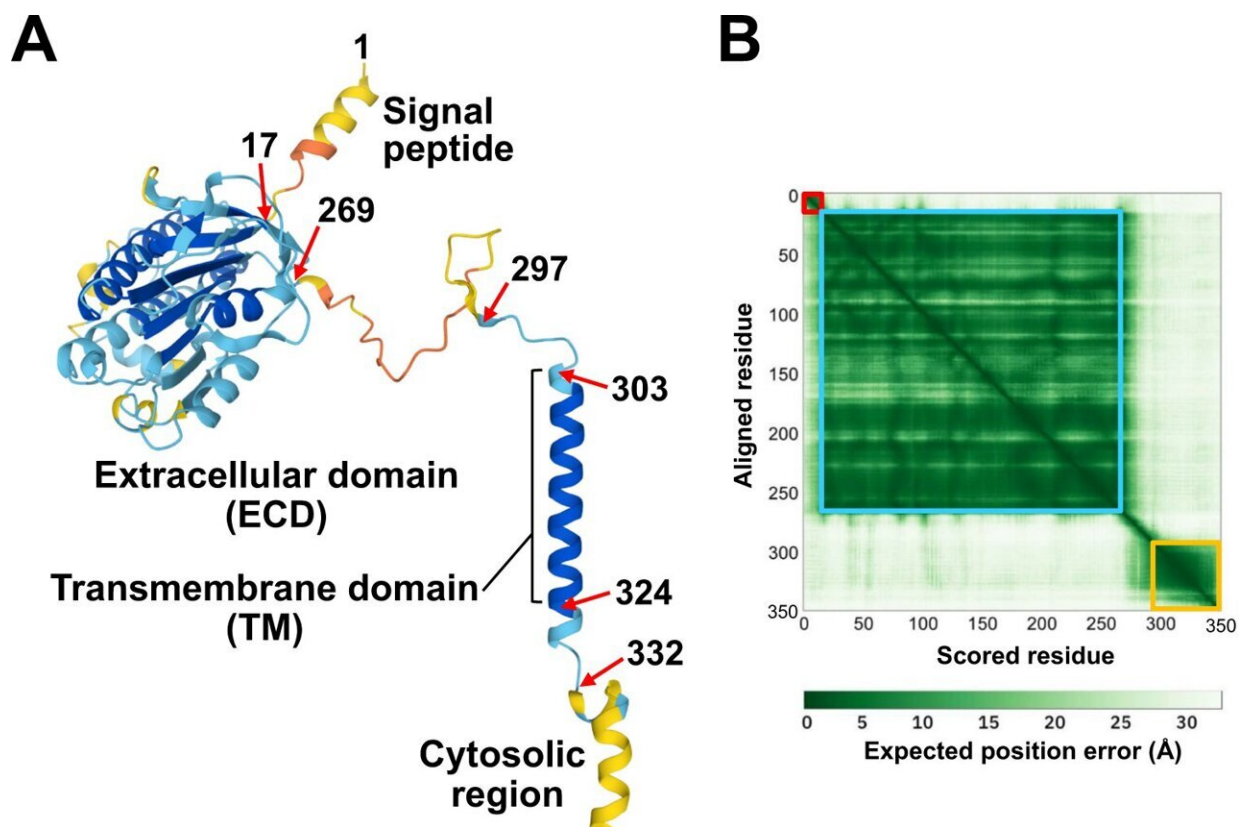


# Highlighting the molecular mechanism underlying pancreatic cancer development

December 9 2022



Predicted structure of human (P)RR. **A** Cartoon representation of the AF2 structural model. The model includes the signal peptide (residues 1–16), extracellular domain (ECD; residues 17–302), transmembrane domain (TM; residues 303–323), and cytosolic region (residues 324–350). The structure is colored according to the predicted local-distance difference test (pLDDT) score in blue, cyan, yellow, and orange for the regions that were most confidently predicted ( $pLDDT > 90$ ), confidently predicted ( $90 > pLDDT > 70$ ), predicted with low confidence ( $70 > pLDDT > 50$ ), and predicted with very low confidence

(pLDDT

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