

## PGC-1 $\alpha$ : a potent transcriptional cofactor involved in the pathogenesis of type 2 diabetes

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Unfortunately the colour of the specific nucleotides in the textbox ‘Human haplotypes and utility of the HapMap

Project’ was incorrect. The correct presentation of the sequence is shown below:

The online version of the original article can be found at: <http://dx.doi.org/10.1007/s00125-006-0268-6>

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### Human haplotypes and utility of the HapMap Project

... The genomic region shown below harbours eight SNPs (red) and may theoretically contain 256 SNP combinations or haplotypes. However, phased chromosome analysis may reveal only five common haplotypes that can be unambiguously identified by the typing of four SNPs (bold red).

```
AGTG..TATG..GCAG..GTAC..ATGC..GTGA..ATAC..GATT  
AGTG..TATG..GCAG..GTAC..ATGC..GTGA..ATAC..GACT  
AGTG..TATG..GCAG..GTCC..ATGC..GTGA..ATGC..GATT  
AGTG..TAGG..GCAG..GTAC..ATGC..GTGA..ATAC..GACT  
AGCG..TATG..GCGG..GTAC..ATCC..GTAA..ATAC..GATT
```