Steady State Inhibition Metric (SSIM) =
$$\frac{C_{\rm SS}}{C_{\rm SS} + IC_{50}}$$

			1	T .
Name	Binding	TMDD	Competitive TMDD	Tissue TMDD
Tissue	In vitro	Circulation	Circulation	Tissue
Ligand	no	no	yes	no
IC ₅₀	K_{ss}	$K_{ m ss} \cdot T_{ m fold,largeD}$	$K_{\mathrm{ss}} \cdot T_{\mathrm{fold,largeD}} \cdot L_{\mathrm{fold,largeD}}$	$\frac{K_{\rm ss} \cdot T_{\rm fold, largeD}}{B}$
Model	D + T ←	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	$\begin{array}{cccccccccccccccccccccccccccccccccccc$
		1 = central compartment2 = peripheral compartment	D_2	$\begin{array}{c c} D_1 \\ \downarrow \\ \downarrow \\ M \end{array} \Leftrightarrow DM.$
		3 = tissue compartment		D_2
M = Membrane-bound target				
		Γ _{fold,largeD} = max fold-increase in total target from baseline at steady state, for large doses		
		L _{fold,largeD} = max fold-increase in ligand from baseline at steady state, for large doses		
		B = tissue biodistribution coefficient		