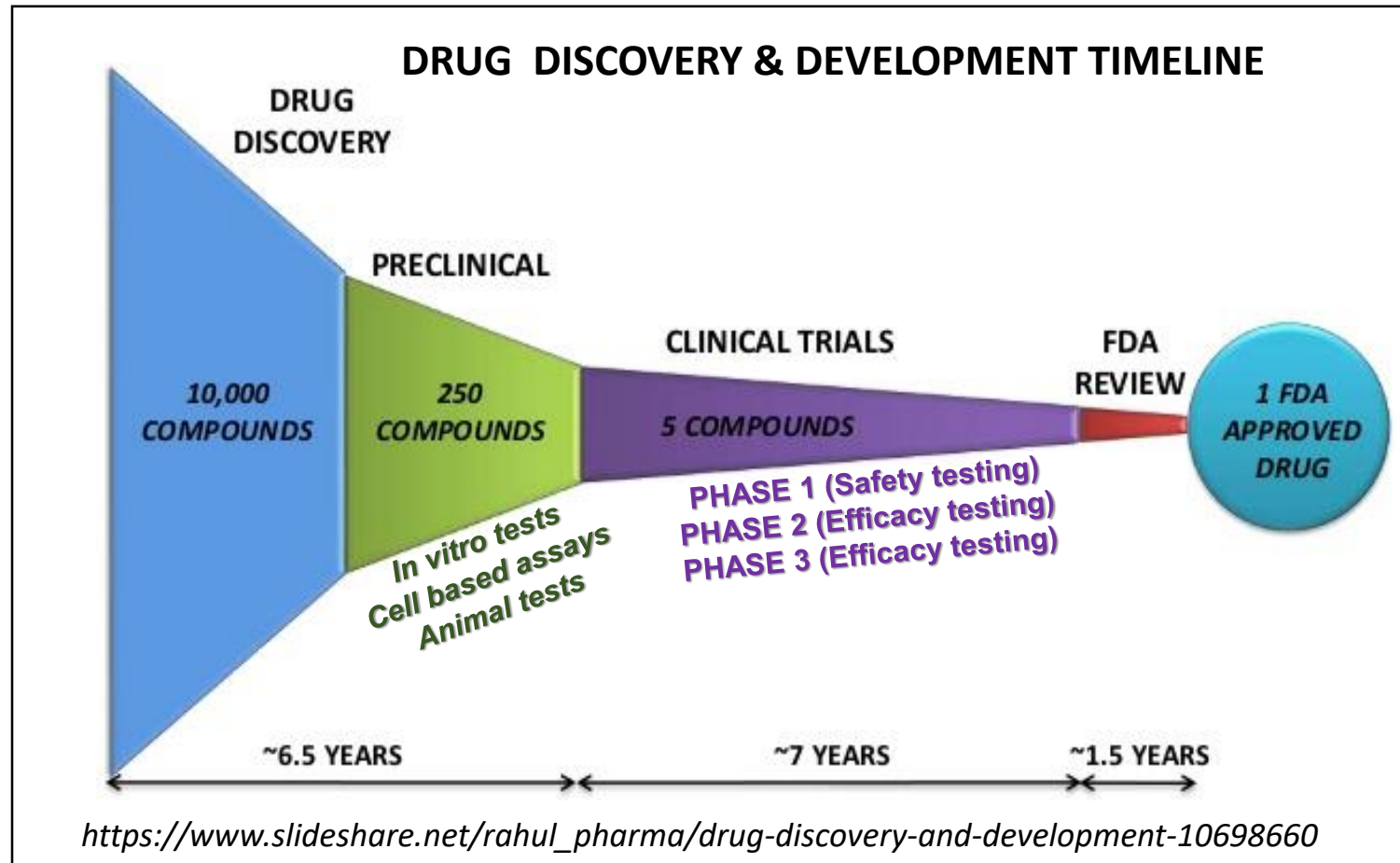


Functional interrogation of Toxcast database for DILI-associated differences in Troglitazone vs Rosiglitazone Maleate

Sricharan Bandhakavi

March 2019

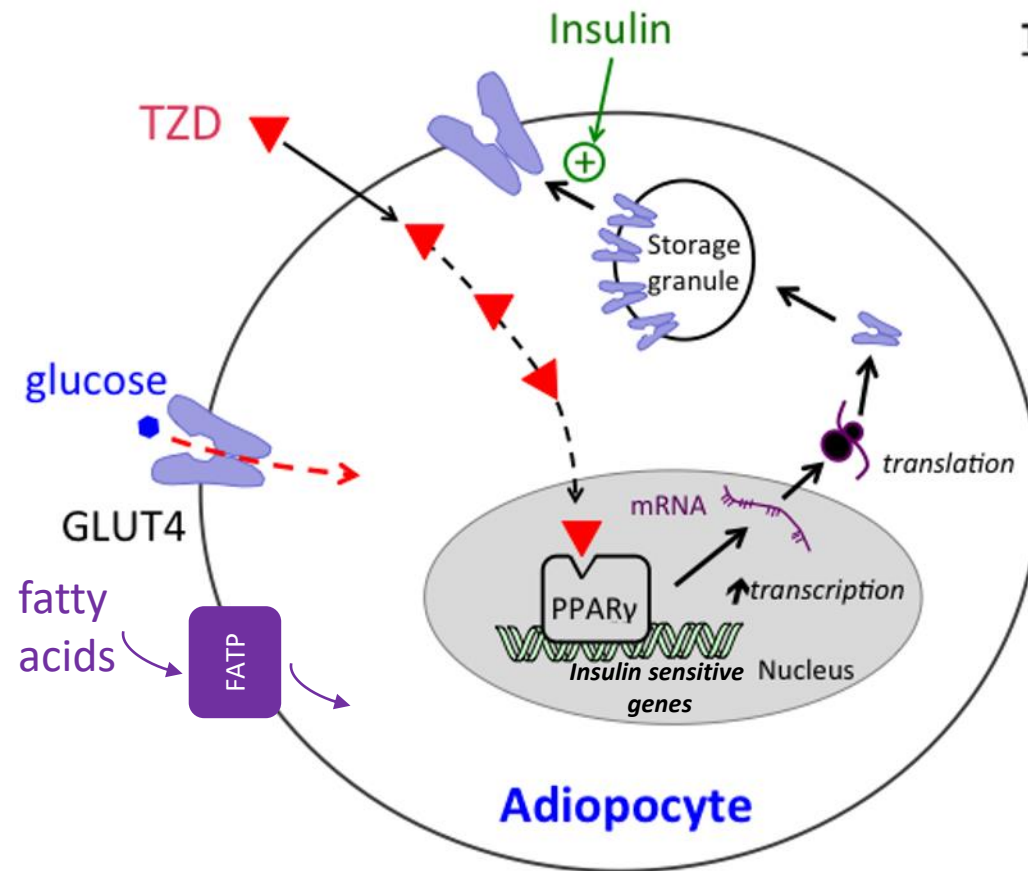
How to better leverage preclinical data for eliminating bad drugs from going into clinical trials/market ?



EBTC GOAL

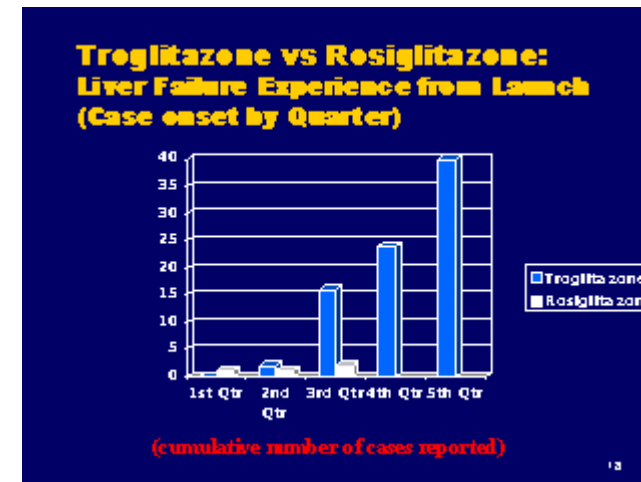
Leverage “preclinical data” for identifying drugs with potential toxicities to liver

Anti-diabetic TZD drugs, Troglitazone and Rosiglitazone, stimulate insulin function by targeting PPAR γ



Thiazolidinediones (TZDs):

(In market since 1999) **Rosiglitazone (maleate)*** – Target: PPAR γ
(withdrawn in year 2000) **Troglitazone**** – Target: PPAR γ > PPAR α



* “lessDILI” drug (LTKB)
** “mostDILI” drug (LTKB)

<https://www.fda.gov/ohrms/dockets/ac/00/slides/3615s1a/sld018.htm>

<http://tmedweb.tulane.edu/pharmwiki/doku.php/thiazolidinediones> &
www.diabetesincontrol.com (Handbook of Diabetes, 4th edition excerpt)

Can we leverage preclinical data (cell-based assays/test results) for Troglitazone to understand potential basis for liver toxicity?

High level project workflow

Extract AC50 information from ToxCast
(Level 5/6) tests for two drugs of interest



From up to 700 cell based assays/tests,
classify all “positive” test results

- Positive with Troglitazone only
- Positive with both Troglitazone and Rosiglitazone Maleate
- Positive with Rosiglitazone Maleate only



Develop schema for differential analysis
of tests/targets affected by each drug

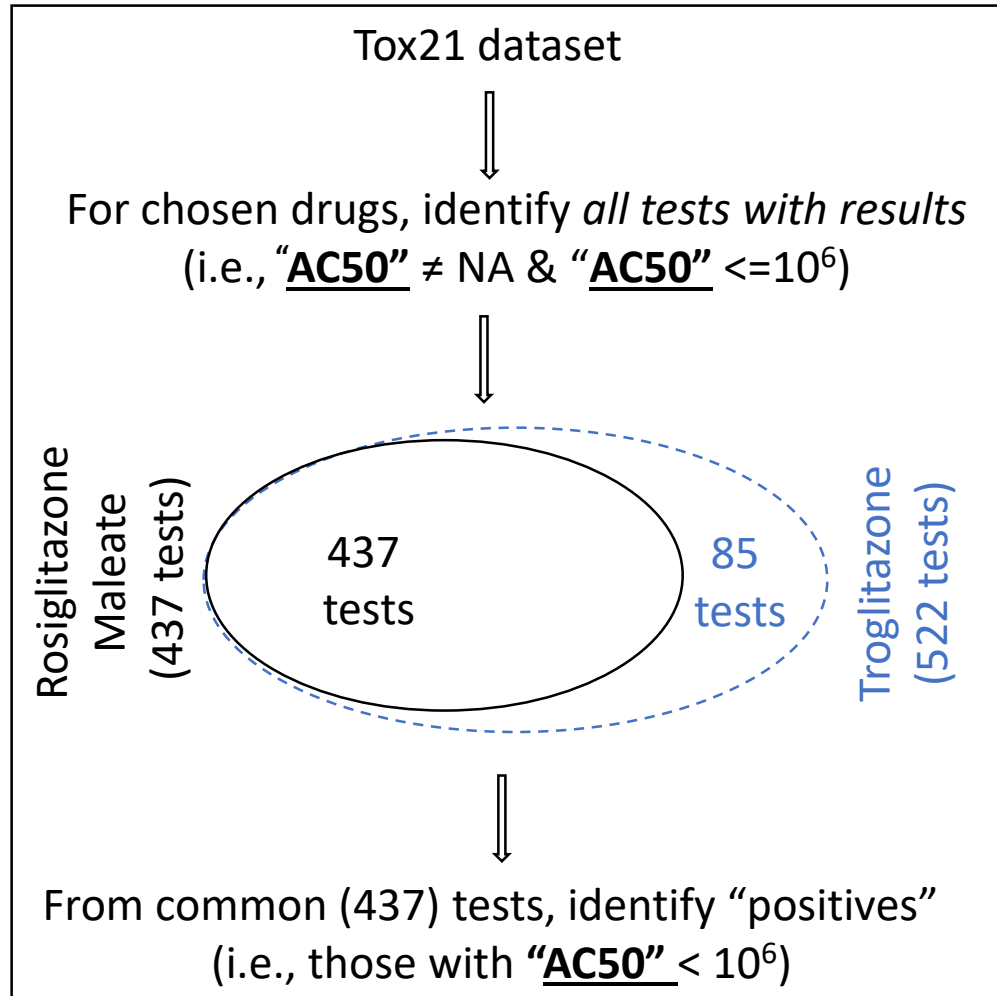
- Generate an “activation score” for potential activation of test “targets” in patients
- Identify differentially affected tests/targets by Troglitazone vs Rosiglitazone Maleate



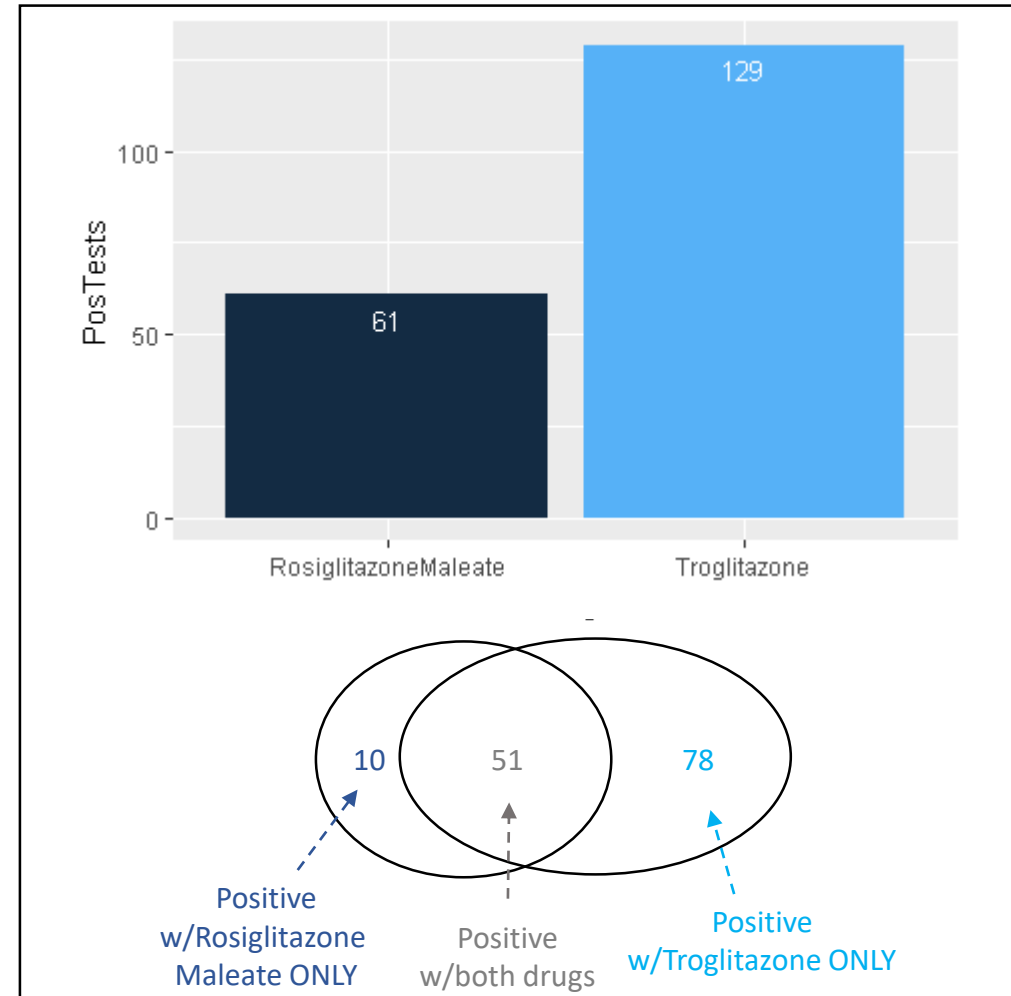
Correlate differentially affected targets with liver toxicity

Higher # of “positive” tests for Troglitazone relative to Rosiglitazone Maleate

Workflow

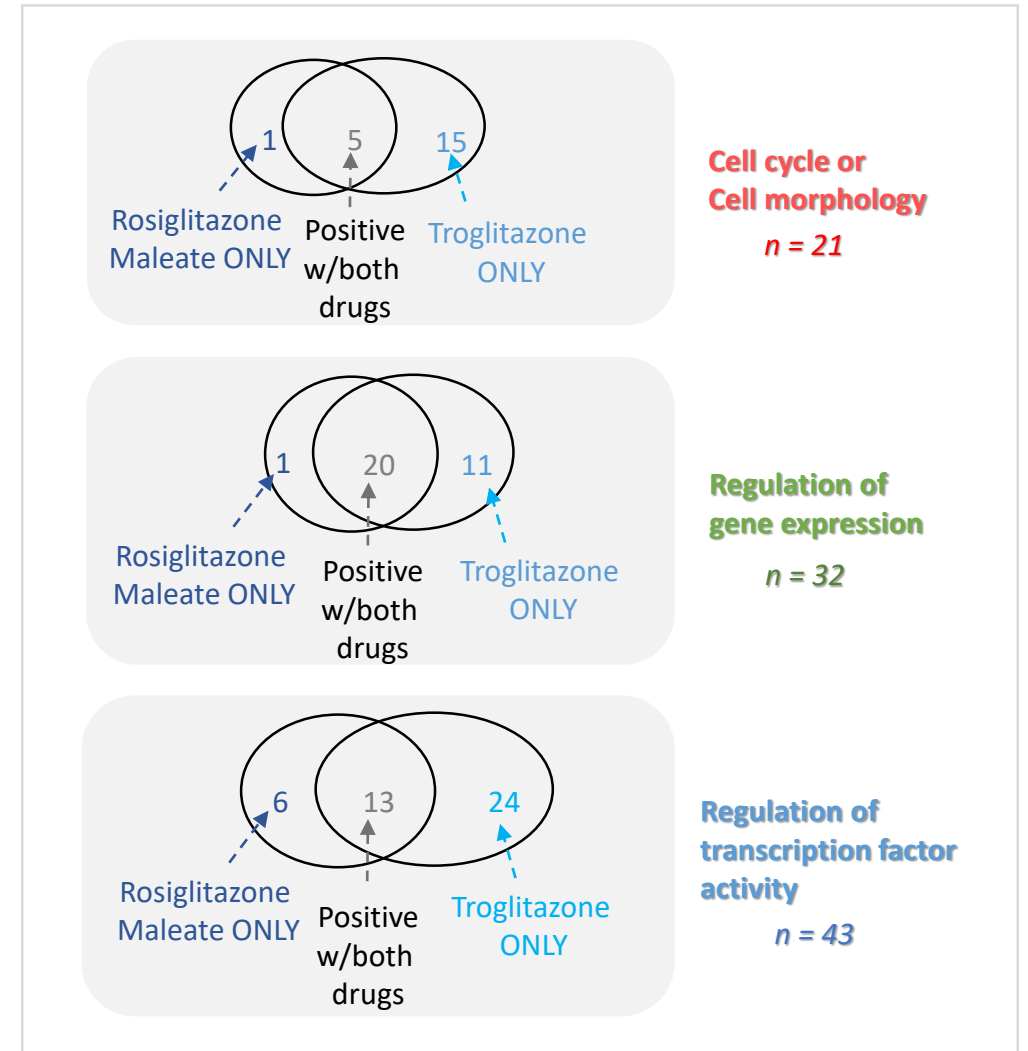
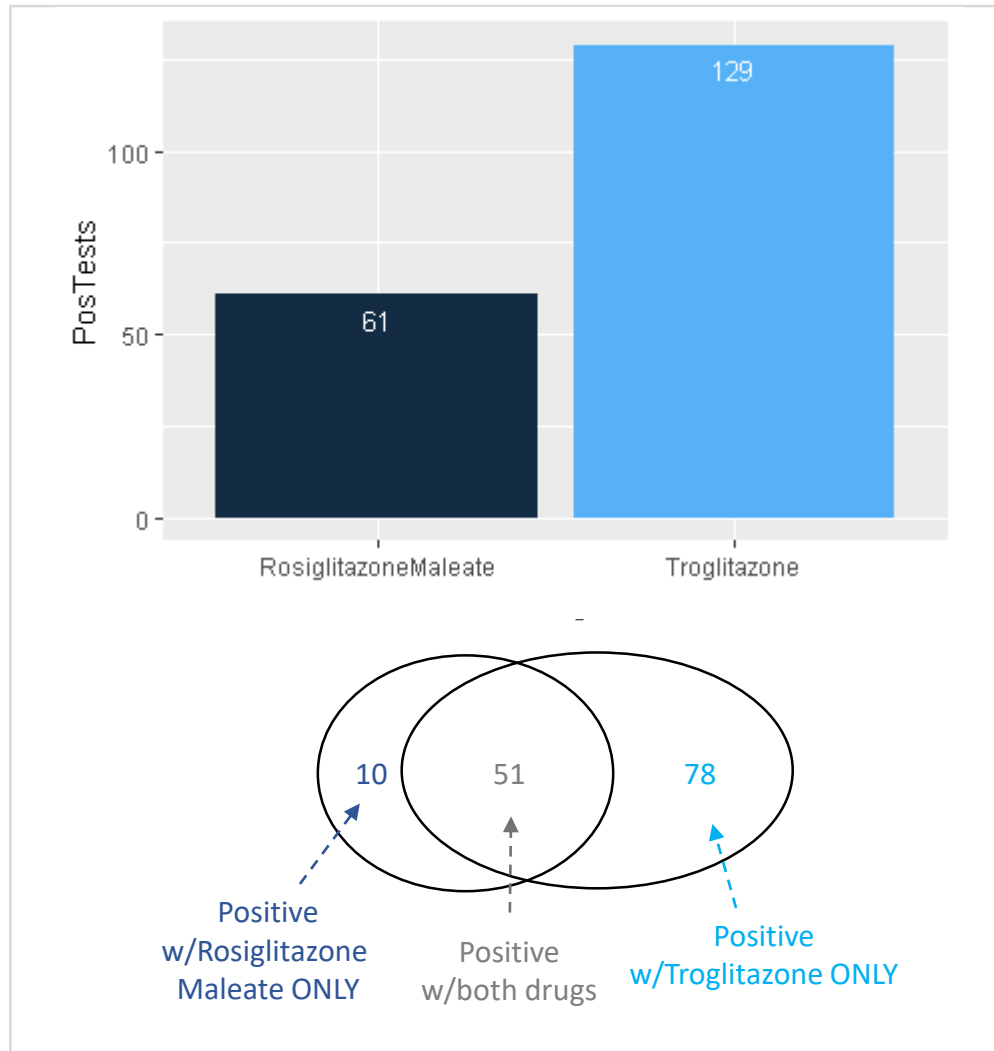


Results



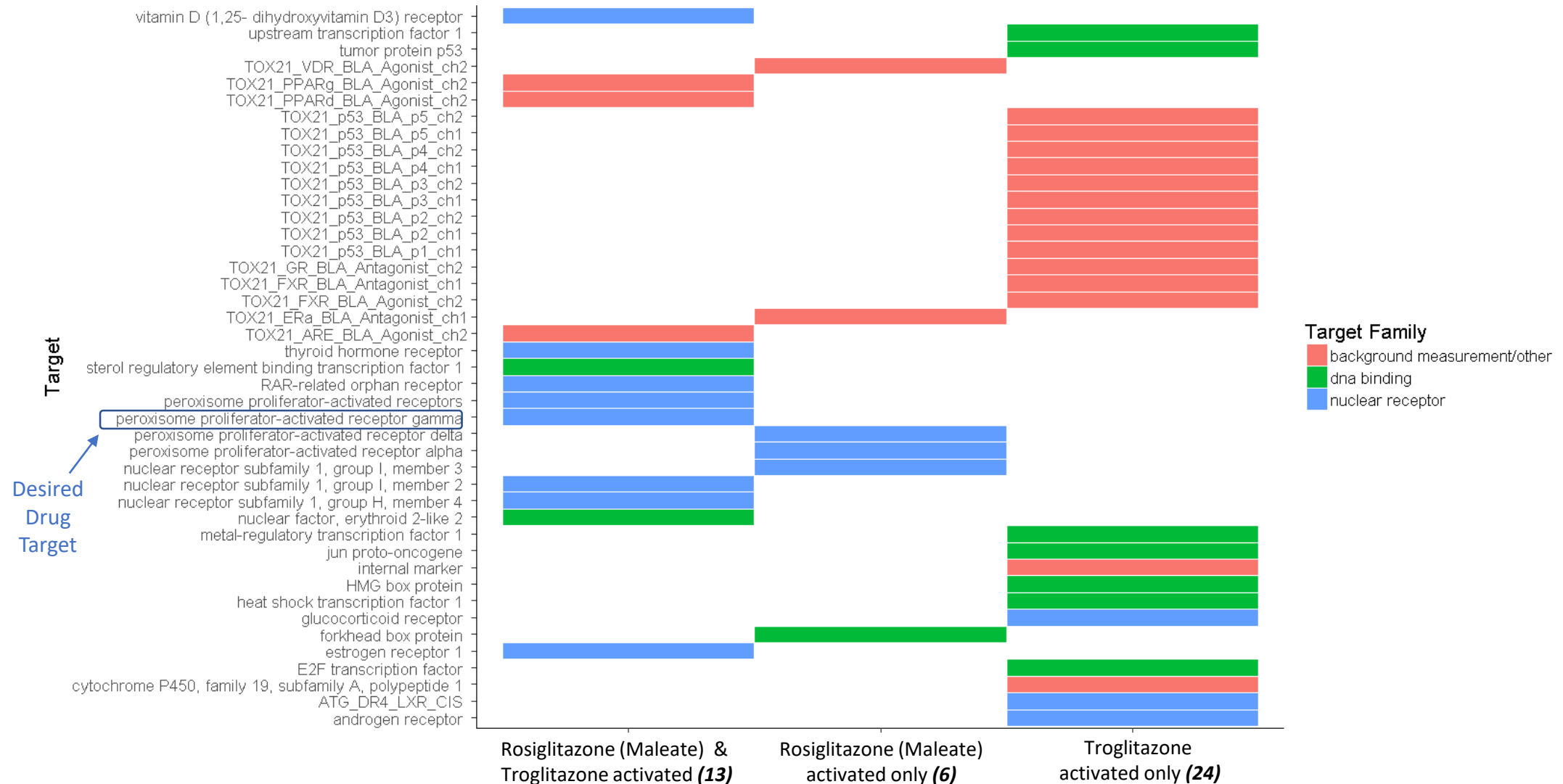
Venn diagrams not drawn to scale

Troglitazone yields higher # of positive tests across biological processes



Venn diagrams not drawn to scale

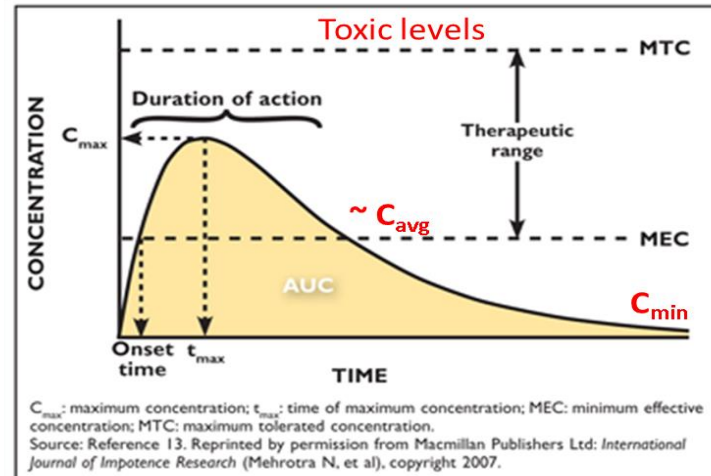
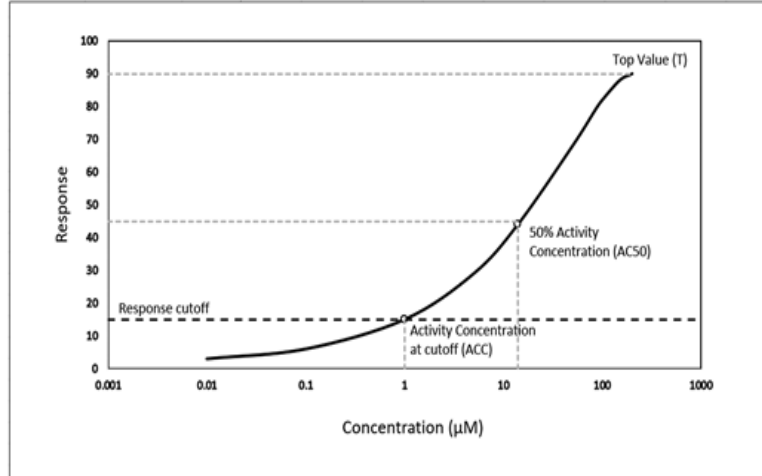
Troglitazone uniquely activates additional targets across target family of transcriptional regulators



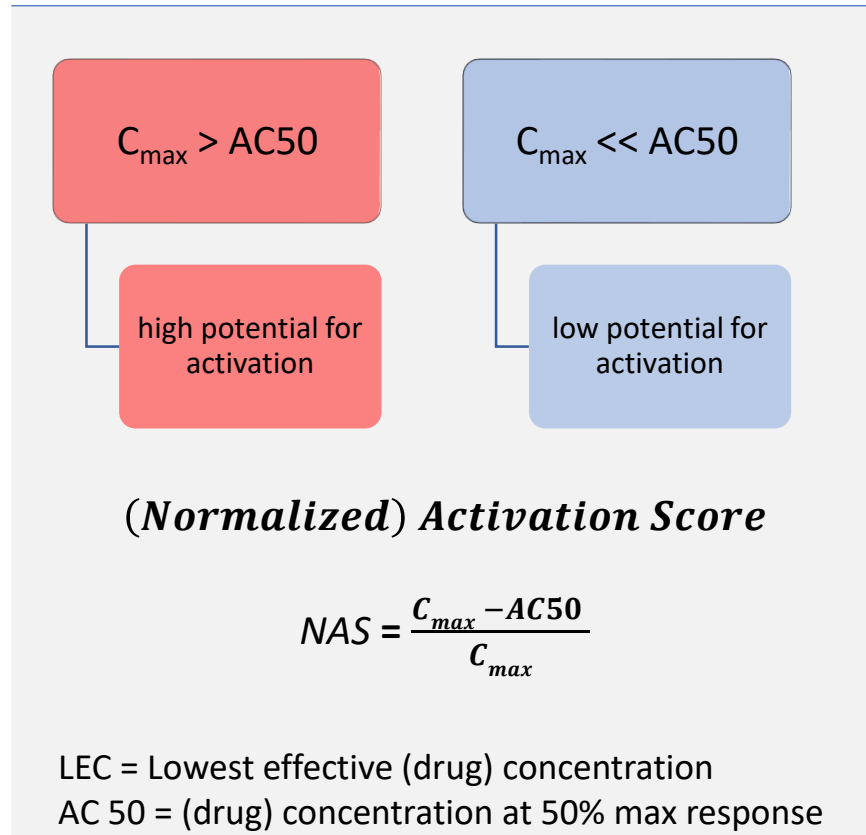
Stratification of targets' activation potential in humans using Normalized Activation Score (NAS)

C_{max} (of drug in humans) is $> AC50$ (for any in vitro/cellular assay) \rightarrow **higher potential for activation (of assay target)**

C_{max} (of drug in humans) $\ll AC50$ (for any in vitro/cellular assays) \rightarrow **lower potential for activation (of assay target)**



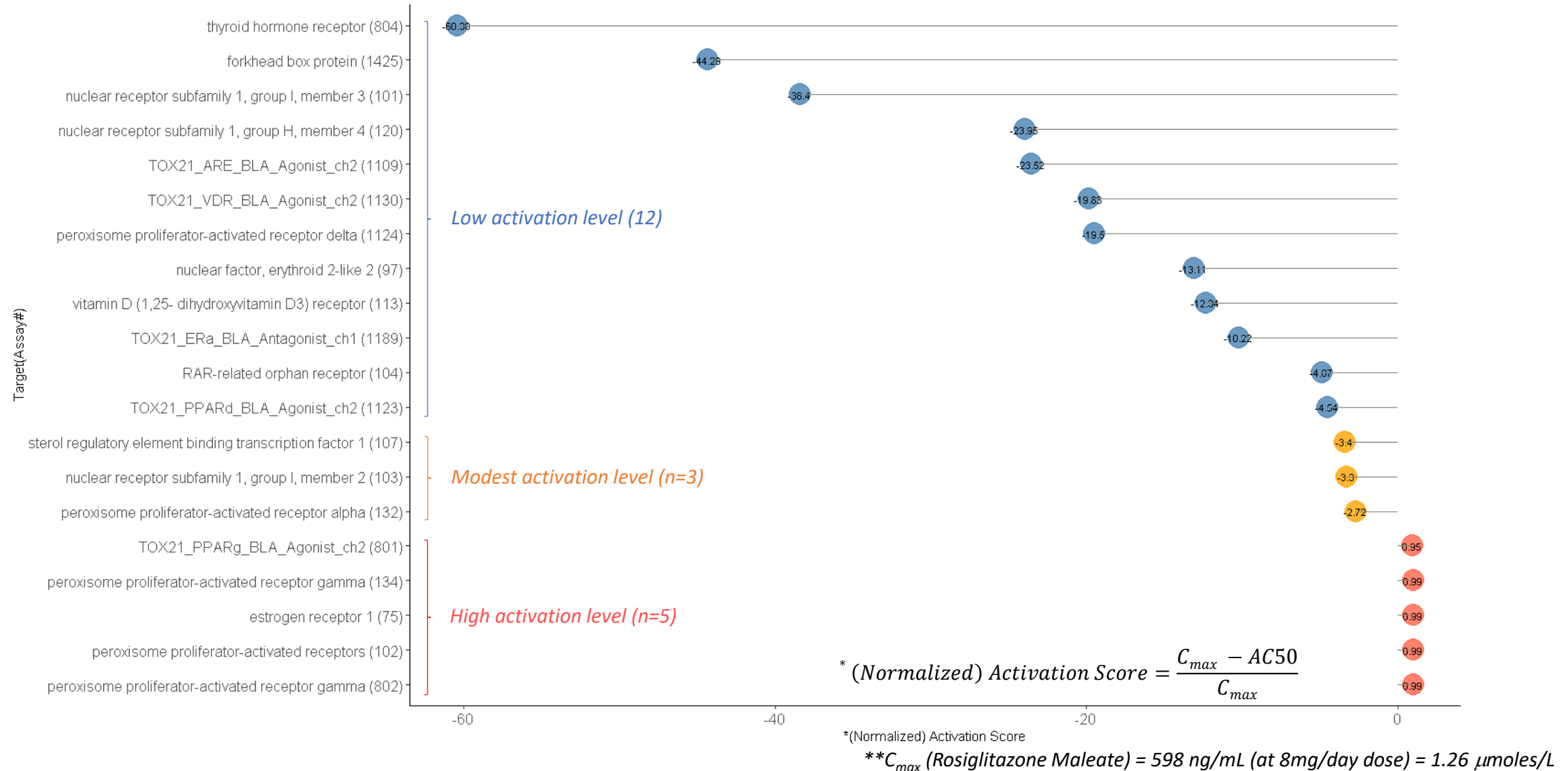
Schema for stratification of targets' activation potential in humans



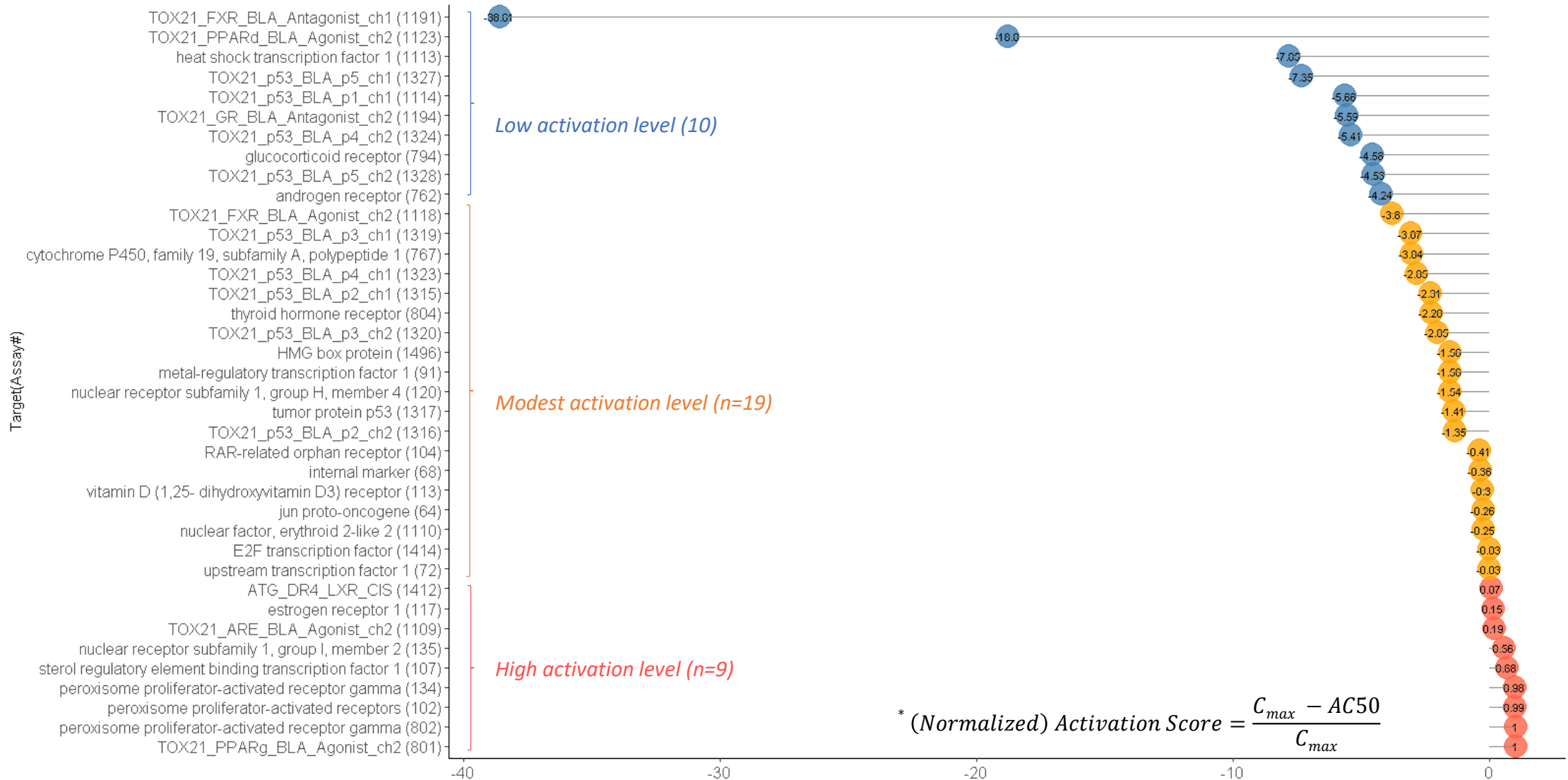
Normalized activation score (NAS) level	Indication
$NAS \geq 0$	$C_{max} > EC50$; highest activation potential
$0 > NAS > -4$	$C_{max} < EC50$; modest activation potential
$NAS < -4$	$C_{max} <<< EC50$; lowest activation potential

For the 437 “common tests” between Rosiglitazone Maleate and Troglitazone :
→ **NAS** values were generated and compared across corresponding biological processes

NAS stratified putative transcriptional regulatory targets of Rosiglitazone Maleate in humans



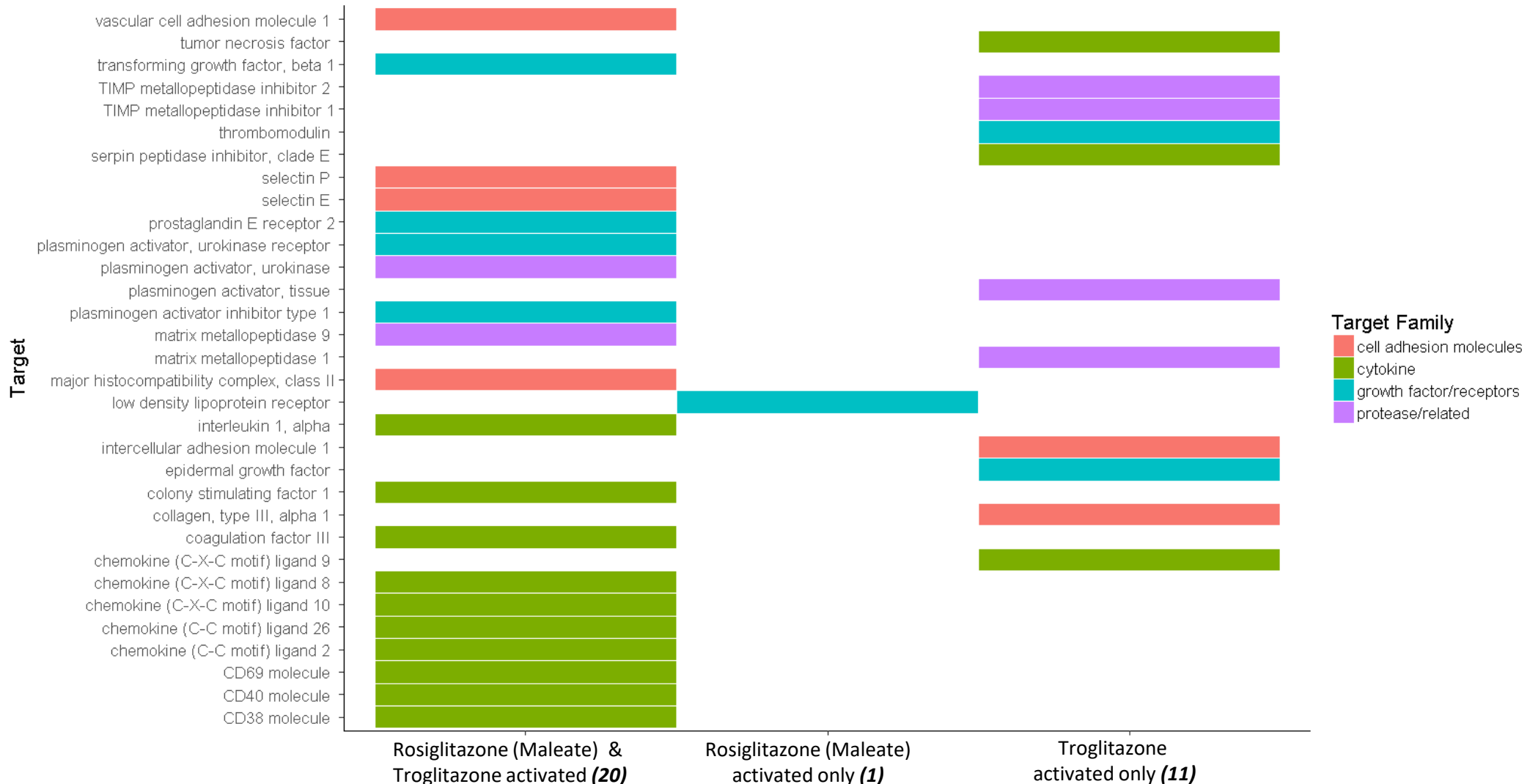
NAS stratified putative transcriptional regulatory targets of Troglitazone in humans



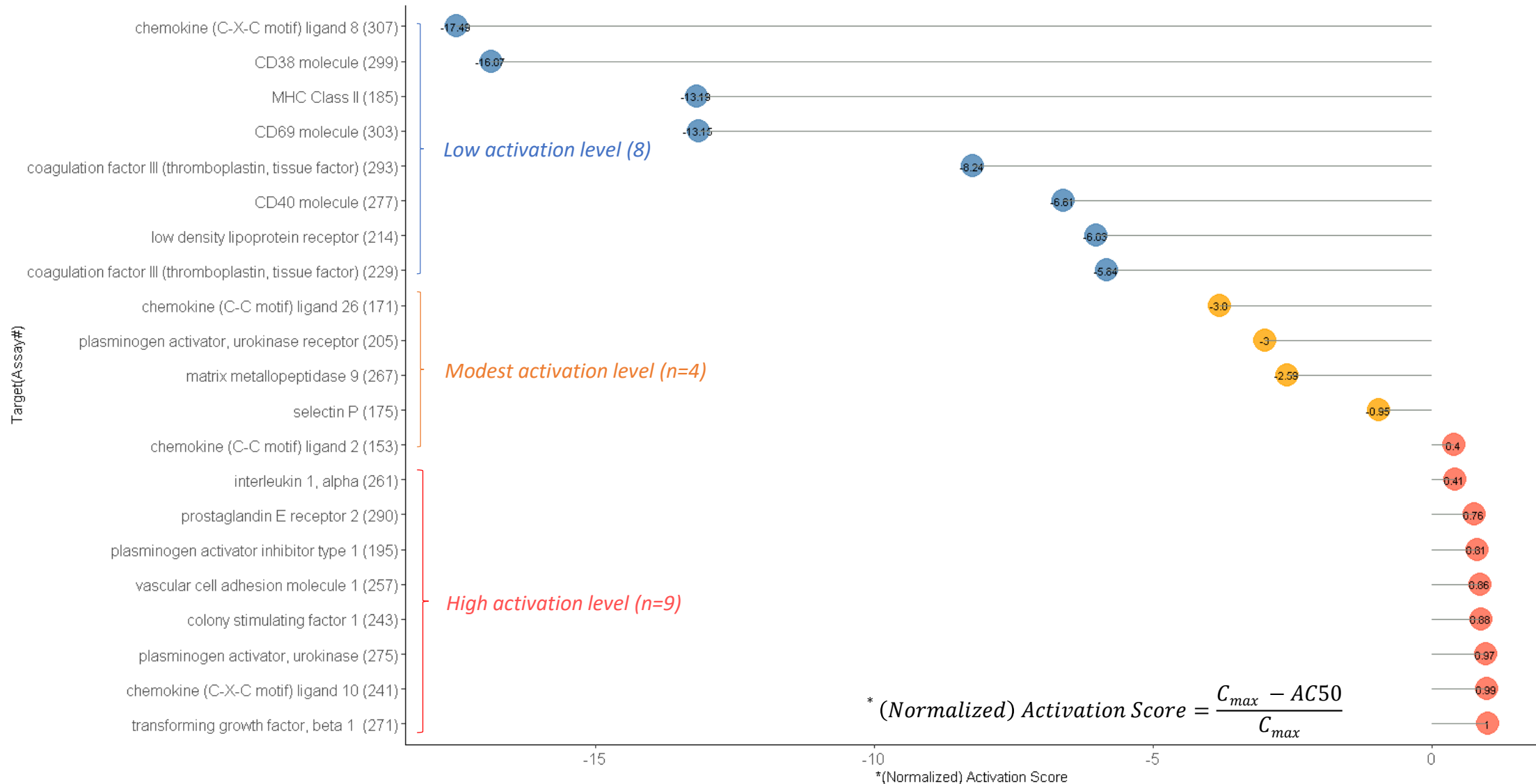
** C_{max} (Troglitazone) = 2.82 $\mu\text{g/mL}$ (at 600mg/day dose) = 6.38 $\mu\text{moles/L}$

*(Normalized) Activation Score

Troglitazone uniquely activates additional targets across target family of gene expression regulators

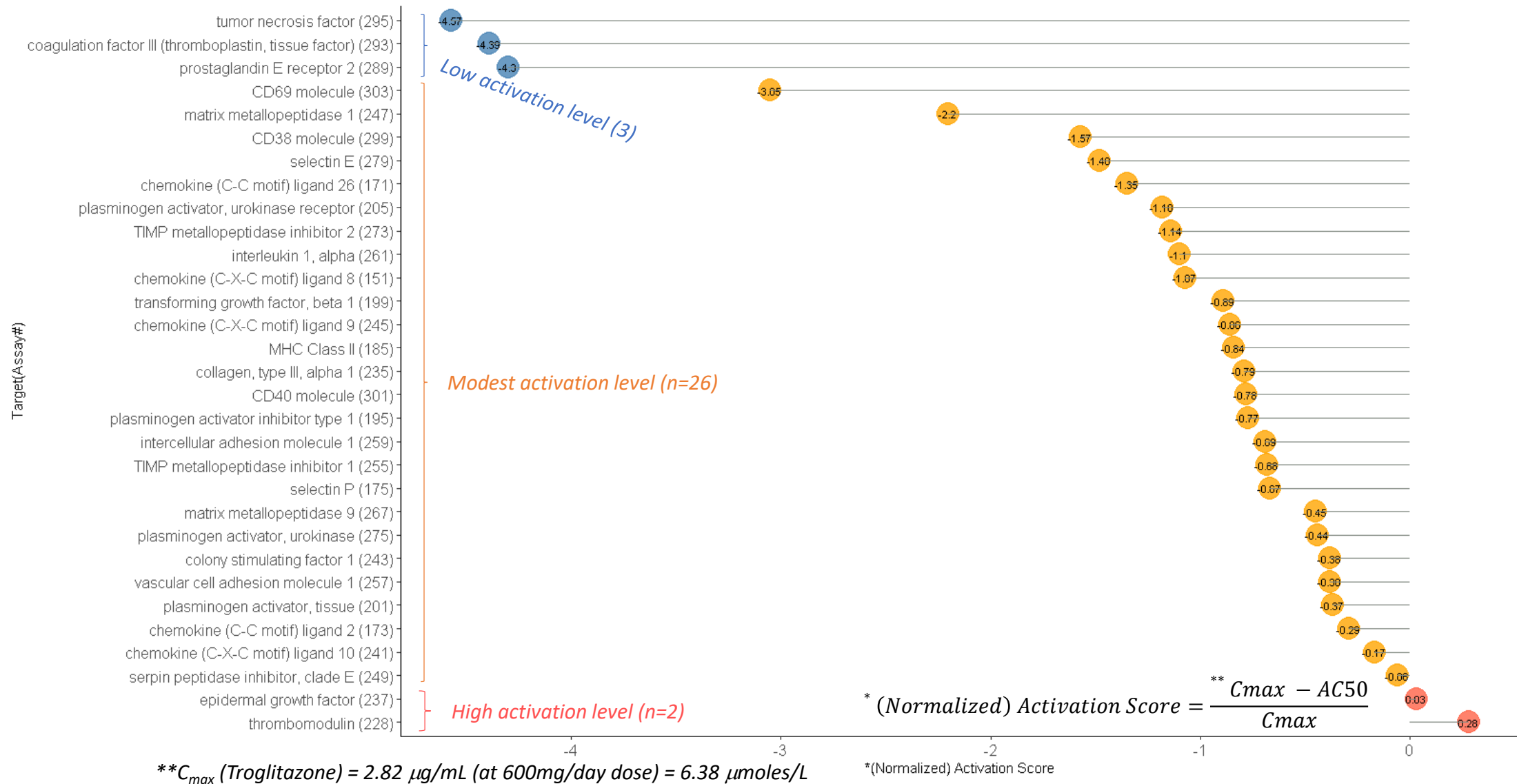


NAS stratified putative gene expression regulatory targets of Rosiglitazone Maleate in humans



** C_{max} (Rosiglitazone Maleate) = 598 ng/mL (at 8mg/day dose) = 1.26 μ moles/L

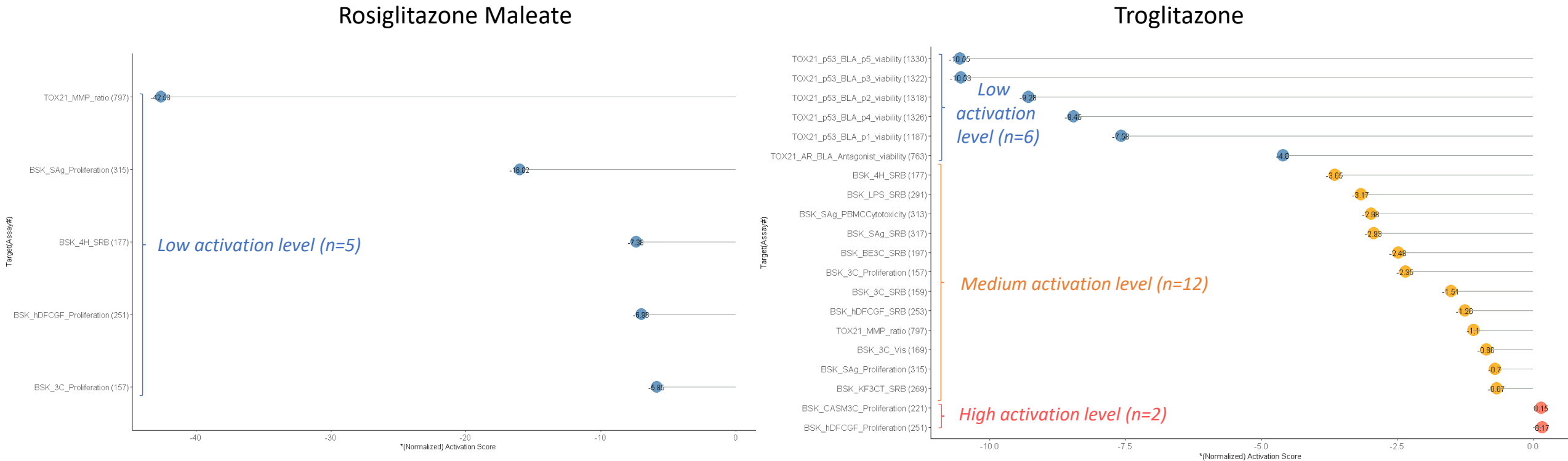
NAS stratified putative gene expression regulatory targets of Troglitazone in humans



Cell cycle/Cell morphology targets



NAS stratified putative cell cycle/morphology targets of Rosiglitazone Maleate and Troglitazone in humans



$$*(\text{Normalized}) \text{ Activation Score} = \frac{C_{\max} - AC50}{C_{\max}}$$

** C_{\max} (Rosiglitazone Maleate) = 598 ng/mL (at 8mg/day dose) = 1.26 μ moles/L

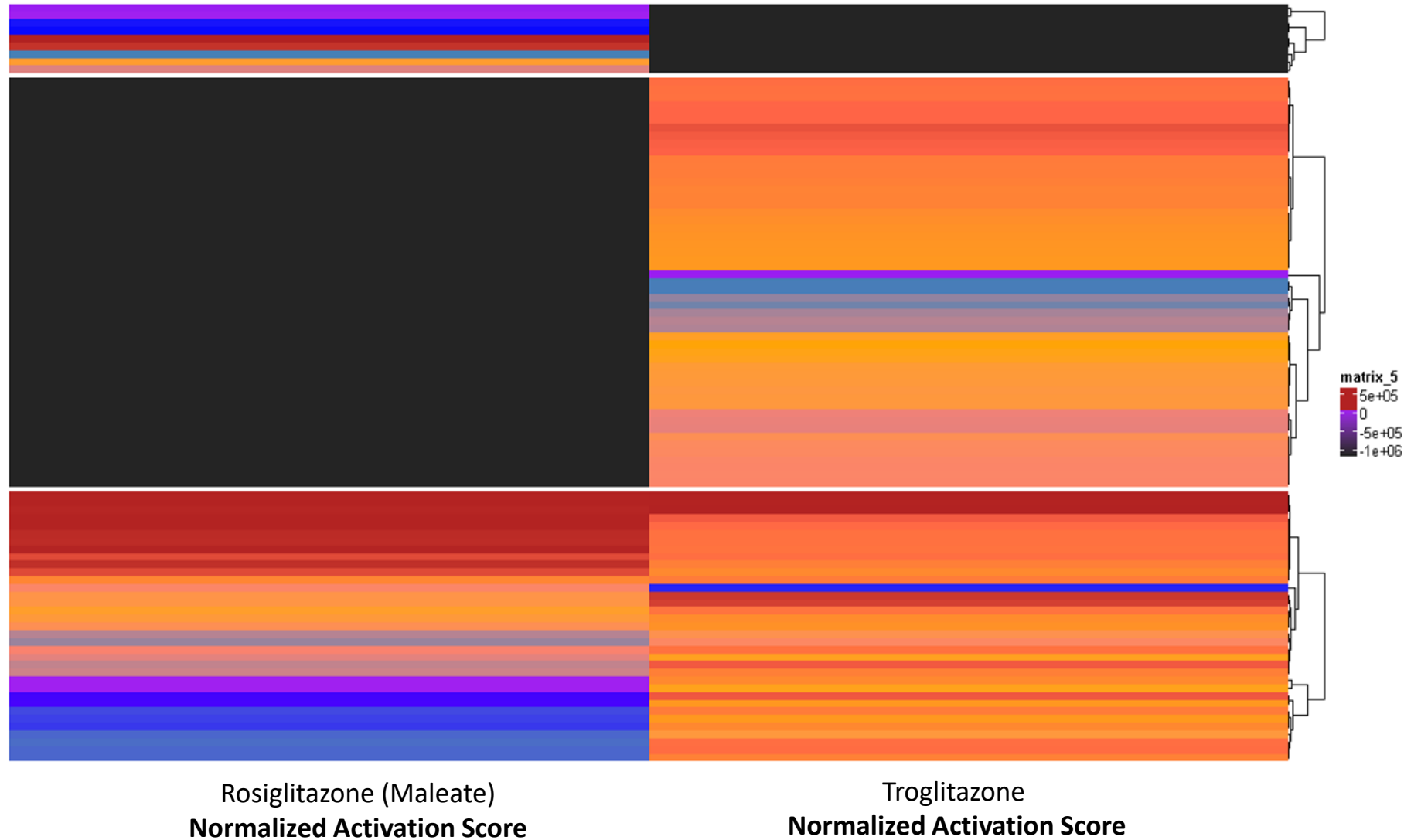
** C_{\max} (Troglitazone) = 2.82 μ g/mL (at 600mg/day dose) = 6.38 μ moles/L

NAS based "clustering" of all tests

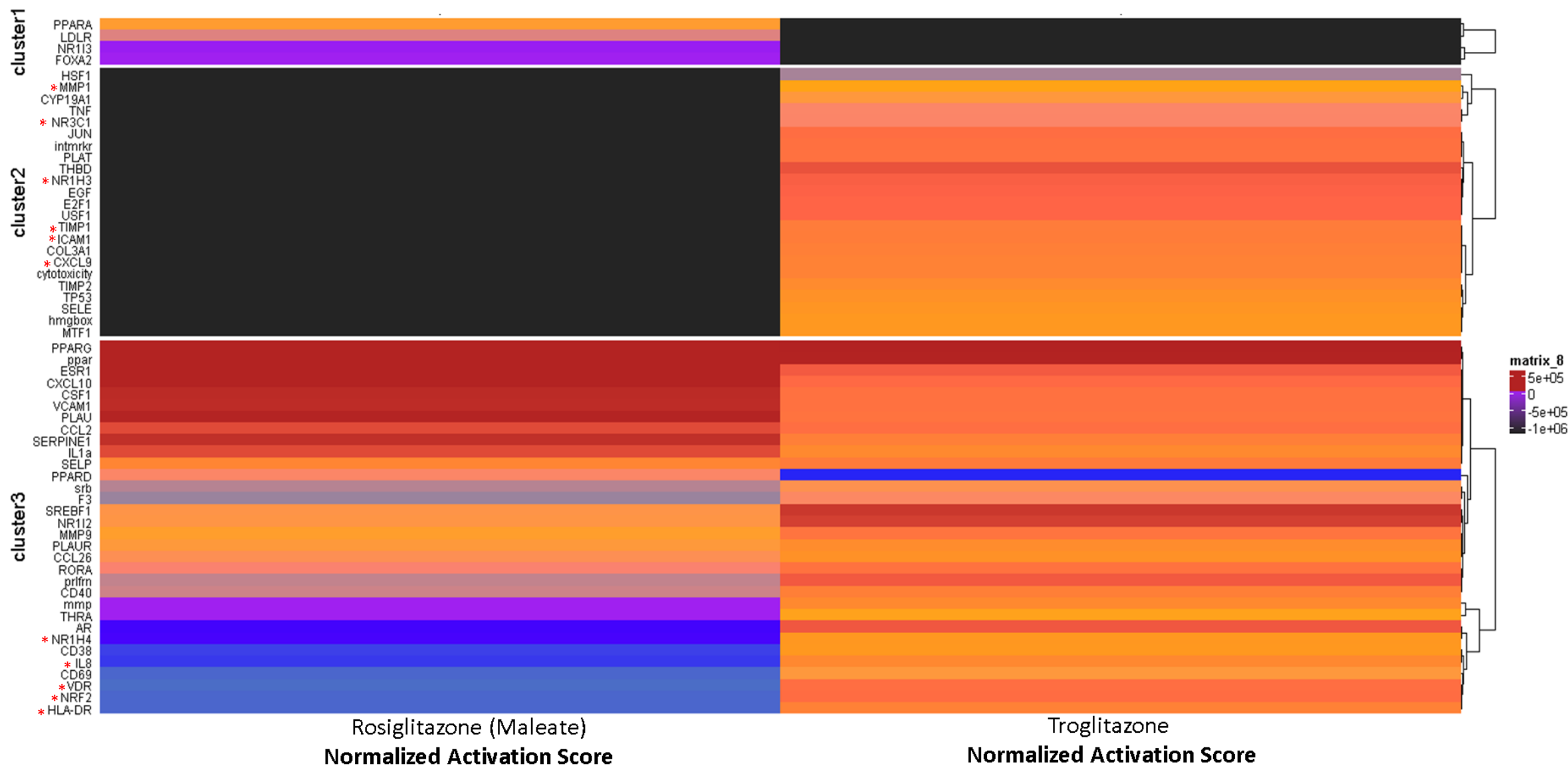
*Tests affected by
Rosiglitazone (Maleate) only*

*Tests affected
by Troglitazone only*

*Tests affected
by both drugs*



NAS based "clustering" of all Targets/Pathways



* Implicated in liver injury/repair pathways - Duarte S et al., 2015; Saiman Y & Friedman SL., 2013; Bohm F et al., 2010; Rudraiah S et al., 2016, Zimmerman HW et al., 2011, Gomex-Ospina N et al., 2016, Jadeja RN et al., 2016, Zuniga S et al., 2011, Tan W et al., 2017

Summary/Conclusions

Normalized activation score (NAS) stratifies molecular actions of drugs

- Targets/pathways with $NAS > 0$ expected to have highest potential for activation in patients and represent potential non-target effects.

Troglitazone associated targets are correlated with DILI risk

- Correlation with Troglitazone-associated targets (higher NAS for Troglitazone vs Rosiglitazone Maleate) with role in liver injury/repair pathways

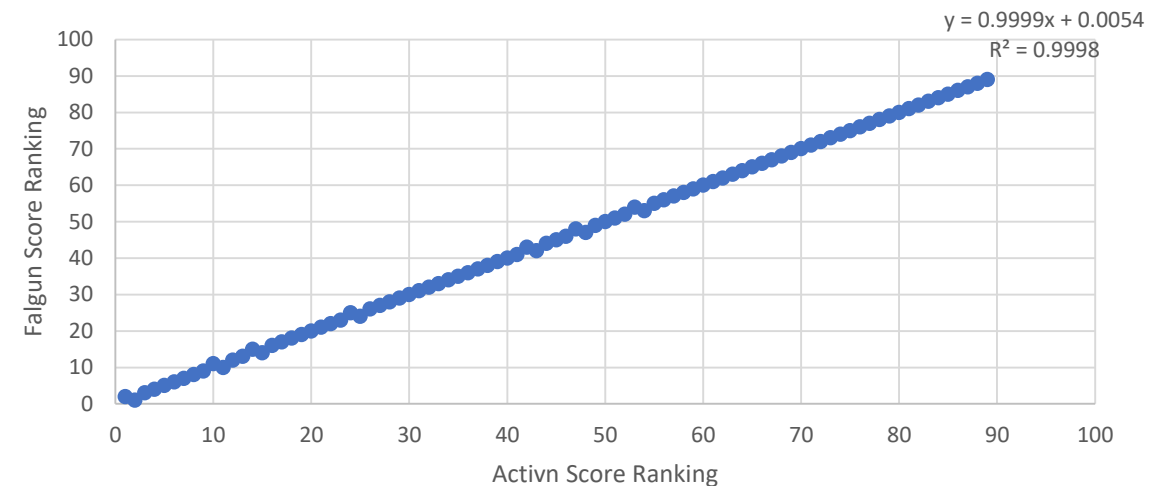
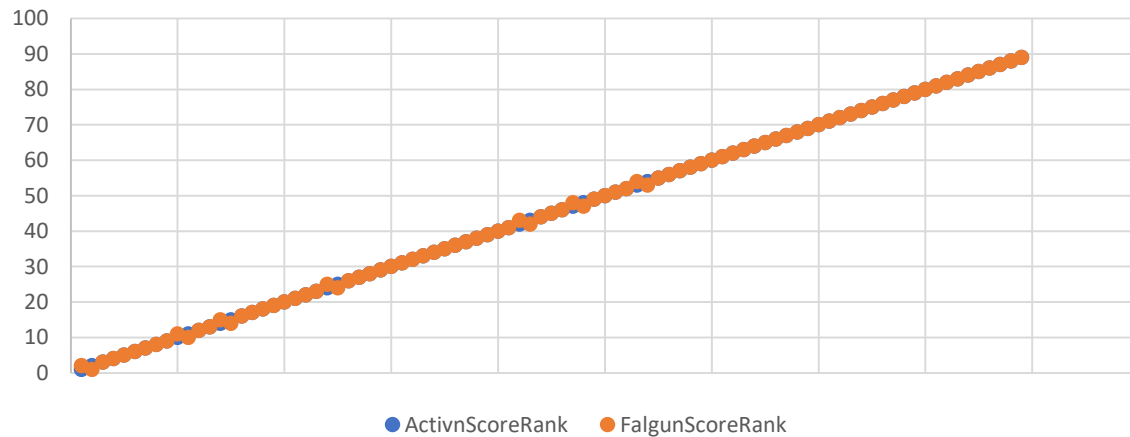
Next step: harness ToxCast for Liver Toxicity modeling

- *Integrate ToxCast targets/pathways data with known predictors of liver toxicity for potentially enhanced models of liver toxicity (initiated)*

Appendix

- “mostDILI” risk for Troglitazone-associated vs Rosiglitazone associated targets
- Comparison of target rankings for activation potential using NAS vs Falgun Shah score
- Count of positive tests for each biological process in Troglitazone vs Rosiglitazone (3 slides)
- Pharmacological activity/targets by DILI class from ToxCast database (3 slides)

Normalized Activation Score* performs nearly identically to Falgun Score** for stratification of Troglitazone affected targets/proteins (n = 89)

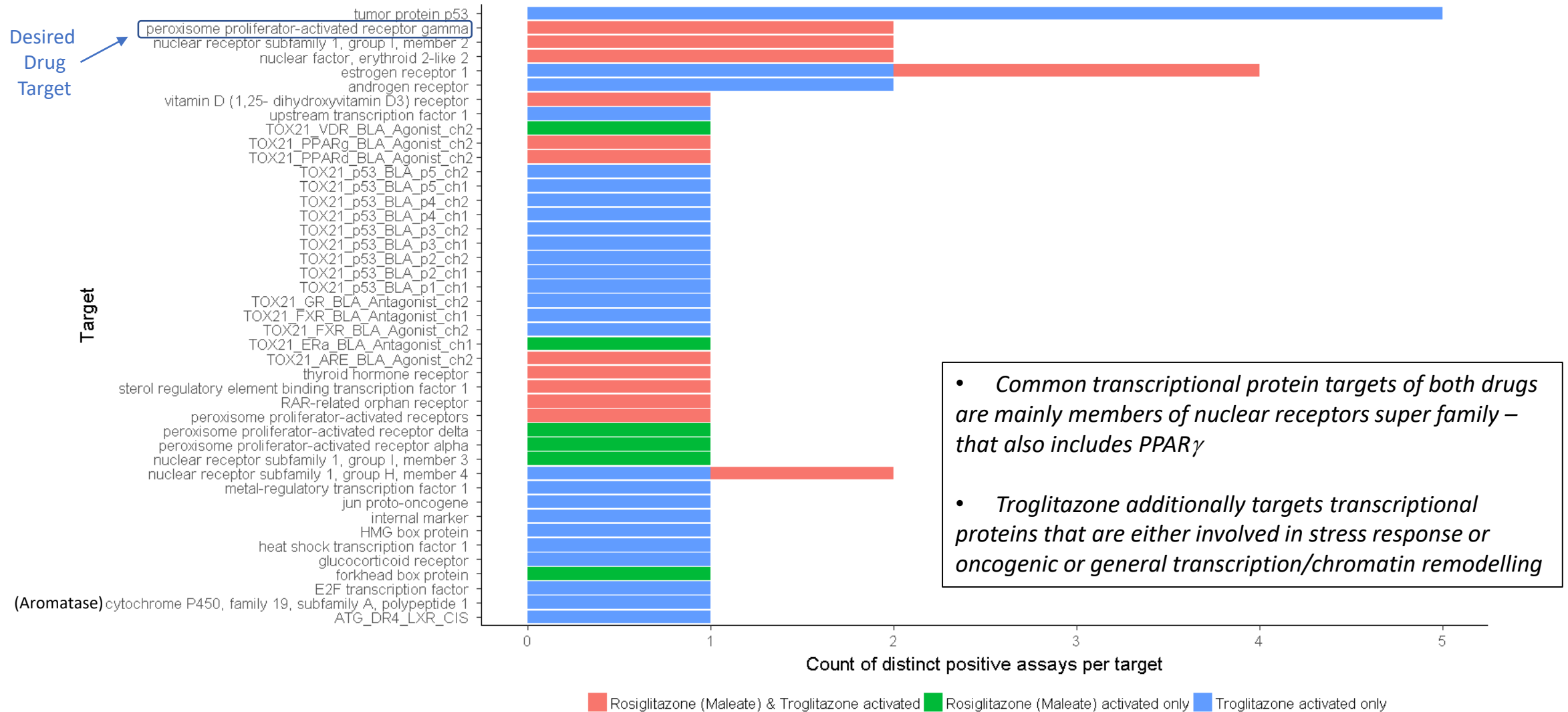


*Normalized Activation Score = $(C_{max} - EC50)/C_{max}$

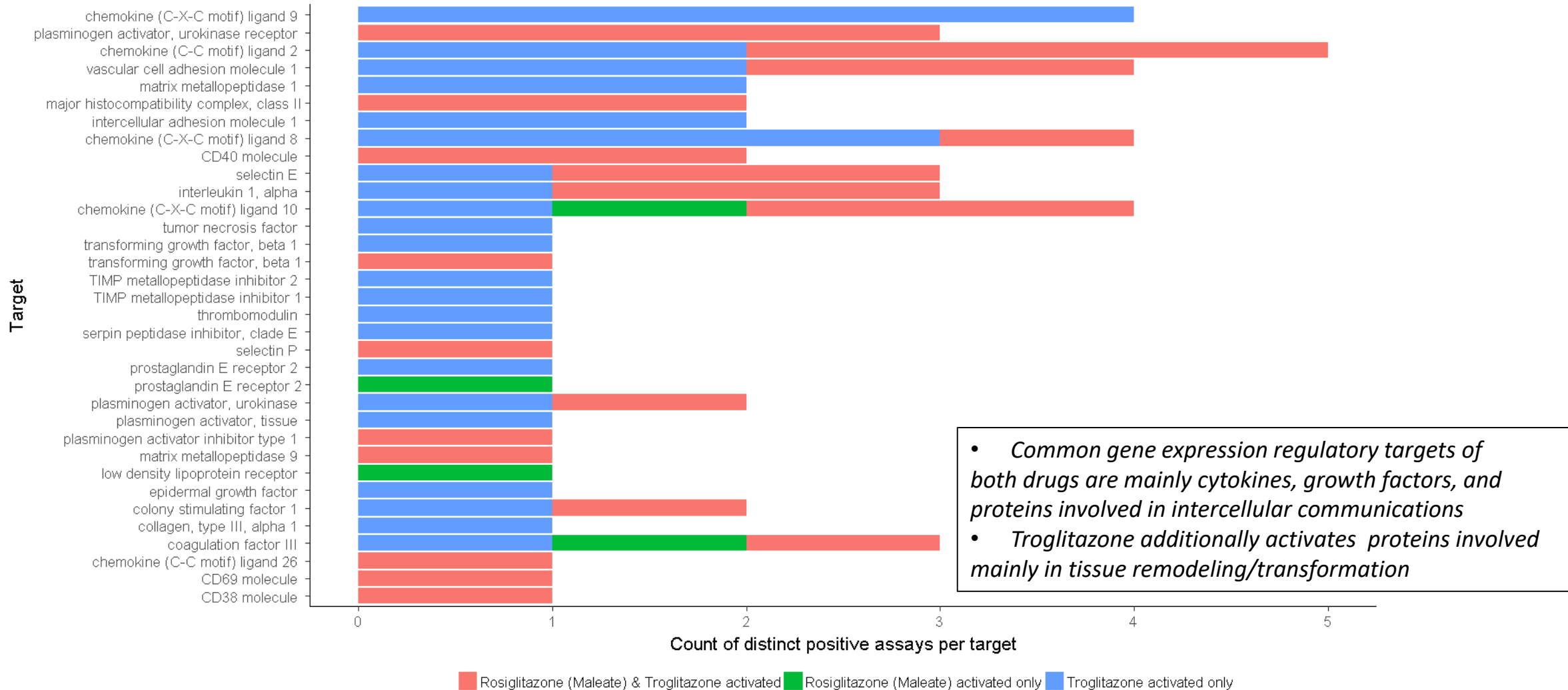
**Falgun Score = $C_{max}/EC50$ (Shah F et al., 2015 Toxicol Sci)

Activation level	Normalized Activation score	Falgun Score ($C_{max}/AC50$)
High activation	1 to ~0	~300 – ~1
Medium activation	0 to -4.0	1 - ~0.20
Low activation	Less than - 4.0	Less than 0.20

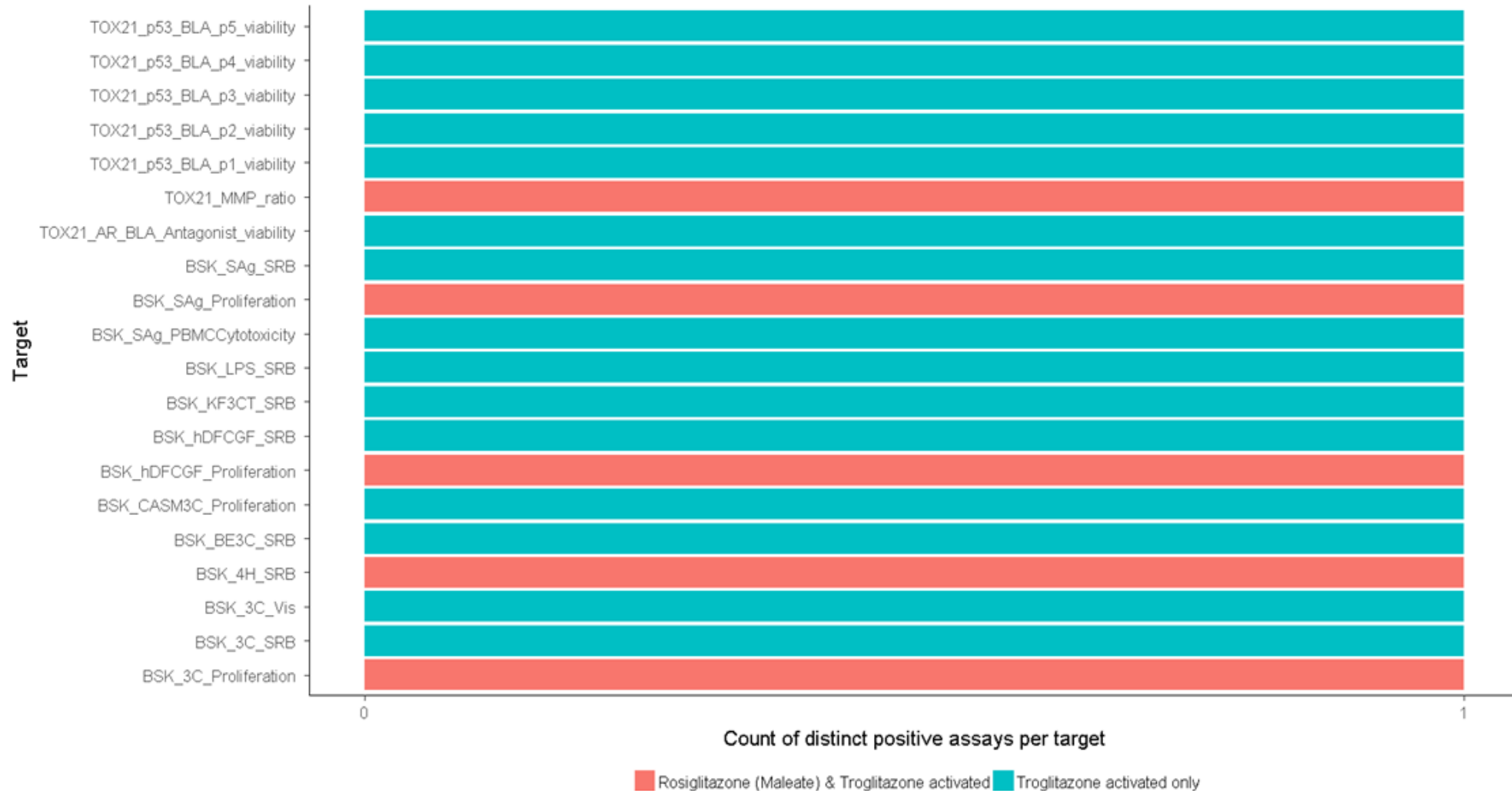
Count of positive tests per transcriptional targets for each drug



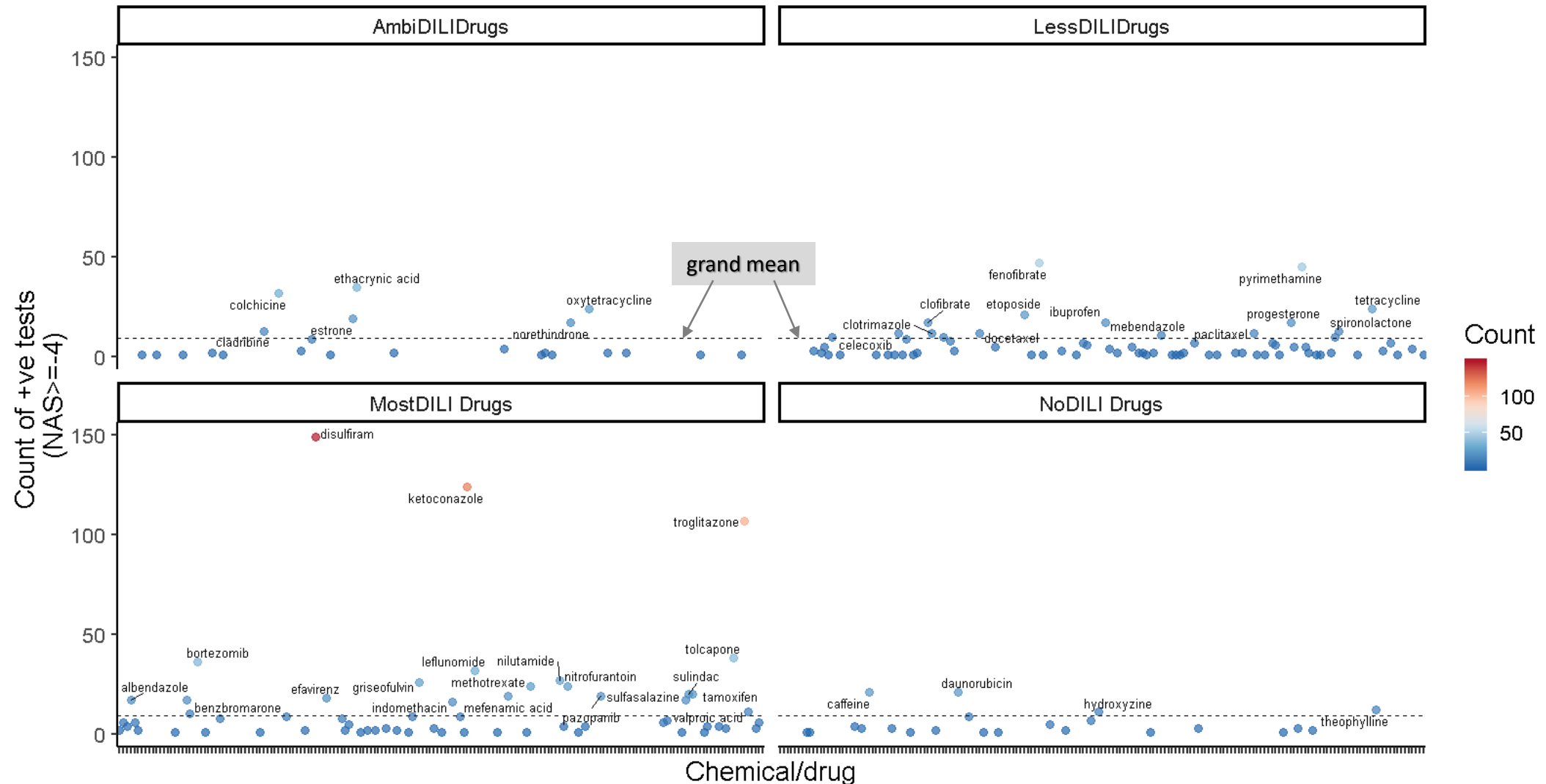
Count of positive assays for gene expression regulatory targets per drug



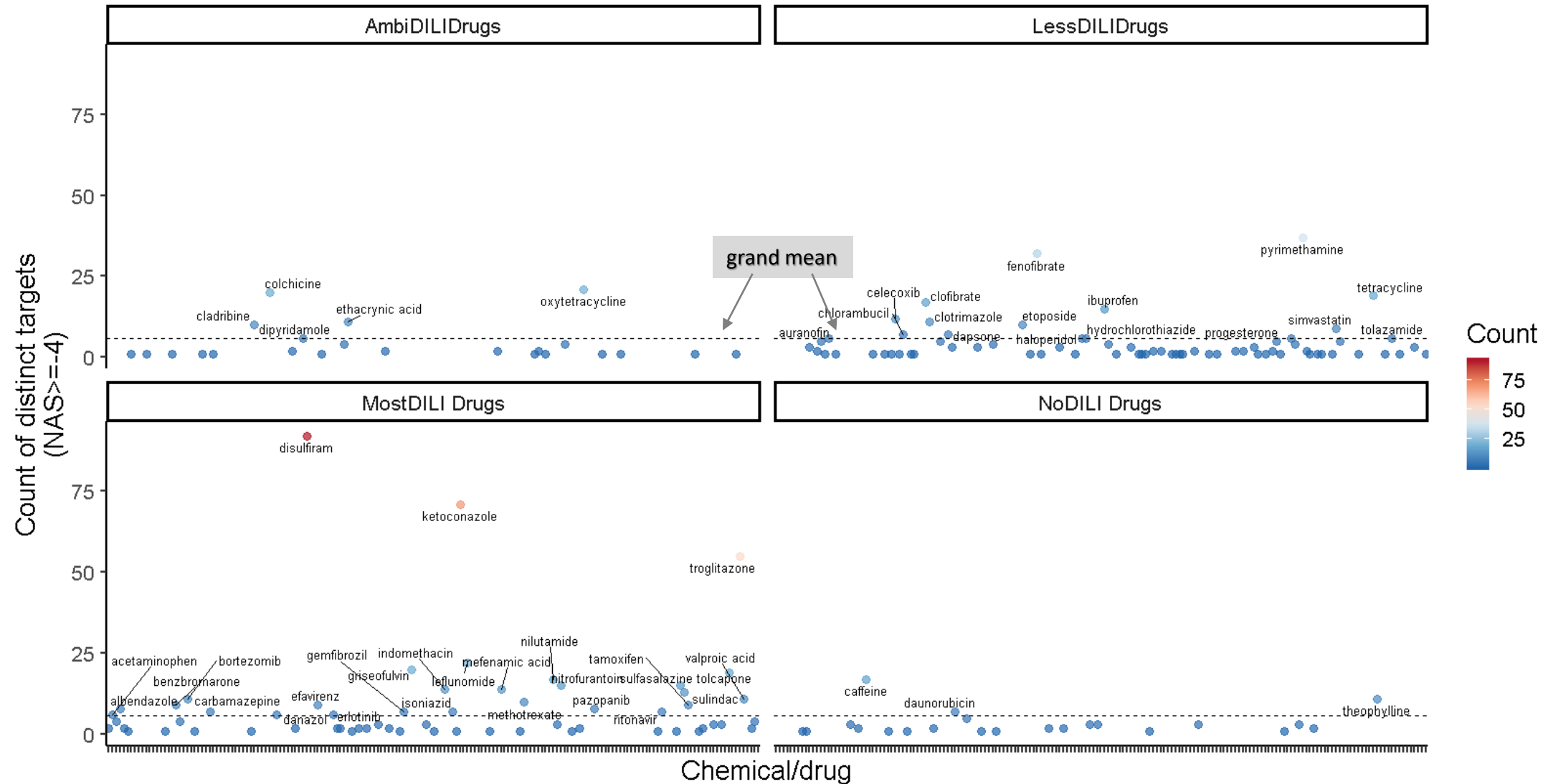
Count of positive assays for cell cycle/cell morphology targets per drug



Subset of drugs are more pharmacologically active in each DILI class



Subset of drugs are more pharmacologically active towards distinct* molecular targets



Highly reactive targets/pathways by DILI class

