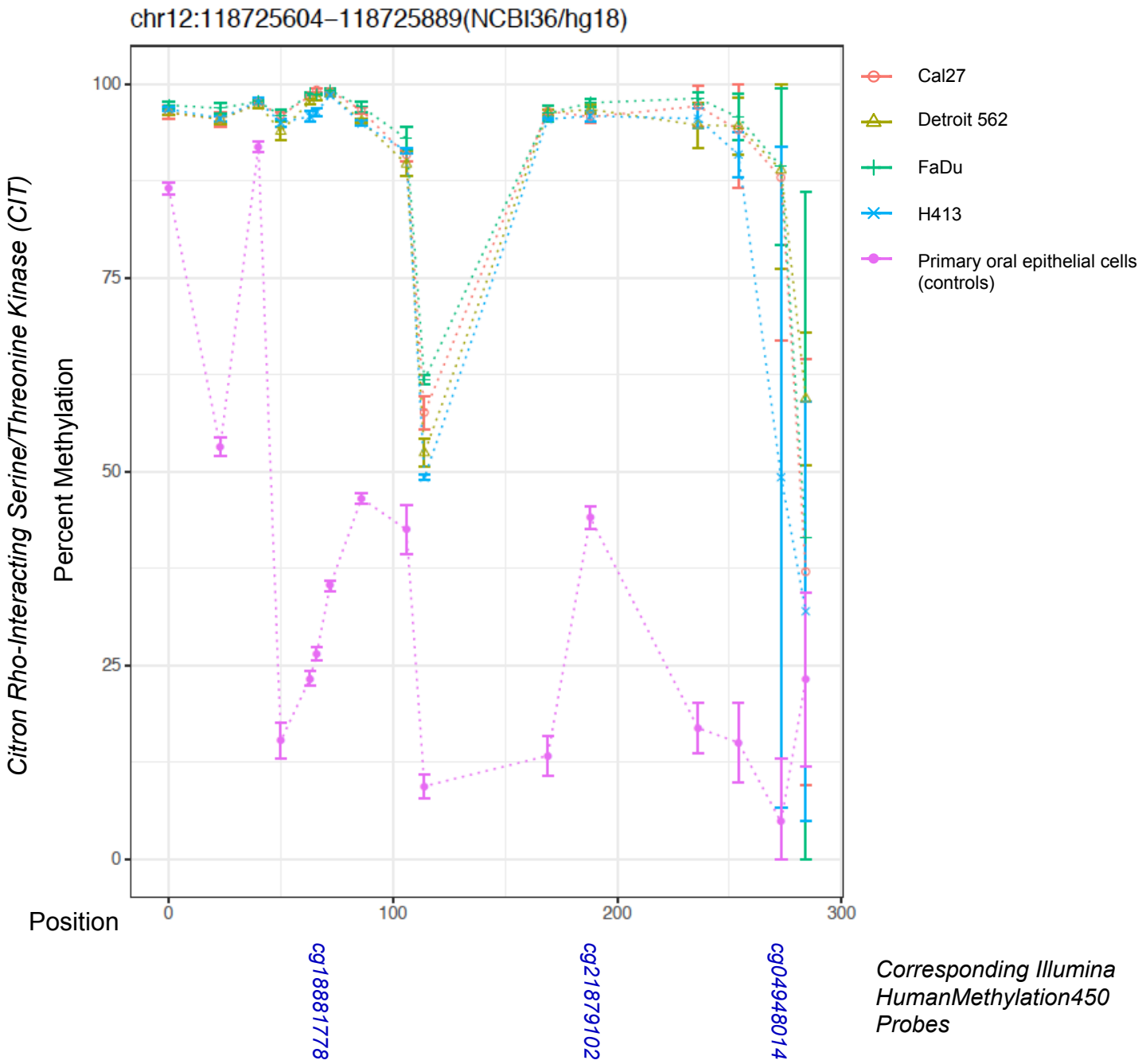
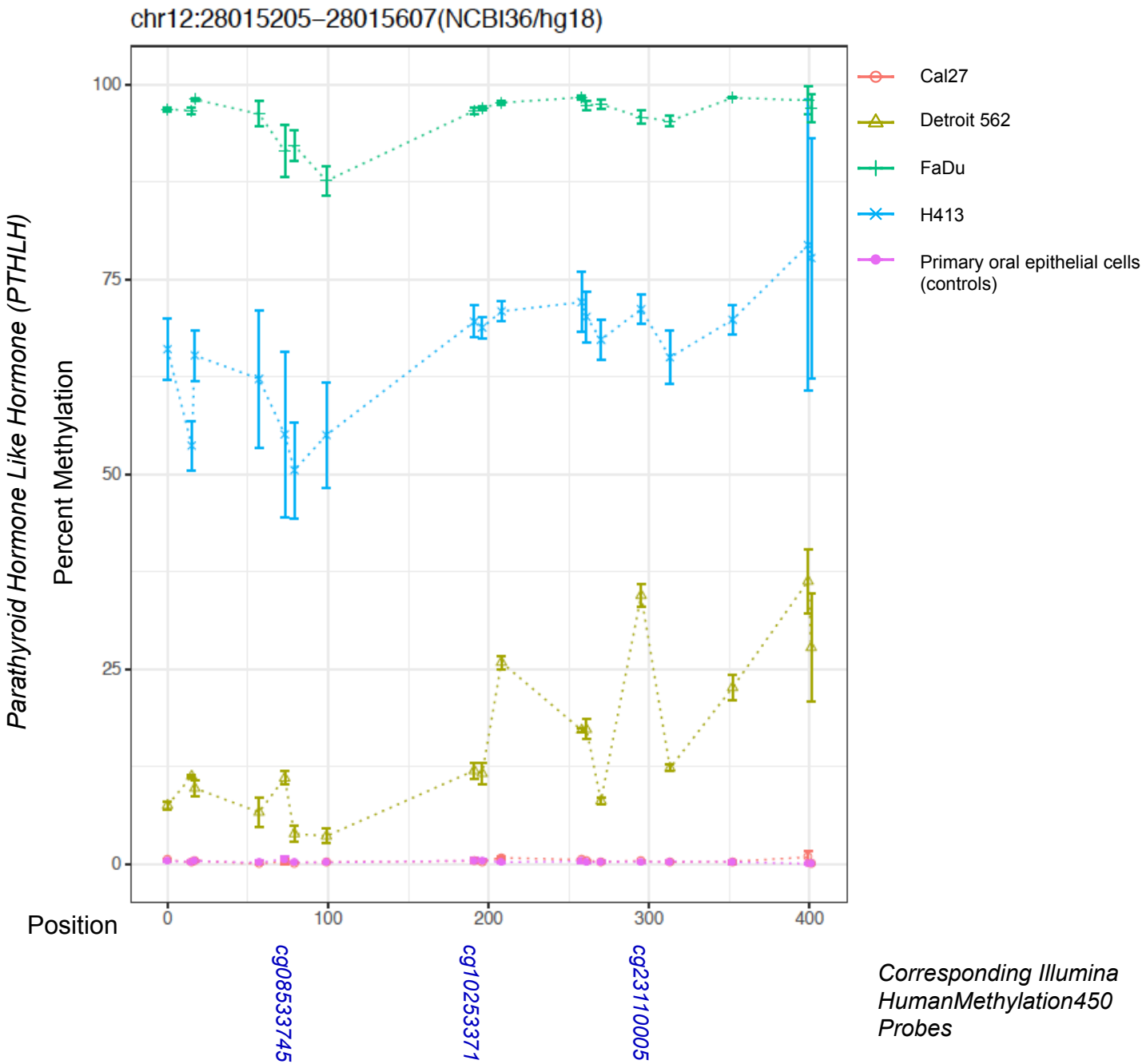
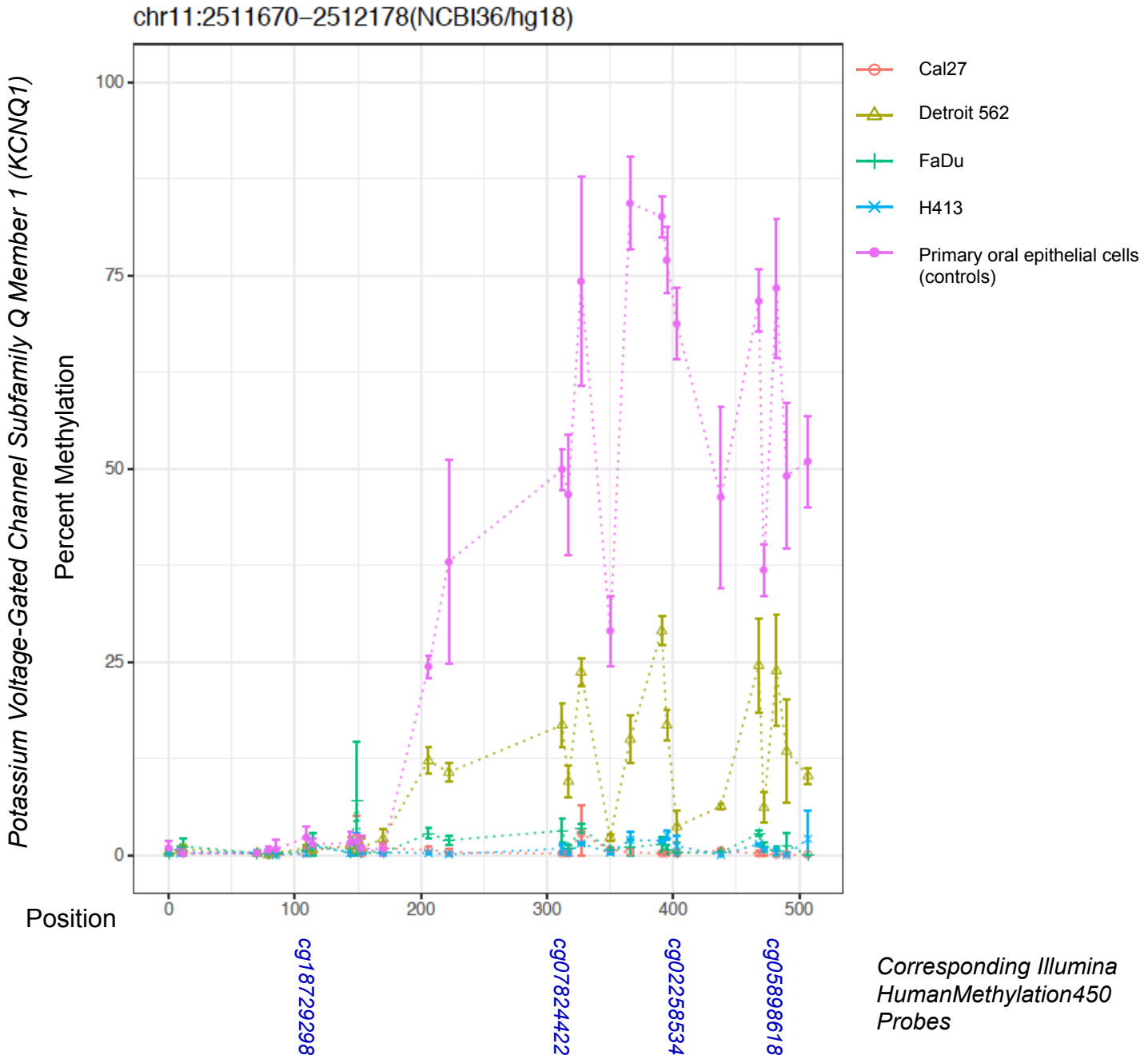
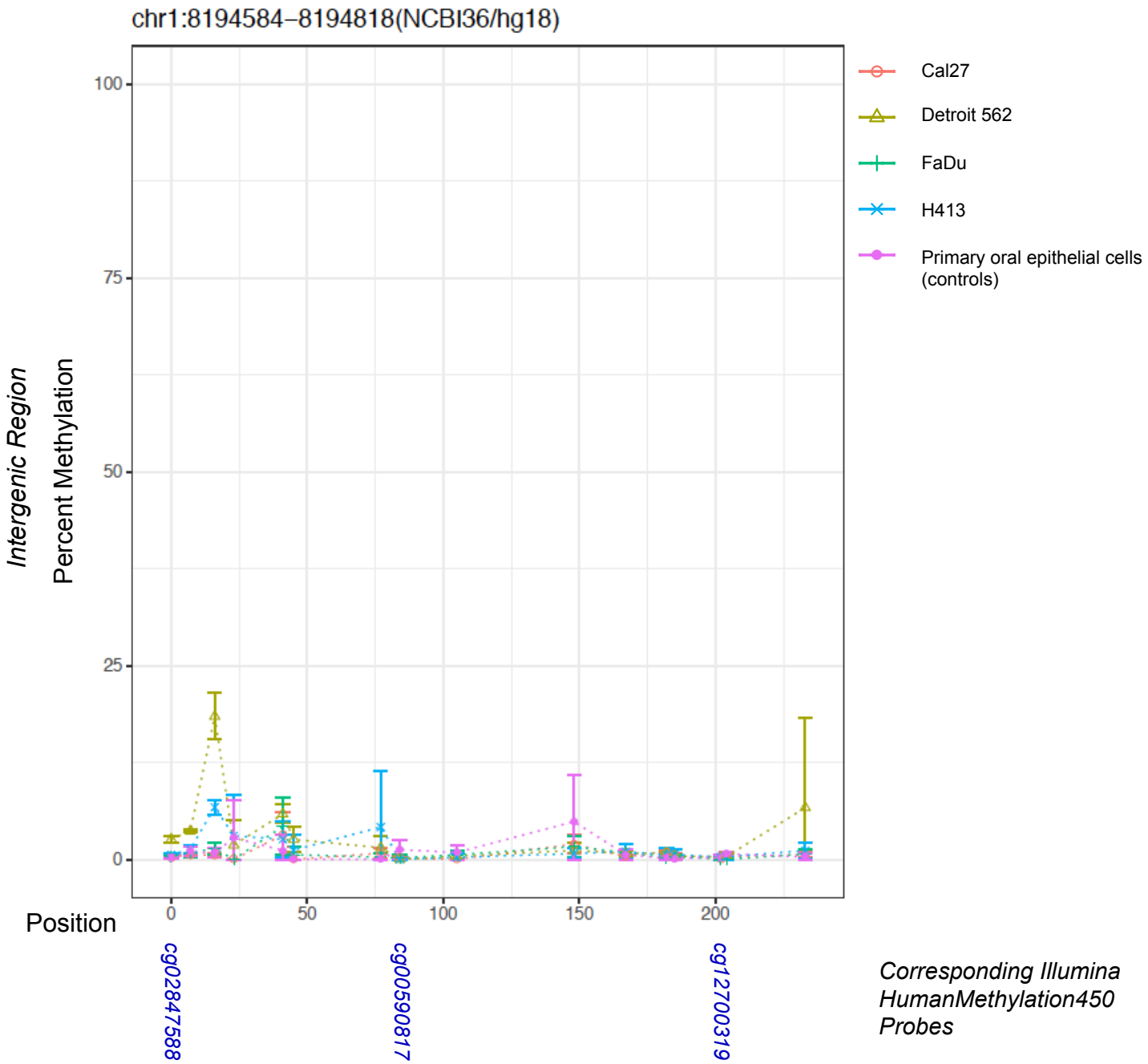
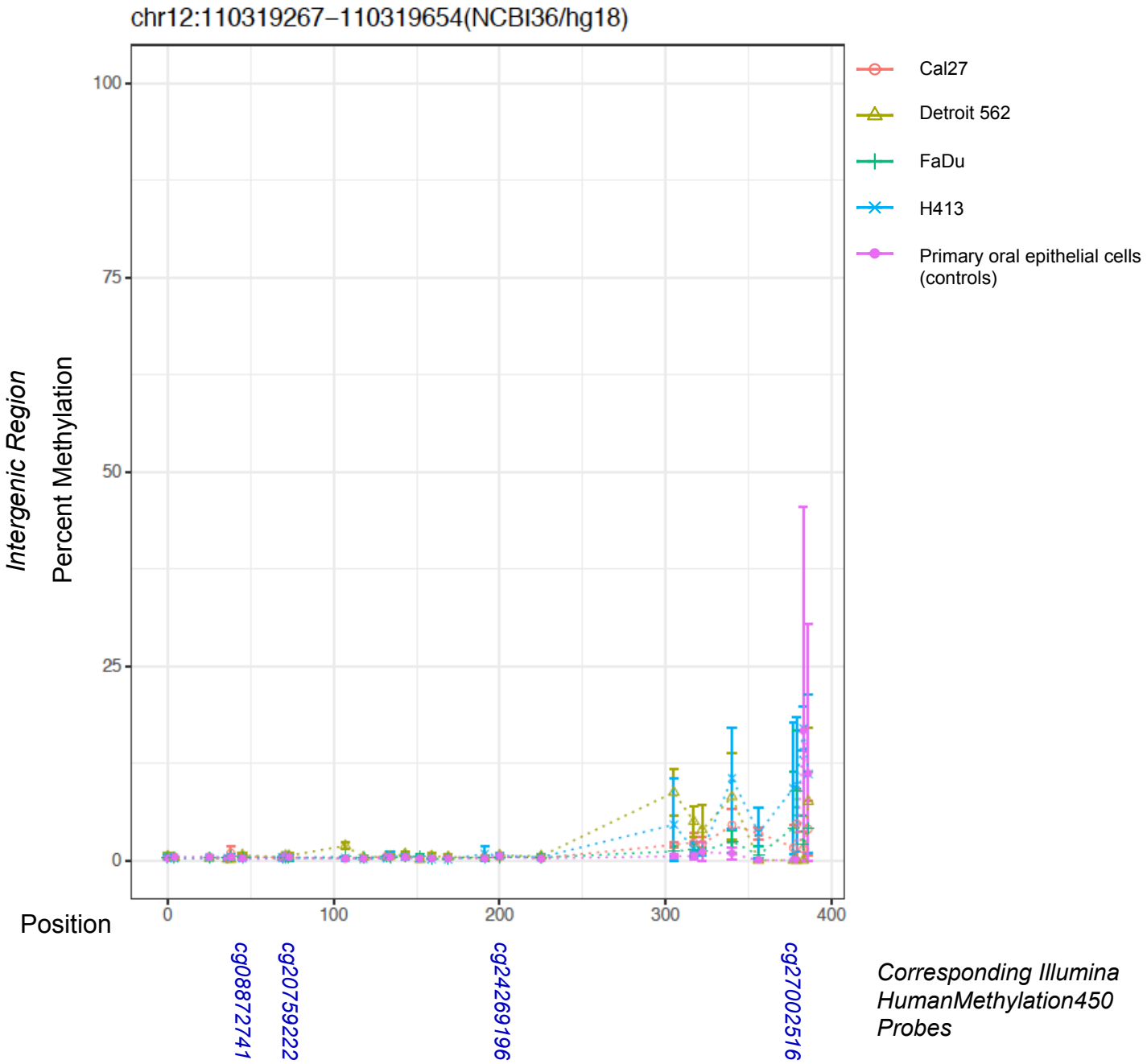
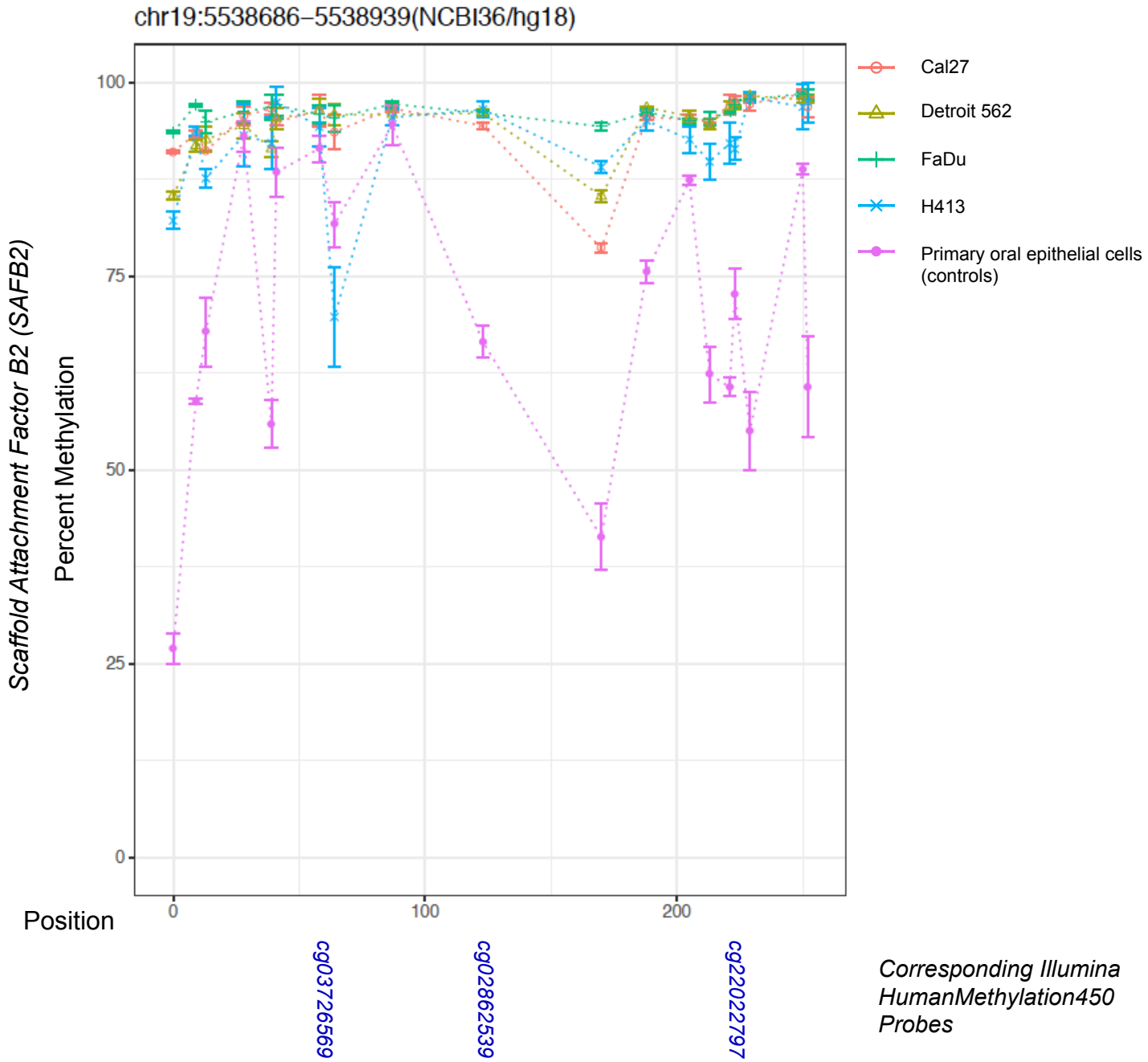

chr12:118725604−118725889(NCBI36/hg18)
Citron Rho-Interacting Serine/Threonine Kinase (CIT)
Percent Methylation
100
75
50
25
0
Position
0
100
200
300
cg18881778
cg21879102
cg04648014
Corresponding Illumina HumanMethylation450 Probes
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)

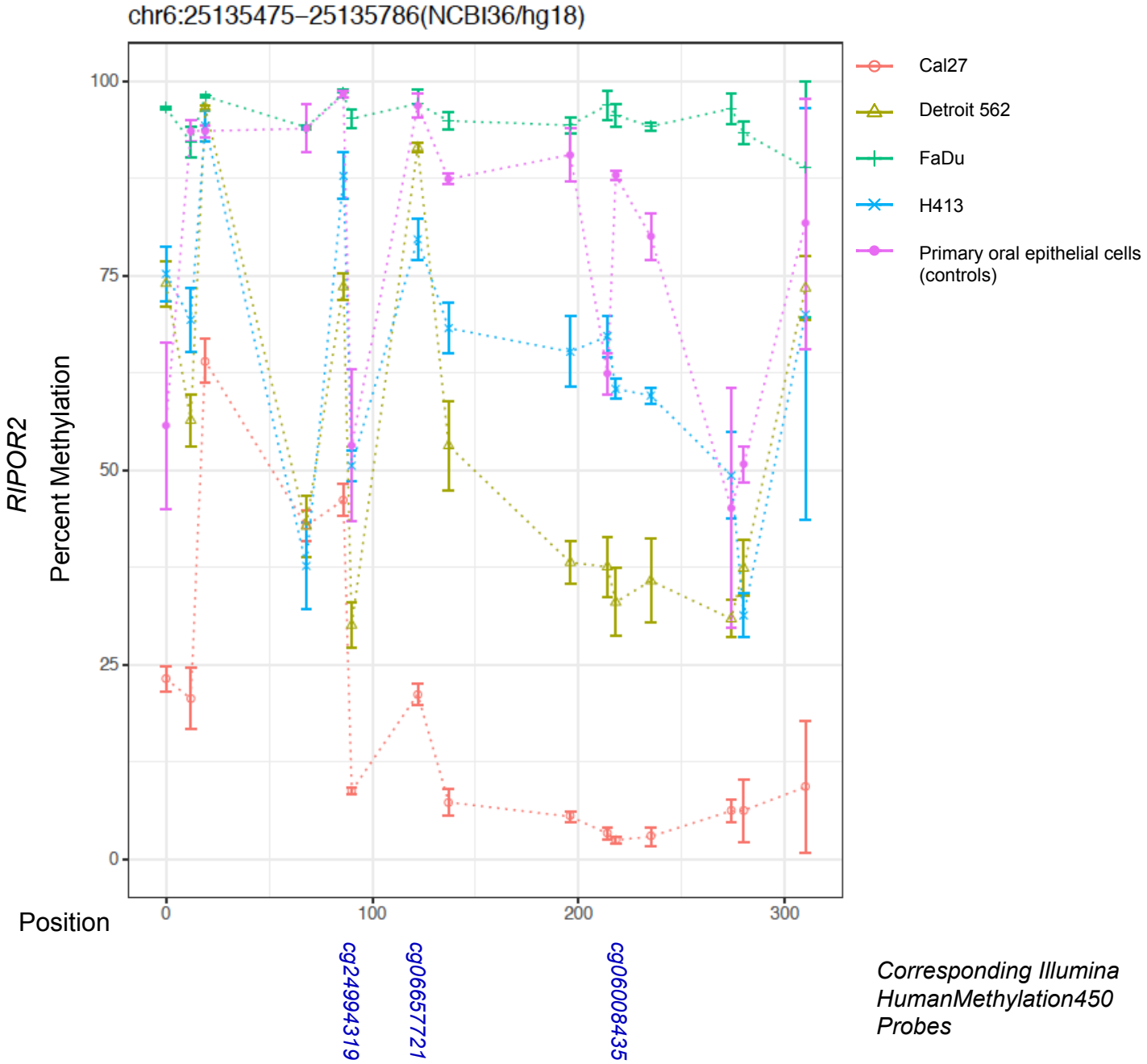

chr12:28015205–28015607(NCBI36/hg18)
Parathyroid Hormone Like Hormone (PTHLH)
Percent Methylation
100
75
50
25
0
Position
0
100
200
300
400
cg08533745
cg10253371
cg23110005
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Corresponding Illumina HumanMethylation450 Probes

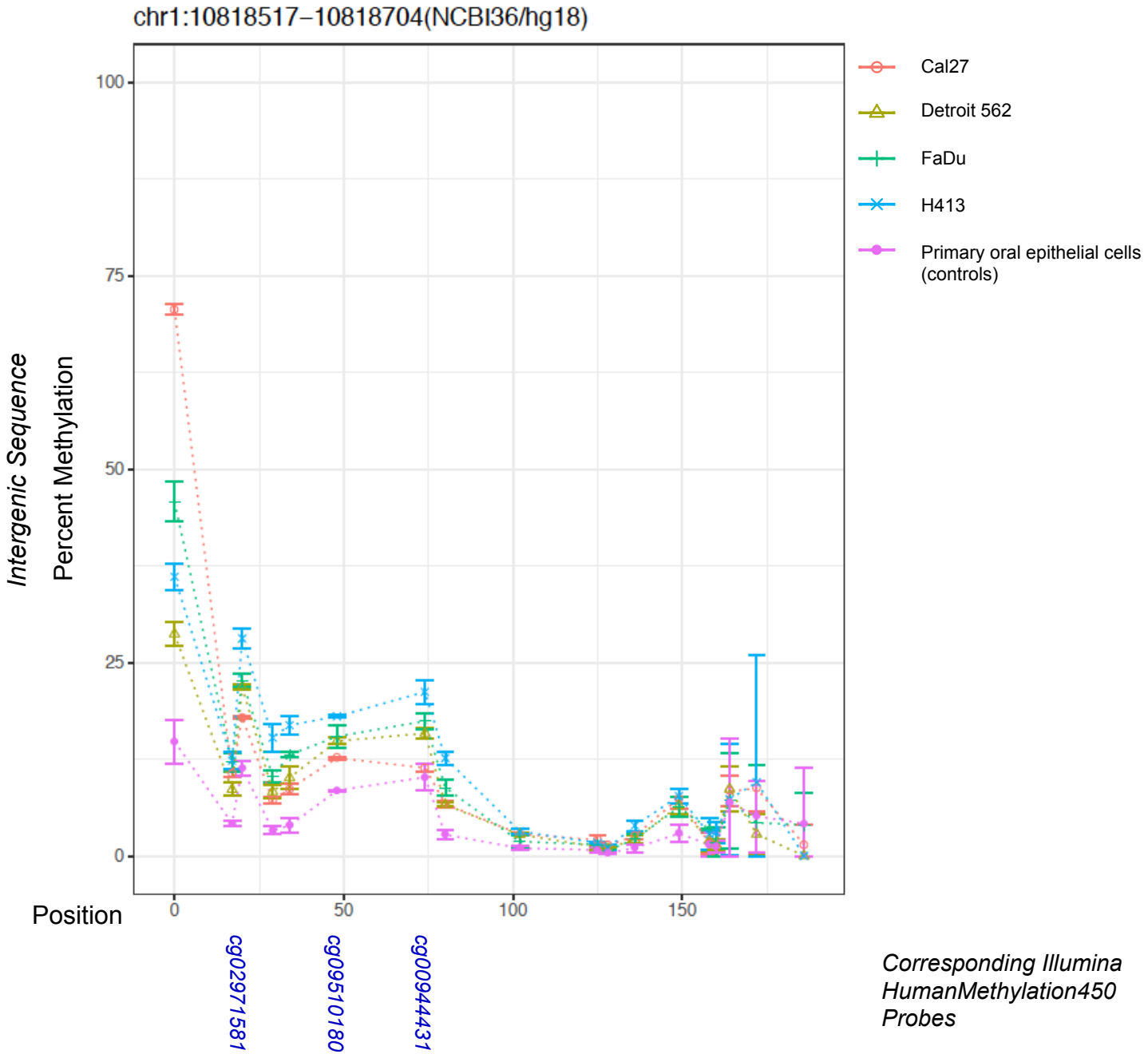

chr11:2511670–2512178(NCBI36/hg18)
Potassium Voltage-Gated Channel Subfamily Q Member 1 (KCNQ1)
Percent Methylation
100
75
50
25
0
0
100
200
300
400
500
Position
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
cg18729298
cg07824422
cg02258534
cg05898618
Corresponding Illumina HumanMethylation450 Probes

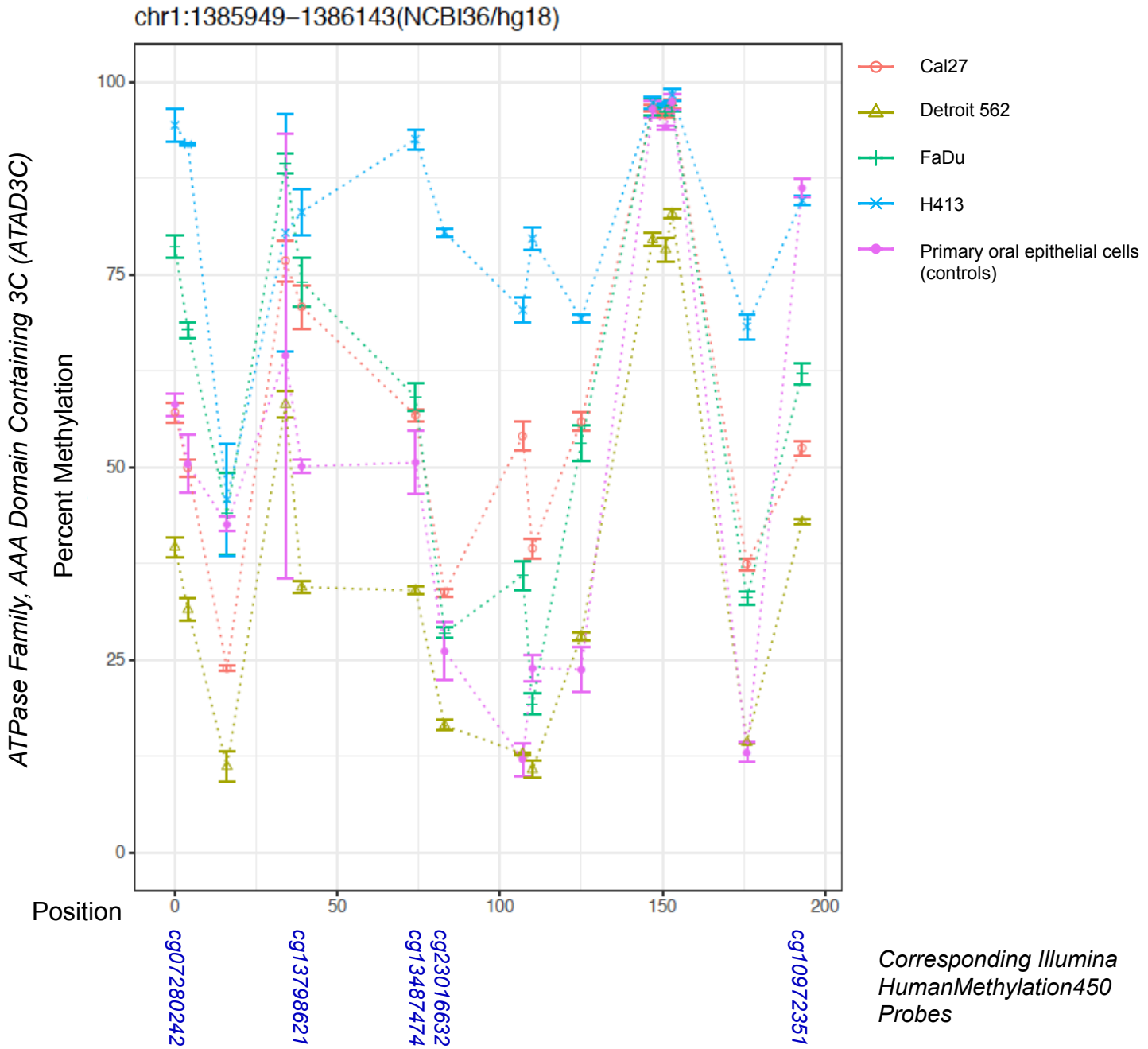

chr1:8194584–8194818(NCBI36/hg18)
Intergenic Region
Percent Methylation
100
75
50
25
0
0
50
100
150
200
Position
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
cg02847588
cg00590817
cg12700319
Corresponding Illumina HumanMethylation450 Probes

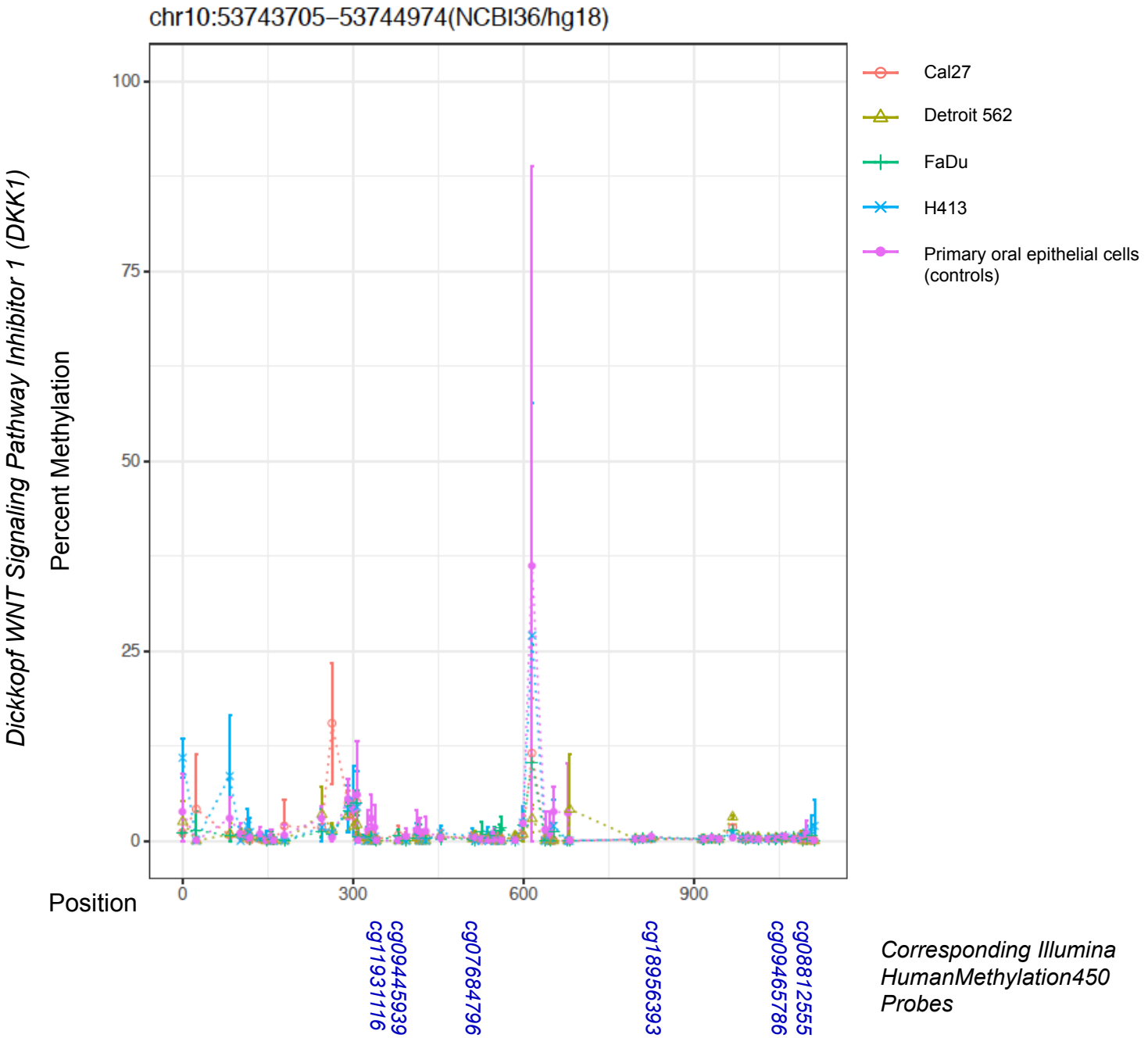

chr12:110319267–110319654(NCBI36/hg18)
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Intergenic Region
Percent Methylation
100
75
50
25
0
0
100
200
300
400
Position
cg08872741
cg20759222
cg24269196
cg27002516
Corresponding Illumina HumanMethylation450 Probes

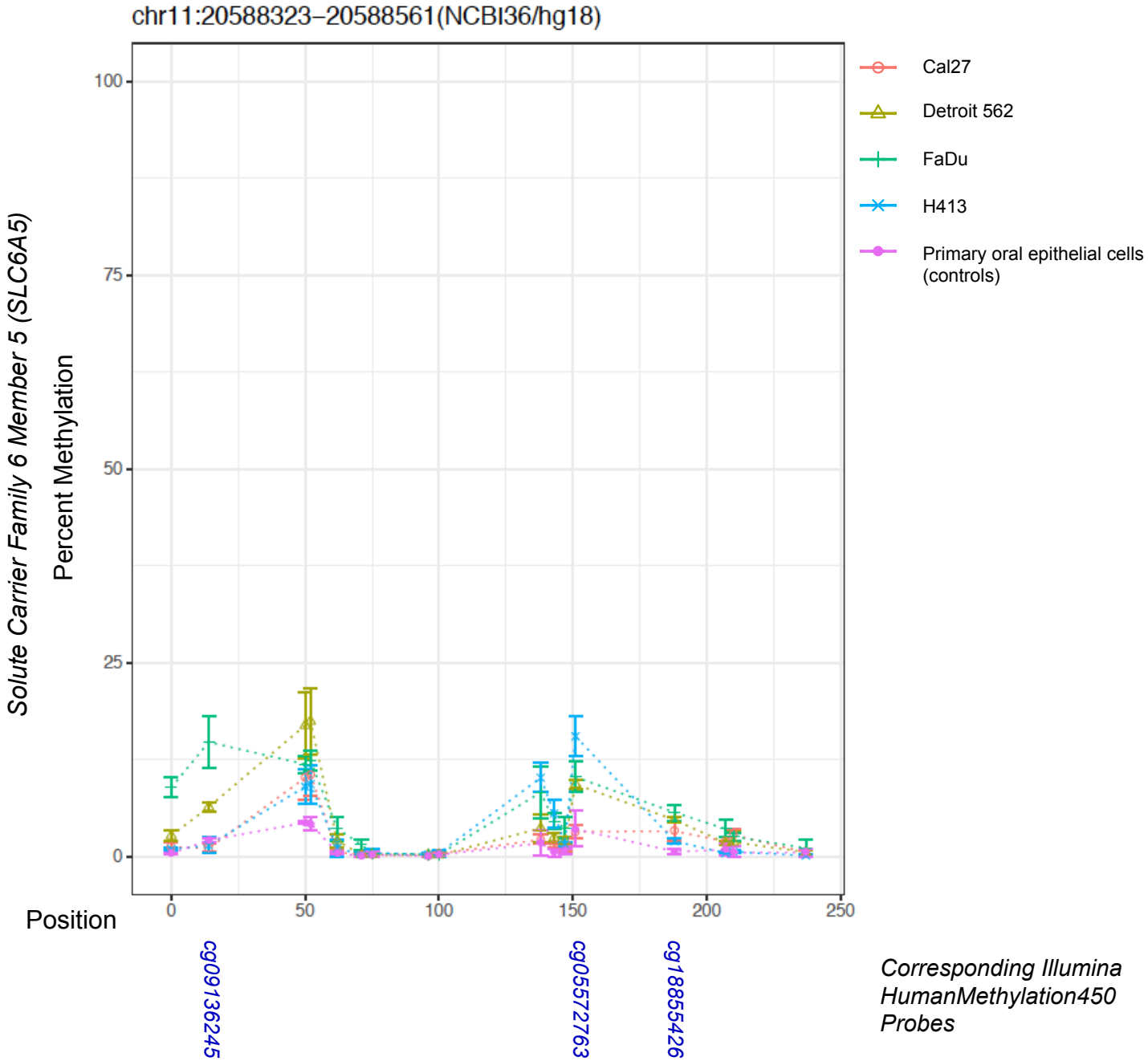

chr19:5538686–5538939(NCBI36/hg18)
Scaffold Attachment Factor B2 (SAFB2)
Percent Methylation
100
75
50
25
0
0
100
200
Position
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
cg03726569
cg02862539
cg22022797
Corresponding Illumina HumanMethylation450 Probes

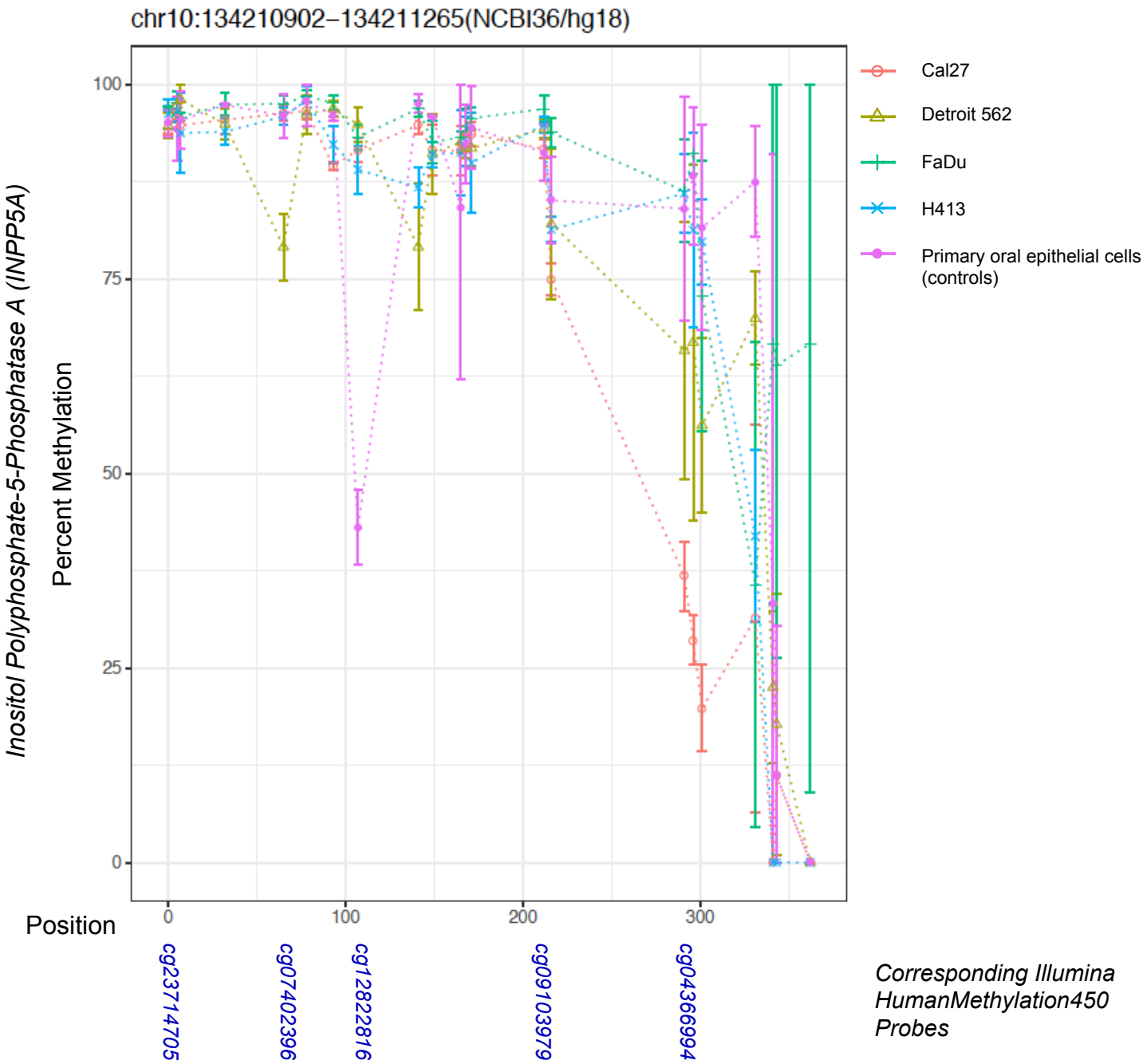

chr6:25135475–25135786(NCBI36/hg18)
RIPOR2
Percent Methylation
100
75
50
25
0
0
100
200
300
Position
cg24994319
cg06657721
cg06008435
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Corresponding Illumina HumanMethylation450 Probes

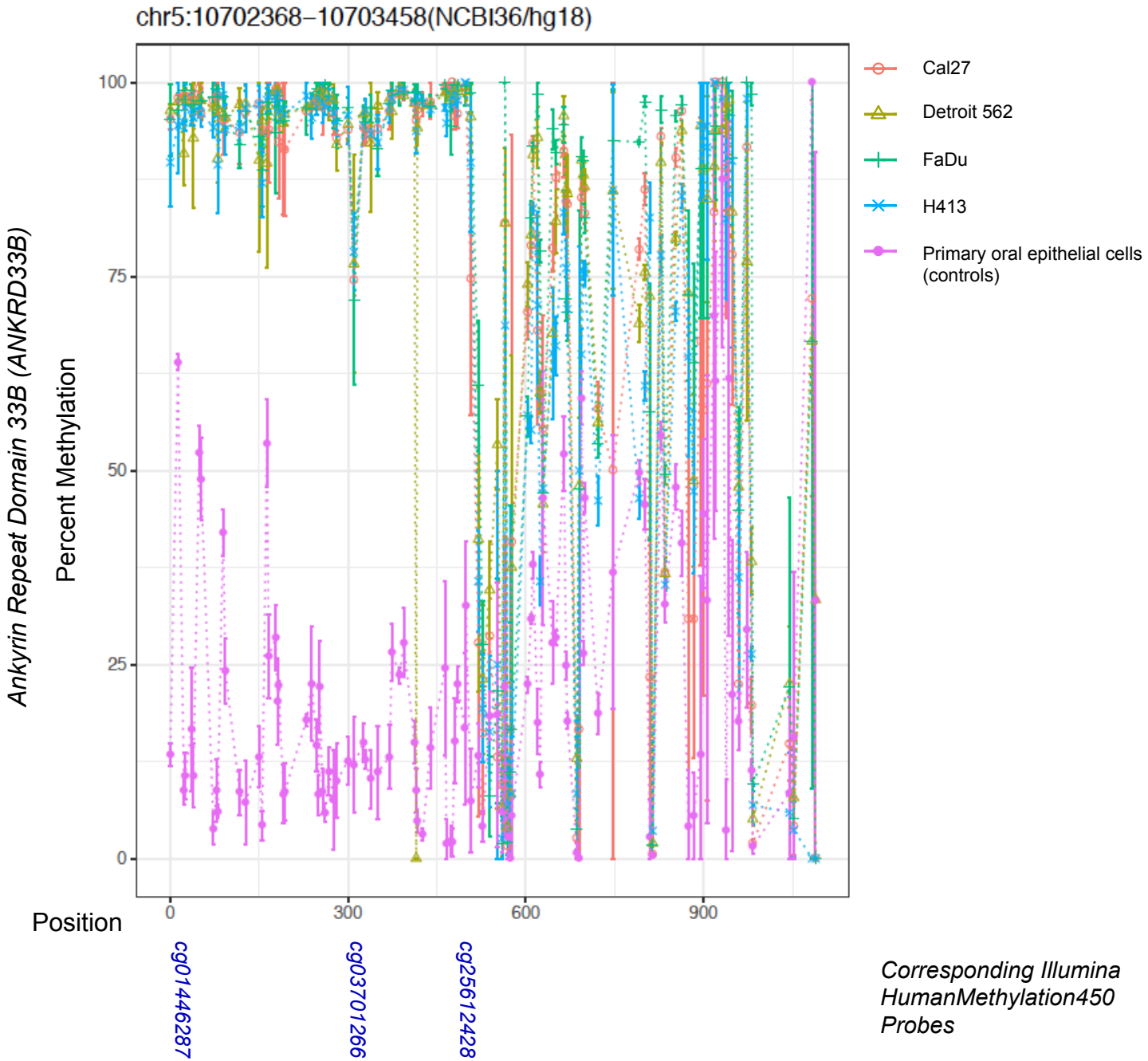

chr1:10818517−10818704(NCBI36/hg18)
Intergenic Sequence
Percent Methylation
100
75
50
25
0
Position
0
50
100
150
cg02971581
cg09510180
cg00944431
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Corresponding Illumina HumanMethylation450 Probes

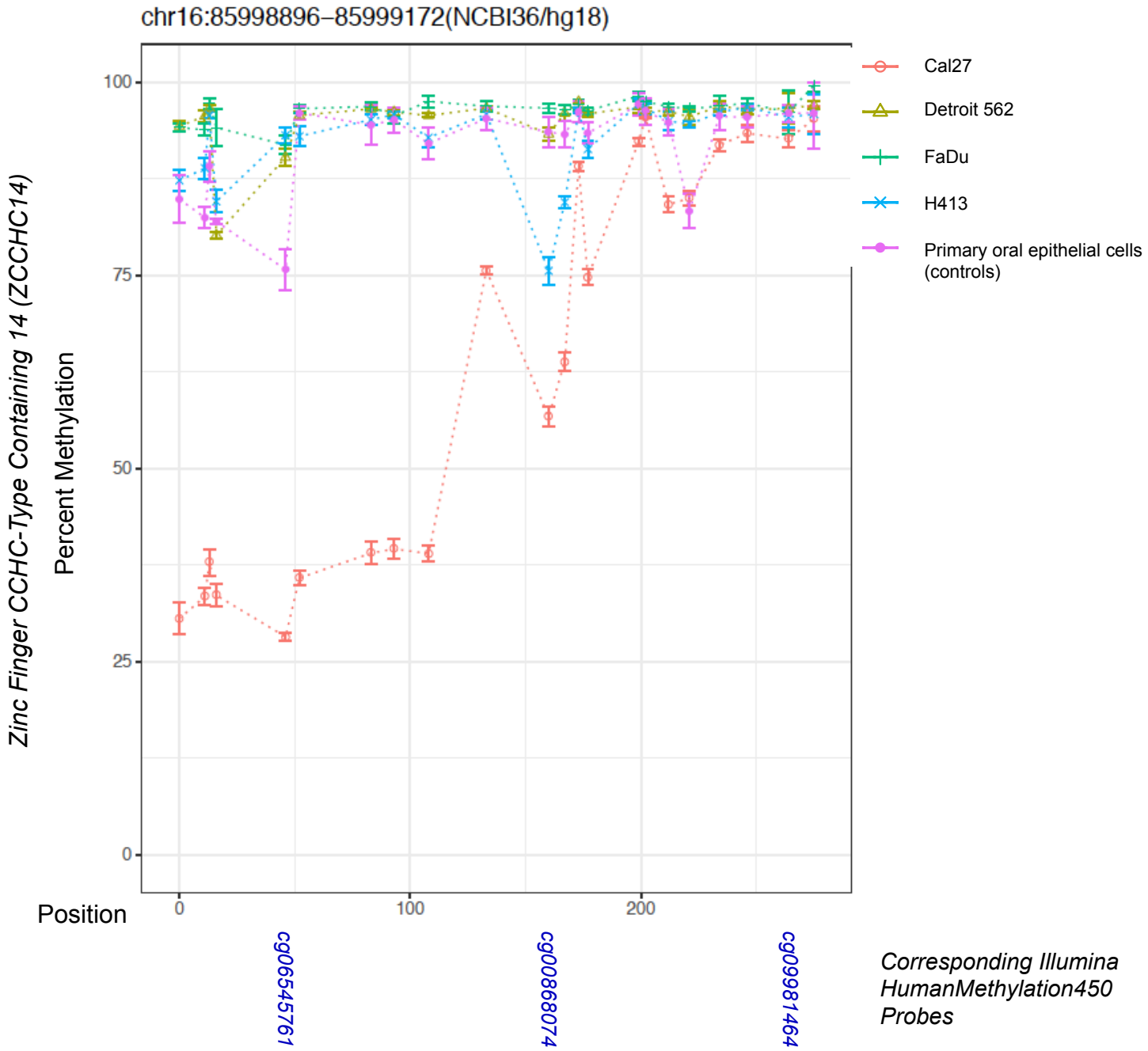

chr1:1385949−1386143(NCBI36/hg18)
ATPase Family, AAA Domain Containing 3C (ATAD3C)
Percent Methylation
100
75
50
25
0
Position
0
50
100
150
200
cg07280242
cg13798621
cg13487474
cg23016632
cg10972351
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Corresponding Illumina HumanMethylation450 Probes

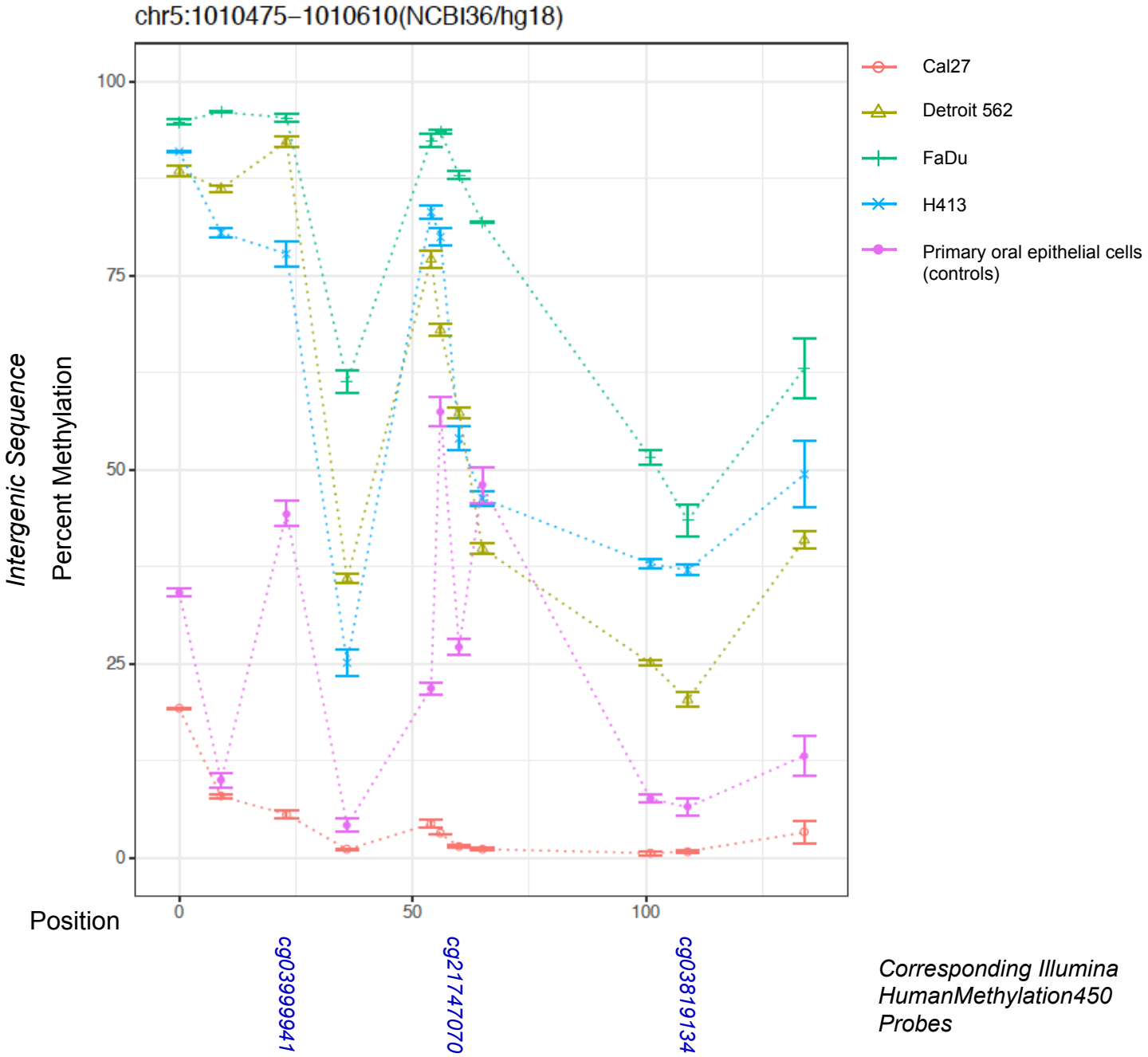

chr10:53743705–53744974(NCBI36/hg18)
Dickkopf WNT Signaling Pathway Inhibitor 1 (DKK1)
Percent Methylation
100
75
50
25
0
0
300
600
900
Position
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
cg09445939
cg11931116
cg07684796
cg18956393
cg09465786
cg08812555
Corresponding Illumina HumanMethylation450 Probes

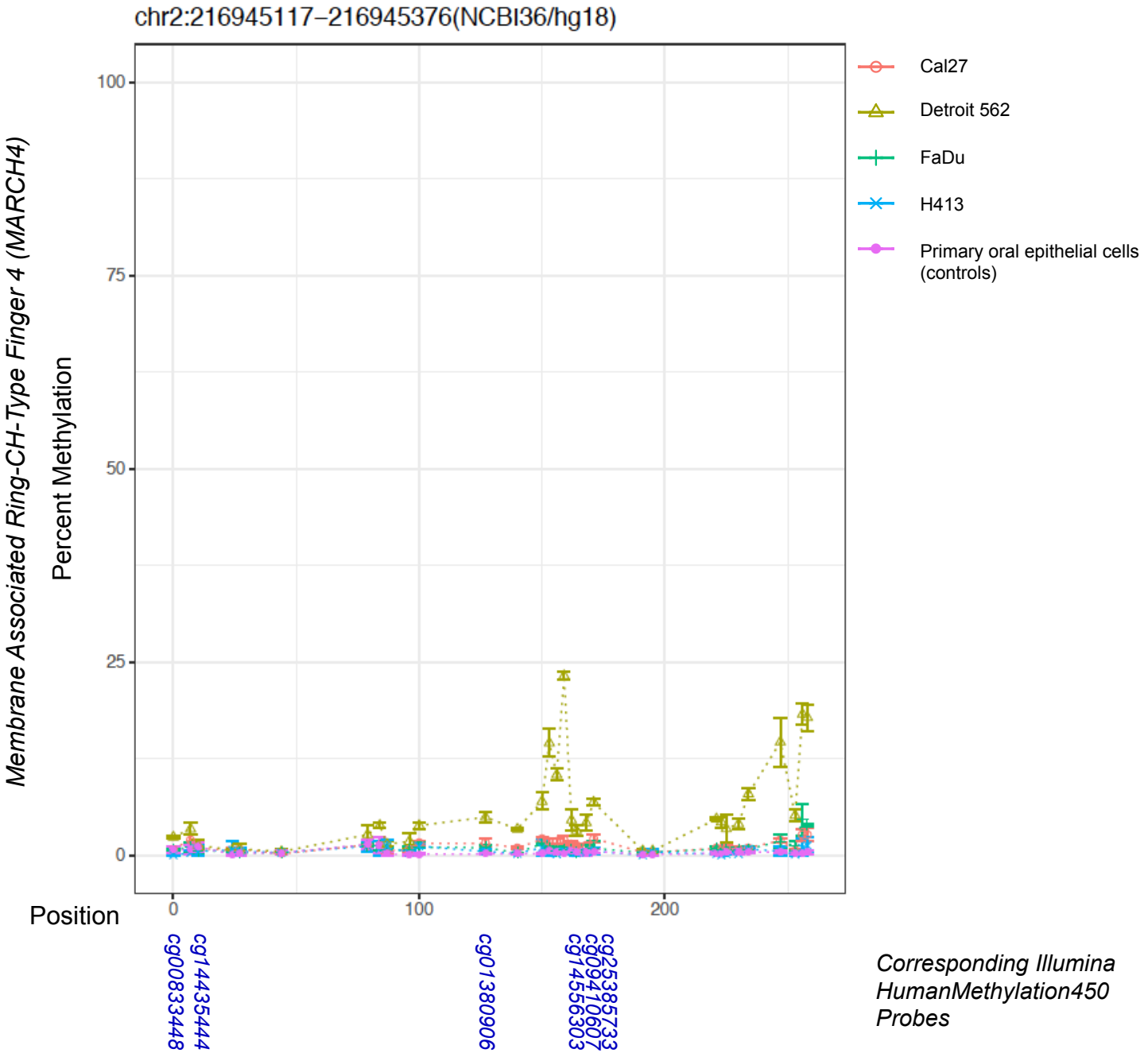

chr11:20588323–20588561(NCBI36/hg18)
Solute Carrier Family 6 Member 5 (SLC6A5)
Percent Methylation
100
75
50
25
0
0
50
100
150
200
250
Position
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
cg09136245
cg05572763
cg18855426
Corresponding Illumina HumanMethylation450 Probes


chr10:134210902−134211265(NCBI36/hg18)
Inositol Polyphosphate-5-Phosphatase A (INPP5A)
Percent Methylation
100
75
50
25
0
Position
0
100
200
300
cg23714705
cg07402396
cg12822816
cg09103979
cg04366994
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Corresponding Illumina HumanMethylation450 Probes


chr5:10702368−10703458(NCBI36/hg18)
Ankyrin Repeat Domain 33B (ANKRD33B)
Percent Methylation
100
75
50
25
0
Position
0
300
600
900
cg01446287
cg03701266
cg25612428
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Corresponding Illumina HumanMethylation450 Probes


chr16:85998896–85999172(NCBI36/hg18)
Zinc Finger CCHC-Type Containing 14 (ZCCHC14)
Percent Methylation
100
75
50
25
0
Position
0
100
200
cg06545761
cg00868074
cg09981464
Corresponding Illumina HumanMethylation450 Probes
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)


chr5:1010475–1010610(NCBI36/hg18)
Intergenic Sequence
Percent Methylation
100
75
50
25
0
Position
0
50
100
cg03999941
cg21747070
cg03819134
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Corresponding Illumina HumanMethylation450 Probes


chr2:216945117–216945376(NCBI36/hg18)
Membrane Associated Ring-CH-Type Finger 4 (MARCH4)
Percent Methylation
100
75
50
25
0
Position
0
100
200
cg00833448
cg14435444
cg01380906
cg14556303
cg09410607
cg25385733
Cal27
Detroit 562
FaDu
H413
Primary oral epithelial cells (controls)
Corresponding Illumina HumanMethylation450 Probes