

## 1.DataSet 2

The second data set was introduced in 2012 by Yamanishi et al[1]. This dataset includes three txt files of drug-target interactions, drug features, and target features, and all data in this data set are numeric.

The second data set is described in two parts; In the first part, information about drug-target interactions and in the second part, information about the drug and target used and how they are represented as feature vectors are described.

### 1-1. Drug-target interaction data

Drug-target interaction file as a two-dimensional matrix  $Y \in \mathbb{R}^{n \times m}$  with n drug rows and m target columns. That is,  $Y_{ij} = 1$  if drug  $d_i$  and target  $t_j$  interact and  $Y_{ij} = 0$  otherwise. In other words, the one indicates the presence of interaction and the zero indicates the lack of interaction between the drug  $d_i$  which is in the matrix row and the target  $t_j$  which is located in the matrix column. Also, the features of drugs are in the form of  $D \in \mathbb{R}^{n \times f}$  matrix and the features of targets are in the form of  $T \in \mathbb{R}^{m \times g}$  matrix, where f and g are the number of features of drugs and targets. The interaction data were obtained from the DrugBank [2, 3]. In total, there are 4809 drug-target interactions between 1862 drugs and their 1554 protein interaction partners.

### 1-2. Drugs and targets information

Chemical structures of drugs were encoded by a chemical fingerprint corresponding to 881 chemical substructures defined in the PubChem database. Chemically identical drugs with the same structures (duplicates) are removed, so structures of all drugs in the above interaction data are unique. Each drug was represented by 881-dimensional binary vector whose elements encode for the presence or absence of each PubChem substructure by 1 or 0, respectively. Genomic information about target proteins was obtained from the UniProt database, and associated protein domains were obtained from the PFAM database [4]. Target proteins in our dataset were associated with 876 PFAM domains. Each target protein was represented by 876-dimensional binary vector whose elements encode for the presence or absence of each of the retained PFAM domain by 1 or 0, respectively

The file of drug features is in the form of  $D \in \mathbb{R}^{n \times f}$  matrix and the target features are in the form of  $T \in \mathbb{R}^{m \times g}$  matrix, where f and g are the number of drug features

and targets features. Formation of drug-target vectors for use in machine learning model as  $[d_1, d_2, \dots, d_p, t_1, t_2, \dots, t_q, I]$ , which label  $I$  is a binary number and indicates the presence or absence of interaction  $I_s$  between the drug and the target in the vector. Label  $I$  is identified by the drug-target interaction matrix. Details of this dataset are given in Table 1. Figures 1 and 2 show a view of the drug-target interaction file and the drug features file, respectively.

Table 1. Statistics of second dataset

Drugs	Targets	Interactions
1862	1554	4809

The file size (5.82 MB) exceeds the configured limit (2.56 MB). Code insight features are not available.

	B11_HUMAN	3BHS1_HUMAN	3BHS2_HUMAN	3HIDH_HUMAN	5HT1A_HUMAN	5HT1B_HUMAN	5HT1D_HUMAN	5HT1E_HUMAN	5HT1F_HUMAN	5HT2A_HUMAN	5HT2B_HUMAN	5HT2C_HUMAN	
1													
2	100252	0	0	0	0	0	0	0	0	0	0	0	0
3	1003	0	0	0	0	0	0	0	0	0	0	0	0
4	1005	0	0	0	0	0	0	0	0	0	0	0	0
5	10062751	0	0	0	0	0	0	0	0	0	0	0	0
6	10132	0	0	0	0	0	0	0	0	0	0	0	0
7	1014	0	0	0	0	0	0	0	0	0	0	0	0
8	10180	0	0	0	0	0	0	0	0	0	0	0	0
9	1021	0	0	0	0	0	0	0	0	0	0	0	0
10	10214	0	0	0	0	0	0	0	0	0	0	0	0
11	10267	0	0	0	0	0	0	0	0	0	0	0	0
12	1030	0	0	0	0	0	0	0	0	0	0	0	0
13	1045	0	0	0	0	0	0	0	0	0	0	0	0
14	1046	0	0	0	0	0	0	0	0	0	0	0	0
15	104741	0	0	0	0	0	0	0	0	0	0	0	0
16	104758	0	0	0	0	0	0	0	0	0	0	0	0
17	104799	0	0	0	0	0	0	0	0	0	0	0	0
18	104850	0	0	0	0	0	0	0	0	0	0	0	0
19	104865	0	0	0	0	0	0	0	0	0	0	0	0
20	1050	0	0	0	0	0	0	0	0	0	0	0	0
21	1051	0	0	0	0	0	0	0	0	0	0	0	0
22	10517	0	0	0	0	0	0	0	0	0	0	0	0
23	1053	0	0	0	0	0	0	0	0	0	0	0	0
24	10531	0	0	0	0	0	1	1	0	0	1	0	0

Figure 1. view of the drug-target interaction file

The file size (3.3 MB) exceeds the configured limit (2.56 MB). Code insight features are not available.

1	SUB1	SUB2	SUB3	SUB4	SUB5	SUB6	SUB7	SUB8	SUB9	SUB10	SUB11	SUB12	SUB13	SUB14	SUB15	SUB16	SUB17	SUB18	✓	
2	100252	1	1	0	0	0	0	0	1	1	1	0	0	1	1	1	0	0	0	0
3	1003	0	0	0	0	0	0	0	0	0	0	0	0	1	1	1	0	0	0	0
4	1005	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
5	10062751	1	1	0	0	0	0	0	0	1	1	1	0	0	1	1	1	0	0	0
6	10132	0	0	0	0	0	0	0	1	0	0	0	0	0	1	1	1	0	0	0
7	1014	1	1	0	0	0	0	0	1	1	0	0	0	1	1	1	0	0	0	0
8	10180	1	1	1	0	0	0	0	0	1	1	1	1	0	1	1	0	0	0	0
9	1021	1	1	0	0	0	0	0	0	1	1	1	0	0	1	1	0	0	0	0
10	10214	1	1	0	0	0	0	0	0	1	1	1	0	0	1	1	1	0	0	0
11	10267	1	1	0	0	0	0	0	0	1	1	0	0	0	0	1	1	1	0	0
12	1030	1	1	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
13	1045	1	1	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	0	0
14	1046	1	0	0	0	0	0	0	0	1	1	0	0	0	1	0	0	0	0	0
15	104741	1	1	1	1	0	0	0	1	1	1	1	1	0	0	0	0	1	1	1
16	104758	1	1	1	0	0	0	0	0	1	1	1	1	0	1	1	1	0	0	0
17	104799	1	1	1	0	0	0	0	0	1	1	1	0	0	1	1	1	0	0	0
18	104850	1	1	1	0	0	0	0	0	1	1	1	0	1	1	1	0	0	0	0
19	104865	1	1	1	0	0	0	0	0	1	1	1	1	0	1	1	1	0	0	0
20	1050	1	1	0	0	0	0	0	0	1	1	1	0	0	1	1	0	0	0	0
21	1051	1	1	0	0	0	0	0	0	1	1	1	0	0	1	1	1	0	0	0
22	10517	1	1	1	0	0	0	0	0	1	1	1	1	0	1	1	0	0	0	0
23	1053	1	1	0	0	0	0	0	0	1	1	1	0	0	1	1	1	0	0	0
24	10531	1	1	1	1	0	0	0	0	1	1	1	1	1	1	1	0	0	0	0
25	1054	1	1	0	0	0	0	0	0	1	1	1	0	0	1	1	0	0	0	0

Figure 2. view of the drug features file

## References:

- [<sup>1</sup>] Y. Yamanishi, E. Pauwels, H. Saigo, and V. Stoven, "Extracting sets of chemical substructures and protein domains governing drug-target interactions," *Journal of chemical information and modeling*, vol. 51, no. 5, pp. 1183-1194, 2011.
- [<sup>2</sup>] C. Knox *et al.*, "DrugBank 3.0: a comprehensive resource for 'omics' research on drugs," *Nucleic acids research*, vol. 39, no. suppl\_1, pp. D1035-D1041, 2010.
- [<sup>3</sup>] D. S. Wishart *et al.*, "DrugBank: a knowledgebase for drugs, drug actions and drug targets," *Nucleic acids research*, vol. 36, no. suppl\_1, pp. D901-D906, 2007.
- [<sup>4</sup>] R. D. Finn *et al.*, "The Pfam protein families database: towards a more sustainable future," *Nucleic acids research*, vol. 44, no. D1, pp. D279-D285, 2016.