

## **Intratumoral heterogeneity of breast cancer detected by epialleles shows association with hypoxic microenvironment**

Yihan Wang<sup>2</sup>, Yan Zhang<sup>1,3</sup>✉, Yan Huang<sup>3</sup>, Chuangeng Chen<sup>3</sup>, Xingda Zhang<sup>4</sup>, Ying Xing<sup>5</sup>, Yue Gu<sup>3</sup>, Mengyan Zhang<sup>3</sup>, Li Cai<sup>5</sup>, Shouping Xu<sup>4</sup>✉, Baoqing Sun<sup>1</sup>✉

<sup>1</sup> State Key Laboratory of Respiratory Disease, Guangzhou Medical University, Guangzhou, 51000, China

<sup>2</sup> College of Bioinformatics Science and Technology, Harbin Medical University, Harbin, 150081, China.

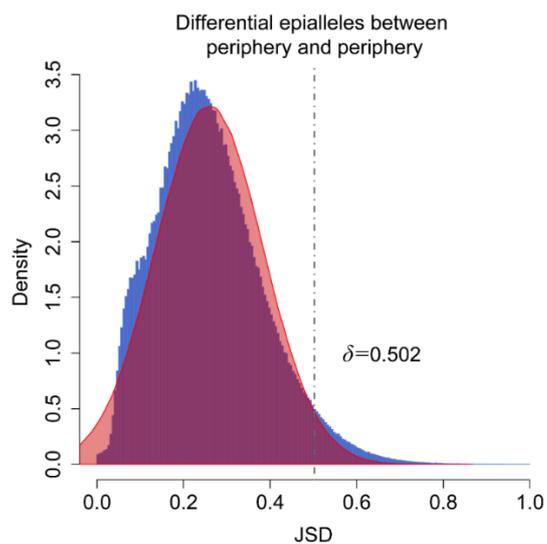
<sup>3</sup> School of Life Science and Technology, Computational Biology Research Center, Harbin Institute of Technology, Harbin, Heilongjiang 150001, China

<sup>4</sup> Department of Breast Surgery, Harbin Medical University Cancer Hospital, Harbin, Heilongjiang 150081, China.

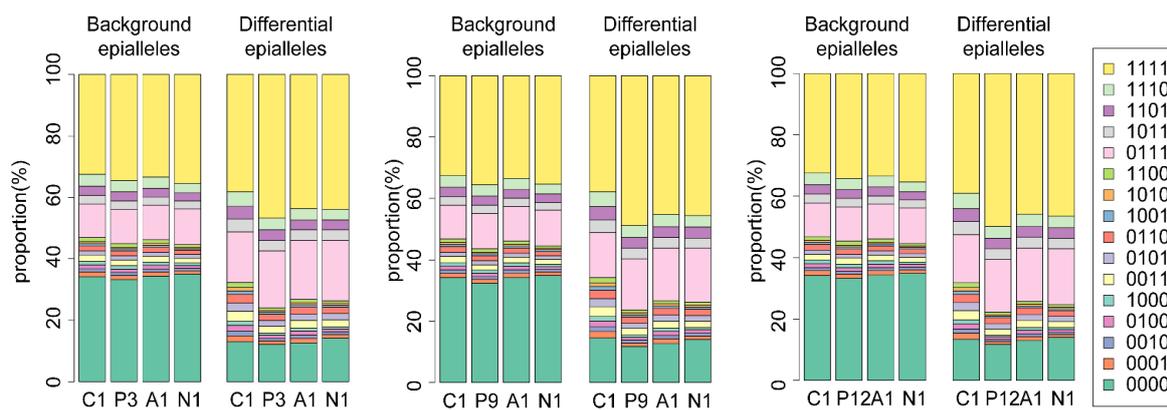
<sup>5</sup> The Fourth Department of Medical Oncology, Harbin Medical University Cancer Hospital, 150 Haping Road, Harbin, 150040, China

✉Corresponding authors: Yan Zhang, Email: zhangtyo@hit.edu.cn; Baoqing Sun, Email: sunbaoqing@vip.163.com; Shouping Xu, Email: shoupingxu@hrbmu.edu.cn

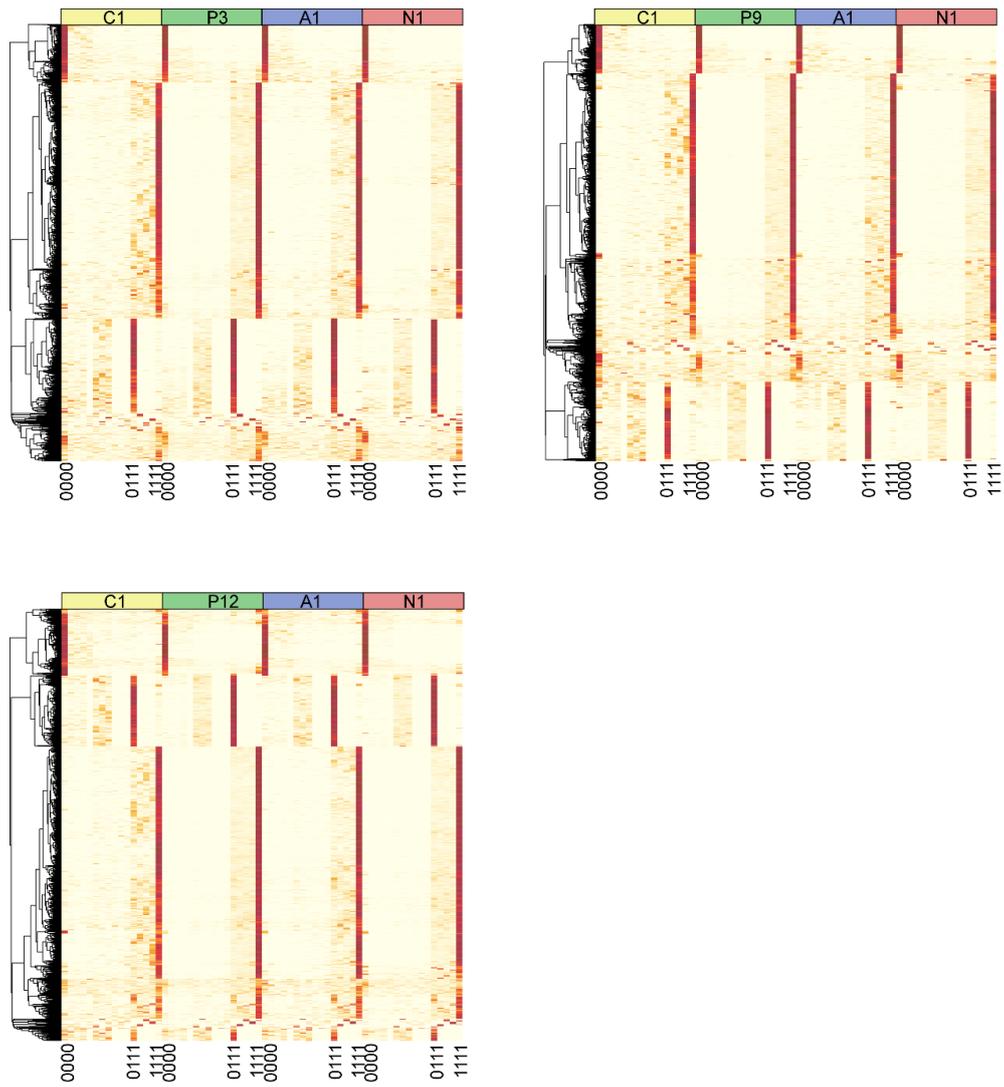
## Supplementary Figures



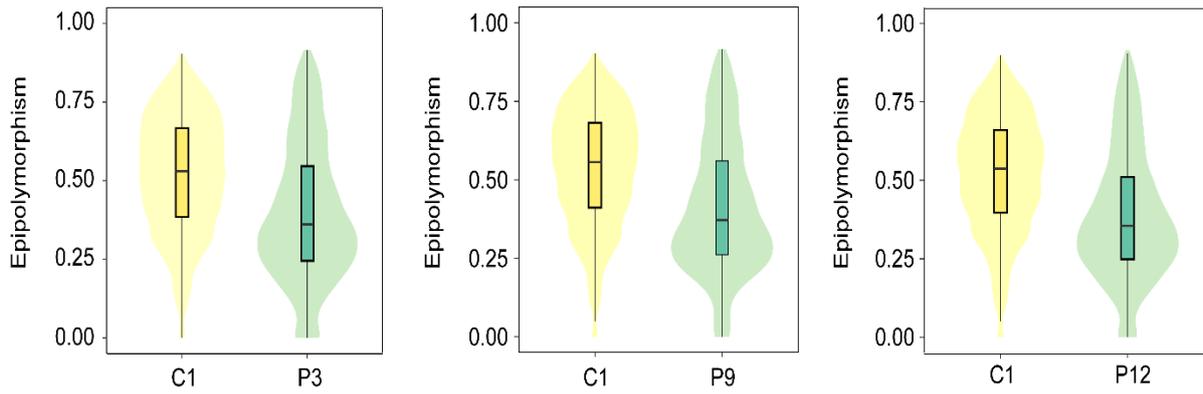
**Figure S1** The distribution of JSD and the determination of threshold for identifying differential epialleles between tumor periphery and tumor periphery.



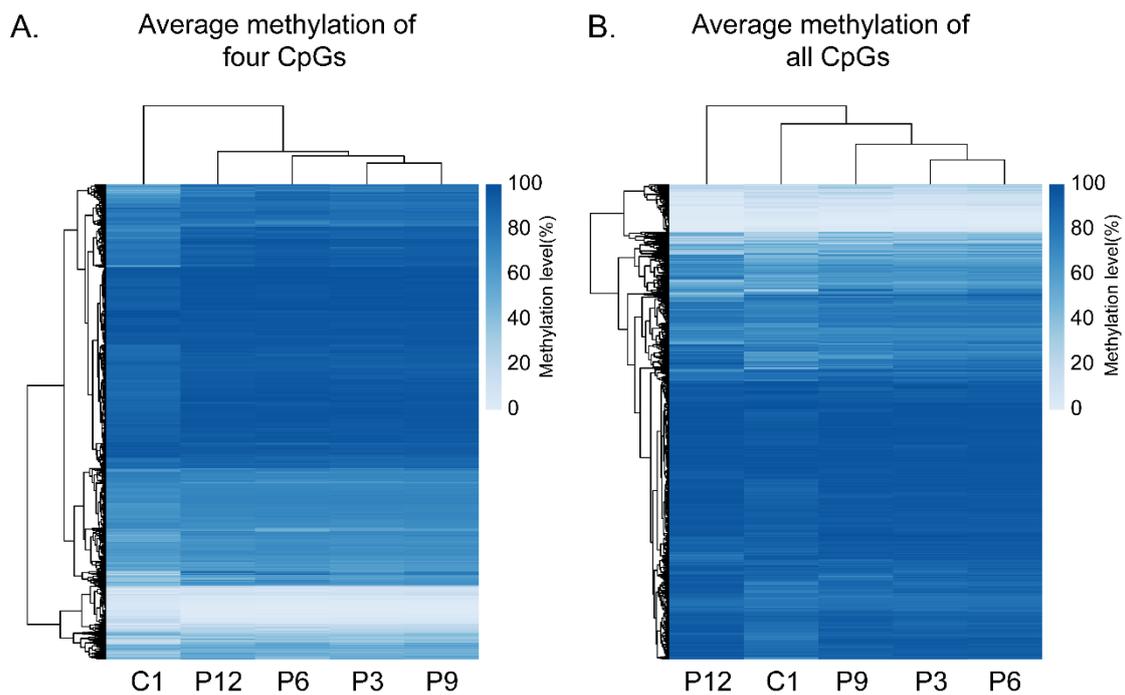
**Figure S2** The compositional changes of CPDEs identified between C1 and P3, P9, P12.



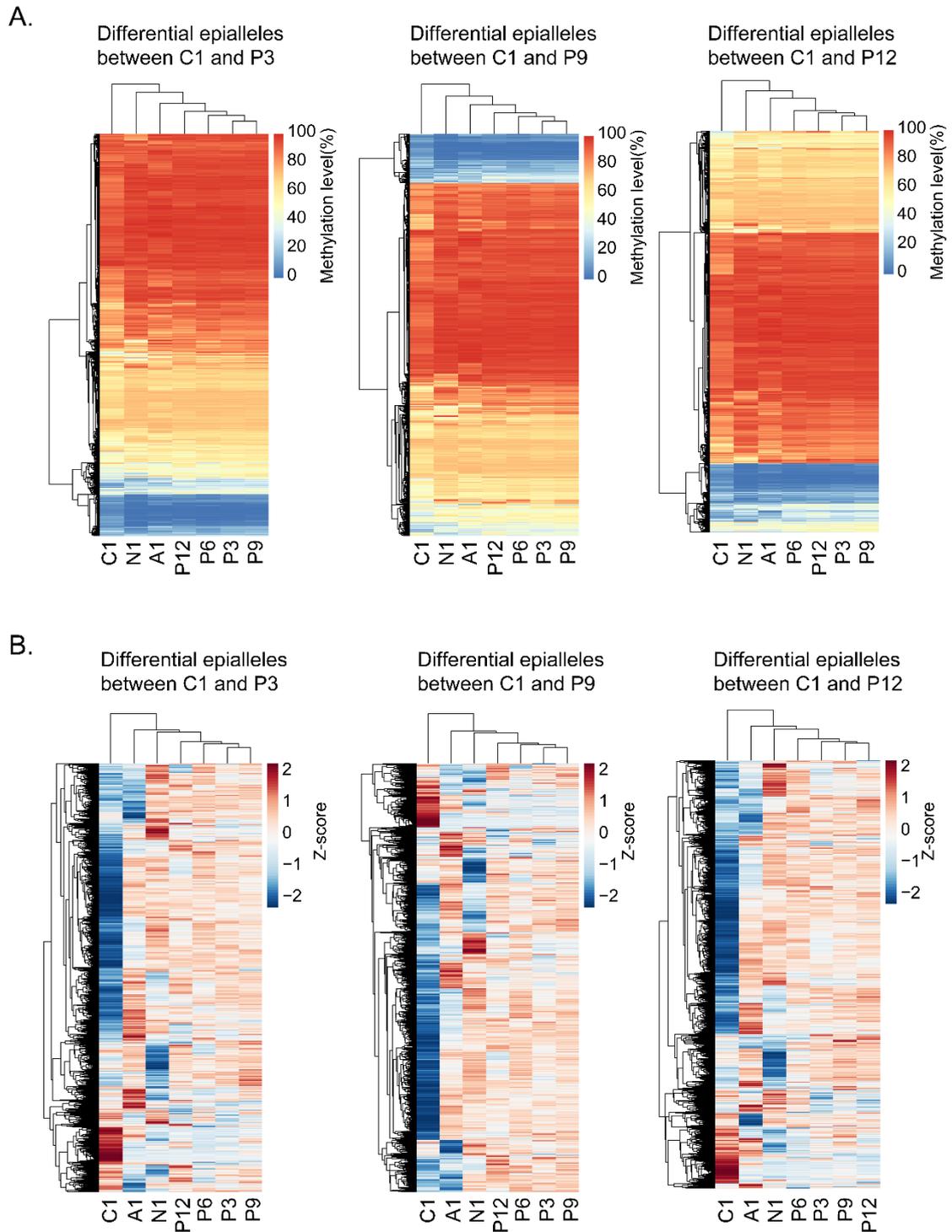
**Figure S3** The compositional changes in the core sample (C1), periphery sample (P3/P9/P12), adjacent sample (A1) and normal sample (N1) for the corresponding CPDEs.



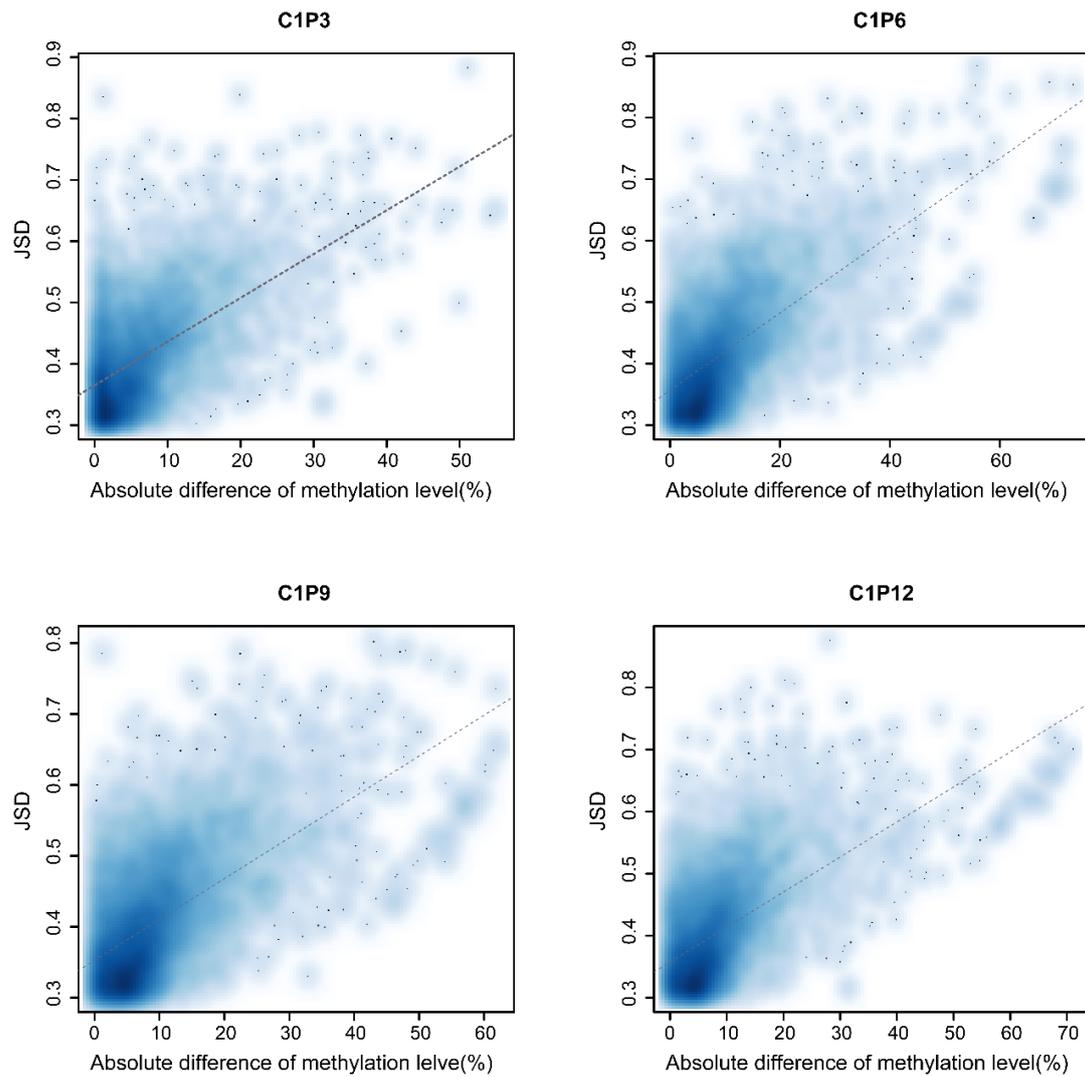
**Figure S4** The epipolymorphism in core sample (C1) and periphery samples (P3/P9/P12) for the corresponding CPDEs (Wilcoxon test,  $p < 0.01$ ).



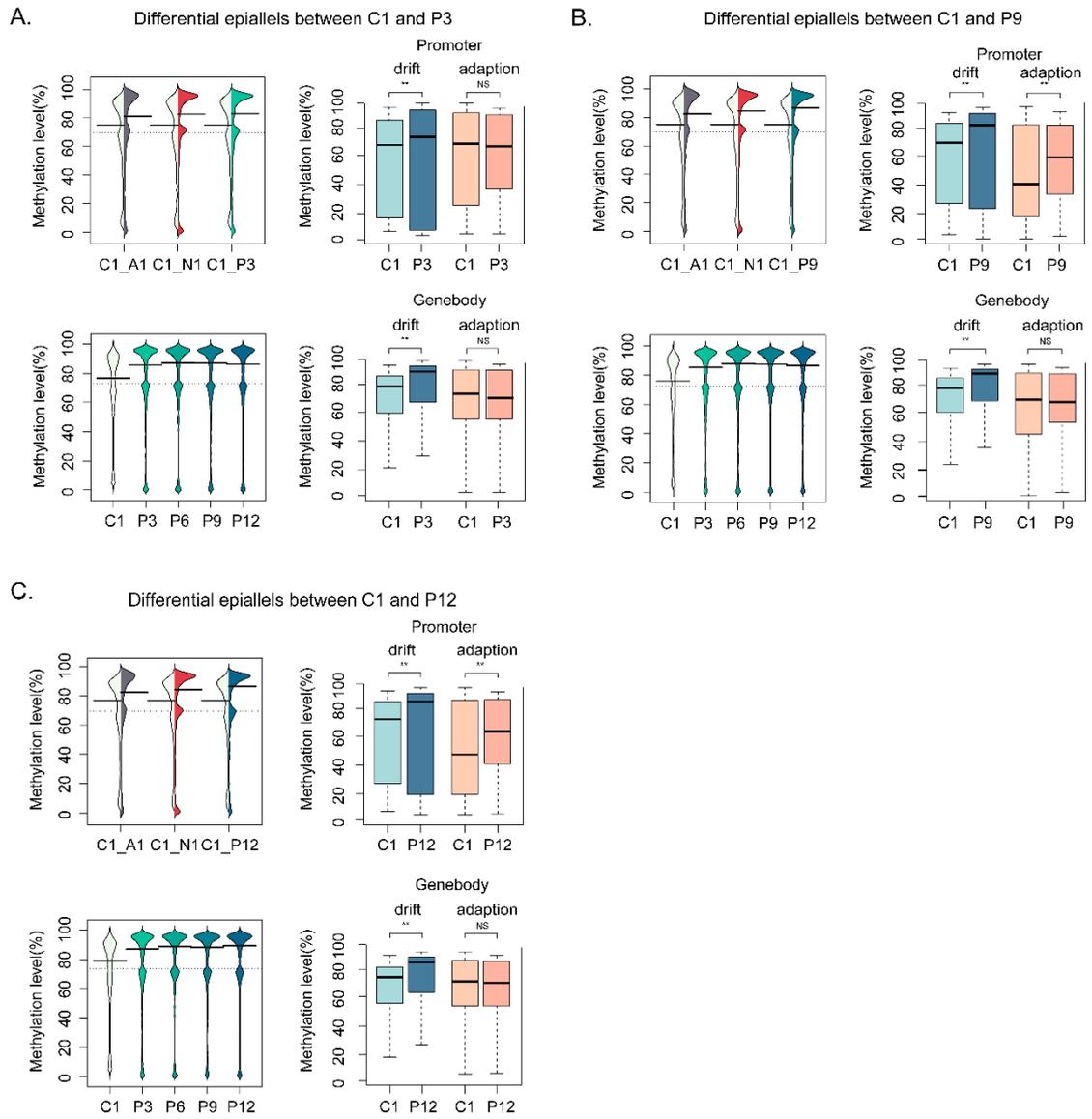
**Figure S5** Comparison of two methods for calculating methylation level of epialleles. (A) The heatmap of CPDEs based on average methylation of four CpG sites within the epiallele. (B) The heatmap of CPDEs based on average methylation of all CpG sites within the epiallele.



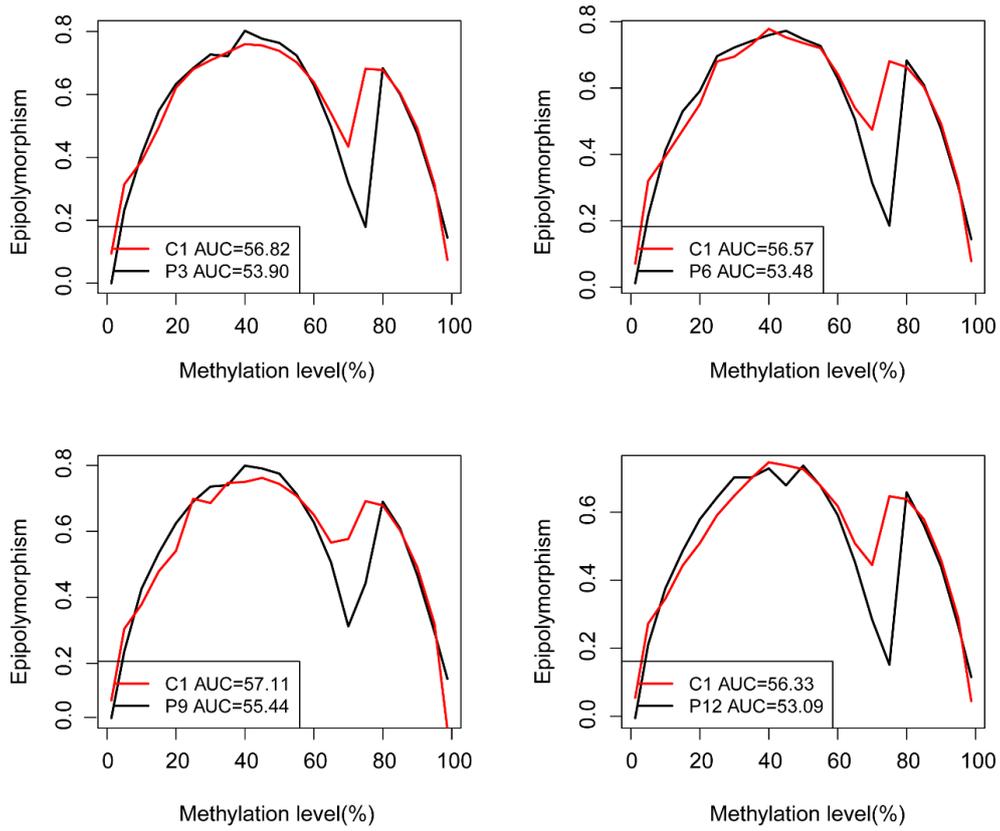
**Figure S6** The heatmaps of differential epialleles. (A) The heatmaps of raw methylation level for CPDEs in C1P3, C1P9 and C1P12. (B) The heatmaps of Z-score methylation level for CPDEs in C1P3, C1P9 and C1P12.



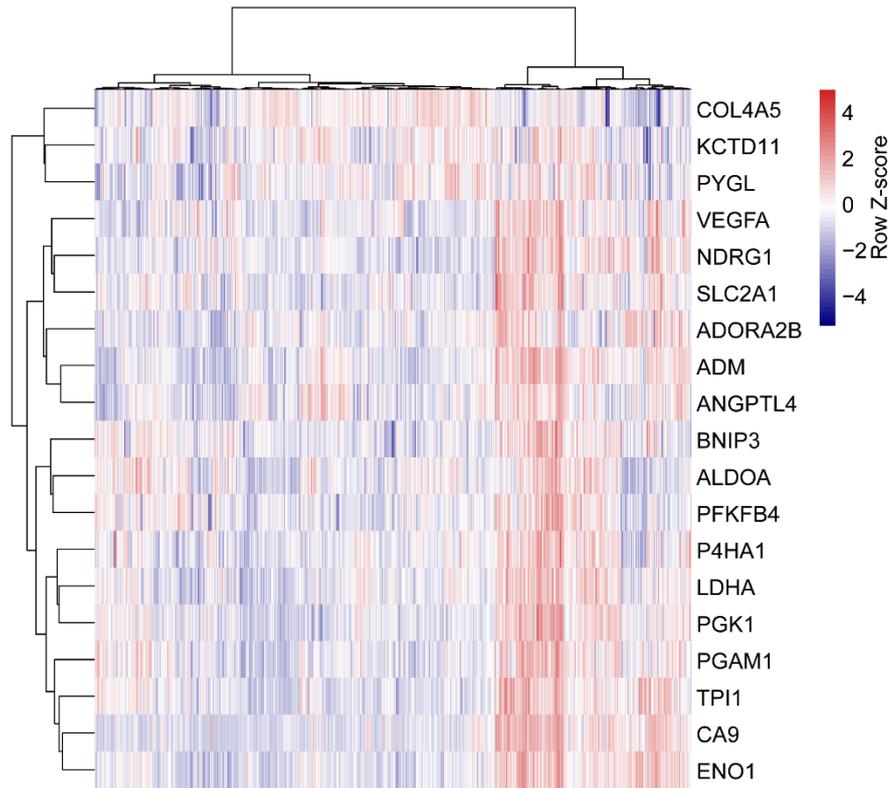
**Figure S7** The correlation between JSD and absolute differences of methylation level.



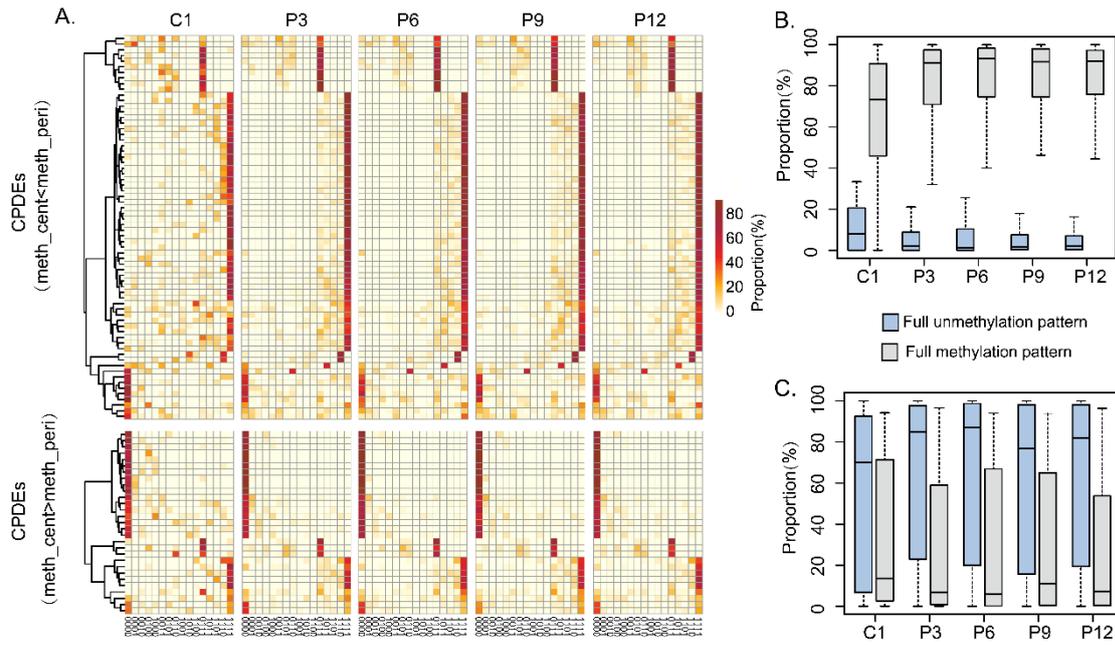
**Figure S8** The methylation-based analysis for CPDEs.



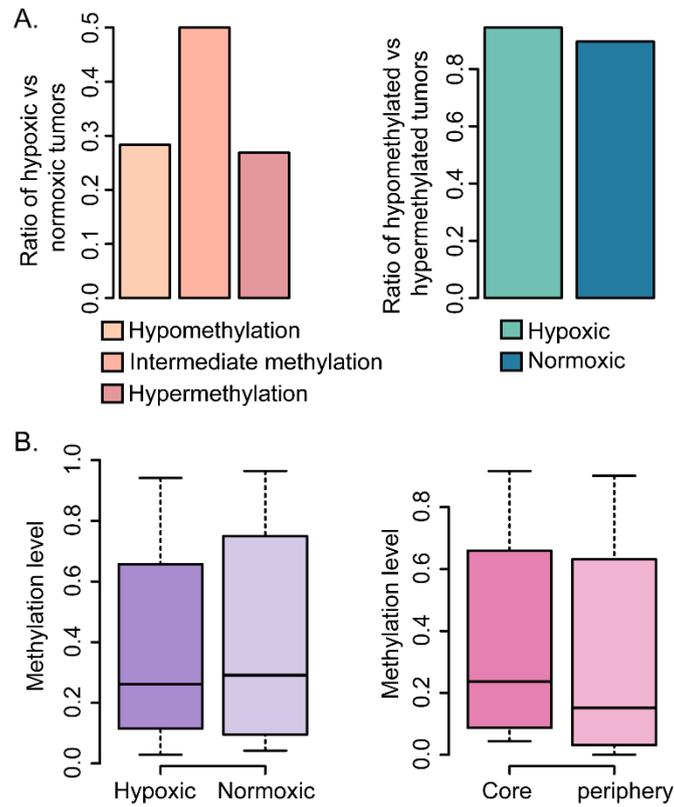
**Figure S9** The AUC of methylation heterogeneity in the core sample and the periphery samples.



**Figure S10** The hierarchical clustering for TCGA BRCA patients based on gene expression of hypoxia signatures.



**Figure S11** The methylation patterns of 97 CPDEs mapped to 450K CpG probes. (A) The heatmap of methylation pattern of 68 CPDEs hypomethylated in tumor core than periphery, and 29 CPDEs hypermethylated in tumor core than periphery. (B) The proportions of full unmethylation pattern ('0000') and full methylation pattern ('1111') in the two classes of CPDEs.



**Figure S12** Effect of hypoxia on methylation in CPDEs with lower methylation in tumor core than tumor periphery. (A) Ratios of hypoxic tumors vs normoxic tumors in the hypomethylation, intermediate methylation and hypermethylation groups (Cochran-Armitage trend chi-square test,  $p=0.805$ ), and ratios of hypomethylated tumors vs hypermethylated tumor in the hypoxic and normoxic clusters (chi-square test,  $p=1$ ). (B) Boxplots of methylation level in hypoxic and normoxic clusters, as well as tumor core and tumor periphery.

## Supplementary Tables

**Table S1** The information of RRBS data for seven samples.

| Sample             | C1         | P3         | P6         | P9          | P12         | A1         | N1         |
|--------------------|------------|------------|------------|-------------|-------------|------------|------------|
| Raw reads          | 58,822,682 | 56,023,350 | 50,470,280 | 109,323,286 | 141,953,654 | 68,383,106 | 56,725,878 |
| Clean reads        | 57,623,782 | 42,174,314 | 42,173,590 | 95,971,184  | 86,778,070  | 67,195,330 | 52,397,042 |
| Depth              | 18.41727   | 18.85564   | 22.8961    | 29.95219    | 43.93936    | 20.03477   | 21.15825   |
| CpGs               | 9,134,184  | 7,318,198  | 6,480,995  | 10,978,312  | 7,981,636   | 9,754,366  | 8,019,834  |
| Mapping efficiency | 74.9%      | 72.4%      | 71.5%      | 76.7%       | 73.4%       | 71.0%      | 75.2%      |

**Table S2** The information of RNA-seq data for seven samples.

| Sample       | C1         | P3         | P6         | P9         | P12        | A1         | N1         |
|--------------|------------|------------|------------|------------|------------|------------|------------|
| Reads        | 40,407,794 | 63,768,476 | 90,734,586 | 73,727,062 | 53,847,362 | 46,418,336 | 43,473,922 |
| Mapping Rate | 84%        | 81.6%      | 66.3%      | 72.2%      | 82.5%      | 87.7%      | 89.1%      |

**Table S3** The number of each class of TCGA BRCA samples.

|                          | Hypoxic | Normoxic |
|--------------------------|---------|----------|
| Hypermethylation         | 32      | 133      |
| Intermediate methylation | 77      | 197      |
| Hypomethylation          | 100     | 145      |

**Table S4** The CPDEs with lower methylation level in tumor core than tumor periphery.

| chr | start     | end       | epiallele                               | CpG        |
|-----|-----------|-----------|---|------------|
| 16  | 85485722  | 85485786  | 85485722:85485745:85485764:85485786     | cg05905731 |
| 2   | 128350569 | 128350595 | 128350569:128350571:128350589:128350595 | cg01577298 |
| X   | 153657332 | 153657363 | 153657332:153657335:153657357:153657363 | cg04334684 |
| 15  | 67073416  | 67073442  | 67073416:67073426:67073433:67073442     | cg03925327 |
| 13  | 114004028 | 114004140 | 114004028:114004039:114004072:114004140 | cg15828353 |
| X   | 153657335 | 153657412 | 153657335:153657357:153657363:153657412 | cg04334684 |
| X   | 153657335 | 153657412 | 153657335:153657357:153657363:153657412 | cg05872808 |
| 11  | 78387281  | 78387323  | 78387281:78387291:78387311:78387323     | cg15006118 |
| 5   | 149546074 | 149546104 | 149546074:149546088:149546099:149546104 | cg25132276 |
| 11  | 719705    | 719729    | 719705:719711:719713:719729             | cg17469934 |
| 10  | 1205650   | 1205691   | 1205650:1205652:1205668:1205691         | cg15057747 |
| 17  | 77901103  | 77901148  | 77901103:77901130:77901144:77901148     | cg11339839 |
| 20  | 49621249  | 49621314  | 49621249:49621297:49621306:49621314     | cg04524933 |
| 17  | 73096446  | 73096503  | 73096446:73096456:73096476:73096503     | cg21853806 |
| 17  | 73096446  | 73096503  | 73096446:73096456:73096476:73096503     | cg24328816 |
| 6   | 168686236 | 168686300 | 168686236:168686247:168686270:168686300 | cg05571645 |
| 6   | 168686236 | 168686300 | 168686236:168686247:168686270:168686300 | cg21701043 |
| 22  | 30950920  | 30950958  | 30950920:30950930:30950934:30950958     | cg22934785 |
| 1   | 2108696   | 2108749   | 2108696:2108706:2108730:2108749         | cg02334669 |
| 8   | 144552717 | 144552771 | 144552717:144552730:144552736:144552771 | cg27413290 |
| 2   | 241827847 | 241827876 | 241827847:241827852:241827864:241827876 | cg24075745 |
| 10  | 1205538   | 1205572   | 1205538:1205563:1205565:1205572         | cg09426810 |
| 10  | 1205538   | 1205572   | 1205538:1205563:1205565:1205572         | cg26293457 |
| 2   | 75720531  | 75720567  | 75720531:75720540:75720546:75720567     | cg00409636 |
| 7   | 813677    | 813751    | 813677:813704:813709:813751             | cg26465987 |
| 7   | 813704    | 813764    | 813704:813709:813751:813764             | cg26465987 |
| 20  | 60969897  | 60969964  | 60969897:60969906:60969929:60969964     | cg23531748 |
| 5   | 2749647   | 2749715   | 2749647:2749649:2749655:2749715         | cg07511080 |
| 12  | 132834295 | 132834340 | 132834295:132834315:132834336:132834340 | cg04774461 |
| 4   | 24801110  | 24801131  | 24801110:24801120:24801125:24801131     | cg10766585 |
| 21  | 15135117  | 15135146  | 15135117:15135134:15135142:15135146     | cg11939992 |
| 16  | 30616268  | 30616334  | 30616268:30616299:30616309:30616334     | cg05045702 |
| 16  | 88738398  | 88738418  | 88738398:88738410:88738413:88738418     | cg03682568 |
| 16  | 88738398  | 88738418  | 88738398:88738410:88738413:88738418     | cg04176488 |
| 12  | 131519904 | 131519996 | 131519904:131519927:131519993:131519996 | cg13586425 |
| 16  | 88018664  | 88018712  | 88018664:88018667:88018681:88018712     | cg26605774 |
| 20  | 60969885  | 60969929  | 60969885:60969897:60969906:60969929     | cg23531748 |
| 6   | 168686216 | 168686270 | 168686216:168686236:168686247:168686270 | cg21701043 |
| X   | 132550050 | 132550132 | 132550050:132550080:132550086:132550132 | cg12600766 |
| 19  | 7677731   | 7677748   | 7677731:7677733:7677740:7677748         | cg02076378 |
| 22  | 30950923  | 30950947  | 30950923:30950930:30950938:30950947     | cg22934785 |
| 19  | 4544068   | 4544095   | 4544068:4544080:4544082:4544095         | cg04848343 |

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|    |           |           |   |            |
|----|-----------|-----------|---|------------|
| 18 | 34340652  | 34340700  | 34340652:34340681:34340691:34340700     | cg23695714 |
| 19 | 7584666   | 7584698   | 7584666:7584674:7584680:7584698         | cg01773515 |
| 1  | 161576928 | 161576989 | 161576928:161576958:161576964:161576989 | cg26022242 |
| 13 | 19412134  | 19412226  | 19412134:19412185:19412222:19412226     | cg08120511 |
| 7  | 23695224  | 23695277  | 23695224:23695247:23695263:23695277     | cg15781397 |
| 17 | 73090153  | 73090195  | 73090153:73090163:73090186:73090195     | cg15190451 |
| 20 | 49626332  | 49626374  | 49626332:49626346:49626348:49626374     | cg03711791 |
| X  | 136648305 | 136648319 | 136648305:136648308:136648315:136648319 | cg22934939 |
| 11 | 67261122  | 67261175  | 67261122:67261166:67261172:67261175     | cg17028218 |
| X  | 100306753 | 100306832 | 100306753:100306781:100306813:100306832 | cg13286902 |
| X  | 100306753 | 100306832 | 100306753:100306781:100306813:100306832 | cg15891447 |
| 11 | 490498    | 490558    | 490498:490525:490552:490558             | cg11004323 |
| 16 | 87873347  | 87873423  | 87873347:87873354:87873369:87873423     | cg10169763 |
| 18 | 34340640  | 34340681  | 34340640:34340642:34340652:34340681     | cg23695714 |
| 22 | 30950930  | 30950961  | 30950930:30950934:30950958:30950961     | cg22934785 |
| 2  | 242608073 | 242608094 | 242608073:242608075:242608087:242608094 | cg06006530 |
| 2  | 242608073 | 242608094 | 242608073:242608075:242608087:242608094 | cg08462942 |
| 17 | 62066571  | 62066673  | 62066571:62066580:62066597:62066673     | cg08387141 |
| 15 | 40751007  | 40751064  | 40751007:40751013:40751016:40751064     | cg02479871 |
| 19 | 39217718  | 39217766  | 39217718:39217722:39217738:39217766     | cg25383568 |
| 6  | 168686247 | 168686309 | 168686247:168686270:168686300:168686309 | cg05571645 |
| 20 | 36532087  | 36532097  | 36532087:36532089:36532091:36532097     | cg11599505 |
| 1  | 5934887   | 5934950   | 5934887:5934913:5934941:5934950         | cg09976670 |
| 3  | 114070427 | 114070444 | 114070427:114070434:114070441:114070444 | cg19622597 |
| 1  | 2108706   | 2108769   | 2108706:2108730:2108749:2108769         | cg02334669 |
| 17 | 77901150  | 77901227  | 77901150:77901179:77901224:77901227     | cg05483571 |
| 7  | 98609840  | 98609873  | 98609840:98609846:98609859:98609873     | cg04212699 |
| 12 | 131519890 | 131519927 | 131519890:131519899:131519904:131519927 | cg13586425 |
| 11 | 68170975  | 68171011  | 68170975:68170999:68171003:68171011     | cg25380755 |
| 2  | 242608075 | 242608104 | 242608075:242608087:242608094:242608104 | cg06006530 |
| 2  | 242608075 | 242608104 | 242608075:242608087:242608094:242608104 | cg08462942 |
| 19 | 4544080   | 4544105   | 4544080:4544082:4544095:4544105         | cg04848343 |
| 16 | 88018667  | 88018719  | 88018667:88018681:88018712:88018719     | cg26605774 |
| 16 | 10274064  | 10274089  | 10274064:10274070:10274086:10274089     | cg10116505 |

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