



Supporting Information

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DeepVISP: Deep learning for virus site integration prediction
and motif discovery

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Supplementary data for “DeepVISP: Deep learning for virus site integration prediction and motif discovery”

Supplementary Table S1. Full list of virus integration sites of hepatitis B virus (HBV).

Supplementary Table S2. Full list of virus integration sites of human herpesvirus (HPV).

Supplementary Table S3. Full list of virus integration sites of Epstein-Barr virus (EBV).

Supplementary Table S4. List of DNA-binding transcription factors matching the PWMs learnt by DeepVISP for the HBV integration.

Supplementary Table S5. List of DNA-binding transcription factors matching the PWMs learnt by DeepVISP for the HPV integration.

Supplementary Table S6. List of DNA-binding transcription factors matching the PWMs learnt by DeepVISP for the EBV integration.