

Short CV
Toby James Gibson

Date of Birth: 06/07/58

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Positions held

- 1977-1981** University of Edinburgh, Scotland (BSc., 1st class Hons. in Molecular Biology)
1981-1986 MRC Laboratory of Molecular Biology, Cambridge, England
1981-1984 PhD. (Supervisor B.G. Barrell) Studies on the Epstein-Barr Virus Genome
1984-1986 Postdoctoral Position with S. Brenner
1986-2015 European Molecular Biology Laboratory (EMBL), Heidelberg, Germany
1986-1991 Postdoctoral Position with P. Argos
1991-1995 Independent Staff Scientist within K. Leonard's group
1996-2015 Computational Biology Team Leader

Grant Consortia

- ELM (FP5, coordinator)
ProteomeBinders (FP6)
EMBRACE (FP6)
DiGtoP (BMBF-NGFN)
SyBoSS (FP7)
SysCilia (FP7)

Editorial Boards

- 2011 - Now Science Signaling (AAAS)
1998 - Now Trends in Biochemical Sciences (Elsevier).
2000 - 2005 Comparative and Functional Genomics (Wiley).
1990 - 2008 DNA Sequence (Harwood Academic Publishers).

Scientific Advisory Board

Jalview alignment software (Dundee)

Selected highly cited publications (Web of Science, 10/4/2015)

- 42,486 Thompson, J. D., Higgins, D. G. and Gibson, T. J. (1994) Clustal-W - Improving The Sensitivity Of Progressive Multiple Sequence Alignment Through Sequence Weighting, Position-Specific Gap Penalties And Weight Matrix Choice. *NAR*, **22**, 4673.
- 26,456 Thompson, J. D. et al. (1997) The CLUSTAL_X windows interface: flexible strategies for multiple sequence alignment aided by quality analysis tools. *NAR*, **25**, 4876.
- 2,538 Biggin, M. D., Gibson, T. J. and Hong, G. F. (1983) Buffer Gradient Gels And S³⁵ Label As An Aid To Rapid DNA-Sequence Determination. *PNAS*, **80**, 3963.
- 1,843 Baer et al. (1984) DNA-Sequence And Expression Of The B95-8 Epstein-Barr Virus Genome. *Nature*, **310**, 207.
- 655 Aasland, R., Gibson, T. J. and Stewart, A. F. (1995) The PHD Finger – implications for chromatin-mediated transcriptional regulation. *TIBS*, **20**, 56.
- 494 Linding, R. et al. GlobPlot: exploring protein sequences for globularity and disorder. *NAR*, **31**, 3701.
- 431 Puntervoll, P. et al. (2003) ELM server: A new resource for investigating short functional sites in modular proteins. *NAR*, **31**, 3625.