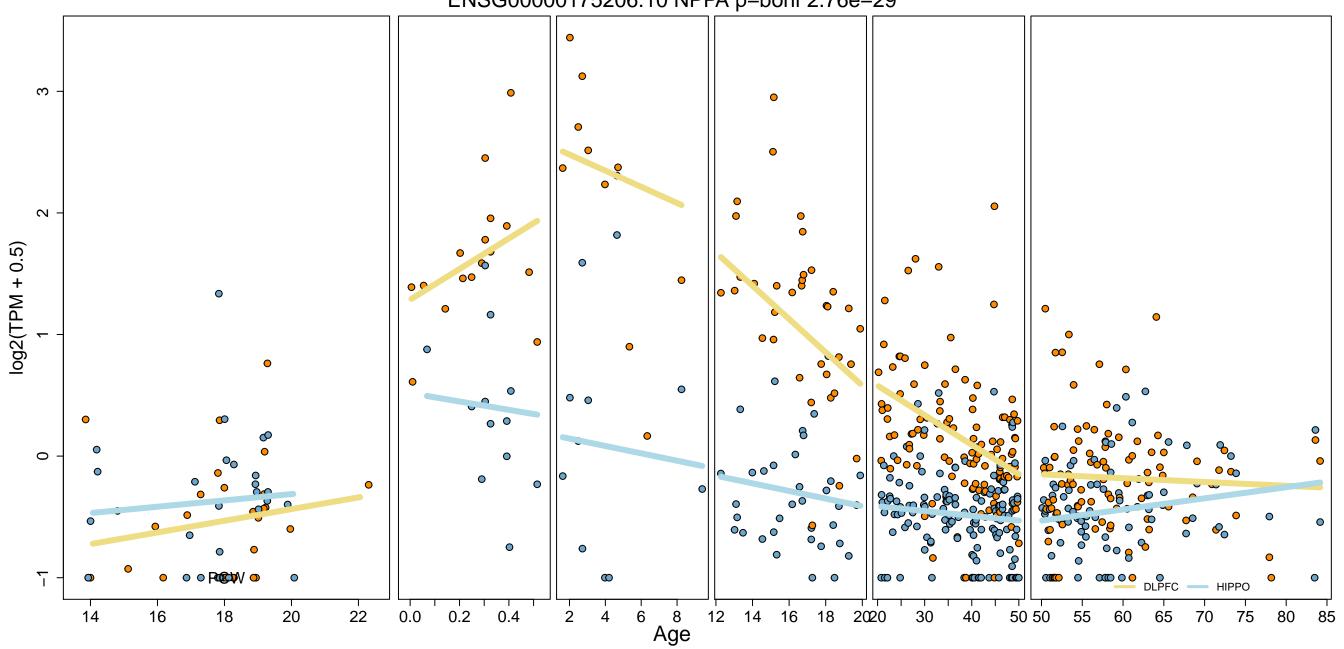
ENSG00000180828.2 BHLHE22 p-bonf 1.19e-29



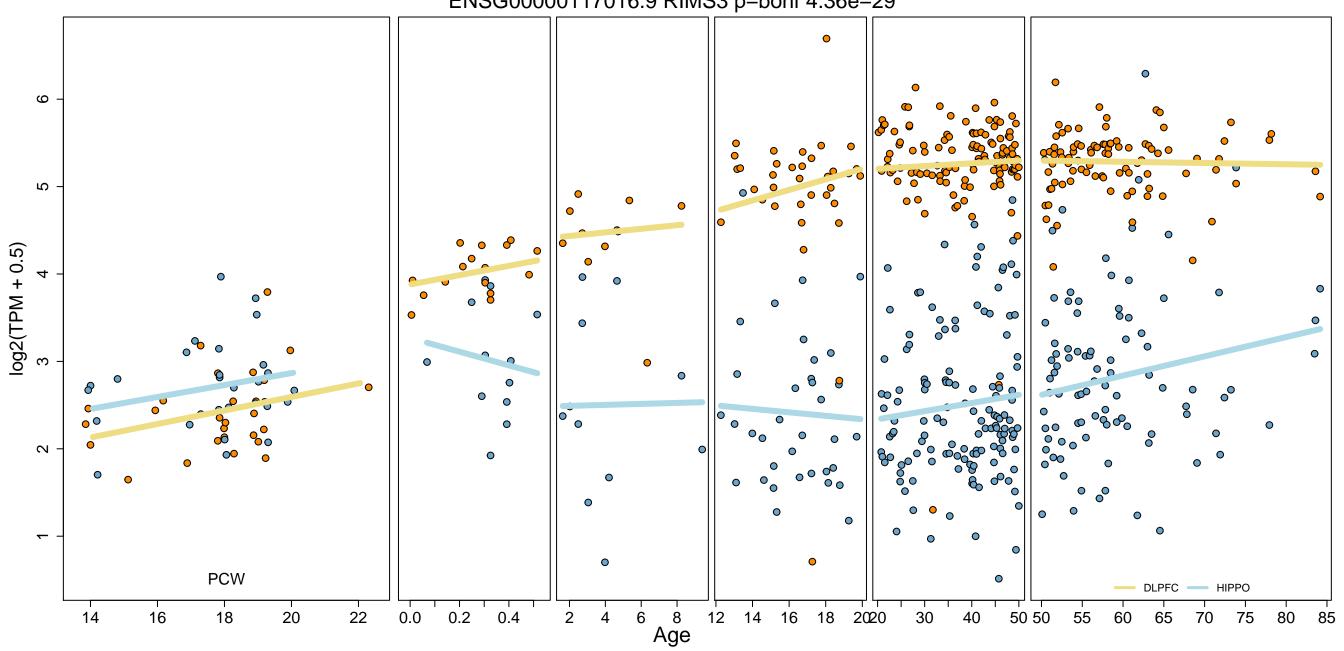
ENSG00000135426.15 TESPA1 p-bonf 1.45e-29



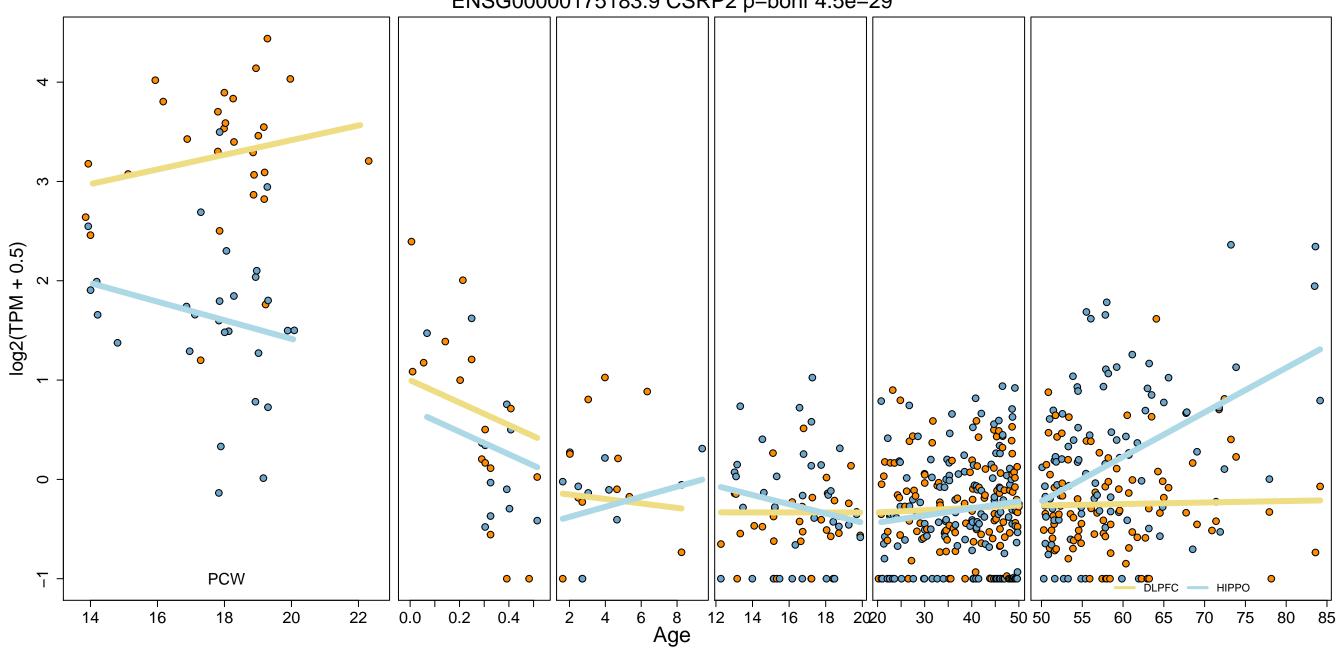
ENSG00000175206.10 NPPA p-bonf 2.76e-29



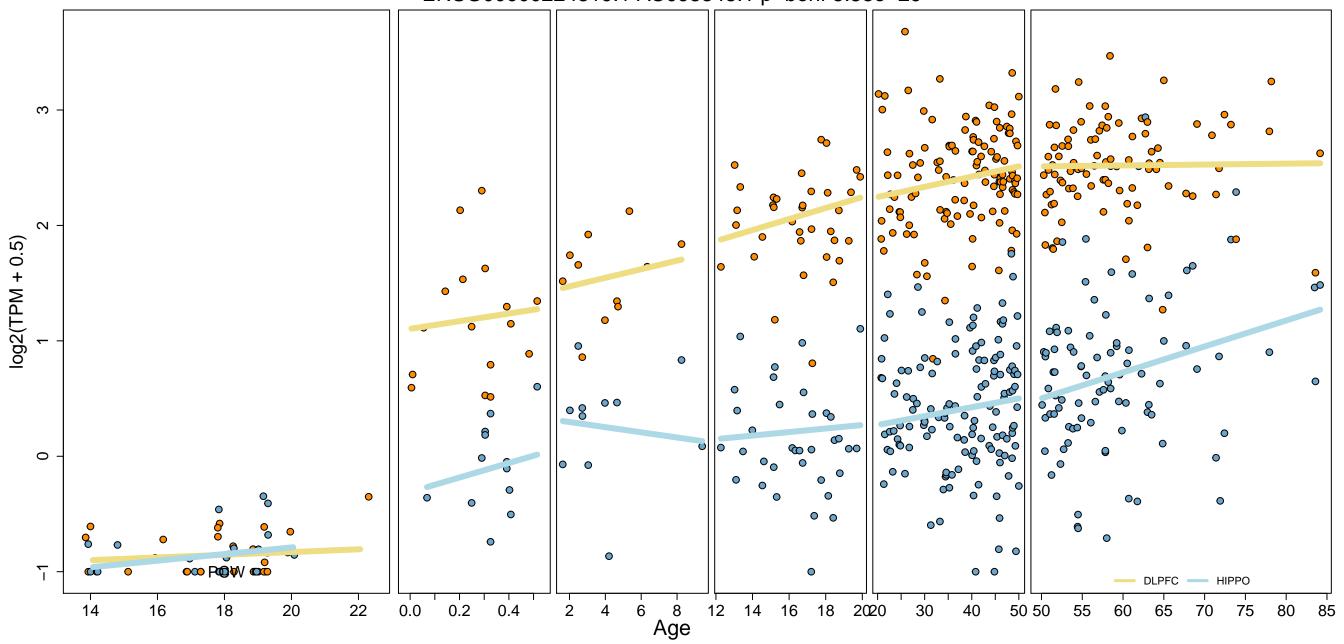
ENSG00000117016.9 RIMS3 p-bonf 4.36e-29



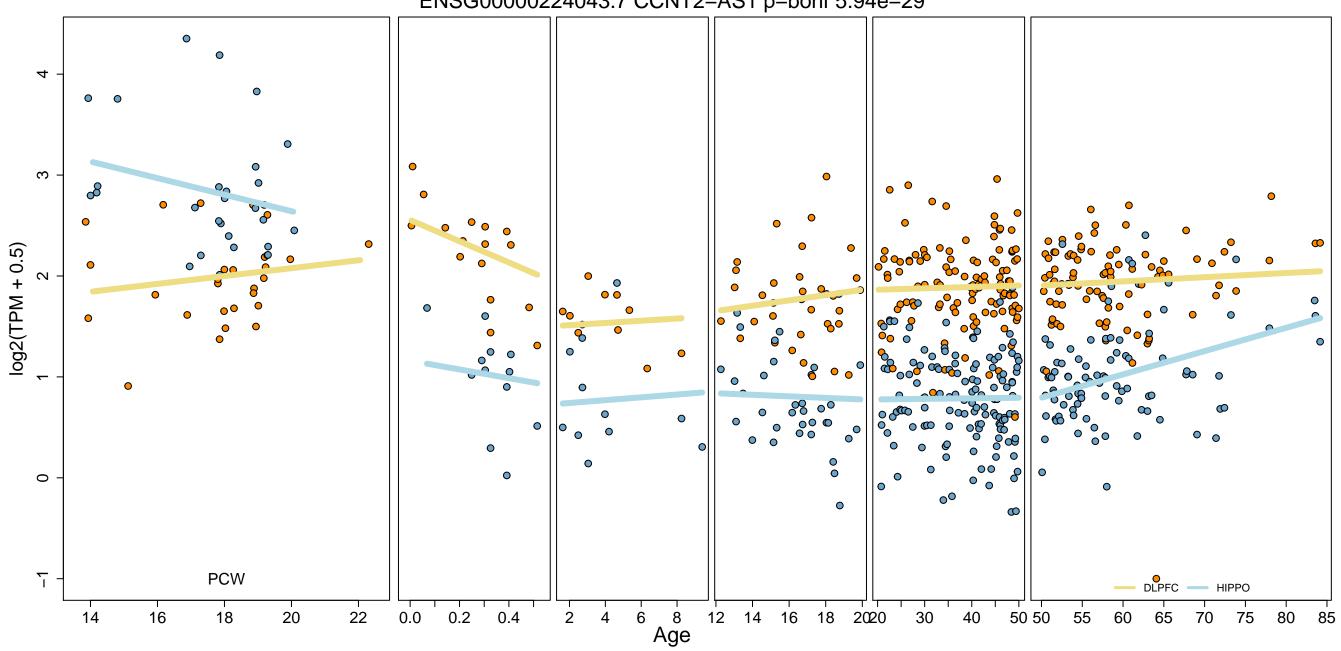
ENSG00000175183.9 CSRP2 p-bonf 4.5e-29



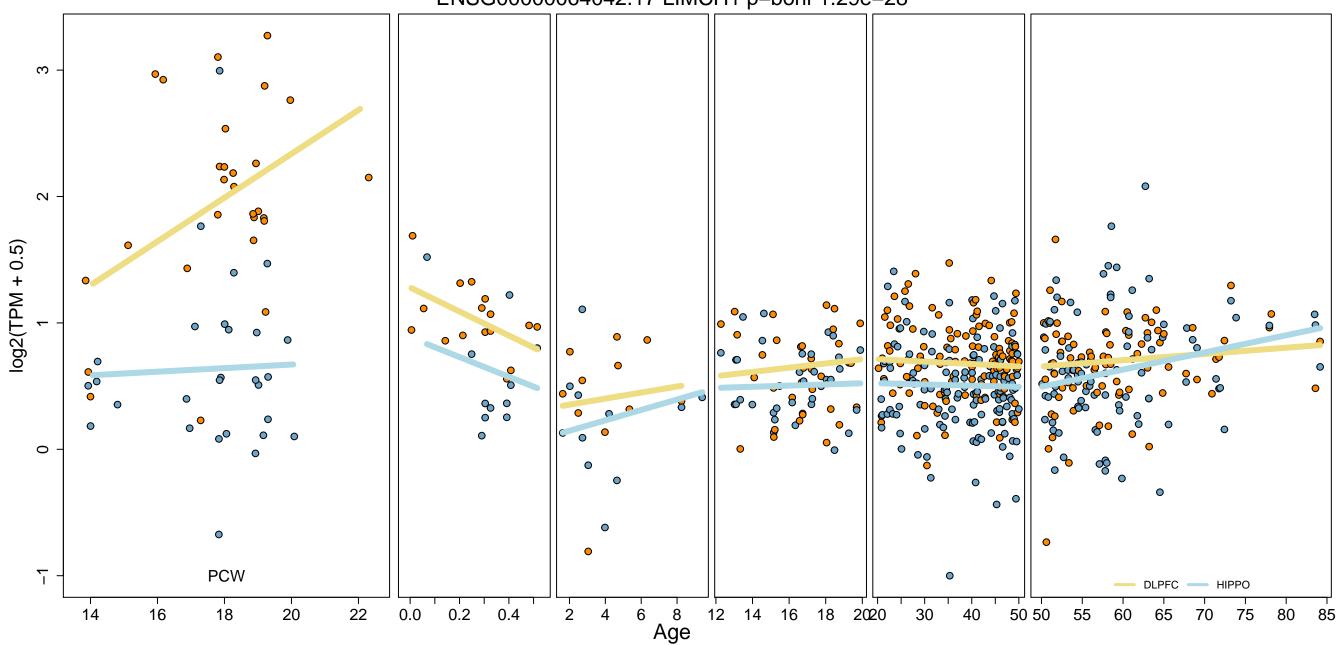
ENSG00000224819.1 AC093843.1 p-bonf 5.58e-29



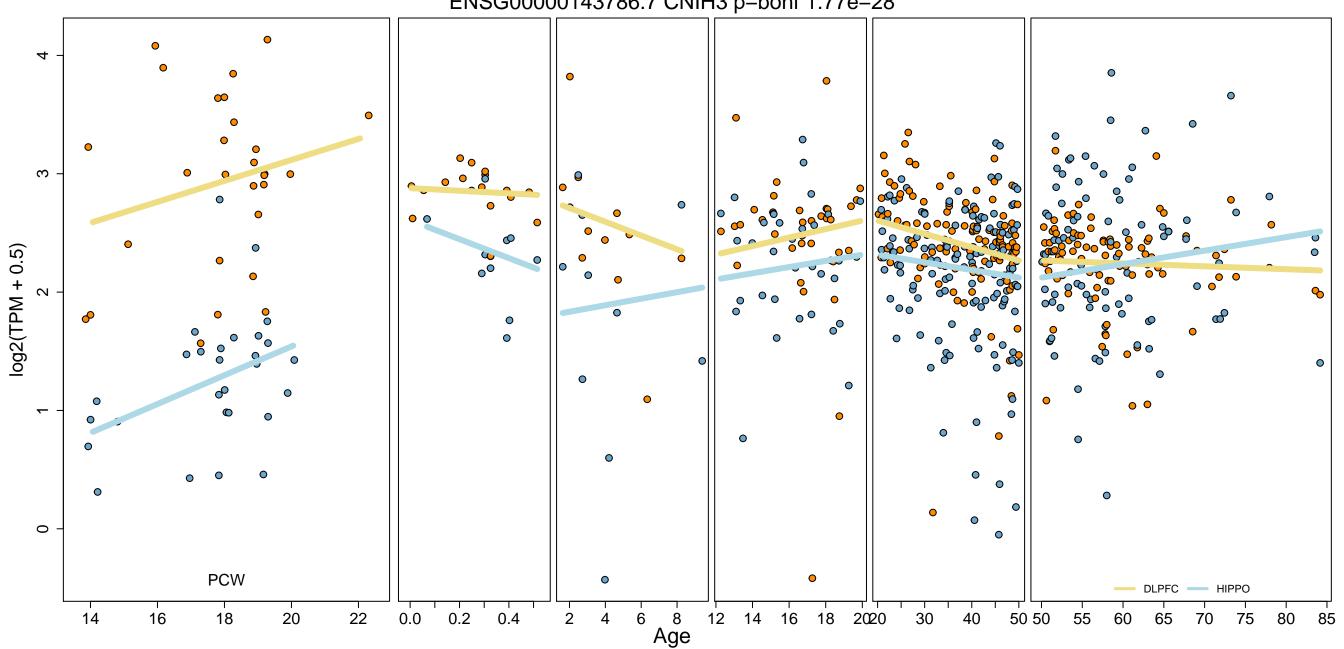
ENSG00000224043.7 CCNT2-AS1 p-bonf 5.94e-29



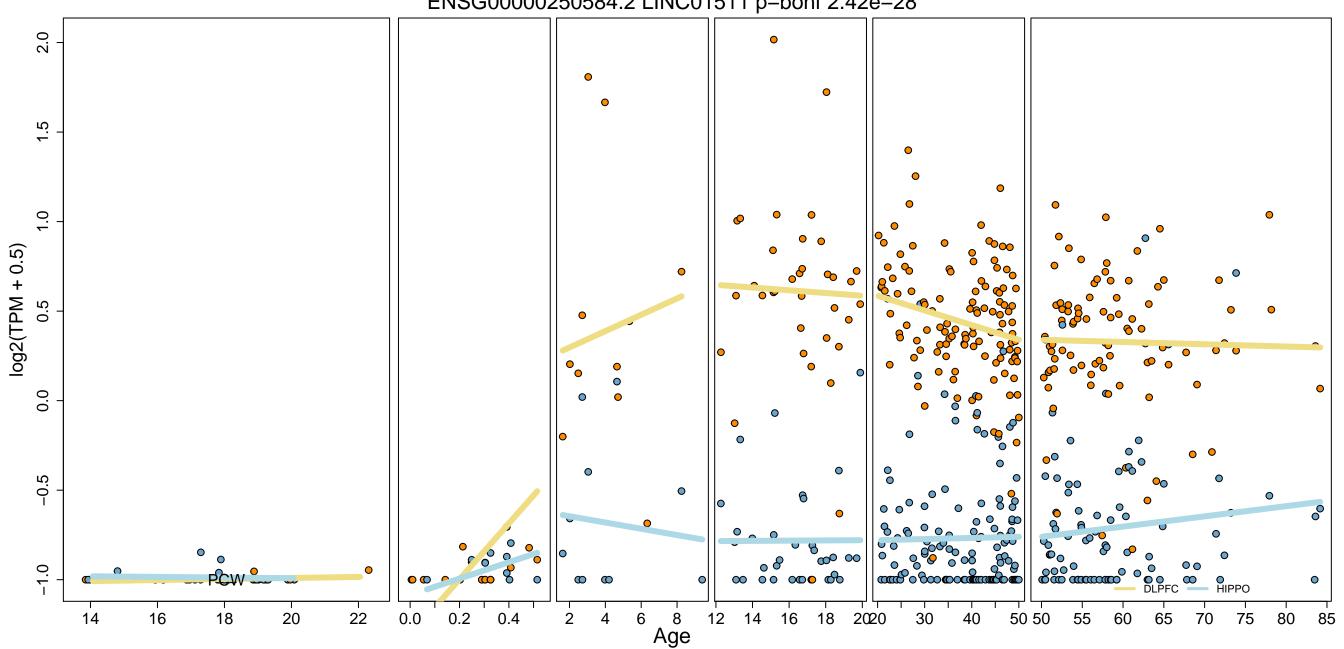
ENSG00000064042.17 LIMCH1 p-bonf 1.29e-28



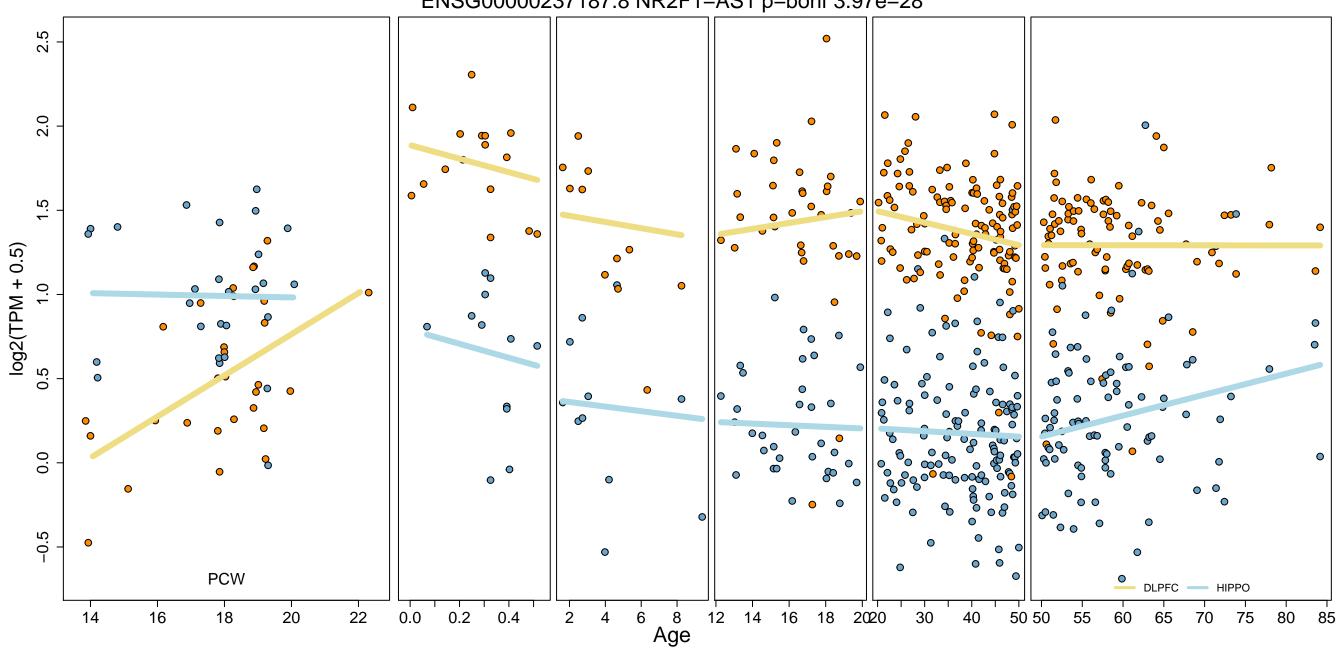
ENSG00000143786.7 CNIH3 p-bonf 1.77e-28



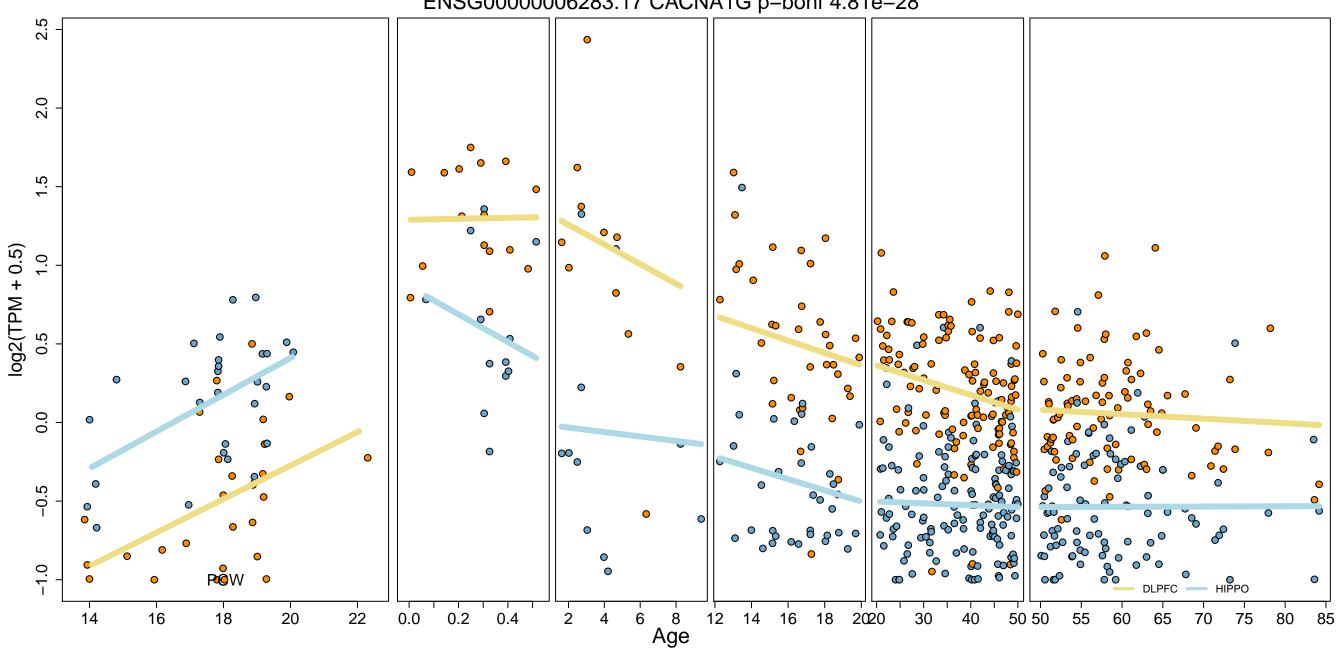
ENSG00000250584.2 LINC01511 p-bonf 2.42e-28



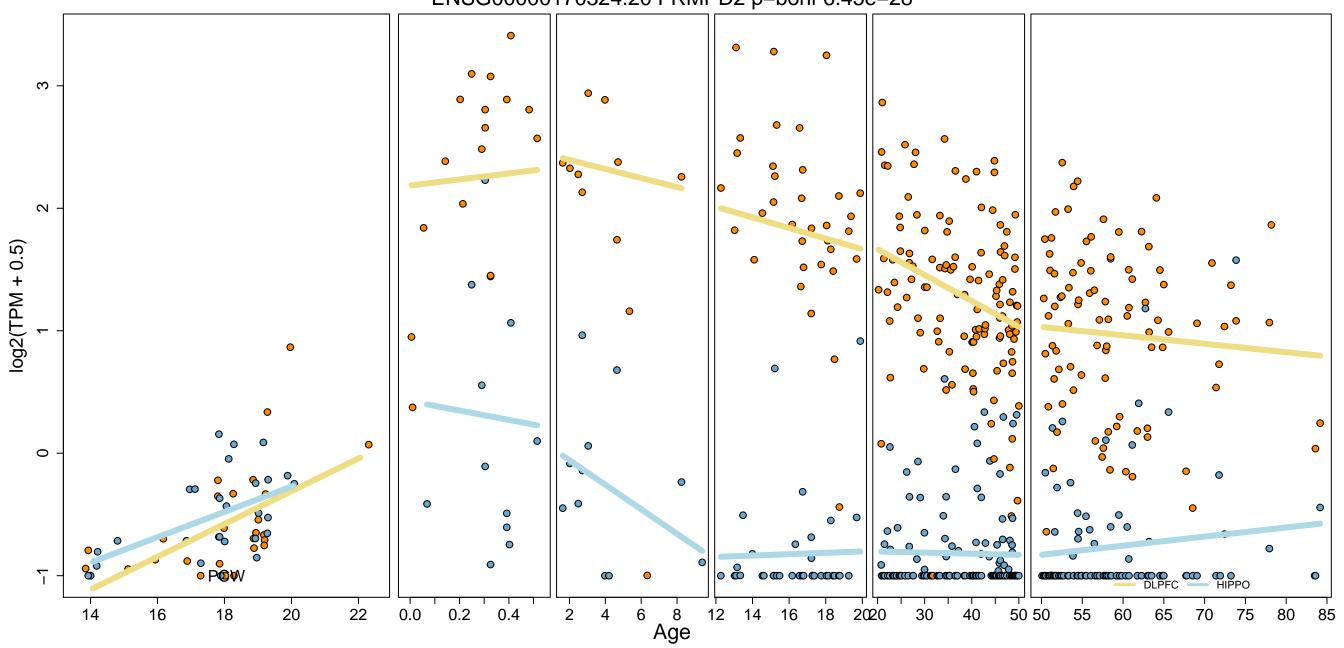
ENSG00000237187.8 NR2F1-AS1 p-bonf 3.97e-28



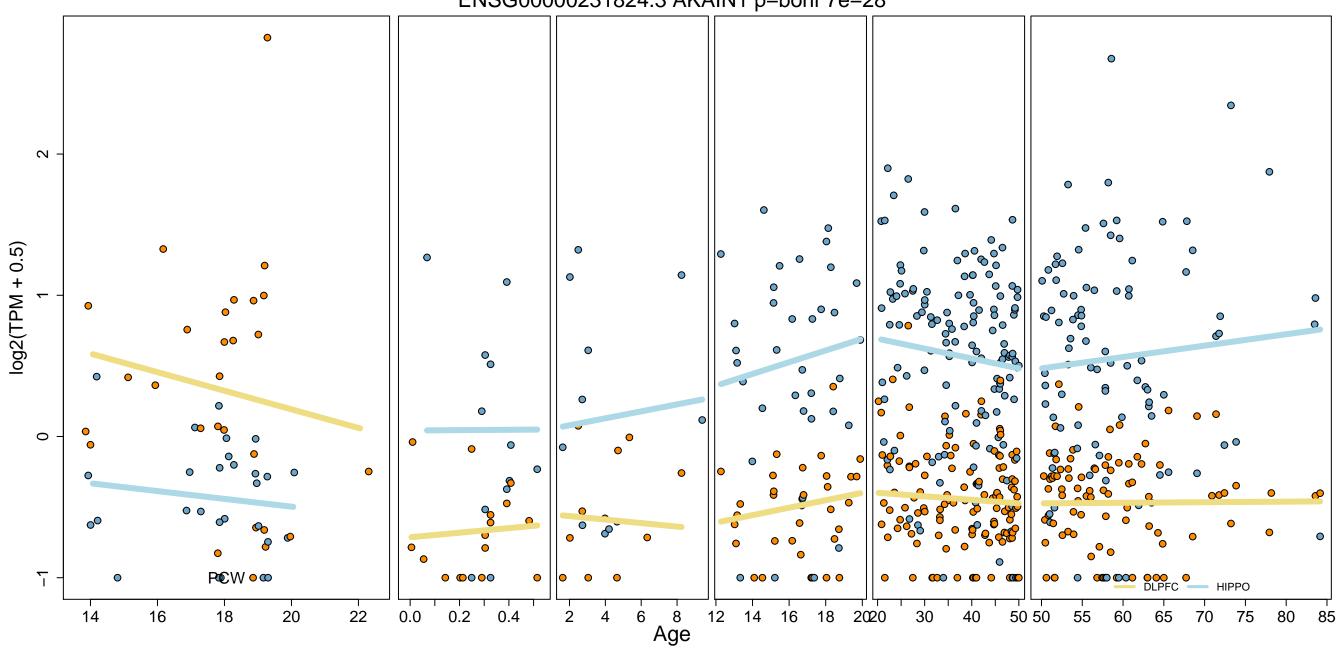
ENSG00000006283.17 CACNA1G p-bonf 4.81e-28



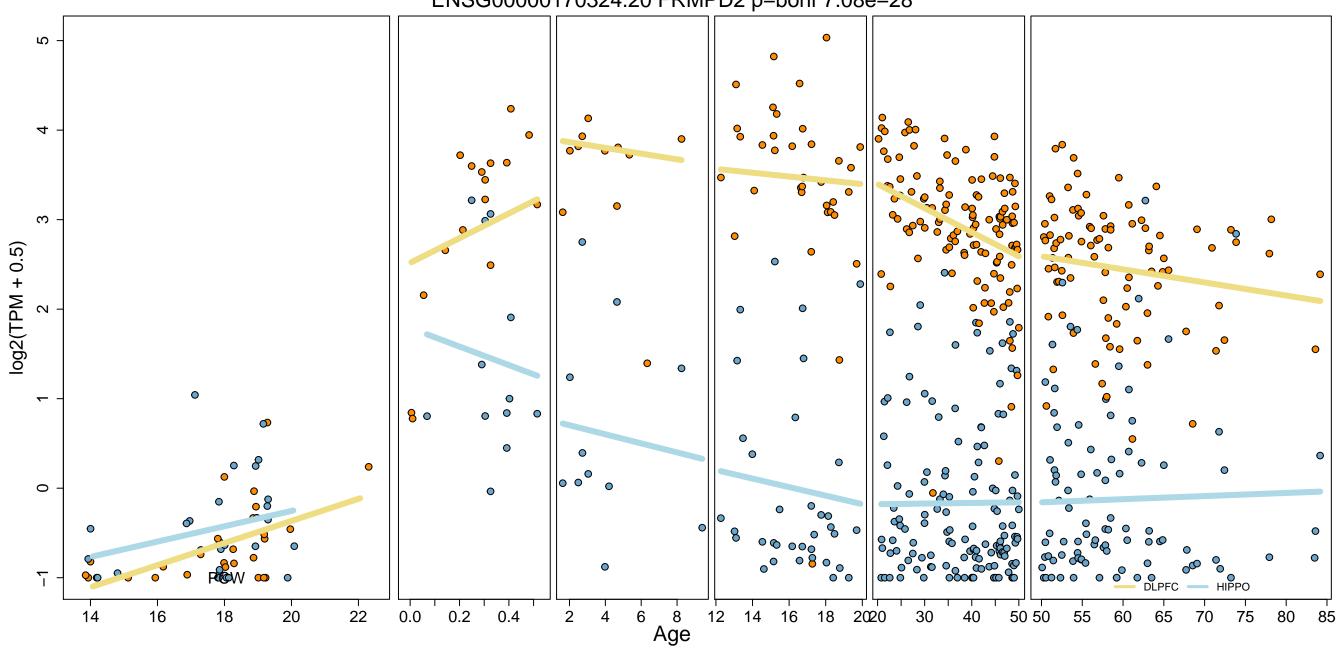
ENSG00000170324.20 FRMPD2 p-bonf 6.45e-28



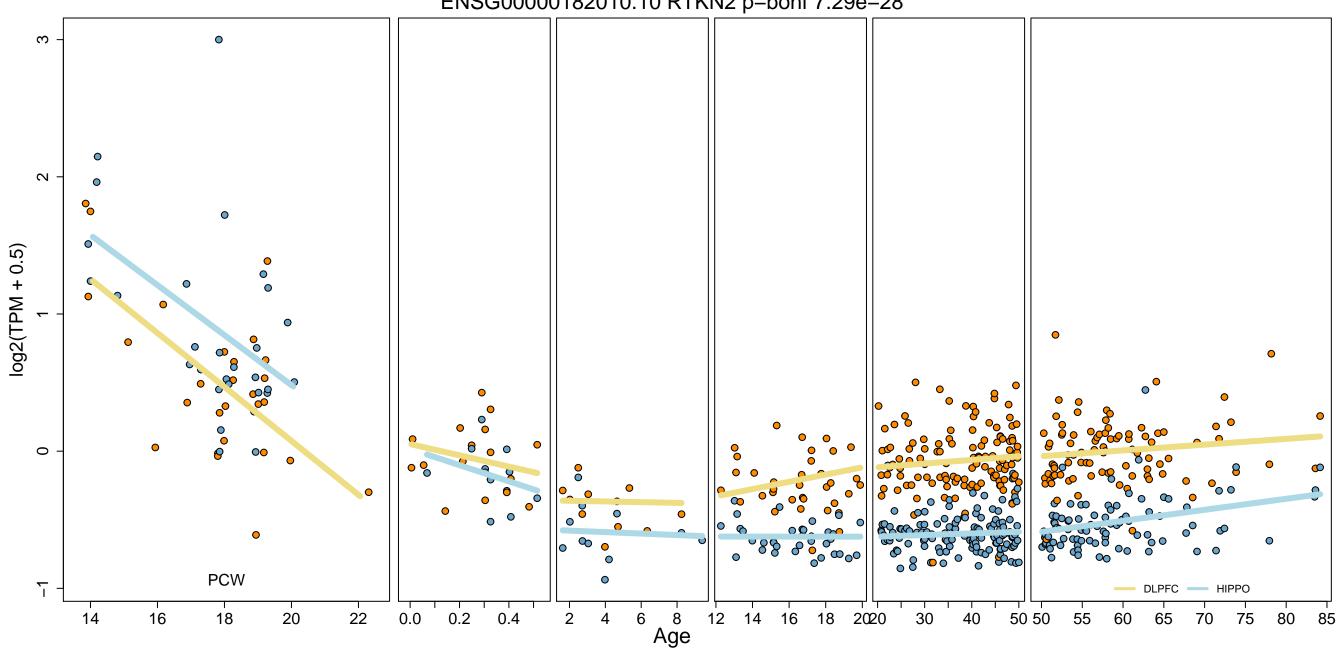
ENSG00000231824.3 AKAIN1 p-bonf 7e-28



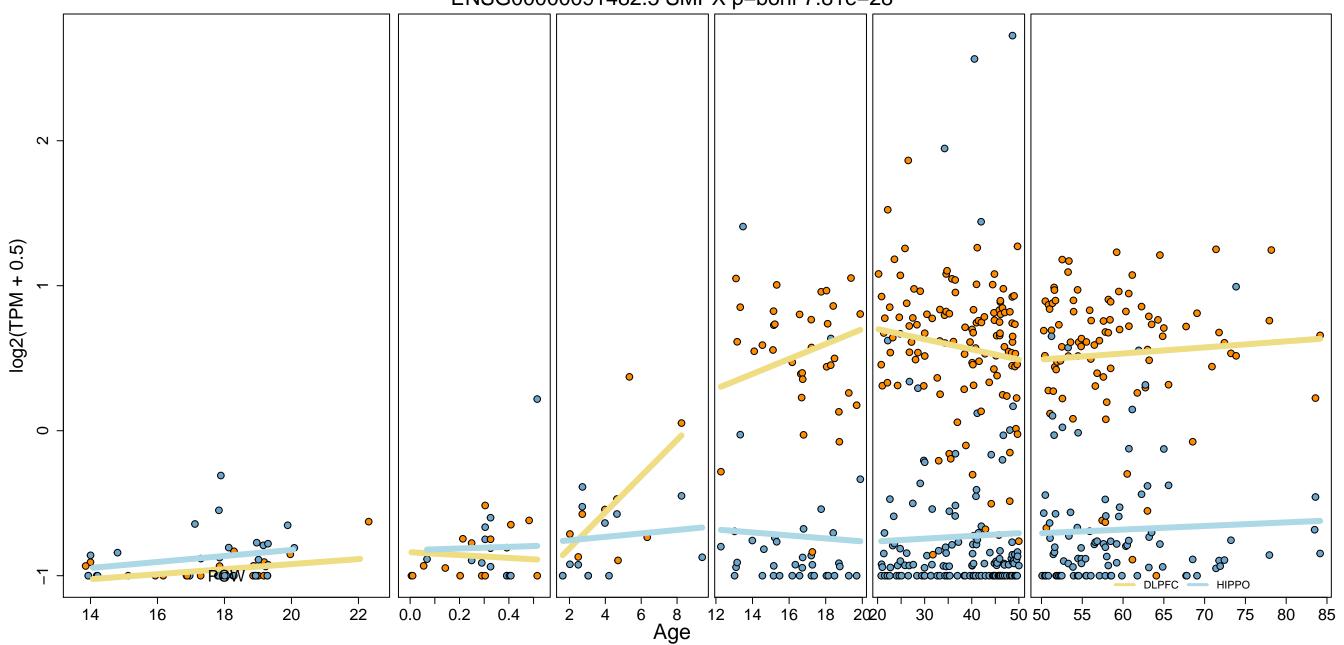
ENSG00000170324.20 FRMPD2 p-bonf 7.08e-28



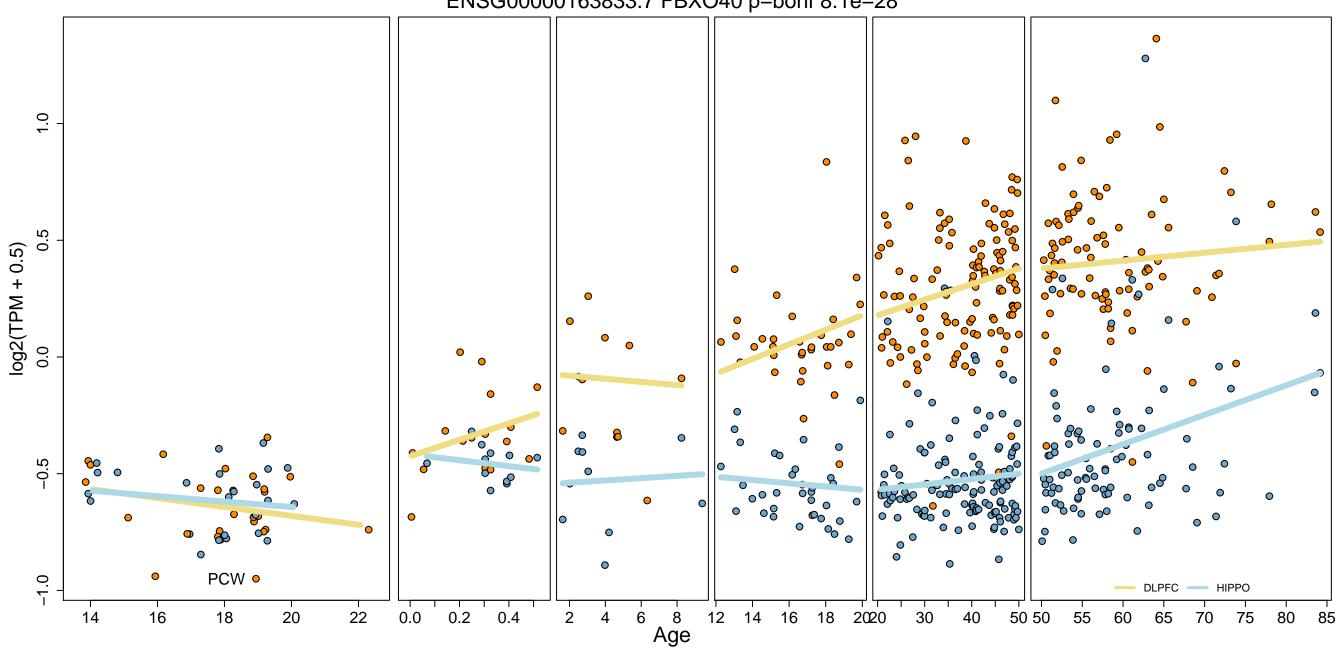
ENSG00000182010.10 RTKN2 p-bonf 7.29e-28



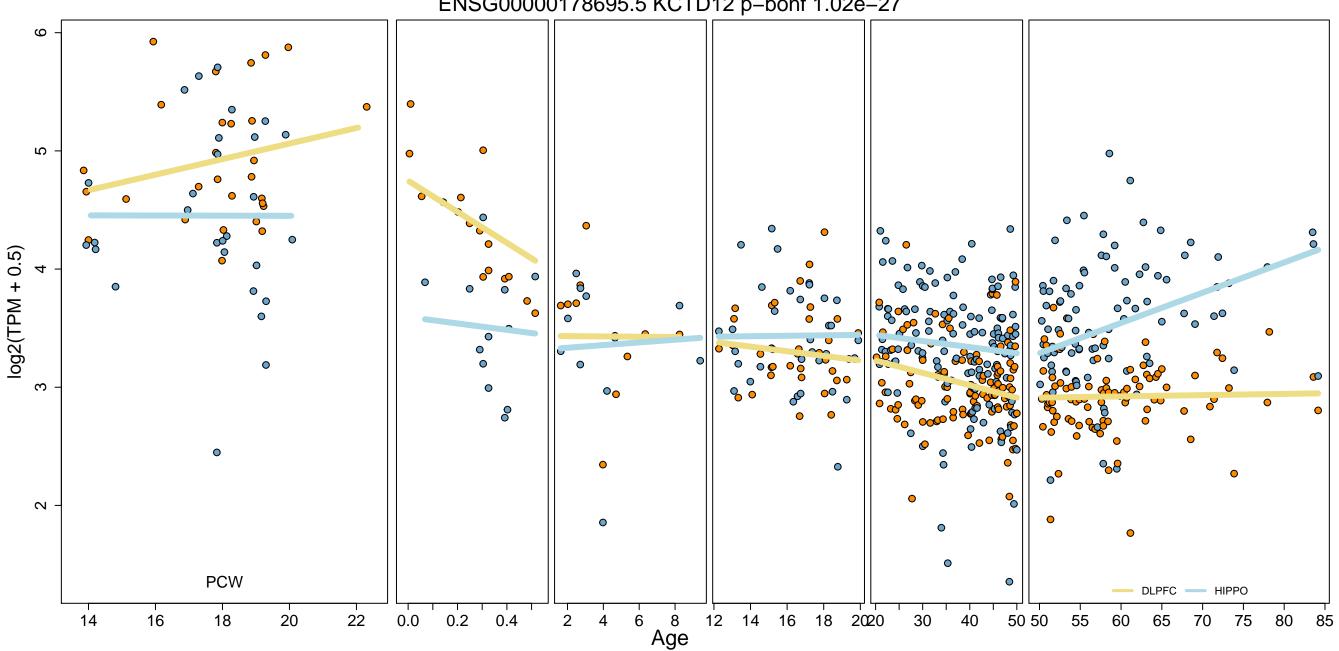
ENSG00000091482.5 SMPX p-bonf 7.81e-28



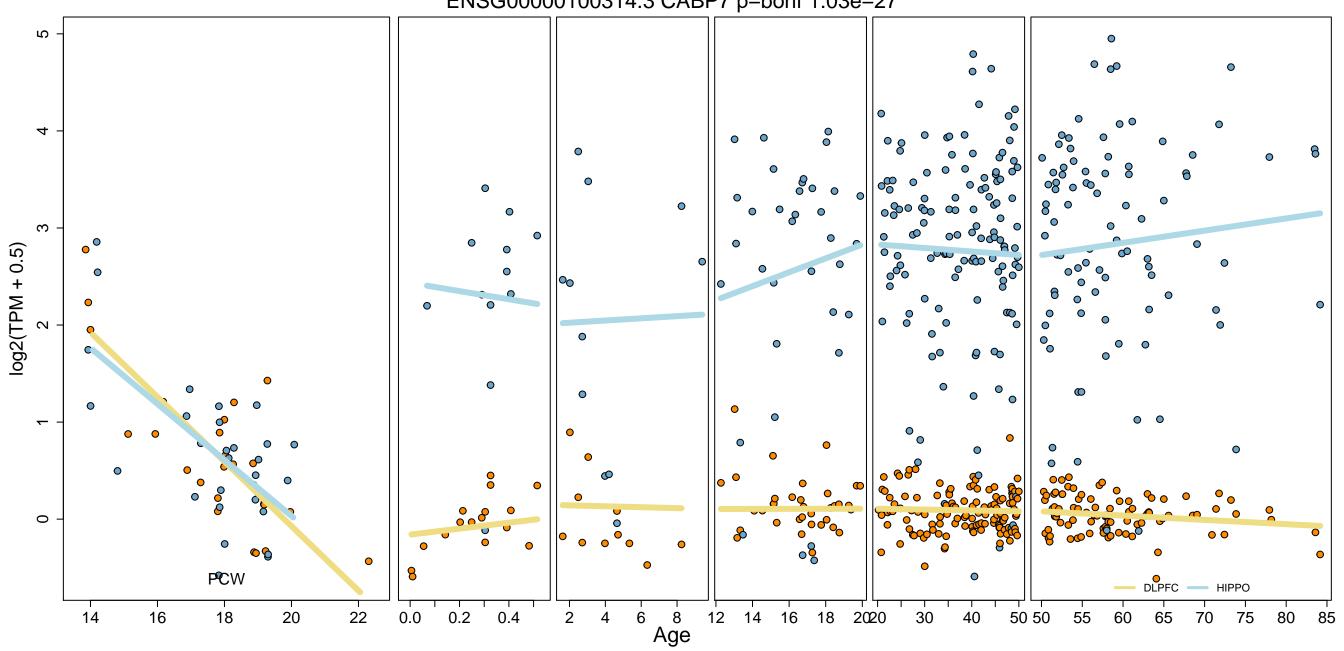
ENSG00000163833.7 FBXO40 p-bonf 8.1e-28



ENSG00000178695.5 KCTD12 p-bonf 1.02e-27



ENSG00000100314.3 CABP7 p-bonf 1.03e-27



ENSG00000231689.2 LINC01090 p-bonf 1.36e-27

