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The image depicts three related structures: a comparison of coactivator and corepressor binding in PPAR α , (left) active on agonist binding, (centre) corepressor binding after helix-12 displacement by an antagonist, and (right) direct corepressor binding. The full figure is presented on page T10 of this issue.

Credit: Fraydoon Rastinejad, Metabolic Signaling and Disease Program, Sanford-Burnham Medical Research Institute, Orlando, Florida, USA.

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