

Leukocyte methylomic imprints of exposure to the genocide against the Tutsi in Rwanda: a pilot epigenome-wide analysis

Musanabaganwa C, Wani AH, Donglasan J, Fatumo S, Jansen S, Mutabaruka J, Rutembesa E, Uwineza A, Hermans EJ, Roozendaal B, Wildman DE, Mutesa L, Uddin M.

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