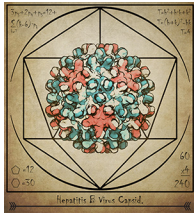


COVER IMAGE



Cover photograph: The hepatitis B virus (HBV) packages its genome in an icosahedral protein shell called a capsid. Eberhard's theorem indicates that a capsid must contain 12 pentamers to form a closed shell (upper left). The HBV capsid is composed of 12 pentamers and 30 hexamers (lower left). The triangulation number of the capsid, $T=4$, indicates that there are four unique positions that constituent proteins can occupy (upper right). There are a total of $T \times 60 = 240$ proteins, altogether 120 dimers, that make up the HBV capsid (lower right). We report mutations in the capsid that enhance normal assembly yet increase antiviral drug resistance. (See related article at e01082-18.) (Copyright © 2018 American Society for Microbiology. All Rights Reserved.)

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AUTHOR CORRECTION

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