## 1 **OPENANNO: annotating genomic regions with chromatin**

### 2 accessibility

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#### 17 Abstract

18 Chromatin accessibility, as a powerful marker of active DNA regulatory elements, 19 provides rich information to understand the regulatory mechanism. The revolution in 20 high-throughput methods has accumulated massive chromatin accessibility profiles in 21 public repositories as a valuable resource for machine learning and integrative studies. 22 Nevertheless, utilization of these data is often hampered by the cumbersome and time-23 consuming collection, processing, and annotation of the chromatin accessibility 24 information. Motivated by the above understanding, we developed a web server, named 25 OPENANNO, to annotate the openness of genomic regions across diverse cell lines, 26 tissues, and systems. The annotation is based on 871 DNase-seq experiments across 27 199 cell lines, 48 tissues, and 11 systems from ENCODE, and openness values 28 rigorously defined by four statistical strategies. Particularly, we designed a parallel 29 program to allow efficient annotation and visualization of the openness of a vast amount 30 of genomic regions. OPENANNO will help users extract and download formulated data in a batch follow-up analysis. Besides, we illustrate the valuable information provided 31 32 by OPENANNO using an enhancer of blood vessels from VISTA Enhancer Browser 33 as an example. Furthermore, we demonstrate three applications of OPENANNO in 34 regulatory mechanism and association studies. We believe that OPENANNO will serve 35 as a comprehensive and user-friendly web server to facilitate methodology 36 development and biological insights discovery, specifically to explore the biological questions and model the regulatory landscape of genome. OPENANNO is freely 37 38 available at http://bioinfo.au.tsinghua.edu.cn/openness/anno or 39 http://159.226.47.242:65424/openness/anno/.

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41 KEYWORDS: Chromatin accessibility; Openness; Annotation; Visualization; Web
42 server

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#### 44 Introduction

45 The era of the personal genome is arriving with the widespread sequencing technologies 46 and the ultimate promise for precision medicine. However, it remains distant in 47 interpreting the context of variations in non-coding DNA sequence associated with 48 disease and other phenotypes, deciphering their biological functions in gene regulation, 49 and further understanding disease mechanism and dynamic response to treatment [1, 2]. 50 Chromatin accessibility is a measure of the ability of nuclear macromolecules to 51 physically contact DNA [3], and plays the role of a powerful marker of active regulatory 52 genomic regions, which have a wide range of effects on the transcription, DNA repair, 53 recombination, and replication [4, 5]. The recent revolution in high-throughput, 54 genome-wide methods invented several biological assays for extracting open chromatin, 55 such as DNase-seq (deoxyribonuclease), FAIRE-seq (formaldehyde-assisted isolation 56 of regulatory elements), ATAC-seq (assay for transposase-accessible chromatin) and 57 MNase-seq (micrococcal nuclease), and thus open a new door for us to make extensive use of chromatin accessibility [6-11]. For example, accessible genomic regions are 58 59 regarded as the primary positions of regulatory elements [12], and thus provide a great 60 opportunity to study transcription factor binding sites, DNA methylation sites, histone 61 modification markers, gene regulation, and regulatory network [13, 14]. In addition, 62 changes in chromatin accessibility have been implicated with different perspectives of 63 human health as a result of the alterations of nucleosome positioning affected by 64 mutations in chromatin remodelers [15-17].

65 The development of high-throughput sequencing techniques has accumulated a vast 66 amount of chromatin profiles across a variety of cell lines. Large collaborative projects, 67 such as Encyclopedia of DNA Elements (ENCODE) [18], have become a part of the major effort. The Roadmap Epigenomics project provides another similar resource for 68 69 human stem cells and tissues [19]. Nevertheless, many experimental biologists may 70 lack the bioinformatics expertise to make full use of these valuable resources efficiently. 71 Cistrome DB, a data portal for ChIP-Seq and chromatin accessibility data, although 72 comprises species, factors, biological source, publication, and other information for 73 their collected ChIP-seq and DNase-seq data [20], limited to containing only a part of 74 currently available transcription factors and histone marks. Therefore, it is still very 75 cumbersome and time-consuming to collect, process, and incorporate the chromatin 76 accessibility information of arbitrary genomic regions into bioinformatics and

epigenetics studies, which thus makes it difficult to full use of the vast amount ofchromatin profiles.

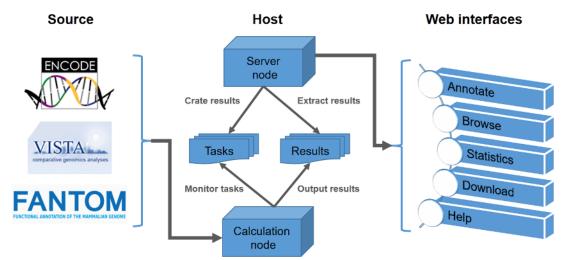
79 We noticed that the epigenome consists of signals from chemical modifications of 80 histones, DNA methylation, non-coding RNA expression, and transcription factors that 81 work in concert to determine the accessibility of the regulatory regions, so-called open 82 region. Then the open regulatory regions can work together with transcription factors, 83 RNA polymerases, and other cellular regulatory machines and produce the final gene 84 expression pattern. In this sense, the chromatin 'openness', *i.e.*, the accessibility of 85 genomic regions, bridges the epigenome and transcriptome and plays an important role 86 in understanding the regulatory mechanism. Motivated by the above demand, we built 87 a web server, named OPENANNO, to **anno**tate the **open**ness of genomic regions across 88 diverse types of cell lines, tissues, and systems. We downloaded the raw sequencing 89 data of 871 DNase-seq experiments across 199 cell lines, 48 tissues and 11 systems 90 from ENCODE data portal [18] and processed by a uniform pipeline. We defined the 91 openness of genomic regions by four statistical strategies, including foreground read 92 count, raw read openness, narrow peak openness, and broad peak openness. 93 Furthermore, we designed a parallel program to enable OPENANNO to efficiently 94 annotate and visualize the openness of a vast amount of genomic regions. We finally 95 demonstrate three applications of OPENANNO in regulatory mechanism and 96 association studies. We believe that this web server will help both the computational 97 and experimental community to facilitate developing methods and discovering 98 important insights, explore the basic biology and various applications, and open a new 99 door to model the regulatory landscape of genome.

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#### 101 Web server content and usage

#### 102 Overall design of OPENANNO

As illustrated in **Figure 1**, the diagram for constructing the OPENANNO web server consists of three main parts, *i.e.*, the source, host, and web interface. In the source part, we deposited the meta-information, raw data, and uniformly processed data of 871 human DNase-seq experiments from the ENCODE project [18]. It also includes the datasets of regulatory elements that have been experimentally validated from FANTOM[21] and VISTA[22]. The second part, calculation node in the host, monitors the file path that contains annotation tasks, and will automatically calculate and store 110 the results in a specific file path once the server node creates a new task. The server 111 node bridges the host and web interfaces by achieving tasks from users and extracting 112 results or other information for visualization and downloading. The third part, web 113 interface, was developed in a concise and easy-to-use mode. The 'Annotate' page 114 provides a service to annotate the openness of a vast amount of genomic regions, and is built on a specially designed infrastructure to extract and visualize big table. The 115 116 'Browse' page enables users to study the openness of a particular genomic region more 117 intuitively. The 'Statistics' page provides detailed information for all the 871 DNase-118 seq experiments, and an intuitive comparison of the number of experiments in different 119 cell lines, tissues or biological systems. The 'Download' page endows users with the 120 ability to directly download the openness of collected experimentally validated 121 regulatory elements. The 'Help' page provides other commonly used information to 122 improve the usability of the web server.



123

124 Figure 1 The diagram for constructing the OPENANNO web server

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#### 126 Web interfaces for annotating

The 'Annotate' page, as the major site of OPENANNO, can annotate the openness of 127 128 genomic regions in batches. As illustrated in **Figure 2**, there are five major steps for 129 this workflow. First of all, we provide a concise task submission approach, which 130 avoids the confusion caused by redundant information to users. By clicking the 'Browse' 131 button, users can upload a bed or bed.gz file (uncompressed or compressed in gzip 132 format, e.g., ENCFF001WKF.bed.gz). The parallel program in our calculation node will extract the first three columns and the sixth column (the chromosomes, starting 133 134 sites, terminating sites and strands, respectively) separated by tabs for calculating the

135 openness. Note that sorting input in advance using other toolkits, such as bedtools 136 (https://bedtools.readthedocs.io), is preferred for speeding up the calculation. In the 137 current web server release, we provide the service to annotate the openness of genomic 138 regions in human reference genome GRCh37 (hg19). We will provide the option of 139 other genomes or species in future releases. Users can choose to calculate the openness 140 of these genomic regions in a particular cell line, or directly calculate the openness in 141 all the 871 cell lines. After saving the data as local files, the users can compare the 142 openness in different cell lines or perform advanced analysis such as the co-openness 143 analysis, which will be demoed in the following section. Furthermore, users can enable 144 the option of Per-base pair annotation to calculate the openness of each base-pair of the 145 genomic regions for some bioinformatics analysis such as machine learning tasks which 146 will be demoed in the following section.

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   | 885 12.13<br>6.50<br>8.285<br>5.07  | INFECTION<br>INTERNATIONAL<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTION<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>INFECTIONI<br>IN  | 10.11<br>9.81<br>0.03  | 61000<br>Rizos<br>Heart<br>434WEP<br>11,70<br>8,51<br>3,19   
   | 221  |
| Covenload Char<br>Char<br>1 10<br>2 10<br>3 10<br>4 10<br>5 10   | Accestate     Accestate   | Drawse<br>Brawse<br>Brawse<br>Box T<br>End<br>120250<br>122570  | 2 program<br>(00) started.<br>(01) started.<br>(01) started.<br>(01) started.<br>(01) 97 45.<br>(01) 97 45.<br>(01) 97 45.<br>(01) 97 45.<br>(01) 97 45.<br>(01) 98.85.<br>(01) 0.05.  | ColorOn •<br>726008<br>14.28<br>9.71  | New Ta<br>2956000<br>14.18<br>21.53  
  | (BOBQU<br>13.01<br>18.88<br>0.00<br>2.28<br>20.82  | 2.05   | 12.50  
   | 9.51<br>15.38  | 16.46<br>7.90   | CO1134:<br>4.59<br>0.70<br>44.54<br>1.55  | 00115AL<br>4.15<br>1.40<br>0.72<br>2.722<br>0.02   
  | CB51572740<br>ICB5397264C2<br>ICB5397264C<br>ICB5397264C<br>ICB5397264C<br>ICB5397264<br>ICB5397260<br>20.02<br>16.38  | ENCSR0011<br>ENCSR0001<br>ENCSR0001<br>940K00<br>22.14<br>7.03  | чое<br>кол<br>зин<br>эслясх<br>7.23<br>8.83   
  | EF0:0002322<br>EF0:0002332<br>CL10002518<br>E5004/0*<br>0.04<br>4.34   | NUY<br>RPA<br>C<br>Hbr  | 108220<br>entrace<br>and ac<br>obliant<br>B827860<br>0.01<br>8.03   | 10.11   
  | elood<br>Diood<br>Heart<br>434WeP<br>11.70<br>8.51   | 224  |
| Church Ch | Chit     Ihr     Thit     Th     | Elizing thicked loader           ead loader           ad writer           Brawse           Bra           120250           122570           17400           19940           19940  | Electrics Deve<br>Charles Constraints<br>Charles Constr   | ColorOn •<br>7260028<br>14.28<br>0.71<br>1.29<br>15.14<br>13.99<br>5.71   | New Ta<br>2555KVW<br>14.18<br>21.53<br>2.05<br>3.23<br>10.61<br>26.95   | (50800)<br>13.01<br>18.98<br>0.00<br>2.28<br>20.92<br>20.92   
  | 2.05<br>5.48<br>10.93<br>0.88<br>5.45<br>17.08   | 12.50<br>10.51<br>10.67<br>0.52<br>14.39<br>21.90  | 9.51<br>15.38<br>0.73<br>22.67<br>28.26<br>10.97   | 16.46<br>7.90<br>0.00<br>12.61<br>24.06<br>20.41  
   | ENCHE<br>DICITI<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCER<br>ENCEN    | BUTISAL<br>4.15<br>1.40<br>2722<br>0.00<br>6.23   | CB915/2H0<br>CD5204K7U<br>CD5204K7U<br>CD539/2EHC<br>CD539/2EHC<br>2002<br>10.38<br>2.20<br>2.73<br>4.5 Ch<br>2.73<br>4.5 Ch<br>2.048   
  | ENCSR0011<br>D4CSR0004<br>ENCSR0004<br>940R00<br>22.14<br>7.03<br>2.07<br>1.96<br>58 12<br>28.67  | 500<br>344<br>3500<br>844<br>3500<br>840<br>840<br>840<br>241<br>3774<br>15.66   | 50000000000000000000000000000000000000   | BIS 12.13<br>BIS 12.13<br>0.00<br>0.28<br>5.37<br>2.68<br>14.76<br>7.16  
  | 102220<br>102220<br>ardiac<br>obliant<br>80210141<br>0.31<br>8.03<br>0.47<br>4.265<br>16.69   | 10.11<br>9.81<br>0.00<br>8.07<br>17.70<br>18.16  | elood<br>Ricod<br>Hoart<br>434We9 <sup>2</sup><br>11,70<br>8,51<br>3,19<br>5,85<br>0,743<br>17,56   
  | 224  |
| Commission         E           0         Chrone           1         10           2         10           3         10           4         10           5         10           6         10           7         10           8         10  | Chit     Ihr     Thi     Th      | Zilzing the           End           Braves           Braves           Braves           120200           120200           174010           179590           101410   | Engine   | ColsrOn •<br>7200008<br>14.28<br>0.71<br>1.29<br>15.14<br>13.98<br>5.71<br>14.99<br>1.14  | New Ta<br>255KVW<br>14.18<br>21.53<br>2.05<br>3.23<br>10.61   
   | (BCBQU<br>13.01<br>18.88<br>0.00<br>2.28<br>20.82  | 2.05<br>5.48<br>10.93<br>0.68<br>5.45  | 12.50<br>10.51<br>10.67<br>0.62<br>14.38<br>21.90<br>10.77<br>0.99  
  | 9.51<br>15.38<br>0.75<br>22.67<br>28.25  | 16.46<br>7.90<br>0.00<br>12.61<br>24.06   | CO1134:<br>4.59<br>0.70<br>44.54<br>1.55  | 00115AL<br>4.15<br>1.40<br>0.72<br>2.722<br>0.02  
   | CB915/2NO<br>ICD8204KRU<br>ICD839/2EHC<br>ICD839/2EHC<br>ICD839/2EHC<br>ICD839/2EHC<br>ICD839/2EHC<br>ICD839/2EHC<br>ICD839/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2EHC<br>ICD829/2E   | ENCSR0011<br>ENCSR0000<br>ENCSR0000<br>9402KUD<br>22.14<br>7.02<br>2.07<br>1.55<br>29.12  | 9000<br>344<br>9000444<br>7,23<br>8,83<br>4,02<br>2,41<br>37,74  
   | EF010002322<br>EF010002322<br>CL10002518<br>2500440<br>0.04<br>0.04<br>0.04<br>0.04<br>0.04<br>0.04<br>0.0   | BIS 1/2,11<br>BIS 1/2,12<br>BIS 1/2,12  | 102200<br>100336<br>ardiac<br>obliaet<br>0.01<br>8.03<br>0.47<br>4.25<br>14.65  | 10.11<br>9.81<br>0.03<br>8.07<br>17.70   | elood<br>Blood<br>Heart<br>434WEP<br>11.70<br>8.51<br>3.19<br>5.85<br>02.43<br>17.56<br>4.78<br>4.78<br>1.06  
  | 224  |
| 200000100000 [ f<br>Clause<br>2 10<br>2 10<br>2 10<br>2 10<br>2 10<br>3 10<br>5 10<br>6 10<br>7 10<br>6 10<br>7 10<br>6 10<br>7 10<br>7 10<br>8 10<br>9 10   | Control      Control     Contro     Contro     Control     Control     Control     Control     Co | Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne<br>Browne    | e program<br>(00) started.<br>(01) started.<br>(01) started.<br>(01) started.<br>(02) started.<br>(03) started.<br>(04) started.<br>(0 | ColarOn •<br>750008<br>14.28<br>0.71<br>1.29<br>15.14<br>13.98<br>5.71<br>14.09<br>1.14<br>2.57   | New Ta<br>2555KVW<br>14.18<br>21.53<br>2.05<br>3.23<br>10.61<br>26.95<br>7.25<br>0.15<br>0.25  
  | (50800<br>13.01<br>18.88<br>0.00<br>2.28<br>20.92<br>20.92<br>10.10<br>1.13<br>1.70  | 2.05<br>5.45<br>10.83<br>0.88<br>5.45<br>17.08<br>10.25<br>0.00<br>0.00<br>0.00  | 12.50<br>10.51<br>10.67<br>0.52<br>14.58<br>21.90<br>10.77<br>0.99<br>1.19   
   | 9.51<br>15.38<br>0.73<br>22.67<br>28.26<br>10.97<br>0.73<br>1.46<br>0.00   | 16.46<br>7.90<br>0.00<br>12.61<br>24.06<br>20.41<br>5.93<br>1.99<br>0.98  | ENCFR<br>ENCFR<br>ENCFR<br>ENCFR<br>ENCFR<br>ENCFR<br>44.59<br>0.00<br>0.76<br>44.54<br>1.55<br>2.34<br>4.69<br>4.59<br>0.20<br>0.76<br>44.54<br>0.234<br>4.69<br>0.78  | 60074H EN<br>8030/02 EN<br>8030/02 EN<br>8030/02 EN<br>8030<br>804<br>804<br>804<br>804<br>804<br>804<br>804<br>80   
  | CB915/200<br>ICB5204K2U<br>ICB5302EHC<br>2002<br>16.38<br>2.20<br>2.73<br>20.48<br>2.73<br>0.46<br>1.62  | ENCSR001<br>ENCSR001<br>ENCSR001<br>ENCSR001<br>22.14<br>7.03<br>2.07<br>1.06<br>2.8.15<br>2.8.67<br>4.30<br>0.00<br>0.05   | 900<br>900<br>900<br>900<br>900<br>800<br>800<br>800<br>800<br>800  
  | 5010012322<br>EF0:0002322<br>CL:0012518<br>501440<br>0.04<br>4.34<br>5.50<br>7.51<br>7.51<br>3.04<br>3.04<br>3.04<br>3.04<br>3.04  | BIS 17.13<br>BIS 17  | 102220<br>102230<br>ardiac<br>bilact<br>0.31<br>8.03<br>0.47<br>4.26<br>14.65<br>16.69<br>3.31<br>1.42<br>0.05  | 10.11<br>9.81<br>0.00<br>8.07<br>17.70<br>18.16<br>2.53<br>1.01<br>9.51  | 434WEP<br>11,70<br>6,51<br>3,19<br>5,243<br>17,56<br>4,79<br>1,06<br>3,10   
  | 224  |
| Commission         E           0         Chrone           1         10           2         10           3         10           4         10           5         10           6         10           7         10           8         10  | Init     Init    | Elizing the           ead loader           ad writer           Brazze           task           task           120200           120200           120200           120200           120200           120200           120200           120201           120202           101410           180200           180200           180200           180200   | Electrics Does<br>Child Started.<br>Child Started.   | ColarOn •<br>7200008<br>14.20<br>9.71<br>1.29<br>15.14<br>13.99<br>5.71<br>14.99<br>1.14<br>2.57<br>14.14<br>0.39   | New Ta<br>2555KVW<br>14.18<br>2.05<br>3.23<br>10.61<br>26.95<br>8.25<br>0.15  
   | (50800<br>13.01<br>18.88<br>0.00<br>2.28<br>20.92<br>20.92<br>10.10<br>1.13  | 2.05<br>5.45<br>10.93<br>0.88<br>5.45<br>17.08<br>10.25<br>0.00  | 12.50<br>10.51<br>10.67<br>0.62<br>14.38<br>21.90<br>10.77<br>0.99   | 9.51<br>15.38<br>0.73<br>22.67<br>28.26<br>10.97<br>0.73<br>1.46   
   | 16.46<br>7.90<br>0.00<br>12.61<br>24.06<br>20.41<br>5.93<br>1.98  | ENCHA<br>ENCIP<br>ENCIP<br>ENCIP<br>ENCIP<br>4.59<br>0.70<br>0.70<br>44.54<br>1.55<br>2.34<br>4.59<br>45.10   | 6000AH EN<br>A039/22 EN<br>1001DAA EN<br>1001DAA EN<br>1001DAA
EN<br>1001DAA<br>1102<br>2003<br>2003<br>1002<br>2003<br>1002<br>2003<br>1002<br>2003<br>2003<br>2003<br>2003<br>2003<br>2003<br>2003<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>2005<br>200   | CB915/2NO<br>ICD8204KRU<br>ICD839/2EHC<br>ICD839/2EHC<br>2002<br>10.38<br>2.20<br>2.78<br>45.60<br>2.78<br>45.60<br>2.78<br>45.60<br>2.78<br>45.60<br>2.78<br>45.60<br>2.74<br>3.046   | ENCSR0011<br>ENCSR0014<br>ENCSR0014<br>ENCSR0014<br>2014<br>2017<br>1.66<br>58 15<br>28 15<br>28 15<br>28 00<br>1.30<br>0.00  | 9006<br>9007<br>9144<br>9579623<br>729<br>8.83<br>4.02<br>2.41<br>37.74<br>15.85<br>4.02<br>2.41<br>3.651<br>4.61<br>0.60<br>2.41<br>3.241<br>0.60<br>2.41<br>0.60   
   | 5010002322<br>EPO-0002322<br>CLI0002518<br>5010002518<br>5010002518<br>5010002518<br>5010002518<br>5010002518<br>5010002518<br>5010002512<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002322<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002312<br>5010002310<br>5010002<br>5010002<br>5010002<br>501000000<br>5010000000000   | BIS 1/2,11<br>BIS 1/2,12<br>BIS 1/2,12  | 102220<br>rm224<br>ardiac<br>bilact<br>8427661<br>8.03<br>0.47<br>4.25<br>14.65<br>16.69<br>3.31<br>1.42  | 10.11<br>9.81<br>0.00<br>8.07<br>17.70<br>18.16<br>2.51<br>1.01  
                                 | elood<br>Blood<br>Heart<br>434WEP<br>11.70<br>8.51<br>3.19<br>5.85<br>02.43<br>17.56<br>4.78<br>4.78<br>1.06   | 224  |
| Commission         F           Class         1         10           2         10         3         10           3         10         3         10           4         10         5         10           5         10         10         10           10         11         10         11   | Initial     I | Drawse<br>and writter<br>and writter<br>and writter<br>brain<br>brain<br>120200<br>1225/0<br>1225/0<br>120200<br>1225/0<br>120200<br>1225/0<br>174010<br>120220<br>12024/0<br>120220<br>12024/0<br>120220<br>12024/0<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120220<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>120200<br>10000<br>100000<br>1000000<br>100000000   | Endina   | ColorOn •<br>7200018<br>14.28<br>9.71<br>14.29<br>15.14<br>13.99<br>5.71<br>14.99<br>1.14<br>2.57<br>14.14<br>9.57<br>14.14<br>0.39<br>0.57   | New Ta<br>2556000<br>14.18<br>21.53<br>2.05<br>5.23<br>10.66<br>0.25<br>0.16<br>0.25<br>0.25<br>0.23<br>0.48<br>2.03<br>0.15                                       
  | 450800<br>13.01<br>15.88<br>0.00<br>2.28<br>20.52<br>20.52<br>10.10<br>1.13<br>1.70<br>6.79<br>2.28<br>1.13  | 2.05<br>5.48<br>10.93<br>0.88<br>5.46<br>17.08<br>10.25<br>0.00<br>0.00<br>0.00<br>2.73<br>0.00<br>0.00  | 12,53<br>10,51<br>10,67<br>0,62<br>14,58<br>21,90<br>10,77<br>0,98<br>1,19<br>5,70<br>0,41<br>0,00   
   | 9.51<br>15.38<br>15.38<br>22.67<br>28.26<br>10.97<br>0.73<br>1.46<br>0.00<br>24.87<br>4.39<br>0.73   | 16.46<br>7.90<br>0.00<br>12.61<br>24.06<br>20.41<br>5.93<br>1.98<br>0.92<br>19.10<br>2.83<br>1.99   | ERCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>4.59<br>0.00<br>0.70<br>44.54<br>1.55<br>2.34<br>4.59<br>0.28<br>2.34<br>4.50<br>0.78<br>2.24<br>4.50<br>0.78<br>2.25<br>2.25<br>2.25<br>2.25<br>1.3   | 001044 EN<br>001044 EN<br>001044 EN<br>001044 EN<br>000<br>000<br>000<br>000<br>000<br>000<br>000<br>0   
  | CB915/200<br>CD8204K7U<br>CB9302E4C<br>2000<br>2000<br>2000<br>2000<br>2000<br>2000<br>2000<br>20  | EHCSR001<br>INCORS44<br>EHCSR000<br>940KUD<br>22.14<br>7.03<br>2.07<br>1.06<br>59.15<br>2.06<br>7.<br>1.00<br>0.03<br>0.03<br>0.03<br>0.03<br>0.03<br>0.03  | 900/98438<br>900/98438<br>7.20<br>8.80<br>4.02<br>2.41<br>37.74<br>16.66<br>1.61<br>0.60<br>2.41<br>0.80<br>2.41<br>0.80<br>0.60  
  | EP010002322<br>EP010002322<br>CL10002510<br>2010002510<br>201002510<br>2010<br>2010  | 805 12.11<br>805 12.11<br>8.50<br>8.50<br>8.50<br>7.16<br>7.16<br>7.16<br>1.73<br>0.46<br>1.73<br>4.90<br>0.00  | 1282200<br>1882200<br>2882201<br>2882201<br>2882201<br>2882201<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>288220<br>28820<br>28820<br>28820<br>28820<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>28920<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>289200<br>28920000000000   | 10.11<br>9.81<br>0.00<br>8.07<br>17.70<br>18.16<br>2.51<br>1.01<br>0.51<br>5.06<br>2.02<br>0.51   
  | 8000<br>Blood<br>Heart<br>11.70<br>8.51<br>3.19<br>5.85<br>0.243<br>17.55<br>4.79<br>1.06<br>3.19<br>7.44<br>4.25<br>0.53  | 225<br>24  |
| Chemilosei         f           1         10           2         10           3         10           4         10           5         10           5         10           7         10           8         10           9         10           10         11  | Chit     Chit    | Drawse<br>ess •<br>120250<br>12050<br>12050<br>12050<br>12050<br>12050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>112050<br>11200 | Energina   | ColarOn •<br>7200008<br>14.20<br>9.71<br>1.29<br>15.14<br>13.99<br>5.71<br>14.99<br>1.14<br>2.57<br>14.14<br>0.39   | New Ta<br>2656/WW<br>14.18<br>21 53<br>2.05<br>5.223<br>10.61<br>26.96<br>n.25<br>0.16<br>0.23<br>0.25<br>0.23<br>0.23<br>0.23<br>0.23<br>0.23                  
   | (300800)<br>13.01<br>18.88<br>0.00<br>2.28<br>20.92<br>20.92<br>10.18<br>1.13<br>1.70<br>6.79<br>2.28  | 2.05<br>5.48<br>10.93<br>0.88<br>5.46<br>17.08<br>40.25<br>0.00<br>0.00<br>0.00<br>0.00<br>2.73<br>0.00  | 12.50<br>10.51<br>10.57<br>0.52<br>14.58<br>21.90<br>10.77<br>0.99<br>1.19<br>5.70<br>0.41  
  | 9.51<br>15.38<br>0.73<br>22.67<br>28.76<br>10.97<br>0.75<br>1.46<br>0.00<br>24.87<br>4.39  | 16.46<br>7.90<br>0.00<br>12.61<br>24.36<br>20.41<br>5.93<br>1.98<br>0.193<br>19.10<br>2.63  | ERCPR<br>EXCPT<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENC<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>ENCPR<br>EN | 6000AH EN<br>A039/22 EN<br>1001DAA EN<br>1001DAA EN<br>1001DAA
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   | 81000<br>Dians<br>Heart<br>11,70<br>8,51<br>3,19<br>02,43<br>17,55<br>02,43<br>17,55<br>1,06<br>3,19<br>1,06<br>3,19<br>1,06<br>3,19<br>1,06<br>4,25   | 224  | | | | | | | | | | | | | | | | | | | | |
| Image: Class of the second s                        | Chit     Chit    | Drazes     icoaler     ic   | E program (00) started.<br>(01) started.<br>(01) started.<br>Essentia: Deve<br>entrio 02 44, v<br>entri 03 44, v<br>entri 03 44, v<br>entri 03 44, v<br>entri 04 44, v entri 04 44, v<br>entri 04 44, v entri 04 44, v en   | ColurOn   | New Ta<br>235K/W<br>14.18<br>2.05<br>3.23<br>10.61<br>20.95<br>0.25<br>0.23<br>0.48<br>2.03<br>0.15<br>0.52<br>1.25<br>0.00   | 490800<br>13,01<br>18,08<br>13,01<br>18,58<br>20,52<br>20,52<br>20,52<br>20,52<br>20,52<br>20,52<br>10,10<br>1,13<br>1,70<br>6,79<br>2,28<br>1,13<br>1,23<br>2,35<br>0,00<br>1,13  | 205<br>545<br>10,93<br>0,88<br>5,45<br>17,08<br>10,28<br>10,28<br>0,00<br>0,00<br>0,00<br>0,00<br>0,00<br>0,00<br>0,00   | 12,50<br>10,51<br>10,57<br>0,52<br>14,78<br>21,90<br>10,77<br>0,98<br>1,19<br>5,70<br>0,41<br>0,40<br>0,021<br>0,72<br>0,31  | 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| Download         I           1         100           2         10           3         10           5         10           5         10           7         10           8         10           10         10           11         10           12         10           13         10           14         10   | Chit     Chit    | Braves     Board     Coader     Sad     Writer     Braves     Exact     Total   | Enrolling and a second  | ColorOn<br>200000<br>14.28<br>2.71<br>1.28<br>5.71<br>1.18<br>5.71<br>1.14<br>2.87<br>14.14<br>0.29<br>0.57<br>14.14<br>2.20  | New Ta<br>2356/VW<br>14.18<br>2.05<br>3.23<br>10.61<br>20.95<br>0.15<br>0.23<br>0.15<br>0.23<br>0.48<br>2.03<br>0.15<br>0.53<br>1.28  |
(10000)<br>(13,01)<br>18,080<br>0,000<br>2,280<br>20,92<br>20,92<br>20,92<br>20,92<br>20,92<br>20,92<br>10,10<br>1,13<br>1,70<br>6,799<br>2,286<br>1,13<br>2,286<br>1,13<br>2,286<br>1,13<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,19<br>1,1     | 2.05<br>5.48<br>10.53<br>0.85<br>5.45<br>17.08<br>10.75<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.88<br>0.65  | 12,50<br>10,51<br>10,67<br>0,52<br>14,76<br>21,90<br>10,77<br>0,99<br>1,19<br>0,99<br>1,19<br>0,72   | 9.01<br>15.38<br>0.73<br>22.87<br>98.26<br>10.97<br>0.73<br>1.46<br>0.00<br>24.87<br>4.39<br>0.73<br>0.73<br>0.75<br>14.63  
  | 16.46<br>7.50<br>0.00<br>12.51<br>74.36<br>20.41<br>8.83<br>1.98<br>0.122<br>1.98<br>0.122<br>1.99<br>0.00<br>13.03   | 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 | 0010AU EN<br>0010AU EN<br>0010AU EN<br>1010AU EN<br>1010AU<br>4.15<br>1.40<br>0.00<br>2.722<br>0.00<br>2.722<br>0.00<br>2.722<br>0.00<br>2.722<br>0.00<br>2.722<br>0.00<br>2.722<br>0.00<br>2.722<br>0.00<br>2.732<br>1.74<br>2.75<br>1.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75<br>2.75  | E25/4240<br>E25/4240<br>20.02<br>10.28<br>2.00<br>20.02<br>10.28<br>2.00<br>20.02<br>10.28<br>2.04<br>2.04<br>2.04<br>2.04<br>2.04<br>2.04<br>2.04<br>2.04   
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| Jowenboard         F           Clears         Clears           1         40           2         10           3         44           5         54           6         10           7         10           8         10           11         14           12         14           13         14           14         15           15         10           17         10  | Chit     Chit    | Breake     Breakee     Breakeee     Breakeeeeeeeeeeeeeeeeeeeeeeeeeeeeeeeeeee  | E program (00) startied.<br>(01) started.<br>(01) started.<br>Elemina Deve<br>Elemina Deve<br>(01) 18 - 18 - 18 - 18 - 18 - 18 - 18 - 18   | ColsrOn + 220006 14,28 0,71 14,28 0,71 14,28 0,71 14,28 0,57 14,36 0,57 14,36 0,57 14,14 0,29 0,557 2,214 0,57 2,214 0,77 2,71 0,71 2,71 0,71 4,71 0,71 0,71 0,71 0,71 0,71   | New Ta<br>2356000 Ta<br>2400 | 4908000<br>18,011<br>18,888<br>0,000<br>2,288<br>20,929<br>20,929<br>10,110<br>1,139<br>1,200<br>6,799<br>2,288<br>0,000<br>1,139<br>2,288<br>0,000<br>1,139<br>2,288<br>0,000<br>1,139<br>2,288<br>0,000<br>1,139<br>2,288<br>0,000<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1,139<br>1, 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  | BHDRUD<br>BHDRUD<br>22.14<br>7.03<br>2.07<br>1.06<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00   | 000<br>000<br>7,244<br>9,5079828<br>7,249<br>15,86<br>1,621<br>0,800<br>0,800<br>0,000<br>0,000<br>0,000<br>0,000<br>0,000<br>0,000<br>0,000<br>0,000<br>0,000   | 2505440*<br>0.0012312<br>CL10002318<br>2505440*<br>0.04<br>4.34<br>5.54<br>1.30<br>28.15<br>13.73<br>3.64<br>0.43<br>3.64<br>0.43<br>3.64<br>0.43<br>3.64<br>0.43<br>3.64<br>0.43<br>0.67<br>0.43<br>0.67<br>0.43<br>0.87<br>0.43<br>0.87<br>0.43<br>0.87<br>0.87<br>0.87<br>0.87<br>0.87<br>0.87<br>0.87<br>0.87  | 885 12231<br>885 12  | Bat2 Beta<br>Bat2 Beta<br>Bat2 Beta<br>Bat3<br>Bat3<br>Bat3<br>Bat3<br>Bat3<br>Bat3<br>Bat3<br>Ba   | 10.11<br>9.81<br>0.03<br>8.07<br>17.70<br>18.16<br>2.751<br>1.01<br>8.51<br>6.06<br>2.02<br>0.51<br>1.01<br>0.00<br>0.00<br>2.02<br>2.80<br>8.00<br>2.07   | 434WEP<br>11.70<br>5.45<br>17.66<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.19<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>3.55<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06<br>1.06   | 225<br>24  |
| Jowenboad         F           Class         Class           1         4           2         11           4         10           5         44           5         44           6         10           6         10           10         44           12         14           13         10           14         15           15         10           16         16           17         17  | Control      Control     Contro     Contro     Control     Control     Control     Control     Co | Breaker     Bander (ander for ander for a   | Enrolman         Color Settield           (0.) Settield         (0.) Settield           (0   | ColorOn + 220E000 + 14.28 - 0.71 + 14.28 - 0.71 + 15.14 + 15.8 + 15.14 + 15.8 + 1.14 + 15.8 + 1.14 + 15.8 + 1.14 + 14.45 + 0.29 + 0.57 + 14.44 + 0.29 + 0.57 + 2.16 + 2.200 + 1.71 + 2.16 + 2.200 + 1.71 + 2.71 + 0.14 + 2.71 + 0.14 + 0.29 + 0.57 + 0.200 + 0.77 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.200 + 0.277 + 0.200 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.277 + 0.200 + 0.20  | New Ta<br>23560/W<br>14.16<br>21.53<br>2.05<br>3.23<br>10.61<br>20.55<br>0.15<br>0.23<br>0.15<br>0.55<br>1.25<br>0.55<br>1.25<br>0.55<br>1.25<br>0.55<br>1.25<br>0.55<br>1.25<br>0.55   | 4908000<br>18,01<br>18,68<br>0,00<br>2,28<br>20,52<br>20,52<br>10,10<br>1,13<br>1,70<br>6,79<br>2,28<br>0,00<br>1,13<br>2,28<br>0,00<br>1,13<br>2,28<br>0,00<br>1,13<br>1,70<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10<br>1,10 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BHCSR000<br>BHCSR000<br>BHCSR000<br>22:14<br>703<br>22:14<br>703<br>22:14<br>703<br>2015<br>106<br>88:57<br>1:30<br>0:00<br>0:52<br>0:00<br>0:52<br>0:00<br>0:52<br>0:00<br>0:52<br>0:00<br>0:52<br>0:00<br>0:52<br>0:00<br>0:52<br>0:00<br>0:52<br>0:00<br>0:52<br>0:00<br>0:52<br>0:52  | 000 000 000 000 000 000 000 000 000 00   | 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   | 9.54<br>15.38<br>0.73<br>22.67<br>29.57<br>10.97<br>0.73<br>1.46<br>0.09<br>24.87<br>4.39<br>0.73<br>0.73<br>8.05<br>3.65<br>3.65<br>3.65<br>1.45  | 16.46<br>7.90<br>0.00<br>12.51<br>24.05<br>20.41<br>5.83<br>1.99<br>0.02<br>19.10<br>2.83<br>1.99<br>0.00<br>10.03<br>4.61<br>2.63<br>1.92<br>2.63<br>1.92<br>2.63  | 001114c<br>enCFR<br>4.59<br>0.070<br>44.56<br>1.55<br>2.34<br>4.59<br>45.10<br>0.76<br>4.55<br>2.34<br>45.10<br>0.77<br>2.35<br>2.57<br>9<br>2.73<br>2.53<br>2.53<br>2.53<br>2.53<br>2.53<br>2.53<br>2.53<br>2.5  | 00004H EH<br>001040 E<br>001044 E<br>001044
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  | PUPUBUCIES<br>INTO MAD2222<br>CUUME2HIE<br>SIGNAP<br>0.64<br>4.34<br>4.34<br>4.34<br>4.34<br>4.34<br>4.34<br>4.34<br>4   | 897 12.11<br>186 12.11  | 98229<br>98279<br>9879<br>987<br>987<br>987<br>987<br>987<br>987  | 10.11<br>9.81<br>0.00<br>17.70<br>18.16<br>2.53<br>1.01<br>0.51<br>0.05<br>1.00<br>0.00<br>2.02<br>0.00<br>2.02<br>0.00<br>2.02<br>0.00<br>2.02<br>0.00<br>2.02<br>0.00<br>2.02<br>0.00<br>2.02<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.000000   
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| Download         I           Class         1           1         1           2         1           3         14           5         14           5         14           5         14           5         14           5         14           5         14           6         17           0         14           12         16           13         14           14         15           15         16           16         16           17         16           18         16           19         16           20         18           22         18           23         18  | Long     Long    | Indiana Market Indiana In   | E program (00) startied.<br>(01) started.<br>(01) started.<br>Elemina Deservices<br>(01) started.<br>Elemina Deservices<br>(01) started.<br>(01) started.<br>(02) starte   | DolmOn 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| 16.45<br>7.90<br>0.00<br>12.51<br>24.35<br>20.41<br>5.83<br>0.92<br>19.10<br>2.15<br>2.15<br>2.15<br>2.15<br>2.15<br>2.15<br>2.15<br>2.15   | ERCPF<br>BACTY<br>ERCFF<br>4.59<br>0.00<br>4459<br>0.00<br>4454<br>1.56<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.070<br>0.00000000 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| Download         I           Class         1           1         1           2         1           3         14           5         14           5         14           5         14           5         14           5         14           5         14           6         17           0         14           12         16           13         14           14         15           15         16           16         16           17         16           18         16           19         16           20         18           22         18           23         18  | Initial         Automatical           11/10         11/10   | Indiana Market Indiana In   | E program (00) startied.<br>(01) started.<br>(01) started.<br>Elemina Deservices<br>(01) started.<br>Elemina Deservices<br>(01) started.<br>(01) started.<br>(02) starte   | DolmOn •<br>2000000<br>14.28<br>0.71<br>14.28<br>15.14<br>13.98<br>5.71<br>14.98<br>1.14<br>2.87<br>14.14<br>0.99<br>0.57<br>2.14<br>2.00<br>1.21<br>2.71<br>0.14<br>2.00<br>1.21<br>0.71<br>0.14<br>1.20<br>0.57<br>2.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.14<br>1.20<br>0.57<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.21<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25<br>1.25 | New Ta<br>2558/00W<br>14.18<br>21.93<br>2.29<br>10.61<br>2.25<br>0.25<br>0.25<br>0.25<br>0.25<br>0.25<br>0.25<br>0.25  
  | 4908024<br>4908024<br>15.01<br>18.88<br>0.00<br>2.38<br>20.92<br>20.92<br>10.10<br>1.70<br>4.70<br>2.38<br>0.00<br>1.13<br>2.38<br>0.00<br>1.19<br>2.58<br>0.00<br>0.00<br>0.00<br>0.00  | 2.05<br>5.45<br>10.55<br>0.85<br>5.45<br>17.08<br>40.25<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00<br>0.00  | 12.53<br>10.51<br>10.51<br>14.56<br>21.59<br>10.59<br>1.19<br>0.99<br>1.19<br>0.99<br>1.19<br>0.99<br>0.21<br>0.21<br>0.21<br>0.21<br>0.21<br>0.21<br>0.21<br>0.21   | 9.51<br>15.38<br>0.75<br>22.67<br>10.27<br>0.75<br>1.45<br>0.09<br>24.39<br>0.73<br>0.73<br>0.73<br>0.73<br>0.75<br>3.025<br>3.05<br>3.05<br>3.05<br>3.05<br>3.05<br>3.05<br>3.05<br>3.0  
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16.45<br>7.90<br>0.00<br>12.51<br>24.55<br>24.55<br>20.41<br>5.83<br>1.99<br>0.02<br>19.10<br>2.153<br>1.99<br>0.02<br>13.03<br>4.51<br>2.53<br>1.99<br>0.00<br>13.03<br>4.51<br>2.53<br>1.99<br>0.00<br>13.03<br>4.51<br>2.53<br>1.99<br>0.00<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>13.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>1.99<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.03<br>14.031 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# Figure 2 Web interfaces for annotating the openness of genomic regions in batches

A. The interface for submitting a new annotation task. B. Display of total calculation
progress, and the interface for sending remarks and download links of results by email.
C. Display of real-time annotation results. Users can browse part of a big table and
scroll to any row and any column smoothly. D. More detailed information about the
experiments and the visualization in UCSC Genome Browser of a specific genomic
region. E. The interface for downloading the result plain text dump files.

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157 After submitting the annotation task, users can follow the links of results sent to 158 their email, and download the results after the calculation is completed. Furthermore, 159 users can directly browse real-time results of each region. Here we demonstrate a 160 seamless integration of hardware and software to hold the big result tables which may 161 contain billions of rows and hundreds of columns. Users can observe part of a big table 162 and scroll to any row and any column smoothly. The computational status in different 163 chromosomes will be updated in real time. Users can arbitrarily switch between 164 different chromosomes or the four different types of openness values. By enabling the 165 Color option, each sample in the table will be colored according to the score of openness 166 for intuitively comparing the openness of different genomic regions in different 167 experiments. More detailed information about the experiments and the visualization of 168 a specific genomic region in UCSC Genome Browser [23] can be obtained by doubleclicking on a row. After the calculation is completed, users can extract the result files 169 170 through the 'Download' button. By clicking the 'New Task' button, users can submit a 171 new task on a new tab in the browser.

172

#### 173 Web interfaces for browse

On the 'Browse' page, users can study the openness of a particular genomic region more intuitively. As illustrated in **Figure 3**, we take an enhancer (chr10: 94,513,996-94,517,989) of the tissue of blood vessels provided by the VISTA Enhancer Browser [22] as an example to demonstrate the service in this page. After submitting the form that contains the chromosome, starting site, terminating site, and strand of a particular genomic region, the web server provides information from three perspectives, including (1) the average openness scores of this genomic region in different biological systems,

- 181 tissues, and cell lines, (2) the visualization of this genomic region in UCSC Genome
- 182 Browser [23], and (3) openness scores and details of the 871 DNase-seq experiments.



183

# Figure 3 The interface for studying the openness of a particular genomic region more intuitively

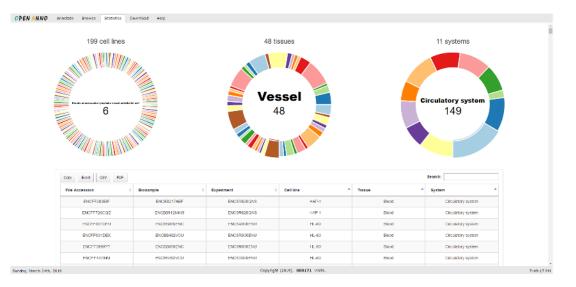
186

First of all, the average openness scores enable users to compare the openness across different biological systems, tissues, and cell lines more intuitively. For example, as shown in **Figure 3**, the average openness score of this enhancer in the vessel tissue is obviously higher than that in other tissues. We calculated z-scores, standard deviations from the mean, the average openness scores of this enhancer in the vessel tissue, and found that the z-scores in all the 4 different types of openness definition 193 beyond the three-sigma limits (3.18 of foreground read count, 4.07 of raw read 194 openness, 3.29 of narrow peak openness and 4.37 of broad peak openness), which 195 demonstrates that all the 4 different types of openness of the enhancer in the vessel 196 tissue are significantly higher than that in other tissues from the statistical perspective. 197 This coincides with the fact that this genomic region is an enhancer in blood vessels according to VISTA annotation. Second, the visualization in UCSC Genome Browser 198 199 provides rapid and reliable displaying of the requested genomic region at any scale, 200 together with dozens of aligned annotation tracks (http://genome.ucsc.edu/) [23]. We 201 also provide a hyperlink to UCSC Genome Browser to facilitate users to achieve more 202 concrete information of the particular genomic region. Finally, openness scores and 203 detailed information of the 871 DNase-seq experiments are filled in a table with 204 advanced features. Users can sort the table according to the openness to find out in 205 which experiment the genomic region has higher openness. Users can also sort the table 206 on the basis of different columns according to their own needs. We also provide a 207 convenient service for searching in the table. Users can quickly query information they 208 are interested in. Furthermore, users can directly copy the table or download tables 209 stored in different formats for other requirements.

210

#### 211 Web interfaces for statistics, downloading, and help

212 On the 'Statistics' page, as shown in Figure 4, users can intuitively compare the number 213 of experiments across different cell lines, tissues, or biological systems. A table with 214 advanced features also provides detailed information, including file accessions, biosamples, experiments, cell lines, tissues, and systems, of all the 871 DNase-seq 215 216 experiments. We collected regulatory element datasets from FANTOM[21] and 217 VISTA[22], including 184,476 FANTOM5 human promoters, 32,693 FANTOM5 218 human enhancers, and 979 VISTA human enhancers. We calculated the openness of these regulatory elements in advance, and provide a download service on the 219 220 'Download' page to allow users to directly download plain text dump files in 221 compressed (gzip) format. We will continue to provide the openness of other public 222 and validated regulatory elements. To improve the usability, we provide a 'Help' page 223 with other commonly used information of the web server, including frequently asked 224 questions, news about the releases of OPENANNO, tutorials of each web interface, 225 citation information, and contact information for help and feedback.



226

Figure 4 The interface for intuitively comparing the number of experiments
 across different cell lines, tissues, or biological systems, and achieving details of all
 the 871 DNase-seq experiments

230

#### 231 **OPENANNO facilitates regulatory mechanism studies**

232 The chromatin 'openness', *i.e.*, the accessibility of genomic regions, calculated using 233 our method has been widely applied to various studies of regulatory mechanism. Here, 234 we show two examples to demonstrate the output of OPENANNO contains valuable 235 information. For example, a model named DeepTACT has been proposed to integrate 236 DNA sequences and chromatin accessibility data for the prediction of chromatin 237 contacts between regulatory elements [24]. Briefly, DeepTACT first performs a onehot encoding strategy and calculates the raw read openness of each site for 238 239 characterizing a given genomic region. DeepTACT takes the sequences of two one-hot 240 encoded regulatory elements, and their chromatin openness scores derived from 241 OPENANNO of a given cell type as input. The output of DeepTACT is the predictive 242 score that represents the probability the two regulatory elements have 3D contact. In 243 the deep neural network of DeepTACT, a sequence module is used to extract features 244 from DNA sequences, an openness module is adopted to learn epigenomic features 245 from chromatin openness scores, and an integration module merges outputs of these two modules and extracts high-level features with an attention-based recurrent neural 246 247 network to predict the probability that the two regulatory elements have 3D contact. Using sequence features and the raw read openness of genomic regions, DeepTACT, 248

as a bootstrapping deep learning model, outperforms existing methods on the task of

inferring both promoter-enhancer and promoter-promoter interactions. In more detail,
with same test sets, DeepTACT achieves a mean auPRC (the area under the precisionrecall curve) score of 0.89 for inferring promoter-promoter interactions compared with
0.76 of SPEID [25] and 0.23 of Rambutan [26]. For inferring promoter-enhancer
interactions, DeepTACT achieves a mean auPRC of 0.82 compared with 0.67 of SPEID
and 0.36 of Rambutan.

Besides, DeepTACT provides a finer mapping of promoter-enhancer and promoterpromoter interactions from high-quality promoter capture Hi-C data. Furthermore, the class of hub promoters identified by DeepTACT, and the integrative analysis of existing GWAS data and chromatin contacts predicted by DeepTACT demonstrate the openness calculated by OPENANNO bridges the epigenome and transcriptome and plays an important role in understanding the regulatory mechanism.

262 In addition to DeepTACT, a model named DeepCAPE is proposed to predict 263 enhancers via the integration of DNA sequences and DNase-seq data [27] with the 264 understanding that DNase I hypersensitivity has been shown to be important to identify 265 active cis-regulatory elements including enhancers, promoters, silencers, insulators, 266 and locus control regions [28]. Briefly, DeepCAPE uses the raw read openness of each 267 site of a genomic region as the information of chromatin accessibility to greatly improve 268 the performance of predicting enhancers. In more detail, when the ratio of positive and 269 negative samples is 1:20, the auROC (the area under the receiver operating 270 characteristic curve) and auPRC scores of DeepCAPE are on average 0.151 and 0.590 271 higher than gkmSVM [29], 0.151 and 0.598 higher than DeepSEA [30], and 0.150 and 272 0.588 higher than DeepEnhancer [31], respectively. One-sided paired-sample 273 Wilcoxon signed rank tests consistently suggest that DeepCAPE consistently achieves 274 higher auPRC scores (*p*-values < 2.2e-16 for all the other three baseline methods), and 275 higher auROC scores than a baseline method (p-values < 2.2e-16 for all the other three 276 baseline methods).

In the model ablation analysis for evaluating the contributions of DNA sequences and DNase-seq data, DeepCAPE illustrates that DNase-seq data provides more information than DNA sequences to greatly improve the performance of prediction. Besides, the information provided by DNA sequences also plays an important role in promoting the performance and making the performance more stable. Because the number of DNase-seq experiments varies between cell lines, the dimensionality of input data varies between cell lines and prevents the use of convolutional neural networks in 284 the cross cell line prediction. DeepCAPE therefore adopts a neural network designed 285 for unsupervised learning of efficient encodings [32], named auto-encoder, to embed 286 chromatin openness scores of a DNA fragment derived from OPENANNO into a vector 287 of fixed length in a low-dimensional latent space. Comparing to the model without an 288 auto-encoder, and other two strategies that average the replicates or randomly select a 289 single replicate, DeepCAPE with an auto-encoder not only makes cross cell line 290 prediction possible, but also maintains superior performance even if the dimensionality 291 of the openness data is reduced. In addition, with a collective scoring strategy, 292 DeepCAPE achieves an average auROC of 0.971 and an average auPRC of 0.862 in 293 the cross cell-line prediction when the ratio of positive and negative samples is 1:20, 294 and thus establishes a landscape of potential enhancers specific to a cell line that still 295 lacks systematic exploration of enhancers.

296 To sum up, DeepCAPE not only achieves superior prediction performance in a cell 297 line-specific manner, but also makes accurate cross cell line predictions possible with 298 the openness scores calculated by OPENANNO. With this understanding, analogous 299 machine learning frameworks can possibly be adapted for the prediction of other 300 functional elements in the genome, including but not limited to promoters, silencers, 301 insulators, repressors, and locus control regions. In addition, the strategy that integrates 302 DNA sequences and chromatin openness can also be generalized for the prioritization 303 of candidate variants in whole-genome sequencing studies, and thus facilitate the 304 regulatory mechanism studies.

305

#### 306 **OPENANNO facilitates association studies**

307 Network-based functional studies play an important role in the identification of disease-308 associated genes and the interpretation of disease mechanism. The functional 309 relationship of a pair of genes is influenced not only by the co-activation of transcripts, 310 but the regulation mechanism [33]. Besides, the consistence of gene chromatin 311 accessibility indicates the tendency of genes being co-regulated. With this 312 understanding, a gene co-opening network has been constructed based on the raw read 313 openness of genes [34]. Briefly, they take alternative promoters of genes into account 314 when calculating the correlation (absolute value of *Pearson*'s correlation coefficient) 315 of the openness scores between two genes, considering the prevalence of the alternative 316 splicing phenomena. By calculating a co-opening score for every pair of genes, they 317 obtain a co-opening matrix for all genes to facilitate the downstream analysis.

318 The results demonstrate that the co-opening network contains new information 319 different from co-expression networks and protein-protein interactions networks. In 320 addition, the genes related to a specific biological process or a specific disease has been 321 demonstrated to tend to be clustered together in the co-opening network, which 322 facilitates detecting functional clusters in the network and predicting new functions for 323 genes. Particularly, through integrative analysis with fruitful genome-wide association 324 studies (GWAS) data, the co-opening network provides a new perspective to the 325 discovery of genes associated with complex diseases, and thus benefits elucidating gene 326 associations and the deciphering of disease mechanisms. For example, by simulating a 327 random walk process on the co-opening network, they use the steady state probability 328 assigned to a gene as a score to measure the likelihood that the gene is associated with 329 the disease under investigation. Applying this strategy to a complex disease named 330 *Psoriasis*, a potentially disfiguring immune-mediated inflammatory disease of skin, 331 they discovered TNFSF14 (TNF super family member 14, a biomarker of Psoriasis 332 [35]), which was ranked second by the random walk model while cannot be identified 333 by GWAS (p-value = 0.1616, ranked 1259 based on the p-value). In general, the co-334 opening network is ready to serve as a useful resource complementary to the widely 335 used co-expression network, and thus shed light on the studies in system biology.

336

#### 337 Perspectives and concluding remarks

338 Chromatin accessibility, which bridges the epigenome and transcriptome, is a very 339 valuable resource for interpreting non-coding genomic region and understanding the 340 regulatory mechanism. In this study, we downloaded raw sequencing data of 871 341 DNase-seq experiments across 199 cell lines, 48 tissues and 11 biological systems from 342 ENCODE, and defined the openness of genomic regions from four perspectives. In 343 addition, we take an enhancer of the tissue of blood vessels provided by the VISTA 344 Enhancer Browser as an example to illustrate the valuable information provided by all 345 the four different types of openness from statistical perspective. Furthermore, we 346 designed a parallel program to endow OPENANNO with the ability to efficiently 347 annotate and visualize openness for a vast amount of genomic regions. Finally, we 348 introduced three examples to demonstrate the output of OPENANNO serves as 349 valuable input for follow-up regulatory mechanism and association studies.

350 Our web server has four main application scenarios. First, one can use our web 351 server to annotate openness of genomic regions, and then integrate the information of 352 openness to a machine learning model for superior performance. Second, one can use 353 our web server to visualize the openness of a specific genomic region to intuitively 354 understand this region has higher openness in which cell lines, tissues, or systems, and 355 thus contribute to the study of functional implications of this genomic region. Third, 356 our web server offers a new opportunity to reinterpret abundant data cumulated by 357 genome-wide association studies, and thus one can characterize variants by integrating 358 upstream openness annotated with our web server and downstream gene expression. 359 Finally, one can use the openness annotated with our web server to construct gene co-360 opening networks which provide a new perspective to association studies.

361 To better serve the academic community, we will continue to collect public data 362 and update OPENANNO regularly in the future. Our next plan is to provide the option 363 of other genomes or species, and the option of annotating using other chromatin 364 accessibility data, such as ATAC-seq data. We will continue to provide the openness 365 of other public and validated regulatory elements for downloading directly. According 366 to users' feedbacks, we will continue to improve the interfaces and performance of 367 OPENANNO. We believe that OPENANNO would serve as a useful tool for both 368 bench scientists and computational biologists, and shed light on studies including but 369 not limited to bioinformatics and system biology.

370

### 371 Materials and methods

#### **Data collection**

373 We first parsed a total of 41,418 JSON files from ENCODE to obtain detailed 374 information about experiments, biosamples, cell lines, tissues, and systems of the 375 DNase-seq data provided by the ENCODE project [18]. Under the constraint that each 376 experiment contains both narrow peaks and broad peaks, we downloaded raw 377 sequencing data of 891 DNase-seq experiments in human reference genome GRCh37 378 (hg19), and then identified experiments corresponding to 199 cell lines, 48 tissues and 379 11 biological systems. We collected datasets of regulatory elements from FANTOM [21] and VISTA[22] that have been experimentally validated, including 184,476 380 381 FANTOM5 human promoters, 32,693 FANTOM5 human enhancers, and 979 VISTA 382 human enhancers.

#### 383

#### 384 **Definition of openness**

385 We defined the openness of given genomic regions from four perspectives, including 386 foreground read count, raw read openness, narrow peak openness, and broad peak 387 openness. Specifically, given the raw sequencing data of a DNase-seq experiment, we 388 provided the number of reads (N), *i.e.*, foreground read count, falling at a specific 389 genomic region to facilitate special applications that may use raw read counts of a 390 DNase-seq experiment directly. To remove the effect of sequencing depth, we defined 391 the raw read openness (S) of a genomic region as the foreground read count (N) divided 392 by the average number of reads falling at a position in a background region of size W 393 surrounding the given region. The raw read openness (S) can be simply calculated as

$$S = \frac{N}{K/W} \tag{1},$$

where *K* is the number of reads falling into the background region of size *W*. The size
of a background region *W* is set to 1 M base pairs, according to the suggestion from
[36].

Analogously, we defined the narrow peak openness (and broad peak openness) of a genomic region as the number of narrow peaks (or broad peaks) overlapping with the genomic region, divided by the average number of narrow peaks (or broad peaks) overlapping with a position in a background region of size *W* surrounding the given genomic region.

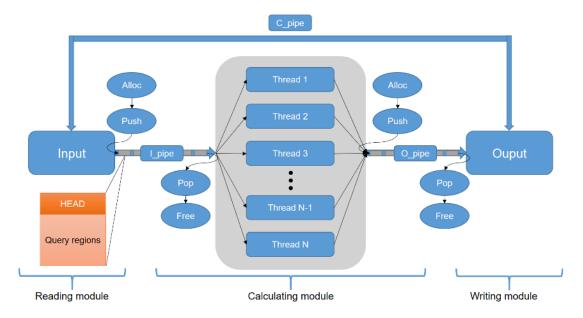
402

#### 403 **Parallel computing and real-time browsing**

To facilitate querying reads and peaks at high frequencies, and the massive demand for 404 405 annotating openness of a large number of genomic regions, we designed a parallel 406 strategy that endows OPENANNO with an ability to efficiently annotate openness of a vast amount of genomic regions. We used C++, a programming language known for its 407 408 high efficiency, to develop a multithreaded program that consists of a reading module, 409 a calculating module, and a writing module. As illustrated in **Figure 5**, the reading 410 module packages the query regions in the input file into data blocks and pushes them 411 into the input pipeline *I pipe*. Each data block contains a portion of the query regions, 412 and header information that contains the number and indexes of these query regions in 413 the input file. When a thread in the calculating module is idle, the data block is 414 automatically extracted from the input pipeline for calculation. The results are pushed

415 into the output pipeline  $O_pipe$  in a similar form of data blocks, which are then popped 416 out and written into the disks by the writing module. Through the communication 417 pipeline  $C_pipe$ , the reading module can respond to the working state of the writing 418 module. When the writing speed is lower than the reading speed, the reading module 419 pauses pushing the data block into  $I_pipe$  to effectively save memory resources. With 420 the parallel program, OPENANNO can calculate the four types of openness across all

421 the 871 DNase-seq experiments of one thousand genomic regions within 1 second.



422

# Figure 5 The multithreaded program for efficiently annotating openness of a vast amount of genomic regions

425

426 Users may need to annotate a large number of genomic regions, and thus results in 427 very large tables of openness, especially when the per-base option is enabled. Loading 428 all the results directly to the front-end of OPENANNO will take a long time and result 429 in an unsmooth experience, and thus is an unrealistic choice. It is, therefore, necessary 430 to provide a highly efficient front-end web application to browse these tables. To 431 accomplish this, we used WebSocket (https://www.websocket.org/), a computer 432 communications protocol, to provide full-duplex communication channels over a single 433 socket over the web and the remote host, realize the real-time browsing of calculation 434 results, and thus users can observe part of a large table and scroll to any row and any 435 column smoothly.

436

#### 437 Web server implementation

438 The whole design of the OPENANNO is shown in Figure 1, with the possible jumps 439 among web pages illustrated. OPENANNO is freely available to all users without a 440 login requirement. The current version of OPENANNO was deployed on a calculation 441 node and a server node of a high-performance computer cluster. The calculation node, 442 whose operating system is CentOS 7.5 (one of the most popular Linux distributions; 443 https://www.centos.org), has 56 hyper-threaded processors to perform efficient parallel 444 computing, RAM of 775 GB to support a large number of reading and writing 445 operations, and storage space of 321 TB to store the vast amounts of data. Incron 446 (http://inotify.aiken.cz/?section=incron) is used to monitor filesystem events and 447 executes predefined commands. Once the server node creates a new task in a specific 448 file path, the parallel program automatically runs and stores the results in another 449 specific file path for the website to read and users to download.

450 In the server node, a Linux-based Nginx server (https://www.nginx.com/), which 451 uses dramatically less memory than Apache (https://www.apache.org/) and can handle 452 roughly four times more requests per second, is deployed to improve the performance, 453 reliability, and security of OPENANNO. PHP (http://www.php.net/) is used for server-454 side scripting. Bootstrap (a popular toolkit for developing with HTML, CSS and 455 JavaScript; https://getbootstrap.com/) and jQuery (a fast and feature-rich library 456 designed to simplify the JavaScript programming; https://jquery.com/) are adopted for 457 the front-end, *i.e.*, the interactive and responsive user interface, of OPENANNO. The 458 user interface of OPENANNO can automatically respond to the devices and browsers 459 with different screen resolutions, and change its structure and shape according to the 460 resolution in order to optimize the visualization. DataTables (https://datatables.net/), as 461 a plug-in for the jQuery and JavaScript library, is used to add advanced features to the 462 tables. The visualizations of bar charts and pie charts are implemented with JavaScript (https://canvasjs.com/) 463 libraries CanvasJS named and morris.js 464 (https://morrisjs.github.io/morris.js/index.html), То obtain respectively. the 465 visualization in UCSC Genome Browser of a query region, we put the information of chromosome, start position, end position and reference genome to the link 466 467 http://genome.ucsc.edu/cgi-

468 bin/hgTracks?db=(reference\_genome)&position=(chromosome):(start\_position)-

469 (end\_position) to finish UCSC link construction.

- 470 All codes are developed using Vim (a highly flexible text editor that supports any
- 471 kind of text; <u>https://www.vim.org/</u>). The performance of OPENANNO has been tested
- 472 in Chrome, Firefox, Opera and Microsoft Edge on Windows 10, Ubuntu 16.04 and
- 473 MacOS 10.12. We hope users could feedback their comments and suggestions through
- the contact page on our website to help us improve OPENANNO.
- 475

### 476 Authors' contributions

477 RJ and YW designed the project. SC and RJ collected data and implemented the web

478 server. SC, YW and RJ wrote the paper. All authors read and approved the final479 manuscript.

480

#### 481 Competing interests

- 482 The authors have declared no competing interests.
- 483

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489 RONG professor at the Institute for Data Science, Tsinghua University.

490

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- 578

### 579 Figure legends

580	Figure 1	The diagram	for constructing	the <b>OPENANNO</b>	web server

581

# 582 Figure 2 Web interfaces for annotating the openness of genomic regions in583 batches

- A. The interface for submitting a new annotation task. B. Display of total calculation
  progress, and the interface for sending remarks and download links of results to email.
  C. Display of real-time annotation results. Users can observe part of a big table and
  scroll to any row and any column smoothly. D. More detailed information about the
  experiments and the visualization in UCSC Genome Browser of a specific genomic
- 589 region. **E.** The interface for downloading the result plain text dump files.
- 590

# 591 Figure 3 The interface for studying the openness of a particular genomic region 592 more intuitively

593

594 Figure 4 The interface for intuitively comparing the number of experiments 595 across different cell lines, tissues or systems, and achieving detail information of 596 all the 871 DNase-seq experiments

597

598 Figure 5 The multithreaded program for efficiently annotating openness of a
599 vast amount of Genomic regions