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Cover legend: Pentameric modeling of the human $\alpha 1 \beta$ glycine receptor (GlyR), viewed from extracellular perspective, showing considerable disruption of Cl^- pore architecture in the Y128C hyperekplexia mutant channel. $\alpha 1$ subunits are in green (Y128C mutant) and β subunits in blue. The mutated residue, C128, is shown in magenta and sections 142–145 and 170–173, which adopt altered α conformation, in red. See the article by Chung et al. in this issue (pages 9612–9620).

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