Abstract Submitted for the DNP12 Meeting of The American Physical Society

Noninvasive Online Measurement of Genome Lengths of Mammalian Tissues in Bulk by 14 MeV Neutron Atometry BOGDAN MAGLICH, ANNA RADOVIC, CHRISTIAN DRUEY, California Science & Engineering Corporation — Genome length, L=, no. of DNA nucleotide base pairs in cell of bovine (b) and porcine (p) tissues, closest to human genome, were hitherto measured by genomic sequencing  $L_b=3$ ,  $L_p=2.7$  Giga base pairs [1,2] (Gbp) errors not given. - We report measurements of  $L_b/L_p$  and  $L_b$ ,  $L_p$  without sequencing by atometry [3,4]. No. of O and C atoms, N, in nucleotide molecules, was obtained from prompt  $\gamma$  rate, G, emitted in inel. scatt. 14 MeV neutrons, with nuclei of C, O, in nucleotide molecule. Since G prop. N,  $L_b/L_p=G_b/G_p$ . p and b meat was irradiated for 30'. From msd G we obtained  $L_b/L_p=1.28\pm0.02$  16% greater than [1,2]. We got absolute  $L_b=1.65/f$ ,  $L_p=1.28/f$  Gbp, 0.3<f<0.6 by irradiating 3 DNA's and calibrating  $\gamma$ /O and  $\gamma$ /C.

[1] Y.Liu et al: Bos Taurus genome assembly; BMC Genomics 2009, 10:180
[2] L.B. Schook et al, Swine Genome Sequencing: Comparative and Functional genomics, V. 6, Issue 4, 251, June 2005
[3] Maglich B. et al, http://whitehousedrugpolicy.gov/ctac99/pdfs/kani.pdf

[4] Maglich B.C. AIP Conf. Proc. 796,431(2005);http://link.aip.org/link/?APCPCS/796/431/1

Bogdan Maglich California Science & Engineering Corporation

Date submitted: 02 Jul 2012

Electronic form version 1.4