Corrigendum

Corrigendum to 'Detecting recombination and diversifying selection in human alpha-papillomavirus' [Inf. Gen. Evol. 8 (2008) 689–692]

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The authors regret that Fig. 1 was incorrectly displayed on the published article. Please find below the corrected Fig. 1.
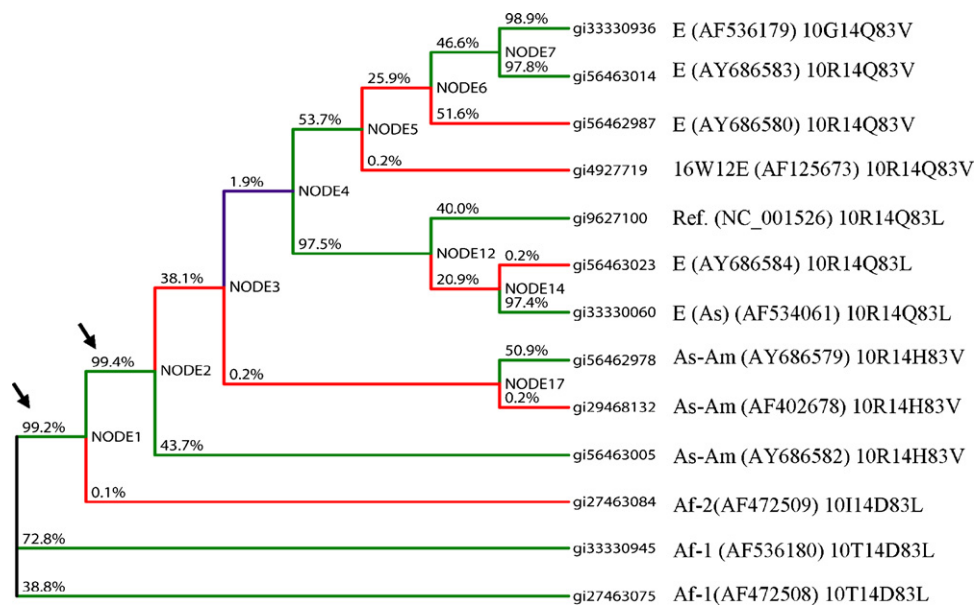

Fig. 1. Variation in selection pressure through time in gene E6 within the HPV16 group. Branch labels represent model averaged probabilities of dN/dS > 1. The composition of the distinct variants for the three positively selected codons (10, 14 and 83) is given.