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On the cover, pictures show transparent human corneas covering the pupil and colored iris. A dataset of 141 distinct proteins identified in the normal human cornea by mass spectrometry is categorized according to the predominant tissue distribution of the proteins (upper middle, EC, extracellular; IC, intracellular; PM, plasma membrane-bound) and function (lower left, extracellular proteins; lower right, intracellular proteins). For details, see the article by Karring et al., pages 1406–1408.
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