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On the cover: Combination of reverse genetic with high-throughput proteomic analysis is a powerful tool for mapping virus-host interactions directly from virus-infected cells. Superposition of (1) background: measles virus-infected cells fuse together to form one single syncytium cell; (2) Vitruvian man and a measles virus particle symbolizing virus-host interactions; and (3) colored balls connected by lines: a map of the human protein interaction sub-network targeted by measles virus V protein. For details, see article by Anastassia V. Komarova *et al.*, pages M110.007443, 1–17.

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