
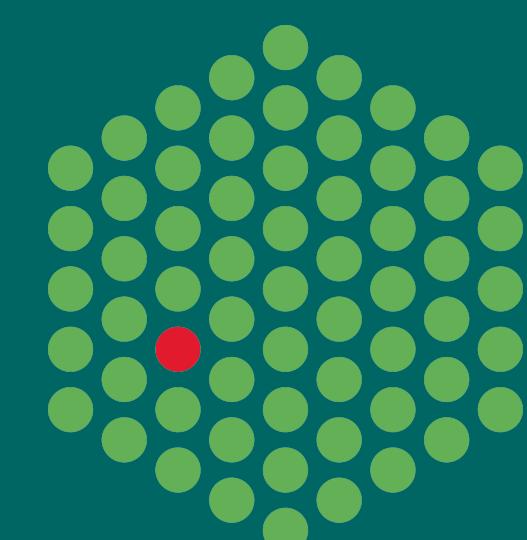


Identifying novel functional linear motifs using host-viral protein interactions and the principle of convergent evolution

**References:**

- All analyses were performed in R. IMEx consortium databases such as IntAct, MINT, DIP, BHF-UCL, MPIDB, MatrixDB, HPIDb, I2D-IMEx, InnateDB-IMEx, MolCon, UniProt, MBLInfo are currently integrated into IntAct [1].
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