# **Receptor activity modifying protein 1 (RAMP1) gene promoter methylation is associated with female** migraine susceptibility

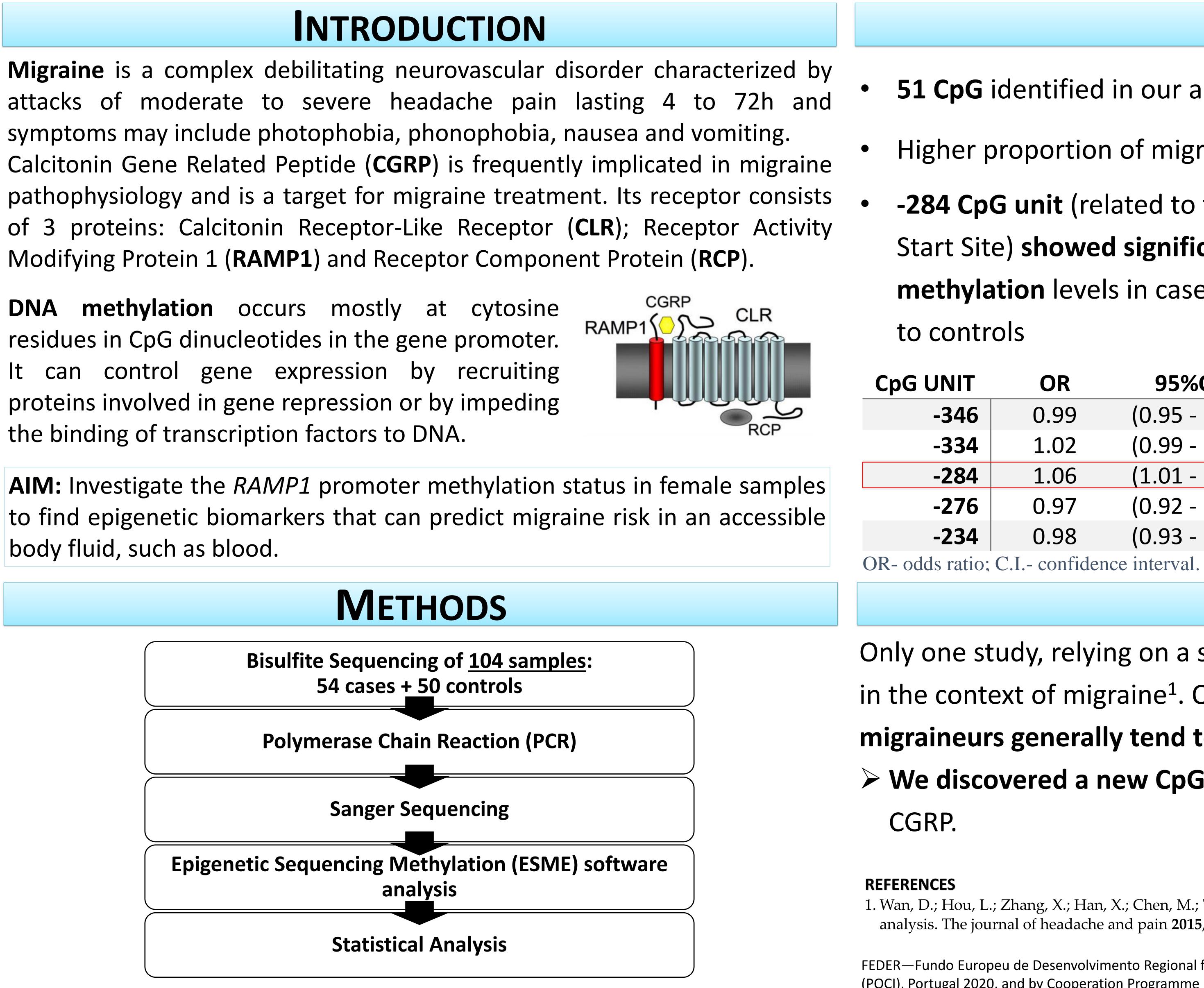
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OR

DNA residues in CpG dinucleotides in the gene promoter. It can control gene expression by recruiting proteins involved in gene repression or by impeding the binding of transcription factors to DNA.

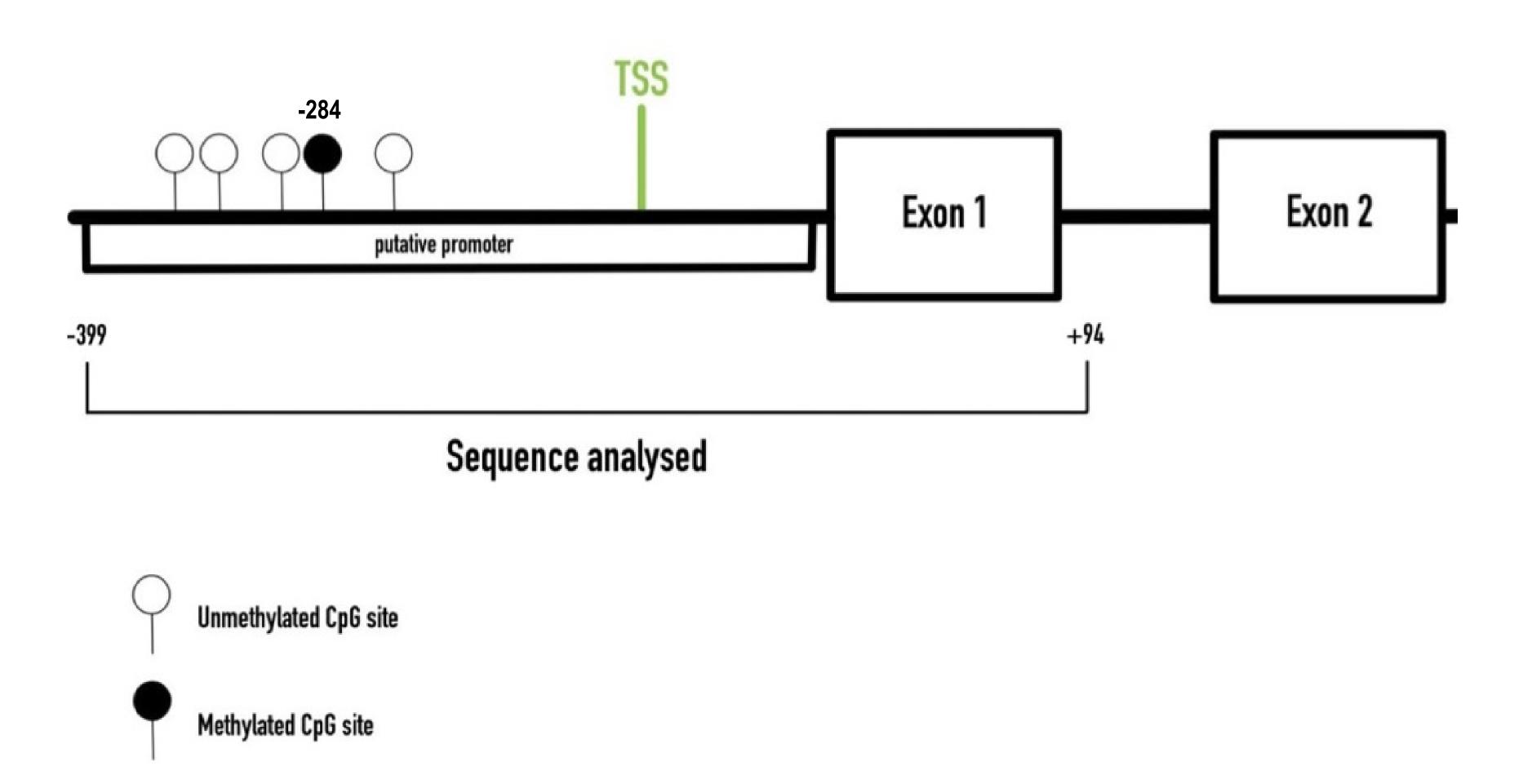
body fluid, such as blood.



- RESULTS
- 51 CpG identified in our analysed sequence only the first 5 showed variability
- Higher proportion of migraine cases with all five CpG units methylated compared to controls (26% vs 16%)

-284 CpG unit (related to the Transcription Start Site) showed significantly higher methylation levels in cases when compared

95%C.I.	<b>P-VALUE</b>
(0.95 - 1.03)	0.582
(0.99 - 1.07)	0.509
(1.01 - 1.12)	0.017*
(0.92 - 1.02)	0.225
(0.93 - 1.03)	0.411
ence interval *n<0.05	



### DISCUSSION

Only one study, relying on a small sample size, has analyzed the methylation of the human RAMP1 promoter in the context of migraine<sup>1</sup>. Our preliminary results seem to contradict that study as we found that **female** migraineurs generally tend to have higher methylation levels than female controls. > We discovered a new CpG unit potentially associated to migraine which may disrupt the transcription of

FEDER—Fundo Europeu de Desenvolvimento Regional funds through the COMPETE 2020—Operational Programme for Competitiveness and Internationalisation (POCI), Portugal 2020, and by Cooperation Programme Interreg V-A Spain–Portugal under project 0702\_MIGRAINEE\_2\_E- POCTEP 2014–2020.





<sup>1.</sup> Wan, D.; Hou, L.; Zhang, X.; Han, X.; Chen, M.; Tang, W.; Liu, R.; Dong, Z.; Yu, S. DNA methylation of RAMP1 gene in migraine: an exploratory analysis. The journal of headache and pain **2015**, 16, doi:10.1186/S10194-015-0576-7.