

A pilot randomised controlled trial of a programme of psychosocial interventions (resettle) for high risk personality disordered offenders

Article

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Figure 1

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1 2 3 4 5 6 7 8 9 10 11

CACTTCACGCAACTCGGCCCGGCGGGCGGCGGGCCACTCACGCAGCTCAGCCGCGGGA

12 13 14 15 16 17 18 19

GGCGCCCCGGCTCTTGTGGCCCGCCCGCTGTCACCCGCAGGGGCACTGGCGGGCGCTTGCCGC

20 21 22 23 24 25

CAAGGGGCAGAGCGAGCTCCCGAGTGGGTCTGGAGCCGCGGAGCTGGGCGGGGGCGGGAAGG

26 27 28 29

AGGTAGCGAGAAAAGAACTGGAGAACTCGGTGGCCCTCTTAACGCCGCCCCAGAGAGACC

Figure 1 Scheme of the human NR3C1 gene analyzed by bisulfite pyrosequencing. The 5'-end of the human NR3C1 gene contains multiple first exons, with multiple transcriptional start sites and mRNA splice variants. The region analyzed by bisulfite pyrosequencing (primer sequences are in bold) contains 29 CpGs and encompasses exon 1-F, which is the human homolog of the rat exon 1-7, previously shown to be differentially methylated [47]