

1. Down TA, Piipari M, Hubbard TJP. Dalliance: interactive genome viewing on the web. *Bioinforma Oxf Engl*. 2011 Mar 15;27(6):889–890. PMID: PMC3051325
2. Yates A, Akanni W, Amode MR, Barrell D, Billis K, Carvalho-Silva D, Cummins C, Clapham P, Fitzgerald S, Gil L, Girón CG, Gordon L, Hourlier T, Hunt SE, Janacek SH, Johnson N, Juettemann T, Keenan S, Lavidas I, Martin FJ, Maurel T, McLaren W, Murphy DN, Nag R, Nuhn M, Parker A, Patricio M, Pignatelli M, Rahtz M, Riat HS, Sheppard D, Taylor K, Thormann A, Vullo A, Wilder SP, Zadissa A, Birney E, Harrow J, Muffato M, Perry E, Ruffier M, Spudich G, Trevanion SJ, Cunningham F, Aken BL, Zerbino DR, Flicek P. Ensembl 2016. *Nucleic Acids Res*. 2016 Jan 4;44(D1):D710-716. PMID: PMC4702834
3. Zhou X, Maricque B, Xie M, Li D, Sundaram V, Martin EA, Koebe BC, Nielsen C, Hirst M, Farnham P, Kuhn RM, Zhu J, Smirnov I, Kent WJ, Haussler D, Madden PAF, Costello JF, Wang T. The Human Epigenome Browser at Washington University. *Nat Methods*. 2011 Dec;8(12):989–990. PMID: PMC3552640
4. Gene Ontology Consortium. Gene Ontology Consortium: going forward. *Nucleic Acids Res*. 2015 Jan;43(Database issue):D1049-1056. PMID: PMC4383973
5. Ashburner M, Ball CA, Blake JA, Botstein D, Butler H, Cherry JM, Davis AP, Dolinski K, Dwight SS, Eppig JT, Harris MA, Hill DP, Issel-Tarver L, Kasarskis A, Lewis S, Matese JC, Richardson JE, Ringwald M, Rubin GM, Sherlock G. Gene ontology: tool for the unification of biology. The Gene Ontology Consortium. *Nat Genet*. 2000 May;25(1):25–29. PMID: PMC3037419
6. Eilbeck K, Lewis SE, Mungall CJ, Yandell M, Stein L, Durbin R, Ashburner M. The Sequence Ontology: a tool for the unification of genome annotations. *Genome Biol*. 2005;6(5):R44. PMID: PMC1175956
7. Clough E, Barrett T. The Gene Expression Omnibus Database. *Methods Mol Biol Clifton NJ*. 2016;1418:93–110. PMID: PMC4944384
8. Smith B, Ashburner M, Rosse C, Bard J, Bug W, Ceusters W, Goldberg LJ, Eilbeck K, Ireland A, Mungall CJ, OBI Consortium, Leontis N, Rocca-Serra P, Ruttenberg A, Sansone S-A, Scheuermann RH, Shah N, Whetzel PL, Lewis S. The OBO Foundry: coordinated evolution of ontologies to support biomedical data integration. *Nat Biotechnol*. 2007 Nov;25(11):1251–1255. PMID: PMC2814061
9. Groza T, Köhler S, Moldenhauer D, Vasilevsky N, Baynam G, Zemojtel T, Schriml LM, Kibbe WA, Schofield PN, Beck T, Vasant D, Brookes AJ, Zankl A, Washington NL, Mungall CJ, Lewis SE, Haendel MA, Parkinson H, Robinson PN. The Human Phenotype Ontology: Semantic Unification of Common and Rare Disease. *Am J Hum Genet*. 2015 Jul 2;97(1):111–124. PMID: PMC4572507
10. Global Alliance for Genomics and Health. GENOMICS. A federated ecosystem for sharing genomic, clinical data. *Science*. 2016 Jun 10;352(6291):1278–1280. PMID: 27284183
11. Terry SF. The global alliance for genomics & health. *Genet Test Mol Biomark*. 2014 Jun;18(6):375–376. PMID: 24896853
12. Paten B, Diekhans M, Druker BJ, Friend S, Guinney J, Gassner N, Guttman M, Kent WJ, Mantey P, Margolin AA, Massie M, Novak AM, Nothhaft F, Pachter L, Patterson D, Smuga-Otto M, Stuart JM, Van't Veer L, Wold B, Haussler D. The NIH BD2K center for big data in translational genomics. *J Am Med Inform Assoc JAMIA*. 2015 Nov;22(6):1143–1147. PMID: 26174866
13. Bejerano G, Pheasant M, Makunin I, Stephen S, Kent WJ, Mattick JS, Haussler D. Ultraconserved elements in the human genome. *Science*. 2004 May 28;304(5675):1321–1325. PMID: 15131266