

## References [ edit ]

1. ^ Barrangou R, Fremaux C, Deveau H, Richards M, Boyaval P, Moineau S, et al. (March 2007). "CRISPR provides acquired resistance against viruses in prokaryotes". *Science*. 315 (5819): 1709–1712. Bibcode:2007Sci...315.1709B. doi:10.1126/science.1138140. PMID 17379808.
2. ^ Pourcel C, Salvignol G, Vergnaud G (March 2005). "CRISPR elements in *Yersinia pestis* acquire repeats by preferential uptake of bacteriophage DNA, provide additional tools for evolutionary study". *Microbiology*. 151 (Pt 3): 653–663. doi:10.1099/mic.0.27437-0. PMID 15758212.
3. ^ Hwang WY, Fu Y, Reyon D, Maeder ML, Tsai SQ, Sander JD, Peterson RT, Yeh JR, Joung JK (March 2013). "Efficient genome editing in zebrafish using a CRISPR-Cas system". *Nature Biotechnology*. 31 (3): 227–229. doi:10.1038/nbt.2501. PMC 3686313. PMID 23360964.
4. ^ Chylinski K, Jinek M, Fonfara I, Hauer M, Doudna JA, Charpentier E (August 2012). "A programmable dual-DNA-guided RNA endonuclease in adaptive bacterial immunity". *Science*. 337 (6096): 816–821. Bibcode:2012Sci...337..816J. doi:10.1126/science.1225829. PMC 6286148. PMID 22745249.
5. ^ Doe J, Dae J, Bolotin A, Quinquis B, Sorokin A, Ehrlich SD (August 2005). "Clustered regularly interspaced short palindromic repeats (CRISPRs) have spacers of extrachromosomal origin". *Microbiology*. 151 (Pt 8): 2551–2561. doi:10.1099/mic.0.28048-0. PMID 16079334.
6. ^ Cong L, Ran FA, Cox D, Lin S, Barretto R, Hsu PD, Habib N, Wu X, Jiang W, Marraffini LA, Zhang F (February 2013). "Multiplex genome engineering using CRISPR/Cas systems". *Science*. 339 (6121): 819–823. Bibcode:2013Sci...339..819C. doi:10.1126/science.1231143. PMC 3795411. PMID 23287718.

Verbatim match for "CRISPR provides acquired resistance against viruses in prokaryotes", "10.1126/science.1138140" or "17379808".

Correctly matching "CRISPR elements in *Yersinia pestis* acquire new repeats by preferential uptake of bacteriophage DNA, and provide additional tools for evolutionary studies" by Pourcel, Salvignol, and Vergnaud with a normalized edit distance  $\leq 0.2$  (0.072).

Incorrectly matching "Efficient genome editing in plants using a CRISPR/Cas system" by Feng, Zhang, Ding, Liu, Yang, Wei, Cao, Zhu, Zhang, Mao, and Zhu with a normalized edit distance  $\leq 0.2$  (0.118).

Looking for "A programmable dual-RNA-guided DNA endonuclease in adaptive bacterial immunity" by Jinek, Chylinski, Fonfara, Hauer, Doudna, and Charpentier. Author ratio score (1.0) and Jaccard index (1.0) ignore inversion of first and second author, whereas author order score penalizes it heavily (-0.479).

Looking for "Clustered regularly interspaced short palindromic repeats (CRISPRs) have spacers of extrachromosomal origin" by Bolotin, Quinquis, Sorokin, and Ehrlich. Author ratio score (1.0) misses incorrect author list, Jaccard index (0.667) and author order score (-0.413) penalize surplus authors and incorrect order respectively.

Looking for "Multiplex genome engineering using CRISPR/Cas systems" by Cong, Ran, Cox, Lin, Barretto, Habib, Hsu, Wu, Jiang, Marraffini, and Zhang. Author ratio score (1.0) and Jaccard index (1.0) are oblivious to change in order, author order score tolerates inversion of sixth and seventh author (0.865).