

Supplementary Material

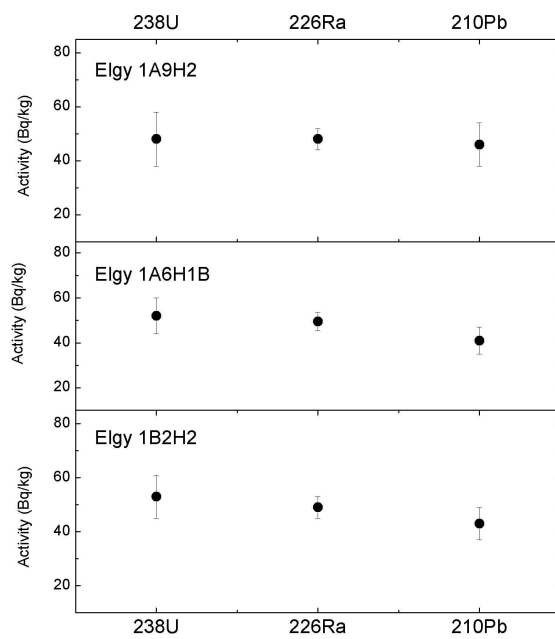


Fig. S1: Activities of the U-series products for the three samples measured at VKTA Rossendorf. If the decay series is in equilibrium the activities of all daughters agree within 2-sigma errors with the activity of the mother. The error bars shown in the graphs represent the 2-sigma errors.

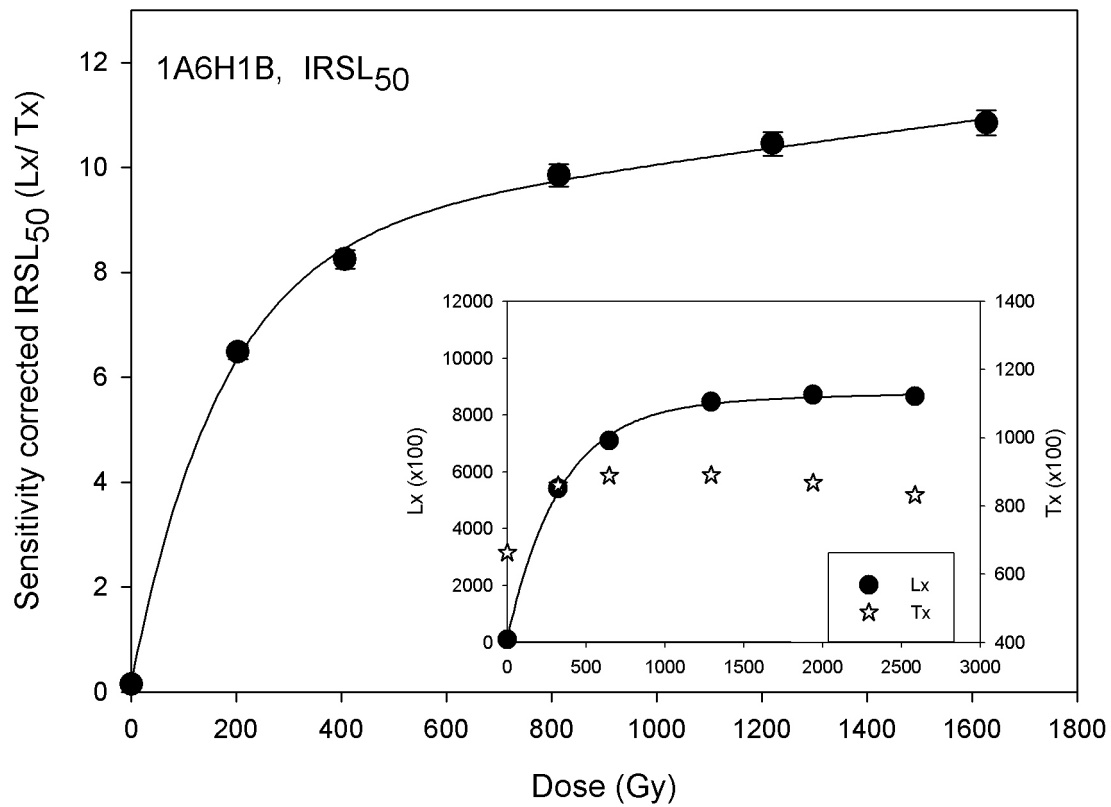


Fig. S2: Dose response curve of sample 1A6H1B fitted to a single exponential plus linear function. The IR₅₀ protocol was applied with 7 dose steps up to a maximum beta dose of ~ 2500 Gy. The deconvolution (inset) of the dose response curve (L_x/T_x) shows a pseudo-increase, i.e the regenerated IRSL₅₀ signal (L_x) is in saturation and the corresponding test dose signal (T_x) is decreasing with each regeneration cycle. The test dose was kept constant.

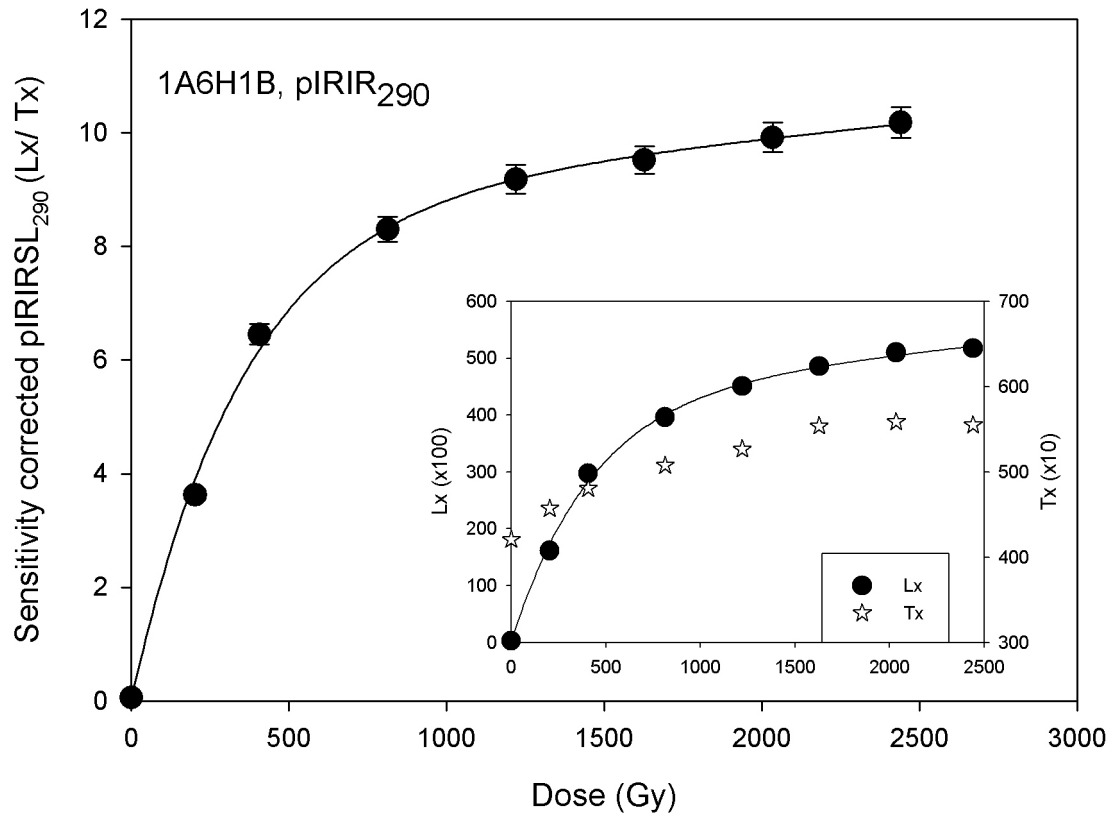


Fig. S3: Saturation dose response curve of sample 1A6H1B fitted to a single exponential plus linear function. The pIRIR₂₉₀ protocol was applied with 7 dose steps up to a maximum beta dose of ~ 2500 Gy. The deconvolution (inset) of the dose response curve (L_x/T_x) shows a real increase with each regeneration cycle, i.e the regenerated pIRIR₂₉₀ signal (L_x) and the corresponding test dose signal (T_x) are increasing. The test dose was kept constant.

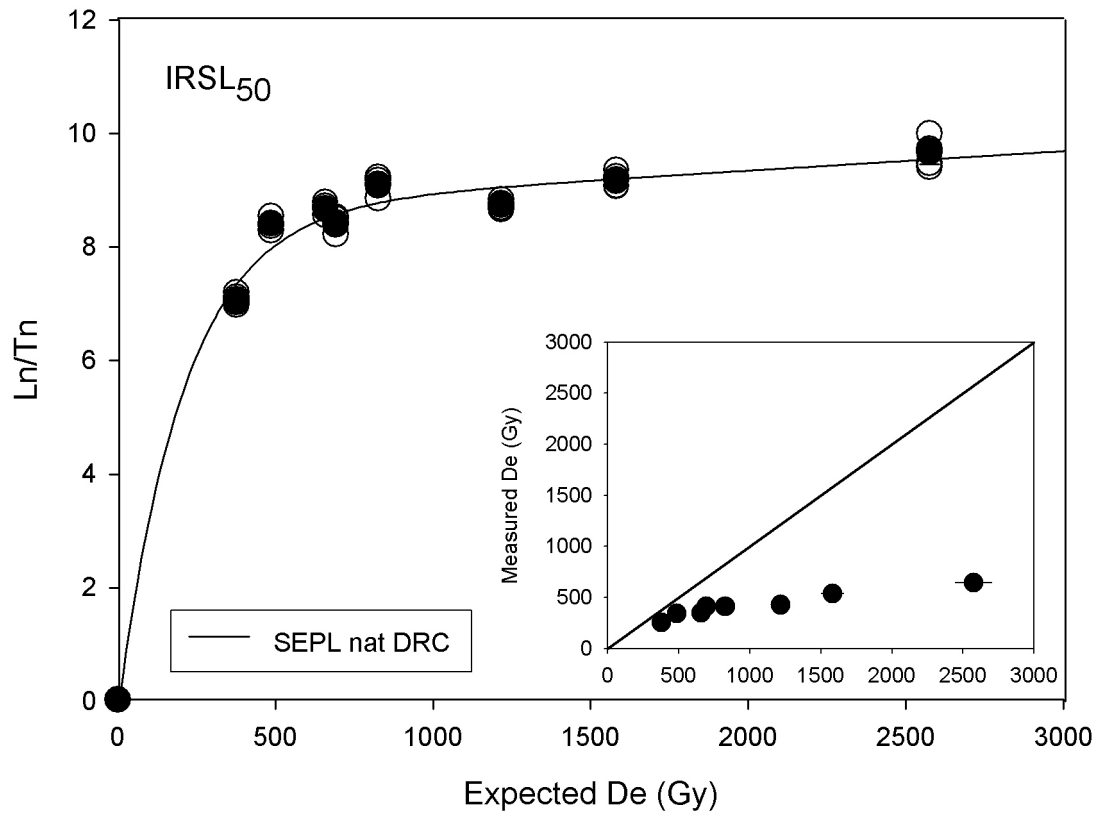


Fig. S4: Test dose normalised natural optically stimulated luminescence signals (L_n/T_n) of the eight samples used for the standard SAR-IRSL protocol plotted against their expected equivalent dose values (D_e) (see Chapot et al., 2012). The expected D_e of each sample was calculated by multiplying the environmental dose rate by the expected age. The inset shows the severe divergence between measured and expected D_e . The solid line is the 1:1 line.

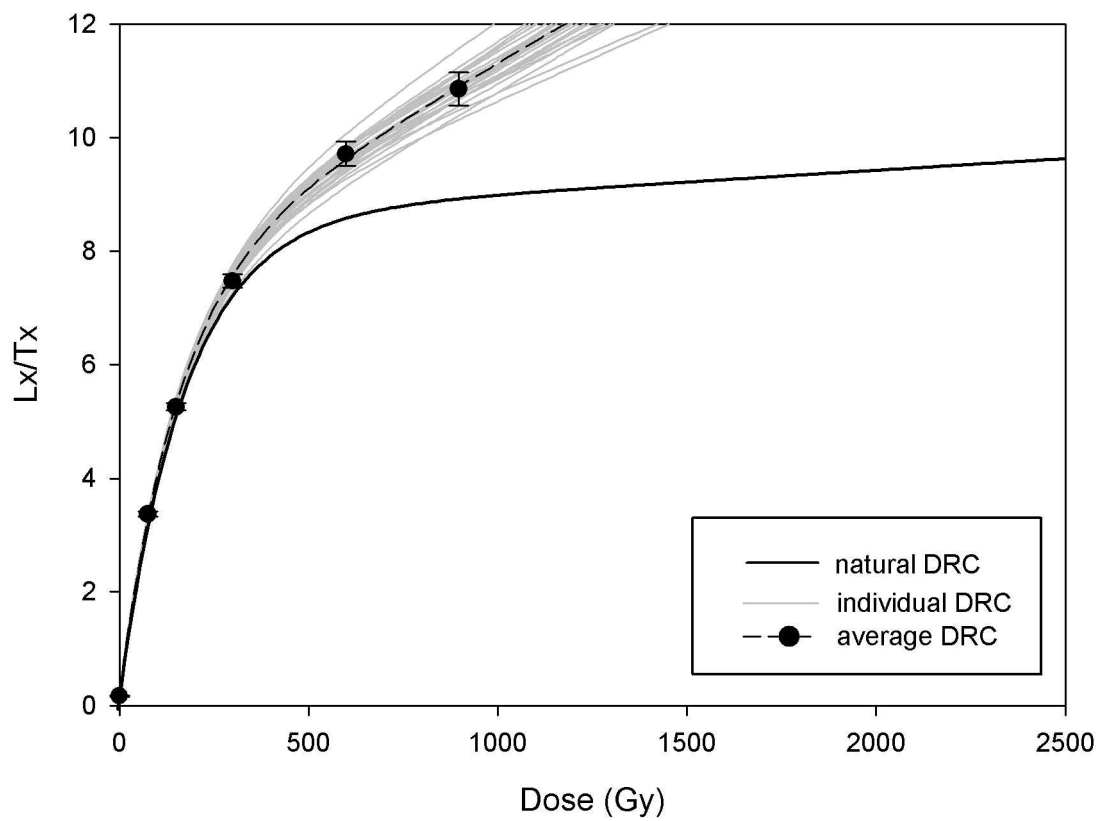


Fig. S5: Comparison of the natural dose response curve (DRC) from Fig. S4 with the single aliquot regenerative DRCs of the aliquots used to construct the natural DRC.