

Publication list, Michael Hofreiter

1. Publications in scientific journals

Peer-reviewed research articles

2016

1. Fortes, G.G., Grandal-d'Anglade, A., Kolbe, B., Fernandes, D., Meleg, I.N., García-Vázquez, A., Pinto-Llona, A.C., Constantin, S., de Torres, T.J., Ortiz, J.E., Frischauf, C., Rabeder, G., **Hofreiter, M** and Barlow, A. (2016) Ancient DNA reveals differences in behaviour and sociality between brown bears and extinct cave bears. *Molecular Ecology*, published online 10th August 2016, doi: 10.1111/mec.13800.
2. Wutke, S., Andersson, L., Benecke, N., Sandoval-Castellanos, E., Gonzalez, J., Hallsson, J.H., Lõugas, L., Magnell, O., Morales-Muniz, A., Orlando, L., Pálsdóttir, A.H., Reissmann, M., Muñoz-Rodríguez, M.B., Ruttkay, M., Trinks, A., **Hofreiter, M.** and Ludwig A. (2016) The origin of ambling horses. *Current Biology*, **26** (15), R697-9 (published online 8th August 2016, doi: 10.1016/j.cub.2016.07.001).
3. Gallego-Llorente, M., Connell, S., Jones, E.R., Merrett, D.C., Jeon, Y., Eriksson, A., Siska, V., Gamba, C., Meiklejohn, C., Beyer, R., Jeon, S., Cho, Y.S., **Hofreiter, M.**, Bhak, J., Manica, A. and Pinhasi R. (2016) The genetics of an early Neolithic pastoralist from the Zagros, Iran. *Scientific Reports*, **6**, 31,326 (published online 9th August 2016, doi: 10.1038/srep31326).
4. Speller, C., van den Hurk, Y., Charpentier, A., Rodrigues, A., Gardeisen, A., Wilkens, B., McGrath, K., Rowsell, K., Spindler, L., Collins, M. and **Hofreiter M.** (2016) Barcoding the largest animals on Earth: ongoing challenges and molecular solutions in the taxonomic identification of ancient cetaceans. *Philos Trans R Soc Lond B Biol Sci.*, **371** (1702), pii: 20150332 (doi: 10.1098/rstb.2015.0332).
5. Elsner, J., Deschler-Erb, S., Stopp, B., **Hofreiter, M.**, Schibler, J. and Schlumbaum, A. (2016) Mitochondrial d-loop variation, coat colour and sex

- identification of Late Iron Age horses in Switzerland. *Journal of Archaeological Science: Reports* **6**, 386–396 (doi: dx.doi.org/10.1016/j.jasrep.2016.03.007).
6. Mohandesan, E., Speller, C.F., Peters, J., Uerpmann, H.P., Uerpmann, M., De Cupere, B., **Hofreiter, M.** and Burger, P.A. (2016) Combined hybridization capture and shotgun sequencing for ancient DNA analysis of extinct wild and domestic dromedary camel. *Molecular Ecology Resources*, **published online 11th June 2016**, doi: **10.1111/1755-0998.12551**.
 7. Almathen, F., Charruau, P., Mohandesan, E., Mwacharo, J.M., Orozco-terWengel, P., Pitt, D., Abdussamad, A.M., Uerpmann, M., Uerpmann, H.P., De Cupere, B., Magee, P., Alnaqeeb, M.A., Salim, B., Raziq, A., Dessie, T., Abdelhadi, O.M., Banabazi, M.H., Al-Eknah, M., Walzer, C., Faye, B., **Hofreiter, M.**, Peters, J., Hanotte, O., Burger, P.A. (2016) Ancient and modern DNA reveal dynamics of domestication and cross-continental dispersal of the dromedary. *Proc Natl Acad Sci USA*, **113** (24), 6707-6712 (published online 9th May 2016, doi: 10.1073/pnas.1519508113).
 8. Rainford, J.L., **Hofreiter, M.** and Mayhew, P.J. (2016) Phylogenetic analyses suggest that diversification and body size evolution are independent in insects. *BMC Evol Biol*, **16**, 8 (doi: 10.1186/s12862-015-0570-3).

2015

9. Jones, E.R., Gonzalez-Fortes, G., Connell, S., Siska, V., Eriksson, A., Martiniano, R., McLaughlin, R.L., Gallego-Llorente, M., Cassidy, L.M., Gamba, C., Meshveliani, T., Bar-Yosef, O., Müller, W., Belfer-Cohen, A., Matskevich, Z., Jakeli, N., Higham, T.F., Currat, M., Lordkipanidze, D., **Hofreiter, M.**, Manica, A., Pinhasi, R. and Bradley, D.G. (2016) Upper Palaeolithic genomes reveal deep roots of modern Eurasians. *Nat Commun* **6**, 8912 (doi: 10.1038/ncomms9912).
10. Gallego-Llorente, M., Jones, E.R., Eriksson, A., Siska, V., Arthur, K.W., Arthur, J.W., Curtis, M.C., Stock, J.T., Coltorti, M., Pieruccini, P., Stretton, S., Brock, F., Higham, T., Park, Y., **Hofreiter, M.**, Bradley, D.G., Bhak, J., Pinhasi, R. and Manica, A. (2015) Ancient Ethiopian genome reveals

extensive Eurasian admixture throughout the African continent. *Science* **350** (6262), 820-822 (published online 8th Oct 2015; doi: 10.1126/science.aad2879).

11. Li, C., Corrigan, S., Yang, L., Straube, N., Harris, M., **Hofreiter, M.**, White, W. and Naylor, G.J.P. (2015) DNA capture reveals transoceanic gene flow in endangered river sharks. *Proc. Natl. Acad. Sci. U S A*, **112** (43), 13,302-13,307 (doi: 10.1073/pnas.1508735112; published online 12th Oct 2015).
12. Le Duc, D., Renaud, G., Krishnan, A., Almén, M.S., Huynen, L., Prohaska, S.J., Ongyerth, M., Bitarello, B.D., Schiöth, H.B., **Hofreiter, M.**, Stadler, P.F., Prüfer, K., Lambert, D., Kelso, J. and Schöneberg, T. (2015) Kiwi genome provides insights into evolution of a nocturnal lifestyle. *Genome Biology*, **16**, 147 (published online 23rd July 2015, doi: 10.1186/s13059-015-0711-4).
13. Pinhasi, R., Fernandes, D., Sirak, K., Novak, M., Connell, S., Alpaslan-Roodenberg, S., Gerritsen, F., Moiseyev, V., Gromov, A., Raczyk, P., Anders, A., Pietruszewsky, M., Rollefson, G., Jovanovic, M., Trinhhoang, H., Bar-Oz, G., Oxenham, M., Matsumura, H. and **Hofreiter, M.** (2015) Optimal Ancient DNA Yields from the Inner Ear Part of the Human Petrous Bone. *PLoS One*, **10** (6), e0129102 (published online 18th June 2015, doi: 10.1371/journal.pone.0129102 eCollection 2015).
14. Springer, M.S., Signore, A.V., Paijmans, J.L., Vélez-Juarbe, J., Domning, D.P., Bauer, C.E., He, K., Crerar, L., Campos, P.F., Murphy, W.J., Meredith, R.W., Gatesy, J., Willerslev, E., MacPhee, R.D., **Hofreiter, M.** and Campbell K.L. (2015) Interordinal gene capture, the phylogenetic position of Steller's sea cow based on molecular and morphological data, and the macroevolutionary history of Sirenia. *Molecular Phylogenetics and Evolution*, **91**, 178-193 (published online 4th June 2015, doi: 10.1016/j.ympev.2015.05.022).
15. Paijmans, J.L.A., Fickel, J., Courtiol, A., **Hofreiter, M.** and Förster, D.W. (2015) Impact of enrichment conditions on cross-species capture of fresh and degraded DNA. *Molecular Ecology Resources*, **16** (1), 42-55 (published online 27th April 2015, doi: 10.1111/1755-0998.12420).
16. Alter, S. E., Meyer M., Post, K., Czechowski, P., Gravlund, P., Gaines, C., Rosenbaum, H., Kaschner, K., Turvey, S.T., van der Plicht, J., Shapiro, B. and **Hofreiter, M.** (2015) Climate impacts on transocean dispersal and habitat in

- gray whales from the Pleistocene to 2100. *Molecular Ecology*, **24** (7), 1510-1522 (published online 9th March 2015, doi: 10.1111/mec.13121).
17. Welcker, F., Collins, M.J., Thomas, J.A., Wadsley, M., Brace, B., Cappellini, E., Turvey, S.T., Reguero, M., Gelfo, J.N., Kramarz, A., Burger, J., Thomas-Oates, J., Ashford, D.A., Ashton, P., Rowsell, K., Porter, D.M., Kessler, B., Fisher, R., Baessmann, C., Kaspar, S., Olsen, J., Kiley, P., Elliot, J., Kelstrup, C., Mullin, V., **Hofreiter, M.**, Willerslev, E., Hublin, J.-J., Orlando, L., Barnes I. & MacPhee, R.D.E. (2015) Ancient proteins resolve the evolutionary history of Darwin's South American ungulates. *Nature*, **522** (7554), 81-84 (published online 18th March 2015, doi: 10.1038/nature14249).
18. Ludwig, A., Reissmann, M., Benecke, N., Bellone, R., Sandoval-Castellanos, E., Cieslak, M., Fortes, G.G., Morales-Muñiz, A., **Hofreiter, M.** and Pruvost, M. (2015) Twenty-five thousand years of fluctuating selection on leopard complex spotting and congenital night blindness in horses. *Phil Trans R Soc B*, **370** (1660), 20130386 (doi: 10.1098/rstb.2013.0386).
19. Teasdale, M.D., van Doorn, N.L., Fiddymment, S., Webb, C., O'Connor, T., **Hofreiter, M.**, Collins, M.J. and Bradley, D.G. (2015) Paging through history: parchment as a reservoir of ancient DNA for next generation sequencing. *Phil Trans R Soc B*, **370** (1660), 20130379 (doi: 10.1098/rstb.2013.0379).
20. Elsner, J., Schibler, J., **Hofreiter, M.** and Schlumbaum, A. (2015) Burial condition is the most important factor for mtDNA PCR amplification success in Palaeolithic equid remains from the Alpine foreland. *Archaeological and Anthropological Sciences*, **1**-11 (DOI 10.1007/s12520-014-0213-4).
21. Elsner, J., Schibler, J., **Hofreiter, M.** and Schlumbaum, A. (2015) Burial condition is the most important factor for mtDNA PCR amplification success in Palaeolithic equid remains from the Alpine foreland. *Archaeological and Anthropological Sciences*, **1**-11 (DOI 10.1007/s12520-014-0213-4).

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22. Schubert, M., Jónsson, H., Chang, D., Der Sarkissian, C., Ermini, L., Ginolhac, A., Albrechtsen, A., Dupanloup, I., Foucal, A., Petersen, B., Fumagalli, M., Raghavan, M., Seguin-Orlando, A., Korneliussen, T.,

- Velazquez, A.M.V., Stenderup, J., Hoover, C.A., Rubin, C.J., Alfarhan, A.H., Alquraishi, S.A., Al-Rasheid, K.A.S., MacHugh, D.E., Kalbfleish, T., MacLeod, J.N., Rubin, E.M., Sicheritz-Ponten, T., Andersson, L., **Hofreiter, M.**, Marques-Bonet, T., Gilbert, M.T.P., Nielsen, R., Excoffier, L., Willerslev, E., Shapiro, B. and Ludovic Orlando, L. (2014) Prehistoric horse genomes reveal the genetic foundation and cost of domestication. *Proc. Natl. Acad. Sci. U S A*, **111** (52), E5661-9 (published online 15th December 2014; doi: 10.1073/pnas.1416991111).
23. Xiang, H., Yuan, J., Gao, J., Yu, B., Luo, Y., Zhou, H., Cai, D., Zhang, Y., Chen, X., Wang, X., **Hofreiter, M.** and Zhao, X. (2014) Early Holocene chicken domestication in northern China. *Proc. Natl. Acad. Sci. U S A*, **111** (49), 17564-9 (published online 24th November 2014; doi: 10.1073/pnas.1411882111).
24. King, T.E., Gonzalez Fortes, G., Balaresque, P., Thomas, M.G., Balding, D., Maisano Delser, P., Neumann, R., Parson, W., Knapp, M., Walsh, S., Tonasso, L., Holt, J., Kayser, M., Appleby, J., Forster, P., Ekserdjian, D., **Hofreiter M.** and Kevin Schürer, K. (2014) Identification of the remains of King Richard III. *Nature Communications*, **5**, 5631 (published online 2nd December; doi: 10.1038/ncomms6631).
25. Rainford, J.L., **Hofreiter, M.**, Nicholson, D.B. and Mayhew, P.J. (2014) Phylogenetic distribution of extant richness suggests metamorphosis is a key innovation driving diversification in insects. *PLoS One*, **9** (9), e109085.
26. Gamba, C., Jones, E.R., Teasdale, M.D., McLaughlin, R.L., Gonzalez-Fortes, G., Mattiangeli, V., Domboróczki, L., Kővári, I., Pap, I., Anders, A., Whittle, A., Dani, J., Raczyk, P., Higham, T.F.G., **Hofreiter, M.**, Bradley, D.G., and Pinhasi, R., (2014) Genome flux and stasis in a five millennium transect of European prehistory. *Nature Communications*, **5**, 5257 (published online 21st Oct 2014; doi: 10.1038/ncomms6257).
27. Mitchell, K.J., Pratt, R.C., Watson, L.N., Gibb, G.C., Llamas, B., Kasper, M., Edson, J., Hopwood, B., Male, D., Armstrong, K., Meyer, M., **Hofreiter, M.**, Austin, J., Donnellan, S.C., Lee, M.S.Y., Phillips, M.J. and Cooper, A. (2014) Molecular phylogeny, biogeography, and habitat preference evolution of

- marsupials. *Molecular Biology and Evolution*, **31** (9), 2322-2330 (published online 30th May 2014; doi: 10.1093/molbev/msu176).
28. Huynen, L., Suzuki, T., Ogura, T., Watanabe, Y., Millar, C.D., **Hofreiter, M.**, Smith, C., Mirmoeini, S. and Lambert, D.M. (2014) Reconstruction and in vivo analysis of the extinct tbx5 gene from ancient wingless moa (Aves: Dinornithiformes). *BMC Evol Biol*, **14**, 75 (published online 14th May 2014; doi: 10.1186/1471-2148-14-75).
 29. Horn, S., Prost, S., Stiller, M., Makowiecki, D., Kusnetsova, T., Benecke, N., Pucher, E., Hufthammer, A.K., Schouwenburg, C., Shapiro, B. and **Hofreiter, M.** (2014) Ancient mitochondrial DNA and the genetic history of Eurasian beaver (*Castor fiber*) in Europe. *Molecular Ecology*, **23**, 1717–1729 (published online 12th February 2014; doi: 10.1111/mec.12691).
 30. Stiller, M., Molak, M., Prost, S., Pacher, M., Rabeder, G., Baryshnikov, G., Rosendahl, W., Münzel, S., Bocherens, H., Grandal-d'Anglade, A., Hilpert, B., Germonpre, M., Stasyk, O., Pinhasi, R., Rohland, N., Mohandesan, E., Ho, S.Y.W., **Hofreiter, M.** and Knapp, M. (2014) Mitochondrial DNA diversity and evolution of the Pleistocene cave bear complex. *Quaternary International*, **339-340**, 224-231 (published online 5th November 2013; <http://dx.doi.org/10.1016/j.quaint.2013.09.023>).

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31. Zhang, H., Paijmans, J.L.A., Chang, F., Wu, X., Chen, G., Lei, C., Yang, X., Wei, Z., Bradley, D.G., Orlando, L., O'Connor, T. and **Hofreiter, M.** (2013) Morphological and genetic evidence for early Holocene cattle management in North-Eastern China. *Nature Communications*, **4**, 2755 (published online 8th November 2013; doi: 10.1038/ncomms3755).
32. Bellone, R.B., Holl, H., Setaluri, V., Devi, S., Maddodi, N., Archer, S., Sandmeyer, L., Ludwig, A., Foerster, D., Pruvost, M., Reissmann, M., Bortfeldt, R., Adelson, D.L., Lim, S.L., Nelson, J., Haase, B., Engensteiner, M., Leeb, T., e Forsyth, G., Mienaltowski, M.J., **Hofreiter, M.**, Paijmans, J.L.A., Gonzalez-Fortes, G., Grahn, B. and Brooks, S.A. (2013) RNA sequencing identifies retroviral insertion in TRPM1 as the cause of congenital

- stationary night blindness and leopard complex spotting in the horse. *PLoS One*, **8** (10), e78280 (published online 22nd October 2013; doi: 10.1371/journal.pone.0078280).
33. Finstermeier, K., Zinner, D., Bräuer, M., Meyer, M., Kreuz E. **Hofreiter, M.** and Roos, C. (2013) A mitogenomic phylogeny of living primates. *PLoS One*, **8** (7), e69504 (published online 16th July 2013; doi: 10.1371/journal.pone.0069504).
 34. Orlando, L., Ginolhac, A., Zhang, G. and 47 co-authors, **Hofreiter, M.**, Nielsen, N., Shapiro, B., Jun, W. and Willerslev, E. (2013) Recalibrating *Equus* evolution using the genome sequence of an Early Middle Pleistocene horse. *Nature*, **499** (7456), 74-78 (published online 26th June 2013; doi: 10.1038/nature12323).
 35. Bocherens, H., Bridault, A., Drucker, D.G., **Hofreiter, M.**, Münzel, S.C., Stiller, M. and van der Plicht, J. (2013) The last of its kind? Radiocarbon, ancient DNA and stable isotope evidence from a late cave bear (*Ursus spelaeus* ROSENmüller, 1794) from Rochedane (France). *Quaternary International*, **339-340**, 179-188 (published online 21st May 2013; <http://dx.doi.org/10.1016/j.quaint.2013.05.021>).
 36. Li, C., **Hofreiter, M.**, Straube, N., Corrigan, S. and Naylor, G.J.P. (2013) Capturing protein-coding genes across highly divergent species. *Biotechniques*, **54**, 321-326 (doi: 10.2144/000114039).
 37. Prost, S., Guralnick, R.P., Waltari, E., Fedorov, V.B., Kuzmina, E., Smirnov, N., van Kolfschoten, T., **Hofreiter, M.** and Vrieling, K. (2013) Losing ground: past history and future fate of Arctic small mammals in a changing climate. *Glob Chang Biol*, **19** (6), 1854-1864 (published online 31st January; doi: 10.1111/gcb.12157).
 38. Prost, S., Klietmann, J., van Kolfschoten, T., Guralnick, R.P., Waltari, E., Vrieling, K., Stiller, M., Nagel, D., Rabeder, G., **Hofreiter, M.** and Sommer, R.S. (2013) Effects of late quaternary climate change on Palearctic shrews. *Glob Chang Biol*, **19** (6), 1865-1874 (published online 28th January; doi: 10.1111/gcb.12153).

- 39.** Haus, T., Akom, E., Agwanda, B., **Hofreiter, M.**, Roos, C. and Zinner, D. (2013) Mitochondrial diversity and distribution of African green monkeys (*Chlorocebus* Gray, 1870). *Am J Primatol.* **75** (4), 350-360 (published online 10th January 2013; doi: 10.1002/ajp.22113).

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- 40.** Bruche, S., Gusset, M., Lippold, S., Eulenberger, K., Junhold, J., Driscoll, C. A. and **Hofreiter, M.** (2012) A genetically distinct lion (*Panthera leo*) population from Ethiopia. *Eur J Wildl Res* **59**, 215–225 (published online 9th October 2012; doi: 10.1007/s10344-012-0668-5).
- 41.** Jenkins, D.L., Davis, L., Stafford, Jr., T.W., Campos, P., Hockett, B., Jones, G.T., Cummings, L.S., Yost, C., Connolly, T.J., Yohe II, R.M., Gibbons, S.C., Raghavan, M., Rasmussen, M., Paijmans, J.L.A., **Hofreiter, M.**, M. Kemp, B.M., Barta, J.L., Cara Monroe, C., Gilbert, M.T.P. and Willerslev, E. (2012) Clovis Age Western Stemmed projectile points and human coprolites at the Paisley Caves. *Science*, **337** (6091), 223-8.
- 42.** Baca, M., Stankovic, A., Marciszak, A., **Hofreiter, M.**, Nadachowski, A., Stefaniak, K., Węgleński, P. and Mackiewicz, P. (2012) Genetic Analyses of Cave Bear specimens from Niedźwiedzia Cave, Sudetes, Poland. *Palaeontologia Electronica*, **15** (2), 21A, 16p.

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- 43.** Gravlund, P., Aaris-Sørensen, K., **Hofreiter, M.**, Meyer, M., Bollback J.P. and Noe-Nygaard N. (2011) Ancient DNA extracted from Danish aurochs (*Bos primigenius*): genetic diversity, genealogical history and preservation. *Annals of Anatomy*, **194** (1), 103-111 (published online 17th November).
- 44.** Lippold, S., Matzke, N., Reißman, M., and **Hofreiter, M.** (2011) Whole mitochondrial genome sequencing of domestic horses reveals incorporation of extensive wild horse diversity during domestication. *BMC Evol Biol*, **11**, 328 (published online 14th November).

45. Lorenzen, E., Noguès-Bravo, D., Orlando, L., Weinstock, J., Binladen, J., Marske, K. A., Haile, J., and 47 co-authors, Wayne, R. K., Cooper, A., **Hofreiter, M.**, Sher, A., Shapiro, B., Rahbek, C. and Willerslev, E. (2011) Individualistic species responses to climate and humans determine Late Quaternary megafaunal extinction and survival. *Nature*, **479** (7373), 359-64 (published online 2nd November; doi: 10.1038/nature10574).
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47. Lerner, H. R. L., Meyer, M., James, H. F., **Hofreiter, M.** and Fleischer, R. C. (2011) Multilocus resolution of phylogeny and timescale in the extant adaptive radiation of Hawaiian honeycreepers. *Current Biology*, **21** (21), 1838-1844 (published online 21st October 2011; doi: 10.1016/j.cub.2011.09.039).
48. Orlando, L., Ginolhac, A., Raghavan, M., Vilstrup, J., Rasmussen, M., Magnussen, K., Steinmann, K., Kapranov, P., Thompson, J. F., Zazula, G., Froese, D., Moltke, I., Shapiro, B., **Hofreiter, M.**, AL-Rasheid, K. A. S., Mundy, J., M. Gilbert1, M. T. P. and Willerslev, E. (2011) True Single-Molecule DNA Sequencing of a Pleistocene Horse bone. *Genome Research*, **21** (10), 1705-19 (published online 29th July 2011).
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50. Yuan, Y., Shen, T. J., Gupta, P., Ho, N. T., Simplaceanu, V., Tam, T. C., **Hofreiter, M.**, Cooper, A., Campbell, K. L. and Ho, C. (2011) A biochemical-biophysical study of hemoglobins from woolly mammoth, asian elephant, and humans. *Biochemistry* **50** (34), 7350-7360 (published online 2nd August 2011).

51. Münzel, S. C., Stiller, M., **Hofreiter, M.**, Mitnik, A., Conard, N. J. and Bocherens, H. (2011) Pleistocene bears in the Swabian Jura (Germany): genetic replacement, ecological displacement, extinctions and survival. *Quaternary International*, **245** (2), 225-237 (published online 22nd April 2011; doi:10.1016/j.quaint.2011.03.060).
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53. Horn, S., Durka, W., Wolf, R., Ermala, A., Stubbe, A., Stubbe, M. and **Hofreiter, M.** (2011) Mitochondrial genomes reveal slow rates of molecular evolution and the timing of speciation in beavers (*Castor*), one of the largest rodent species. *PLoS One*, **6** (1): e14622 (published online 28th January 2011, doi: 10.1371/journal.pone.0014622).

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55. Cieslak, M., Pruvost, M., Benecke, N., **Hofreiter, M.**, Morales, A., Reißmann M. and Ludwig, A. (2010) Origin and history of mitochondrial DNA lineages in domestic horses. *PLoS One*, **5** (12): e15311 (published online 20th December 2010, doi: 10.1371/journal.pone.0015311).
56. Rohland, N., Reich, D., Mallick, S., Meyer, M., Green, R.E., Georgiadis, N.J., Roca, A., and **Hofreiter, M.** (2010) Genomic DNA sequences from mastodon and woolly mammoth reveal deep speciation of forest and savanna elephants. *PLoS Biology*, **8** (12), e1000564 (published online 21st December 2010, doi:10.1371/journal.pbio.1000564).

- 57.** Campos, P.F., Kristensen, T., Orlando, L., Sher, A., Kholodova, M.V., Götherström, A., **Hofreiter, M.**, Drucker, D.G., Kosintsev, P., Tikhonov, A., Baryshnikov, G. F., Willerslev, E. and Gilbert, M.T.P. (2010) Ancient DNA sequences point to a large loss of genetic diversity in the saiga antelope (*Saiga tatarica*) since the Pleistocene. *Molecular Ecology*, **19** (22), 4863-4875 (published online 27th September 2010, doi: 10.1111/j.1365-294X.2010.04826.x).
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