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Association of adeno-associated virus type 2 with melanomas of squamous cell origin, but not those of epithelial origin. A number of human malignancies, including melanomas, are associated with an increased prevalence of adeno-associated viruses (AAVs). The AAV genome consists of non-recombinant integrated episomal elements. The objectives of this study were to determine the prevalence of these elements and to determine whether DNA from AAV was detected in human cutaneous malignant melanomas of squamous cell origin and in melanomas of epithelial origin. DNA from melanoma and non-melanoma samples was obtained from archival paraffin-embedded material and assayed for the presence of AAV sequences. This was followed by a polymerase chain reaction (PCR) for the recovery and sequencing of AAV sequences. The cutaneous malignancy cohort consisted of one cohort of 20 primary melanomas of non-squamous cell origin (MNP), and five cohorts of squamous cell origin (MNP), and five cohorts of squamous cell origin (MNP), and five cohorts of squamous cell origin (MNP), and 11 melanomas of squamous cell origin (MSPC). DNA from AAV was detected in 70% (14/20) of MSPC and in 0% of NMNP. No amplification products were detected from any of the melanomas of squamous cell origin (MSPC). DNA from AAV was detected in 70% (14/20) of NSPC and in 0% of NMNP. No amplification products were detected from any of the melanomas of squamous cell origin (MSPC). DNA from AAV was detected in 70% (14/20) of NSPC and in 0% of NMNP. No amplification products were detected from any of the melanomas of squamous cell origin (MSPC). DNA from AAV was detected in 70% (14/20) of NSPC and in 0% of NMNP. No amplification products were detected from any of the melanomas of squamous cell origin (MSPC). DNA from AAV was detected in 70% (14/20) of NSPC and in 0% of NMNP. No amplification products were detected from any of the melanomas of squamous cell origin (MSPC). DNA from AAV was detected in 70% (14/20) of NSPC and in 0% of NMNP. No amplification products were

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